Dossier zur Nutzenbewertung gemäß § 35a SGB V

Olaparib (Lynparza®)

AstraZeneca GmbH

Anhang 4-G

Erhaltungstherapie in Kombination mit Bevacizumab von erwachsenen Patientinnen mit einem fortgeschrittenen (FIGO-Stadien III und IV) high-grade epithelialen Ovarialkarzinom, Eileiterkarzinom oder primären Peritonealkarzinom, die nach einer abgeschlossenen Platin-basierten Erstlinien-Chemotherapie in Kombination mit Bevacizumab ein Ansprechen (vollständig oder partiell) haben und deren Tumor mit einem positiven Status der homologen Rekombinations-Defizienz (HRD) assoziiert ist

Stand: 28.10.2022

Table 2.1 PAOLA1: Summary of observation period (months) for PRO endpoints Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=132)
EORTC-QLQ-C30	n	255	132
	Median	24.18	24.05
	Min	0.0	0.0
	Max	52.5	41.2
EORTC-QLQ-OV28	n	255	132
	Median	24.18	24.05
	Min	0.0	0.0
	Max	52.5	41.2
EO-5D-5L	n	255	132
10 3D 3E	Median	24.18	24.05
	Min		
		0.0	0.0
	Max	52.5	41.2

Table 2.2.1 PAOLA1: Summary of analysis of time to worsening in EORTC QLQ-C30 global QoL/health status, functional, symptom and single item scales

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=132)	
	Number (%) Median time of patients (95% CI) n with events (months) [a]	Number (%) Median time of patients (95% CI) n with events (months) [a]	Hazard ratio 2-sided [b] 95% CI [b] p-value [c]
EORTC QLQ-C30 Global QoL/health status	255 146 (57.3) 16.6 (11.5,21.8)	132 81 (61.4) 13.8 (9.3,17.2)	0.85 0.65, 1.12 0.2343
EORTC QLQ-C30 Functional scale: Physical	255 125 (49.0) 20.0 (13.9,52.5)	132 74 (56.1) 16.4 (11.5,22.4)	0.85 0.64, 1.14 0.2790
EORTC QLQ-C30 Functional scale: Role	255 167 (65.5) 8.4 (5.8,11.2)	132 82 (62.1) 9.3 (6.1,16.2)	1.11 0.85, 1.46 0.4501
EORTC QLQ-C30 Functional scale: Cognitive	255 174 (68.2) 11.1 (8.5,14.0)	132 85 (64.4) 8.5 (5.9,13.6)	0.91 0.70, 1.19 0.4835
EORTC QLQ-C30 Functional scale: Emotional	255 158 (62.0) 13.8 (9.0,19.3)	132 85 (64.4) 11.1 (8.3,13.8)	0.93 0.71, 1.22 0.5708
EORTC QLQ-C30 Functional scale: Social	255 148 (58.0) 13.5 (8.6,19.6)	132 81 (61.4) 11.3 (8.5,16.4)	0.91 0.69, 1.20 0.4710
EORTC QLQ-C30 Single item symptom scale: Loss of appetite	255 146 (57.3) 13.6 (11.1,22.1)	132 65 (49.2) 22.3 (16.6,28.7)	1.42 1.06, 1.92 0.0227*
EORTC QLQ-C30 Single item symptom scale: Constipation	255 133 (52.2) 19.9 (16.6,23.4)	132 69 (52.3) 19.7 (14.0,22.3)	1.03 0.77, 1.39 0.8313
EORTC QLQ-C30 Single item symptom scale: Diarrhoea	255 124 (48.6) 24.0 (16.6,25.9)	132 58 (43.9) 23.5 (19.9,35.0)	1.15 0.84, 1.58 0.4093

Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation.

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Table 2.2.1 PAOLA1: Summary of analysis of time to worsening in EORTC QLQ-C30 global QoL/health status, functional, symptom and single item scales

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=132)		
	Number (%) Median time of patients (95% CI) n with events (months) [a]	Number (%) Median time of patients (95% CI) n with events (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [c]
EORTC QLQ-C30 Single item symptom scale: Dyspnoea	255 125 (49.0) 20.7 (16.0,52.5)	132 67 (50.8) 18.7 (12.3,24.9)	0.92 0.68, 1.25	0.5796
EORTC QLQ-C30 Symptom scale: Fatigue	255 199 (78.0) 5.6 (3.1, 6.0)	132 98 (74.2) 5.7 (5.5,11.1)	1.10 0.86, 1.41	0.4815
EORTC QLQ-C30 Single item symptom scale: Financial difficulties	255 77 (30.2) 38.4 (38.4, NE)	132 48 (36.4) NE (NE, NE)	0.72 0.50, 1.04	0.0709
EORTC QLQ-C30 Symptom scale: Nausea and vomiting	255 178 (69.8) 5.8 (5.6, 8.7)	132 70 (53.0) 19.2 (12.7,23.5)	1.81 1.37, 2.42	<0.0001*
EORTC QLQ-C30 Symptom scale: Pain	255 183 (71.8) 5.8 (5.6, 8.3)	132 95 (72.0) 5.6 (3.0, 8.1)	0.92 0.72, 1.19	0.5505
EORTC QLQ-C30 Single item symptom scale: Insomnia	255 159 (62.4) 11.3 (8.4,14.0)	132 91 (68.9) 8.3 (5.6,11.1)	0.73 0.56, 0.95	0.0185*

Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation.

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

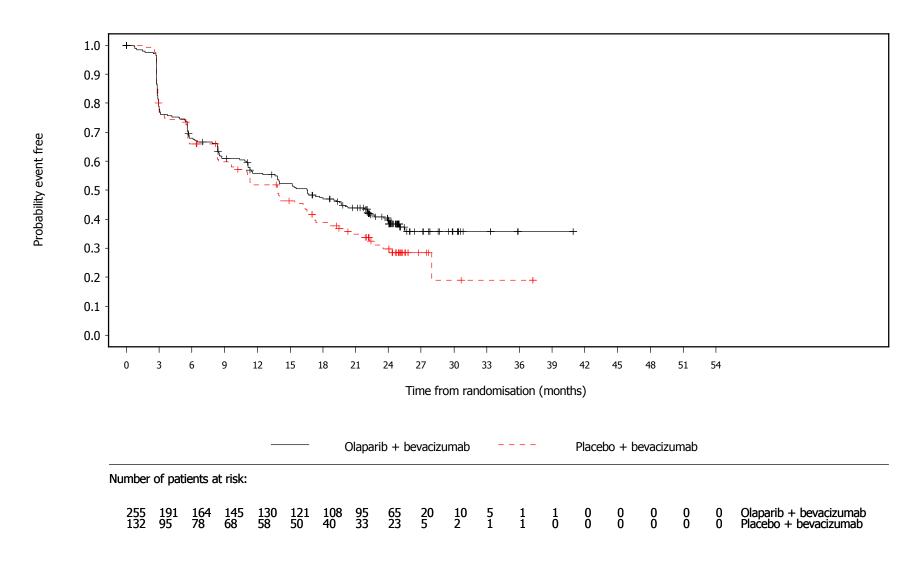
[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Figure 2.2.2.1 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Global QoL/health status time to clinically meaningful worsening (first occurrence)

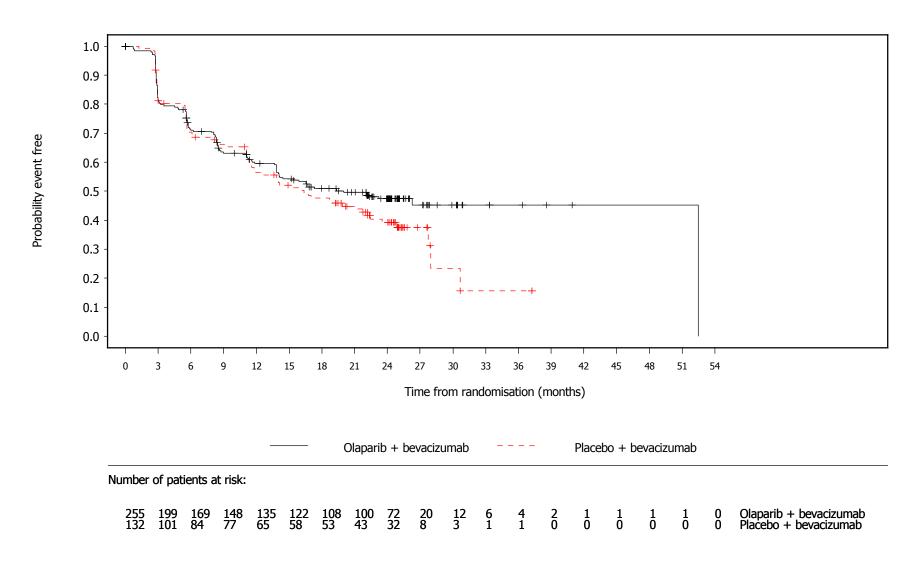
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr.sas ettemainprfaa 25NOV2020:15:58 kvbv306

Figure 2.2.2.2 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Physical time to clinically meaningful worsening (first occurrence)

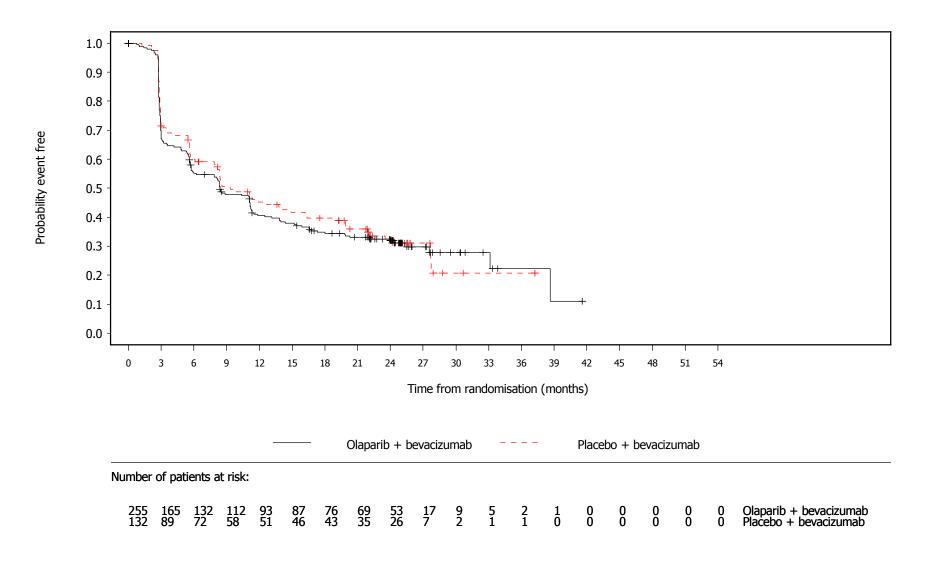
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.3 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Role time to clinically meaningful worsening (first occurrence)

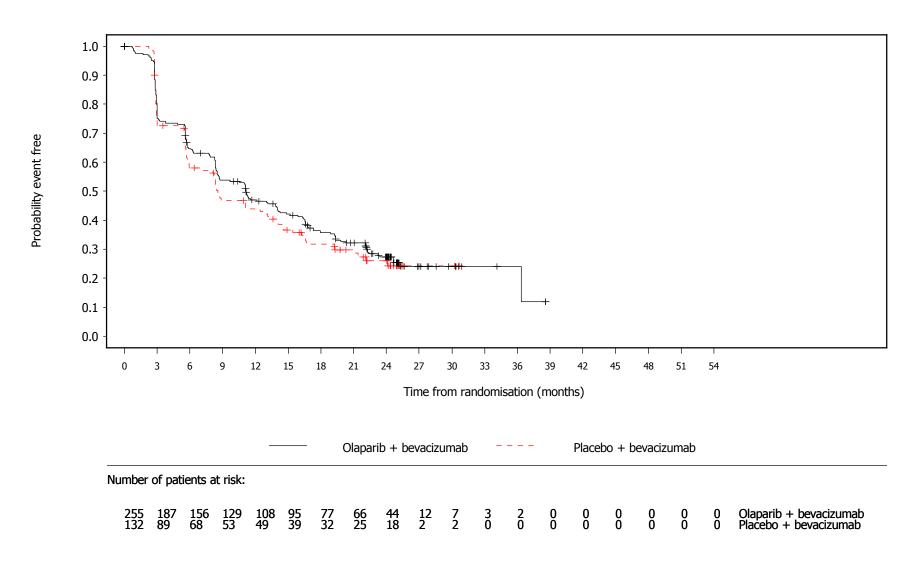
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.4 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Cognitive time to clinically meaningful worsening (first occurrence)

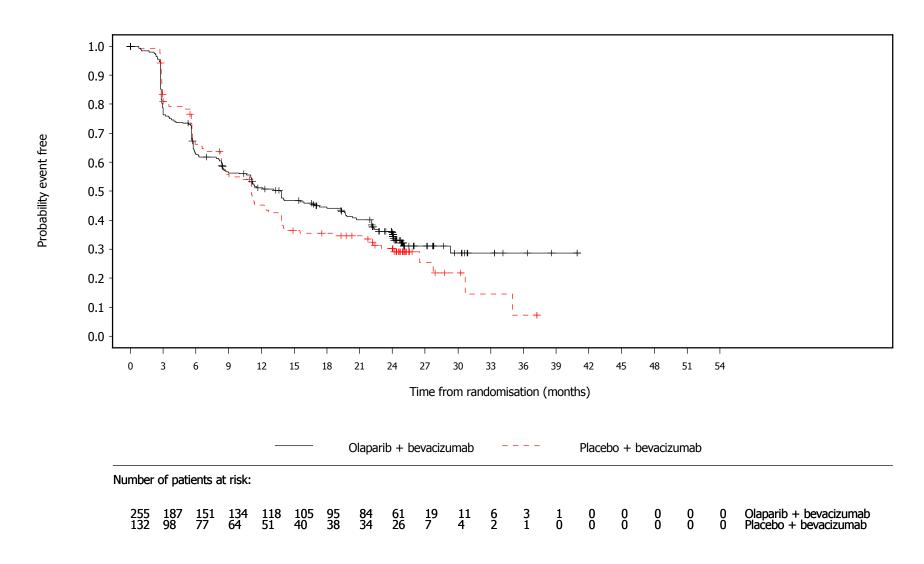
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.5 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Emotional time to clinically meaningful worsening (first occurrence)

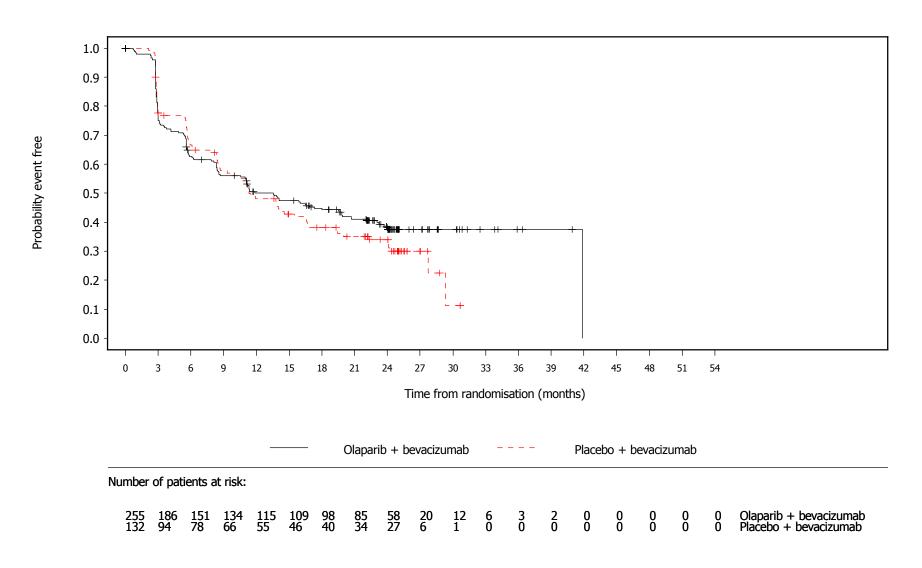
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.6 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Social time to clinically meaningful worsening (first occurrence)

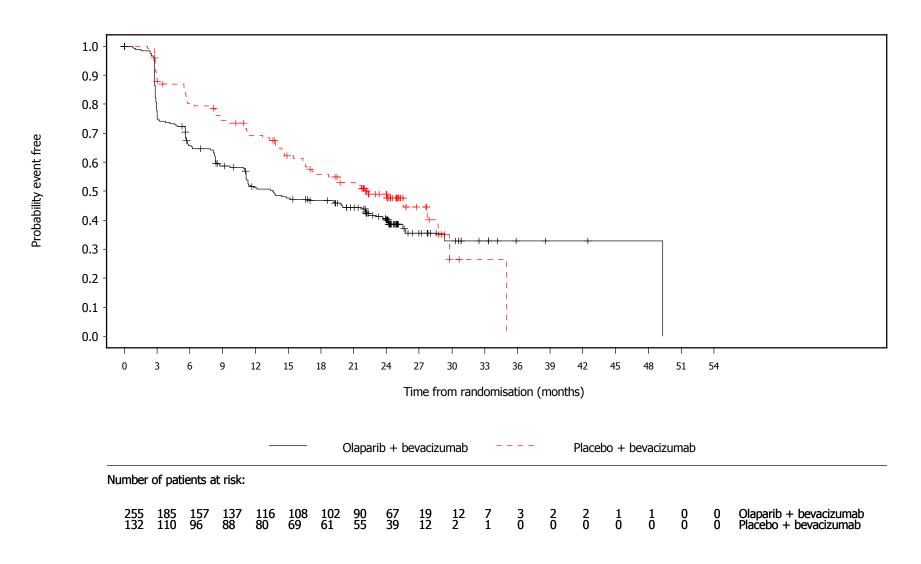
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.7 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Loss of appetite time to clinically meaningful worsening (first occurrence)

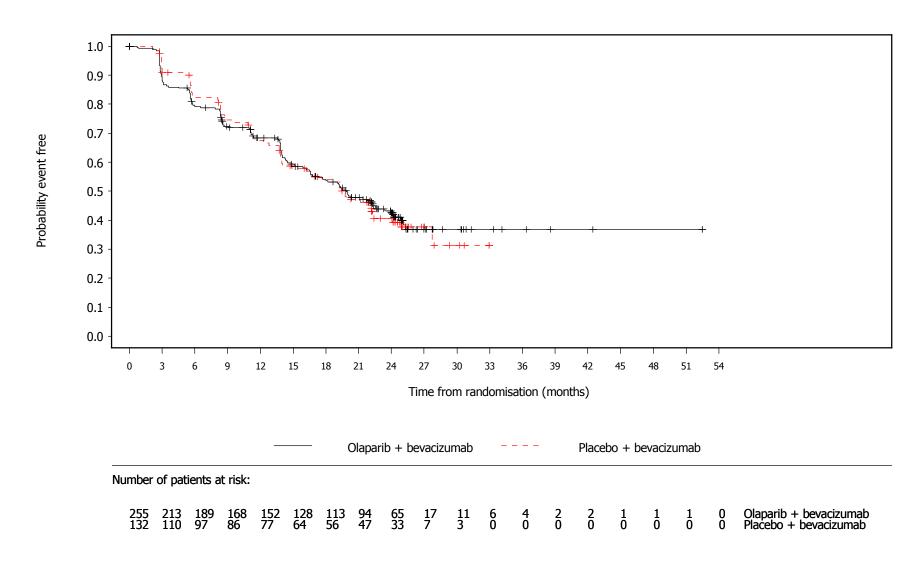
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.8 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Constipation time to clinically meaningful worsening (first occurrence)

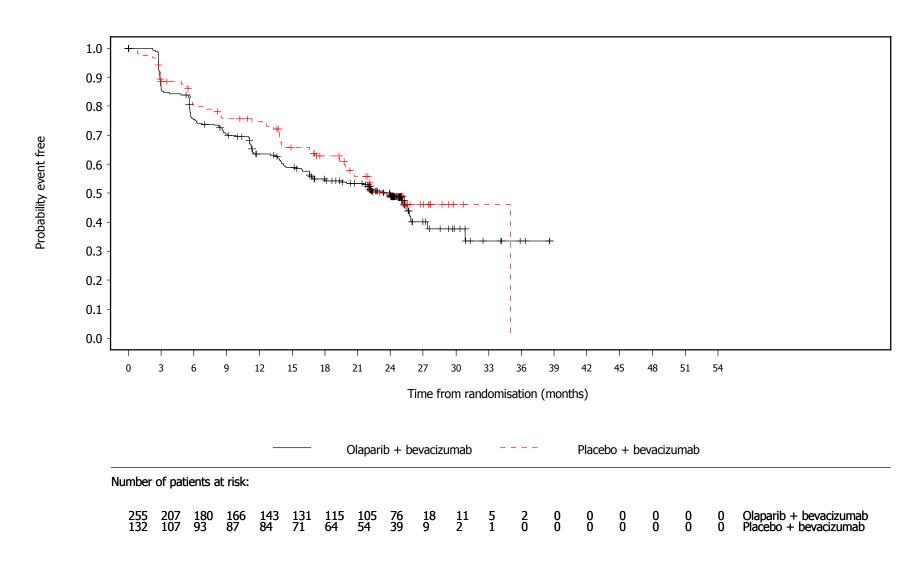
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.9 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Diarrhoea time to clinically meaningful worsening (first occurrence)

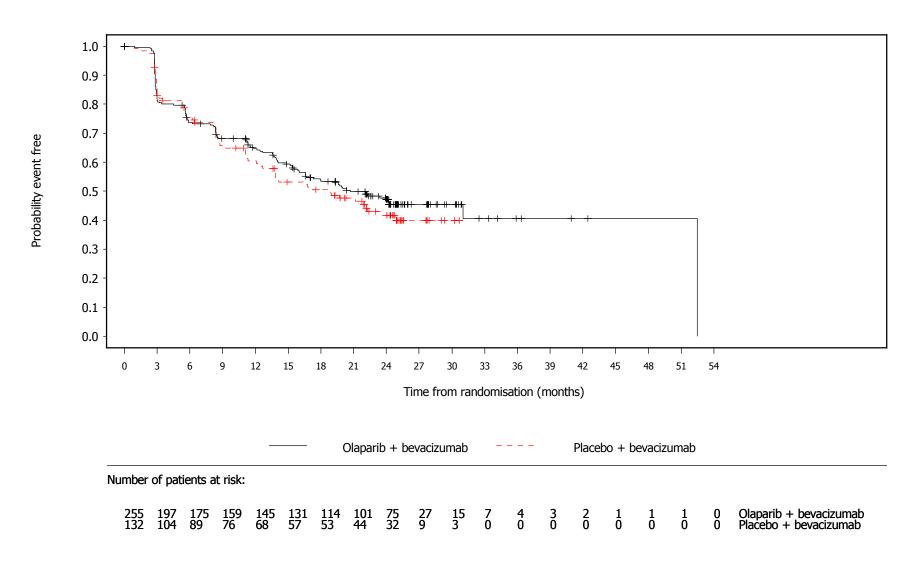
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.10 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Dyspnoea time to clinically meaningful worsening (first occurrence)

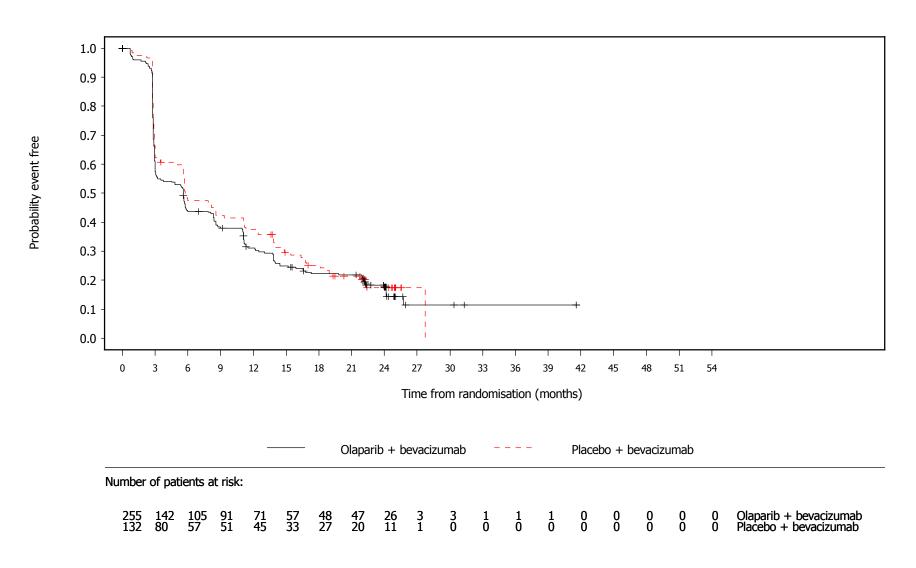
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.11 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Fatigue time to clinically meaningful worsening (first occurrence)

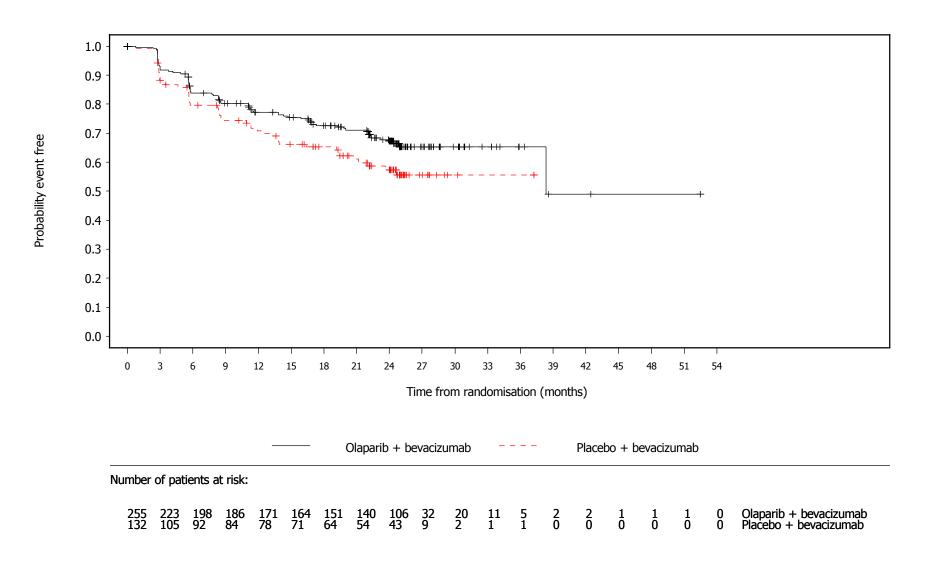
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.12 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Financial difficulties time to clinically meaningful worsening (first occurrence)

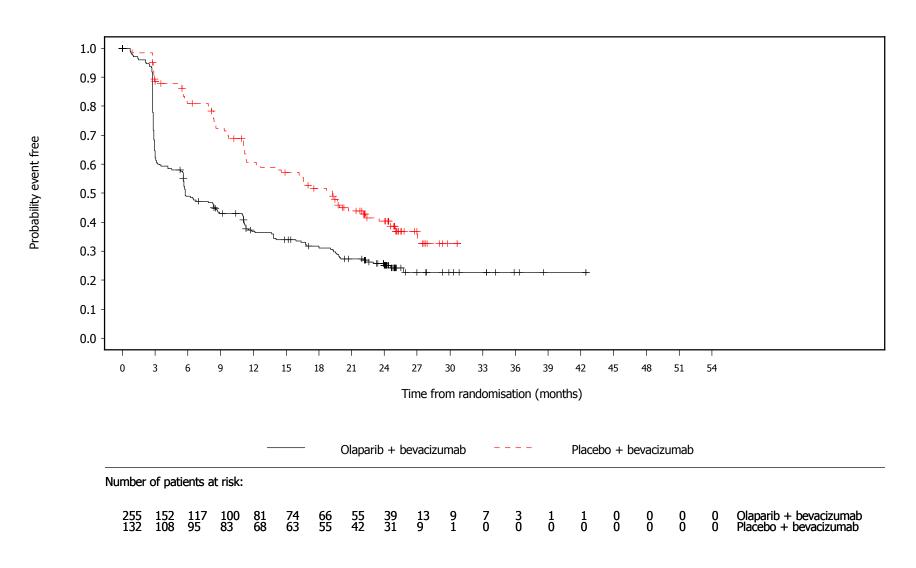
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.13 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Nausea and vomiting time to clinically meaningful worsening (first occurrence)

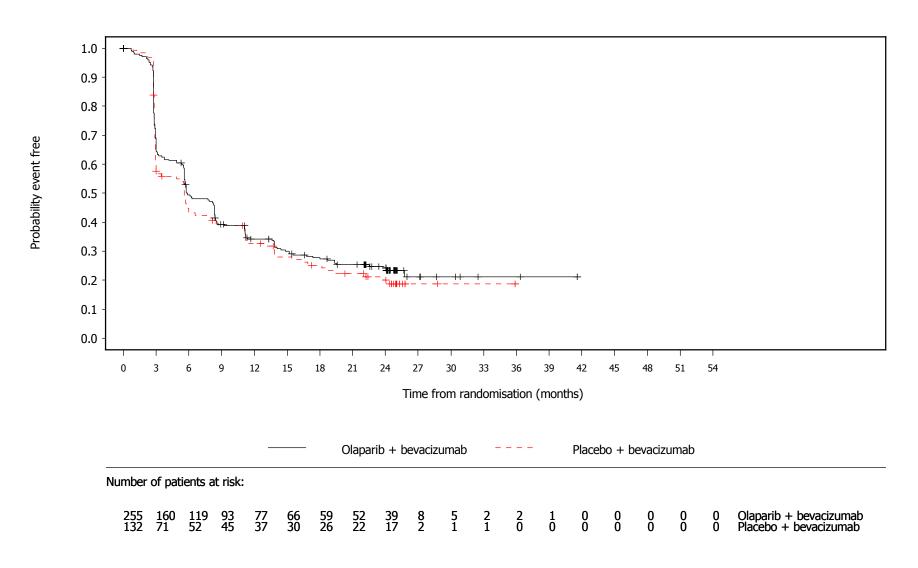
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.14 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Pain time to clinically meaningful worsening (first occurrence)

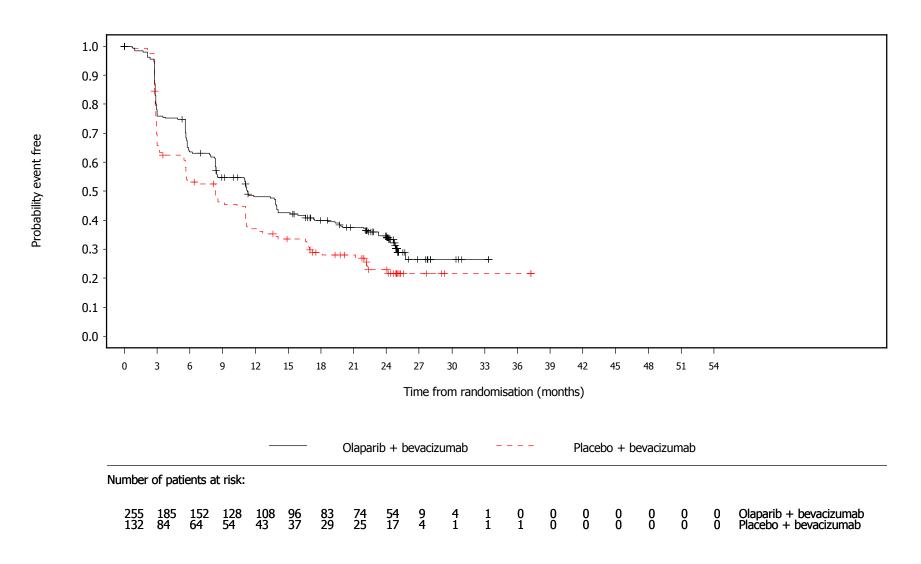
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.2.15 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Insomnia time to clinically meaningful worsening (first occurrence)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Table 2.2.3.1 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Global QoL/health status time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olap	arib + (N=2		zumab		Placebo + (N=	bevaciz 132)	umab				
Subgroup	n	of pa	er (%) atients events	(dian time 95% CI) nths) [a]	(Number (%) of patients with events	(9	ian time 5% CI) ths) [a]	Hazard ratio [b] 95% C	95% CI	I [b]	2-sided p-value [b]
First line treatment outc	ome (IV	RS)											
NED [PDS]	92	49	(53.3)	20.0	(13.6, NE)	48	28 (58.3)	14.0 (5.6,22.7)	0.72	0.46,	1.16	0.1747
NED/CR [IDS]	74	46	(62.2)	11.1	(6.5,16.6)	38	20 (52.6)	13.8 (5.5, NE)	1.17	0.70,	2.02	0.5567
NED/CR [Chemo]	40	23	(57.5)	19.7	(8.3, NE)	20	15 (75.0)	9.8 (3.5,22.3)	0.68	0.36,	1.34	0.2578
PR	49	28	(57.1)	15.3	(5.6,24.0)	26	18 (69.2)	14.0 (5.6,19.9)	0.75	0.42,	1.38	0.3464
Interaction p-value													0.4751
Screening laboratory tBRC	A statı	ıs (IV	/RS)										
tBRCAm	150	78	(52.0)	19.7	(14.0,25.5)	65	40 (61.5)	11.3 (8.3,21.4)	0.75	0.51,	1.11	0.1446
non-tBRCAm	105	68	(64.8)	11.4	(8.4,17.7)	67	41 (61.2)	13.9 (8.3,17.4)	0.94	0.64,	1.40	0.7568
Interaction p-value													0.4128
First line treatment outc	ome (e0	CRF)											
NED [PDS]	89	50	(56.2)	18.0	(11.0, NE)	47	28 (59.6)	13.8 (5.6,22.7)	0.77	0.49,	1.24	0.2767
NED/CR [IDS]	74	44	(59.5)	11.5	(6.5,22.1)	32	17 (53.1)	13.8 (5.6, NE)	1.04	0.61,	1.88	0.8796
NED/CR [Chemo]	39	19	(48.7)	22.1	(15.2, NE)	18	14 (77.8)	8.3 (3.0,17.2)	0.44	0.22,	0.90	0.0255*
PR	50	32	(64.0)	10.9	(4.9,23.6)	34	22 (64.7)	14.0 (8.3,23.5)	1.03	0.60,	1.80	0.9087
Interaction p-value													0.2199
Screening laboratory tBRC	A statı	ıs (e0	CRF)										
tBRCAm	147	76	(51.7)	20.3	(13.8,25.5)	67	40 (59.7)	11.3 (8.3,22.3)	0.77	0.52,	1.13	0.1773
non-tBRCAm	108	70	(64.8)	13.6	(8.4,16.7)	65	41 (63.1)	13.9 (8.3,17.2)	0.91	0.62,	1.35	0.6255
Interaction p-value													0.5368
Age group													
<65 years	185	109	(58.9)	15.2	(11.0,19.7)	98	56 (57.1)	16.2 (9.3,20.8)	0.97	0.70,	1.34	0.8427
>=65 years	70	37	(52.9)	22.1	(11.3, NE)	34	25 (73.5)	9.9 (5.5,15.4)	0.51	0.31,	0.86	0.0126*
Interaction p-value													0.0411*

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aaa 25NOV2020:12:08 khcs324

Table 2.2.3.1 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Global QoL/health status time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			pevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b] 95% CI [b]	2-sided p-value [b]	
FIGO Stage (Disease state)									
III	182	105 (57.7)	16.6 (11.5,21.8)	90	54 (60.0)	11.5 (8.3,17.4)	0.80	0.58, 1.12	0.1990
IV	73	41 (56.2)	15.6 (8.4,25.0)	42	27 (64.3)	14.0 (8.3,21.4)	0.85	0.53, 1.40	0.5261
Interaction p-value									0.8411
Region									
Europe	245	141 (57.6)	16.6 (11.4,20.3)	126	77 (61.1)	13.8 (9.3,17.2)	0.83	0.63, 1.10	
Japan	10	5 (50.0)	24.0 (2.8,24.0)	6	4 (66.7)	10.1 (2.8, NE)	0.60	0.16, 2.42	0.4524
Interaction p-value									0.6370
ECOG performance status at	Basel	ine							
(0) Normal activity	190	110 (57.9)	15.2 (11.1,19.7)	100	66 (66.0)	11.2 (5.8,16.4)	0.77	0.57, 1.05	0.1012
(1) Restricted activity	61	33 (54.1)	22.1 (11.4, NE)	31	15 (48.4)	18.6 (11.3, NE)	0.99	0.55, 1.88	0.9734
Interaction p-value									0.4725
Baseline CA-125 value									
<=ULN	228	125 (54.8)	19.0 (13.8,22.8)	118	73 (61.9)	13.9 (8.3,17.2)	0.74	0.56, 0.99	0.0440*
>ULN	27	21 (77.8)	5.9 (2.9,11.2)	14	8 (57.1)	11.3 (5.5, NE)	1.90	0.88, 4.58	0.1065
Interaction p-value									0.0253*
Histological grade									
High grade	255	146 (57.3)	16.6 (11.5,21.8)	132	81 (61.4)	13.8 (9.3,17.2)	0.82	0.63, 1.08	0.1557
Interaction p-value									NC
Cytoreductive surgery outco	me								
No residue	166	95 (57.2)	15.2 (11.1,20.3)	80	45 (56.3)	14.0 (8.3,19.3)	0.88	0.62, 1.27	0.5001
Residue	79	45 (57.0)	21.8 (11.3,25.0)	44	29 (65.9)	12.6 (8.3,17.2)	0.76	0.48, 1.23	0.2577
Interaction p-value									0.6172

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aaa 25NOV2020:12:08 khcs324

Table 2.2.3.1 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Global QoL/health status time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olaparib + 1 (N=2				bevacizumab 132)				2-sided p-value [b]
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	,	Hazard ratio [b] 95%	95% C]	CI [b]	
Timing of cytoreductive s	surgery									
Upfront	146	81 (55.5)	20.3 (15.3,25.0)	79	47 (59.5)	13.8 (8.3,19.3)	0.73	0.51,	1.06	0.0957
Interval	99	59 (59.6)	11.1 (6.1,16.6)	45	27 (60.0)	14.0 (5.6,23.5)	1.04	0.67,	1.67	0.8544
Interaction p-value										0.2306
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	81 (51.3)	21.8 (13.8, NE)	77	45 (58.4)	11.3 (8.3,21.4)	0.76	0.53,	1.10	0.1390
Non-tBRCAm	97	65 (67.0)	11.4 (8.4,17.7)	55	36 (65.5)	13.9 (8.3,17.2)	0.93	0.62,	1.41	0.7284
Interaction p-value										0.4575
Status somatic BRCA mutat	ions									
sBRCAm	22	9 (40.9)	25.5 (3.0, NE)	7	3 (42.9)	NE (NE, NE)	0.81	0.24,	3.64	0.7526
gBRCAm	66	37 (56.1)	13.8 (5.7, NE)	31	17 (54.8)	19.3 (8.3, NE)	1.10	0.63,	2.01	0.7389
Non-BRCAm	41	26 (63.4)	16.6 (8.3,22.1)	22	16 (72.7)	11.5 (3.5,16.6)	0.72	0.39,	1.38	0.3146
Interaction p-value										0.6127

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aaa 25NOV2020:12:08 khcs324

Table 2.2.3.2 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Physical time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)				
Subgroup	(Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% C	95% CI	I [b]	2-sided p-value [b]
First line treatment outo	come (IVI	RS)								
NED [PDS]	92	41 (44.6)	52.5 (13.8,52.5)	48	30 (62.5)	11.5 (6.2,22.4)	0.57	0.36,	0.92	0.0230*
NED/CR [IDS]	74	37 (50.0)	20.0 (8.5, NE)	38	20 (52.6)	18.9 (8.5, NE)	0.98	0.58,	1.72	0.9444
NED/CR [Chemo]	40	24 (60.0)	11.3 (5.5,22.1)	20	9 (45.0)	NE (NE, NE)	1.80	0.87,	4.10	0.1170
PR	49	23 (46.9)	19.4 (11.9, NE)	26	15 (57.7)	19.9 (5.6,28.0)	0.67	0.35,	1.32	0.2432
Interaction p-value										0.0595
Screening laboratory tBRC	CA status	s (IVRS)								
tBRCAm	150	76 (50.7)	19.4 (11.5, NE)	65	37 (56.9)	14.2 (11.2,24.9)	0.86	0.58,	1.28	0.4449
non-tBRCAm	105	49 (46.7)	22.3 (13.8,52.5)	67	37 (55.2)	18.7 (8.3,28.0)	0.78	0.51,	1.21	0.2641
Interaction p-value										0.7581
First line treatment outo	come (eCI	RF)								
NED [PDS]	89	40 (44.9)	52.5 (13.8,52.5)	47	30 (63.8)	11.5 (5.7,22.4)	0.56	0.35,	0.90	0.0177*
NED/CR [IDS]	74	35 (47.3)	23.2 (8.6, NE)	32	18 (56.3)	17.0 (6.1, NE)	0.84	0.48,	1.52	0.5532
NED/CR [Chemo]	39	25 (64.1)	11.2 (4.5,16.7)	18	8 (44.4)	19.9 (5.7, NE)	1.87	0.88,	4.44	0.1058
PR	50	23 (46.0)	22.3 (11.0, NE)	34	18 (52.9)	21.7 (11.3, NE)	0.82	0.44,	1.53	0.5183
Interaction p-value										0.0670
Screening laboratory tBRC	CA status	s (eCRF)								
tBRCAm	147	75 (51.0)	17.3 (11.3, NE)	67	38 (56.7)	15.4 (11.2,23.5)	0.87	0.59,	1.29	0.4790
non-tBRCAm	108	50 (46.3)	22.3 (13.8,52.5)	65	36 (55.4)	18.9 (8.1,28.0)	0.77	0.50,	1.19	0.2310
Interaction p-value										0.6781
Age group										
<65 years	185	89 (48.1)	20.0 (13.8, NE)	98	55 (56.1)	16.4 (11.5,27.8)	0.84	0.60,	1.18	0.3016
>=65 years	70	36 (51.4)	19.4 (11.3,52.5)	34	19 (55.9)	15.4 (6.1,22.2)	0.78	0.45,	1.39	0.3869
Interaction p-value										0.8299

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aab 25NOV2020:12:08 khcs324

Table 2.2.3.2 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Physical time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2				oevacizumab 132)			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95%	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)									
III	182	91 (50.0)	19.4 (13.8,52.5)	90	51 (56.7)	14.0 (11.1,22.2)	0.80	0.57, 1.14	0.2089
IV	73	34 (46.6)	26.3 (7.9, NE)	42	23 (54.8)	21.7 (11.2,30.7)	0.87	0.52, 1.50	0.6163
Interaction p-value									0.7873
Region									
Europe	245	123 (50.2)	16.7 (13.8,52.5)	126	72 (57.1)	15.4 (11.3,21.7)	0.82	0.61, 1.10	0.1791
Japan	10	2 (20.0)	NE (NE, NE)	6	2 (33.3)	NE (NE, NE)	0.67	0.08, 5.61	0.6931
Interaction p-value									0.8477
ECOG performance status at	Basel	ine							
(0) Normal activity	190	97 (51.1)	16.6 (13.8,52.5)	100	57 (57.0)	16.8 (11.1,24.9)	0.88	0.64, 1.23	0.4528
(1) Restricted activity	61	26 (42.6)	26.3 (13.8, NE)	31	17 (54.8)	15.4 (11.2, NE)	0.67	0.37, 1.26	0.2056
Interaction p-value									0.4389
Baseline CA-125 value									
<=ULN	228	109 (47.8)	23.2 (14.0,52.5)	118	67 (56.8)	16.2 (11.5,23.5)	0.77	0.57, 1.05	0.0952
>ULN	27	16 (59.3)	9.0 (5.6,20.0)	14	7 (50.0)	21.2 (9.3, NE)	1.48	0.63, 3.86	0.3763
Interaction p-value									0.1607
Histological grade									
High grade	255	125 (49.0)	20.0 (13.9,52.5)	132	74 (56.1)	16.4 (11.5,22.4)	0.82	0.62, 1.10	0.1913
Interaction p-value									NC
Cytoreductive surgery outco	me								
No residue	166	77 (46.4)	26.3 (14.2,52.5)	80	48 (60.0)	13.8 (8.5,21.2)	0.67	0.47, 0.97	0.0360*
Residue	79	43 (54.4)	14.1 (8.5, NE)	44	21 (47.7)	24.9 (13.9, NE)	1.27	0.76, 2.19	0.3597
Interaction p-value									0.0478*

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aab 25NOV2020:12:08 khcs324

Table 2.2.3.2 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Physical time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2				pevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events		Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
Timing of cytoreductive s	urgery									
Upfront	146	69 (47.3)	22.3 (13.9,52.5)	79	44 (55.7)	13.9 (11.2,27.8)	0.73	0.50,	1.07	0.1046
Interval	99	51 (51.5)	17.3 (8.5, NE)	45	25 (55.6)	21.2 (9.3, NE)	1.05	0.66,	1.72	0.8480
Interaction p-value										0.2393
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	78 (49.4)	19.4 (13.7, NE)	77	42 (54.5)	16.8 (11.3,27.8)	0.85	0.59,	1.25	0.4114
Non-tBRCAm	97	47 (48.5)	22.1 (11.9,52.5)	55	32 (58.2)	16.4 (8.1,28.0)	0.78	0.50,	1.24	0.2957
Interaction p-value										0.7782
Status somatic BRCA mutat	ions									
sBRCAm	22	5 (22.7)	NE (NE, NE)	7	2 (28.6)	NE (NE, NE)	0.63	0.14,	4.42	0.5984
gBRCAm	66	36 (54.5)	15.9 (8.5, NE)	31	17 (54.8)	16.8 (11.1, NE)	0.98	0.56,	1.79	0.9503
Non-BRCAm	41	18 (43.9)	52.5 (14.2,52.5)	22	9 (40.9)	NE (NE, NE)	1.04	0.47,	2.44	0.9246
Interaction p-value										0.8724

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[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aab 25NOV2020:12:08 khcs324

Table 2.2.3.3 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Role time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olapa		bevacizumab 255)		Pl		bevacizumab 132)	:			
Subgroup	n	of pa	er (%) tients events	Median time (95% CI) (months) [a]		of	nber (%) patients h events		Hazard ratio [b]	95% CI	5% CI [b]	2-sided p-value [b]
First line treatment outo	come (IV	RS)										
NED [PDS]	92	58	(63.0)	11.1 (5.8,18.	0) 4	8 2	9 (60.4)	11.3 (3.2,24.1)	0.95	0.61,	1.50	0.8191
NED/CR [IDS]	74	51	(68.9)	5.6 (3.0,11.	1) 3	8 2	0 (52.6)	9.6 (8.3, NE)	1.70	1.03,	2.92	0.0374*
NED/CR [Chemo]	40	23	(57.5)	8.5 (3.3, 1	IE) 2	0 1	4 (70.0)	4.6 (2.9,13.8)	0.76	0.39,	1.51	0.4181
PR	49	35	(71.4)	8.5 (4.9,13.	8) 2	6 1	9 (73.1)	11.1 (2.9,19.9)	0.88	0.51,	1.57	0.6567
Interaction p-value												0.1710
Screening laboratory tBRO	CA statu	ıs (IV	RS)									
tBRCAm	150	92	(61.3)	8.9 (5.8,16.	8) 6	5 3	7 (56.9)	11.9 (5.7,27.7)	1.12	0.77,	1.66	0.5529
non-tBRCAm	105	75	(71.4)	8.3 (4.8,11.	1) 6	7 4	5 (67.2)	8.3 (5.6,14.9)	1.06	0.74,	1.55	0.7386
Interaction p-value												0.8488
First line treatment outo	come (eC	RF)										
NED [PDS]	89	54	(60.7)	11.1 (8.1,24.	4) 4	7 2	8 (59.6)	8.5 (3.0,24.1)	0.88	0.56,	1.41	0.5802
NED/CR [IDS]	74	52	(70.3)	5.5 (2.9,11.	1) 3	2 1	7 (53.1)	9.6 (8.3, NE)	1.79	1.06,	3.19	0.0297*
NED/CR [Chemo]	39	25	(64.1)	8.2 (3.0,15.	2) 1	8 1	3 (72.2)	5.7 (2.9,13.8)	0.86	0.45,	1.74	0.6711
PR	50	35	(70.0)	8.9 (5.7,15.	4) 3	4 2	3 (67.6)	11.1 (3.0,22.1)	0.97	0.58,	1.66	0.9064
Interaction p-value												0.1821
Screening laboratory tBRO	CA statu	ıs (eC	RF)									
tBRCAm	147	89	(60.5)	8.9 (5.8,17.	3) 6	7 3	8 (56.7)	11.3 (5.7,27.7)	1.08	0.75,	1.60	0.6790
non-tBRCAm	108	78	(72.2)	8.3 (4.8,11.	1) 6	5 4	4 (67.7)	8.3 (5.6,14.9)	1.09	0.76,	1.60	0.6343
Interaction p-value												0.9713
Age group												
<65 years	185	121	(65.4)	8.5 (6.2,11.	3) 9	8 6	1 (62.2)	8.5 (5.6,14.9)	1.03	0.76,	1.41	0.8594
>=65 years	70	46	(65.7)	5.8 (3.3,11.	9) 3	4 2	1 (61.8)	13.8 (5.6,20.2)	1.17	0.71,	2.00	0.5526
Interaction p-value												0.6768

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aac 25NoV2020:12:08 khcs324

Table 2.2.3.3 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Role time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			oevacizumab 132)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95%	95% CI [CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	117 (64.3)	8.4 (5.7,11.2)	90	56 (62.2)	11.1 (5.7,19.9)	1.08	0.79, 1	L.50	0.6282
IV	73	50 (68.5)	10.4 (5.4,14.4)	42	26 (61.9)	8.3 (5.5,22.1)	1.02	0.64, 1	L.67	0.9201
Interaction p-value										0.8527
Region										
Europe	245	163 (66.5)	8.3 (5.8,11.1)	126	79 (62.7)	9.3 (5.7,16.2)	1.07	0.82, 1	L.40	0.6370
Japan	10	4 (40.0)	NE (NE, NE)	6	3 (50.0)	NE (NE, NE)	0.84	0.18, 4	1.25	0.8163
Interaction p-value										0.7558
ECOG performance status at	Basel	ine								
(0) Normal activity	190	129 (67.9)	8.4 (5.8,11.1)	100	64 (64.0)	8.5 (5.6,16.4)	1.09	0.81, 1	L.48	0.5740
(1) Restricted activity	61	36 (59.0)	7.2 (3.0,25.3)	31	18 (58.1)	9.6 (5.5,27.7)	1.04	0.60, 1	L.88	0.8851
Interaction p-value										0.8931
Baseline CA-125 value										
<=ULN	228		8.4 (5.8,11.2)	118		8.5 (5.7,18.7)	1.07	0.81, 1		0.6454
>ULN	27	18 (66.7)	10.4 (3.5,19.9)	14	9 (64.3)	9.6 (2.8, NE)	1.03	0.47, 2	2.40	0.9481
Interaction p-value										0.9282
Histological grade										
High grade	255	167 (65.5)	8.4 (5.8,11.2)	132	82 (62.1)	9.3 (6.1,16.2)	1.06	0.82, 1	L.39	0.6480
Interaction p-value										NC
Cytoreductive surgery outco	me									
No residue	166	107 (64.5)	8.3 (5.6,11.3)	80	46 (57.5)	9.3 (8.1,21.9)	1.17	0.83, 1	L.66	0.3804
Residue	79	53 (67.1)	8.5 (5.8,13.8)	44	30 (68.2)	7.9 (3.5,19.9)	0.92	0.59, 1	L.46	0.7197
Interaction p-value										0.4156

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aac 25NOV2020:12:08 khcs324

Table 2.2.3.3 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Role time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	•	Olaparib + 1 (N=2				oevacizumab 132)			
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]	Number (%) of patients n with events		Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	urgery								
Upfront	146	92 (63.0)	11.1 (8.2,14.0)	79	50 (63.3)	8.2 (3.5,16.4)	0.86	0.61, 1.22	0.4003
Interval	99	68 (68.7)	5.7 (3.1,11.1)	45	26 (57.8)	9.6 (8.3, NE)	1.52	0.98, 2.42	0.0636
Interaction p-value									0.0490*
Myriad tumour BRCA mutati	on stat	tus							
tBRCAm	158	100 (63.3)	8.5 (5.8,13.1)	77	45 (58.4)	11.1 (5.6,22.1)	1.06	0.75, 1.53	0.7257
Non-tBRCAm	97	67 (69.1)	8.3 (4.8,11.2)	55	37 (67.3)	8.3 (5.6,18.7)	1.08	0.73, 1.64	0.6910
Interaction p-value									0.9458
Status somatic BRCA mutat	ions								
sBRCAm	22	11 (50.0)	11.1 (3.0, NE)	7	2 (28.6)	NE (NE, NE)	1.90	0.51, 12.26	0.3718
gBRCAm	66	46 (69.7)	5.6 (3.0,12.5)	31	16 (51.6)	23.5 (8.3, NE)	1.70	0.98, 3.10	0.0583
Non-BRCAm	41	27 (65.9)	9.7 (3.0,19.9)	22	16 (72.7)	8.2 (3.5,21.9)	0.94	0.51, 1.78	0.8444
Interaction p-value									0.3462

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aac 25NOV2020:12:08 khcs324

Table 2.2.3.4 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Cognitive time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olapa	arib + 1 (N=2	pevacizumab 55)			pevacizumab 132)				2-sided p-value [b]
Subgroup		er (%) ntients events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	ratio		
First line treatment outco	ome (IVRS)									
NED [PDS]	92 63	(68.5)	11.1 (6.3,14.0)	48	31 (64.6)	8.5 (5.6,16.4)	0.91	0.60,	1.42	0.6812
NED/CR [IDS]	74 47	(63.5)	14.8 (8.3,19.4)	38	21 (55.3)	16.3 (5.6, NE)	1.07	0.65,	1.83	0.7821
NED/CR [Chemo]	40 28	(70.0)	15.2 (8.3,19.4)	20	17 (85.0)	5.6 (2.9, 8.5)	0.52	0.29,	0.97	0.0406*
PR	49 36	(73.5)	8.4 (3.7,11.3)	26	16 (61.5)	12.7 (5.5,15.4)	1.12	0.63,	2.08	0.6976
Interaction p-value										0.2601
Screening laboratory tBRC	A status (IV	RS)								
tBRCAm	150 104	(69.3)	11.3 (8.7,16.8)	65	44 (67.7)	8.3 (5.7,13.6)	0.84	0.59,	1.20	0.3237
non-tBRCAm	105 70	(66.7)	8.4 (5.8,14.8)	67	41 (61.2)	11.1 (5.7,14.6)	1.01	0.69,	1.50	0.9397
Interaction p-value										0.4651
First line treatment outco	ome (eCRF)									
NED [PDS]	89 59	(66.3)	11.1 (6.2,14.0)	47	30 (63.8)	8.7 (5.8,16.4)	0.92	0.60,	1.45	0.7268
NED/CR [IDS]	74 48	(64.9)	11.5 (8.3,17.3)	32	19 (59.4)	14.1 (5.6,22.2)	1.03	0.62,	1.80	0.9067
NED/CR [Chemo]	39 27	(69.2)	15.2 (8.3,22.1)	18	13 (72.2)	8.2 (5.5,13.1)	0.67	0.35,	1.34	0.2472
PR	50 37	(74.0)	8.5 (4.9,13.8)	34	22 (64.7)	8.6 (3.0,15.4)	1.03	0.61,	1.77	0.9151
Interaction p-value										0.7522
Screening laboratory tBRC	A status (eC	RF)								
tBRCAm	147 102	(69.4)	11.3 (8.7,16.8)	67	45 (67.2)	8.3 (5.7,13.6)	0.83	0.59,	1.20	0.3161
non-tBRCAm	108 72	(66.7)	8.4 (5.8,14.8)	65	40 (61.5)	11.1 (5.6,16.4)	1.02	0.70,	1.51	0.9299
Interaction p-value										0.4546
Age group										
<65 years	185 127	(68.6)	11.2 (8.5,14.8)	98	65 (66.3)	8.5 (5.7,13.1)	0.89	0.67,	1.21	0.4654
>=65 years	70 47	(67.1)	10.3 (5.8,16.6)	34	20 (58.8)	12.7 (5.7,21.4)	0.97	0.58,	1.67	0.9001
Interaction p-value										0.7981

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aad 25NOV2020:12:08 khcs324

Table 2.2.3.4 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Cognitive time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95	95% CI [}	2-sided o] p-value[b]
FIGO Stage (Disease state)									
III	182	128 (70.3)	11.0 (8.3,13.8)	90	56 (62.2)	8.7 (5.7,14.4)	1.04	0.76, 1	.43 0.8185
IV	73	46 (63.0)	14.8 (8.4,22.3)	42	29 (69.0)	8.3 (5.6,15.4)	0.67	0.42, 1	.08 0.1014
Interaction p-value									0.1331
Region									
Europe	245	170 (69.4)	11.1 (8.4,13.8)	126	82 (65.1)	8.5 (5.8,12.7)	0.91	0.70, 1	
Japan	10	4 (40.0)	NE (NE, NE)	6	3 (50.0)	NE (NE, NE)	0.78	0.17, 3	.94 0.7407
Interaction p-value									0.8416
ECOG performance status at	Basel	ine							
(0) Normal activity	190	125 (65.8)	11.4 (8.5,16.3)	100	67 (67.0)	8.3 (5.7,12.5)	0.81	0.60, 1	.09 0.1579
(1) Restricted activity	61	46 (75.4)	8.6 (5.6,14.0)	31	18 (58.1)	14.1 (5.6,21.4)	1.31	0.77, 2	.31 0.3266
Interaction p-value									0.1194
Baseline CA-125 value									
<=ULN	228		11.2 (8.5,14.3)	118		8.5 (5.9,13.1)	0.88	0.67, 1	
>ULN	27	19 (70.4)	6.1 (3.0,16.6)	14	8 (57.1)	21.2 (2.8, NE)	1.23	0.56, 2	
Interaction p-value									0.4413
Histological grade									
High grade	255	174 (68.2)	11.1 (8.5,14.0)	132	85 (64.4)	8.5 (5.9,13.6)	0.91	0.71, 1	.19 0.4856
Interaction p-value									NC
Cytoreductive surgery outco	ome								
No residue	166	, ,	11.1 (8.3,15.9)	80	, ,	8.7 (5.8,16.4)	0.96	0.69, 1	
Residue	79	56 (70.9)	12.1 (8.5,16.6)	44	29 (65.9)	8.3 (5.6,13.1)	0.85	0.55, 1	
Interaction p-value									0.6809

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aad 25NOV2020:12:08 khcs324

Table 2.2.3.4 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Cognitive time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	;	Olaparib + 1 (N=2				oevacizumab 132)				
	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	,	Hazard ratio [b] 95% ([b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	100 (68.5)	11.1 (8.3,14.0)	79	52 (65.8)	8.5 (5.6,12.5)	0.86	0.62,	1.21	0.3738
Interval	99	66 (66.7)	13.0 (8.3,16.8)	45	27 (60.0)	11.1 (5.7,21.2)	1.03	0.67,	1.64	0.8939
Interaction p-value										0.5177
Myriad tumour BRCA mutati	on stat	tus								
tBRCAm	158	110 (69.6)	11.2 (8.7,16.3)	77	50 (64.9)	8.3 (5.8,14.1)	0.86	0.62,	1.21	0.3790
Non-tBRCAm	97	64 (66.0)	8.6 (5.8,15.2)	55	35 (63.6)	11.1 (5.6,14.6)	1.00	0.66,	1.52	0.9839
Interaction p-value										0.5854
Status somatic BRCA mutat	ions									
sBRCAm	22	16 (72.7)	8.5 (2.9,19.4)	7	4 (57.1)	9.5 (2.8, NE)	1.10	0.40,	3.88	0.8596
gBRCAm	66	43 (65.2)	16.8 (8.7,20.3)	31	20 (64.5)	8.7 (5.6,19.2)	0.80	0.48,	1.40	0.4260
Non-BRCAm	41	31 (75.6)	8.3 (5.6,16.6)	22	14 (63.6)	14.0 (5.6,22.2)	1.21	0.66,	2.35	0.5436
Interaction p-value										0.6004

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aad 25NOV2020:12:08 khcs324

Table 2.2.3.5 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Emotional time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	·	Olapar	rib + 1 (N=2	bevacizumab 255)			bevacizumab 132)				
Subgroup	n	Number of pata with e	ients	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	ratio	2-sided p-value [b]	
First line treatment out	come (IV	RS)									
NED [PDS]	92	53 (!	57.6)	13.9 (8.7,24.2)	48	30 (62.5)	11.3 (5.7,27.8)	0.86	0.55,	1.36	0.5120
NED/CR [IDS]	74	50 (6	67.6)	8.3 (5.6,15.6)	38	27 (71.1)	8.7 (5.9,13.9)	0.92	0.58,	1.50	0.7442
NED/CR [Chemo]	40	23 (!	57.5)	19.7 (8.5,25.0)	20	11 (55.0)	22.0 (8.3, NE)	1.13	0.56,	2.41	0.7347
PR	49	32 (65.3)	12.1 (5.7,22.1)	26	17 (65.4)	9.0 (2.9,13.9)	0.72	0.41,	1.33	0.2881
Interaction p-value											0.8112
Screening laboratory tBR0	CA statı	ıs (IVR	S)								
tBRCAm	150	92 (61.3)	15.6 (10.6,22.1)	65	40 (61.5)	11.2 (6.9,22.2)	0.93	0.64,	1.36	0.6835
non-tBRCAm	105	66 (6	62.9)	11.1 (5.7,16.7)	67	45 (67.2)	11.1 (8.3,13.9)	0.90	0.61,	1.32	0.5703
Interaction p-value											0.9043
First line treatment out	come (e0	CRF)									
NED [PDS]	89	54 (6	60.7)	13.9 (8.7,22.1)	47	29 (61.7)	11.2 (5.7,27.8)	0.94	0.60,	1.50	0.7918
NED/CR [IDS]	74	48 (64.9)	10.6 (5.6,19.8)	32	23 (71.9)	11.0 (6.6,14.1)	0.87	0.54,	1.46	0.5876
NED/CR [Chemo]	39	22 (56.4)	19.6 (8.5,25.0)	18	10 (55.6)	22.0 (3.5, NE)	0.98	0.48,	2.17	0.9595
PR	50	33 (6	66.0)	9.0 (5.7,19.7)	34	22 (64.7)	11.1 (5.6,13.9)	0.86	0.51,	1.50	0.5942
Interaction p-value											0.9885
Screening laboratory tBR0	CA statı	ıs (eCRI	F)								
tBRCAm	147	91 (61.9)	14.0 (8.5,20.7)	67	41 (61.2)	11.2 (8.3,21.4)	0.94	0.66,	1.38	0.7560
non-tBRCAm	108	67 (6	62.0)	11.2 (5.8,19.7)	65	44 (67.7)	10.2 (8.3,13.9)	0.86	0.59,	1.27	0.4427
Interaction p-value											0.7358
Age group											
<65 years	185	122 (65.9)	10.0 (5.9,14.0)	98	65 (66.3)	10.2 (6.9,11.4)	0.96	0.71,	1.30	0.7748
>=65 years	70	36 (!	51.4)	22.4 (12.1, NE)	34	20 (58.8)	14.0 (9.0,26.5)	0.75	0.44,	1.33	0.3183
Interaction p-value											0.4570

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aae 25NOV2020:12:08 khcs324

Table 2.2.3.5 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Emotional time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			bevacizumab 132)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	- Hazard ratio [b] 95% CI [b	[b]	2-sided] p-value [b]	
FIGO Stage (Disease state)										
III	182	115 (63.2)	13.8 (8.7,19.4)	90	56 (62.2)	11.1 (8.3,14.0)	0.96	0.70,	1.33	0.7829
IV	73	43 (58.9)	12.1 (6.3,24.1)	42	29 (69.0)	11.0 (6.9,13.9)	0.77	0.48,	1.24	0.2787
Interaction p-value										0.4544
Region										
Europe	245	154 (62.9)	13.1 (8.5,18.0)	126	82 (65.1)	11.1 (8.3,12.5)	0.87	0.67,	1.15	0.3215
Japan	10	4 (40.0)	NE (NE, NE)	6	3 (50.0)	24.0 (13.8, NE)	1.09	0.24,	5.51	0.9146
Interaction p-value										0.7774
ECOG performance status at	Basel	ine								
(0) Normal activity	190	118 (62.1)	13.1 (8.4,18.0)	100	63 (63.0)	11.2 (8.3,13.9)	0.96	0.71,	1.31	0.7748
(1) Restricted activity	61	37 (60.7)	13.8 (8.4,24.0)	31	22 (71.0)	8.4 (6.9,14.1)	0.71	0.42,	1.22	0.2053
Interaction p-value										0.3345
Baseline CA-125 value										
<=ULN	228		13.8 (10.0,19.3)	118		11.1 (8.4,13.9)	0.90	0.68,		0.4456
>ULN	27	16 (59.3)	9.0 (5.7,29.3)	14	9 (64.3)	6.9 (5.6,21.4)	0.87	0.39,	2.05	0.7314
Interaction p-value										0.9369
Histological grade										
High grade	255	158 (62.0)	13.8 (9.0,19.3)	132	85 (64.4)	11.1 (8.3,13.8)	0.89	0.69,	1.17	0.4051
Interaction p-value										NC
Cytoreductive surgery outco	me									
No residue	166	103 (62.0)	11.4 (7.9,19.8)	80	53 (66.3)	11.1 (6.9,13.9)	0.90	0.65,	1.26	0.5355
Residue	79	49 (62.0)	16.7 (8.5,22.1)	44	26 (59.1)	10.0 (5.6,23.0)	0.89	0.56,	1.46	0.6446
Interaction p-value										0.9808

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aae 25NoV2020:12:08 khcs324

Table 2.2.3.5 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Emotional time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olaparib + 1 (N=2				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	,	Hazard ratio [b] 95% CI [b]		2-sided p-value [b]	
Timing of cytoreductive s	surgery									
Upfront	146	87 (59.6)	16.9 (11.4,22.1)	79	47 (59.5)	11.2 (8.3,23.0)	0.89	0.62,	1.27	0.5061
Interval	99	65 (65.7)	8.4 (5.7,15.6)	45	32 (71.1)	8.7 (5.9,13.9)	0.90	0.60,	1.40	0.6435
Interaction p-value										0.9421
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	98 (62.0)	13.9 (10.6,19.8)	77	47 (61.0)	11.2 (8.3,21.4)	0.94	0.67,	1.34	0.7195
Non-tBRCAm	97	60 (61.9)	11.1 (5.7,20.4)	55	38 (69.1)	9.0 (6.2,13.8)	0.85	0.57,	1.28	0.4251
Interaction p-value										0.7069
Status somatic BRCA mutat	ions									
sBRCAm	22	12 (54.5)	11.1 (5.6, NE)	7	5 (71.4)	5.6 (2.8, NE)	0.53	0.19,	1.67	0.2569
gBRCAm	66	42 (63.6)	13.8 (6.0,22.1)	31	18 (58.1)	11.1 (5.7,35.0)	1.05	0.61,	1.88	0.8547
Non-BRCAm	41	23 (56.1)	19.7 (5.7, NE)	22	15 (68.2)	11.1 (8.3,13.9)	0.77	0.41,	1.51	0.4382
Interaction p-value										0.4989

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aae 25NoV2020:12:08 khcs324

Table 2.2.3.6 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Social time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olap	arib + (N=2		zumab		Placebo + (N=	bevaci 132)	zumab				2-sided p-value [b]
Subgroup	n	of p	er (%) atients events	(9	ian time 95% CI) nths) [a]		Number (%) of patients with events	3	dian time (95% CI) onths) [a]	Hazard ratio [b]	ratio		
First line treatment outc	ome (IV	/RS)											
NED [PDS]	92	54	(58.7)	13.9	(8.3,23.1)	48	34 (70.8)	10.4	(6.2,14.2)	0.70	0.46,	1.08	0.1088
NED/CR [IDS]	74	47	(63.5)	11.1	(5.7,19.8)	38	21 (55.3)	13.9	(5.9,29.3)	1.27	0.77,	2.16	0.3596
NED/CR [Chemo]	40	19	(47.5)	24.0	(8.6, NE)	20	13 (65.0)	14.3	(3.5, NE)	0.72	0.36,	1.49	0.3631
PR	49	28	(57.1)	8.5	(5.3, NE)	26	13 (50.0)	13.9	(2.9, NE)	0.94	0.50,	1.88	0.8546
Interaction p-value													0.3281
Screening laboratory tBRC	A statı	ıs (I	JRS)										
tBRCAm	150	86	(57.3)	15.9	(8.6,23.1)	65	37 (56.9)	14.0	(11.1,24.1)	1.01	0.69,	1.50	0.9627
non-tBRCAm	105	62	(59.0)	11.2	(5.8,19.8)	67	44 (65.7)	9.9	(5.7,14.6)	0.78	0.53,	1.16	0.2221
Interaction p-value													0.3641
First line treatment outc	ome (e0	CRF)											
NED [PDS]	89	53	(59.6)	13.5	(8.3,23.1)	47	33 (70.2)	10.4	(6.2,16.4)	0.73	0.47,	1.14	0.1589
NED/CR [IDS]	74	46	(62.2)	11.1	(5.6,19.9)	32	19 (59.4)	11.5	(5.8,29.3)	1.10	0.66,	1.92	0.7216
NED/CR [Chemo]	39	19	(48.7)	23.7	(8.6, NE)	18	10 (55.6)	16.7	(8.3, NE)	0.86	0.41,	1.93	0.7066
PR	50	28	(56.0)	8.5	(5.5, NE)	34	18 (52.9)	13.9	(3.0, NE)	0.99	0.55,	1.82	0.9670
Interaction p-value													0.6684
Screening laboratory tBRC	A statı	ıs (e	CRF)										
tBRCAm	147	83	(56.5)	15.9	(8.5,23.2)	67	38 (56.7)	14.0	(9.3,24.1)	0.98	0.67,	1.45	0.9046
non-tBRCAm	108	65	(60.2)	11.3	(5.8,19.6)	65	43 (66.2)	9.9	(5.7,15.5)	0.81	0.55,	1.19	0.2803
Interaction p-value													0.4913
Age group													
<65 years	185	105	(56.8)	13.6	(8.4,19.6)	98	60 (61.2)	13.9	(8.5,19.4)	0.92	0.67,	1.27	0.5964
>=65 years	70	43	(61.4)	11.3	(6.3,23.2)	34	21 (61.8)	9.9	(5.8,15.5)	0.77	0.46,	1.33	0.3483
Interaction p-value													0.5912

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aaf 25NoV2020:12:08 khcs324

Table 2.2.3.6 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Social time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			bevacizumab 132)		Hazard ratio [b] 95% CI [b]	2-sided p-value [b]
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]			
FIGO Stage (Disease state)									
III	182	108 (59.3)	11.3 (8.4,18.0)	90	54 (60.0)	11.5 (8.5,20.0)	0.95	0.69, 1.32	0.7495
IV	73	40 (54.8)	16.0 (7.9, NE)	42	27 (64.3)	11.3 (5.8,19.4)	0.74	0.46, 1.22	0.2392
Interaction p-value									0.4186
Region									
Europe	245	145 (59.2)	11.4 (8.5,18.0)	126	79 (62.7)	11.2 (8.5,15.5)	0.87	0.66, 1.14	0.3071
Japan	10	3 (30.0)	NE (NE, NE)	6	2 (33.3)	NE (NE, NE)	1.02	0.17, 7.77	0.9800
Interaction p-value									0.8557
ECOG performance status at	Basel	ine							
(0) Normal activity	190	105 (55.3)	16.0 (11.2,23.1)	100	67 (67.0)	11.2 (8.3,14.6)	0.72	0.53, 0.99	0.0421*
(1) Restricted activity	61	41 (67.2)	5.7 (3.0,11.9)	31	14 (45.2)	19.4 (6.1, NE)	1.74	0.97, 3.31	0.0628
Interaction p-value									0.0088*
Baseline CA-125 value									
<=ULN	228		14.1 (11.1,20.7)	118		13.8 (8.7,16.7)	0.86	0.65, 1.16	0.3268
>ULN	27	19 (70.4)	5.7 (3.5,16.6)	14	9 (64.3)	5.6 (2.8, NE)	1.00	0.46, 2.32	0.9984
Interaction p-value									0.7354
Histological grade									
High grade	255	148 (58.0)	13.5 (8.6,19.6)	132	81 (61.4)	11.3 (8.5,16.4)	0.88	0.67, 1.16	0.3632
Interaction p-value									NC
Cytoreductive surgery outco	me								
No residue	166	101 (60.8)	13.5 (8.3,17.3)	80	53 (66.3)	10.4 (8.3,14.2)	0.85	0.61, 1.19	0.3403
Residue	79	41 (51.9)	19.6 (8.4, NE)	44	24 (54.5)	16.6 (11.1, NE)	0.92	0.56, 1.54	0.7403
Interaction p-value									0.7998

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aaf 25NoV2020:12:08 khcs324

Table 2.2.3.6 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Social time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	,	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery								
Upfront	146	82 (56.2)	16.0 (10.6,24.0)	79	51 (64.6)	11.3 (8.5,16.6)	0.76	0.54, 1.08	0.1257
Interval	99	60 (60.6)	11.1 (5.8,19.9)	45	26 (57.8)	13.9 (6.1,29.3)	1.12	0.71, 1.80	0.6296
Interaction p-value									0.1835
Myriad tumour BRCA mutati	ion stat	us							
tBRCAm	158	89 (56.3)	15.9 (10.6,23.2)	77	44 (57.1)	14.0 (9.3,24.1)	0.93	0.65, 1.35	0.6942
Non-tBRCAm	97	59 (60.8)	11.1 (5.6,16.6)	55	37 (67.3)	9.9 (5.7,14.6)	0.83	0.56, 1.27	0.3941
Interaction p-value									0.6998
Status somatic BRCA mutat	cions								
sBRCAm	22	8 (36.4)	NE (NE, NE)	7	1 (14.3)	NE (NE, NE)	2.94	0.54, 54.64	0.2436
gBRCAm	66	44 (66.7)	8.5 (4.2,14.1)	31	15 (48.4)	24.1 (11.3, NE)	1.79	1.02, 3.33	0.0425*
Non-BRCAm	41	26 (63.4)	11.9 (5.4,41.9)	22	16 (72.7)	11.0 (5.6,15.5)	0.70	0.38, 1.35	0.2807
Interaction p-value									0.0693

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aaf 25NOV2020:12:08 khcs324

Table 2.2.3.7 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Loss of appetite time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olap	arib + (N=2	bevaci 255)	zumab		Plac	ebo + l (N=)	bevaci 132)	zumab				
Subgroup	n	of pa	er (%) atients events	(lian time 95% CI) nths) [a]		of pa	er (%) atients events		dian time (95% CI) onths) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)												
NED [PDS]	92	44	(47.8)	25.5	(11.2,49.3)	48	23	(47.9)	25.6	(16.4,35.0)	1.01	0.62,	1.71	0.9650
NED/CR [IDS]	74	50	(67.6)	9.7	(6.2,19.3)	38	17	(44.7)	29.8	(13.9, NE)	2.08	1.23,	3.72	0.0059*
NED/CR [Chemo]	40	24	(60.0)	11.3	(8.1,24.2)	20	10	(50.0)	22.0	(8.7, NE)	1.51	0.74,	3.32	0.2585
PR	49	28	(57.1)	12.1	(5.8,25.7)	26	15	(57.7)	14.0	(6.4,21.7)	0.97	0.52,	1.87	0.9265
Interaction p-value														0.1822
Screening laboratory tBR0	CA statu	ıs (IV	JRS)											
tBRCAm	150	86	(57.3)	13.6	(10.9,23.1)	65	30	(46.2)	25.6	(16.7,35.0)	1.50	1.003,	2.31	0.0484*
non-tBRCAm	105	60	(57.1)	13.8	(8.3,24.2)	67	35	(52.2)	17.4	(14.4,28.7)	1.18	0.78,	1.80	0.4476
Interaction p-value														0.4151
First line treatment out	come (eC	RF)												
NED [PDS]	89	43	(48.3)	24.2	(11.3,49.3)	47	23	(48.9)	25.6	(16.4,35.0)	0.99	0.60,	1.68	0.9836
NED/CR [IDS]	74	52	(70.3)	8.3	(5.6,13.6)	32	14	(43.8)	29.8	(11.3,29.8)	2.39	1.36,	4.49	0.0019*
NED/CR [Chemo]	39	21	(53.8)	11.3	(8.2, NE)	18	7	(38.9)	NE	(NE, NE)	1.67	0.74,	4.24	0.2214
PR	50	28	(56.0)	13.7	(5.8,25.7)	34	21	(61.8)	14.7	(8.3,21.7)	0.90	0.51,	1.60	0.7041
Interaction p-value														0.0571
Screening laboratory tBR0	CA statu	ıs (e0	CRF)											
tBRCAm	147	84	(57.1)	13.6	(8.7,23.1)	67	30	(44.8)	25.6	(16.7,35.0)	1.53	1.02,	2.36	0.0395*
non-tBRCAm	108	62	(57.4)	13.8	(8.3,24.2)	65	35	(53.8)	17.4	(14.4,28.7)	1.15	0.76,	1.76	0.5118
Interaction p-value														0.3402
Age group														
<65 years	185	104	(56.2)	13.8	(11.0,22.1)	98	47	(48.0)	24.0	(16.7,28.7)	1.34	0.95,	1.90	0.0941
>=65 years	70	42	(60.0)	11.5	(7.9,24.2)	34	18	(52.9)	16.6	(12.7,29.8)	1.28	0.75,	2.29	0.3743
Interaction p-value														0.9004

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aag 25NoV2020:12:08 khcs324

Table 2.2.3.7 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Loss of appetite time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olapa	rib + (N=2		zumab			- bev N=132	vacizumab 2)				
Subgroup	n	Number of pat with e	ients	(!	ian time 95% CI) nths) [a]		Number (% of patient with event	s	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	100 (54.9)	19.2	(11.1,24.2)	90	43 (47.8	3) 25	5.6 (16.7,35.0)	1.32	0.93,	1.90	0.1250
IV	73	46 (63.0)	11.2	(6.1,14.8)	42	22 (52.4	1) 19	9.7 (11.2,28.7)	1.36	0.83,	2.31	0.2245
Interaction p-value													0.9137
Region													
Europe	245	140 (57.1)	13.4	(11.0,21.7)	126	61 (48.4	1) 22	2.3 (16.4,29.8)	1.33	0.99,		0.0560
Japan	10	6 (60.0)	22.1	(2.8, NE)	6	4 (66.5	7) 23	3.0 (2.8,24.0)	1.15	0.33,	4.49	0.8293
Interaction p-value													0.8219
ECOG performance status at	Basel	ine											
(0) Normal activity	190	106 (55.8)	13.6	(11.0,22.1)	100	48 (48.0)) 25	5.6 (16.6,29.8)	1.38	0.99,	1.96	0.0591
(1) Restricted activity	61	36 (59.0)	12.8	(5.6,25.7)	31	17 (54.8	3) 16	6.7 (8.4, NE)	1.10	0.63,	2.01	0.7395
Interaction p-value													0.5129
Baseline CA-125 value													
<=ULN	228	129 (56.6)	13.7	(11.1,22.4)	118	59 (50.0)) 22	2.3 (16.4,28.7)	1.26	0.93,	1.73	0.1390
>ULN	27	17 (63.0)	11.2	(5.6,22.1)	14	6 (42.9	9) 21	1.7 (9.3, NE)	2.03	0.84,	5.62	0.1185
Interaction p-value													0.3276
Histological grade													
High grade	255	146 (57.3)	13.6	(11.1,22.1)	132	65 (49.2	2) 22	2.3 (16.6,28.7)	1.32	0.99,	1.78	0.0571
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	97 (58.4)	13.6	(9.7,22.1)	80	37 (46.3	3) 25	5.6 (17.4,29.8)	1.49	1.03,	2.21	0.0333*
Residue	79	41 (51.9)	22.1	(10.9, NE)	44	21 (47.7	7) 23	1.7 (14.6, NE)	1.18	0.70,	2.03	0.5373
Interaction p-value													0.4773

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aag 25NoV2020:12:08 khcs324

Table 2.2.3.7 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Loss of appetite time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	·	Olaparib + 1 (N=2				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	,	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	70 (47.9)	24.2 (15.2,49.3)	79	37 (46.8)	25.6 (17.4,35.0)	1.02	0.69,	1.53	0.9306
Interval	99	68 (68.7)	8.3 (5.7,11.5)	45	21 (46.7)	21.7 (14.6, NE)	2.17	1.36,	3.63	0.0010*
Interaction p-value										0.0172*
Myriad tumour BRCA mutati	ion stat	us								
tBRCAm	158	92 (58.2)	11.5 (8.7,22.4)	77	34 (44.2)	27.8 (19.7,35.0)	1.60	1.09,	2.40	0.0160*
Non-tBRCAm	97	54 (55.7)	15.2 (8.4,25.7)	55	31 (56.4)	16.4 (12.7,22.0)	1.02	0.66,	1.61	0.9283
Interaction p-value										0.1393
Status somatic BRCA mutat	ions									
sBRCAm	22	12 (54.5)	11.0 (2.9, NE)	7	1 (14.3)	NE (NE, NE)	5.15	1.01,	93.99	0.0478*
gBRCAm	66	49 (74.2)	8.3 (4.6,11.3)	31	15 (48.4)	24.0 (11.3,35.0)	2.16	1.24,	3.99	0.0059*
Non-BRCAm	41	24 (58.5)	11.9 (4.8,49.3)	22	14 (63.6)	14.7 (8.7, NE)	1.03	0.54,	2.06	0.9239
Interaction p-value										0.1189

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aag 25NoV2020:12:08 khcs324

Table 2.2.3.8 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Constipation time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2			Placebo + h	pevacizumab 132)			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IV)	RS)							
NED [PDS]	92	42 (45.7)	NE (NE, NE)	48	27 (56.3)	19.4 (11.3,27.8)	0.72	0.44, 1.18	0.1839
NED/CR [IDS]	74	46 (62.2)	19.8 (14.8,22.5)	38	19 (50.0)	19.7 (13.7, NE)	1.21	0.72, 2.11	0.4885
NED/CR [Chemo]	40	22 (55.0)	13.8 (8.3, NE)	20	9 (45.0)	NE (NE, NE)	1.61	0.76, 3.68	0.2177
PR	49	23 (46.9)	19.5 (14.0, NE)	26	14 (53.8)	19.9 (14.0,24.9)	0.84	0.44, 1.67	0.6017
Interaction p-value									0.2597
Screening laboratory tBR0	CA statu	s (IVRS)							
tBRCAm	150	76 (50.7)	20.0 (16.3,25.3)	65	33 (50.8)	19.7 (13.7, NE)	0.99	0.67, 1.52	0.9781
non-tBRCAm	105	57 (54.3)	19.9 (13.9,24.2)	67	36 (53.7)	19.4 (13.9,24.9)	0.98	0.65, 1.50	0.9325
Interaction p-value									0.9670
First line treatment out	come (eCl	RF)							
NED [PDS]	89	41 (46.1)	NE (NE, NE)	47	26 (55.3)	19.4 (11.3,27.8)	0.78	0.48, 1.29	0.3285
NED/CR [IDS]	74	43 (58.1)	19.8 (15.9,24.0)	32	17 (53.1)	17.7 (10.8, NE)	0.97	0.57, 1.75	0.9223
NED/CR [Chemo]	39	20 (51.3)	19.2 (8.6, NE)	18	8 (44.4)	14.0 (6.0, NE)	1.22	0.56, 2.95	0.6276
PR	50	27 (54.0)	15.2 (13.8, NE)	34	18 (52.9)	22.2 (14.0, NE)	1.13	0.63, 2.08	0.6917
Interaction p-value									0.7305
Screening laboratory tBR0	CA statu	s (eCRF)							
tBRCAm	147	74 (50.3)	20.0 (16.3,25.3)	67	33 (49.3)	19.7 (13.7, NE)	1.01	0.68, 1.54	0.9603
non-tBRCAm	108	59 (54.6)	19.8 (14.0,24.2)	65	36 (55.4)	19.4 (13.9,24.9)	0.96	0.64, 1.47	0.8536
Interaction p-value									0.8679
Age group									
<65 years	185	99 (53.5)	19.1 (14.8,22.5)	98	52 (53.1)	19.7 (13.8,24.9)	1.04	0.75, 1.47	0.8048
>=65 years	70	34 (48.6)	22.6 (16.6, NE)	34	17 (50.0)	18.7 (12.3, NE)	0.81	0.46, 1.49	0.4959
Interaction p-value									0.4748

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aah 25NOV2020:12:08 khcs324

Table 2.2.3.8 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Constipation time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			pevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI [b	2-sided] p-value [b]
FIGO Stage (Disease state)									
III	182	103 (56.6)	17.3 (14.0,22.1)	90	47 (52.2)	19.4 (13.7,24.1)	1.06	0.75, 1.	50 0.7580
IV	73	30 (41.1)	24.9 (19.4, NE)	42	22 (52.4)	19.7 (13.8, NE)	0.77	0.45, 1.	35 0.3561
Interaction p-value									0.3431
Region									
Europe	245	128 (52.2)	19.8 (16.5,23.4)	126	65 (51.6)	19.7 (15.4,24.1)	1.01	0.75, 1.	
Japan	10	5 (50.0)	25.1 (5.6,25.1)	6	4 (66.7)	8.3 (2.8, NE)	0.53	0.14, 2.	15 0.3557
Interaction p-value									0.3624
ECOG performance status at	Basel	ine							
(0) Normal activity	190	100 (52.6)	19.8 (16.3,22.5)	100	51 (51.0)	22.1 (14.0,27.8)	1.09	0.78, 1.	0.6077
(1) Restricted activity	61	30 (49.2)	23.4 (13.8, NE)	31	18 (58.1)	15.4 (11.1,21.2)	0.63	0.35, 1.	15 0.1261
Interaction p-value									0.1123
Baseline CA-125 value									
<=ULN	228	118 (51.8)	20.0 (16.5,24.2)	118	63 (53.4)	19.4 (13.9,22.3)	0.92	0.68, 1.	25 0.5822
>ULN	27	15 (55.6)	19.9 (11.1,24.2)	14	6 (42.9)	22.1 (8.3, NE)	1.70	0.69, 4.	78 0.2543
Interaction p-value									0.2087
Histological grade									
High grade	255	133 (52.2)	19.9 (16.6,23.4)	132	69 (52.3)	19.7 (14.0,22.3)	0.98	0.73, 1.	31 0.8786
Interaction p-value									NC
Cytoreductive surgery outco	me								
No residue	166	86 (51.8)	20.0 (16.8,25.1)	80	43 (53.8)	19.4 (13.7,22.3)	0.88	0.61, 1.	28 0.5011
Residue	79	43 (54.4)	15.2 (13.8,24.2)	44	23 (52.3)	22.2 (13.9, NE)	1.19	0.72, 2.	00 0.5005
Interaction p-value									0.3464

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aah 25NOV2020:12:08 khcs324

Table 2.2.3.8 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Constipation time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	•	Olaparib + k (N=2				bevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	urgery									
Upfront	146	72 (49.3)	20.0 (14.5, NE)	79	41 (51.9)	19.7 (13.9,27.8)	0.96	0.66,	1.42	0.8331
Interval	99	57 (57.6)	19.8 (16.3,22.6)	45	25 (55.6)	19.4 (11.1, NE)	0.99	0.63,	1.61	0.9674
Interaction p-value										0.9190
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	80 (50.6)	19.5 (15.2,25.3)	77	37 (48.1)	21.2 (13.7, NE)	1.02	0.70,	1.53	0.9119
Non-tBRCAm	97	53 (54.6)	19.9 (14.5,24.2)	55	32 (58.2)	18.7 (13.8,24.1)	0.93	0.61,	1.46	0.7645
Interaction p-value										0.7658
Status somatic BRCA mutat	ions									
sBRCAm	22	10 (45.5)	15.2 (7.9, NE)	7	4 (57.1)	9.6 (2.8, NE)	0.73	0.24,	2.67	0.6046
gBRCAm	66	38 (57.6)	21.4 (14.5,25.1)	31	18 (58.1)	16.2 (11.3, NE)	0.91	0.53,	1.64	0.7568
Non-BRCAm	41	22 (53.7)	20.0 (13.8, NE)	22	14 (63.6)	13.8 (8.3, NE)	0.74	0.38,	1.48	0.3863
Interaction p-value										0.8729

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aah 25NOV2020:12:08 khcs324

Table 2.2.3.9 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Diarrhoea time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment outo	come (IVI	RS)								
NED [PDS]	92	41 (44.6)	25.5 (16.8, NE)	48	22 (45.8)	22.3 (14.0,35.0)	0.94	0.57,	1.61	0.8216
NED/CR [IDS]	74	32 (43.2)	NE (NE, NE)	38	17 (44.7)	22.1 (13.8, NE)	0.93	0.52,	1.71	0.8080
NED/CR [Chemo]	40	21 (52.5)	22.1 (8.3, NE)	20	8 (40.0)	25.2 (17.1, NE)	1.72	0.79,	4.14	0.1744
PR	49	30 (61.2)	14.2 (11.3,21.6)	26	11 (42.3)	23.5 (8.3, NE)	1.36	0.70,	2.84	0.3765
Interaction p-value										0.5236
Screening laboratory tBRO	CA status	s (IVRS)								
tBRCAm	150	74 (49.3)	24.0 (16.8,27.4)	65	29 (44.6)	25.2 (20.0,35.0)	1.13	0.75,	1.77	0.5671
non-tBRCAm	105	50 (47.6)	21.6 (11.5, NE)	67	29 (43.3)	22.1 (14.0, NE)	1.11	0.71,	1.78	0.6406
Interaction p-value										0.9611
First line treatment outo	come (eCI	RF)								
NED [PDS]	89	39 (43.8)	25.9 (16.6, NE)	47	20 (42.6)	35.0 (19.4,35.0)	1.02	0.60,	1.78	0.9448
NED/CR [IDS]	74	34 (45.9)	24.0 (13.0, NE)	32	15 (46.9)	22.1 (13.8, NE)	1.00	0.55,	1.89	0.9961
NED/CR [Chemo]	39	19 (48.7)	24.0 (8.7, NE)	18	7 (38.9)	25.2 (14.0, NE)	1.44	0.63,	3.69	0.3984
PR	50	30 (60.0)	15.3 (11.3,25.5)	34	15 (44.1)	23.5 (11.3, NE)	1.32	0.72,	2.53	0.3727
Interaction p-value										0.8377
Screening laboratory tBRO	CA status	s (eCRF)								
tBRCAm	147	72 (49.0)	24.0 (16.9,27.4)	67	29 (43.3)	25.2 (20.0,35.0)	1.14	0.75,	1.78	0.5531
non-tBRCAm	108	52 (48.1)	21.6 (11.4, NE)	65	29 (44.6)	22.1 (14.0, NE)	1.11	0.71,	1.77	0.6480
Interaction p-value										0.9406
Age group										
<65 years	185	86 (46.5)	25.3 (16.9,27.4)	98	40 (40.8)	35.0 (20.7,35.0)	1.15	0.80,	1.69	0.4605
>=65 years	70	38 (54.3)	16.6 (11.3,30.9)	34	18 (52.9)	19.9 (12.7,22.2)	1.03	0.59,	1.84	0.9316
Interaction p-value						<u> </u>				0.7381

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aai 25NoV2020:12:08 khcs324

Table 2.2.3.9 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Diarrhoea time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [[b] p-	2-sided -value [b]
FIGO Stage (Disease state)										
III	182	91 (50.0)	23.3 (15.9,25.9)	90	40 (44.4)	22.2 (19.8,35.0)	1.16	0.81, 1	L.70	0.4296
IV	73	33 (45.2)	25.0 (11.4, NE)	42	18 (42.9)	25.2 (13.9, NE)	1.02	0.58, 1	1.85	0.9529
Interaction p-value										0.7081
Region										
Europe	245	119 (48.6)	24.0 (16.6,25.9)	126	54 (42.9)	25.2 (19.8,35.0)	1.13	0.82, 1	L.57	0.4584
Japan	10	5 (50.0)	16.6 (2.8, NE)	6	4 (66.7)	22.1 (5.5, NE)	1.01	0.27, 4	1.08	0.9893
Interaction p-value										0.8716
ECOG performance status at	Basel	ine								
(0) Normal activity	190	88 (46.3)	24.0 (16.8, NE)	100	44 (44.0)	25.2 (20.7,35.0)	1.07	0.75, 1	L.55	0.7142
(1) Restricted activity	61	35 (57.4)	14.5 (8.7,25.7)	31	14 (45.2)	19.4 (12.3, NE)	1.32	0.72, 2	2.53	0.3741
Interaction p-value										0.5665
Baseline CA-125 value										
<=ULN	228		25.5 (21.6,30.9)	118		25.2 (19.8,35.0)	1.04	0.75, 1		0.8282
>ULN	27	20 (74.1)	11.1 (5.7,16.6)	14	7 (50.0)	20.7 (5.7, NE)	1.96	0.87, 5	5.01	0.1079
Interaction p-value										0.1626
Histological grade										
High grade	255	124 (48.6)	24.0 (16.6,25.9)	132	58 (43.9)	23.5 (19.9,35.0)	1.12	0.82, 1	1.54	0.4826
Interaction p-value										NC
Cytoreductive surgery outco	me									
No residue	166	75 (45.2)	25.9 (16.8, NE)	80	36 (45.0)	22.2 (19.4,35.0)	1.00	0.68, 1	1.51	0.9954
Residue	79	43 (54.4)	21.9 (11.3,25.7)	44	17 (38.6)	25.2 (17.1, NE)	1.46	0.85, 2	2.64	0.1759
Interaction p-value										0.2785

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aai 25NoV2020:12:08 khcs324

Table 2.2.3.9 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Diarrhoea time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2		P.	lacebo + k (N=1	pevacizumab 132)			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	of	umber (%) patients th events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	urgery								
Upfront	146	71 (48.6)	24.0 (16.6,25.9)	79	32 (40.5)	25.2 (20.0,35.0)	1.23	0.82, 1.89	0.3246
Interval	99	47 (47.5)	24.0 (13.4, NE)	45	21 (46.7)	22.2 (13.9, NE)	1.00	0.61, 1.71	NC
Interaction p-value									0.5412
Myriad tumour BRCA mutati	on stat	us							
tBRCAm	158	74 (46.8)	25.3 (18.0,30.9)	77	32 (41.6)	25.2 (20.0,35.0)	1.08	0.72, 1.66	0.7041
Non-tBRCAm	97	50 (51.5)	15.3 (11.1, NE)	55	26 (47.3)	22.1 (14.0, NE)	1.20	0.76, 1.96	0.4420
Interaction p-value									0.7450
Status somatic BRCA mutat	ions								
sBRCAm	22	10 (45.5)	21.0 (5.6, NE)	7	2 (28.6)	NE (NE, NE)	2.18	0.57, 14.22	0.2788
gBRCAm	66	31 (47.0)	25.3 (14.5, NE)	31	13 (41.9)	23.5 (19.4,35.0)	1.29	0.69, 2.56	0.4346
Non-BRCAm	41	24 (58.5)	14.0 (5.7, NE)	22	13 (59.1)	19.4 (12.7, NE)	1.11	0.58, 2.25	0.7533
Interaction p-value									0.7106

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aai 25NoV2020:12:08 khcs324

Table 2.2.3.10 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Dyspnoea time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI) (months) [)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment outo	come (IV	RS)									
NED [PDS]	92	39 (42.4)	52.5 (17.9,52.5)	48	26 (54.2)	16.7 (8.5,	NE)	0.63	0.39,	1.06	0.0797
NED/CR [IDS]	74	41 (55.4)	17.3 (11.5,24.2)	38	19 (50.0)	18.9 (11.1,	NE)	1.15	0.67,	2.02	0.6226
NED/CR [Chemo]	40	20 (50.0)	22.1 (8.2, NE)	20	8 (40.0)	NE (NE,	NE)	1.59	0.73,	3.84	0.2522
PR	49	25 (51.0)	15.4 (11.3, NE)	26	14 (53.8)	11.1 (2.9,	NE)	0.71	0.37,	1.40	0.3087
Interaction p-value											0.1644
Screening laboratory tBRO	CA statu	s (IVRS)									
tBRCAm	150	67 (44.7)	31.0 (17.3, NE)	65	34 (52.3)	17.0 (11.1,	NE)	0.76	0.51,	1.16	0.1967
non-tBRCAm	105	58 (55.2)	15.4 (11.3,23.4)	67	33 (49.3)	18.9 (11.1,	NE)	1.12	0.73,	1.73	0.6156
Interaction p-value											0.2043
First line treatment outo	come (eC	RF)									
NED [PDS]	89	37 (41.6)	52.5 (18.0,52.5)	47	24 (51.1)	17.0 (8.6,	NE)	0.68	0.41,	1.15	0.1444
NED/CR [IDS]	74	41 (55.4)	14.8 (8.3,31.0)	32	17 (53.1)	18.9 (8.5,	NE)	1.13	0.65,	2.04	0.6772
NED/CR [Chemo]	39	21 (53.8)	15.2 (4.5, NE)	18	8 (44.4)	11.1 (5.6,	NE)	1.29	0.59,	3.09	0.5383
PR	50	24 (48.0)	22.3 (12.1, NE)	34	17 (50.0)	19.7 (9.0,	NE)	0.82	0.45,	1.56	0.5463
Interaction p-value											0.4539
Screening laboratory tBRO	CA statu	s (eCRF)									
tBRCAm	147	64 (43.5)	31.0 (18.0, NE)	67	35 (52.2)	17.0 (11.1,	NE)	0.72	0.48,	1.10	0.1250
non-tBRCAm	108	61 (56.5)	15.3 (11.2,22.1)	65	32 (49.2)	18.9 (11.1,	NE)	1.18	0.77,	1.83	0.4562
Interaction p-value											0.1061
Age group											
<65 years	185	89 (48.1)	22.1 (16.6, NE)	98	49 (50.0)	19.7 (11.3,	NE)	0.90	0.64,	1.28	0.5527
>=65 years	70	36 (51.4)	19.7 (11.3,52.5)	34	18 (52.9)	13.8 (8.4,	NE)	0.87	0.50,	1.57	0.6407
Interaction p-value											0.9293

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aaj 25NoV2020:12:08 khcs324

Table 2.2.3.10 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Dyspnoea time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)									
III	182	85 (46.7)	24.0 (17.9,52.5)	90	47 (52.2)	16.7 (11.1, NE)	0.81	0.57, 1.17	0.2532
IV	73	40 (54.8)	14.8 (8.3, NE)	42	20 (47.6)	21.2 (13.8, NE)	1.13	0.67, 1.98	0.6425
Interaction p-value									0.3026
Region									
Europe	245	121 (49.4)	20.7 (16.0,52.5)	126	64 (50.8)	18.7 (12.3,24.9)	0.90	0.66, 1.22	0.4878
Japan	10	4 (40.0)	NE (NE, NE)	6	3 (50.0)	NE (NE, NE)	0.81	0.18, 4.10	0.7814
Interaction p-value									0.8931
ECOG performance status at	Basel	ine							
(0) Normal activity	190	96 (50.5)	19.4 (15.2,52.5)	100	53 (53.0)	18.7 (11.1, NE)	0.92	0.66, 1.30	0.6406
(1) Restricted activity	61	27 (44.3)	NE (NE, NE)	31	14 (45.2)	18.9 (8.5, NE)	0.82	0.44, 1.61	0.5549
Interaction p-value									0.7549
Baseline CA-125 value									
<=ULN	228	110 (48.2)	22.3 (16.0,52.5)	118	58 (49.2)	19.7 (11.1, NE)	0.90	0.66, 1.25	0.5363
>ULN	27	15 (55.6)	19.7 (5.6, NE)	14	9 (64.3)	13.8 (8.4,21.2)	0.85	0.38, 2.01	0.6936
Interaction p-value									0.8839
Histological grade									
High grade	255	125 (49.0)	20.7 (16.0,52.5)	132	67 (50.8)	18.7 (12.3,24.9)	0.89	0.67, 1.21	0.4678
Interaction p-value									NC
Cytoreductive surgery outco	ome								
No residue	166	80 (48.2)	20.7 (16.6,52.5)	80	42 (52.5)	17.0 (12.1,24.9)	0.85	0.59, 1.24	0.3948
Residue	79	38 (48.1)	23.4 (12.1, NE)	44	21 (47.7)	19.7 (8.7, NE)	0.97	0.57, 1.68	0.8997
Interaction p-value									0.6959

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aaj 25NoV2020:12:08 khcs324

Table 2.2.3.10 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Dyspnoea time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2				oevacizumab 132)					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events)	Hazard ratio [b]	95% C	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	66 (45.2)	52.5 (16.7,52.5)	79	40 (50.6)	18.7 (8.7,	NE)	0.79	0.53,	1.17	0.2355
Interval	99	52 (52.5)	19.9 (12.5,31.0)	45	23 (51.1)	18.9 (11.1,	NE)	1.06	0.66,	1.76	0.8160
Interaction p-value											0.3489
Myriad tumour BRCA mutati	ion stat	us									
tBRCAm	158	73 (46.2)	24.2 (16.6, NE)	77	38 (49.4)	19.7 (12.1,	NE)	0.85	0.58,	1.27	0.4165
Non-tBRCAm	97	52 (53.6)	17.9 (12.5,24.2)	55	29 (52.7)	13.8 (8.5,	NE)	0.98	0.62,	1.56	0.9207
Interaction p-value											0.6458
Status somatic BRCA mutat	cions										
sBRCAm	22	7 (31.8)	NE (NE, NE)	7	4 (57.1)	19.5 (2.8,	NE)	0.54	0.16,	2.05	0.3379
gBRCAm	66	33 (50.0)	17.3 (11.3, NE)	31	17 (54.8)	16.7 (3.0,	NE)	0.80	0.45,	1.47	0.4661
Non-BRCAm	41	21 (51.2)	23.4 (14.8,52.5)	22	10 (45.5)	NE (NE,	NE)	1.09	0.52,	2.43	0.8225
Interaction p-value											0.6141

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aaj 25NoV2020:12:08 khcs324

Table 2.2.3.11 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Fatigue time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k			Placebo + k (N=1	pevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IV	TRS)							
NED [PDS]	92	68 (73.9)	5.6 (2.9,11.1)	48	36 (75.0)	5.7 (3.0, 8.5)	0.95	0.64, 1.43	0.7942
NED/CR [IDS]	74	58 (78.4)	5.4 (2.9, 8.3)	38	26 (68.4)	11.1 (5.5,16.8)	1.49	0.95, 2.41	0.0829
NED/CR [Chemo]	40	33 (82.5)	3.9 (2.9, 5.8)	20	15 (75.0)	9.8 (3.0,22.3)	1.61	0.89, 3.05	0.1171
PR	49	40 (81.6)	8.6 (3.0,11.3)	26	21 (80.8)	2.9 (2.8,13.8)	0.64	0.38, 1.11	0.1093
Interaction p-value									0.0569
Screening laboratory tBRO	CA statı	ıs (IVRS)							
tBRCAm	150	122 (81.3)	5.6 (3.0, 8.4)	65	50 (76.9)	5.7 (3.0, 9.3)	1.11	0.80, 1.55	0.5437
non-tBRCAm	105	77 (73.3)	5.6 (3.0, 8.3)	67	48 (71.6)	8.1 (3.5,13.9)	1.10	0.77, 1.59	0.5939
Interaction p-value									0.9885
First line treatment outo	come (e0	CRF)							
NED [PDS]	89	65 (73.0)	5.6 (2.9, 8.4)	47	36 (76.6)	5.6 (3.0, 8.2)	0.89	0.60, 1.35	0.5792
NED/CR [IDS]	74	56 (75.7)	3.8 (2.8, 8.3)	32	22 (68.8)	10.2 (3.0,16.9)	1.45	0.90, 2.42	0.1311
NED/CR [Chemo]	39	32 (82.1)	5.2 (2.9, 8.6)	18	12 (66.7)	11.2 (3.0,22.3)	1.76	0.93, 3.57	0.0818
PR	50	43 (86.0)	8.5 (3.0,11.1)	34	28 (82.4)	5.5 (2.8,13.8)	0.83	0.52, 1.34	0.4354
Interaction p-value									0.1284
Screening laboratory tBRO	CA statı	ıs (eCRF)							
tBRCAm	147	119 (81.0)	5.6 (3.0, 8.6)	67	50 (74.6)	5.7 (3.0,11.1)	1.12	0.81, 1.58	0.4830
non-tBRCAm	108	80 (74.1)	5.5 (3.0, 8.1)	65	48 (73.8)	7.9 (3.0,12.5)	1.09	0.77, 1.57	0.6264
Interaction p-value									0.9078
Age group									
<65 years	185	147 (79.5)	5.6 (3.0, 5.8)	98	71 (72.4)	5.7 (3.0, 8.5)	1.14	0.87, 1.53	0.3465
>=65 years	70	52 (74.3)	5.8 (3.0, 8.7)	34	27 (79.4)	11.1 (5.7,14.0)	1.03	0.65, 1.66	0.9081
Interaction p-value									0.6995

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aak 25NOV2020:12:08 khcs324

Table 2.2.3.11 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Fatigue time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2				oevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI []	2-sided b] p-value[b]
FIGO Stage (Disease state)									
III	182	138 (75.8)	5.6 (3.2, 8.3)	90	66 (73.3)	6.0 (3.0,11.1)	1.09	0.82, 1	.47 0.5492
IV	73	61 (83.6)	4.2 (2.9, 8.4)	42	32 (76.2)	5.7 (3.0,13.9)	1.18	0.77, 1	.83 0.4529
Interaction p-value									0.7814
Region									
Europe	245	194 (79.2)	5.6 (3.1, 5.8)	126	94 (74.6)	5.7 (5.5, 9.3)	1.13	0.89, 1	.45 0.3285
Japan	10	5 (50.0)	8.3 (2.8, NE)	6	4 (66.7)	8.4 (2.8, NE)	0.69	0.18, 2	.79 0.5834
Interaction p-value									0.4753
ECOG performance status at	Basel	ine							
(0) Normal activity	190	153 (80.5)	5.4 (3.0, 5.7)	100	77 (77.0)	5.7 (5.6, 8.5)	1.22	0.93, 1	.62 0.1439
(1) Restricted activity	61	43 (70.5)	8.5 (3.0,22.0)	31	21 (67.7)	11.2 (2.9,16.6)	0.88	0.53, 1	.52 0.6370
Interaction p-value									0.2810
Baseline CA-125 value									
<=ULN	228		5.6 (3.1, 7.9)	118		5.7 (5.0,11.1)	1.12	0.87, 1	
>ULN	27	23 (85.2)	3.0 (2.9,11.1)	14	12 (85.7)	8.5 (2.8,11.3)	1.03	0.52, 2	
Interaction p-value									0.8274
Histological grade									
High grade	255	199 (78.0)	5.6 (3.1, 6.0)	132	98 (74.2)	5.7 (5.5,11.1)	1.11	0.88, 1	.42 0.3865
Interaction p-value									NC
Cytoreductive surgery outco	ome								
No residue	166	124 (74.7)	5.1 (3.0, 5.7)	80	58 (72.5)	5.8 (5.0,11.1)	1.13	0.83, 1	.55 0.4514
Residue	79	67 (84.8)	5.8 (3.0,11.0)	44	33 (75.0)	8.5 (3.0,14.0)	1.20	0.79, 1	.84 0.3974
Interaction p-value									0.8231

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aak 25NOV2020:12:08 khcs324

Table 2.2.3.11 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Fatigue time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			oevacizumab 132)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	,	Hazard ratio [b]	95% C	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	114 (78.1)	5.6 (3.1, 8.4)	79	59 (74.7)	5.7 (3.0, 8.5)	1.01	0.74,	1.39	0.9691
Interval	99	77 (77.8)	4.8 (2.9, 8.3)	45	32 (71.1)	11.1 (5.6,14.9)	1.43	0.96,	2.19	0.0832
Interaction p-value										0.1827
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	127 (80.4)	5.6 (3.1, 8.4)	77	55 (71.4)	5.7 (3.2,11.1)	1.16	0.85,	1.61	0.3426
Non-tBRCAm	97	72 (74.2)	5.4 (3.0, 5.8)	55	43 (78.2)	7.9 (3.0,12.5)	1.05	0.72,	1.54	0.8163
Interaction p-value										0.6706
Status somatic BRCA mutat	ions									
sBRCAm	22	14 (63.6)	11.0 (5.6,22.1)	7	4 (57.1)	10.5 (2.8, NE)	1.03	0.37,	3.63	0.9621
gBRCAm	66	55 (83.3)	4.3 (2.9,11.1)	31	24 (77.4)	5.7 (2.9,13.9)	1.13	0.71,	1.85	0.6228
Non-BRCAm	41	29 (70.7)	5.4 (2.8,11.3)	22	19 (86.4)	5.7 (2.8,13.9)	0.84	0.47,	1.52	0.5485
Interaction p-value										0.7388

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aak 25NOV2020:12:08 khcs324

Table 2.2.3.12 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Financial difficulties time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2					bevacizumab 132)					
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)										
NED [PDS]	92	24 (26.1)	NE (NE,	NE)	48	17 (35.4)	NE (NE,	NE)	0.59	0.32,	1.11	0.1007
NED/CR [IDS]	74	24 (32.4)	NE (NE,	NE)	38	9 (23.7)	NE (NE,	NE)	1.36	0.65,	3.09	0.4205
NED/CR [Chemo]	40	14 (35.0)	NE (NE,	NE)	20	11 (55.0)	13.9 (5.7,	NE)	0.63	0.28,	1.41	0.2517
PR	49	15 (30.6)	38.4 (22.3,	NE)	26	11 (42.3)	19.2 (4.7,	NE)	0.52	0.24,	1.15	0.1046
Interaction p-value												0.2519
Screening laboratory tBR	CA statu	s (IVRS)										
tBRCAm	150	50 (33.3)	38.4 (38.4,	NE)	65	25 (38.5)	NE (NE,	NE)	0.79	0.49,	1.30	0.3424
non-tBRCAm	105	27 (25.7)	NE (NE,	NE)	67	23 (34.3)	NE (NE,	NE)	0.61	0.35,	1.08	0.0872
Interaction p-value												0.4950
First line treatment out	come (eC	RF)										
NED [PDS]	89	25 (28.1)	NE (NE,	NE)	47	16 (34.0)	NE (NE,	NE)	0.68	0.37,	1.30	0.2407
NED/CR [IDS]	74	24 (32.4)	NE (NE,	NE)	32	7 (21.9)	NE (NE,	NE)	1.49	0.68,	3.75	0.3352
NED/CR [Chemo]	39	10 (25.6)	NE (NE,	NE)	18	10 (55.6)	13.3 (5.7,	NE)	0.38	0.15,	0.92	0.0326*
PR	50	17 (34.0)	38.4 (22.3,	NE)	34	15 (44.1)	19.4 (8.8,	NE)	0.62	0.31,	1.27	0.1869
Interaction p-value												0.1401
Screening laboratory tBR	CA statu	s (eCRF)										
tBRCAm	147	50 (34.0)	38.4 (38.4,	NE)	67	25 (37.3)	NE (NE,	NE)	0.83	0.52,	1.36	0.4409
non-tBRCAm	108	27 (25.0)	NE (NE,	NE)	65	23 (35.4)	NE (NE,	NE)	0.58	0.33,	1.02	0.0563
Interaction p-value												0.3381
Age group												
<65 years	185	58 (31.4)	NE (NE,	NE)	98	38 (38.8)	NE (NE,	NE)	0.72	0.48,	1.10	0.1269
>=65 years	70	19 (27.1)	38.4 (38.4,	NE)	34	10 (29.4)	NE (NE,	NE)	0.72	0.34,	1.62	0.4114
Interaction p-value												0.9903

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aal 25NoV2020:12:08 khcs324

Table 2.2.3.12 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Financial difficulties time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)				bevacizumab 132)					
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events)	Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	58 (31.9)	NE (NE,	NE)	90	32 (35.6)	NE (NE,	NE)	0.77	0.50,	1.20	0.2466
IV	73	19 (26.0)	38.4 (38.4,	NE)	42	16 (38.1)	NE (NE,	NE)	0.59	0.30,	1.16	0.1236
Interaction p-value												0.5027
Region												
Europe	245	74 (30.2)	38.4 (38.4,	NE)	126	47 (37.3)	NE (NE,	NE)	0.68	0.48,	0.99	0.0450*
Japan	10	3 (30.0)	NE (NE,	NE)	6	1 (16.7)	NE (NE,	NE)	2.31	0.30, 4	16.77	0.4406
Interaction p-value												0.2612
ECOG performance status at	Basel	ine										
(0) Normal activity	190	58 (30.5)	38.4 (38.4,	NE)	100	38 (38.0)	NE (NE,	NE)	0.72	0.48,	1.09	0.1191
(1) Restricted activity	61	17 (27.9)	NE (NE,	NE)	31	10 (32.3)	NE (NE,	NE)	0.67	0.31,	1.51	0.3186
Interaction p-value												0.8661
Baseline CA-125 value												
<=ULN	228		38.4 (38.4,		118	42 (35.6)		NE)	0.72	0.49,		0.1036
>ULN	27	8 (29.6)	NE (NE,	NE)	14	6 (42.9)	21.2 (11.3,	NE)	0.68	0.24,	2.07	0.4808
Interaction p-value												0.9153
Histological grade												
High grade	255	77 (30.2)	38.4 (38.4,	NE)	132	48 (36.4)	NE (NE,	NE)	0.72	0.50,	1.03	0.0746
Interaction p-value												NC
Cytoreductive surgery outco	me											
No residue	166	50 (30.1)	NE (NE,	NE)	80	23 (28.8)	NE (NE,	NE)	0.94	0.58,	1.57	0.8038
Residue	79	24 (30.4)	38.4 (38.4,	NE)	44	21 (47.7)	19.4 (11.3,	NE)	0.53	0.30,	0.97	0.0399*
Interaction p-value												0.1497

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aal 25NoV2020:12:08 khcs324

Table 2.2.3.12 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Financial difficulties time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		-	bevacizumab 255)			Placebo + bo (N=1		ab				
Subgroup		Number (%) of patients with events)		Number (%) of patients with events	(95%	n time CI) s) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	urgery											
Upfront	146	40 (27.4)	NE (NE,	NE)	79	32 (40.5)	NE (NE, NE)	0.54	0.34,	0.86	0.0107*
Interval	99	34 (34.3)	NE (NE,	NE)	45	12 (26.7)	NE (NE, NE)	1.31	0.70,	2.64	0.4063
Interaction p-value												0.0264*
Myriad tumour BRCA mutati	on stat	us										
tBRCAm	158	54 (34.2)	38.4 (38.4,	NE)	77	31 (40.3)	NE (NE, NE)	0.72	0.47,	1.14	0.1617
Non-tBRCAm	97	23 (23.7)	NE (NE,	NE)	55	17 (30.9)	NE (NE, NE)	0.67	0.36,	1.28	0.2187
Interaction p-value												0.8434
Status somatic BRCA mutat	ions											
sBRCAm	22	7 (31.8)	38.4 (11.1,	NE)	7	3 (42.9)	NE (NE, NE)	0.63	0.17,	2.95	0.5192
gBRCAm	66	22 (33.3)	NE (NE,	NE)	31	9 (29.0)	NE (NE, NE)	1.11	0.53,	2.55	0.7828
Non-BRCAm	41	10 (24.4)	NE (NE,	NE)	22	7 (31.8)	NE (NE, NE)	0.72	0.28,	1.98	0.5072
Interaction p-value												0.6861

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aal 25NoV2020:12:08 khcs324

Table 2.2.3.13 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Nausea and vomiting time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		bevacizumab 255)			pevacizumab 132)				
Subgroup	Number (%) of patients n with events		(Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment outc	ome (IVRS)								
NED [PDS]	92 66 (71.7)	5.7 (3.0,11.1)	48	26 (54.2)	16.9 (9.7, NE)	1.75	1.12,	2.80	0.0125*
NED/CR [IDS]	74 54 (73.0)	5.6 (2.9, 8.7)	38	20 (52.6)	19.2 (11.1, NE)	2.05	1.25,	3.51	0.0042*
NED/CR [Chemo]	40 27 (67.5)	5.7 (3.1,13.8)	20	11 (55.0)	19.3 (8.3, NE)	1.87	0.95,	3.94	0.0694
PR	49 31 (63.3)	11.1 (3.1,16.9)	26	13 (50.0)	19.9 (9.7,27.0)	1.53	0.82,	3.02	0.1899
Interaction p-value									0.9149
Screening laboratory tBRC	A status (IVRS)								
tBRCAm	150 104 (69.3)	5.8 (3.5,11.0)	65	34 (52.3)	19.7 (11.4,27.0)	1.90	1.31,	2.84	0.0006*
non-tBRCAm	105 74 (70.5)	5.7 (5.4,11.1)	67	36 (53.7)	17.4 (9.7,25.0)	1.72	1.16,	2.59	0.0062*
Interaction p-value									0.7212
First line treatment outc	ome (eCRF)								
NED [PDS]	89 65 (73.0)	5.6 (2.9,11.0)	47	25 (53.2)	17.4 (9.7, NE)	1.90	1.21,	3.06	0.0047*
NED/CR [IDS]	74 53 (71.6)	5.6 (2.9, 8.7)	32	19 (59.4)	16.2 (9.6,24.4)	1.74	1.05,	3.01	0.0322*
NED/CR [Chemo]	39 25 (64.1)	7.1 (4.5,16.8)	18	9 (50.0)	19.3 (11.1, NE)	1.84	0.89,	4.16	0.1034
PR	50 32 (64.0)	11.0 (3.0,19.4)	34	17 (50.0)	19.9 (9.7, NE)	1.57	0.88,	2.89	0.1256
Interaction p-value									0.9678
Screening laboratory tBRC	A status (eCRF)								
tBRCAm	147 102 (69.4)	5.8 (3.4,11.0)	67	34 (50.7)	20.7 (14.3, NE)	1.95	1.34,	2.91	0.0004*
non-tBRCAm	108 76 (70.4)	5.7 (5.5,11.1)	65	36 (55.4)	16.6 (9.7,24.4)	1.67	1.13,	2.51	0.0093*
Interaction p-value									0.5883
Age group									
<65 years	185 133 (71.9)	5.7 (3.2, 8.4)	98	54 (55.1)	19.2 (11.2,24.4)	1.81	1.32,	2.50	0.0002*
>=65 years	70 45 (64.3)	11.1 (5.6,14.1)	34	16 (47.1)	19.9 (11.3, NE)	1.83	1.06,	3.34	0.0303*
Interaction p-value									0.9661

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aam 25NOV2020:12:08 khcs324

Table 2.2.3.13 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Nausea and vomiting time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				pevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)									
III	182	125 (68.7)	6.2 (5.6,11.1)	90	46 (51.1)	19.7 (12.7,25.0)	1.87	1.35, 2.65	0.0002*
IV	73	53 (72.6)	5.6 (3.0,11.1)	42	24 (57.1)	16.6 (8.3,27.0)	1.69	1.05, 2.78	0.0294*
Interaction p-value									0.7267
Region									
Europe	245	171 (69.8)	5.8 (5.5, 8.7)	126	66 (52.4)	18.7 (11.4,23.5)	1.80	1.36, 2.41	<0.0001*
Japan	10	7 (70.0)	5.7 (2.8, NE)	6	4 (66.7)	23.6 (8.1,25.0)	1.83	0.55, 7.00	0.3252
Interaction p-value									0.9780
ECOG performance status at	Basel	ine							
(0) Normal activity	190	140 (73.7)	5.7 (4.2, 8.3)	100	55 (55.0)	18.7 (11.3,25.0)	2.02	1.49, 2.79	<0.0001*
(1) Restricted activity	61	36 (59.0)	10.1 (4.5,25.7)	31	15 (48.4)	19.7 (11.1, NE)	1.33	0.74, 2.51	0.3406
Interaction p-value									0.2385
Baseline CA-125 value									
<=ULN	228		5.7 (5.5, 8.6)	118		18.7 (12.3,22.3)	1.74	1.31, 2.34	<0.0001*
>ULN	27	17 (63.0)	11.1 (3.0,20.0)	14	5 (35.7)	NE (NE, NE)	2.55	1.01, 7.76	0.0482*
Interaction p-value									0.4614
Histological grade									
High grade	255	178 (69.8)	5.8 (5.6, 8.7)	132	70 (53.0)	19.2 (12.7,23.5)	1.80	1.37, 2.39	<0.0001*
Interaction p-value									NC
Cytoreductive surgery outco	ome								
No residue	166	, ,	5.6 (3.1, 8.4)	80		17.4 (11.1,22.1)	1.85	1.32, 2.65	0.0003*
Residue	79	52 (65.8)	8.3 (5.6,12.1)	44	23 (52.3)	18.7 (11.1, NE)	1.62	1.005, 2.70	0.0478*
Interaction p-value									0.6613

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aam 25NOV2020:12:08 khcs324

Table 2.2.3.13 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Nausea and vomiting time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			bevacizumab 132)				
Subgroup	n	Number (%) of patients with events	. ,		Number (%) of patients with events	,	Hazard ratio [b]	95% C	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	105 (71.9)	5.8 (4.5,11.1)	79	41 (51.9)	18.7 (11.2, NE)	1.82	1.28,	2.64	0.0008*
Interval	99	68 (68.7)	5.7 (3.1,11.1)	45	26 (57.8)	16.2 (11.1,24.4)	1.72	1.11,	2.75	0.0152*
Interaction p-value										0.8490
Myriad tumour BRCA mutati	ion stat	cus								
tBRCAm	158	109 (69.0)	5.8 (3.5,11.0)	77	37 (48.1)	22.1 (16.6, NE)	2.05	1.43,	3.02	<0.0001*
Non-tBRCAm	97	69 (71.1)	5.7 (5.4,11.1)	55	33 (60.0)	12.7 (8.5,19.9)	1.52	1.01,	2.33	0.0424*
Interaction p-value										0.2939
Status somatic BRCA mutat	ions									
sBRCAm	22	11 (50.0)	13.8 (2.8, NE)	7	4 (57.1)	22.6 (2.8, NE)	1.02	0.35,	3.69	0.9713
gBRCAm	66	48 (72.7)	3.8 (2.9, 6.2)	31	18 (58.1)	19.2 (11.3,27.0)	1.90	1.13,	3.36	0.0156*
Non-BRCAm	41	33 (80.5)	5.6 (2.8, 8.3)	22	14 (63.6)	12.3 (8.1, NE)	1.91	1.04,	3.70	0.0352*
Interaction p-value										0.6275

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aam 25NOV2020:12:08 khcs324

Table 2.2.3.14 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Pain time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	·	Olaparib + k (N=2			Placebo + b (N=1				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IV	RS)							
NED [PDS]	92	66 (71.7)	8.3 (5.6,11.2)	48	33 (68.8)	3.1 (2.9,11.5)	0.86	0.57, 1.33	0.4991
NED/CR [IDS]	74	58 (78.4)	5.6 (2.9, 5.9)	38	28 (73.7)	5.6 (2.9, 7.6)	1.12	0.72, 1.78	0.6241
NED/CR [Chemo]	40	29 (72.5)	5.6 (3.0,11.2)	20	13 (65.0)	7.3 (3.0, NE)	1.34	0.71, 2.67	0.3711
PR	49	30 (61.2)	8.3 (5.5,19.5)	26	21 (80.8)	5.6 (2.8,12.9)	0.55	0.31, 0.97	0.0382*
Interaction p-value									0.1467
Screening laboratory tBRO	CA statu	ıs (IVRS)							
tBRCAm	150	105 (70.0)	5.8 (4.2, 8.4)	65	45 (69.2)	6.0 (4.9,13.8)	1.07	0.76, 1.53	0.7137
non-tBRCAm	105	78 (74.3)	5.9 (5.6, 8.4)	67	50 (74.6)	3.0 (2.9, 6.0)	0.77	0.55, 1.11	0.1652
Interaction p-value									0.2083
First line treatment outo	come (eC	RF)							
NED [PDS]	89	66 (74.2)	8.1 (5.3,11.1)	47	33 (70.2)	3.0 (2.9,11.2)	0.90	0.60, 1.38	0.6175
NED/CR [IDS]	74	59 (79.7)	5.5 (2.8, 5.9)	32	24 (75.0)	5.7 (2.9, 8.1)	1.23	0.77, 2.01	0.3920
NED/CR [Chemo]	39	25 (64.1)	8.5 (3.0,15.2)	18	11 (61.1)	8.7 (3.5, NE)	1.05	0.53, 2.22	0.8981
PR	50	32 (64.0)	8.3 (5.5,18.7)	34	26 (76.5)	5.6 (2.9,12.9)	0.65	0.39, 1.10	0.1050
Interaction p-value									0.3445
Screening laboratory tBRO	CA statu	ıs (eCRF)							
tBRCAm	147	102 (69.4)	5.8 (4.2, 8.4)	67	46 (68.7)	5.9 (4.9,13.8)	1.05	0.74, 1.50	0.7960
non-tBRCAm	108	81 (75.0)	5.9 (5.6, 8.4)	65	49 (75.4)	3.0 (2.9, 8.1)	0.79	0.55, 1.13	0.1890
Interaction p-value									0.2586
Age group									
<65 years	185	138 (74.6)	5.8 (4.9, 8.3)	98	70 (71.4)	5.6 (3.0, 8.7)	0.97	0.73, 1.31	0.8627
>=65 years	70	45 (64.3)	5.8 (5.6,11.1)	34	25 (73.5)	5.6 (2.9,11.1)	0.75	0.47, 1.25	0.2664
Interaction p-value									0.3795

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aan 25NoV2020:12:08 khcs324

Table 2.2.3.14 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Pain time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2				oevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)									
III	182	127 (69.8)	7.8 (5.6, 8.5)	90	67 (74.4)	5.6 (3.0, 7.6)	0.81	0.60, 1.0	9 0.1571
IV	73	56 (76.7)	5.6 (3.0, 8.3)	42	28 (66.7)	5.6 (3.0,13.9)	1.23	0.79, 1.9	7 0.3571
Interaction p-value									0.1194
Region									
Europe	245	179 (73.1)	5.8 (5.6, 8.3)	126	91 (72.2)	5.6 (3.0, 7.6)	0.93	0.72, 1.2	0.5599
Japan	10	4 (40.0)	NE (NE, NE)	6	4 (66.7)	13.5 (2.8, NE)	0.51	0.12, 2.1	0.3469
Interaction p-value									0.4102
ECOG performance status at	Basel	ine							
(0) Normal activity	190	140 (73.7)	5.7 (5.6, 8.3)	100	73 (73.0)	5.6 (3.0, 7.6)	0.94	0.71, 1.2	0.6922
(1) Restricted activity	61	40 (65.6)	8.3 (3.0,14.5)	31	22 (71.0)	5.6 (3.0,16.9)	0.85	0.51, 1.4	5 0.5429
Interaction p-value									0.7276
Baseline CA-125 value									
<=ULN	228		6.2 (5.6, 8.4)	118		5.6 (3.0, 8.1)	0.92	0.71, 1.2	
>ULN	27	17 (63.0)	5.6 (3.0, NE)	14	9 (64.3)	6.6 (2.8, NE)	0.85	0.39, 2.0	
Interaction p-value									0.8553
Histological grade									
High grade	255	183 (71.8)	5.8 (5.6, 8.3)	132	95 (72.0)	5.6 (3.0, 8.1)	0.91	0.71, 1.1	.7 0.4717
Interaction p-value									NC
Cytoreductive surgery outco	ome								
No residue	166	126 (75.9)	5.7 (4.2, 8.3)	80	58 (72.5)	5.5 (3.0, 7.6)	0.99	0.73, 1.3	0.9652
Residue	79	50 (63.3)	8.5 (5.7,15.2)	44	30 (68.2)	6.0 (3.0,11.2)	0.78	0.50, 1.2	0.2779
Interaction p-value									0.3806

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aan 25NoV2020:12:08 khcs324

Table 2.2.3.14 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Pain time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

_	:	Olaparib + N=2			Placebo + b (N=1				
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]	Number (%) of patients n with events		Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery								
Upfront	146	102 (69.9)	8.4 (5.7,11.2)	79	53 (67.1)	5.7 (3.0,11.1)	0.90	0.65, 1.27	0.5516
Interval	99	74 (74.7)	5.6 (3.2, 6.3)	45	35 (77.8)	5.6 (2.9, 8.1)	0.94	0.63, 1.42	0.7522
Interaction p-value									0.8914
Myriad tumour BRCA mutati	on stat	us							
tBRCAm	158	111 (70.3)	6.2 (4.9, 8.4)	77	50 (64.9)	6.0 (5.2,13.8)	1.11	0.80, 1.57	0.5234
Non-tBRCAm	97	72 (74.2)	5.8 (5.6, 8.3)	55	45 (81.8)	2.9 (2.9, 5.8)	0.68	0.47, 0.998	0.0488*
Interaction p-value									0.0557
Status somatic BRCA mutat	ions								
sBRCAm	22	14 (63.6)	7.9 (2.9,19.4)	7	3 (42.9)	NE (NE, NE)	1.63	0.53, 7.09	0.4207
gBRCAm	66	48 (72.7)	4.9 (3.0,11.1)	31	21 (67.7)	11.5 (5.8,22.2)	1.26	0.76, 2.15	0.3716
Non-BRCAm	41	28 (68.3)	5.9 (4.9,14.8)	22	20 (90.9)	2.8 (2.8, 2.9)	0.39	0.22, 0.70	0.0021*
Interaction p-value									0.0062*

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aan 25NOV2020:12:08 khcs324

Table 2.2.3.15 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Insomnia time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

Subgroup		Olap	arib + (N=2		zumab		Placebo + bevacizumab (N=132)						
	n	of pa	er (%) atients events	(lian time 95% CI) nths) [a]		Number of pat: with e	ients	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% C	[d] I	2-sided p-value [b]	
First line treatment outco	ome (IV	RS)											
NED [PDS]	92	57	(62.0)	11.0	(5.8,16.6)	48	36 (75.0)	5.6 (3.0, 9.2)	0.60	0.40,	0.92	0.0199*
NED/CR [IDS]	74	41	(55.4)	14.0	(7.9, NE)	38	26 (6	58.4)	8.5 (3.1,21.2)	0.72	0.44,	1.18	0.1886
NED/CR [Chemo]	40	29	(72.5)	11.3	(5.7,21.9)	20	15 (75.0)	7.8 (2.9,22.0)	0.89	0.48,	1.70	0.7120
PR	49	32	(65.3)	11.3	(5.8,22.3)	26	14 (53.8)	11.1 (5.6, NE)	1.05	0.57,	2.04	0.8717
Interaction p-value													0.4678
Screening laboratory tBRCA	A statu	ıs (IV	RS)										
tBRCAm	150	91	(60.7)	11.3	(8.4,16.6)	65	43 (6	56.2)	10.3 (5.6,13.9)	0.85	0.60,	1.23	0.3873
non-tBRCAm	105	68	(64.8)	11.2	(8.3,16.6)	67	48 (71.6)	5.7 (3.0,11.1)	0.67	0.46,	0.97	0.0351*
Interaction p-value													0.3588
First line treatment outco	ome (eC	RF)											
NED [PDS]	89	53	(59.6)	11.0	(5.8,23.3)	47	35 (74.5)	5.6 (3.0, 8.5)	0.57	0.37,	0.88	0.0113*
NED/CR [IDS]	74	41	(55.4)	13.9	(8.3, NE)	32	22 (6	58.8)	8.4 (3.1,21.2)	0.69	0.42,	1.19	0.1766
NED/CR [Chemo]	39	27	(69.2)	11.3	(5.6,21.9)	18	13 (72.2)	10.3 (3.0,11.2)	0.81	0.42,	1.62	0.5310
PR	50	35	(70.0)	11.0	(5.6,17.4)	34	20 (58.8)	11.3 (5.6,22.3)	1.19	0.70,	2.10	0.5280
Interaction p-value													0.2071
Screening laboratory tBRCA	A statu	ıs (eC	CRF)										
tBRCAm	147	88	(59.9)	13.4	(8.4,19.8)	67	44 (6	55.7)	10.3 (5.6,14.1)	0.83	0.58,	1.21	0.3282
non-tBRCAm	108	71	(65.7)	11.2	(8.3,14.0)	65	47 (72.3)	5.6 (2.9,11.1)	0.68	0.47,	0.99	0.0430*
Interaction p-value													0.4370
Age group													
<65 years	185	117	(63.2)	11.3	(8.4,15.2)	98	69 (70.4)	5.6 (3.0,10.3)	0.71	0.53,	0.96	0.0285*
>=65 years	70	42	(60.0)	11.3	(5.8,19.3)	34	22 (6	54.7)	11.1 (8.3,17.0)	0.87	0.53,	1.49	0.6127
Interaction p-value													0.4997

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aao 25NOV2020:12:08 khcs324

Table 2.2.3.15 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Insomnia time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

Subgroup			bevacizumab 255)			pevacizumab 132)			2-sided p-value [b]
	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI [b]	
FIGO Stage (Disease state)									
III	182	117 (64.3)	11.3 (8.3,15.2)	90	61 (67.8)	8.5 (5.6,12.7)	0.85	0.62, 1.16	0.2989
IV	73	42 (57.5)	11.2 (8.3,22.3)	42	30 (71.4)	8.3 (2.8,11.2)	0.56	0.35, 0.91	0.0193*
Interaction p-value									0.1579
Region									
Europe	245	156 (63.7)	11.1 (8.4,13.9)	126	86 (68.3)	8.3 (5.6,11.1)	0.77	0.60, 1.01	0.0589
Japan	10	3 (30.0)	NE (NE, NE)	6	5 (83.3)	22.1 (2.8,24.0)	0.30	0.06, 1.22	0.0928
Interaction p-value									0.1942
ECOG performance status at	Basel	ine							
(0) Normal activity	190	119 (62.6)	11.4 (8.4,16.6)	100	70 (70.0)	7.4 (5.2,11.1)	0.74	0.55, 1.001	0.0509
(1) Restricted activity	61	36 (59.0)	11.1 (5.8,25.7)	31	21 (67.7)	11.1 (3.2,16.9)	0.73	0.43, 1.28	0.2691
Interaction p-value									0.9756
Baseline CA-125 value									
<=ULN	228	143 (62.7)	11.4 (8.4,14.1)	118	81 (68.6)	, , ,	0.76	0.58, 1.00002	0.0500
>ULN	27	16 (59.3)	11.0 (5.7,20.0)	14	10 (71.4)	8.3 (2.9,21.2)	0.70	0.32, 1.61	0.3906
Interaction p-value									0.8612
Histological grade									
High grade	255	159 (62.4)	11.3 (8.4,14.0)	132	91 (68.9)	8.3 (5.6,11.1)	0.75	0.58, 0.98	0.0330*
Interaction p-value									NC
Cytoreductive surgery outco	me								
No residue	166	97 (58.4)	11.3 (8.4,17.3)	80	58 (72.5)	5.7 (3.0, 8.5)	0.62	0.45, 0.86	0.0045*
Residue	79	58 (73.4)	11.1 (5.7,14.0)	44	29 (65.9)	11.1 (3.5,17.0)	1.07	0.69, 1.69	0.7705
Interaction p-value									0.0490*

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aao 25NOV2020:12:08 khcs324

Table 2.2.3.15 PAOLA1: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Insomnia time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + N=2				oevacizumab 132)				
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]	Number (%) of patients n with events		,	Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	96 (65.8)	11.0 (8.3,13.8)	79	56 (70.9)	5.7 (3.0,10.3)	0.74	0.53,	1.03	0.0748
Interval	99	59 (59.6)	13.8 (8.3,18.6)	45	31 (68.9)	11.1 (5.6,16.9)	0.77	0.50,	1.20	0.2473
Interaction p-value										0.8742
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	95 (60.1)	11.3 (8.4,15.9)	77	50 (64.9)	8.5 (5.2,11.3)	0.79	0.56,	1.12	0.1840
Non-tBRCAm	97	64 (66.0)	11.2 (5.8,17.4)	55	41 (74.5)	8.3 (3.2,11.1)	0.71	0.48,	1.06	0.0909
Interaction p-value										0.6831
Status somatic BRCA mutat	ions									
sBRCAm	22	15 (68.2)	8.5 (3.0,14.1)	7	5 (71.4)	2.8 (2.8, NE)	0.46	0.18,	1.42	0.1613
gBRCAm	66	43 (65.2)	8.4 (5.8,14.1)	31	20 (64.5)	13.9 (5.6,22.3)	1.09	0.65,	1.89	0.7512
Non-BRCAm	41	26 (63.4)	13.8 (8.5,25.7)	22	17 (77.3)	8.3 (2.8,12.7)	0.61	0.33,	1.15	0.1225
Interaction p-value										0.2111

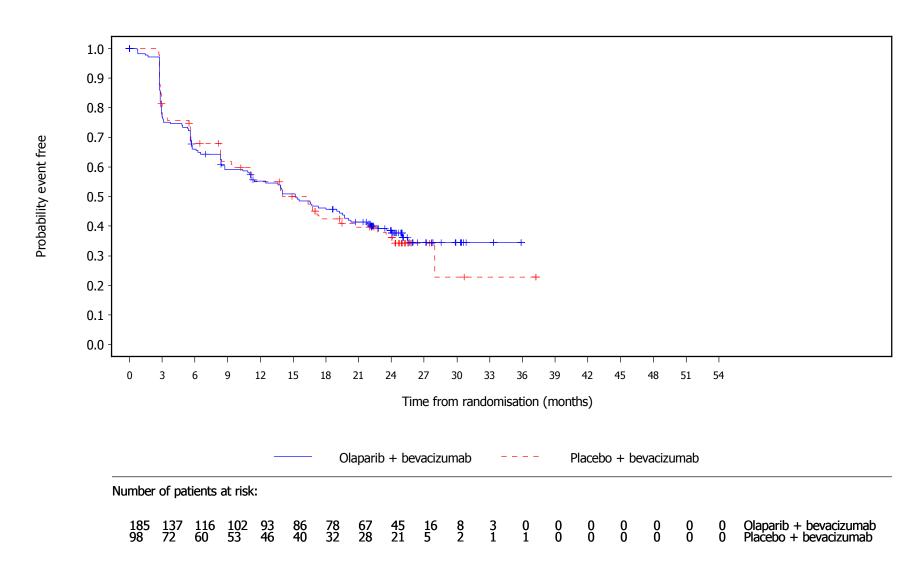
Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3aao 25NOV2020:12:08 khcs324

Figure 2.2.4.1 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Global QoL/health status time to clinically meaningful deterioration for Age group=<65 years

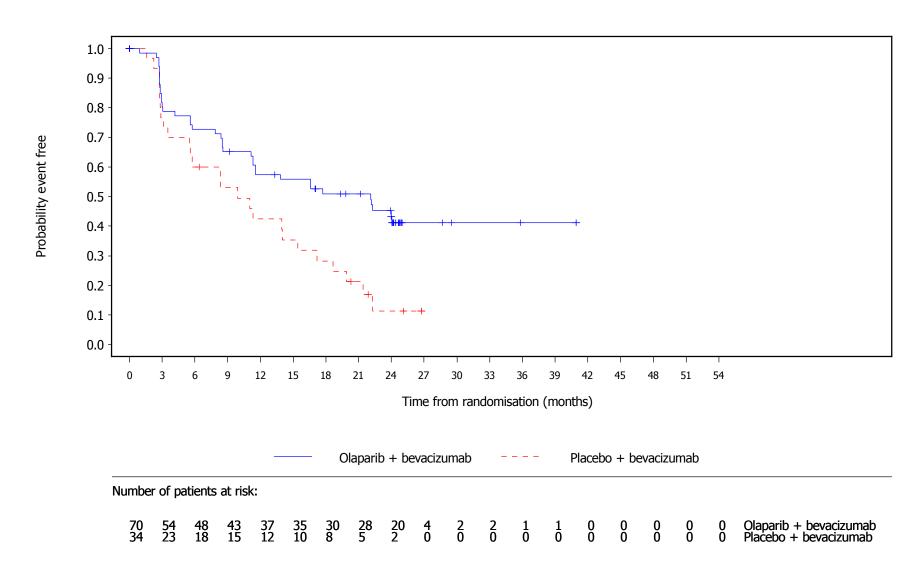
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3daa 25NOV2020:12:08 khcs324

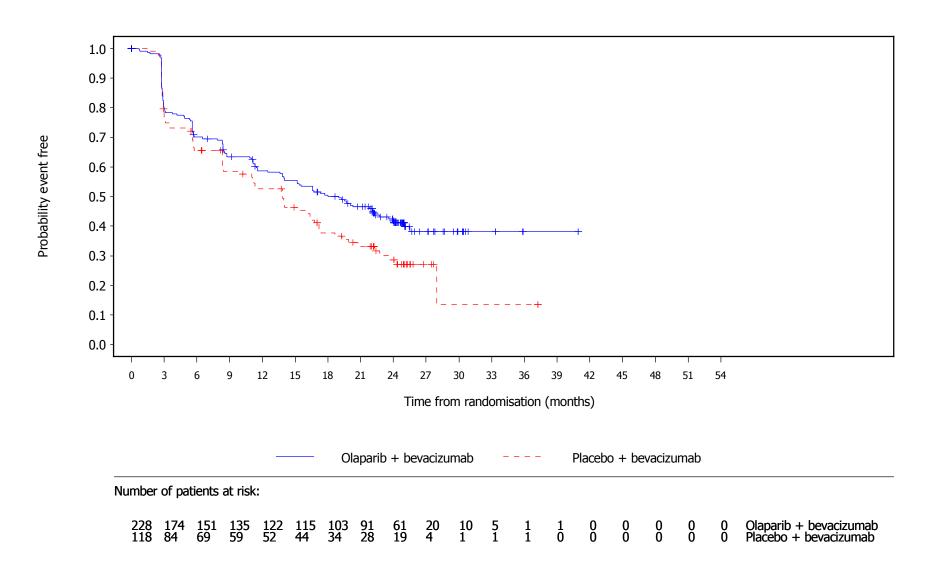
Figure 2.2.4.2 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Global QoL/health status time to clinically meaningful deterioration for Age group=>=65 years

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



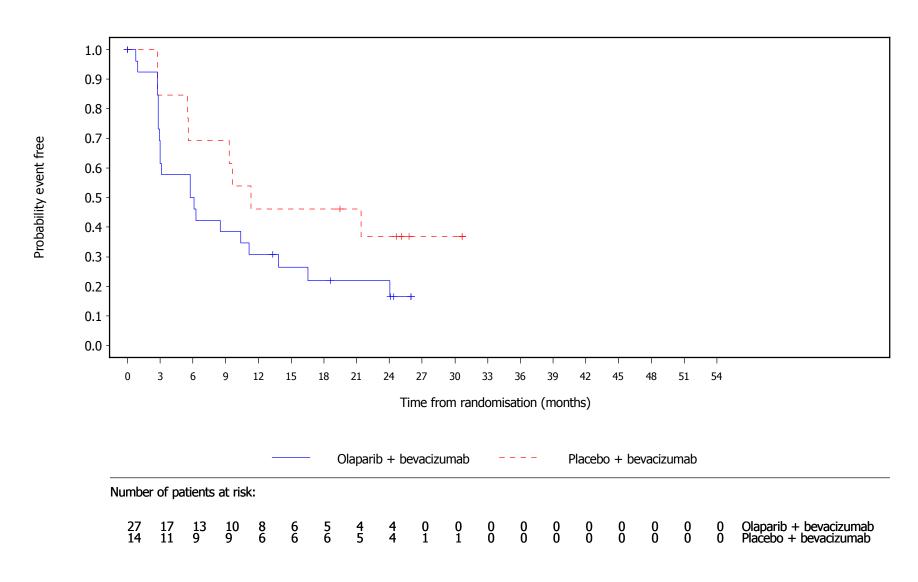
Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dab 25NOV2020:12:08 khcs324

Figure 2.2.4.3 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Global QoL/health status time to clinically meaningful deterioration for Baseline CA-125 value=<=ULN
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dac 25NOV2020:12:08 khcs324

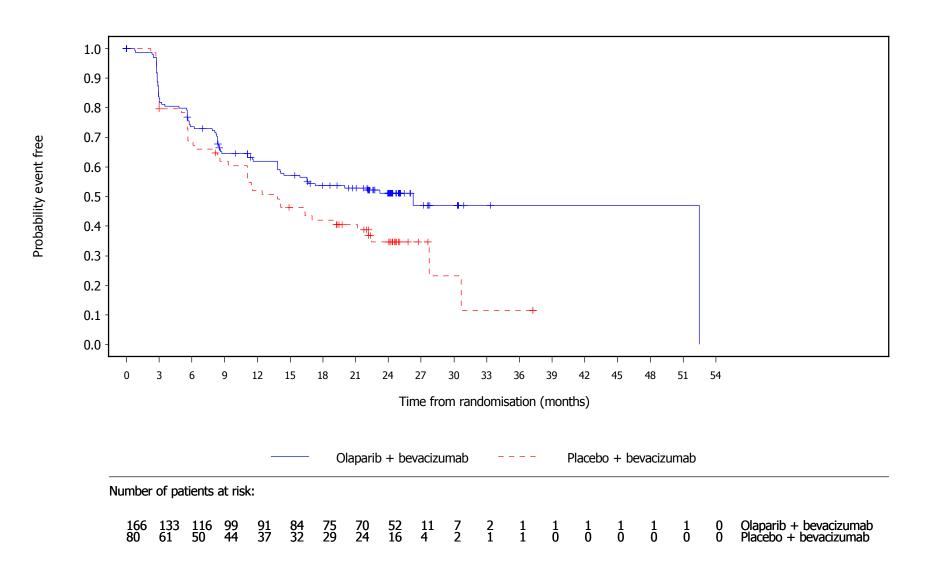
Figure 2.2.4.4 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Global QoL/health status time to clinically meaningful deterioration for Baseline CA-125 value=>ULN
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dad 25NOV2020:12:08 khcs324

Figure 2.2.4.5 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Physical time to clinically meaningful deterioration for Cytoreductive surgery outcome=No residue

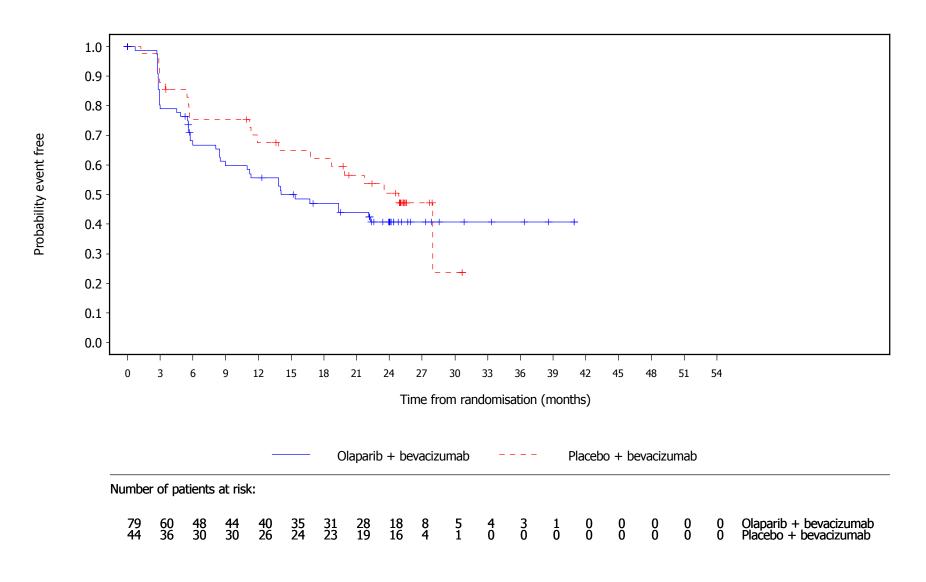
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dae 25NOV2020:12:08 khcs324

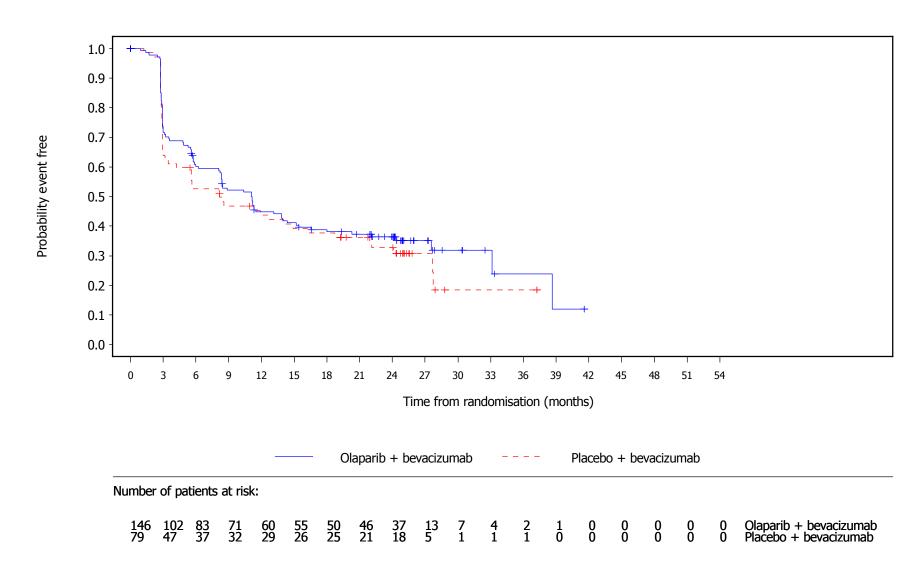
Figure 2.2.4.6 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Physical time to clinically meaningful deterioration for Cytoreductive surgery outcome=Residue

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3daf 25NOV2020:12:08 khcs324

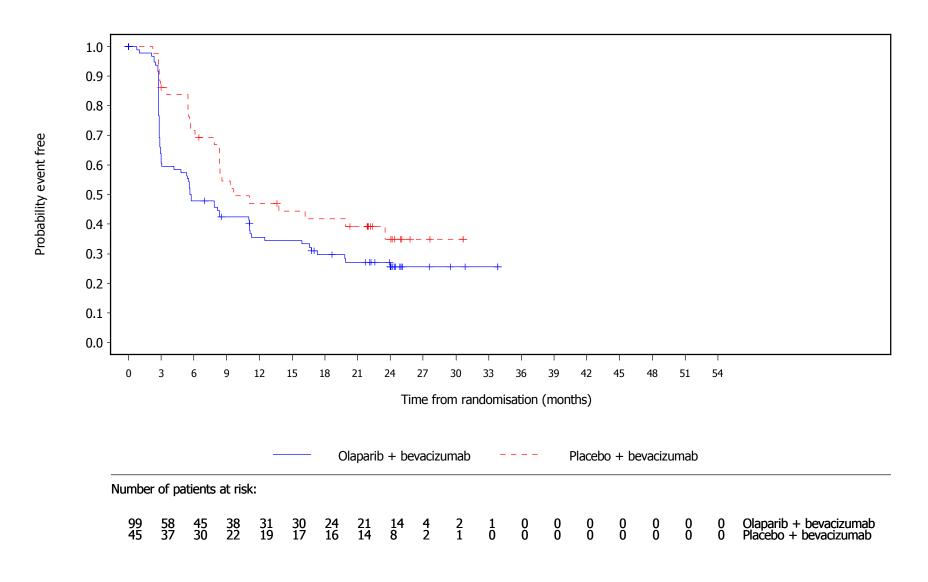
Figure 2.2.4.7 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Role time to clinically meaningful deterioration for Timing of cytoreductive surgery=Upfront
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dag 25NOV2020:12:08 khcs324

Figure 2.2.4.8 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Role time to clinically meaningful deterioration for Timing of cytoreductive surgery=Interval

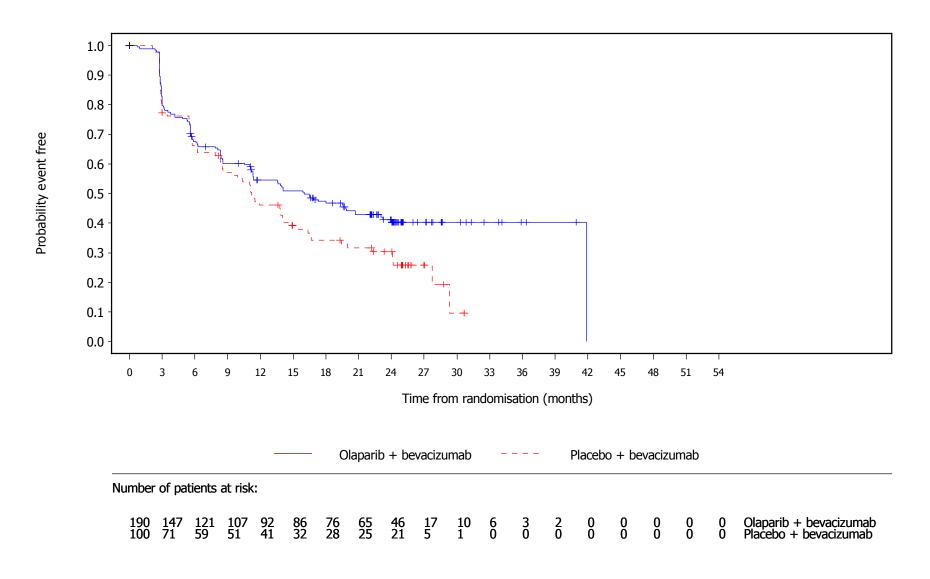
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dah 25NOV2020:12:08 khcs324

Figure 2.2.4.9 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Social time to clinically meaningful deterioration for ECOG performance status at Baseline=(0) Normal activity

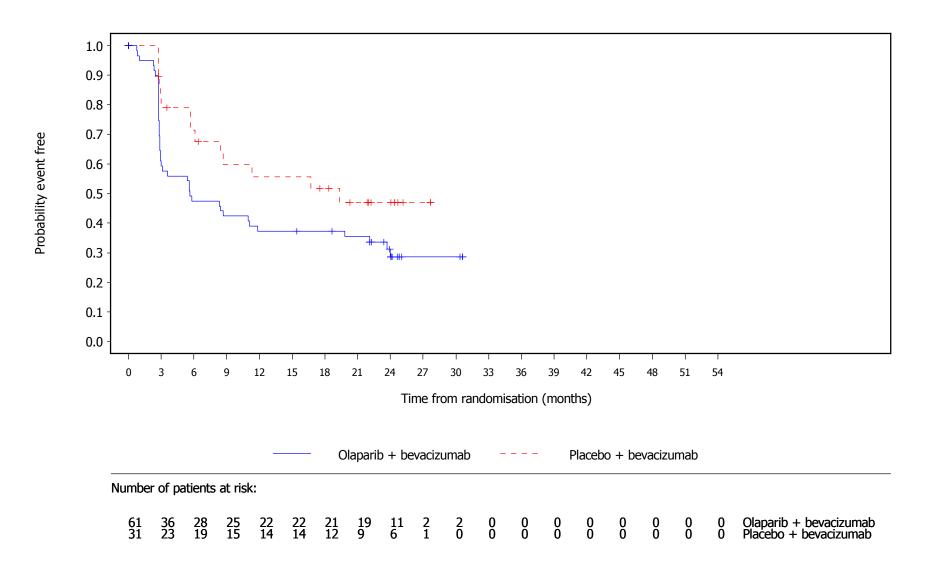
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dai 25NOV2020:12:08 khcs324

Figure 2.2.4.10 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Social time to clinically meaningful deterioration for ECOG performance status at Baseline=(1) Restricted activity

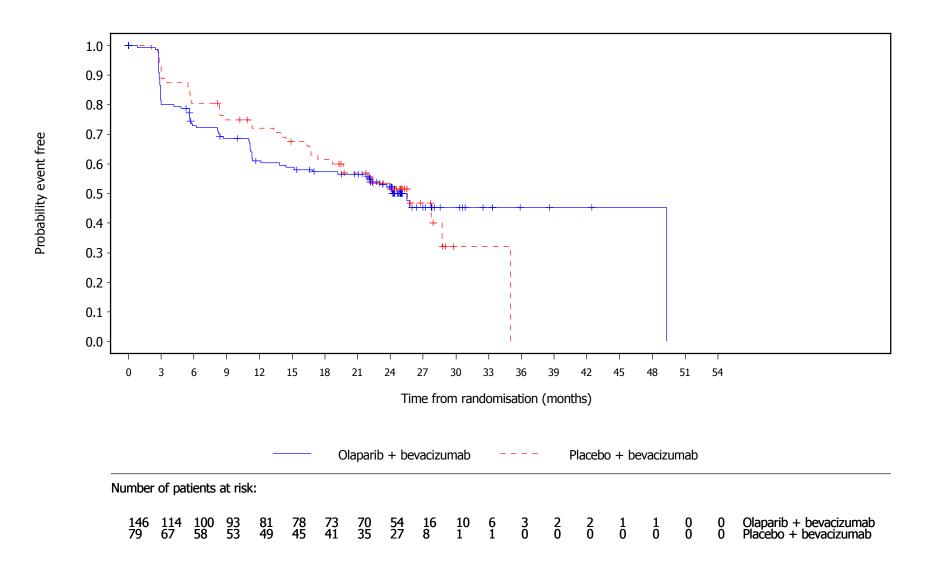
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3daj 25NOV2020:12:08 khcs324

Figure 2.2.4.11 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Loss of appetite time to clinically meaningful deterioration for Timing of cytoreductive surgery=Upfront

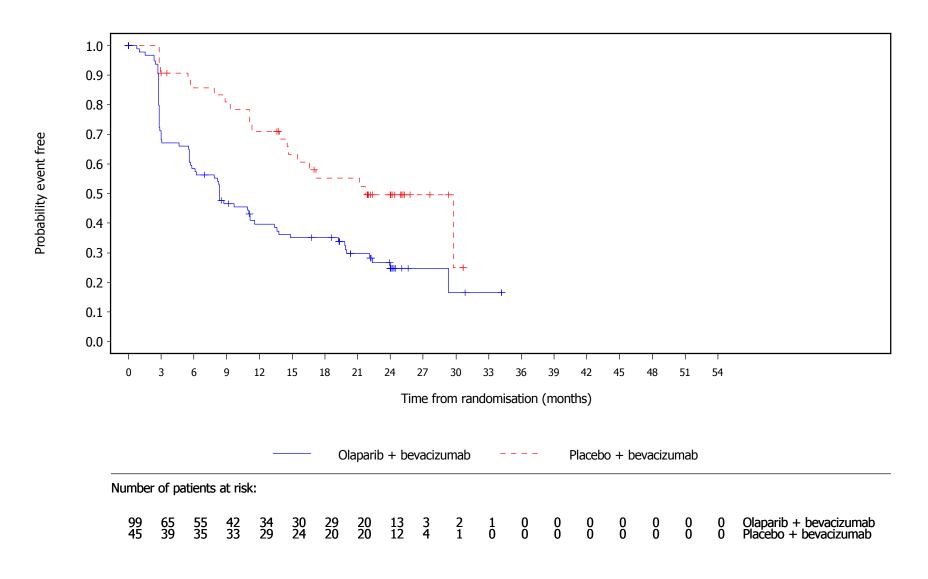
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dak 25NOV2020:12:08 khcs324

Figure 2.2.4.12 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Loss of appetite time to clinically meaningful deterioration for Timing of cytoreductive surgery=Interval

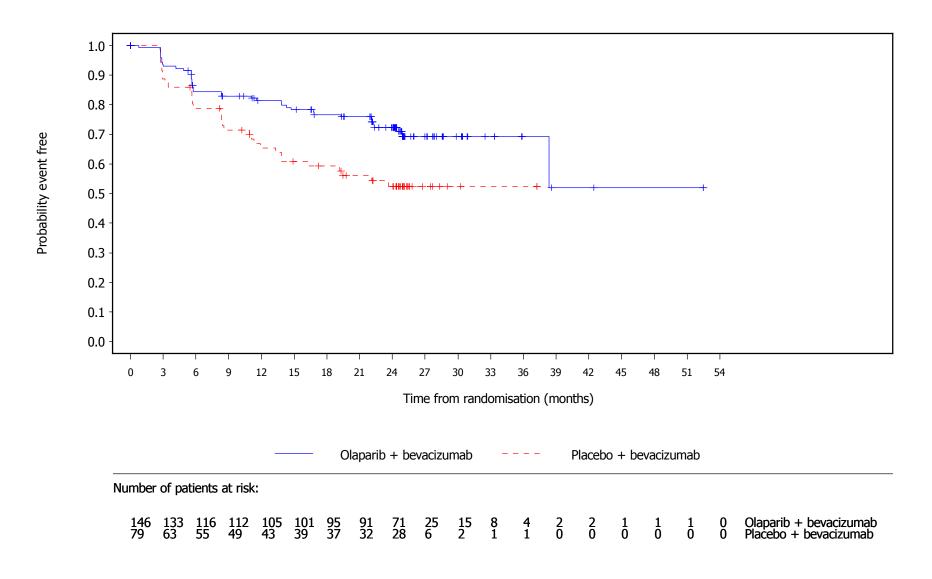
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dal 25NOV2020:12:08 khcs324

Figure 2.2.4.13 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Financial difficulties time to clinically meaningful deterioration for Timing of cytoreductive surgery=Upfront

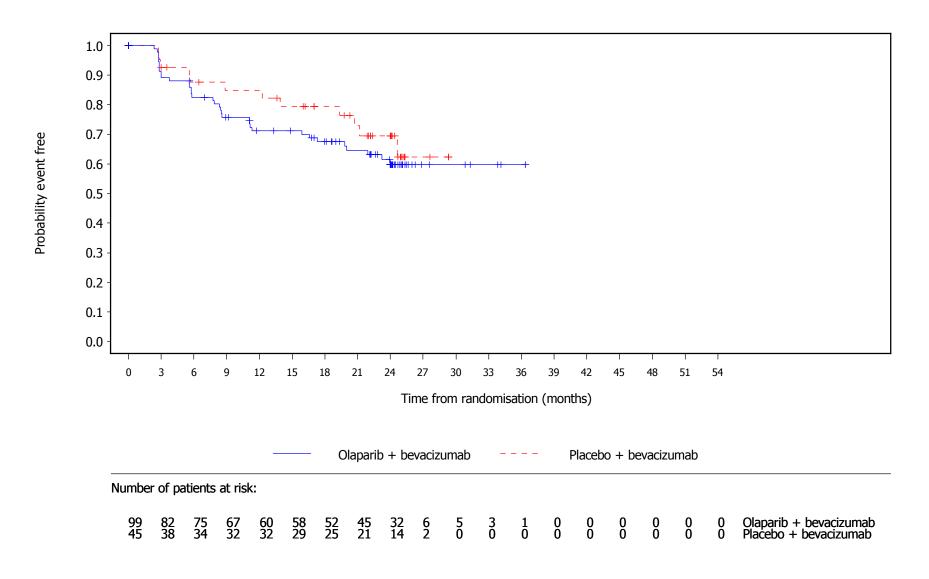
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dam 25NOV2020:12:08 khcs324

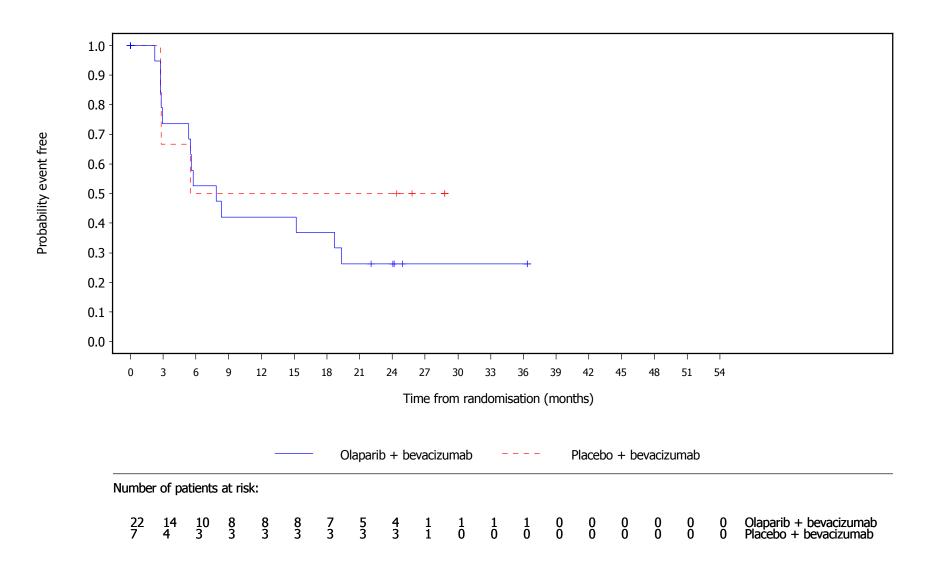
Figure 2.2.4.14 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Financial difficulties time to clinically meaningful deterioration for Timing of cytoreductive surgery=Interval

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



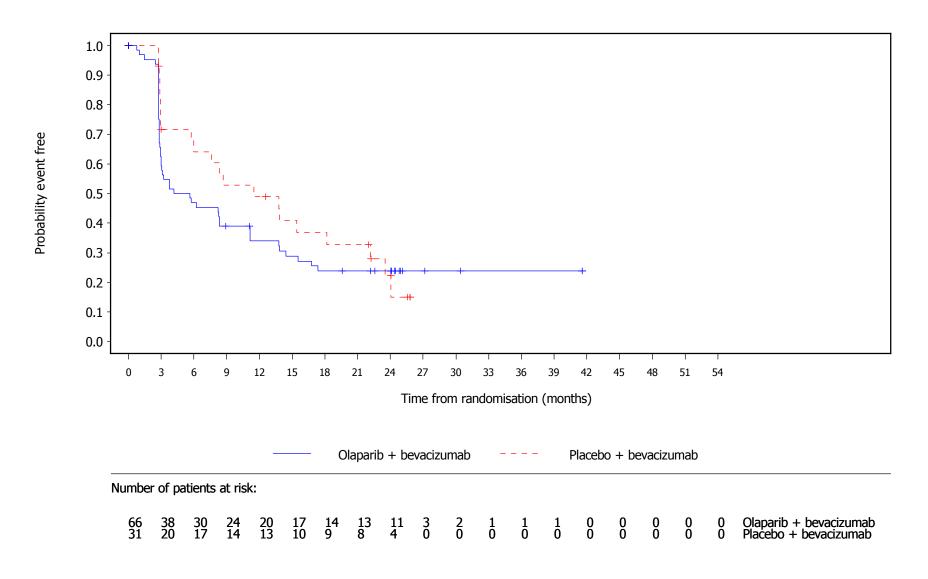
Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dan 25NOV2020:12:08 khcs324

Figure 2.2.4.15 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Pain time to clinically meaningful deterioration for Status somatic BRCA mutations=sBRCAm
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dao 25NOV2020:12:08 khcs324

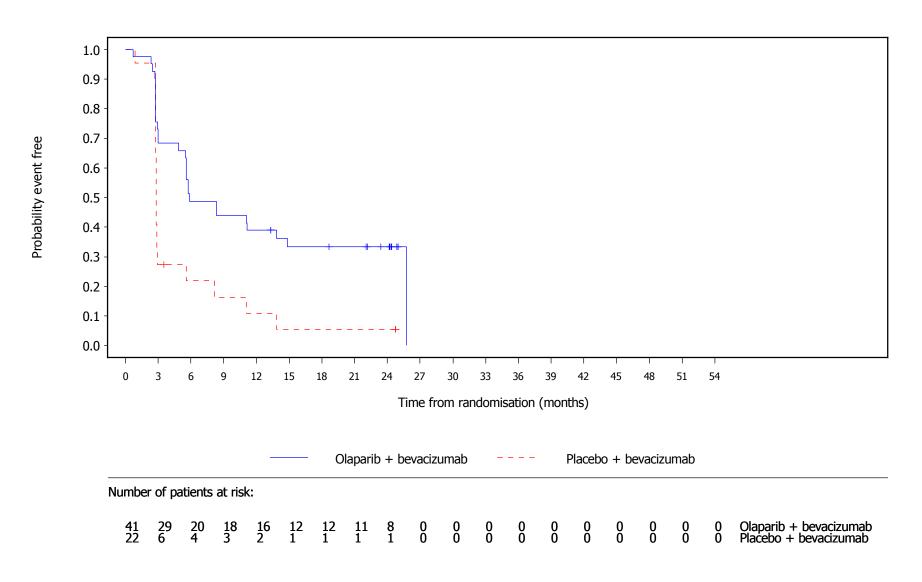
Figure 2.2.4.16 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Pain time to clinically meaningful deterioration for Status somatic BRCA mutations=gBRCAm
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dap 25NOV2020:12:08 khcs324

Figure 2.2.4.17 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Pain time to clinically meaningful deterioration for Status somatic BRCA mutations=Non-BRCAm

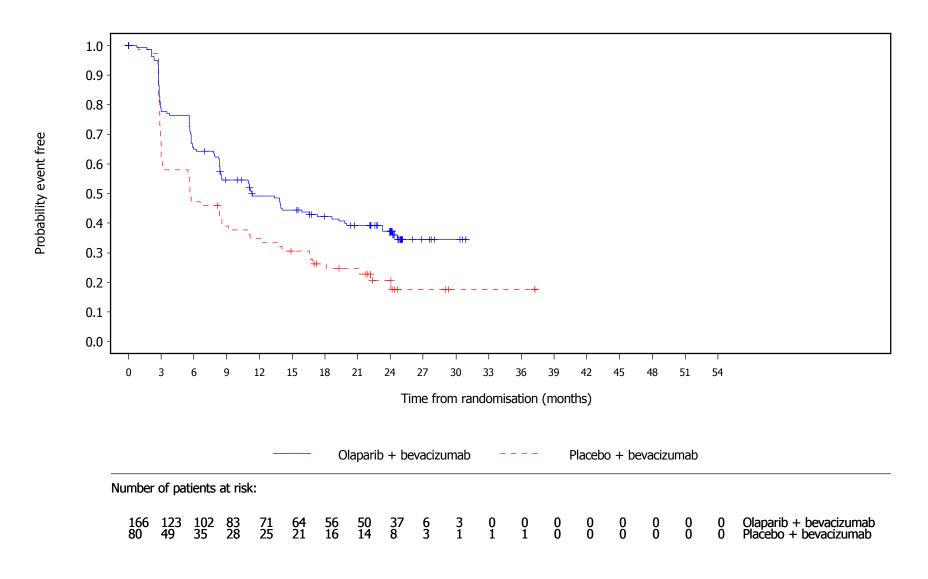
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.2.4.18 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Insomnia time to clinically meaningful deterioration for Cytoreductive surgery outcome=No residue

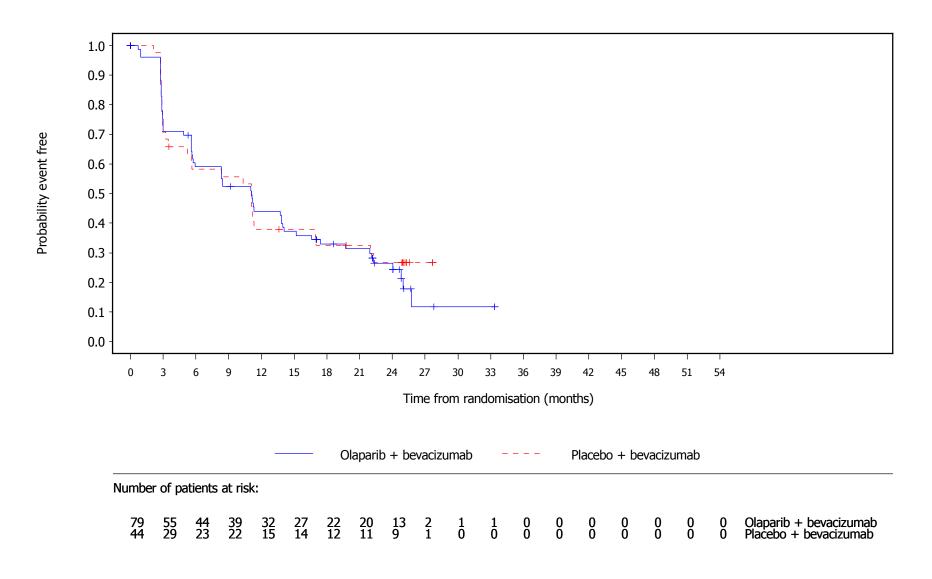
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3dar 25NOV2020:12:08 khcs324

Figure 2.2.4.19 PAOLA1: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Insomnia time to clinically meaningful deterioration for Cytoreductive surgery outcome=Residue

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3das 25NOV2020:12:08 khcs324

Table 2.3.1 PAOLA1: Summary of analysis of time to worsening in EORTC QLQ-OV28 symptom and single item scales Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)		Placebo + b (N=1				
	Number (%) Median to of patients (95% CI n with events (months))	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI	255 169 (66.3) 11.1 (8.3,	14.0) 1	132 89 (67.4)	8.3 (5.7,11.3)	0.88	0.68, 1.15	0.3509
EORTC QLQ-OV28 Symptom scale/items: Body image	255 126 (49.4) 21.9 (12.7,	NE) 1	132 71 (53.8)	18.7 (11.5,25.1)	0.93	0.70, 1.26	0.6383
EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects	255 135 (52.9) 17.9 (12.0,	24.6) 1	132 82 (62.1)	11.1 (8.3,16.6)	0.75	0.57, 0.997	0.0450*
EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment	255 134 (52.5) 12.2 (8.3,	24.1) 1	132 65 (49.2)	17.5 (11.2, NE)	1.15	0.86, 1.57	0.3624
EORTC QLQ-OV28 Symptom scale/items: Hormonal	255 135 (52.9) 19.1 (14.3,	24.2) 1	132 76 (57.6)	11.3 (5.6,19.1)	0.75	0.56, 0.996	0.0462*
EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy	255 114 (44.7) 25.3 (18.6,	NE) 1	132 58 (43.9)	23.0 (12.7, NE)	0.93	0.68, 1.29	0.6541
EORTC QLQ-OV28 Symptom scale/items: Other single items	255 127 (49.8) 21.9 (16.6,	25.7) 1	132 64 (48.5)	19.4 (16.4, NE)	1.01	0.75, 1.38	0.9536

Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation.

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

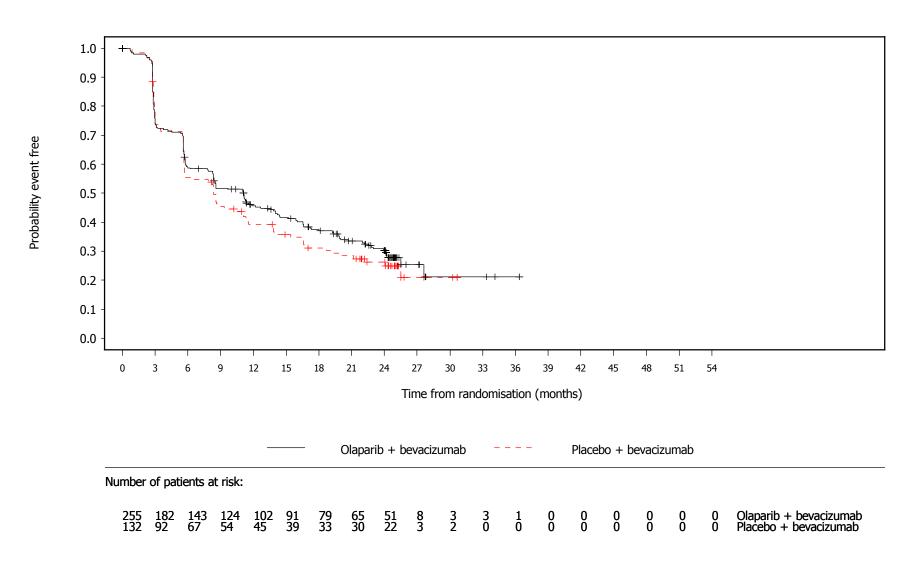
[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Figure 2.3.2.1 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI time to clinically meaningful worsening (first occurrence)

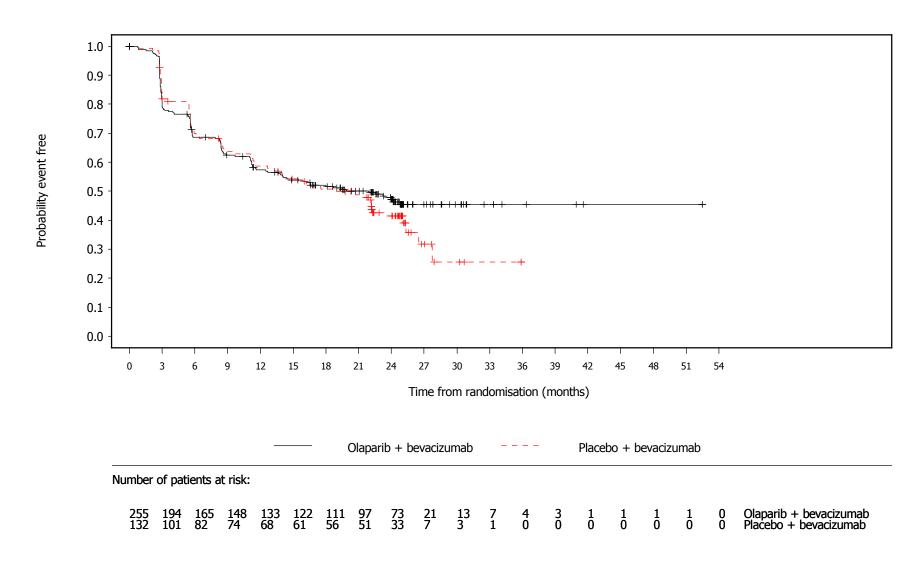
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr.sas ettemainprgaa 25NOV2020:15:58 kvbv306

Figure 2.3.2.2 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Body image time to clinically meaningful worsening (first occurrence)

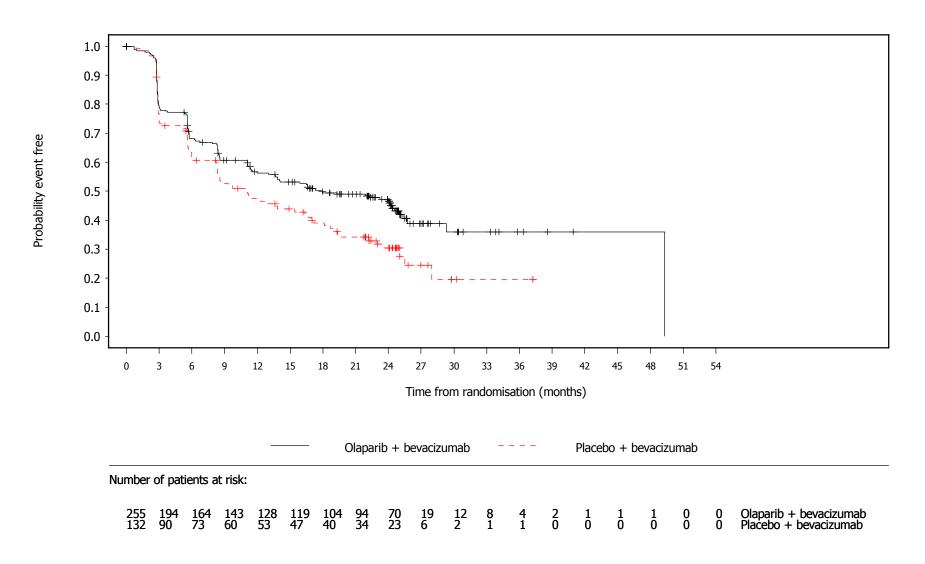
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr.sas ettemainprgab 25NOV2020:15:58 kvbv306

Figure 2.3.2.3 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects time to clinically meaningful worsening (first occurrence)

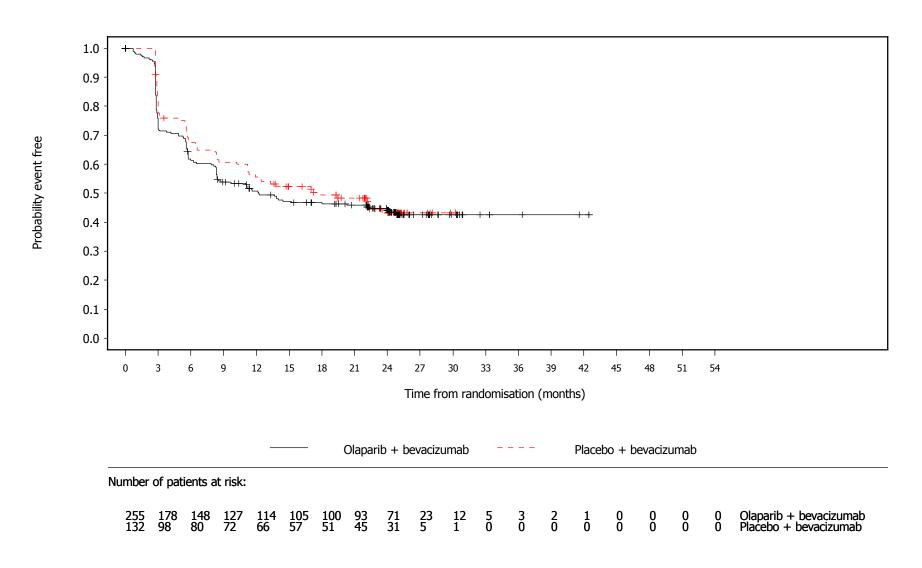
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr.sas ettemainprgac 25NOV2020:15:58 kvbv306

Figure 2.3.2.4 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful worsening (first occurrence)

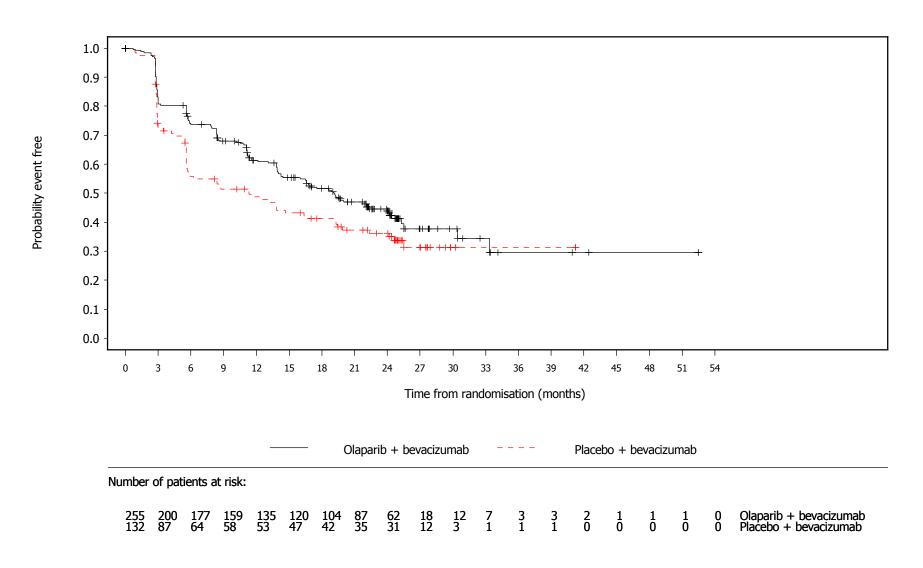
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr.sas ettemainprgad 25NOV2020:15:58 kvbv306

Figure 2.3.2.5 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Hormonal time to clinically meaningful worsening (first occurrence)

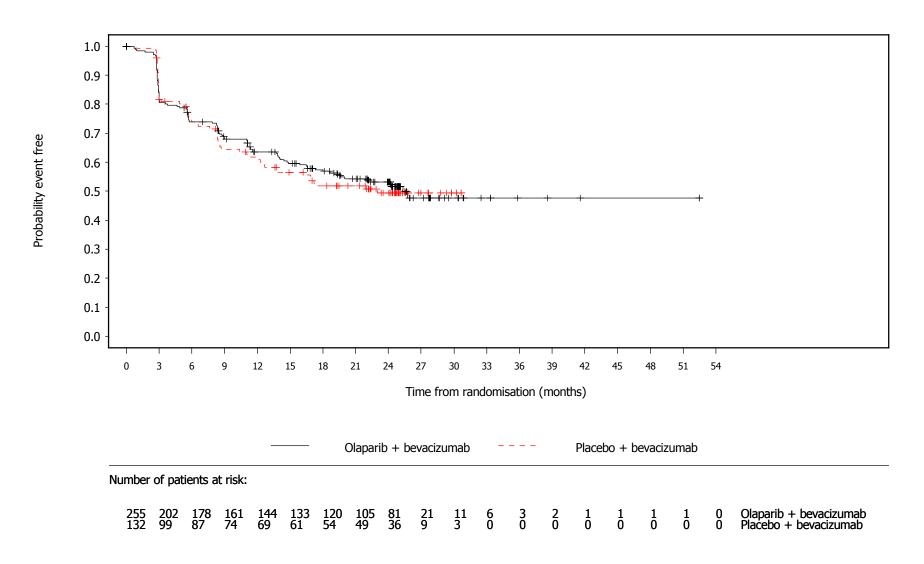
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr.sas ettemainprgae 25NOV2020:15:58 kvbv306

Figure 2.3.2.6 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy time to clinically meaningful worsening (first occurrence)

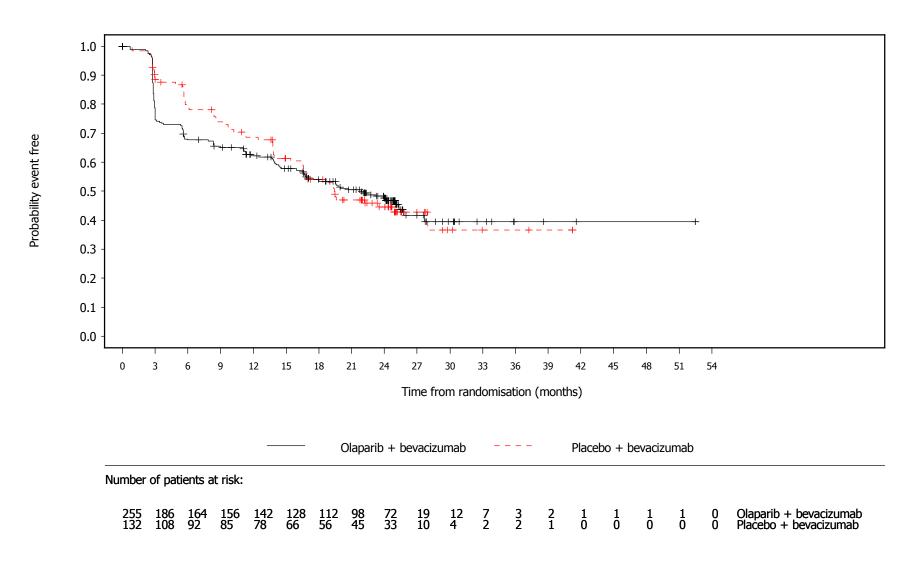
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr.sas ettemainprgaf 25NOV2020:15:58 kvbv306

Figure 2.3.2.7 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Other single items time to clinically meaningful worsening (first occurrence)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr.sas ettemainprgag 25NOV2020:15:58 kvbv306

Table 2.3.3.1 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		bevacizumab 255)		Placebo + h (N=1	oevacizumab 132)			
Subgroup	Number (%) of patients n with events	(95% CI)	(Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IVRS)							
NED [PDS]	92 59 (64.1)	11.4 (5.8,18.0)	48	37 (77.1)	5.8 (5.6,11.1)	0.62	0.41, 0.94	0.0254*
NED/CR [IDS]	74 47 (63.5)	8.3 (5.6,20.0)	38	21 (55.3)	13.8 (5.6, NE)	1.28	0.78, 2.19	0.3372
NED/CR [Chemo]	40 27 (67.5)	11.3 (5.6,19.2)	20	13 (65.0)	8.5 (5.6, NE)	1.13	0.59, 2.25	0.7253
PR	49 36 (73.5)	8.5 (5.6,12.2)	26	18 (69.2)	7.9 (2.9,16.6)	0.89	0.51, 1.60	0.6861
Interaction p-value								0.1444
Screening laboratory tBRC	A status (IVRS)							
tBRCAm	150 104 (69.3)	8.5 (5.7,13.6)	65	47 (72.3)	8.3 (5.6,11.5)	0.96	0.68, 1.36	0.8116
non-tBRCAm	105 65 (61.9)	11.9 (8.3,19.8)	67	42 (62.7)	8.5 (5.6,16.4)	0.82	0.56, 1.22	0.3288
Interaction p-value								0.5637
First line treatment outo	come (eCRF)							
NED [PDS]	89 56 (62.9)	12.7 (5.8,18.0)	47	35 (74.5)	7.0 (5.6,11.1)	0.65	0.43, 1.01	0.0546
NED/CR [IDS]	74 49 (66.2)	8.3 (5.6,17.3)	32	16 (50.0)	13.9 (5.6, NE)	1.61	0.94, 2.93	0.0847
NED/CR [Chemo]	39 25 (64.1)	11.2 (5.6,19.2)	18	10 (55.6)	10.3 (5.6, NE)	1.30	0.64, 2.83	0.4797
PR	50 37 (74.0)	9.8 (5.7,16.6)	34	27 (79.4)	7.9 (2.9,11.3)	0.72	0.44, 1.20	0.2082
Interaction p-value								0.0406*
Screening laboratory tBRC	CA status (eCRF)							
tBRCAm	147 103 (70.1)	8.4 (5.7,12.7)	67	47 (70.1)	8.3 (5.7,11.5)	1.01	0.72, 1.43	0.9721
non-tBRCAm	108 66 (61.1)	12.2 (8.5,19.9)	65	42 (64.6)	8.3 (5.6,14.0)	0.78	0.53, 1.16	0.2125
Interaction p-value								0.3352
Age group								
<65 years	185 130 (70.3)	8.5 (5.7,12.2)	98	67 (68.4)	8.3 (5.7,11.4)	1.01	0.75, 1.36	0.9708
>=65 years	70 39 (55.7)	16.6 (8.5, NE)	34	22 (64.7)	8.5 (3.5,18.7)	0.68	0.41, 1.16	0.1523
Interaction p-value								0.2016

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3baa 25NoV2020:12:08 khcs324

Table 2.3.3.1 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			pevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b] 95% CI	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)									
III	182	118 (64.8)	11.1 (8.3,15.2)	90	61 (67.8)	8.5 (5.6,11.3)	0.89	0.66, 1.2	2 0.4623
IV	73	51 (69.9)	8.5 (5.6,13.6)	42	28 (66.7)	8.3 (5.6,15.4)	0.94	0.60, 1.5	1 0.7992
Interaction p-value									0.8415
Region									
Europe	245	162 (66.1)	11.1 (8.3,14.0)	126	86 (68.3)	8.3 (5.7,11.1)	0.87	0.67, 1.1	
Japan	10	7 (70.0)	8.3 (3.1, NE)	6	3 (50.0)	24.0 (5.5, NE)	2.01	0.56, 9.3	
Interaction p-value									0.2136
ECOG performance status at	Basel	ine							
(0) Normal activity	190	131 (68.9)	8.6 (6.0,12.2)	100	69 (69.0)	8.3 (5.6,11.1)	0.94	0.71, 1.2	6 0.6801
(1) Restricted activity	61	35 (57.4)	14.2 (8.3, NE)	31	20 (64.5)	11.3 (5.6,21.2)	0.79	0.46, 1.3	
Interaction p-value									0.5823
Baseline CA-125 value									
<=ULN	228	151 (66.2)	11.1 (8.3,14.0)	118		8.3 (5.7,11.5)	0.90	0.69, 1.1	
>ULN	27	18 (66.7)	11.2 (5.6,20.0)	14	9 (64.3)	11.1 (2.9, NE)	0.97	0.45, 2.2	
Interaction p-value									0.8488
Histological grade									
High grade	255	169 (66.3)	11.1 (8.3,14.0)	132	89 (67.4)	8.3 (5.7,11.3)	0.90	0.70, 1.1	7 0.4429
Interaction p-value									NC
Cytoreductive surgery outco	ome								
No residue	166	107 (64.5)	11.1 (6.2,15.9)	80	52 (65.0)	8.5 (5.6,13.9)	0.93	0.67, 1.3	
Residue	79	57 (72.2)	11.1 (5.7,15.2)	44	31 (70.5)	8.3 (5.6,14.0)	0.94	0.61, 1.4	8 0.7985
Interaction p-value									0.9425

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3baa 25NOV2020:12:08 khcs324

Table 2.3.3.1 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2			Placebo + 1 (N=1					
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	99 (67.8)	11.1 (6.0,14.2)	79	57 (72.2)	8.3 (5.6,11.1)	0.79	0.57,	1.10	0.1636
Interval	99	65 (65.7)	9.7 (5.7,19.4)	45	26 (57.8)	11.1 (5.6, NE)	1.23	0.79,	1.98	0.3598
Interaction p-value										0.1164
Myriad tumour BRCA mutat	ion stat	us								
tBRCAm	158	105 (66.5)	11.0 (6.0,14.4)	77	51 (66.2)	8.5 (5.7,13.9)	0.94	0.68,	1.32	0.7189
Non-tBRCAm	97	64 (66.0)	11.1 (5.8,16.0)	55	38 (69.1)	8.1 (5.5,14.0)	0.86	0.58,	1.29	0.4480
Interaction p-value										0.7221
Status somatic BRCA mutat	ions									
sBRCAm	22	12 (54.5)	16.6 (2.8, NE)	7	3 (42.9)	NE (NE, NE)	1.42	0.45,	6.25	0.5714
gBRCAm	66	48 (72.7)	6.0 (5.6,13.6)	31	24 (77.4)	8.8 (5.6,13.9)	0.93	0.58,	1.55	0.7875
Non-BRCAm	41	30 (73.2)	11.0 (5.6,19.8)	22	18 (81.8)	5.7 (2.9,13.8)	0.74	0.42,	1.36	0.3256
Interaction p-value										0.6128

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3baa 25NoV2020:12:08 khcs324

Table 2.3.3.2 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Body image time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2			Placebo + h					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IVI	RS)								
NED [PDS]	92	37 (40.2)	NE (NE, NE)	48	26 (54.2)	11.3 (3.0, NE)	0.55	0.33,	0.92	0.0224*
NED/CR [IDS]	74	41 (55.4)	14.8 (8.3, NE)	38	19 (50.0)	22.2 (8.8,26.5)	1.31	0.77,	2.31	0.3184
NED/CR [Chemo]	40	19 (47.5)	23.4 (11.6, NE)	20	12 (60.0)	12.4 (5.2, NE)	0.82	0.40,	1.74	0.6015
PR	49	29 (59.2)	11.2 (5.6,24.8)	26	14 (53.8)	17.5 (11.1,25.4)	1.24	0.67,	2.42	0.5046
Interaction p-value										0.0873
Screening laboratory tBRC	CA status	s (IVRS)								
tBRCAm	150	73 (48.7)	22.5 (11.6, NE)	65	33 (50.8)	21.2 (11.3, NE)	0.95	0.63,	1.45	0.8049
non-tBRCAm	105	53 (50.5)	19.5 (11.1, NE)	67	38 (56.7)	14.0 (8.5,25.4)	0.85	0.56,	1.30	0.4504
Interaction p-value										0.7143
First line treatment outo	come (eCI	RF)								
NED [PDS]	89	36 (40.4)	NE (NE, NE)	47	25 (53.2)	11.5 (5.6, NE)	0.58	0.35,	0.98	0.0405*
NED/CR [IDS]	74	38 (51.4)	19.9 (8.3, NE)	32	17 (53.1)	22.0 (8.3,26.5)	1.08	0.62,	1.96	0.7991
NED/CR [Chemo]	39	20 (51.3)	21.9 (5.6, NE)	18	10 (55.6)	13.3 (3.5, NE)	0.92	0.44,	2.05	0.8303
PR	50	30 (60.0)	11.2 (5.7,24.8)	34	18 (52.9)	18.7 (12.7, NE)	1.36	0.76,	2.48	0.2999
Interaction p-value										0.1661
Screening laboratory tBRC	CA status	s (eCRF)								
tBRCAm	147	71 (48.3)	23.2 (12.5, NE)	67	34 (50.7)	21.2 (11.2, NE)	0.92	0.62,	1.40	0.6935
non-tBRCAm	108	55 (50.9)	18.7 (11.1, NE)	65	37 (56.9)	16.6 (8.5,25.4)	0.88	0.58,	1.34	0.5332
Interaction p-value										0.8653
Age group										
<65 years	185	90 (48.6)	22.5 (11.6, NE)	98	53 (54.1)	20.7 (11.1,25.4)	0.86	0.61,	1.21	0.3724
>=65 years	70	36 (51.4)	16.6 (8.4, NE)	34	18 (52.9)	18.7 (6.4, NE)	0.99	0.57,	1.78	0.9717
Interaction p-value										0.6649

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bab 25NOV2020:12:08 khcs324

Table 2.3.3.2 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Body image time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2				Placebo + N=1					
Subgroup		Number (%) of patients with events	Median tin (95% CI) (months) [Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	87 (47.8)	23.4 (13.7,	NE)	90	48 (53.3)	18.7 (8.8,27.8)	0.85	0.60,	1.22	0.3652
IV	73	39 (53.4)	14.8 (8.4,	NE)	42	23 (54.8)	20.7 (9.7,25.4)	1.00	0.60,	1.70	0.9950
Interaction p-value											0.6090
Region											
Europe	245	122 (49.8)	19.9 (12.5,	NE)	126	68 (54.0)	17.5 (11.5,25.1)	0.89	0.66,	1.20	0.4241
Japan	10	4 (40.0)	NE (NE,	NE)	6	3 (50.0)	NE (NE, NE)	0.98	0.22,	4.96	0.9759
Interaction p-value											0.8990
ECOG performance status at	Basel	ine									
(0) Normal activity	190	95 (50.0)	18.0 (11.3,	NE)	100	55 (55.0)	20.7 (11.2,25.4)	0.92	0.66,	1.29	0.6149
(1) Restricted activity	61	30 (49.2)	23.4 (8.4,	NE)	31	16 (51.6)	17.5 (6.1, NE)	0.87	0.48,	1.64	0.6597
Interaction p-value											0.8836
Baseline CA-125 value											
<=ULN	228	112 (49.1)	22.5 (13.7,	NE)	118	64 (54.2)	16.6 (11.1,25.1)	0.85	0.63,	1.16	0.3098
>ULN	27	14 (51.9)	11.3 (3.1,	NE)	14	7 (50.0)	21.2 (2.9, NE)	1.30	0.54,	3.44	0.5632
Interaction p-value											0.3776
Histological grade											
High grade	255	126 (49.4)	21.9 (12.7,	NE)	132	71 (53.8)	18.7 (11.5,25.1)	0.89	0.67,	1.20	0.4356
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	76 (45.8)	NE (NE,	NE)	80	43 (53.8)	20.7 (8.3,26.5)	0.76	0.52,	1.11	0.1522
Residue	79	44 (55.7)	16.4 (8.4,2	4.8)	44	24 (54.5)	18.7 (11.2,25.4)	1.12	0.69,	1.86	0.6618
Interaction p-value											0.2202

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bab 25NOV2020:12:08 khcs324

Table 2.3.3.2 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Body image time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			Olaparib + bevacizumab (N=255)			bevacizumab 132)			
ubgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	,	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]	
Timing of cytoreductive s	surgery								
Upfront	146	67 (45.9)	NE (NE, NE)	79	42 (53.2)	14.0 (8.3, NE)	0.74	0.50, 1.09	0.1248
Interval	99	53 (53.5)	16.6 (9.7,24.0)	45	25 (55.6)	22.1 (8.8,25.4)	1.11	0.70, 1.81	0.6735
Interaction p-value									0.1894
Myriad tumour BRCA mutati	on stat	us							
tBRCAm	158	78 (49.4)	21.9 (11.3, NE)	77	39 (50.6)	21.2 (11.2,27.8)	0.94	0.64, 1.39	0.7375
Non-tBRCAm	97	48 (49.5)	19.9 (9.7, NE)	55	32 (58.2)	16.6 (9.7,25.4)	0.84	0.54, 1.32	0.4354
Interaction p-value									0.7067
Status somatic BRCA mutat	ions								
sBRCAm	22	5 (22.7)	NE (NE, NE)	7	3 (42.9)	NE (NE, NE)	0.43	0.11, 2.11	0.2737
gBRCAm	66	37 (56.1)	12.5 (8.3, NE)	31	16 (51.6)	22.3 (11.3, NE)	1.23	0.70, 2.28	0.4784
Non-BRCAm	41	21 (51.2)	19.9 (5.8, NE)	22	13 (59.1)	16.6 (9.7, NE)	0.93	0.47, 1.91	0.8398
Interaction p-value									0.4216

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bab 25NOV2020:12:08 khcs324

Table 2.3.3.3 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	•	Olaparib + 1 (N=2			Placebo + N=1					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	95% CI [b]	2-sided p-value [b]
First line treatment outo	ome (IV	RS)								
NED [PDS]	92	51 (55.4)	17.9 (11.4,49.3)	48	26 (54.2)	11.4 (5.7,25.5)	0.81	0.51,	1.31	0.3798
NED/CR [IDS]	74	40 (54.1)	17.3 (6.3,29.3)	38	23 (60.5)	8.3 (5.5,25.1)	0.83	0.50,	1.41	0.4900
NED/CR [Chemo]	40	19 (47.5)	25.0 (5.6, NE)	20	16 (80.0)	7.1 (2.9,19.1)	0.50	0.26,	0.98	0.0442*
PR	49	25 (51.0)	13.7 (8.3, NE)	26	17 (65.4)	13.8 (8.3,19.4)	0.65	0.35,	1.22	0.1765
Interaction p-value										0.6138
Screening laboratory tBRC	'A statu	s (IVRS)								
tBRCAm	150	71 (47.3)	24.3 (16.5, NE)	65	40 (61.5)	16.6 (5.8,22.2)	0.64	0.44,	0.95	0.0269*
non-tBRCAm	105	64 (61.0)	11.3 (5.8,22.3)	67	42 (62.7)	9.7 (6.0,12.7)	0.86	0.59,	1.28	0.4652
Interaction p-value										0.2823
First line treatment outc	ome (eCl	RF)								
NED [PDS]	89	47 (52.8)	18.0 (11.4,49.3)	47	25 (53.2)	16.4 (5.8, NE)	0.79	0.49,	1.30	0.3493
NED/CR [IDS]	74	39 (52.7)	17.3 (6.3, NE)	32	19 (59.4)	8.8 (5.5, NE)	0.83	0.49,	1.47	0.5075
NED/CR [Chemo]	39	17 (43.6)	25.0 (8.6, NE)	18	13 (72.2)	11.1 (3.5,19.1)	0.50	0.25,	1.06	0.0706
PR	50	30 (60.0)	11.3 (8.3,25.7)	34	24 (70.6)	9.7 (5.6,19.4)	0.71	0.42,	1.23	0.2177
Interaction p-value										0.7295
Screening laboratory tBRC	'A statu	s (eCRF)								
tBRCAm	147	70 (47.6)	24.3 (15.9, NE)	67	41 (61.2)	15.4 (5.8,22.2)	0.64	0.44,	0.95	0.0271*
non-tBRCAm	108	65 (60.2)	11.3 (6.8,22.3)	65	41 (63.1)	8.6 (5.7,13.8)	0.85	0.57,	1.26	0.4087
Interaction p-value										0.3210
Age group										
<65 years	185	96 (51.9)	18.0 (11.4,25.4)	98	59 (60.2)	11.4 (6.0,19.4)	0.75	0.55,	1.05	0.0911
>=65 years	70	39 (55.7)	16.7 (8.4,49.3)	34	23 (67.6)	8.3 (5.8,13.8)	0.63	0.38,	1.07	0.0835
Interaction p-value										0.5529

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bac 25NoV2020:12:08 khcs324

Table 2.3.3.3 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			oevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)									
III	182	103 (56.6)	16.5 (11.2,24.1)	90	54 (60.0)	12.3 (8.3,19.4)	0.87	0.63, 1.21	0.3975
IV	73	32 (43.8)	25.0 (8.6, NE)	42	28 (66.7)	8.3 (3.0,18.1)	0.45	0.27, 0.76	0.0027*
Interaction p-value									0.0358*
Region									
Europe	245	130 (53.1)	17.9 (12.0,24.6)	126	78 (61.9)	11.2 (8.3,17.0)	0.72	0.55, 0.96	0.0259*
Japan	10	5 (50.0)	8.4 (2.8, NE)	6	4 (66.7)	5.7 (2.8, NE)	0.65	0.17, 2.62	0.5243
Interaction p-value									0.8759
ECOG performance status at	Basel	ine							
(0) Normal activity	190	99 (52.1)	19.3 (12.0,25.0)	100	65 (65.0)	9.3 (6.0,16.6)	0.68	0.50, 0.94	0.0194*
(1) Restricted activity	61	32 (52.5)	13.8 (6.8, NE)	31	17 (54.8)	12.3 (5.7,22.9)	0.79	0.45, 1.46	0.4454
Interaction p-value									0.6628
Baseline CA-125 value									
<=ULN	228		21.9 (13.6,25.0)	118		11.4 (8.2,18.7)	0.73	0.55, 0.99	0.0403*
>ULN	27	16 (59.3)	8.4 (5.6,29.3)	14	11 (78.6)	9.3 (2.8,17.0)	0.63	0.29, 1.39	0.2454
Interaction p-value									0.7205
Histological grade									
High grade	255	135 (52.9)	17.9 (12.0,24.6)	132	82 (62.1)	11.1 (8.3,16.6)	0.72	0.55, 0.95	0.0205*
Interaction p-value									NC
Cytoreductive surgery outco	ome								
No residue	166	88 (53.0)	17.9 (11.5,29.3)	80	45 (56.3)	11.2 (5.8,19.7)	0.80	0.56, 1.16	0.2410
Residue	79	40 (50.6)	24.2 (8.6, NE)	44	33 (75.0)	8.6 (5.6,18.7)	0.54	0.34, 0.86	0.0104*
Interaction p-value									0.1848

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bac 25NOV2020:12:08 khcs324

Table 2.3.3.3 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bevacizumab (N=255)			Placebo + 1 (N=1					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	urgery									
Upfront	146	73 (50.0)	24.1 (13.8,49.3)	79	49 (62.0)	8.6 (5.7,18.7)	0.62	0.43,	0.89	0.0110*
Interval	99	55 (55.6)	16.6 (8.3,24.3)	45	29 (64.4)	11.1 (5.6,23.5)	0.82	0.53,	1.31	0.4046
Interaction p-value										0.3282
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	75 (47.5)	24.3 (15.9, NE)	77	43 (55.8)	17.0 (8.3,22.9)	0.74	0.51,	1.08	0.1204
Non-tBRCAm	97	60 (61.9)	11.2 (6.8,22.3)	55	39 (70.9)	8.3 (3.5,12.3)	0.71	0.47,	1.07	0.0990
Interaction p-value										0.8760
Status somatic BRCA mutat	ions									
sBRCAm	22	12 (54.5)	11.1 (2.8, NE)	7	5 (71.4)	12.6 (2.8, NE)	0.78	0.29,	2.44	0.6400
gBRCAm	66	35 (53.0)	24.2 (8.5,29.3)	31	17 (54.8)	17.2 (5.6,25.1)	0.82	0.46,	1.50	0.5020
Non-BRCAm	41	26 (63.4)	11.1 (5.7,49.3)	22	18 (81.8)	2.9 (2.8,12.3)	0.46	0.25,	0.86	0.0163*
Interaction p-value										0.3840

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bac 25NOV2020:12:08 khcs324

Table 2.3.3.4 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2			Placebo + bevacizumab (N=132)						
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI) (months))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment outo	ome (IVI	RS)									
NED [PDS]	92	35 (38.0)	NE (NE, NE)	48	25 (52.1)	11.3 (5.6,	NE)	0.60	0.36,	1.01	0.0525
NED/CR [IDS]	74	47 (63.5)	5.7 (3.0, 8.7)	38	15 (39.5)	NE (NE,	NE)	2.34	1.34,	4.33	0.0023*
NED/CR [Chemo]	40	22 (55.0)	8.3 (3.1, NE)	20	12 (60.0)	12.6 (5.7,	NE)	1.18	0.59,	2.46	0.6456
PR	49	30 (61.2)	12.1 (6.2,22.1)	26	13 (50.0)	17.0 (3.0,	NE)	1.03	0.55,	2.04	0.9310
Interaction p-value											0.0059*
Screening laboratory tBRC	!A statu:	s (IVRS)									
tBRCAm	150	79 (52.7)	13.7 (8.3, NE)	65	33 (50.8)	19.4 (8.2,	NE)	1.07	0.72,	1.63	0.7322
non-tBRCAm	105	55 (52.4)	12.1 (6.0, NE)	67	32 (47.8)	17.0 (8.3,	NE)	1.16	0.76,	1.82	0.4935
Interaction p-value											0.7907
First line treatment outo	ome (eCI	RF)									
NED [PDS]	89	37 (41.6)	NE (NE, NE)	47	25 (53.2)	10.7 (3.0,	NE)	0.64	0.38,	1.07	0.0855
NED/CR [IDS]	74	43 (58.1)	7.9 (5.4,24.0)	32	12 (37.5)	NE (NE,	NE)	2.06	1.12,	4.09	0.0185*
NED/CR [Chemo]	39	19 (48.7)	11.2 (4.9, NE)	18	10 (55.6)	13.3 (5.5,	NE)	1.00	0.47,	2.23	0.9933
PR	50	34 (68.0)	8.4 (5.7,13.7)	34	18 (52.9)	17.0 (8.3,	NE)	1.39	0.79,	2.51	0.2551
Interaction p-value											0.0285*
Screening laboratory tBRC	'A statu	s (eCRF)									
tBRCAm	147	78 (53.1)	13.7 (8.3, NE)	67	33 (49.3)	19.4 (8.2,	NE)	1.11	0.75,	1.69	0.6103
non-tBRCAm	108	56 (51.9)	12.1 (6.0, NE)	65	32 (49.2)	13.9 (8.3,	NE)	1.11	0.73,	1.74	0.6281
Interaction p-value											0.9954
Age group											
<65 years	185	97 (52.4)	12.3 (8.3, NE)	98	47 (48.0)	22.1 (11.2,	NE)	1.15	0.82,	1.65	0.4163
>=65 years	70	37 (52.9)	11.5 (5.7, NE)	34	18 (52.9)	12.3 (6.4,	NE)	0.98	0.57,	1.76	0.9476
Interaction p-value											0.6326

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bad 25NOV2020:12:08 khcs324

Table 2.3.3.4 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

-			bevacizumab 255)			oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	94 (51.6)	13.8 (8.4, NE)	90	43 (47.8)	22.1 (11.3,	NE)	1.18	0.83, 1.71	0.3533
IV	73	40 (54.8)	8.5 (5.7, NE)	42	22 (52.4)	11.2 (5.6,	NE)	0.96	0.57, 1.64	0.8661
Interaction p-value										0.5091
Region										
Europe	245	129 (52.7)	12.2 (8.3,24.1)	126	63 (50.0)	17.0 (10.2,	NE)	1.09	0.81, 1.48	0.5805
Japan	10	5 (50.0)	22.1 (2.8, NE)	6	2 (33.3)	NE (NE,	NE)	1.68	0.36, 11.70	0.5236
Interaction p-value										0.6022
ECOG performance status at	Basel	ine								
(0) Normal activity	190	102 (53.7)	11.2 (8.3,24.9)	100	52 (52.0)	17.0 (6.6,	NE)	1.07	0.77, 1.51	0.6879
(1) Restricted activity	61	30 (49.2)	22.5 (8.3, NE)	31	13 (41.9)	NE (NE,	NE)	1.27	0.68, 2.52	0.4674
Interaction p-value										0.6475
Baseline CA-125 value										
<=ULN	228	119 (52.2)	12.3 (8.3,24.9)	118	60 (50.8)	17.0 (8.5,	NE)	1.06	0.78, 1.45	0.7246
>ULN	27	15 (55.6)	11.2 (5.3, NE)	14	5 (35.7)	NE (NE,	NE)	1.72	0.67, 5.29	0.2750
Interaction p-value										0.3540
Histological grade										
High grade	255	134 (52.5)	12.2 (8.3,24.1)	132	65 (49.2)	17.5 (11.2,	NE)	1.11	0.83, 1.50	0.4970
Interaction p-value										NC
Cytoreductive surgery outco	me									
No residue	166	81 (48.8)	22.5 (8.4, NE)	80	37 (46.3)	22.1 (6.6,	NE)	1.06	0.73, 1.59	0.7564
Residue	79	46 (58.2)	11.1 (5.7,22.1)	44	24 (54.5)	17.0 (8.3,	NE)	1.21	0.74, 2.01	0.4522
Interaction p-value										0.6934

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bad 25NOV2020:12:08 khcs324

Table 2.3.3.4 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	•	Olaparib + k (N=2				oevacizumab 132)					
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]	Number (%) of patients n with events				Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
Timing of cytoreductive s	urgery										
Upfront	146	69 (47.3)	24.1 (11.6, NE)	79	41 (51.9)	12.5 (8.2,	NE)	0.84	0.57,	1.24	0.3763
Interval	99	58 (58.6)	8.1 (5.6,12.2)	45	20 (44.4)	23.5 (8.5,	NE)	1.71	1.05,	2.91	0.0320*
Interaction p-value											0.0268*
Myriad tumour BRCA mutati	on stat	us									
tBRCAm	158	83 (52.5)	13.7 (8.3, NE)	77	38 (49.4)	19.4 (8.2,	NE)	1.07	0.73,	1.58	0.7377
Non-tBRCAm	97	51 (52.6)	12.1 (6.2, NE)	55	27 (49.1)	13.9 (8.3,	NE)	1.17	0.74,	1.89	0.5012
Interaction p-value											0.7614
Status somatic BRCA mutat	ions										
sBRCAm	22	9 (40.9)	24.1 (5.8, NE)	7	2 (28.6)	NE (NE,	NE)	1.71	0.44,	11.23	0.4679
gBRCAm	66	42 (63.6)	8.3 (5.6,14.5)	31	11 (35.5)	NE (NE,	NE)	2.23	1.19,	4.56	0.0113*
Non-BRCAm	41	17 (41.5)	NE (NE, NE)	22	12 (54.5)	12.3 (5.6,	NE)	0.76	0.37,	1.63	0.4734
Interaction p-value											0.1031

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bad 25NoV2020:12:08 khcs324

Table 2.3.3.5 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Hormonal time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	·		bevacizumab 255)		Placebo + h (N=1	pevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]) ratio		2-sided p-value [b]
First line treatment out	come (IV	RS)							
NED [PDS]	92	50 (54.3)	17.0 (13.8,33.4)	48	29 (60.4)	8.5 (5.5,19.3)	0.69	0.44, 1.10	0.1124
NED/CR [IDS]	74	39 (52.7)	19.3 (11.1, NE)	38	18 (47.4)	19.4 (5.6, NE)	1.07	0.62, 1.92	0.8006
NED/CR [Chemo]	40	20 (50.0)	21.9 (14.0, NE)	20	14 (70.0)	11.0 (2.9,24.6)	0.61	0.31, 1.23	0.1587
PR	49	26 (53.1)	19.5 (11.2,30.4)	26	15 (57.7)	5.6 (2.9,19.9)	0.62	0.33, 1.20	0.1480
Interaction p-value									0.4717
Screening laboratory tBRG	CA statu	ıs (IVRS)							
tBRCAm	150	83 (55.3)	19.3 (13.9,24.2)	65	37 (56.9)	11.8 (5.6,24.1)	0.79	0.54, 1.18	0.2472
non-tBRCAm	105	52 (49.5)	18.6 (12.1, NE)	67	39 (58.2)	11.1 (5.6,19.3)	0.69	0.46, 1.06	0.0873
Interaction p-value									0.6425
First line treatment out	come (eC	CRF)							
NED [PDS]	89	46 (51.7)	18.9 (13.8, NE)	47	29 (61.7)	6.2 (5.5,19.3)	0.61	0.38, 0.97	0.0388*
NED/CR [IDS]	74	40 (54.1)	16.8 (8.7,25.3)	32	15 (46.9)	19.4 (5.6, NE)	1.09	0.62, 2.04	0.7742
NED/CR [Chemo]	39	19 (48.7)	21.9 (14.3, NE)	18	10 (55.6)	13.9 (3.0, NE)	0.74	0.35, 1.66	0.4521
PR	50	28 (56.0)	19.3 (10.9,30.4)	34	22 (64.7)	11.1 (4.7,19.9)	0.69	0.39, 1.21	0.1937
Interaction p-value									0.4754
Screening laboratory tBR0	CA statu	ıs (eCRF)							
tBRCAm	147	81 (55.1)	19.2 (13.9,24.2)	67	38 (56.7)	11.3 (5.6,24.1)	0.78	0.54, 1.16	0.2201
non-tBRCAm	108	54 (50.0)	19.1 (12.1,33.4)	65	38 (58.5)	11.1 (5.6,19.4)	0.70	0.46, 1.07	0.1001
Interaction p-value									0.7082
Age group									
<65 years	185	101 (54.6)	17.5 (13.9,22.1)	98	60 (61.2)	8.4 (5.5,13.9)	0.68	0.50, 0.95	0.0227*
>=65 years	70	34 (48.6)	24.0 (11.5, NE)	34	16 (47.1)	19.9 (5.7, NE)	0.97	0.54, 1.80	0.9188
Interaction p-value									0.3066

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bae 25NOV2020:12:08 khcs324

Table 2.3.3.5 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Hormonal time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1				oevacizumab 132)			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	95% CI) of patients		Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [}	2-sided o] p-value[b]
FIGO Stage (Disease state)									
III	182	99 (54.4)	16.8 (13.9,24.0)	90	52 (57.8)	11.3 (5.7,19.4)	0.81	0.58, 1	.14 0.2241
IV	73	36 (49.3)	22.3 (13.9, NE)	42	24 (57.1)	13.9 (3.0,24.6)	0.61	0.37, 1	.04 0.0706
Interaction p-value									0.3820
Region									
Europe	245	130 (53.1)	19.1 (14.0,24.2)	126	72 (57.1)	11.3 (5.7,16.6)	0.75	0.57, 1	
Japan	10	5 (50.0)	22.1 (2.8, NE)	6	4 (66.7)	12.5 (2.9, NE)	0.62	0.16, 2	.52 0.4866
Interaction p-value									0.7836
ECOG performance status at	Basel	ine							
(0) Normal activity	190	94 (49.5)	20.0 (16.0,30.4)	100	60 (60.0)	11.1 (5.6,19.3)	0.68	0.49, 0	.95 0.0226*
(1) Restricted activity	61	37 (60.7)	11.5 (8.4,24.0)	31	16 (51.6)	11.3 (5.6, NE)	0.94	0.53, 1	.73 0.8269
Interaction p-value									0.3471
Baseline CA-125 value									
<=ULN	228	124 (54.4)	17.5 (13.9,24.0)	118	66 (55.9)	12.3 (5.7,19.4)	0.82	0.61, 1	
>ULN	27	11 (40.7)	25.3 (8.5,25.3)	14	10 (71.4)	5.6 (2.8,24.6)	0.33	0.14, 0	.79 0.0142*
Interaction p-value									0.0543
Histological grade									
High grade	255	135 (52.9)	19.1 (14.3,24.2)	132	76 (57.6)	11.3 (5.6,19.1)	0.75	0.57, 0	.99 0.0460*
Interaction p-value									NC
Cytoreductive surgery outco	ome								
No residue	166	88 (53.0)	17.5 (13.8,25.3)	80	44 (55.0)	11.8 (5.6,22.3)	0.79	0.55, 1	.14 0.1994
Residue	79	42 (53.2)	21.9 (14.3,25.5)	44	28 (63.6)	11.3 (5.6,19.1)	0.68	0.42, 1	.11 0.1233
Interaction p-value									0.6431

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bae 25NOV2020:12:08 khcs324

Table 2.3.3.5 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Hormonal time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)				
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]	Number (%) of patients n with events		,	Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	77 (52.7)	19.1 (14.0,30.4)	79	49 (62.0)	8.7 (5.6,16.4)	0.64	0.45,	0.92	0.0174*
Interval	99	53 (53.5)	19.3 (11.1,24.4)	45	23 (51.1)	14.7 (5.6, NE)	0.97	0.60,	1.62	0.9141
Interaction p-value										0.1740
Myriad tumour BRCA mutati	ion stat	us								
tBRCAm	158	87 (55.1)	19.3 (13.9,24.2)	77	42 (54.5)	11.3 (5.6,24.1)	0.81	0.56,	1.18	0.2711
Non-tBRCAm	97	48 (49.5)	18.6 (12.1, NE)	55	34 (61.8)	11.1 (5.6,19.3)	0.66	0.43,	1.03	0.0695
Interaction p-value										0.4860
Status somatic BRCA mutat	cions									
sBRCAm	22	9 (40.9)	24.4 (8.3, NE)	7	2 (28.6)	NE (NE, NE)	1.05	0.27,	6.92	0.9455
gBRCAm	66	42 (63.6)	11.2 (5.8,19.5)	31	18 (58.1)	8.5 (4.2,24.1)	0.94	0.55,	1.68	0.8286
Non-BRCAm	41	22 (53.7)	16.6 (7.8, NE)	22	16 (72.7)	12.3 (4.7,19.3)	0.63	0.33,	1.23	0.1706
Interaction p-value										0.6185

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[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bae 25NoV2020:12:08 khcs324

Table 2.3.3.6 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bevacizumab (N=255)							oevacizumab 132)					
Subgroup	Number (%) of patients n with events		S	Median time (95% CI) (months) [a]		Number (%) of patients n with events		Median time (95% CI) (months) [a]		Hazard ratio [b]	95% CI [b]		2-sided p-value [b]	
First line treatment outo	ome (IV	RS)												
NED [PDS]	92	33 (35.9) NE	(NE,	NE)	48	22 (4	5.8)	16.4 (8.4,	NE)	0.63	0.37,	1.10	0.1055
NED/CR [IDS]	74	37 (50.0) 18.6	(13.8,	NE)	38	13 (3	4.2)	NE (NE,	NE)	1.62	0.88,	3.16	0.1218
NED/CR [Chemo]	40	21 (52.5) 19.4	(3.0,	NE)	20	11 (5	5.0)	17.2 (3.0,	NE)	1.07	0.52,	2.30	0.8612
PR	49	23 (46.9) 25.3	(11.1,	NE)	26	12 (4	6.2)	12.7 (3.0,	NE)	0.80	0.41,	1.67	0.5401
Interaction p-value														0.1485
Screening laboratory tBRC	!A statu	s (IVRS)												
tBRCAm	150	73 (48.7) 22.1	(14.5,	NE)	65	33 (5	0.8)	13.9 (8.3,	NE)	0.87	0.58,	1.33	0.5024
non-tBRCAm	105	41 (39.0) NE	(NE,	NE)	67	25 (3	7.3)	NE (NE,	NE)	1.00	0.62,	1.67	0.9865
Interaction p-value														0.6566
First line treatment outc	ome (eC	RF)												
NED [PDS]	89	30 (33.7) NE	(NE,	NE)	47	21 (4	4.7)	16.9 (8.3,	NE)	0.61	0.35,	1.09	0.0931
NED/CR [IDS]	74	37 (50.0) 16.6	(11.5,	NE)	32	11 (3	4.4)	NE (NE,	NE)	1.62	0.86,	3.34	0.1422
NED/CR [Chemo]	39	24 (61.5) 13.7	(3.0,	24.2)	18	10 (5	5.6)	17.2 (3.0,	NE)	1.19	0.58,	2.61	0.6425
PR	50	21 (42.0) 25.7	(11.1,	NE)	34	16 (4	7.1)	17.0 (8.3,	NE)	0.77	0.41,	1.50	0.4416
Interaction p-value														0.1290
Screening laboratory tBRC	'A statu	s (eCRF)												
tBRCAm	147	71 (48.3) 22.2	(14.5,	NE)	67	33 (4	9.3)	16.9 (8.3,	NE)	0.88	0.59,	1.34	0.5382
non-tBRCAm	108	43 (39.8) NE	(NE,	NE)	65	25 (3	8.5)	NE (NE,	NE)	1.01	0.62,	1.67	0.9767
Interaction p-value														0.6736
Age group														
<65 years	185	90 (48.6) 22.2	(15.9,	NE)	98	46 (4	6.9)	17.2 (10.4,	NE)	0.95	0.67,	1.36	0.7637
>=65 years	70	24 (34.3) NE	(NE,	NE)	34	12 (3	5.3)	NE (NE,	NE)	0.96	0.49,	1.99	0.9099
Interaction p-value														0.9708

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3baf 25NoV2020:12:08 khcs324

Table 2.3.3.6 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2					oevacizumab 132)					
Subgroup	n	Number (%) of patients with events	(95% CI	Median time (95% CI) (months) [a]		Number (%) of patients with events			Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	79 (43.4)	25.7 (19.4,	NE)	90	39 (43.3)	NE (NE,	NE)	0.95	0.65,	1.41	0.7936
IV	73	35 (47.9)	19.6 (9.0,	NE)	42	19 (45.2)	17.5 (8.7,	NE)	0.96	0.56,	1.71	0.8834
Interaction p-value												0.9781
Region												
Europe	245	110 (44.9)	25.3 (18.0,	NE)	126	56 (44.4)	21.9 (11.9,	NE)	0.92	0.67,	1.28	0.6165
Japan	10	4 (40.0)	NE (NE,	NE)	6	2 (33.3)	NE (NE,	NE)	1.78	0.35,	12.82	0.4957
Interaction p-value												0.4436
ECOG performance status at	Basel	ine										
(0) Normal activity	190	87 (45.8)	24.2 (15.9,	NE)	100	46 (46.0)	21.9 (11.0,	NE)	0.97	0.68,	1.40	0.8810
(1) Restricted activity	61	25 (41.0)	25.7 (16.6,	NE)	31	12 (38.7)	NE (NE,	NE)	0.89	0.46,	1.85	0.7529
Interaction p-value												0.8324
Baseline CA-125 value												
<=ULN	228	100 (43.9)	NE (NE,	NE)	118	51 (43.2)	NE (NE,	NE)	0.93	0.67,	1.31	0.6756
>ULN	27	14 (51.9)	19.6 (5.6,	25.3)	14	7 (50.0)	17.0 (6.6,	NE)	1.14	0.47,	3.00	0.7781
Interaction p-value												0.6804
Histological grade												
High grade	255	114 (44.7)	25.3 (18.6,	NE)	132	58 (43.9)	23.0 (12.7,	NE)	0.95	0.70,	1.31	0.7478
Interaction p-value												NC
Cytoreductive surgery outco	me											
No residue	166	69 (41.6)	NE (NE,	NE)	80	32 (40.0)	NE (NE,	NE)	0.97	0.64,	1.49	0.8808
Residue	79	40 (50.6)	22.1 (8.9,	NE)	44	21 (47.7)	17.2 (6.0,	NE)	1.04	0.62,	1.79	0.8981
Interaction p-value												0.8465

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[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3baf 25NoV2020:12:08 khcs324

Table 2.3.3.6 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	•	Olaparib + 1 (N=2				bevacizumab 132)						
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	urgery											
Upfront	146	58 (39.7)	NE (NE,	NE)	79	35 (44.3)	23.0 (8.6,	NE)	0.77	0.51,	1.18	0.2307
Interval	99	51 (51.5)	16.6 (11.5,	NE)	45	18 (40.0)	NE (NE,	NE)	1.41	0.84,	2.47	0.2020
Interaction p-value												0.0806
Myriad tumour BRCA mutati	on stat	us										
tBRCAm	158	77 (48.7)	22.1 (14.1,	NE)	77	34 (44.2)	17.5 (8.7,	NE)	1.02	0.69,	1.55	0.9132
Non-tBRCAm	97	37 (38.1)	NE (NE,	NE)	55	24 (43.6)	NE (NE,	NE)	0.82	0.49,	1.39	0.4545
Interaction p-value												0.5099
Status somatic BRCA mutat	ions											
sBRCAm	22	10 (45.5)	8.5 (2.8,	NE)	7	3 (42.9)	NE (NE,	NE)	1.40	0.43,	6.24	0.5995
gBRCAm	66	32 (48.5)	24.2 (13.9,	NE)	31	14 (45.2)	17.2 (8.7,	NE)	0.89	0.49,	1.73	0.7254
Non-BRCAm	41	19 (46.3)	25.7 (13.7,	NE)	22	9 (40.9)	NE (NE,	NE)	1.11	0.52,	2.59	0.7881
Interaction p-value												0.7942

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3baf 25NOV2020:12:08 khcs324

Table 2.3.3.7 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Other single items time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	(Olaparib + h (N=2				oevacizumab 132)			
Subgroup	0	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)							
NED [PDS]	92	40 (43.5)	NE (NE, NE)	48	22 (45.8)	19.4 (13.8, NE)	0.97	0.58, 1.66	0.9172
NED/CR [IDS]	74	43 (58.1)	14.3 (8.3,24.0)	38	18 (47.4)	19.6 (9.3, NE)	1.42	0.83, 2.53	0.2007
NED/CR [Chemo]	40	20 (50.0)	21.9 (11.3, NE)	20	8 (40.0)	NE (NE, NE)	1.34	0.61, 3.23	0.4789
PR	49	24 (49.0)	19.6 (11.9, NE)	26	16 (61.5)	16.6 (4.9,24.9)	0.65	0.35, 1.25	0.1899
Interaction p-value									0.2895
Screening laboratory tBR0	CA status	(IVRS)							
tBRCAm	150	80 (53.3)	20.0 (14.5,25.3)	65	31 (47.7)	19.6 (13.9, NE)	1.14	0.76, 1.75	0.5309
non-tBRCAm	105	47 (44.8)	25.7 (11.1, NE)	67	33 (49.3)	19.4 (13.8, NE)	0.95	0.61, 1.50	0.8311
Interaction p-value									0.5619
First line treatment out	come (eCR	F)							
NED [PDS]	89	37 (41.6)	NE (NE, NE)	47	21 (44.7)	19.4 (13.8, NE)	0.96	0.57, 1.67	0.8875
NED/CR [IDS]	74	41 (55.4)	14.3 (8.3,27.7)	32	16 (50.0)	19.4 (9.3, NE)	1.26	0.72, 2.31	0.4238
NED/CR [Chemo]	39	22 (56.4)	19.6 (8.3,25.0)	18	7 (38.9)	NE (NE, NE)	1.50	0.67, 3.80	0.3319
PR	50	25 (50.0)	25.3 (11.9, NE)	34	20 (58.8)	18.7 (8.3,28.0)	0.76	0.42, 1.38	0.3561
Interaction p-value									0.4953
Screening laboratory tBR0	CA status	(eCRF)							
tBRCAm	147	78 (53.1)	20.7 (14.5,25.3)	67	32 (47.8)	19.5 (15.4, NE)	1.13	0.76, 1.73	0.5626
non-tBRCAm	108	49 (45.4)	25.7 (13.8, NE)	65	32 (49.2)	19.4 (13.8, NE)	0.96	0.62, 1.52	0.8693
Interaction p-value									0.6094
Age group									
<65 years	185	95 (51.4)	19.6 (14.3,25.7)	98	46 (46.9)	22.2 (16.6, NE)	1.19	0.84, 1.70	0.3311
>=65 years	70	32 (45.7)	27.7 (13.7, NE)	34	18 (52.9)	14.0 (6.1, NE)	0.74	0.42, 1.35	0.3238
Interaction p-value									0.1802

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[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bag 25NOV2020:12:08 khcs324

Table 2.3.3.7 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Other single items time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2				Placebo + (N=	bevacizuma 132)	ıb				
Subgroup	n	Number (%) of patients with events	Median t (95% CI (months)	:)		Number (%) of patients with events		CI)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	86 (47.3)	24.0 (16.7,	NE)	90	38 (42.2)	NE (I	NE, NE)	1.20	0.83,	1.78	0.3425
IV	73	41 (56.2)	17.5 (8.3,	25.3)	42	26 (61.9)	15.4 (8	.3,22.2)	0.84	0.52,	1.38	0.4798
Interaction p-value												0.2565
Region												
Europe	245	123 (50.2)	21.9 (16.6,	25.7)	126	60 (47.6)	19.4 (16	.4, NE)	1.07	0.79,	1.47	0.6495
Japan	10	4 (40.0)	NE (NE,	NE)	6	4 (66.7)	20.8 (2	.8, NE)	0.69	0.16,	2.92	0.6030
Interaction p-value												0.5449
ECOG performance status at	Basel	ine										
(0) Normal activity	190	97 (51.1)	19.6 (12.5,	27.7)	100	50 (50.0)	19.4 (16	.6, NE)	1.13	0.81,	1.61	0.4708
(1) Restricted activity	61	30 (49.2)	24.0 (16.0,	NE)	31	14 (45.2)	19.5 (10	.2, NE)	0.95	0.51,	1.85	0.8811
Interaction p-value												0.6394
Baseline CA-125 value												
<=ULN	228	116 (50.9)	20.7 (14.6,	25.7)	118	57 (48.3)	19.4 (16	.6, NE)	1.11	0.81,	1.53	0.5319
>ULN	27	11 (40.7)	25.3 (9.0,	NE)	14	7 (50.0)	11.3 (5	.5, NE)	0.67	0.26,	1.83	0.4184
Interaction p-value												0.3366
Histological grade												
High grade	255	127 (49.8)	21.9 (16.6,	25.7)	132	64 (48.5)	19.4 (16	.4, NE)	1.05	0.78,	1.43	0.7355
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	80 (48.2)	23.2 (14.1,	NE)	80	37 (46.3)	19.5 (16	.4, NE)	1.12	0.76,	1.67	0.5677
Residue	79	40 (50.6)	21.9 (14.6,	NE)	44	22 (50.0)	23.5 (11	.2,28.0)	0.97	0.58,	1.65	0.8953
Interaction p-value												0.6563

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[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bag 25NoV2020:12:08 khcs324

Table 2.3.3.7 PAOLA1: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Other single items time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	•	Olaparib + b (N=2				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	urgery									
Upfront	146	69 (47.3)	25.0 (16.7, NE)	79	35 (44.3)	19.5 (16.4, NE)	1.07	0.71,	1.62	0.7582
Interval	99	51 (51.5)	20.0 (12.5, NE)	45	24 (53.3)	19.4 (9.3, NE)	1.05	0.65,	1.73	0.8523
Interaction p-value										0.9563
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	81 (51.3)	21.9 (16.0,27.7)	77	33 (42.9)	23.5 (16.6, NE)	1.22	0.82,	1.85	0.3367
Non-tBRCAm	97	46 (47.4)	25.7 (9.0, NE)	55	31 (56.4)	16.7 (10.2,28.0)	0.88	0.56,	1.40	0.5848
Interaction p-value										0.2981
Status somatic BRCA mutat	ions									
sBRCAm	22	11 (50.0)	16.7 (3.6, NE)	7	3 (42.9)	NE (NE, NE)	1.39	0.43,	6.16	0.5997
gBRCAm	66	37 (56.1)	16.6 (11.3,27.7)	31	12 (38.7)	NE (NE, NE)	1.57	0.84,	3.15	0.1588
Non-BRCAm	41	22 (53.7)	19.5 (3.0, NE)	22	17 (77.3)	9.7 (5.6,13.9)	0.63	0.34,	1.21	0.1640
Interaction p-value										0.1276

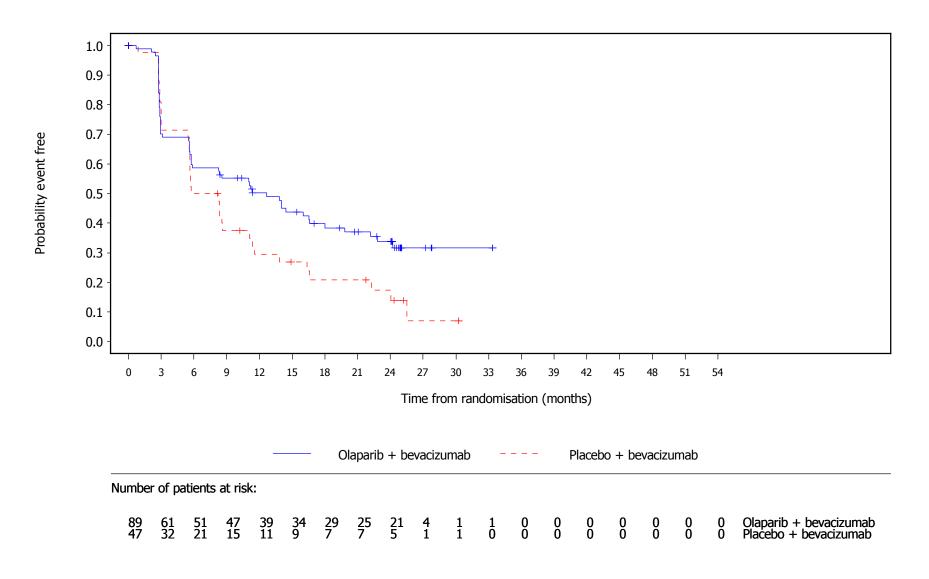
Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3bag 25NoV2020:12:08 khcs324

Figure 2.3.4.1 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI time to clinically meaningful deterioration for First line treatment outcome (eCRF)=NED [PDS]

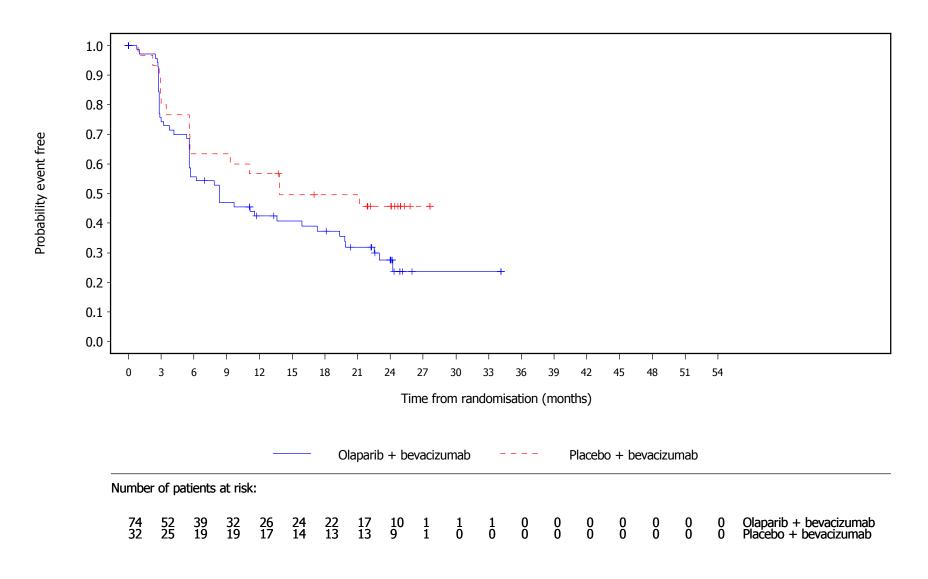
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3eaa 25NOV2020:12:08 khcs324

Figure 2.3.4.2 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI time to clinically meaningful deterioration for First line treatment outcome (eCRF)=NED/CR [IDS]

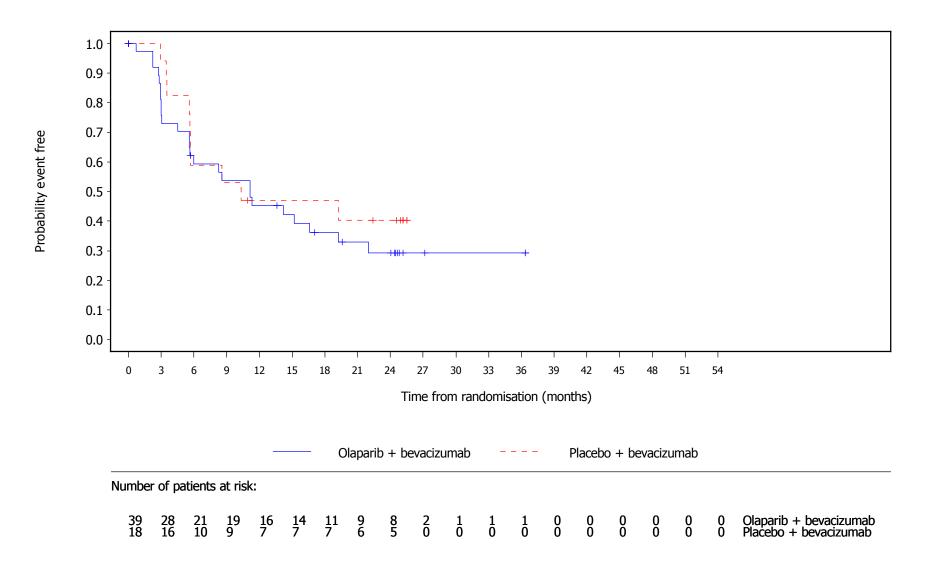
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.3.4.3 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI time to clinically meaningful deterioration for First line treatment outcome (eCRF)=NED/CR [Chemo]

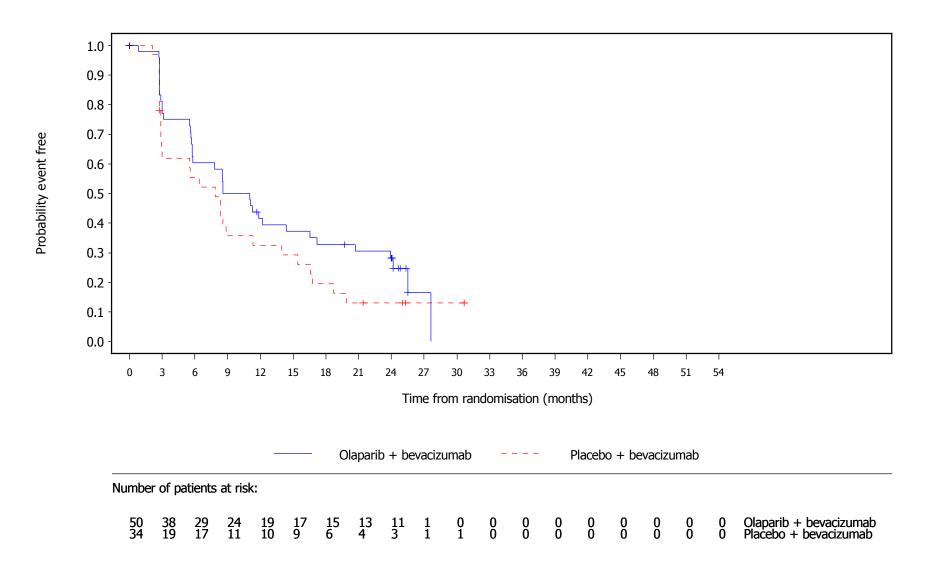
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.3.4.4 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI time to clinically meaningful deterioration for First line treatment outcome (eCRF)=PR

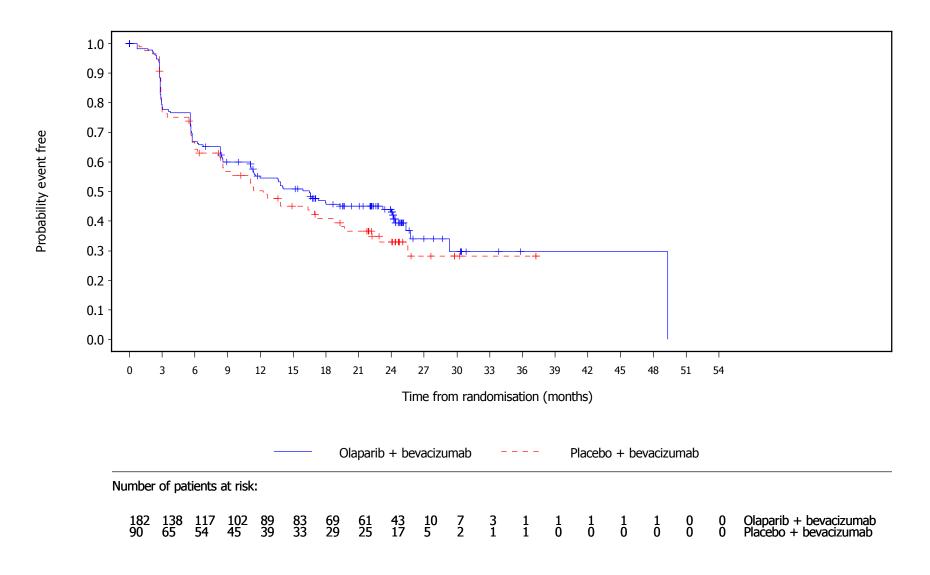
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.3.4.5 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects time to clinically meaningful deterioration for FIGO Stage (Disease state)=III

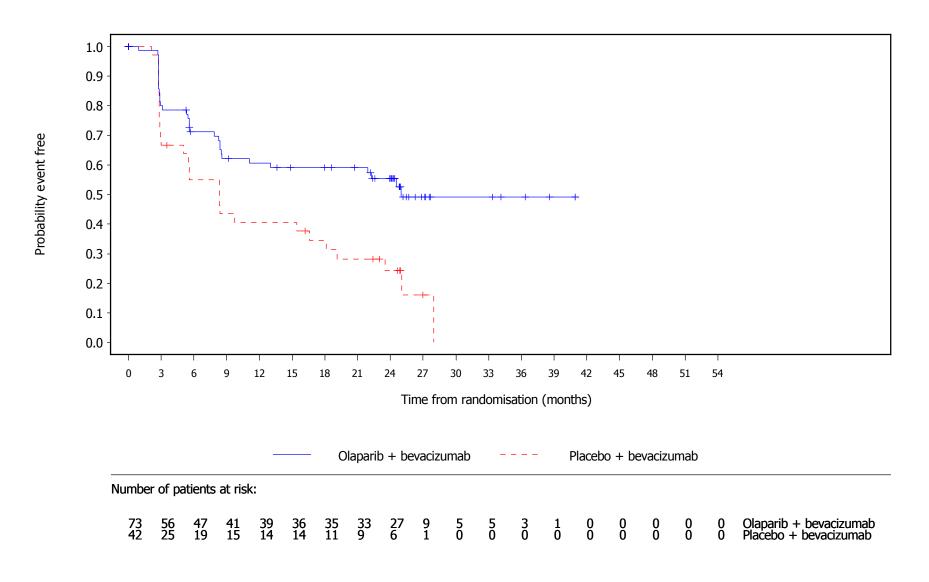
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3eae 25NOV2020:12:08 khcs324

Figure 2.3.4.6 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects time to clinically meaningful deterioration for FIGO Stage (Disease state)=IV

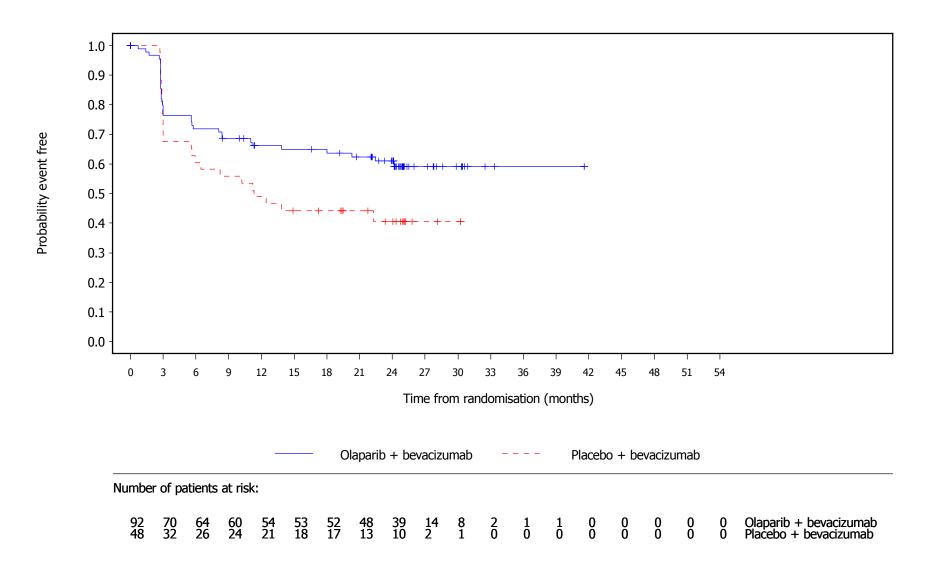
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.3.4.7 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration for First line treatment outcome (IVRS)=NED [PDS]

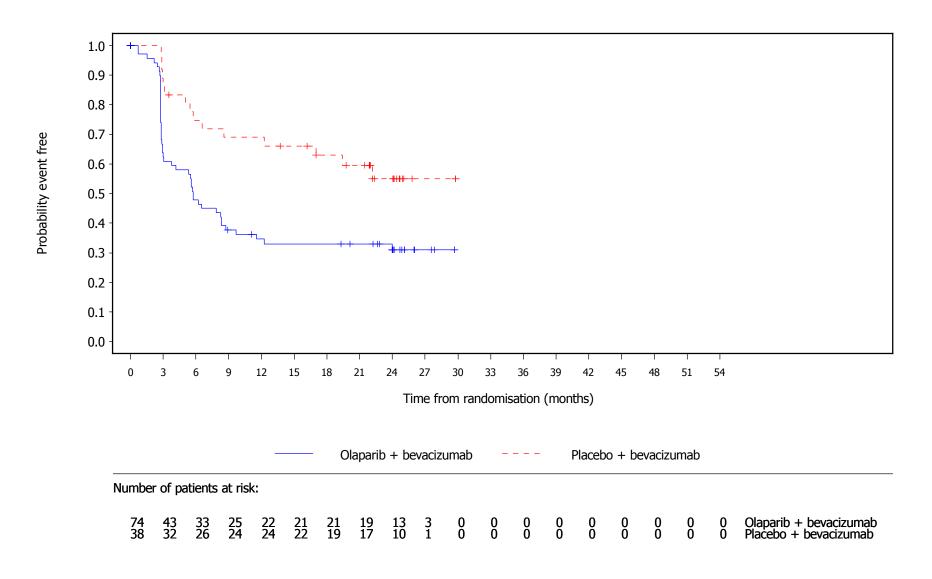
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.3.4.8 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration for First line treatment outcome (IVRS)=NED/CR [IDS]

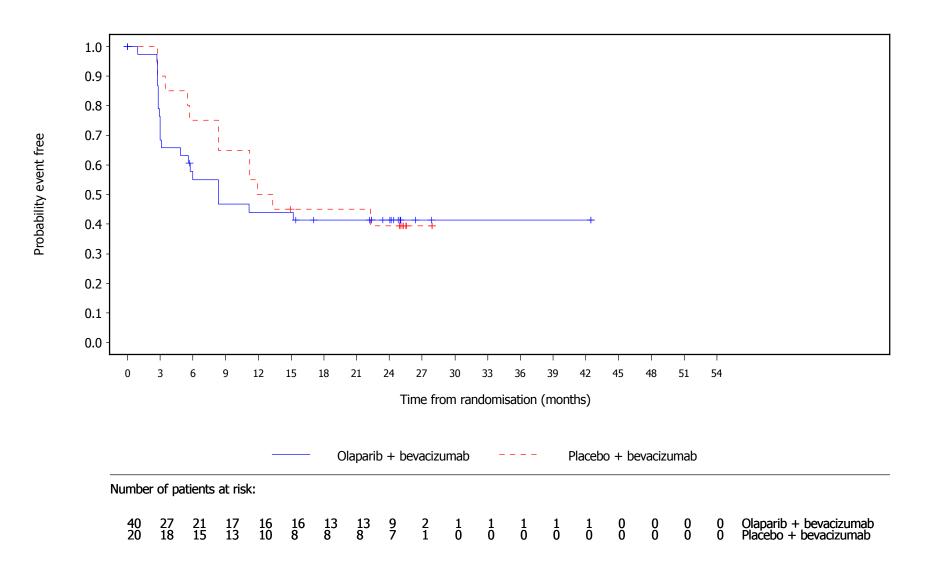
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.3.4.9 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration for First line treatment outcome (IVRS)=NED/CR [Chemo]

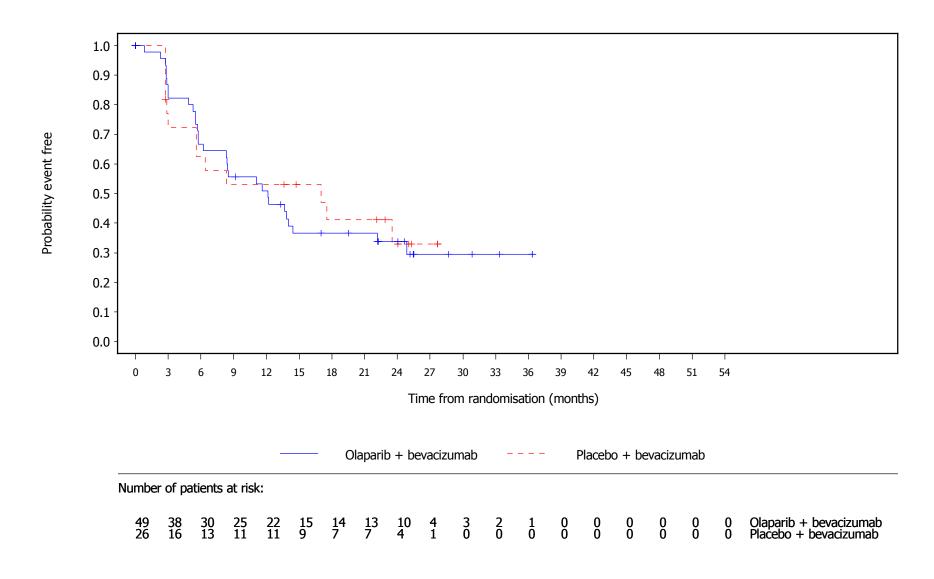
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.3.4.10 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration for First line treatment outcome (IVRS)=PR

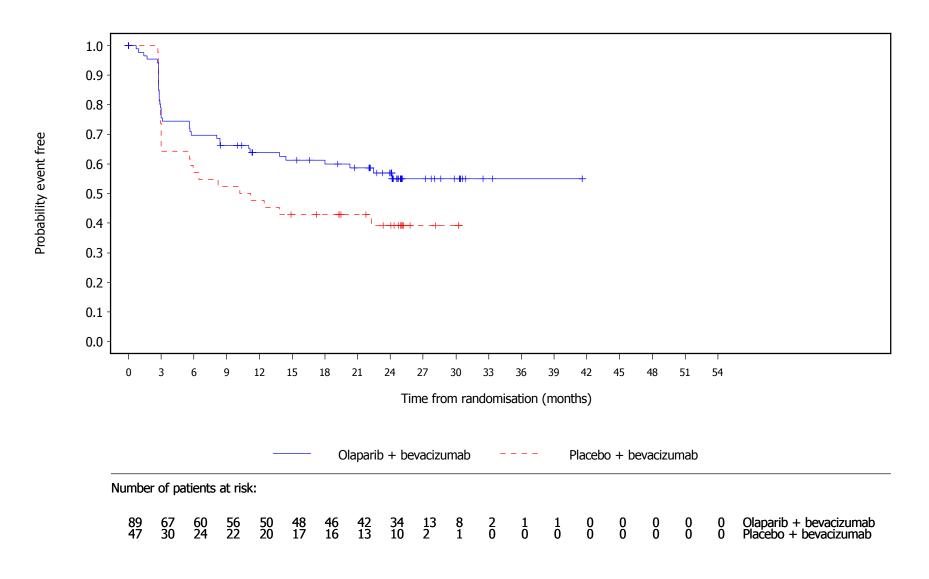
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.3.4.11 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration for First line treatment outcome (eCRF)=NED [PDS]

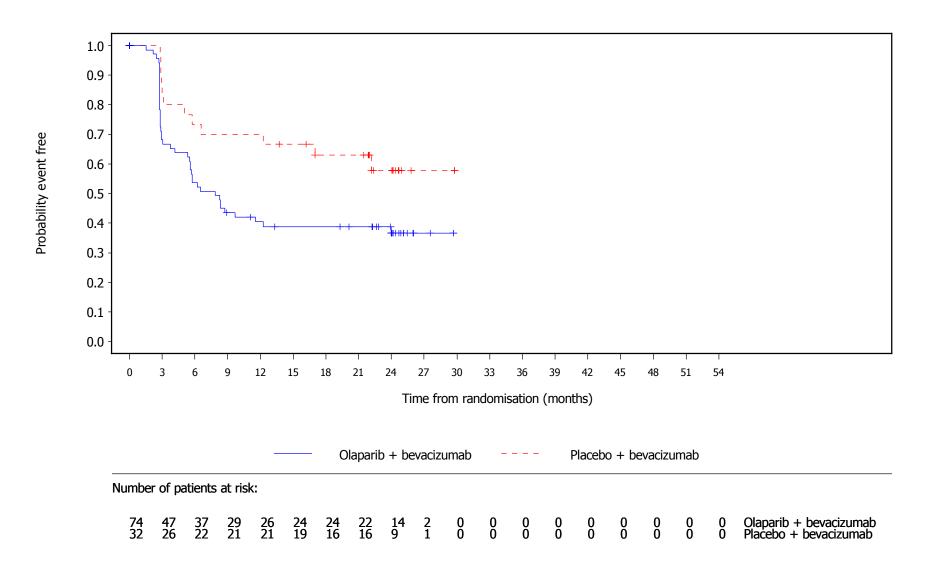
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.3.4.12 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration for First line treatment outcome (eCRF)=NED/CR [IDS]

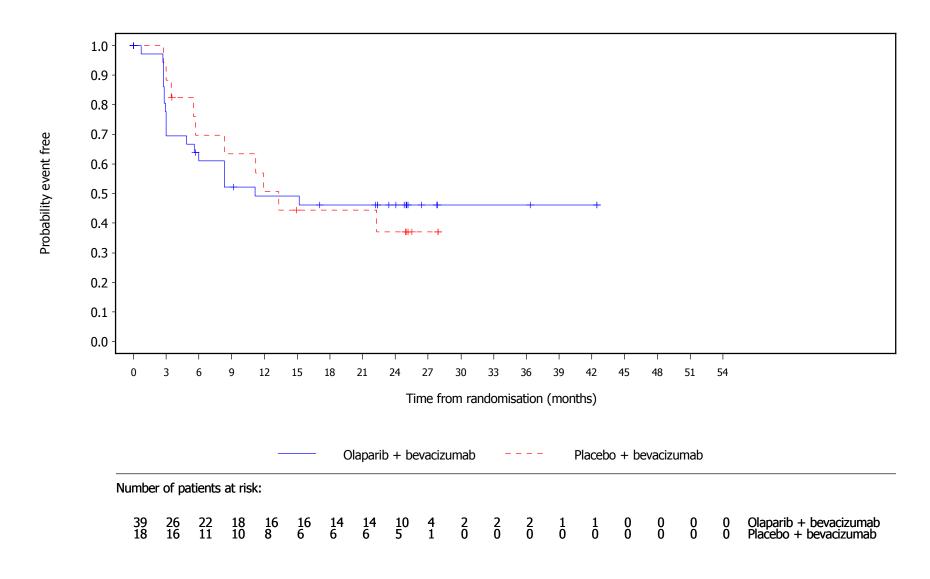
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.3.4.13 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration for First line treatment outcome (eCRF)=NED/CR [Chemo]

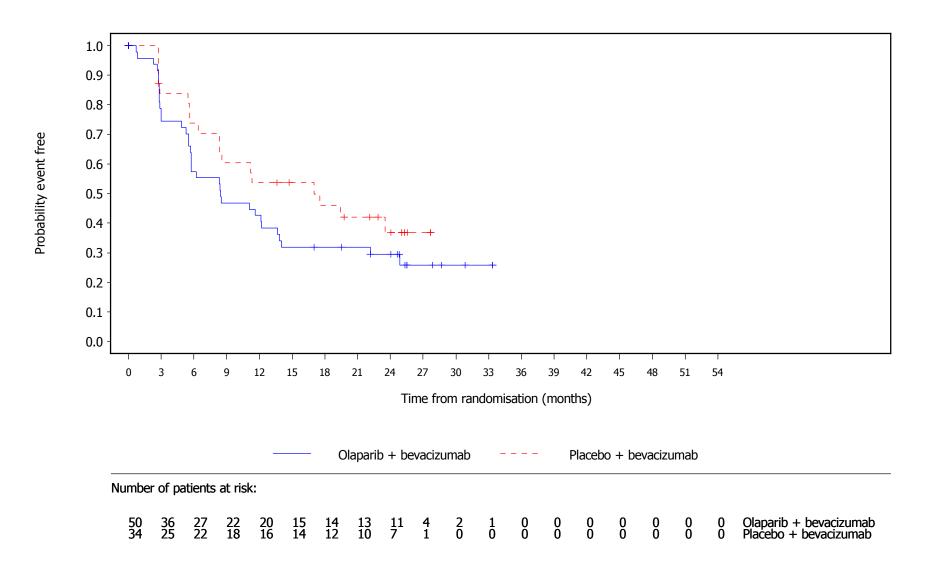
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3eam 25NOV2020:12:08 khcs324

Figure 2.3.4.14 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration for First line treatment outcome (eCRF)=PR

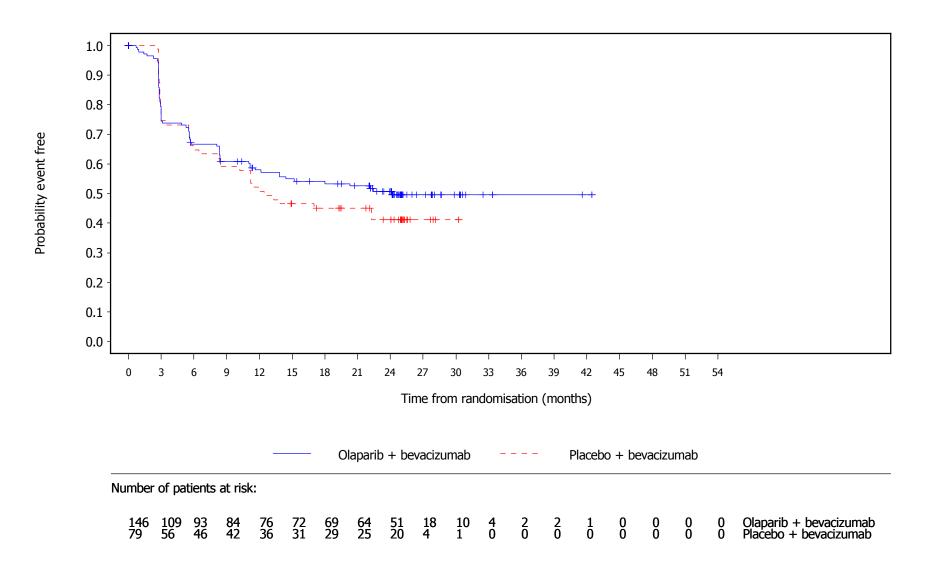
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



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Figure 2.3.4.15 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration for Timing of cytoreductive surgery=Upfront

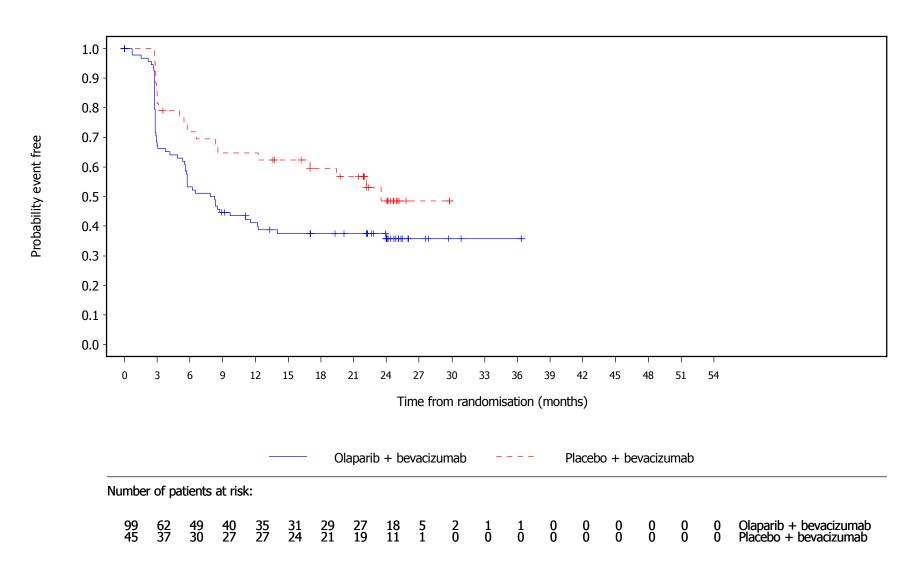
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3eao 25NOV2020:12:08 khcs324

Figure 2.3.4.16 PAOLA1: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment time to clinically meaningful deterioration for Timing of cytoreductive surgery=Interval

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3eap 25NOV2020:12:08 khcs324

Table 2.4.1.1 PAOLA1: Summary of analysis of time to worsening in EQ-5D VAS (MID = 10) Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)				pevacizumab 132)			
	Number (%) of patients n with events	Median time (95% CI) (months) [a]	of	Jumber (%) f patients ith events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
EQ-5D-5L Visual analogue scale (MID = 10)	255 156 (61.2)	11.1 (8.3,13.9)	132	78 (59.1)	16.4 (9.6,21.9)	1.15	0.87, 1.52	0.3464

Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation.

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Table 2.4.1.2 PAOLA1: Summary of analysis of time to worsening in EQ-5D VAS (MID = 7) Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + b (N=25			bevacizumab =132)			
	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	(95% CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
EQ-5D-5L Visual analogue scale (MID = 7)	255 156 (61.2)	11.1 (8.3,13.9)	132 78 (59.1)	16.4 (9.6,21.9)	1.15	0.88, 1.52	0.3326

Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation.

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

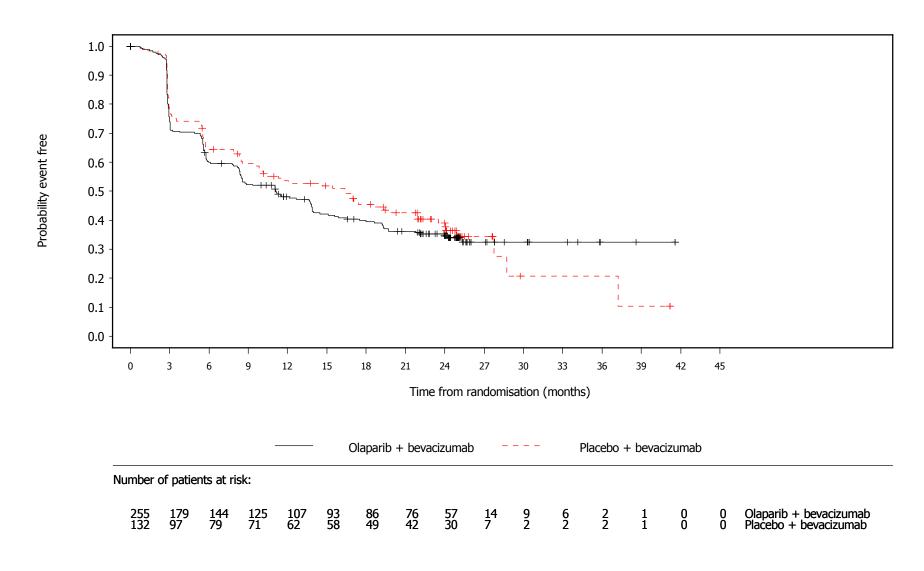
[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Figure 2.4.2.1 PAOLA1: Kaplan-Meier plot of EQ-5D-5L Visual analogue scale (MID = 10) time to clinically meaningful worsening (first occurrence)

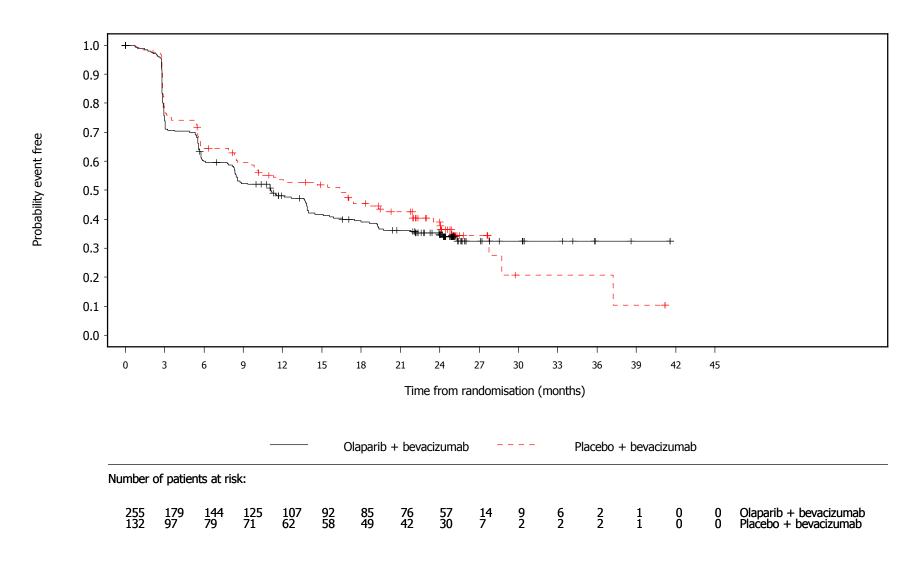
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr.sas ettemainprhaa 25NOV2020:15:58 kvbv306

Figure 2.4.2.2 PAOLA1: Kaplan-Meier plot of EQ-5D-5L Visual analogue scale (MID = 7) time to clinically meaningful worsening (first occurrence)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr.sas ettemainpriaa 25NOV2020:15:58 kvbv306

Table 2.4.3.1 PAOLA1: Summary of subgroup analysis of EQ-5D-5L Visual analogue scale (MID = 10) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	:		+ b N=25	evacizumab 55)			bevacizumab 132)				
Subgroup	n	Number (of patien with even	nts	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)									
NED [PDS]	92	47 (51	1)	13.8 (8.3, NE)	48	30 (62.5)	17.0 (8.5,27.8)	0.85	0.54,	1.36	0.5034
NED/CR [IDS]	74	49 (66	2)	11.1 (5.6,15.9)	38	19 (50.0)	14.7 (5.6, NE)	1.44	0.86,	2.51	0.1652
NED/CR [Chemo]	40	24 (60	0)	13.9 (5.6, NE)	20	14 (70.0)	10.0 (3.0,19.4)	0.84	0.44,	1.66	0.6023
PR	49	36 (73	5)	5.9 (3.0,13.7)	26	15 (57.7)	18.7 (2.9,24.9)	1.43	0.80,	2.70	0.2330
Interaction p-value											0.3113
Screening laboratory tBR0	CA stati	ıs (IVRS)									
tBRCAm	150	89 (59	3)	11.5 (8.3,15.9)	65	36 (55.4)	15.4 (8.5, NE)	1.15	0.79,	1.72	0.4729
non-tBRCAm	105	67 (63	8)	8.5 (5.7,13.9)	67	42 (62.7)	16.4 (5.6,21.9)	1.09	0.74,	1.61	0.6704
Interaction p-value											0.8381
First line treatment out	come (e0	CRF)									
NED [PDS]	89	48 (53	9)	13.6 (8.3, NE)	47	30 (63.8)	17.0 (8.3,24.1)	0.88	0.56,	1.41	0.5945
NED/CR [IDS]	74	49 (66	2)	11.1 (5.6,15.9)	32	16 (50.0)	12.1 (3.1, NE)	1.39	0.81,	2.53	0.2377
NED/CR [Chemo]	39	22 (56	4)	15.1 (5.6, NE)	18	11 (61.1)	10.3 (3.5, NE)	0.92	0.45,	1.96	0.8122
PR	50	36 (72	0)	6.0 (3.0,13.7)	34	21 (61.8)	14.7 (5.5,24.9)	1.34	0.79,	2.33	0.2871
Interaction p-value											0.5097
Screening laboratory tBR0	CA stati	ıs (eCRF)									
tBRCAm	147	87 (59	2)	11.4 (8.3,18.0)	67	36 (53.7)	15.4 (8.5, NE)	1.18	0.81,	1.77	0.3924
non-tBRCAm	108	69 (63	9)	9.3 (5.7,14.0)	65	42 (64.6)	16.4 (5.6,21.9)	1.04	0.71,	1.54	0.8237
Interaction p-value											0.6556
Age group											
<65 years	185	110 (59	5)	12.7 (8.4,15.9)	98	57 (58.2)	17.0 (9.6,24.9)	1.08	0.79,	1.50	0.6362
>=65 years	70	46 (65	7)	8.4 (5.6,13.7)	34	21 (61.8)	14.7 (5.5,21.9)	1.12	0.68,	1.92	0.6605
Interaction p-value							<u> </u>				0.9024

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3caa 25NOV2020:12:08 khcs324

Table 2.4.3.1 PAOLA1: Summary of subgroup analysis of EQ-5D-5L Visual analogue scale (MID = 10) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			bevacizumab 132)				
Subgroup	n	Number (%) of patients with events			Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	108 (59.3)	13.6 (8.4,16.6)	90	51 (56.7)	17.4 (9.9,27.8)	1.13	0.82,	1.60	0.4546
IV	73	48 (65.8)	7.9 (3.0,15.1)	42	27 (64.3)	9.6 (5.6,19.4)	1.04	0.65,	1.68	0.8804
Interaction p-value										0.7604
Region										
Europe	245	151 (61.6)	11.1 (8.3,13.9)	126	75 (59.5)	15.4 (8.5,21.9)	1.08	0.82,	1.43	0.5863
Japan	10	5 (50.0)	13.9 (2.8, NE)	6	3 (50.0)	24.0 (2.9, NE)	1.36	0.33,	6.65	0.6681
Interaction p-value										0.7520
ECOG performance status at	Basel	ine								
(0) Normal activity	190	119 (62.6)	11.0 (7.9,13.8)	100	63 (63.0)	11.8 (7.9,19.9)	1.06	0.78,	1.44	0.7210
(1) Restricted activity	61	34 (55.7)	13.7 (6.0, NE)	31	14 (45.2)	21.9 (9.8, NE)	1.30	0.71,	2.50	0.4039
Interaction p-value										0.5589
Baseline CA-125 value										
<=ULN	228		12.1 (8.5,15.1)	118		17.4 (9.8,24.1)	1.11	0.83,		0.4959
>ULN	27	20 (74.1)	3.0 (2.9, 6.1)	14	11 (78.6)	9.6 (2.8,14.7)	1.06	0.52,	2.29	0.8760
Interaction p-value										0.9161
Histological grade										
High grade	255	156 (61.2)	11.1 (8.3,13.9)	132	78 (59.1)	16.4 (9.6,21.9)	1.10	0.84,	1.44	0.5076
Interaction p-value										NC
Cytoreductive surgery outco	me									
No residue	166	98 (59.0)	11.2 (8.3,15.9)	80	46 (57.5)	17.0 (8.3,24.1)	1.08	0.77,	1.55	0.6676
Residue	79	49 (62.0)	8.9 (5.7,19.2)	44	26 (59.1)	17.1 (8.1,24.9)	1.19	0.75,	1.94	0.4740
Interaction p-value										0.7503

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3caa 25NOV2020:12:08 khcs324

Table 2.4.3.1 PAOLA1: Summary of subgroup analysis of EQ-5D-5L Visual analogue scale (MID = 10) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	·	Olaparib + bevacizumab (N=255)				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% C	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	82 (56.2)	12.1 (8.3,19.4)	79	47 (59.5)	17.0 (9.9,24.1)	1.00	0.70,	1.44	0.9818
Interval	99	65 (65.7)	8.7 (5.6,15.6)	45	25 (55.6)	13.4 (5.7, NE)	1.31	0.84,	2.12	0.2417
Interaction p-value										0.3545
Myriad tumour BRCA mutati	ion stat	us								
tBRCAm	158	96 (60.8)	11.4 (8.3,15.6)	77	41 (53.2)	17.0 (9.6,28.7)	1.22	0.85,	1.77	0.2833
Non-tBRCAm	97	60 (61.9)	9.3 (5.7,17.5)	55	37 (67.3)	16.4 (5.6,19.9)	0.97	0.64,	1.47	0.8730
Interaction p-value										0.4097
Status somatic BRCA mutat	ions									
sBRCAm	22	12 (54.5)	8.1 (2.8, NE)	7	4 (57.1)	12.6 (5.6, NE)	1.22	0.42,	4.37	0.7260
gBRCAm	66	39 (59.1)	13.8 (8.4,25.3)	31	17 (54.8)	23.5 (9.6, NE)	1.24	0.71,	2.25	0.4567
Non-BRCAm	41	25 (61.0)	13.9 (5.7, NE)	22	17 (77.3)	7.7 (2.8,21.9)	0.69	0.37,	1.29	0.2382
Interaction p-value										0.3567

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3caa 25NoV2020:12:08 khcs324

Table 2.4.3.2 PAOLA1: Summary of subgroup analysis of EQ-5D-5L Visual analogue scale (MID = 7) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		+ bevacizumab (N=255)	·		oevacizumab 132)			
Subgroup	Number (of patien n with even	nts (95% CI)	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IVRS)							
NED [PDS]	92 47 (51	.1) 13.8 (8.3, NE)	48	30 (62.5)	17.0 (8.5,27.8)	0.86	0.55, 1.37	0.5156
NED/CR [IDS]	74 49 (66	.2) 11.1 (5.6,15.9)	38	19 (50.0)	14.7 (5.6, NE)	1.44	0.86, 2.51	0.1658
NED/CR [Chemo]	40 24 (60	.0) 13.9 (5.6, NE)	20	14 (70.0)	10.0 (3.0,19.4)	0.84	0.44, 1.66	0.6032
PR	49 36 (73	.5) 5.9 (3.0,13.7)	26	15 (57.7)	18.7 (2.9,24.9)	1.43	0.80, 2.70	0.2326
Interaction p-value								0.3168
Screening laboratory tBRO	CA status (IVRS)							
tBRCAm	150 89 (59	.3) 11.5 (8.3,15.9)	65	36 (55.4)	15.4 (8.5, NE)	1.15	0.79, 1.72	0.4724
non-tBRCAm	105 67 (63	.8) 8.5 (5.7,13.9)	67	42 (62.7)	16.4 (5.6,21.9)	1.09	0.75, 1.62	0.6553
Interaction p-value								0.8491
First line treatment outo	come (eCRF)							
NED [PDS]	89 48 (53	.9) 13.6 (8.3, NE)	47	30 (63.8)	17.0 (8.3,24.1)	0.89	0.56, 1.41	0.6076
NED/CR [IDS]	74 49 (66	.2) 11.1 (5.6,15.9)	32	16 (50.0)	12.1 (3.1, NE)	1.39	0.81, 2.53	0.2386
NED/CR [Chemo]	39 22 (56	.4) 15.1 (5.6, NE)	18	11 (61.1)	10.3 (3.5, NE)	0.91	0.45, 1.96	0.8111
PR	50 36 (72	.0) 6.0 (3.0,13.7)	34	21 (61.8)	14.7 (5.5,24.9)	1.34	0.79, 2.33	0.2853
Interaction p-value								0.5159
Screening laboratory tBRO	CA status (eCRF)							
tBRCAm	147 87 (59	.2) 11.4 (8.3,18.0)	67	36 (53.7)	15.4 (8.5, NE)	1.18	0.81, 1.77	0.3912
non-tBRCAm	108 69 (63	.9) 9.3 (5.7,14.0)	65	42 (64.6)	16.4 (5.6,21.9)	1.05	0.72, 1.55	0.8091
Interaction p-value								0.6641
Age group								
<65 years	185 110 (59	.5) 12.7 (8.4,15.6)	98	57 (58.2)	17.0 (9.6,24.9)	1.08	0.79, 1.50	0.6265
>=65 years	70 46 (65	.7) 8.4 (5.6,13.7)	34	21 (61.8)	14.7 (5.5,21.9)	1.12	0.68, 1.92	0.6593
Interaction p-value								0.9069

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3cab 25NoV2020:12:08 khcs324

Table 2.4.3.2 PAOLA1: Summary of subgroup analysis of EQ-5D-5L Visual analogue scale (MID = 7) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			oevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI [b	2-sided p-value [b]
FIGO Stage (Disease state)									
III	182	108 (59.3)	13.6 (8.4,15.9)	90	51 (56.7)	17.4 (9.9,27.8)	1.14	0.82, 1.	0.4470
IV	73	48 (65.8)	7.9 (3.0,15.1)	42	27 (64.3)	9.6 (5.6,19.4)	1.04	0.65, 1.	68 0.8782
Interaction p-value									0.7565
Region									
Europe	245	151 (61.6)	11.1 (8.3,13.9)	126	75 (59.5)	15.4 (8.5,21.9)	1.08	0.82, 1.	
Japan	10	5 (50.0)	13.9 (2.8, NE)	6	3 (50.0)	24.0 (2.9, NE)	1.36	0.33, 6.	0.6667
Interaction p-value									0.7525
ECOG performance status at	Basel	ine							
(0) Normal activity	190	119 (62.6)	11.0 (7.9,13.8)	100	63 (63.0)	11.8 (7.9,19.9)	1.06	0.78, 1.	45 0.7125
(1) Restricted activity	61	34 (55.7)	13.7 (6.0, NE)	31	14 (45.2)	21.9 (9.8, NE)	1.30	0.71, 2.	0.4011
Interaction p-value									0.5595
Baseline CA-125 value									
<=ULN	228	136 (59.6)	12.1 (8.5,14.5)	118	67 (56.8)	17.4 (9.8,24.1)	1.11	0.83, 1.	49 0.4887
>ULN	27	20 (74.1)	3.0 (2.9, 6.1)	14	11 (78.6)	9.6 (2.8,14.7)	1.06	0.52, 2.	30 0.8706
Interaction p-value									0.9178
Histological grade									
High grade	255	156 (61.2)	11.1 (8.3,13.9)	132	78 (59.1)	16.4 (9.6,21.9)	1.10	0.84, 1.	45 0.4994
Interaction p-value									NC
Cytoreductive surgery outco	ome								
No residue	166	98 (59.0)	11.2 (8.3,15.6)	80	46 (57.5)	17.0 (8.3,24.1)	1.08	0.77, 1.	55 0.6577
Residue	79	49 (62.0)	8.9 (5.7,19.2)	44	26 (59.1)	17.1 (8.1,24.9)	1.19	0.75, 1.	94 0.4728
Interaction p-value									0.7552

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3cab 25NoV2020:12:08 khcs324

Table 2.4.3.2 PAOLA1: Summary of subgroup analysis of EQ-5D-5L Visual analogue scale (MID = 7) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	urgery									
Upfront	146	82 (56.2)	12.1 (8.3,19.3)	79	47 (59.5)	17.0 (9.9,24.1)	1.00	0.70,	1.44	0.9956
Interval	99	65 (65.7)	8.7 (5.6,15.6)	45	25 (55.6)	13.4 (5.7, NE)	1.31	0.84,	2.12	0.2428
Interaction p-value										0.3612
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	96 (60.8)	11.4 (8.3,15.6)	77	41 (53.2)	17.0 (9.6,28.7)	1.22	0.85,	1.78	0.2826
Non-tBRCAm	97	60 (61.9)	9.3 (5.7,14.0)	55	37 (67.3)	16.4 (5.6,19.9)	0.97	0.65,	1.48	0.8891
Interaction p-value										0.4179
Status somatic BRCA mutat	ions									
sBRCAm	22	12 (54.5)	8.1 (2.8, NE)	7	4 (57.1)	12.6 (5.6, NE)	1.22	0.43,	4.38	0.7235
gBRCAm	66	39 (59.1)	13.8 (8.4,25.3)	31	17 (54.8)	23.5 (9.6, NE)	1.24	0.71,	2.26	0.4544
Non-BRCAm	41	25 (61.0)	13.9 (5.7, NE)	22	17 (77.3)	7.7 (2.8,21.9)	0.69	0.38,	1.31	0.2508
Interaction p-value										0.3678

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr_v3.sas ettesubpr_v3cab 25NoV2020:12:08 khcs324

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			Result							
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max		
EORTC QLQ-C30 Global QoL/health status	Olaparib + bevacizumab (N=255)	Baseline [a]	245	69.32	17.460	16.7	66.67	100.		
		Wk 12 (Day 85)	225	65.70	17.457	0.0	66.67	100.		
		Wk 24 (Day 169)	201	67.91	17.732	0.0	66.67	100.		
		Wk 36 (Day 253)	178	69.38	16.867	25.0	70.83	100.		
		Wk 48 (Day 337)	175	68.57	17.989	0.0	66.67	100.		
		Wk 60 (Day 421)	162	71.35	15.951	33.3	70.83	100.		
		Wk 72 (Day 505)	158	71.52	16.646	33.3	75.00	100.		
		Wk 84 (Day 589)	138	72.64	16.266	33.3	75.00	100.		
		Wk 96 (Day 673)	137	70.74	18.249	16.7	66.67	100.		
		Wk 108 (Day 757)	110	73.11	19.638	16.7	83.33	100.		
		Wk 120 (Day 841)	1	66.67	NC	66.7	66.67	66.5		
		Wk 132 (Day 925)	1	66.67	NC	66.7	66.67	66.		
		Wk 144 (Day 1009)	1	66.67	NC	66.7	66.67	66.		
		Wk 156 (Day 1093)	1	83.33	NC	83.3	83.33	83.		
		End of Treatment	131	69.85	19.460	0.0	66.67	100		
		30 day Follow-up	61	66.94	21.299	8.3	66.67	100		
	Placebo + bevacizumab (N=132)	Baseline [a]	124	68.75	15.855	33.3	66.67	100		
		Wk 12 (Day 85)	117	67.31	16.957	25.0	66.67	100		
		Wk 24 (Day 169)	103	68.53	17.501	25.0	66.67	100		
		Wk 36 (Day 253)	97	67.44	16.583	16.7	66.67	100		
		Wk 48 (Day 337)	86	68.70	18.957	16.7	70.83	100		
		Wk 60 (Day 421)	71	72.30	15.544	25.0	66.67	100		
		Wk 72 (Day 505)	67	71.14	17.062	25.0	75.00	100		
		Wk 84 (Day 589)	50	71.67	15.430	33.3	75.00	100		
		Wk 96 (Day 673)	41	71.34	17.584	16.7	66.67	100		
		Wk 108 (Day 757)	32	75.00	16.667	33.3	83.33	100		
		End of Treatment	70	65.36	17.528	16.7	66.67	100		
		30 day Follow-up	24	69.10	22.045	33.3	75.00	100		

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			Result							
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max		
EORTC QLQ-C30 Functional scale: Physical	Olaparib + bevacizumab (N=255)	Baseline [a]	245	79.27	16.865	20.0	80.00	100.		
		Wk 12 (Day 85)	225	77.47	18.485	6.7	80.00	100.		
		Wk 24 (Day 169)	202	78.28	19.534	0.0	86.67	100.		
		Wk 36 (Day 253)	177	78.99	19.241	6.7	80.00	100.		
		Wk 48 (Day 337)	174	79.25	18.870	0.0	80.00	100.		
		Wk 60 (Day 421)	162	80.20	17.123	13.3	86.67	100.		
		Wk 72 (Day 505)	156	81.53	19.063	6.7	86.67	100		
		Wk 84 (Day 589)	138	83.44	17.991	13.3	86.67	100		
		Wk 96 (Day 673)	136	84.14	16.804	20.0	86.67	100		
		Wk 108 (Day 757)	110	83.80	17.676	26.7	86.67	100		
		Wk 120 (Day 841)	1	100.00	NC	100.0	100.00	100		
		Wk 132 (Day 925)	1	100.00	NC	100.0	100.00	100		
		Wk 144 (Day 1009)	1	100.00	NC	100.0	100.00	100		
		Wk 156 (Day 1093)	1	100.00	NC	100.0	100.00	100		
		End of Treatment	131	80.28	18.569	20.0	86.67	100		
		30 day Follow-up	61	80.08	18.857	26.7	80.00	100		
	Placebo + bevacizumab (N=132)	Baseline [a]	126	76.67	18.614	20.0	80.00	100		
		Wk 12 (Day 85)	118	76.91	18.890	20.0	80.00	100		
		Wk 24 (Day 169)	103	78.95	16.547	26.7	86.67	100		
		Wk 36 (Day 253)	97	78.95	16.835	33.3	86.67	100		
		Wk 48 (Day 337)	86	78.80	17.723	26.7	80.00	100		
		Wk 60 (Day 421)	71	82.42	16.556	40.0	86.67	100		
		Wk 72 (Day 505)	67	80.55	18.743	26.7	86.67	100		
		Wk 84 (Day 589)	51	83.66	16.935	40.0	86.67	100		
		Wk 96 (Day 673)	41	82.80	17.169	13.3	86.67	100		
		Wk 108 (Day 757)	31	82.37	19.667	33.3	86.67	100		
		End of Treatment	69	79.64	18.985	25.0	86.67	100		
		30 day Follow-up	24	82.78	18.328	40.0	86.67	100		

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Result							
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max	
EORTC QLQ-C30 Functional scale: Role	Olaparib + bevacizumab (N=255)	Baseline [a]	245	72.93	26.556	0.0	66.67	100.	
		Wk 12 (Day 85)	224	68.90	25.548	0.0	66.67	100.	
		Wk 24 (Day 169)	202	73.93	26.172	0.0	66.67	100.	
		Wk 36 (Day 253)	178	72.19	24.912	0.0	66.67	100.	
		Wk 48 (Day 337)	175	72.67	25.904	0.0	66.67	100.	
		Wk 60 (Day 421)	162	75.41	25.143	0.0	83.33	100.	
		Wk 72 (Day 505)	158	77.11	25.826	0.0	83.33	100.	
		Wk 84 (Day 589)	140	79.29	23.980	0.0	83.33	100.	
		Wk 96 (Day 673)	137	79.68	25.140	0.0	83.33	100	
		Wk 108 (Day 757)	110	79.09	25.386	0.0	83.33	100	
		Wk 120 (Day 841)	1	100.00	NC	100.0	100.00	100	
		Wk 132 (Day 925)	1	100.00	NC	100.0	100.00	100	
		Wk 144 (Day 1009)	1	100.00	NC	100.0	100.00	100	
		Wk 156 (Day 1093)	1	100.00	NC	100.0	100.00	100	
		End of Treatment	131	74.81	27.298	0.0	83.33	100	
		30 day Follow-up	61	69.67	29.425	0.0	66.67	100	
	Placebo + bevacizumab (N=132)	Baseline [a]	126	72.35	27.338	0.0	66.67	100	
		Wk 12 (Day 85)	118	69.35	25.688	0.0	66.67	100	
		Wk 24 (Day 169)	103	71.52	25.633	16.7	66.67	100	
		Wk 36 (Day 253)	97	71.48	23.809	33.3	66.67	100	
		Wk 48 (Day 337)	86	76.36	23.910	0.0	75.00	100	
		Wk 60 (Day 421)	71	76.29	22.122	16.7	66.67	100	
		Wk 72 (Day 505)	68	78.19	21.405	16.7	75.00	100	
		Wk 84 (Day 589)	51	82.35	21.963	33.3	100.00	100	
		Wk 96 (Day 673)	41	79.67	21.574	16.7	83.33	100	
		Wk 108 (Day 757)	32	79.69	22.294	33.3	83.33	100	
		End of Treatment	70	72.62	23.058	16.7	66.67	100	
		30 day Follow-up	24	78.47	28.009	0.0	100.00	100	

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			Result						
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max	
EORTC QLQ-C30 Functional scale: Cognitive	Olaparib + bevacizumab (N=255)	Baseline [a]	246	81.44	20.393	0.0	83.33	100.	
		Wk 12 (Day 85)	224	81.25	20.351	0.0	83.33	100.	
		Wk 24 (Day 169)	202	81.85	20.577	0.0	83.33	100.	
		Wk 36 (Day 253)	178	80.15	21.968	0.0	83.33	100.	
		Wk 48 (Day 337)	175	80.10	21.789	0.0	83.33	100.	
		Wk 60 (Day 421)	163	79.75	22.433	0.0	83.33	100.	
		Wk 72 (Day 505)	158	78.80	23.768	0.0	83.33	100.	
		Wk 84 (Day 589)	138	78.14	22.765	0.0	83.33	100.	
		Wk 96 (Day 673)	137	78.71	22.842	0.0	83.33	100.	
		Wk 108 (Day 757)	110	81.21	22.475	0.0	83.33	100	
		Wk 120 (Day 841)	1	83.33	NC	83.3	83.33	83.	
		Wk 132 (Day 925)	1	66.67	NC	66.7	66.67	66.	
		Wk 144 (Day 1009)	1	83.33	NC	83.3	83.33	83.	
		Wk 156 (Day 1093)	1	83.33	NC	83.3	83.33	83.	
		End of Treatment	131	79.64	23.323	0.0	83.33	100	
		30 day Follow-up	61	77.05	20.450	16.7	83.33	100	
	Placebo + bevacizumab (N=132)	Baseline [a]	124	81.45	19.034	16.7	83.33	100	
		Wk 12 (Day 85)	117	78.92	21.374	0.0	83.33	100	
		Wk 24 (Day 169)	103	80.26	19.067	33.3	83.33	100	
		Wk 36 (Day 253)	97	80.07	19.490	0.0	83.33	100	
		Wk 48 (Day 337)	86	80.04	20.101	33.3	83.33	100	
		Wk 60 (Day 421)	71	84.27	18.875	16.7	83.33	100	
		Wk 72 (Day 505)	67	82.09	19.530	33.3	83.33	100	
		Wk 84 (Day 589)	51	83.99	16.654	50.0	83.33	100	
		Wk 96 (Day 673)	41	82.11	19.857	33.3	83.33	100	
		Wk 108 (Day 757)	32	82.29	18.422	33.3	83.33	100	
		End of Treatment	70	79.76	21.960	0.0	83.33	100	
		30 day Follow-up	24	81.25	23.215	33.3	83.33	100	

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			Result							
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max		
EORTC QLQ-C30 Functional scale: Emotional	Olaparib + bevacizumab (N=255)	Baseline [a]	246	76.10	22.102	0.0	83.33	100.		
		Wk 12 (Day 85)	224	75.25	21.725	0.0	75.00	100.		
		Wk 24 (Day 169)	202	74.16	22.572	0.0	75.00	100.		
		Wk 36 (Day 253)	178	75.81	21.422	0.0	75.00	100.		
		Wk 48 (Day 337)	175	73.41	22.013	0.0	75.00	100.		
		Wk 60 (Day 421)	163	73.16	22.869	0.0	75.00	100.		
		Wk 72 (Day 505)	158	73.80	23.448	0.0	75.00	100		
		Wk 84 (Day 589)	138	73.37	22.646	0.0	75.00	100		
		Wk 96 (Day 673)	137	72.61	24.125	0.0	75.00	100		
		Wk 108 (Day 757)	110	75.38	25.023	0.0	83.33	100		
		Wk 120 (Day 841)	1	100.00	NC	100.0	100.00	100		
		Wk 132 (Day 925)	1	100.00	NC	100.0	100.00	100		
		Wk 144 (Day 1009)	1	100.00	NC	100.0	100.00	100		
		Wk 156 (Day 1093)	1	91.67	NC	91.7	91.67	91.		
		End of Treatment	131	71.06	26.076	0.0	75.00	100		
		30 day Follow-up	61	70.90	22.237	0.0	75.00	100		
	Placebo + bevacizumab (N=132)	Baseline [a]	124	77.87	17.939	16.7	83.33	100		
		Wk 12 (Day 85)	117	73.79	21.422	0.0	75.00	100		
		Wk 24 (Day 169)	103	73.79	22.184	16.7	75.00	100		
		Wk 36 (Day 253)	97	76.23	20.222	8.3	83.33	100		
		Wk 48 (Day 337)	86	76.20	20.419	8.3	75.00	100		
		Wk 60 (Day 421)	71	76.84	19.653	8.3	83.33	100		
		Wk 72 (Day 505)	67	77.78	16.973	33.3	75.00	100		
		Wk 84 (Day 589)	51	79.36	16.899	41.7	83.33	100		
		Wk 96 (Day 673)	41	76.63	19.651	8.3	75.00	100		
		Wk 108 (Day 757)	32	78.39	19.499	33.3	83.33	100		
		End of Treatment	70	70.08	22.056	0.0	75.00	100		
		30 day Follow-up	24	72.92	21.598	25.0	79.17	100		

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			Result							
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max		
EORTC QLQ-C30 Functional scale: Social	Olaparib + bevacizumab (N=255)	Baseline [a]	246	73.85	27.814	0.0	83.33	100.0		
		Wk 12 (Day 85)	224	76.12	24.931	0.0	83.33	100.0		
		Wk 24 (Day 169)	202	76.49	24.684	0.0	83.33	100.0		
		Wk 36 (Day 253)	178	78.09	25.004	0.0	83.33	100.0		
		Wk 48 (Day 337)	174	75.77	25.315	0.0	83.33	100.0		
		Wk 60 (Day 421)	163	80.27	24.931	0.0	83.33	100.0		
		Wk 72 (Day 505)	158	82.07	22.694	0.0	100.00	100.0		
		Wk 84 (Day 589)	138	80.07	23.385	0.0	83.33	100.0		
		Wk 96 (Day 673)	137	78.10	26.099	0.0	83.33	100.0		
		Wk 108 (Day 757)	110	80.76	25.956	0.0	100.00	100.		
		Wk 120 (Day 841)	1	100.00	NC	100.0	100.00	100.		
		Wk 132 (Day 925)	1	100.00	NC	100.0	100.00	100.		
		Wk 144 (Day 1009)	1	100.00	NC	100.0	100.00	100.		
		Wk 156 (Day 1093)	1	100.00	NC	100.0	100.00	100.		
		End of Treatment	131	77.10	27.059	0.0	83.33	100.		
		30 day Follow-up	61	71.58	29.556	0.0	66.67	100.		
	Placebo + bevacizumab (N=132)	Baseline [a]	124	73.52	24.694	0.0	66.67	100.		
		Wk 12 (Day 85)	117	76.50	25.348	0.0	83.33	100.		
		Wk 24 (Day 169)	103	79.29	22.563	0.0	83.33	100.		
		Wk 36 (Day 253)	97	79.04	22.983	0.0	83.33	100.		
		Wk 48 (Day 337)	86	79.84	21.407	16.7	83.33	100.		
		Wk 60 (Day 421)	71	85.92	18.181	33.3	100.00	100.		
		Wk 72 (Day 505)	67	84.83	20.665	16.7	100.00	100.		
		Wk 84 (Day 589)	51	83.33	22.361	33.3	100.00	100.		
		Wk 96 (Day 673)	41	86.99	18.451	33.3	100.00	100.		
		Wk 108 (Day 757)	32	82.81	20.515	33.3	100.00	100.		
		End of Treatment	70	75.71	25.010	0.0	83.33	100.		
		30 day Follow-up	24	77.78	30.163	0.0	100.00	100.		

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-C30 Single item symptom scale: Loss of appetite	Olaparib + bevacizumab (N=255)	Baseline [a]	245	8.03	17.177	0.0	0.00	66.7
		Wk 12 (Day 85)	225	18.81	27.577	0.0	0.00	100.
		Wk 24 (Day 169)	201	15.09	25.144	0.0	0.00	100.
		Wk 36 (Day 253)	176	14.02	23.487	0.0	0.00	100.
		Wk 48 (Day 337)	175	14.48	23.839	0.0	0.00	100.
		Wk 60 (Day 421)	162	11.32	19.363	0.0	0.00	66.7
		Wk 72 (Day 505)	156	11.32	22.239	0.0	0.00	100.
		Wk 84 (Day 589)	138	6.04	15.196	0.0	0.00	66.7
		Wk 96 (Day 673)	137	11.19	20.721	0.0	0.00	100
		Wk 108 (Day 757)	110	7.88	17.445	0.0	0.00	100
		Wk 120 (Day 841)	1	0.00	NC	0.0	0.00	0.0
		Wk 132 (Day 925)	1	0.00	NC	0.0	0.00	0.0
		Wk 144 (Day 1009)	1	0.00	NC	0.0	0.00	0.0
		Wk 156 (Day 1093)	1	0.00	NC	0.0	0.00	0.0
		End of Treatment	131	13.74	23.341	0.0	0.00	100
		30 day Follow-up	61	14.75	25.477	0.0	0.00	100
	Placebo + bevacizumab (N=132)	Baseline [a]	126	8.73	17.476	0.0	0.00	66.
		Wk 12 (Day 85)	118	9.60	21.396	0.0	0.00	100
		Wk 24 (Day 169)	103	9.71	20.147	0.0	0.00	100
		Wk 36 (Day 253)	95	8.77	18.963	0.0	0.00	100
		Wk 48 (Day 337)	86	8.91	19.419	0.0	0.00	100
		Wk 60 (Day 421)	71	6.57	16.542	0.0	0.00	100
		Wk 72 (Day 505)	68	7.35	15.068	0.0	0.00	66.
		Wk 84 (Day 589)	51	5.23	12.243	0.0	0.00	33.
		Wk 96 (Day 673)	41	8.13	16.297	0.0	0.00	66.
		Wk 108 (Day 757)	32	7.29	16.361	0.0	0.00	66.
		End of Treatment	70	13.33	21.535	0.0	0.00	100
		30 day Follow-up	24	8.33	20.264	0.0	0.00	66.

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-C30 Single item symptom scale: Constipation	Olaparib + bevacizumab (N=255)	Baseline [a]	243	17.83	27.311	0.0	0.00	100.
		Wk 12 (Day 85)	221	13.42	24.532	0.0	0.00	100.
		Wk 24 (Day 169)	200	15.33	25.414	0.0	0.00	100.
		Wk 36 (Day 253)	178	14.98	25.557	0.0	0.00	100.
		Wk 48 (Day 337)	175	14.48	25.395	0.0	0.00	100
		Wk 60 (Day 421)	161	20.29	29.621	0.0	0.00	100
		Wk 72 (Day 505)	155	20.00	27.287	0.0	0.00	100
		Wk 84 (Day 589)	139	21.82	29.410	0.0	0.00	100
		Wk 96 (Day 673)	136	19.61	26.441	0.0	0.00	100
		Wk 108 (Day 757)	110	20.00	27.539	0.0	0.00	100
		Wk 120 (Day 841)	1	33.33	NC	33.3	33.33	33.
		Wk 132 (Day 925)	1	0.00	NC	0.0	0.00	0.0
		Wk 144 (Day 1009)	1	0.00	NC	0.0	0.00	0.0
		Wk 156 (Day 1093)	1	0.00	NC	0.0	0.00	0.0
		End of Treatment	130	18.46	27.550	0.0	0.00	100
		30 day Follow-up	61	24.04	31.110	0.0	0.00	100
	Placebo + bevacizumab (N=132)	Baseline [a]	124	14.78	24.529	0.0	0.00	100
		Wk 12 (Day 85)	117	11.97	22.515	0.0	0.00	100
		Wk 24 (Day 169)	100	15.00	23.391	0.0	0.00	100
		Wk 36 (Day 253)	97	12.37	20.593	0.0	0.00	66.
		Wk 48 (Day 337)	85	14.90	23.852	0.0	0.00	100
		Wk 60 (Day 421)	70	13.81	20.059	0.0	0.00	66.
		Wk 72 (Day 505)	67	15.92	24.862	0.0	0.00	100
		Wk 84 (Day 589)	50	16.67	25.422	0.0	0.00	100
		Wk 96 (Day 673)	41	24.39	27.913	0.0	33.33	100
		Wk 108 (Day 757)	32	18.75	25.312	0.0	0.00	100
		End of Treatment	69	20.29	26.945	0.0	0.00	100
		30 day Follow-up	24	19.44	27.657	0.0	0.00	100

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-C30 Single item symptom scale: Diarrhoea	Olaparib + bevacizumab (N=255)	Baseline [a]	243	9.60	19.146	0.0	0.00	100.
		Wk 12 (Day 85)	223	12.71	22.661	0.0	0.00	100.
		Wk 24 (Day 169)	199	13.23	25.693	0.0	0.00	100.
		Wk 36 (Day 253)	178	8.99	19.896	0.0	0.00	100.
		Wk 48 (Day 337)	174	9.20	16.955	0.0	0.00	66.7
		Wk 60 (Day 421)	162	10.49	19.824	0.0	0.00	100.
		Wk 72 (Day 505)	158	9.70	19.295	0.0	0.00	100.
		Wk 84 (Day 589)	138	7.73	18.583	0.0	0.00	100.
		Wk 96 (Day 673)	137	9.49	19.364	0.0	0.00	100.
		Wk 108 (Day 757)	110	6.06	15.099	0.0	0.00	66.5
		Wk 120 (Day 841)	1	0.00	NC	0.0	0.00	0.0
		Wk 132 (Day 925)	1	0.00	NC	0.0	0.00	0.0
		Wk 144 (Day 1009)	1	0.00	NC	0.0	0.00	0.0
		Wk 156 (Day 1093)	1	0.00	NC	0.0	0.00	0.0
		End of Treatment	131	10.18	20.628	0.0	0.00	100
		30 day Follow-up	61	9.29	18.390	0.0	0.00	100
	Placebo + bevacizumab (N=132)	Baseline [a]	125	13.07	24.647	0.0	0.00	100
		Wk 12 (Day 85)	117	10.83	22.244	0.0	0.00	100
		Wk 24 (Day 169)	102	12.75	25.279	0.0	0.00	100
		Wk 36 (Day 253)	95	13.33	24.982	0.0	0.00	100
		Wk 48 (Day 337)	86	14.34	26.341	0.0	0.00	100
		Wk 60 (Day 421)	70	11.90	22.725	0.0	0.00	100
		Wk 72 (Day 505)	66	9.60	21.693	0.0	0.00	100
		Wk 84 (Day 589)	51	10.46	22.598	0.0	0.00	100
		Wk 96 (Day 673)	41	13.01	22.209	0.0	0.00	66.
		Wk 108 (Day 757)	32	8.33	22.401	0.0	0.00	100
		End of Treatment	70	11.43	23.320	0.0	0.00	100
		30 day Follow-up	24	9.72	23.008	0.0	0.00	100

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-C30 Single item symptom scale: Dyspnoea	Olaparib + bevacizumab (N=255)	Baseline [a]	242	23.97	26.723	0.0	33.33	100.
		Wk 12 (Day 85)	223	26.31	28.040	0.0	33.33	100.
		Wk 24 (Day 169)	200	23.33	26.524	0.0	33.33	100.
		Wk 36 (Day 253)	178	23.22	27.631	0.0	16.67	100.
		Wk 48 (Day 337)	175	23.43	26.809	0.0	33.33	100.
		Wk 60 (Day 421)	162	24.49	28.728	0.0	33.33	100.
		Wk 72 (Day 505)	157	23.14	26.057	0.0	33.33	100.
		Wk 84 (Day 589)	138	21.01	23.853	0.0	33.33	100.
		Wk 96 (Day 673)	137	22.14	25.654	0.0	33.33	100.
		Wk 108 (Day 757)	109	21.41	24.226	0.0	33.33	100.
		Wk 120 (Day 841)	1	0.00	NC	0.0	0.00	0.0
		Wk 132 (Day 925)	1	0.00	NC	0.0	0.00	0.0
		Wk 144 (Day 1009)	1	0.00	NC	0.0	0.00	0.0
		Wk 156 (Day 1093)	1	0.00	NC	0.0	0.00	0.0
		End of Treatment	130	24.10	26.255	0.0	33.33	100
		30 day Follow-up	61	28.96	28.203	0.0	33.33	100
	Placebo + bevacizumab (N=132)	Baseline [a]	126	23.28	27.087	0.0	16.67	100
		Wk 12 (Day 85)	118	20.62	25.382	0.0	0.00	100
		Wk 24 (Day 169)	103	21.68	25.866	0.0	0.00	100
		Wk 36 (Day 253)	97	23.02	27.370	0.0	0.00	100
		Wk 48 (Day 337)	86	19.77	24.722	0.0	0.00	100
		Wk 60 (Day 421)	71	22.07	26.394	0.0	0.00	100
		Wk 72 (Day 505)	67	17.41	21.987	0.0	0.00	66.
		Wk 84 (Day 589)	51	16.99	25.274	0.0	0.00	100
		Wk 96 (Day 673)	41	22.76	24.082	0.0	33.33	66.
		Wk 108 (Day 757)	32	21.87	21.767	0.0	33.33	66.
		End of Treatment	70	20.48	23.599	0.0	16.67	100
		30 day Follow-up	24	23.61	26.882	0.0	33.33	100

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-C30 Symptom scale: Fatigue	Olaparib + bevacizumab (N=255)	Baseline [a]	244	33.15	23.035	0.0	33.33	100.
		Wk 12 (Day 85)	225	39.70	23.039	0.0	33.33	100.
		Wk 24 (Day 169)	202	35.04	21.919	0.0	33.33	100.
		Wk 36 (Day 253)	177	35.00	23.025	0.0	33.33	100
		Wk 48 (Day 337)	175	35.02	23.536	0.0	33.33	100
		Wk 60 (Day 421)	162	33.57	23.984	0.0	33.33	100
		Wk 72 (Day 505)	157	31.00	24.020	0.0	33.33	100
		Wk 84 (Day 589)	139	31.65	22.329	0.0	33.33	100
		Wk 96 (Day 673)	137	30.25	23.771	0.0	33.33	100
		Wk 108 (Day 757)	109	29.66	22.696	0.0	22.22	100
		Wk 120 (Day 841)	1	22.22	NC	22.2	22.22	22.
		Wk 132 (Day 925)	1	22.22	NC	22.2	22.22	22.
		Wk 144 (Day 1009)	1	22.22	NC	22.2	22.22	22.
		Wk 156 (Day 1093)	1	22.22	NC	22.2	22.22	22.
		End of Treatment	130	33.33	23.973	0.0	33.33	100
		30 day Follow-up	61	37.25	24.540	0.0	33.33	100
	Placebo + bevacizumab (N=132)	Baseline [a]	125	34.31	22.357	0.0	33.33	100
		Wk 12 (Day 85)	118	37.19	24.347	0.0	33.33	100
		Wk 24 (Day 169)	103	34.57	20.946	0.0	33.33	88.
		Wk 36 (Day 253)	97	34.25	20.549	0.0	33.33	88.
		Wk 48 (Day 337)	86	31.85	22.245	0.0	33.33	100
		Wk 60 (Day 421)	71	28.95	21.164	0.0	33.33	88
		Wk 72 (Day 505)	68	31.54	20.686	0.0	33.33	88
		Wk 84 (Day 589)	51	27.45	21.006	0.0	22.22	66
		Wk 96 (Day 673)	41	27.24	20.100	0.0	22.22	88.
		Wk 108 (Day 757)	32	28.30	18.686	0.0	33.33	66
		End of Treatment	69	34.14	23.207	0.0	33.33	100
		30 day Follow-up	24	31.25	21.907	0.0	33.33	66.

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-C30 Single item symptom scale: Financial difficulties	Olaparib + bevacizumab (N=255)	Baseline [a]	244	17.76	29.686	0.0	0.00	100.
		Wk 12 (Day 85)	222	16.82	29.704	0.0	0.00	100.
		Wk 24 (Day 169)	199	17.42	29.743	0.0	0.00	100.
		Wk 36 (Day 253)	176	16.67	28.508	0.0	0.00	100.
		Wk 48 (Day 337)	173	16.57	28.217	0.0	0.00	100.
		Wk 60 (Day 421)	162	15.64	28.579	0.0	0.00	100.
		Wk 72 (Day 505)	154	16.23	28.325	0.0	0.00	100.
		Wk 84 (Day 589)	138	15.70	27.373	0.0	0.00	100.
		Wk 96 (Day 673)	137	15.57	28.311	0.0	0.00	100.
		Wk 108 (Day 757)	110	13.64	25.660	0.0	0.00	100.
		Wk 120 (Day 841)	1	0.00	NC	0.0	0.00	0.0
		Wk 132 (Day 925)	1	0.00	NC	0.0	0.00	0.0
		Wk 144 (Day 1009)	1	0.00	NC	0.0	0.00	0.0
		Wk 156 (Day 1093)	1	0.00	NC	0.0	0.00	0.0
		End of Treatment	130	14.10	25.875	0.0	0.00	100
		30 day Follow-up	59	13.56	27.065	0.0	0.00	100
	Placebo + bevacizumab (N=132)	Baseline [a]	122	19.13	29.665	0.0	0.00	100
		Wk 12 (Day 85)	117	19.66	29.084	0.0	0.00	100
		Wk 24 (Day 169)	102	17.32	27.643	0.0	0.00	100
		Wk 36 (Day 253)	97	15.46	25.029	0.0	0.00	100
		Wk 48 (Day 337)	86	17.05	26.442	0.0	0.00	100
		Wk 60 (Day 421)	71	15.49	26.922	0.0	0.00	100
		Wk 72 (Day 505)	66	14.65	23.482	0.0	0.00	100
		Wk 84 (Day 589)	51	12.42	24.001	0.0	0.00	100
		Wk 96 (Day 673)	41	12.20	20.758	0.0	0.00	66.
		Wk 108 (Day 757)	32	12.50	20.302	0.0	0.00	66.
		End of Treatment	70	18.10	28.762	0.0	0.00	100
		30 day Follow-up	24	15.28	27.766	0.0	0.00	100

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-C30 Symptom scale: Nausea and vomiting	Olaparib + bevacizumab (N=255)	Baseline [a]	245	4.35	11.750	0.0	0.00	100.0
		Wk 12 (Day 85)	225	14.89	20.637	0.0	0.00	100.
		Wk 24 (Day 169)	202	10.97	16.518	0.0	0.00	83.3
		Wk 36 (Day 253)	178	9.64	14.897	0.0	0.00	83.3
		Wk 48 (Day 337)	175	11.71	17.893	0.0	0.00	100.
		Wk 60 (Day 421)	162	10.70	16.986	0.0	0.00	83.3
		Wk 72 (Day 505)	157	9.77	17.300	0.0	0.00	100.
		Wk 84 (Day 589)	139	8.03	15.195	0.0	0.00	100.
		Wk 96 (Day 673)	137	7.06	12.414	0.0	0.00	66.7
		Wk 108 (Day 757)	110	5.91	13.656	0.0	0.00	100.
		Wk 120 (Day 841)	1	0.00	NC	0.0	0.00	0.0
		Wk 132 (Day 925)	1	0.00	NC	0.0	0.00	0.0
		Wk 144 (Day 1009)	1	0.00	NC	0.0	0.00	0.0
		Wk 156 (Day 1093)	1	0.00	NC	0.0	0.00	0.0
		End of Treatment	131	9.16	17.198	0.0	0.00	100.
		30 day Follow-up	61	9.56	24.430	0.0	0.00	100.
	Placebo + bevacizumab (N=132)	Baseline [a]	126	3.17	8.873	0.0	0.00	50.0
		Wk 12 (Day 85)	118	4.52	11.251	0.0	0.00	66.7
		Wk 24 (Day 169)	103	3.07	7.286	0.0	0.00	33.3
		Wk 36 (Day 253)	97	5.33	10.904	0.0	0.00	50.0
		Wk 48 (Day 337)	86	4.84	11.395	0.0	0.00	66.7
		Wk 60 (Day 421)	71	2.11	5.585	0.0	0.00	16.7
		Wk 72 (Day 505)	68	1.96	5.410	0.0	0.00	16.7
		Wk 84 (Day 589)	51	3.27	8.178	0.0	0.00	33.3
		Wk 96 (Day 673)	41	6.91	17.864	0.0	0.00	83.3
		Wk 108 (Day 757)	32	5.73	16.725	0.0	0.00	66.7
		End of Treatment	70	6.19	15.061	0.0	0.00	66.7
		30 day Follow-up	24	11.11	25.380	0.0	0.00	100.

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-C30 Symptom scale: Pain	Olaparib + bevacizumab (N=255)	Baseline [a]	247	22.81	23.268	0.0	16.67	100.
		Wk 12 (Day 85)	225	28.89	25.394	0.0	33.33	100.
		Wk 24 (Day 169)	202	28.30	26.354	0.0	16.67	100.
		Wk 36 (Day 253)	178	29.03	25.738	0.0	33.33	100.
		Wk 48 (Day 337)	175	29.71	26.375	0.0	33.33	100.
		Wk 60 (Day 421)	164	25.81	24.790	0.0	16.67	100.
		Wk 72 (Day 505)	159	25.05	23.109	0.0	16.67	100.
		Wk 84 (Day 589)	139	21.22	22.948	0.0	16.67	100.
		Wk 96 (Day 673)	137	20.68	23.788	0.0	16.67	100.
		Wk 108 (Day 757)	110	18.64	24.076	0.0	0.00	100.
		Wk 120 (Day 841)	1	33.33	NC	33.3	33.33	33.3
		Wk 132 (Day 925)	1	16.67	NC	16.7	16.67	16.
		Wk 144 (Day 1009)	1	16.67	NC	16.7	16.67	16.
		Wk 156 (Day 1093)	1	33.33	NC	33.3	33.33	33.
		End of Treatment	131	21.88	23.758	0.0	16.67	100
		30 day Follow-up	61	26.23	26.080	0.0	16.67	100
	Placebo + bevacizumab (N=132)	Baseline [a]	126	23.94	22.949	0.0	16.67	100
		Wk 12 (Day 85)	118	32.77	26.414	0.0	33.33	100
		Wk 24 (Day 169)	104	30.13	22.646	0.0	33.33	100
		Wk 36 (Day 253)	97	30.76	24.807	0.0	33.33	100
		Wk 48 (Day 337)	86	29.26	25.041	0.0	33.33	100
		Wk 60 (Day 421)	71	27.23	20.362	0.0	33.33	83.
		Wk 72 (Day 505)	68	25.49	24.344	0.0	16.67	83.
		Wk 84 (Day 589)	51	23.20	20.569	0.0	16.67	66.
		Wk 96 (Day 673)	41	18.29	21.668	0.0	0.00	66.
		Wk 108 (Day 757)	32	19.27	23.988	0.0	8.33	83.
		End of Treatment	70	27.14	23.938	0.0	33.33	83.
		30 day Follow-up	24	31.94	31.051	0.0	33.33	100

Table 2.5.1 PAOLA1: Summary of EORTC QLQ-C30 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-C30 Single item symptom scale: Insomnia	Olaparib + bevacizumab (N=255)	Baseline [a]	243	28.40	30.049	0.0	33.33	100.
		Wk 12 (Day 85)	226	30.97	30.051	0.0	33.33	100.
		Wk 24 (Day 169)	202	33.17	31.805	0.0	33.33	100.
		Wk 36 (Day 253)	178	30.90	30.279	0.0	33.33	100.
		Wk 48 (Day 337)	174	30.27	30.043	0.0	33.33	100.
		Wk 60 (Day 421)	160	33.33	33.017	0.0	33.33	100.
		Wk 72 (Day 505)	157	28.03	31.011	0.0	33.33	100.
		Wk 84 (Day 589)	139	31.89	29.725	0.0	33.33	100.
		Wk 96 (Day 673)	136	32.11	33.063	0.0	33.33	100.
		Wk 108 (Day 757)	109	32.11	30.404	0.0	33.33	100
		Wk 120 (Day 841)	1	0.00	NC	0.0	0.00	0.0
		Wk 132 (Day 925)	1	0.00	NC	0.0	0.00	0.0
		Wk 144 (Day 1009)	1	0.00	NC	0.0	0.00	0.0
		Wk 156 (Day 1093)	1	33.33	NC	33.3	33.33	33.
		End of Treatment	130	35.64	30.834	0.0	33.33	100
		30 day Follow-up	61	32.24	31.604	0.0	33.33	100
	Placebo + bevacizumab (N=132)	Baseline [a]	126	21.96	25.002	0.0	33.33	100
		Wk 12 (Day 85)	118	32.49	29.706	0.0	33.33	100
		Wk 24 (Day 169)	103	28.80	30.625	0.0	33.33	100
		Wk 36 (Day 253)	97	28.87	29.513	0.0	33.33	100
		Wk 48 (Day 337)	86	29.07	28.375	0.0	33.33	100
		Wk 60 (Day 421)	71	30.99	26.018	0.0	33.33	100
		Wk 72 (Day 505)	68	26.96	25.273	0.0	33.33	100
		Wk 84 (Day 589)	51	28.10	32.912	0.0	33.33	100
		Wk 96 (Day 673)	40	30.00	23.631	0.0	33.33	100
		Wk 108 (Day 757)	32	33.33	26.774	0.0	33.33	100
		End of Treatment	69	34.30	27.399	0.0	33.33	100
		30 day Follow-up	24	30.56	30.954	0.0	33.33	100

Table 2.5.2.1 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Global QoL/health status (mixed model for repeated measures) Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=255		Placebo + bev		Difference betwee	n groups
Timepoint	n Mean (SD) at	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE)	Estimated difference (95% CI)	p-value
Wk 12 (Day 85)	203 70.40 (17.296)	-4.17 (1.064)	111 69.37 (15.495)	-2.05 (1.438)	-2.12 (-5.639, 1.40	2) 0.2373
Wk 24 (Day 169)	186 70.74 (17.656)	-0.97 (1.032)	98 68.20 (16.203)	-0.47 (1.422)	-0.50 (-3.962, 2.96	0) 0.7758
Wk 36 (Day 253)	165 69.29 (17.138)	-0.23 (1.109)	91 68.68 (15.681)	-2.54 (1.509)	2.31 (-1.376, 5.99	7) 0.2184
Wk 48 (Day 337)	163 69.33 (18.090)	-0.35 (1.177)	76 69.85 (16.157)	-0.13 (1.709)	-0.22 (-4.306, 3.86	8) 0.9160
Wk 60 (Day 421)	151 68.65 (17.995)	2.18 (1.071)	67 69.78 (16.969)	0.20 (1.569)	1.98 (-1.761, 5.72	2) 0.2983
Wk 72 (Day 505)	147 69.27 (17.713)	2.39 (1.181)	59 70.90 (16.035)	2.25 (1.827)	0.13 (-4.155, 4.41	9) 0.9517
Wk 84 (Day 589)	131 68.77 (17.883)	2.91 (1.179)	45 73.52 (15.416)	-0.40 (1.905)	3.31 (-1.106, 7.73	6) 0.1410
Wk 96 (Day 673)	129 69.06 (17.927)	1.13 (1.396)	37 70.50 (17.524)	0.63 (2.429)	0.49 (-5.031, 6.02	0.8600
Average over all visits	219 70.02 (17.125)	0.36 (0.783)	118 68.50 (15.981)	-0.31 (1.136)	0.67 (-2.042, 3.39	1) 0.6256
Hedges' g SMD					0.06 (-0.167, 0.28	1) 0.6194

model within-subject error.

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.2.2 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Functional scale: Physical (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be		Placebo + bev		Difference between	groups
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Wk 12 (Day 85)	203 79.48 (17.057)	-1.04 (0.969)	114 76.84 (18.235)	-0.39 (1.299)	-0.64 (-3.837, 2.548) 0.6916
Wk 24 (Day 169)	188 78.94 (17.352)	0.14 (1.021)	100 76.27 (19.182)	1.79 (1.400)	-1.65 (-5.057, 1.767) 0.3435
Wk 36 (Day 253)	164 79.43 (17.334)	0.68 (1.110)	93 76.45 (19.393)	0.73 (1.501)	-0.05 (-3.724, 3.633) 0.9807
Wk 48 (Day 337)	164 79.09 (17.415)	0.43 (1.057)	78 78.16 (18.993)	0.59 (1.498)	-0.16 (-3.768, 3.454) 0.9318
Wk 60 (Day 421)	152 79.59 (16.997)	1.73 (1.047)	68 76.79 (20.001)	3.01 (1.516)	-1.28 (-4.912, 2.354	0.4889
Wk 72 (Day 505)	146 78.90 (18.044)	3.67 (1.139)	61 77.02 (20.605)	2.76 (1.697)	0.91 (-3.117, 4.936) 0.6570
Wk 84 (Day 589)	132 79.38 (17.512)	4.54 (1.103)	47 76.88 (20.470)	5.55 (1.688)	-1.01 (-4.982, 2.966	0.6180
Wk 96 (Day 673)	129 79.48 (17.293)	5.28 (1.066)	37 75.86 (21.277)	5.09 (1.760)	0.19 (-3.866, 4.249	0.9260
Average over all visits	219 79.52 (16.897)	1.93 (0.833)	120 76.50 (18.826)	2.39 (1.174)	-0.46 (-3.295, 2.375	0.7499
Hedges' g SMD					-0.04 (-0.259, 0.186	0.7470

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.2.3 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Functional scale: Role (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=255		Placebo + bev		Difference between o	groups
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Wk 12 (Day 85)	202 72.61 (26.557)	-3.46 (1.469)	114 72.95 (27.010)	-2.87 (1.962)	-0.59 (-5.412, 4.231)	0.8098
Wk 24 (Day 169)	188 72.52 (27.002)	2.24 (1.558)	100 71.67 (27.677)	0.09 (2.135)	2.15 (-3.050, 7.351)	0.4165
Wk 36 (Day 253)	165 72.12 (26.872)	0.67 (1.597)	93 71.51 (27.207)	-1.57 (2.146)	2.25 (-3.019, 7.511)	0.4019
Wk 48 (Day 337)	165 73.43 (27.540)	0.55 (1.603)	78 74.15 (25.432)	2.72 (2.295)	-2.17 (-7.675, 3.344)	0.4397
Wk 60 (Day 421)	152 72.26 (27.859)	4.19 (1.562)	68 72.30 (26.953)	1.04 (2.288)	3.14 (-2.312, 8.600)	0.2575
Wk 72 (Day 505)	148 71.51 (27.784)	4.83 (1.616)	62 72.85 (26.522)	3.47 (2.435)	1.36 (-4.394, 7.116)	0.6419
Wk 84 (Day 589)	134 72.76 (28.506)	6.58 (1.620)	47 72.70 (26.564)	6.29 (2.607)	0.30 (-5.751, 6.346)	0.9229
Wk 96 (Day 673)	129 72.48 (28.386)	7.29 (1.707)	37 71.17 (25.954)	3.83 (2.912)	3.46 (-3.191, 10.111)	0.3065
Average over all visits	219 72.53 (26.806)	2.86 (1.121)	120 72.50 (27.809)	1.62 (1.598)		0.5264
Hedges' g SMD					0.07 (-0.150, 0.296)	0.5205

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.2.4 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Functional scale: Cognitive (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=255		Placebo + bev		Difference between groups		
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Wk 12 (Day 85)	204 81.94 (19.975)	-1.17 (1.089)	111 81.53 (19.378)	-2.82 (1.479)	1.66 (-1.958, 5.269)	0.3681	
Wk 24 (Day 169)	188 81.83 (19.932)	0.35 (1.127)	98 82.48 (17.968)	-2.18 (1.559)	2.53 (-1.261, 6.311)	0.1903	
Wk 36 (Day 253)	166 81.93 (19.455)	-2.11 (1.375)	91 83.88 (16.565)	-3.97 (1.869)	1.86 (-2.709, 6.430)	0.4235	
Wk 48 (Day 337)	164 81.71 (19.988)	-2.40 (1.329)	76 84.43 (15.714)	-2.24 (1.932)	-0.15 (-4.773, 4.465)	0.9477	
Wk 60 (Day 421)	153 81.05 (20.507)	-0.90 (1.273)	67 85.32 (15.495)	-1.91 (1.882)	1.01 (-3.465, 5.494)	0.6560	
Wk 72 (Day 505)	148 82.21 (20.171)	-2.71 (1.393)	59 83.62 (15.000)	-2.08 (2.153)	-0.63 (-5.687, 4.417)	0.8047	
Wk 84 (Day 589)	131 81.30 (19.723)	-2.68 (1.317)	46 82.97 (14.694)	-1.46 (2.098)	-1.21 (-6.096, 3.668)	0.6247	
Wk 96 (Day 673)	129 81.40 (19.613)	-3.35 (1.546)	37 81.53 (14.585)	-1.78 (2.657)	-1.56 (-7.621, 4.494)	0.6116	
Average over all visits	220 81.52 (20.440)	-1.87 (0.930)	118 81.92 (19.069)	-2.31 (1.348)	0.44 (-2.787, 3.660)	0.7903	
Hedges' g SMD					0.03 (-0.193, 0.255)	0.7868	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.2.5 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Functional scale: Emotional (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=25		Placebo + bev		Difference between groups		
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Wk 12 (Day 85)	204 76.44 (22.199)	-1.44 (1.152)	111 77.60 (18.011)	-3.36 (1.562)	1.92 (-1.897, 5.742)	0.3227	
Wk 24 (Day 169)	188 75.72 (22.689)	-1.30 (1.246)	98 78.00 (17.876)	-3.97 (1.722)	2.67 (-1.519, 6.851)	0.2110	
Wk 36 (Day 253)	166 76.32 (22.469)	-0.55 (1.278)	91 79.37 (16.309)	-2.98 (1.744)	2.44 (-1.823, 6.696)	0.2612	
Wk 48 (Day 337)	164 76.13 (22.711)	-2.77 (1.238)	76 79.68 (16.907)	-1.64 (1.807)	-1.13 (-5.447, 3.185)	0.6064	
Wk 60 (Day 421)	153 75.62 (23.432)	-2.30 (1.360)	67 80.35 (17.597)	-4.41 (2.030)	2.11 (-2.714, 6.925)	0.3904	
Wk 72 (Day 505)	148 75.06 (23.536)	-2.44 (1.423)	59 79.19 (18.005)	-1.38 (2.175)	-1.06 (-6.190, 4.060)	0.6828	
Wk 84 (Day 589)	131 73.54 (23.958)	-1.74 (1.410)	46 79.35 (17.457)	-0.80 (2.229)	-0.94 (-6.147, 4.266)	0.7222	
Wk 96 (Day 673)	129 74.55 (23.595)	-3.28 (1.612)	37 78.60 (18.424)	-2.26 (2.737)	-1.02 (-7.285, 5.252)	0.7496	
Average over all visits	220 75.73 (22.447)	-1.98 (0.961)	118 78.23 (17.836)	-2.60 (1.396)	0.62 (-2.717, 3.961)	0.7141	
Hedges' g SMD					0.04 (-0.181, 0.266)	0.7091	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.2.6 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Functional scale: Social (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bev		Placebo + be		Difference between groups		
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Wk 12 (Day 85)	204 74.51 (28.046)	2.55 (1.359)	111 73.72 (25.181)	3.52 (1.844)	-0.97 (-5.475, 3.538)	0.6728	
Wk 24 (Day 169)	188 74.29 (27.434)	4.39 (1.399)	98 73.64 (24.397)	5.61 (1.935)	-1.22 (-5.920, 3.478)	0.6096	
Wk 36 (Day 253)	166 74.30 (27.865)	3.79 (1.526)	91 73.08 (24.441)	5.13 (2.080)	-1.34 (-6.418, 3.742)	0.6045	
Wk 48 (Day 337)	163 73.52 (28.493)	2.02 (1.533)	76 75.44 (23.170)	6.14 (2.228)	-4.12 (-9.445, 1.206)	0.1289	
Wk 60 (Day 421)	153 72.11 (28.793)	7.08 (1.447)	67 73.38 (24.110)	9.92 (2.144)	-2.84 (-7.937, 2.250)	0.2726	
Wk 72 (Day 505)	148 72.41 (28.816)	8.51 (1.537)	59 75.14 (23.030)	9.50 (2.380)	-0.99 (-6.576, 4.591)	0.7265	
Wk 84 (Day 589)	131 73.03 (27.616)	5.57 (1.604)	46 75.36 (23.502)	7.27 (2.557)	-1.70 (-7.652, 4.244)	0.5730	
Wk 96 (Day 673)	129 73.51 (27.998)	4.69 (1.771)	37 70.72 (23.042)	13.72 (3.054)	-9.03 (-15.991, -2.079)	0.0111*	
Average over all visits	220 73.48 (28.160)	4.82 (1.055)	118 73.59 (25.067)	7.60 (1.529)	-2.78 (-6.433, 0.877)	0.1358	
Hedges' g SMD					-0.17 (-0.397, 0.051)	0.1293	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.2.7 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Loss of appetite (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)			Placebo + bev		Difference between groups	
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Wk 12 (Day 85)	204 8.50 (17.624)	10.95 (1.665)	114	8.19 (16.922)	1.56 (2.233)	9.39 (3.908, 14.868)	0.0008*
Wk 24 (Day 169)	187 8.02 (17.312)	6.12 (1.543)	100	8.67 (18.115)	2.54 (2.107)	3.58 (-1.559, 8.718)	0.1715
Wk 36 (Day 253)	163 7.98 (17.698)	5.02 (1.603)	91	8.06 (15.979)	1.54 (2.157)	3.48 (-1.809, 8.772)	0.1962
Wk 48 (Day 337)	165 7.68 (17.509)	6.32 (1.544)	78	6.84 (15.532)	1.83 (2.219)	4.49 (-0.832, 9.812)	0.0979
Wk 60 (Day 421)	152 8.11 (17.992)	2.30 (1.360)	68	6.37 (14.406)	0.52 (2.005)	1.78 (-2.994, 6.549)	0.4638
Wk 72 (Day 505)	146 9.13 (18.578)	2.27 (1.519)	62	6.99 (16.125)	0.25 (2.303)	2.02 (-3.421, 7.454)	0.4658
Wk 84 (Day 589)	132 8.84 (18.355)	-1.82 (1.191)	47	4.26 (13.219)	-2.35 (1.961)	0.54 (-3.994, 5.066)	0.8158
Wk 96 (Day 673)	129 8.01 (17.567)	2.02 (1.589)	37	7.21 (15.977)	0.77 (2.822)	1.25 (-5.138, 7.631)	0.7007
Average over all visits	219 8.07 (17.219)	4.15 (0.958)	120	8.89 (17.677)	0.83 (1.370)	3.31 (0.026, 6.603)	0.0483*
Hedges' g SMD						0.23 (0.005, 0.452)	0.0450*

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.2.8 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Constipation (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be		Placebo + bev		Difference between groups		
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Wk 12 (Day 85)	198 16.84 (27.233)	-1.87 (1.420)	112 14.58 (24.824)	-2.53 (1.891)	0.66 (-3.996, 5.312)	0.7810	
Wk 24 (Day 169)	185 16.40 (27.610)	-1.31 (1.496)	95 13.33 (24.982)	0.25 (2.077)	-1.56 (-6.604, 3.478)	0.5422	
Wk 36 (Day 253)	164 16.06 (26.490)	-0.60 (1.582)	92 10.87 (21.602)	-0.19 (2.139)	-0.41 (-5.652, 4.833)	0.8778	
Wk 48 (Day 337)	163 16.97 (26.802)	-0.84 (1.649)	77 12.12 (22.884)	1.73 (2.362)	-2.57 (-8.247, 3.111)	0.3741	
Wk 60 (Day 421)	149 16.55 (27.020)	4.14 (1.863)	67 11.44 (23.608)	1.22 (2.778)	2.92 (-3.678, 9.519)	0.3842	
Wk 72 (Day 505)	143 17.72 (27.636)	2.89 (1.969)	61 12.02 (24.368)	1.28 (2.980)	1.61 (-5.436, 8.666)	0.6522	
Wk 84 (Day 589)	131 17.05 (27.848)	5.32 (2.039)	46 12.32 (25.684)	2.54 (3.342)	2.78 (-4.944, 10.511)	0.4784	
Wk 96 (Day 673)	126 15.61 (25.884)	3.52 (2.037)	37 17.12 (28.997)	7.66 (3.678)	-4.14 (-12.432, 4.161)	0.3266	
Average over all visits	216 17.44 (27.832)	1.41 (1.100)	118 14.12 (24.424)	1.49 (1.611)	-0.09 (-3.931, 3.757)	0.9643	
Hedges' g SMD					-0.01 (-0.230, 0.219)	0.9635	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.2.9 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Diarrhoea (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be		Placebo + bev		Difference between groups		
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95%	CI) p-value	
Wk 12 (Day 85)	200 9.00 (17.905)	4.09 (1.394)	112 13.69 (25.529)	-1.25 (1.872)	5.34 (0.744, 9	.942) 0.0229*	
Wk 24 (Day 169)	184 8.88 (18.099)	2.81 (1.601)	98 14.29 (26.216)	0.36 (2.204)	2.46 (-2.918, 7	.830) 0.3692	
Wk 36 (Day 253)	163 8.38 (18.274)	0.57 (1.361)	90 14.07 (26.899)	2.17 (1.843)	-1.60 (-6.126, 2	.924) 0.4867	
Wk 48 (Day 337)	162 8.44 (16.744)	1.15 (1.376)	77 11.69 (24.041)	2.56 (1.982)	-1.41 (-6.173, 3	.349) 0.5596	
Wk 60 (Day 421)	149 8.95 (18.434)	0.75 (1.401)	67 11.44 (24.310)	2.16 (2.079)	-1.40 (-6.349, 3	.543) 0.5767	
Wk 72 (Day 505)	145 8.05 (16.328)	0.54 (1.455)	59 11.30 (22.836)	0.45 (2.246)	0.08 (-5.200, 5	.366) 0.9755	
Wk 84 (Day 589)	130 9.49 (18.670)	-2.04 (1.420)	46 10.14 (22.077)	3.07 (2.300)	-5.11 (-10.439, 0	.224) 0.0603	
Wk 96 (Day 673)	126 10.05 (19.446)	0.14 (1.606)	37 8.11 (21.380)	2.50 (2.831)	-2.37 (-8.788, 4	.057) 0.4685	
Average over all visits	217 9.37 (18.956)	1.00 (0.927)	119 13.17 (25.012)	1.50 (1.341)	-0.50 (-3.715, 2	.714) 0.7593	
Hedges' g SMD					-0.04 (-0.259, 0	.188) 0.7546	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.2.10 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Dyspnoea (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=255		Placebo + bev		Difference between groups	
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Wk 12 (Day 85)	198 24.41 (27.369)	2.12 (1.649)	114 22.22 (26.110)	-2.27 (2.184)	4.39 (-0.994, 9.776)	0.1096
Wk 24 (Day 169)	183 24.77 (27.617)	-1.21 (1.642)	100 22.67 (27.167)	-1.59 (2.224)	0.38 (-5.056, 5.825)	0.8895
Wk 36 (Day 253)	163 23.72 (27.895)	-2.27 (1.848)	93 23.66 (27.619)	-0.38 (2.467)	-1.89 (-7.960, 4.175)	0.5397
Wk 48 (Day 337)	164 24.59 (28.328)	-1.10 (1.703)	78 21.37 (25.750)	-5.65 (2.436)	4.55 (-1.303, 10.402)	0.1271
Wk 60 (Day 421)	151 24.28 (28.004)	-1.14 (1.838)	68 23.53 (28.829)	-0.53 (2.673)	-0.61 (-6.997, 5.782)	0.8516
Wk 72 (Day 505)	146 24.89 (28.718)	-1.63 (1.708)	61 24.59 (27.823)	-7.09 (2.574)	5.46 (-0.621, 11.546)	0.0782
Wk 84 (Day 589)	131 24.68 (28.819)	-3.22 (1.776)	47 24.11 (27.542)	-8.14 (2.855)	4.93 (-1.700, 11.552)	0.1443
Wk 96 (Day 673)	128 24.74 (28.137)	-3.44 (1.779)	37 29.73 (29.169)	-1.55 (3.106)	-1.89 (-8.945, 5.159)	0.5973
Average over all visits	216 24.23 (27.174)	-1.48 (1.194)	120 22.78 (26.633)	-3.40 (1.697)	1.92 (-2.167, 5.997)	0.3567
Hedges' g SMD					0.11 (-0.117, 0.330)	0.3495

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[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.2.11 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Fatigue (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=255		Placebo + bev		Difference between groups		
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Wk 12 (Day 85)	202 32.89 (22.878)	6.90 (1.383)	113 34.37 (22.756)	3.27 (1.854)	3.63 (-0.921, 8.18	3) 0.1175	
Wk 24 (Day 169)	187 32.86 (23.303)	1.10 (1.234)	99 33.84 (22.351)	0.66 (1.694)	0.44 (-3.680, 4.56	8) 0.8324	
Wk 36 (Day 253)	164 32.59 (22.573)	1.79 (1.407)	93 34.47 (22.486)	2.02 (1.895)	-0.23 (-4.877, 4.41	7) 0.9224	
Wk 48 (Day 337)	165 32.53 (23.468)	2.11 (1.440)	78 33.40 (21.665)	-1.58 (2.061)	3.69 (-1.254, 8.64	3) 0.1428	
Wk 60 (Day 421)	152 32.53 (22.744)	0.44 (1.446)	68 34.31 (22.138)	-2.73 (2.121)	3.17 (-1.883, 8.23	0) 0.2177	
Wk 72 (Day 505)	147 33.26 (23.686)	-2.81 (1.479)	62 33.06 (21.980)	-0.74 (2.227)	-2.07 (-7.339, 3.19	3) 0.4389	
Wk 84 (Day 589)	133 33.00 (23.046)	-1.69 (1.474)	47 30.85 (21.354)	-5.42 (2.337)	3.73 (-1.715, 9.16	9) 0.1786	
Wk 96 (Day 673)	129 32.30 (23.009)	-3.15 (1.587)	37 31.98 (21.573)	-6.40 (2.730)	3.24 (-2.979, 9.46	6) 0.3055	
Average over all visits	218 32.77 (22.890)	0.59 (1.022)	119 34.50 (22.536)	-1.36 (1.459)	1.95 (-1.554, 5.45	7) 0.2743	
Hedges' g SMD					0.13 (-0.097, 0.35	0) 0.2675	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.2.12 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Financial difficulties (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be		Placebo + bev		Difference between groups		
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE)	Estimated difference (95%	cI)	p-value
Wk 12 (Day 85)	201 18.24 (29.796)	-0.89 (1.388)	110 18.79 (29.445)	0.74 (1.879)	-1.63 (-6.226,	2.968)	0.4863
Wk 24 (Day 169)	183 20.04 (31.242)	-1.55 (1.467)	97 19.24 (29.594)	-2.02 (2.024)	0.47 (-4.451,	5.386)	0.8516
Wk 36 (Day 253)	162 18.31 (29.029)	-1.79 (1.592)	90 19.26 (30.386)	-3.92 (2.170)	2.13 (-3.167,	7.426)	0.4293
Wk 48 (Day 337)	162 18.52 (29.488)	-1.92 (1.554)	76 20.18 (30.346)	-3.55 (2.231)	1.63 (-3.720,	6.986)	0.5486
Wk 60 (Day 421)	150 18.89 (29.768)	-2.73 (1.677)	65 18.46 (28.886)	-1.56 (2.463)	-1.17 (-7.042,	4.694)	0.6940
Wk 72 (Day 505)	144 19.21 (30.172)	-1.77 (1.720)	58 18.39 (29.401)	-1.62 (2.618)	-0.15 (-6.319,	6.013)	0.9611
Wk 84 (Day 589)	130 17.95 (28.801)	-2.34 (1.712)	46 18.84 (31.152)	-1.61 (2.665)	-0.73 (-6.965,	5.508)	0.8183
Wk 96 (Day 673)	128 19.27 (29.463)	-3.64 (1.708)	37 25.23 (33.707)	-6.26 (2.877)	2.62 (-3.966,	9.214)	0.4336
Average over all visits	218 18.50 (30.006)	-2.08 (1.206)	117 18.80 (29.809)	-2.47 (1.730)	0.40 (-3.752,	4.544)	0.8510
Hedges' g SMD					0.02 (-0.203,	0.246)	0.8490

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.2.13 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Nausea and vomiting (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)			Placebo + bev		Difference between groups	
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Wk 12 (Day 85)	203 4.52 (12.192)	10.64 (1.147)	114	3.22 (9.109)	0.33 (1.537)	10.31 (6.534, 14.084)	<0.0001*
Wk 24 (Day 169)	188 4.52 (12.499)	5.80 (0.889)	100	3.00 (8.335)	-0.70 (1.223)	6.50 (3.525, 9.479)	<0.0001*
Wk 36 (Day 253)	165 4.14 (10.477)	6.03 (0.985)	93	3.05 (8.130)	1.80 (1.327)	4.23 (0.975, 7.483)	0.0111*
Wk 48 (Day 337)	165 4.75 (12.858)	7.90 (1.138)	78	1.92 (5.995)	1.86 (1.639)	6.05 (2.111, 9.981)	0.0027*
Wk 60 (Day 421)	152 4.82 (13.098)	6.41 (1.063)	68	2.45 (6.607)	-0.33 (1.550)	6.74 (3.033, 10.445)	0.0004*
Wk 72 (Day 505)	147 5.10 (13.460)	5.21 (1.027)	62	2.15 (6.390)	-1.12 (1.543)	6.33 (2.671, 9.988)	0.0008*
Wk 84 (Day 589)	133 5.14 (13.481)	4.47 (1.093)	47	2.13 (6.609)	0.91 (1.772)	3.56 (-0.550, 7.677)	0.0892
Wk 96 (Day 673)	129 5.17 (13.944)	3.28 (1.148)	37	2.70 (7.363)	3.57 (2.057)	-0.29 (-4.940, 4.366)	0.9033
Average over all visits Hedges' g SMD	219 4.41 (11.873)	6.22 (0.709)	120	3.33 (9.065)	0.79 (1.013)	5.43 (2.995, 7.863) 0.51 (0.280, 0.732)	<0.0001* <0.0001*

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.2.14 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Pain (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=255		Placebo + bev (N=132		Difference between g	Difference between groups	
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from	n Mean (SD) at	Mean (SE)		p-value	
Wk 12 (Day 85)	205 21.95 (23.056)	7.42 (1.541)	114 23.54 (22.916)	9.13 (2.072)	-1.70 (-6.784, 3.376)	0.5098	
Wk 24 (Day 169)	188 22.78 (23.560)	4.71 (1.630)	101 23.27 (21.868)	6.85 (2.226)	-2.14 (-7.571, 3.287)	0.4381	
Wk 36 (Day 253)	166 22.29 (22.738)	6.81 (1.766)	93 23.12 (22.392)	9.01 (2.378)	-2.20 (-8.027, 3.634)	0.4590	
Wk 48 (Day 337)	165 22.63 (23.277)	6.66 (1.800)	78 22.44 (22.606)	5.77 (2.578)	0.89 (-5.299, 7.081)	0.7771	
Wk 60 (Day 421)	155 22.80 (23.264)	2.68 (1.622)	68 23.04 (22.310)	5.78 (2.400)	-3.10 (-8.804, 2.604)	0.2855	
Wk 72 (Day 505)	150 23.33 (23.412)	1.98 (1.653)	62 22.85 (23.422)	2.85 (2.499)	-0.87 (-6.774, 5.028)	0.7710	
Wk 84 (Day 589)	133 22.81 (24.228)	-0.80 (1.737)	47 22.34 (23.384)	2.25 (2.817)	-3.05 (-9.573, 3.470)	0.3575	
Wk 96 (Day 673)	130 22.44 (22.989)	-0.64 (1.752)	37 23.42 (23.060)	-4.03 (3.081)	3.39 (-3.593, 10.382)	0.3393	
Average over all visits	221 22.78 (23.173)	3.60 (1.166)	120 23.47 (22.917)	4.70 (1.669)	-1.10 (-5.103, 2.908)		
Hedges' g SMD					-0.06 (-0.284, 0.160)	0.5847	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.2.15 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Insomnia (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bev		Placebo + bev		Difference between groups		
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Wk 12 (Day 85)	202 27.72 (29.392)	4.11 (1.809)	114 23.10 (25.524)	7.72 (2.408)	-3.61 (-9.540, 2.329)	0.2329	
Wk 24 (Day 169)	187 28.70 (30.369)	5.55 (1.968)	100 22.00 (24.718)	4.22 (2.679)	1.33 (-5.225, 7.892)	0.6894	
Wk 36 (Day 253)	164 27.03 (29.439)	5.42 (2.022)	93 21.15 (23.460)	4.92 (2.718)	0.50 (-6.183, 7.183)	0.8830	
Wk 48 (Day 337)	164 26.02 (28.850)	4.90 (2.017)	78 21.37 (22.776)	3.46 (2.907)	1.44 (-5.536, 8.417)	0.6846	
Wk 60 (Day 421)	149 27.07 (29.093)	7.41 (2.119)	68 20.10 (23.843)	7.79 (3.092)	-0.38 (-7.781, 7.018)	0.9192	
Wk 72 (Day 505)	146 26.94 (29.378)	2.50 (1.994)	62 22.04 (24.100)	4.75 (3.021)	-2.25 (-9.392, 4.892)	0.5355	
Wk 84 (Day 589)	132 27.53 (30.958)	4.01 (2.161)	47 20.57 (24.626)	5.76 (3.485)	-1.74 (-9.844, 6.354)	0.6715	
Wk 96 (Day 673)	127 26.25 (30.176)	4.62 (2.331)	37 25.23 (26.534)	7.36 (3.935)	-2.73 (-11.755, 6.287)	0.5509	
Average over all visits	217 28.26 (30.258)	4.81 (1.421)	120 22.50 (25.258)	5.74 (2.031)	-0.93 (-5.818, 3.957)	0.7083	
Hedges' g SMD					-0.04 (-0.266, 0.180)	0.7031	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Figure 2.5.3.1 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Global QoL/health status across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

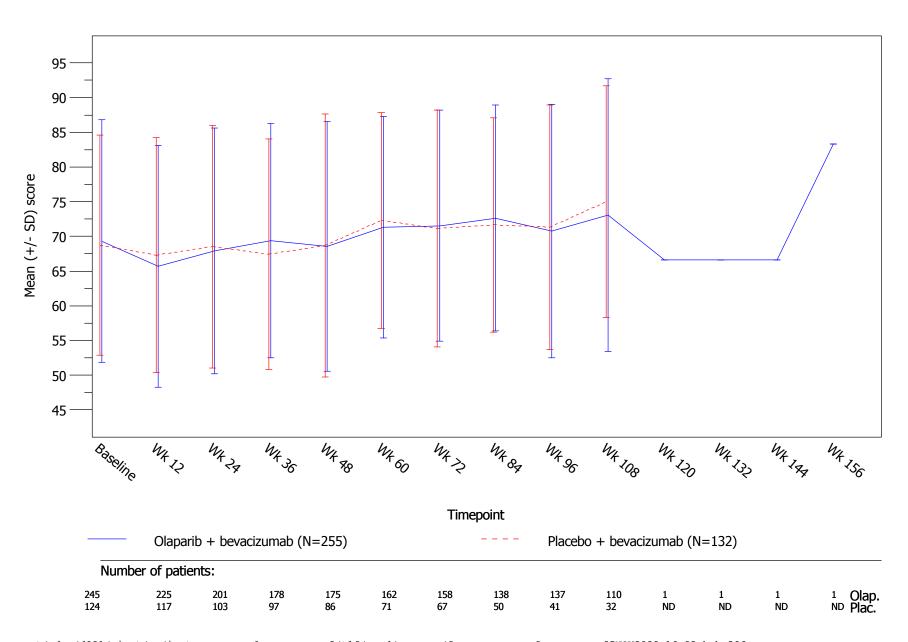


Figure 2.5.3.2 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Functional scale: Physical across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

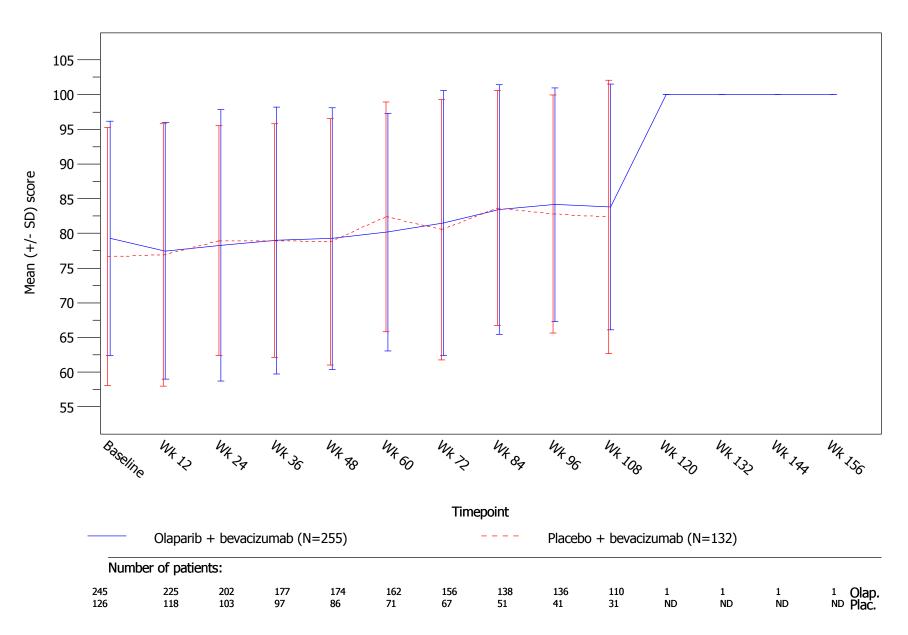


Figure 2.5.3.3 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Functional scale: Role across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

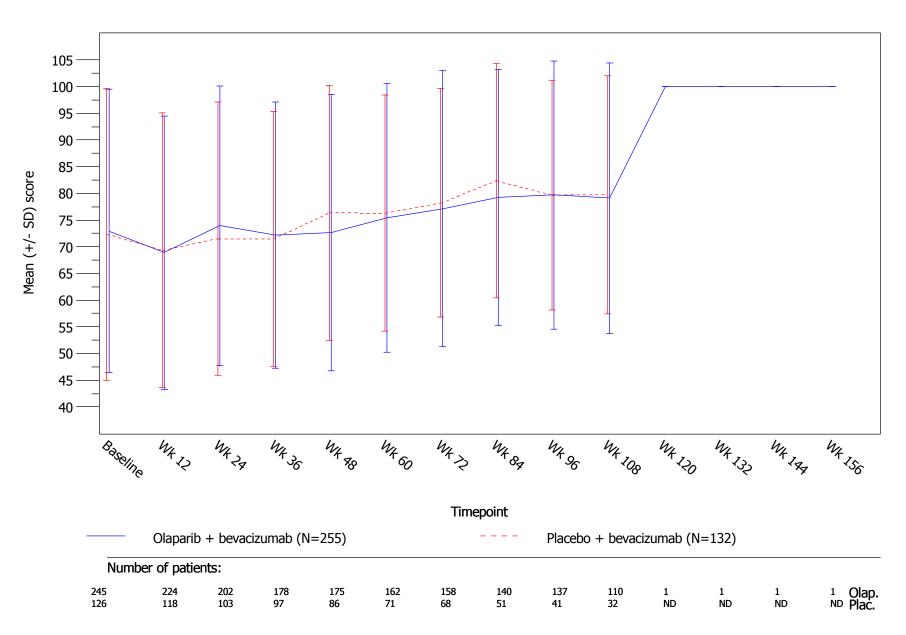


Figure 2.5.3.4 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Functional scale: Cognitive across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

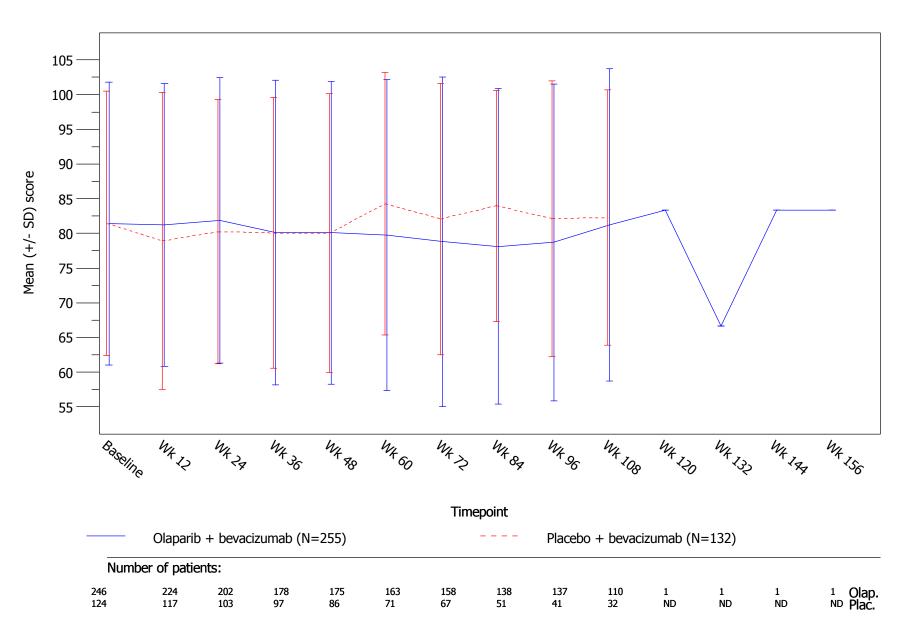


Figure 2.5.3.5 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Functional scale: Emotional across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

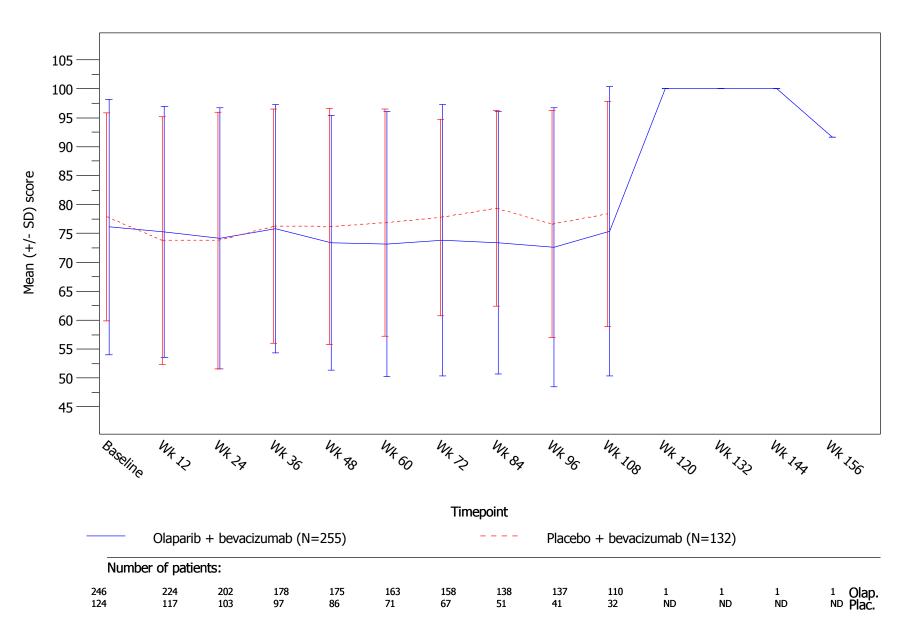


Figure 2.5.3.6 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Functional scale: Social across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

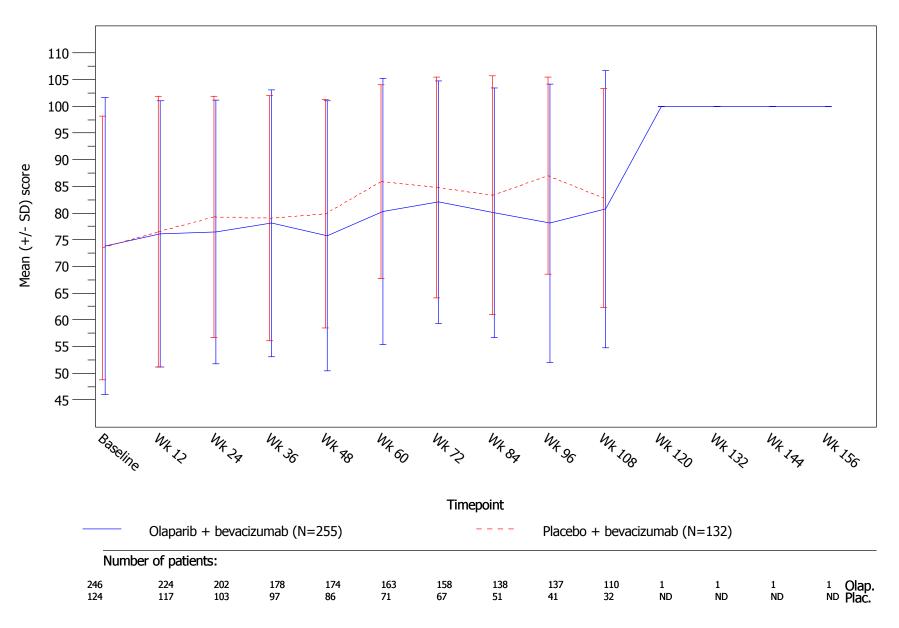


Figure 2.5.3.7 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Single item symptom scale: Loss of appetite across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

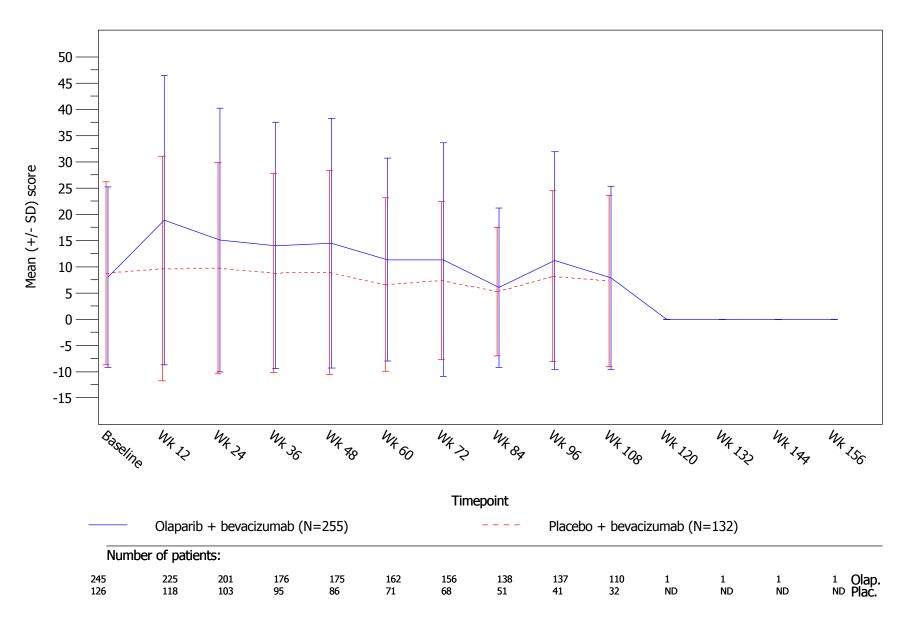


Figure 2.5.3.8 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Single item symptom scale: Constipation across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

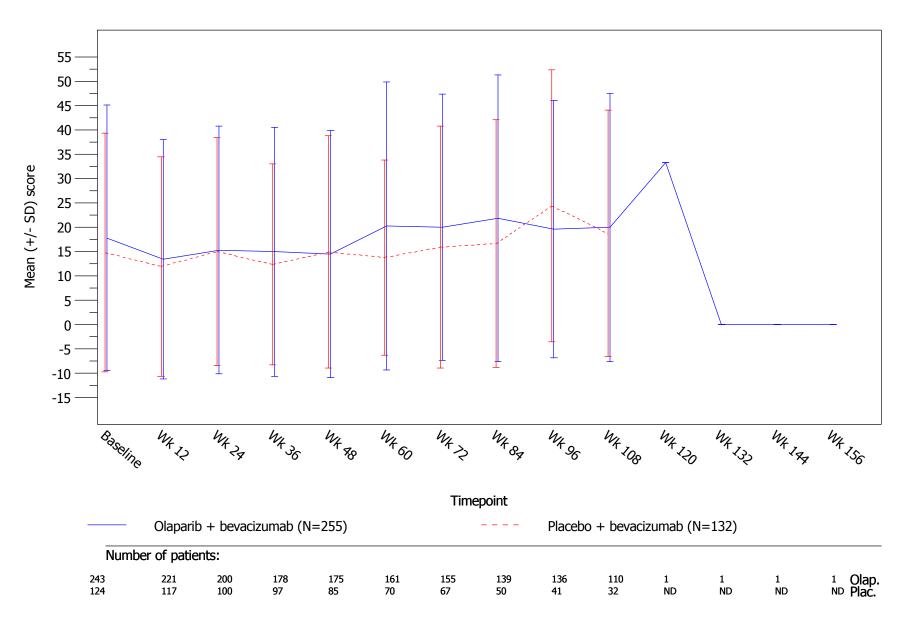


Figure 2.5.3.9 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Single item symptom scale: Diarrhoea across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

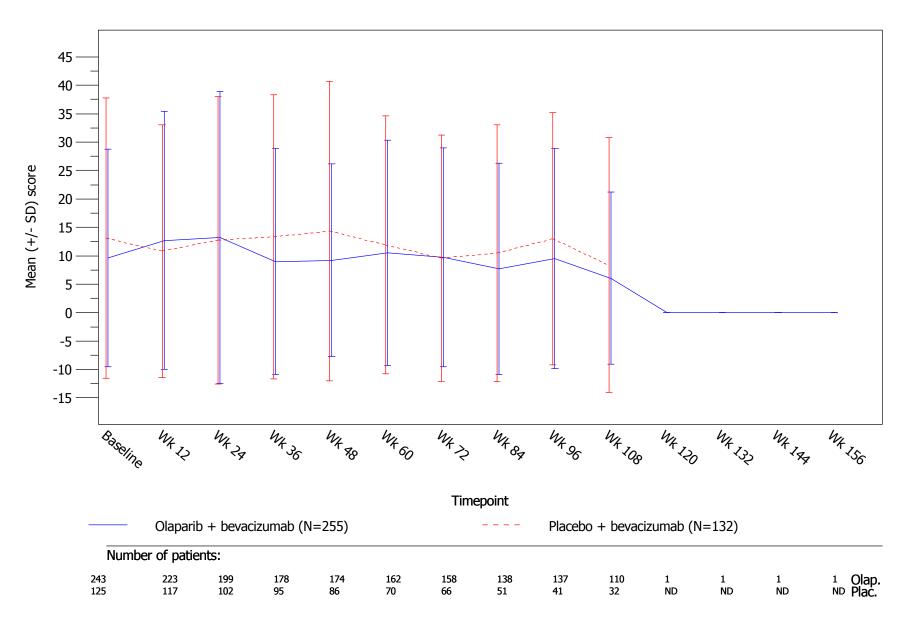


Figure 2.5.3.10 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Single item symptom scale: Dyspnoea across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

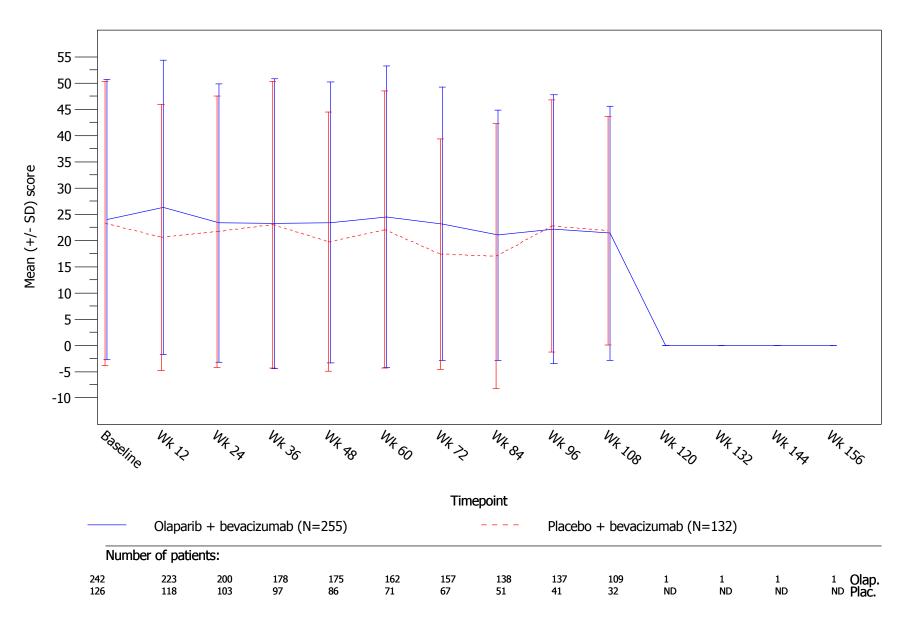


Figure 2.5.3.11 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Symptom scale: Fatigue across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

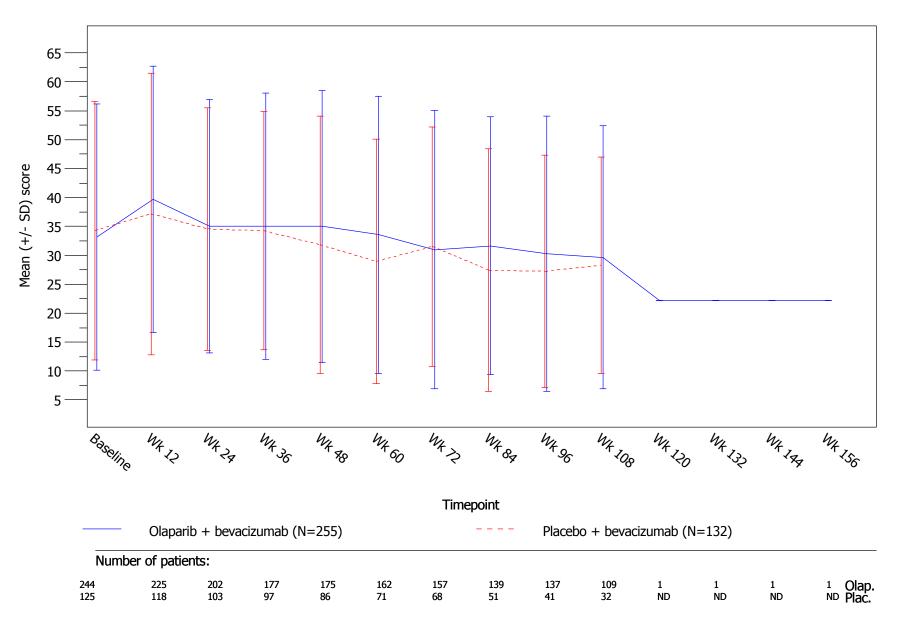


Figure 2.5.3.12 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Single item symptom scale: Financial difficulties across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

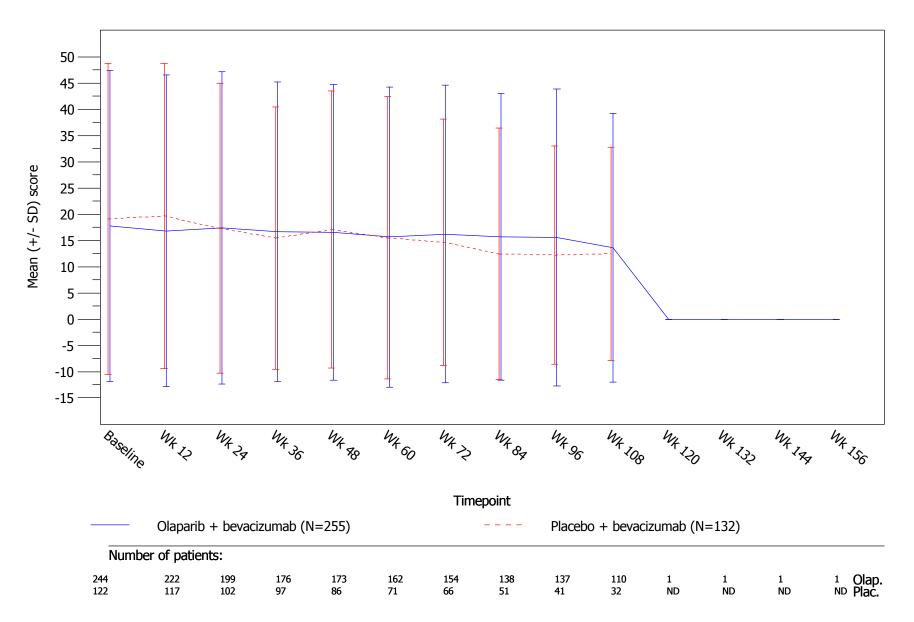


Figure 2.5.3.13 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Symptom scale: Nausea and vomiting across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

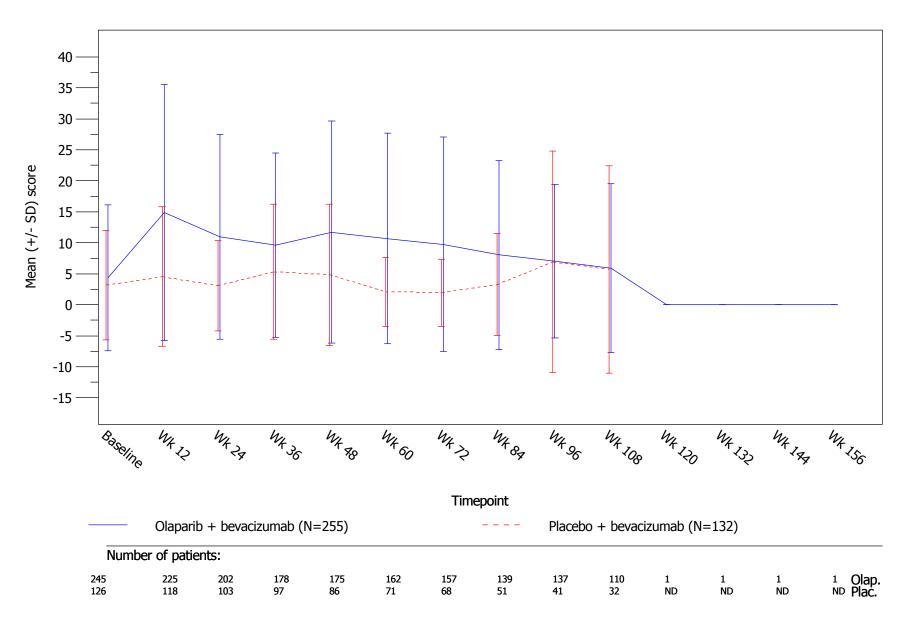


Figure 2.5.3.14 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Symptom scale: Pain across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

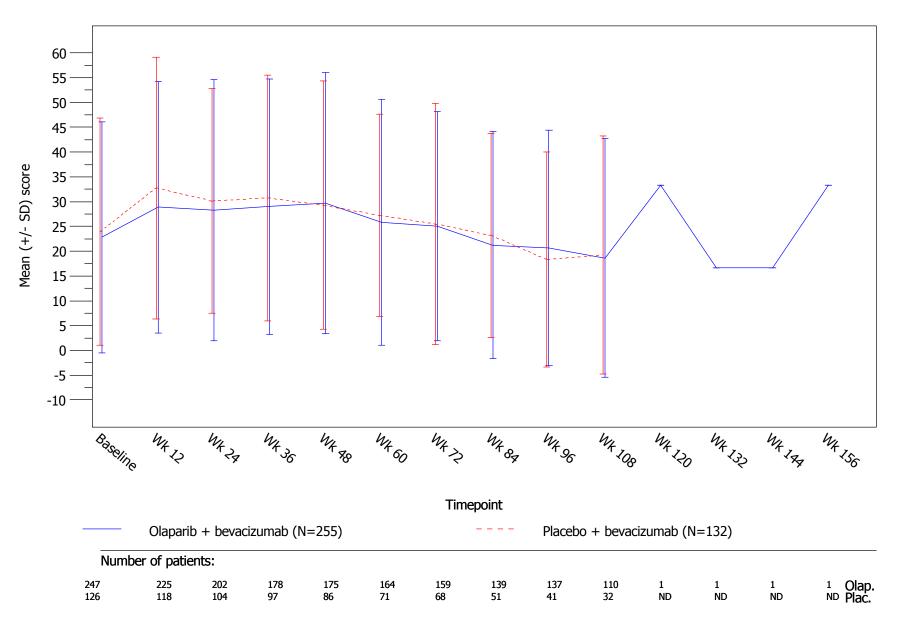


Figure 2.5.3.15 PAOLA1: Mean (+/- SD) score for EORTC QLQ-C30 Single item symptom scale: Insomnia across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

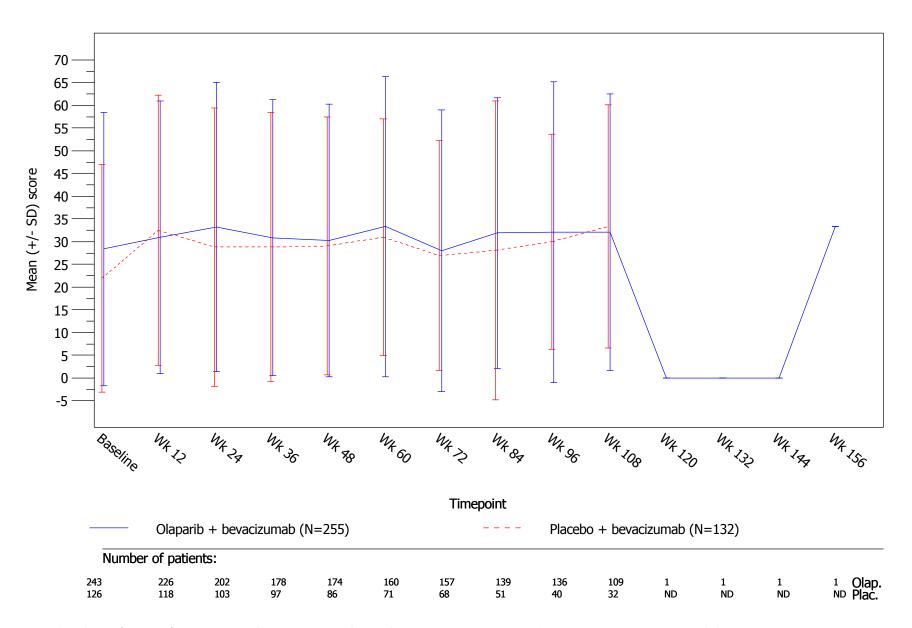


Table 2.5.4.1 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Global QoL/health status (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=25!		Placebo + bev		Difference between g	roups
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
First line treat	ment outcome (IVRS)					
NED [PDS]	81 70.16 (17.523)	2.58 (1.201)	43 66.47 (16.008)	1.05 (1.732)	1.52 (-2.658, 5.706)	0.4717
Hedges' g SMD					0.14 (-0.233, 0.508)	0.4660
NED/CR [IDS]	61 69.95 (17.165)	-1.26 (1.412)	33 70.20 (17.057)	2.76 (2.089)	-4.02 (-9.043, 0.995)	0.1146
Hedges' g SMD					-0.35 (-0.777, 0.076)	0.1071
NED/CR [Chemo]	37 66.22 (16.776)	1.08 (2.265)	20 69.58 (12.174)	-2.89 (3.235)	3.97 (-4.024, 11.965)	0.3214
Hedges' g SMD					0.28 (-0.267, 0.826)	0.3165
PR	40 73.33 (16.473)	-3.40 (2.243)	22 68.94 (17.851)	-6.99 (3.290)	3.59 (-4.391, 11.572)	0.3713
Hedges' g SMD					0.24 (-0.280, 0.764)	0.3631
Int. p-value						0.0723
tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value	atory tBRCA status (I 129 67.05 (17.112) 90 74.26 (16.319)	2.26 (1.046)	59 68.50 (15.246) 59 68.50 (16.815)		1.89 (-1.957, 5.728) 0.16 (-0.153, 0.464) -0.78 (-4.752, 3.185) -0.07 (-0.395, 0.262)	0.3341 0.3244 0.6965 0.6912 0.5770
First line treat	tment outcome (eCRF)					
NED [PDS]	79 69.62 (18.004)	2.48 (1.218)	42 66.87 (15.991)	0.83 (1.728)	1.66 (-2.539, 5.849)	0.4357
Hedges' g SMD					0.15 (-0.225, 0.525)	0.4325
NED/CR [IDS]	60 71.11 (17.390)	-0.58 (1.347)	28 70.83 (18.356)	2.69 (2.209)	-3.27 (-8.422, 1.877)	0.2095
Hedges' g SMD					-0.30 (-0.750, 0.152)	0.1934
NED/CR [Chemo]	34 65.93 (17.087)	1.73 (2.505)	17 72.06 (12.127)	-5.35 (4.048)		0.1482
Hedges' g SMD					0.46 (-0.134, 1.045)	0.1301
PR	43 72.48 (15.594)	-3.76 (2.129)	30 67.22 (15.618)	-3.60 (2.723)		0.9631
Hedges' g SMD					-0.01 (-0.477, 0.455)	0.9628
Int. p-value						0.0327

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.1 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Global QoL/health status (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + be		Difference between groups		
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Screening labor	atory tE	BRCA status (e	CRF)						
tBRCAm	126 66	.87 (17.239)	2.23 (1.064)	60	67.92 (15.783)	0.56 (1.632)	1.67 (-2.184, 5.514)	0.3943	
Hedges' g SMD							0.14 (-0.171, 0.444)	0.3855	
non-tBRCAm	93 74	.28 (16.098)	-2.07 (1.170)	58	69.11 (16.298)	-1.51 (1.606)	-0.57 (-4.516, 3.378)	0.7760	
Hedges' g SMD							-0.05 (-0.377, 0.279)	0.7717	
Int. p-value								0.7156	
Age group									
<65 years	160 69	.69 (17.296)	0.16 (0.961)	89	67.98 (15.686)	0.00 (1.368)	0.16 (-3.133, 3.460)	0.9224	
Hedges' g SMD							0.01 (-0.246, 0.272)	0.9211	
>=65 years	59 70	.90 (16.765)	0.56 (1.330)	29	70.11 (17.037)	-2.23 (2.072)	2.79 (-2.116, 7.695)	0.2612	
Hedges' g SMD							0.26 (-0.184, 0.709)	0.2489	
Int. p-value								0.3867	
FIGO Stage (Dis	ease sta	ate)							
III	157 69	.80 (17.838)	0.09 (0.856)	81	70.06 (16.914)	1.55 (1.254)	-1.46 (-4.457, 1.531)	0.3366	
Hedges' g SMD							-0.13 (-0.402, 0.135)	0.3294	
IV	62 70	.56 (15.293)	1.00 (1.718)	37	65.09 (13.298)	-5.41 (2.561)	6.41 (0.254, 12.559)	0.0414	
Hedges' g SMD							0.44 (0.032, 0.856)	0.0348	
Int. p-value								0.1424	
Region									
Europe	210 69	.88 (17.136)	0.41 (0.800)	112	68.38 (16.177)	-0.12 (1.177)	0.53 (-2.274, 3.331)	0.7106	
Hedges' g SMD							0.04 (-0.185, 0.274)	0.7049	
Japan	9	NC	NC	6	NC	NC	NC	NC	
Int. p-value								NC	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.1 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Global QoL/health status (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		+ bevacizumab (N=255)	Placebo + be(N=13		Difference between groups		
Subgroup	n Mean (SD)		n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
ECOG performance	e status at Base	eline					
(0) Normal activity	162 71.04 (17.	106) -0.56 (0.920)	90 70.93 (14.741)	-1.70 (1.287)	1.15 (-1.974,	4.265)	0.4699
Hedges' g SMD					0.10 (-0.162,		0.4653
(1) Restricted activity	54 66.51 (17.	169) 4.12 (1.519)	28 60.71 (17.547)	3.80 (2.391)	0.33 (-5.331,	5.990)	0.9080
Hedges' g SMD Int. p-value					0.03 (-0.429,	0.484)	0.9046 0.5934
Baseline CA-125	value						
<=ULN	194 69.76 (17.	0.98 (0.813)	105 68.49 (16.137)	-0.43 (1.181)	, ,		0.3275
Hedges' g SMD >ULN Int. p-value	25 NC	NC	13 NC	NC	0.12 (-0.117, NC	0.359)	0.3184 NC NC
Histological gra	ade						
High grade	219 70.02 (17.	125) 0.36 (0.783)	118 68.50 (15.981)	-0.31 (1.136)	0.67 (-2.042,	3.391)	0.6256
Hedges' g SMD Int. p-value					0.06 (-0.167,	0.281)	0.6194 ID
Cytoreductive s	urgery outcome						
No residue	142 70.25 (17.	514) 1.16 (0.878)	71 68.19 (16.980)	1.39 (1.314)	-0.23 (-3.346,	2.892)	0.8858
Hedges' g SMD					-0.02 (-0.306,	0.264)	0.8839
Residue	70 68.57 (16.	125) -0.90 (1.607)	39 70.51 (14.287)	-2.70 (2.277)	, ,		0.5214
Hedges' g SMD					0.13 (-0.262,	0.522)	0.5155
Int. p-value							0.5274

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.1 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Global QoL/health status (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + 1 (N=2		Placebo + ber		Difference between groups		
Subgroup	n Mean (SD) at [a] Baseline [b]		n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Timing of cytore	eductive surgery						
Upfront	130 69.68 (17.207)	1.66 (0.955)	70 67.86 (15.689)	-0.36 (1.356)	2.02 (-1.253, 5.297	0.2246	
Hedges' g SMD					0.18 (-0.109, 0.473	0.2196	
Interval	82 69.72 (16.897)	-1.17 (1.325)	40 71.04 (16.665)	1.50 (2.084)	-2.68 (-7.574, 2.216	0.2806	
Hedges' g SMD					-0.21 (-0.594, 0.164	0.2667	
Int. p-value						0.0265	
Myriad tumour BI	RCA mutation status						
tBRCAm	135 67.28 (17.116)	1.80 (1.065)	68 67.89 (15.458)	0.70 (1.590)	1.10 (-2.674, 4.881	0.5650	
Hedges' g SMD					0.09 (-0.204, 0.379	0.5583	
Non-tBRCAm	84 74.40 (16.300)	-1.96 (1.165)	50 69.33 (16.789)	-1.96 (1.626)	0.00 (-3.976, 3.970	0.9988	
Hedges' g SMD					0.00 (-0.350, 0.350	0.9988	
Int. p-value						0.9527	
Status somatic E	BRCA mutations						
sBRCAm	18 68.06 (18.358)	0.64 (2.309)	6 69.44 (18.758)	4.30 (3.976)	-3.65 (-13.239, 5.934	0.4358	
Hedges' g SMD					-0.36 (-1.291, 0.570	0.4477	
gBRCAm	55 68.48 (16.873)	-0.24 (1.543)	28 67.56 (15.272)	2.12 (2.342)	-2.36 (-7.942, 3.224	0.4028	
Hedges' g SMD					-0.20 (-0.655, 0.257	0.3934	
Non-BRCAm	37 72.97 (14.083)	-1.99 (1.816)	21 69.44 (15.884)	-3.52 (2.675)	1.54 (-4.959, 8.035	0.6364	
Hedges' g SMD					0.13 (-0.404, 0.668	0.6294	
Int. p-value						0.5435	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.2 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Physical (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be [.] (N=255		Placebo + bev (N=132		Difference between groups			
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value	
First line treat	tment outcome (IVRS)							
NED [PDS]	81 78.81 (17.299)	2.25 (1.360)	43 76.71 (17.700)	0.99 (1.943)	1.26 (-3.441,	5.956)	0.5972	
Hedges' g SMD					0.10 (-0.269,	0.471)	0.5939	
NED/CR [IDS]	61 81.80 (14.673)	1.69 (1.474)	34 78.63 (19.816)	3.52 (2.061)	-1.82 (-6.869,	3.224)	0.4748	
Hedges' g SMD					-0.15 (-0.575,	0.266)	0.4708	
NED/CR [Chemo]	36 77.96 (18.573)	1.52 (2.111)	20 75.17 (19.749)	5.44 (3.002)	-3.92 (-11.331,	3.495)	0.2923	
Hedges' g SMD					-0.30 (-0.848,	0.251)	0.2870	
PR	41 78.88 (17.920)	2.35 (2.050)	23 74.13 (19.455)	0.99 (2.881)	1.37 (-5.706,	8.444)	0.7003	
Hedges' g SMD					0.10 (-0.410,	0.612)	0.6982	
Int. p-value							0.5920	
Screening labora tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value	atory tBRCA status (I 130 78.27 (16.483) 89 81.34 (17.417)	7RS) 2.32 (1.111) 1.27 (1.272)	59 75.62 (19.512) 61 77.35 (18.258)		0.34 (-3.691, 0.03 (-0.281, -1.64 (-5.683, -0.13 (-0.460,	0.334) 2.410)	0.8667 0.8650 0.4252 0.4223 0.9008	
First line treat	tment outcome (eCRF)							
NED [PDS]	79 77.09 (17.414)	2.13 (1.401)	42 78.25 (17.611)	0.93 (1.973)	1.20 (-3.594,	5.994)	0.6210	
Hedges' g SMD					0.09 (-0.280,	0.469)	0.6195	
NED/CR [IDS]	60 83.28 (14.304)	1.36 (1.451)	29 79.54 (21.077)	2.65 (2.194)	-1.29 (-6.536,	3.954)	0.6258	
Hedges' g SMD					-0.11 (-0.556,	0.332)	0.6208	
NED/CR [Chemo]	33 79.39 (18.361)	0.28 (2.344)	17 76.08 (20.284)	4.80 (3.555)	-4.52 (-13.158,	4.117)	0.2958	
Hedges' g SMD					-0.32 (-0.910,	0.268)	0.2857	
PR	44 78.65 (17.911)	2.97 (1.924)	31 72.10 (17.422)	3.51 (2.434)	-0.54 (-6.746,	5.665)	0.8626	
Hedges' g SMD					-0.04 (-0.501,	0.419)	0.8616	
Int. p-value							0.7329	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.2 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Physical (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bev (N=255			Placebo + bev (N=132		Difference between groups			
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value	
Screening labor	atory tE	BRCA status (e0	CRF)							
tBRCAm	127 78	.12 (16.622)	2.25 (1.129)	60	75.92 (19.481)	2.02 (1.708)	0.23 (-3.813,	4.269)	0.9114	
Hedges' g SMD							0.02 (-0.289,	0.325)	0.9104	
non-tBRCAm	92 81	.44 (17.175)	1.44 (1.243)	60	77.08 (18.293)	2.90 (1.601)	-1.46 (-5.484,	2.556)	0.4725	
Hedges' g SMD							-0.12 (-0.446,	0.205)	0.4689	
Int. p-value									0.9720	
Age group										
<65 years	161 79	.02 (17.756)	1.96 (0.955)	90	76.61 (18.107)	3.07 (1.326)	-1.11 (-4.330,	2.112)	0.4982	
Hedges' g SMD							-0.09 (-0.348,	0.168)	0.4942	
>=65 years	58 80	.89 (14.294)	1.51 (1.754)	30	76.17 (21.164)	1.07 (2.573)	0.44 (-5.769,	6.654)	0.8876	
Hedges' g SMD							0.03 (-0.409,	0.473)	0.8860	
Int. p-value									0.7919	
FIGO Stage (Dis	ease sta	ate)								
III	156 78	.72 (17.033)	1.59 (1.024)	83	77.15 (20.089)	2.93 (1.452)	-1.35 (-4.847,	2.156)	0.4499	
Hedges' g SMD							-0.10 (-0.370,	0.163)	0.4460	
IV	63 81	.48 (16.524)	2.98 (1.319)	37	75.05 (15.784)	1.13 (1.906)	1.85 (-2.762,	6.464)	0.4275	
Hedges' g SMD							0.17 (-0.238,	0.575)	0.4164	
Int. p-value									0.7442	
Region										
Europe	210 79	.59 (16.904)	1.71 (0.857)	114	76.73 (18.725)	1.80 (1.220)	-0.10 (-3.032,	2.840)	0.9486	
Hedges' g SMD							-0.01 (-0.236,	0.220)	0.9479	
Japan	9	NC	NC	6	NC	NC	NC		NC	
Int. p-value									NC	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.2 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Physical (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

n Mean (SD) at [a] Baseline [b] status at Baseline [62 81.93 (15.225)	Mean (SE) Change from baseline 0.73 (0.926)	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95	% CI)	p-value
162 81.93 (15.225)	0.73 (0.926)	92 78.80 (17.575)	1.09 (1.264)	-0.36 (-3.453,	0 500	
162 81.93 (15.225)	0.73 (0.926)	92 78.80 (17.575)	1.09 (1.264)	-0.36 (-3.453,	0. 500:	
,	0.73 (0.926)	92 78.80 (17.575)	1.09 (1.264)	-0.36 (-3.453,	0 500:	
53 72.71 (19.653)				,	2.730)	0.8181
53 72.71 (19.653)				-0.03 (-0.286,		0.8168
	6.74 (1.873)	28 68.93 (21.058)	6.63 (2.808)	0.11 (-6.618,	6.842)	0.9737
				0.01 (-0.450,	0.466)	0.9731
						0.9550
alue						
194 79.80 (16.894)	2.18 (0.859)	107 77.21 (18.742)	1.95 (1.208)	, ,		0.8737
25 NC	NC	13 NC	NC		0.255)	0.8723 NC
25 NC	IVC	15 110	INC	IVC		NC
2						
	1.93 (0.833)	120 76.50 (18.826)	2.39 (1.174)	-0.46 (-3.295,	2.375)	0.7499
				-0.04 (-0.259,	0.186)	0.7470
						ID
gery outcome						
142 79.81 (16.332)	1.97 (0.995)	72 78.50 (18.987)	1.56 (1.447)			0.8156
				0.03 (-0.250,	0.318)	0.8139
70 78.39 (18.443)	1.89 (1.603)	40 73.96 (18.036)	4.75 (2.237)	, ,	,	0.3003
				-0.21 (-0.598,	0.181)	0.2947 0.2580
	lue 94 79.80 (16.894) 25 NC 19 79.52 (16.897)	lue 94 79.80 (16.894) 2.18 (0.859) 25 NC NC 19 79.52 (16.897) 1.93 (0.833) Hery outcome 42 79.81 (16.332) 1.97 (0.995)	lue 94 79.80 (16.894) 2.18 (0.859) 107 77.21 (18.742) 25 NC NC 13 NC 19 79.52 (16.897) 1.93 (0.833) 120 76.50 (18.826) erry outcome 42 79.81 (16.332) 1.97 (0.995) 72 78.50 (18.987)	lue 94 79.80 (16.894) 2.18 (0.859) 107 77.21 (18.742) 1.95 (1.208) 25 NC NC 13 NC NC 19 79.52 (16.897) 1.93 (0.833) 120 76.50 (18.826) 2.39 (1.174) Hery outcome 42 79.81 (16.332) 1.97 (0.995) 72 78.50 (18.987) 1.56 (1.447)	1ue 94 79.80 (16.894) 2.18 (0.859) 107 77.21 (18.742) 2.5 NC NC 13 NC NC 1.97 (0.995) 1.93 (0.833) 1.97 (0.995) 1.97 (0.995) 1.98 (18.987) 1.98 (1.447) 1.96 (1.447) 1.97 (0.41 (-3.052, 0.03 (-0.250, 70 78.39 (18.443)) 1.89 (1.603) 40 73.96 (18.036) 4.75 (2.237) -2.87 (-8.336,	104

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.2 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Physical (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be ⁻ (N=255			Placebo + bev		Difference between groups			
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value		
Timing of cytore	oduati	uo gurgoru								
Upfront Hedges' g SMD		77.55 (18.124)	2.80 (1.113)		76.20 (18.008)	2.03 (1.545)	0.77 (-2.989, 4.526) 0.06 (-0.230, 0.350)	0.6852		
Interval Hedges' g SMD Int. p-value	83	82.13 (14.846)	0.56 (1.295)	41	78.05 (20.014)	4.24 (1.946)	-3.68 (-8.317, 0.955) -0.30 (-0.681, 0.072)			
Myriad tumour BI	RCA mu	tation status								
tBRCAm Hedges' g SMD		78.54 (16.318)	2.36 (1.071)	68	76.40 (18.934)	2.62 (1.577)	-0.26 (-4.017, 3.504) -0.02 (-0.311, 0.271)			
Non-tBRCAm Hedges' g SMD Int. p-value	83	81.11 (17.790)	1.15 (1.351)	52	76.63 (18.866)	2.18 (1.772)	-1.03 (-5.459, 3.392) -0.08 (-0.429, 0.265)			
Status somatic H	BRCA m	utations								
sBRCAm Hedges' g SMD		80.00 (14.642)	5.49 (2.490)	6	80.00 (20.221)	5.17 (4.404)	0.32 (-10.341, 10.972) 0.03 (-0.895, 0.953)			
gBRCAm Hedges' g SMD	57	78.45 (16.832)	-0.32 (1.644)	28	75.95 (20.314)	4.01 (2.460)	-4.33 (-10.213, 1.559) -0.34 (-0.795, 0.115)	0.1475		
Non-BRCAm Hedges' g SMD Int. p-value	36	82.79 (17.525)	2.83 (1.759)	22	79.32 (16.893)	5.44 (2.350)	-2.61 (-8.520, 3.307) -0.24 (-0.772, 0.293)			

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.3 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Role (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + b (N=2!		Placebo + be		Difference between groups		
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
First line treat	ment outcome (IVRS)						
NED [PDS]	81 70.37 (27.639)	4.17 (1.868)	43 73.26 (24.704)	1.51 (2.663)	2.65 (-3.791, 9.100)	0.4165	
Hedges' g SMD					0.15 (-0.216, 0.525)	0.4126	
NED/CR [IDS]	61 75.14 (23.692)	1.64 (2.064)	34 74.51 (28.790)	6.27 (2.941)	-4.64 (-11.785, 2.513)	0.2008	
Hedges' g SMD					-0.28 (-0.700, 0.143)	0.1947	
NED/CR [Chemo]	36 71.76 (29.498)	3.18 (3.008)	20 66.67 (33.333)	-0.03 (4.549)	3.21 (-7.770, 14.194)	0.5591	
Hedges' g SMD					0.17 (-0.380, 0.715)	0.5483	
PR	41 NC	NC	23 NC	NC	NC	NC	
Int. p-value						0.2178	
tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value	130 67.44 (28.139) 89 79.96 (22.916)	,	59 71.19 (29.333) 61 73.77 (26.433)	2.93 (2.466) 0.48 (2.010)	2.79 (-2.994, 8.575) 0.15 (-0.156, 0.460) -1.81 (-6.857, 3.227) -0.12 (-0.446, 0.206)	0.3424 0.3346 0.4777 0.4714 0.6045	
	ment outcome (eCRF)		40.55.50.400.450.	1 22 / 2 (25)	0.50 / 0.000 0.105	0.4050	
NED [PDS]	79 68.78 (28.665)	3.91 (1.909)	42 75.79 (22.452)	1.33 (2.685)	,	0.4363	
Hedges' g SMD	60 76 67 (04 006)	0.70 / 0.000	00 77 01 (00 000)	4 51 / 2 104)	0.15 (-0.225, 0.525)	0.4332	
NED/CR [IDS]	60 76.67 (24.006)	0.70 (2.068)	29 77.01 (29.009)	4.71 (3.194)	-4.02 (-11.592, 3.556)	0.2944	
Hedges' g SMD NED/CR [Chemo]	33 73.23 (29.149)	2.98 (3.041)	17 72.55 (30.585)	0.06 (5.188)	-0.24 (-0.688, 0.202) 2.92 (-9.228, 15.071)	0.2848	
Hedges' q SMD	33 /3.23 (29.149)	2.98 (3.041)	1/ /2.55 (30.585)	0.00 (5.188)	0.15 (-0.434, 0.738)	0.6295	
PR	44 74.24 (25.021)	2.81 (2.598)	31 65.05 (30.839)	-0.94 (3.359)	3.75 (-4.758, 12.263)	0.8102	
Hedges' g SMD	11 /1.21 (23.021)	2.01 (2.390)	31 03.03 (30.039)	0.74 (3.339)	0.21 (-0.253, 0.669)	0.3763	
Int. p-value					0.21 (0.255, 0.005)	0.4672	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.3 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Role (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be N=255			Placebo + bev		Difference between groups		
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Screening labor	atory tE	BRCA status (e	CRF)						
tBRCAm	127 66	.93 (28.249)	5.71 (1.604)	60	71.67 (29.321)	3.21 (2.451)	2.50 (-3.288, 8.284)	0.3954	
Hedges' g SMD							0.14 (-0.172, 0.443)	0.3879	
non-tBRCAm	92 80	.25 (22.636)	-1.26 (1.518)	60	73.33 (26.433)	0.45 (2.019)	-1.71 (-6.730, 3.311)	0.5018	
Hedges' g SMD							-0.11 (-0.439, 0.212)	0.4945	
Int. p-value								0.6724	
Age group									
<65 years	161 70	.39 (27.762)	3.33 (1.319)	90	71.30 (28.270)	2.63 (1.858)	0.70 (-3.792, 5.186)	0.7599	
Hedges' g SMD							0.04 (-0.217, 0.299)	0.7569	
>=65 years	58 78	.45 (23.155)	1.66 (2.135)	30	76.11 (26.509)	-0.85 (3.161)	2.52 (-5.069, 10.102)	0.5110	
Hedges' g SMD							0.15 (-0.291, 0.591)	0.5051	
Int. p-value								0.8915	
FIGO Stage (Dis	ease sta	ate)							
III	156 71	.26 (27.276)	2.02 (1.363)	83	74.90 (28.200)	3.44 (1.952)	-1.42 (-6.111, 3.275)	0.5522	
Hedges' g SMD							-0.08 (-0.348, 0.185)	0.5475	
IV	63 75	.66 (25.545)	5.08 (1.879)	37	67.12 (26.494)	-3.31 (2.824)	8.39 (1.615, 15.166)	0.0158	
Hedges' g SMD							0.53 (0.115, 0.940)	0.0123	
Int. p-value								0.0551	
Region									
Europe	210 72	.46 (26.929)	2.79 (1.158)	114	72.08 (28.265)	1.37 (1.669)	1.42 (-2.578, 5.416)	0.4854	
Hedges' g SMD							0.08 (-0.146, 0.311)	0.4782	
Japan	9	NC	NC	6	NC	NC	NC	NC	
Int. p-value								NC	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.3 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Role (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=25		Placebo + bev		Difference between groups		
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
ECOG performance	e status at Baseline						
(0) Normal	162 74.59 (25.143)	1.31 (1.269)	92 74.46 (26.778)	0.00 (1.745)	1.32 (-2.934, 5.567)	0.5423	
activity Hedges' g SMD					0.08 (-0.176, 0.336)	0.5390	
(1) Restricted	53 66.98 (30.573)	8.11 (2.530)	28 66.07 (30.592)	6.39 (3.854)	1.72 (-7.455, 10.899)	0.7098	
activity	33 00.70 (30.373)	0.11 (2.550)	20 00107 (3013)27	0.00 (0.001)	11.12 (0.7050	
Hedges' g SMD					0.09 (-0.369, 0.547)	0.7026	
Int. p-value						0.7172	
Baseline CA-125	value						
<=ULN	194 73.02 (26.927)	2.66 (1.163)	107 74.30 (26.430)	0.96 (1.657)	1.70 (-2.282, 5.689)	0.4009	
Hedges' g SMD					0.10 (-0.133, 0.339)	0.3939	
>ULN Int. p-value	25 NC	NC	13 NC	NC	NC	NC NC	
Histological gra	ade						
High grade	219 72.53 (26.806)	2.86 (1.121)	120 72.50 (27.809)	1.62 (1.598)	1.24 (-2.602, 5.077)	0.5264	
Hedges' g SMD					0.07 (-0.150, 0.296)	0.5205	
Int. p-value						ID	
Cytoreductive s	urgery outcome						
No residue	142 71.83 (26.939)	2.81 (1.372)	72 75.69 (25.475)	2.66 (2.001)	0.15 (-4.635, 4.935)	0.9509	
Hedges' g SMD					0.01 (-0.275, 0.293)	0.9503	
Residue	70 72.86 (26.945)	3.09 (2.044)	40 66.25 (31.687)	1.23 (2.930)	1.85 (-5.249, 8.953)	0.6056	
Hedges' g SMD					0.10 (-0.284, 0.493)	0.5988	
Int. p-value						0.9361	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.3 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Role (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	C)laparib + be (N=255			Placebo + bev			Difference between groups			
Subgroup		ean (SD) at aseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (S Change f baseli	rom	Estimated difference (99		p-value	
Timing of cytore	ductive	surgery									
Upfront Hedges' g SMD	129 70.	54 (28.376)	4.34 (1.464)	71 '	70.66 (27.526)	0.48 (2	.048)	3.87 (-1.100, 0.23 (-0.062,		0.1263 0.1231	
Interval Hedges' g SMD Int. p-value	83 74.	70 (24.329)	0.42 (1.770)	41 '	75.20 (29.136)	5.38 (2	.711)	-4.96 (-11.371, -0.30 (-0.674,	•	0.1283 0.1204 0.0094	
Myriad tumour BF	RCA mutat	ion status									
tBRCAm Hedges' g SMD	136 68.	75 (27.699)	4.57 (1.516)	68 '	72.30 (28.595)	2.55 (2	.257)	2.02 (-3.343, 0.11 (-0.179,	,	0.4583 0.4513	
Non-tBRCAm Hedges' g SMD Int. p-value	83 78.	71 (24.182)	-0.05 (1.640)	52 '	72.76 (27.022)	0.79 (2	.203)	-0.84 (-6.297, -0.05 (-0.401,		0.7600 0.7567 0.7174	
Status somatic E	BRCA muta	tions									
sBRCAm Hedges' g SMD	18 72.	22 (23.570)	7.67 (3.977)	6 '	72.22 (32.773)	2.93 (7	.435)	4.74 (-13.189, 0.27 (-0.661,	,	0.5821 0.5736	
gBRCAm Hedges' g SMD	57 70.	47 (27.280)	-0.38 (2.482)	28	67.86 (30.065)	5.86 (3	.773)	-6.24 (-15.227, -0.32 (-0.778,	2.745)	0.1708 0.1641	
Non-BRCAm Hedges' g SMD Int. p-value	36 82.	41 (21.434)	1.20 (2.223)	22 '	78.03 (18.819)	-1.13 (3	.122)	2.33 (-5.413, 0.17 (-0.365,	10.069)	0.5475 0.5405 0.2241	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.4 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Cognitive (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		bevacizumab 255)	Placebo + ber (N=13)		Difference between groups			
Subgroup	n Mean (SD) at	Mean (SE) Change from	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value		
First line treat	ment outcome (IVRS)						
NED [PDS]	81 83.33 (18.066) -2.20 (1.519)	43 82.56 (18.882)	-3.34 (2.179)	1.14 (-4.121, 6.394)	0.6694		
Hedges' g SMD					0.08 (-0.289, 0.451)	0.6664		
NED/CR [IDS]	61 77.05 (19.992) -0.85 (2.003)	33 83.84 (17.423)	-0.72 (2.904)	-0.13 (-7.187, 6.922)	0.9702		
Hedges' g SMD					-0.01 (-0.432, 0.415)	0.9696		
NED/CR [Chemo]	37 78.38 (25.112) -0.76 (1.838)	20 80.00 (21.357)	-4.42 (2.798)	3.66 (-3.125, 10.446)	0.2821		
Hedges' g SMD					0.31 (-0.237, 0.857)	0.2673		
PR	41 87.40 (19.643) -3.08 (2.134)	22 79.55 (20.530)	-1.01 (3.125)	-2.07 (-9.681, 5.547)	0.5888		
Hedges' g SMD					-0.15 (-0.664, 0.373)	0.5820		
Int. p-value						0.6725		
Screening labora tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value	tory tBRCA status 130 78.59 (21.789 90 85.74 (17.589			-4.59 (2.043) 0.52 (1.698)	3.28 (-1.522, 8.087) 0.22 (-0.092, 0.525) -3.46 (-7.674, 0.764) -0.28 (-0.605, 0.055)	0.1793 0.1688 0.1076 0.1019 0.1058		
First line treat	ment outcome (eCRF	')						
NED [PDS]	79 80.80 (22.343		42 83.33 (18.405)	-2.89 (2.232)	0.77 (-4.653, 6.189)	0.7795		
Hedges' g SMD	, , , , , ,			,	0.05 (-0.321, 0.428)	0.7780		
NED/CR [IDS]	60 80.00 (18.866) -0.35 (1.916)	28 84.52 (18.104)	-2.32 (3.044)	1.97 (-5.206, 9.155)	0.5854		
Hedges' g SMD	,	, , , ,		,	0.13 (-0.321, 0.577)	0.5754		
NED/CR [Chemo]	34 80.88 (20.569) -1.10 (1.838)	17 81.37 (21.955)	-4.74 (3.076)	3.63 (-3.662, 10.927)	0.3191		
Hedges' g SMD		,		. , ,	0.31 (-0.272, 0.900)	0.2935		
PR	44 85.61 (19.553) -3.82 (2.031)	30 78.89 (19.045)	-1.23 (2.595)	-2.59 (-9.188, 4.016)	0.4371		
Hedges' g SMD	,	, , , ,		,,	-0.19 (-0.651, 0.279)	0.4337		
Int. p-value					,,,	0.6482		

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.4 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Cognitive (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be			Placebo + bev (N=132		Difference between groups			
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95	% CI)	p-value	
Screening labor	atory tE	BRCA status (e	CRF)							
tBRCAm	127 78	.35 (21.954)	-1.23 (1.314)	60	85.28 (15.674)	-4.61 (2.024)	3.38 (-1.410,	8.176)	0.1654	
Hedges' g SMD							0.22 (-0.085,	0.531)	0.1551	
non-tBRCAm	93 85	.84 (17.367)	-2.99 (1.264)	58	78.45 (21.631)	0.62 (1.725)	-3.61 (-7.858,	0.639)	0.0953	
Hedges' g SMD							-0.29 (-0.616,	0.044)	0.0890	
Int. p-value									0.0862	
Age group										
<65 years	161 79	.61 (21.407)	-1.79 (1.147)	89	82.40 (18.181)	-3.24 (1.635)	1.45 (-2.490,	5.391)	0.4692	
Hedges' g SMD							0.10 (-0.162,	0.356)	0.4620	
>=65 years	59 86	.72 (16.603)	-2.25 (1.424)	29	80.46 (21.853)	0.74 (2.185)	-2.99 (-8.205,	2.222)	0.2562	
Hedges' g SMD							-0.26 (-0.711,	0.182)	0.2455	
Int. p-value									0.3850	
FIGO Stage (Dis	ease sta	ate)								
III	157 80	.36 (21.558)	-2.33 (1.121)	81	81.28 (20.135)	-0.33 (1.630)	-1.99 (-5.893,	1.905)	0.3146	
Hedges' g SMD							-0.14 (-0.408,		0.3087	
IV	63 84	.39 (17.163)	-1.08 (1.546)	37	83.33 (16.667)	-7.80 (2.339)	6.72 (1.149,		0.0186*	
Hedges' g SMD							0.51 (0.100,	0.925)	0.0149*	
Int. p-value									0.0575	
Region										
Europe	211 81	.44 (20.743)	-2.00 (0.964)	112	81.70 (19.240)	-2.47 (1.416)	0.48 (-2.896,	3.847)	0.7816	
Hedges' g SMD							0.03 (-0.196,	0.262)	0.7774	
Japan	9	NC	NC	6	NC	NC	NC		NC	
Int. p-value									NC	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.4 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Cognitive (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

				(N=132	2)	Difference between groups			
[a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value	
stati	us at Baseline								
163 8	82.62 (19.097)	-1.65 (1.099)	90	82.78 (19.101)	-3.53 (1.535)			0.3201	
54 5	78.09 (24.190)	-1.93 (1.719)	28	79.17 (19.043)	1.59 (2.787)			0.3157 0.2867	
						-0.26 (-0.719,	0.198)	0.2655 0.0452*	
alue		1 21 / 0 062)	105	01 E0 (10 464)	2 52 / 1 205\	1 21 / 2 120	4 E42)	0.4771	
195 (51.71 (20.696)	-1.31 (0.962)	105	01.59 (19.404)	-2.52 (1.395)			0.4771	
25	NC	NC	13	NC	NC	NC		NC NC	
le									
220 8	81.52 (20.440)	-1.87 (0.930)	118	81.92 (19.069)	-2.31 (1.348)			0.7903	
						0.03 (-0.193,	0.255)	0.7868 ID	
gery	outcome								
142 8	80.40 (20.657)	-1.51 (1.199)	71	83.33 (18.473)	-2.43 (1.769)	0.92 (-3.299,	5.139)	0.6676	
						, ,	,	0.6635	
71 8	82.39 (20.488)	-2.63 (1.380)	39	80.34 (17.882)	-2.75 (2.009)	, ,	,	0.9613	
						0.01 (-0.381,	0.401)	0.9605 0.5964	
	stat 163 : 54 : 54 : 54 : 54 : 54 : 54 : 54 : 5	status at Baseline 163 82.62 (19.097) 54 78.09 (24.190) salue 195 81.71 (20.696) 25 NC de 220 81.52 (20.440)	status at Baseline 163 82.62 (19.097) -1.65 (1.099) 54 78.09 (24.190) -1.93 (1.719) ralue 195 81.71 (20.696) -1.31 (0.962) 25 NC NC le 220 81.52 (20.440) -1.87 (0.930) regery outcome 142 80.40 (20.657) -1.51 (1.199)	status at Baseline 163 82.62 (19.097) -1.65 (1.099) 90 54 78.09 (24.190) -1.93 (1.719) 28 value 195 81.71 (20.696) -1.31 (0.962) 105 25 NC NC 13 de 220 81.52 (20.440) -1.87 (0.930) 118	status at Baseline 163 82.62 (19.097) -1.65 (1.099) 90 82.78 (19.101) 54 78.09 (24.190) -1.93 (1.719) 28 79.17 (19.043) Falue 195 81.71 (20.696) -1.31 (0.962) 105 81.59 (19.464) 25 NC NC 13 NC Rele 220 81.52 (20.440) -1.87 (0.930) 118 81.92 (19.069) Regery outcome 142 80.40 (20.657) -1.51 (1.199) 71 83.33 (18.473)	status at Baseline 163 82.62 (19.097) -1.65 (1.099) 90 82.78 (19.101) -3.53 (1.535) 54 78.09 (24.190) -1.93 (1.719) 28 79.17 (19.043) 1.59 (2.787) **alue 195 81.71 (20.696) -1.31 (0.962) 105 81.59 (19.464) -2.52 (1.395) 25 NC NC 13 NC NC **le 220 81.52 (20.440) -1.87 (0.930) 118 81.92 (19.069) -2.31 (1.348) **gery outcome 142 80.40 (20.657) -1.51 (1.199) 71 83.33 (18.473) -2.43 (1.769)	status at Baseline 163 82.62 (19.097) -1.65 (1.099) 90 82.78 (19.101) -3.53 (1.535) 1.88 (-1.838, 54 78.09 (24.190) -1.93 (1.719) 28 79.17 (19.043) 1.59 (2.787) -3.52 (-10.054, -0.26 (-0.719, **alue 195 81.71 (20.696) -1.31 (0.962) 105 81.59 (19.464) -2.52 (1.395) 1.21 (-2.130, 25 NC NC 13 NC NC NC NC **le 220 81.52 (20.440) -1.87 (0.930) 118 81.92 (19.069) -2.31 (1.348) 0.44 (-2.787, 0.03 (-0.193, **gery outcome 142 80.40 (20.657) -1.51 (1.199) 71 83.33 (18.473) -2.43 (1.769) 0.92 (-3.299, 0.06 (-0.222, 71 82.39 (20.488) -2.63 (1.380) 39 80.34 (17.882) -2.75 (2.009) 0.12 (-4.718,	status at Baseline 163 82.62 (19.097) -1.65 (1.099) 90 82.78 (19.101) -3.53 (1.535) 1.88 (-1.838, 5.601) 54 78.09 (24.190) -1.93 (1.719) 28 79.17 (19.043) 1.59 (2.787) -3.52 (-10.054, 3.020) -0.26 (-0.719, 0.198) **alue 195 81.71 (20.696) -1.31 (0.962) 105 81.59 (19.464) -2.52 (1.395) 1.21 (-2.130, 4.543) 0.09 (-0.150, 0.325) **DRC NC 13 NC NC NC NC NC **Recompleted by the state of the state	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.4 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Cognitive (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + bev		Difference k	oetween g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
Timing of cytor		9 1	1 20 / 1 145)	70	01 10 (10 010)	2 10 / 1 625)	1 07 / 2 065	F 010\	0 2402
Upfront	130	81.67 (21.250)	-1.32 (1.145)	70	81.19 (19.018)	-3.19 (1.635)	1.87 (-2.065,	5.812)	0.3493
Hedges' g SMD Interval	0.2	00 10 /10 FF0\	-2.25 (1.654)	4.0	04 17 (16 050)	0 10 / 0 570)	0.14 (-0.150,	0.431) 5.942)	0.3435 0.9667
Hedges' q SMD	83	80.12 (19.558)	-2.25 (1.054)	40	84.17 (16.858)	-2.12 (2.572)	-0.13 (-6.198, -0.01 (-0.385,	0.369)	0.9659
Int. p-value							-0.01 (-0.383,	0.309)	0.4849
Myriad tumour B	RCA mu	tation status							
tBRCAm	136	79.04 (21.372)	-1.89 (1.227)	68	85.05 (16.577)	-4.29 (1.846)	2.39 (-2.006,	6.795)	0.2845
Hedges' g SMD							0.16 (-0.128,	0.455)	0.2725
Non-tBRCAm	84	85.52 (18.254)	-1.86 (1.403)	50	77.67 (21.458)	0.66 (1.945)	-2.52 (-7.297,	2.254)	0.2983
Hedges' g SMD							-0.19 (-0.541,	0.161)	0.2886
Int. p-value									0.2114
Status somatic	BRCA m	utations							
sBRCAm	18	NC	NC	6	NC	NC	NC		NC
gBRCAm	56	74.11 (21.770)	-1.50 (2.046)	28	82.74 (16.026)	-2.67 (3.093)	1.17 (-6.266,	8.609)	0.7544
Hedges' g SMD							0.07 (-0.380,	0.528)	0.7489
Non-BRCAm	37	83.33 (18.002)	1.60 (1.670)	21	76.19 (21.455)	-0.71 (2.528)	2.31 (-3.800,	8.412)	0.4526
Hedges' g SMD							0.21 (-0.324,	0.750)	0.4374
Int. p-value									0.9429

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.5 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Emotional (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + b (N=25		Placebo + be (N=13		Difference between groups			
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value		
First line treat	ment outcome (IVRS)							
NED [PDS]	81 76.85 (22.973)	-0.82 (1.411)	43 70.54 (20.521)	-1.29 (2.033)	0.47 (-4.450, 5.382)	0.8514		
Hedges' g SMD					0.04 (-0.334, 0.406)	0.8493		
NED/CR [IDS]	61 75.09 (21.636)	-4.67 (2.181)	33 85.94 (13.992)	-4.34 (3.194)	-0.33 (-8.171, 7.504)	0.9328		
Hedges' g SMD					-0.02 (-0.442, 0.405)	0.9303		
NED/CR [Chemo]	37 69.67 (26.895)	1.60 (2.360)	20 80.83 (14.833)	1.42 (3.529)	0.18 (-8.519, 8.878)	0.9669		
Hedges' g SMD					0.01 (-0.532, 0.556)	0.9658		
PR	41 79.95 (17.163)	-3.51 (2.138)	22 79.29 (14.516)	-5.39 (3.216)	1.88 (-5.869, 9.622)	0.6294		
Hedges' g SMD					0.13 (-0.388, 0.649)	0.6215		
Int. p-value						0.2656		
Screening labora	itory tBRCA status (IVRS)						
tBRCAm	130 72.78 (24.509)	-1.44 (1.328)	59 79.10 (17.219)	-5.79 (2.088)	4.35 (-0.553, 9.247)	0.0817		
Hedges' g SMD					0.28 (-0.028, 0.590)	0.0750		
non-tBRCAm	90 80.00 (18.384)	-2.51 (1.313)	59 77.35 (18.540)	0.56 (1.736)	-3.08 (-7.381, 1.228)	0.1598		
Hedges' g SMD					-0.24 (-0.568, 0.090)	0.1551		
Int. p-value						0.1786		
First line treat	ment outcome (eCRF)							
NED [PDS]	79 74.26 (23.952)	-0.60 (1.465)	42 71.03 (20.515)	-1.00 (2.079)	0.40 (-4.642, 5.443)	0.8752		
Hedges' g SMD	, , , ,	, , , ,	, ,		0.03 (-0.344, 0.404)	0.8744		
NED/CR [IDS]	60 77.73 (19.798)	-4.15 (2.124)	28 86.31 (14.910)	-4.53 (3.370)	0.39 (-7.642, 8.419)	0.9236		
Hedges' g SMD	, ,	, ,	, , ,	/	0.02 (-0.426, 0.471)	0.9207		
NED/CR [Chemo]	34 72.63 (25.924)	1.56 (2.482)	17 82.84 (13.330)	-1.18 (4.143)	2.74 (-7.467, 12.945)	0.5811		
Hedges' g SMD	,	, - ,	,,	/	0.18 (-0.408, 0.759)	0.5555		
PR	44 77.34 (20.752)	-4.16 (2.059)	30 78.52 (15.108)	-2.27 (2.747)		0.5851		
Hedges' g SMD	, , ,	, , , ,	,,	, ,	-0.13 (-0.596, 0.333)	0.5795		
Int. p-value					, , , , , , , , , , , , , , , , , , , ,	0.2062		

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.5 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Emotional (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	(Olaparib + be (N=255			Placebo + bev (N=132		Difference between groups			
Subgroup		ean (SD) at aseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value	
Screening labor	atory tBI	RCA status (e	CRF)							
tBRCAm	127 72.	.59 (24.762)	-1.71 (1.343)	60	79.44 (17.285)	-5.77 (2.071)	4.06 (-0.833,	8.951)	0.1034	
Hedges' g SMD							0.26 (-0.046,	0.570)	0.0957	
non-tBRCAm	93 80.	.02 (18.096)	-2.15 (1.292)	58	76.96 (18.455)	0.74 (1.751)	-2.89 (-7.195,	1.415)	0.1866	
Hedges' g SMD							-0.22 (-0.554,	0.104)	0.1808	
Int. p-value									0.2417	
Age group										
<65 years	161 74.	.86 (23.302)	-3.16 (1.171)	89	77.15 (17.316)	-3.06 (1.673)	-0.09 (-4.120,	3.930)	0.9631	
Hedges' g SMD							-0.01 (-0.265,	0.253)	0.9625	
>=65 years	59 78.	.11 (19.918)	1.23 (1.509)	29	81.51 (19.292)	-1.64 (2.321)	2.87 (-2.675,	8.412)	0.3064	
Hedges' g SMD							0.24 (-0.207,	0.685)	0.2934	
Int. p-value									0.8160	
FIGO Stage (Dis	ease stat	te)								
III	157 75.	.05 (23.969)	-2.29 (1.138)	81	77.50 (19.482)	-1.43 (1.662)	-0.86 (-4.831,	3.117)	0.6711	
Hedges' g SMD							-0.06 (-0.327,	0.209)	0.6667	
IV	63 77.	.43 (18.171)	-1.07 (1.853)	37	79.80 (13.666)	-6.91 (2.681)	5.84 (-0.649,		0.0771	
Hedges' g SMD							0.38 (-0.031,	0.788)	0.0700	
Int. p-value									0.2033	
Region										
Europe	211 75.	.21 (22.644)	-2.07 (0.990)	112	77.88 (18.129)	-3.07 (1.458)	1.01 (-2.464,	4.478)	0.5685	
Hedges' g SMD							0.07 (-0.161,	0.297)	0.5602	
Japan	9	NC	NC	6	NC	NC	NC		NC	
Int. p-value									NC	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.5 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Emotional (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + bev (N=132		Difference l	oetween g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at	Mean (SE) Change from baseline	Estimated difference (95		p-value
ECOG performance	e stat	us at Baseline							
(0) Normal activity	163	77.45 (20.719)	-2.44 (1.119)	90	77.90 (17.759)	-3.09 (1.562)	0.65 (-3.134,	4.438)	0.7347
Hedges' g SMD							0.04 (-0.213,	0.302)	0.7324
(1) Restricted activity	54	69.50 (26.465)	0.47 (1.956)	28	79.27 (18.371)	-1.05 (3.212)	1.53 (-6.031,	9.090)	0.6886
Hedges' g SMD Int. p-value							0.10 (-0.358,	0.556)	0.6711 0.7726
Baseline CA-125									
<=ULN	195	74.93 (23.293)	-1.65 (0.993)	105	78.62 (18.048)	-2.52 (1.445)	0.87 (-2.583,	,	0.6190
Hedges' g SMD >ULN Int. p-value	25	NC	NC	13	NC	NC	0.06 (-0.176, NC	0.299)	0.6119 NC NC
Histological gra	ade								
High grade		75.73 (22.447)	-1.98 (0.961)	118	78.23 (17.836)	-2.60 (1.396)	0.62 (-2.717,	3.961)	0.7141
Hedges' g SMD Int. p-value							0.04 (-0.181,	0.266)	0.7091 ID
Cytoreductive s	urgery	z outcome							
No residue		75.98 (22.154)	-2.23 (1.208)	71	77.00 (19.744)	-2.13 (1.788)	-0.10 (-4.359,	4.151)	0.9617
Hedges' g SMD							-0.01 (-0.292,	0.278)	0.9612
Residue	71	73.94 (23.567)	-0.66 (1.755)	39	79.49 (13.422)	-3.51 (2.564)	2.85 (-3.358,	9.065)	0.3640
Hedges' g SMD							0.19 (-0.205,	0.578)	0.3515
Int. p-value								,	0.377

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.5 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Emotional (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + be		nab	Differenc	e between g	groups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Cha	an (SE) nge from seline	Estima difference		p-value
Timing of cytore			0.10 (1.165)		TO TO (10 041)			1 01 / 0 00	5 005	0 0401
Upfront	T30	74.62 (23.726)	0.19 (1.167)	70	73.53 (18.241)	-1.7.	1 (1.657)			0.3481
Hedges' g SMD	00.1	76 25 (00 005)	4 50 / 1 505)	4.0	05 40 (14 045)	4 2		0.14 (-0.15		0.3433
Interval	83	76.37 (20.807)	-4.59 (1.787)	40	85.49 (14.045)	-4.3	3 (2.824)	•		0.9506
Hedges' g SMD Int. p-value								-0.01 (-0.39	0, 0.365)	0.9484
ine. p varae										0.0717
Myriad tumour BF	RCA mu	tation status								
tBRCAm	136 '	73.92 (23.437)	-1.87 (1.290)	68	79.53 (17.089)	-4.3	4 (1.938)	2.48 (-2.13	2, 7.088)	0.2904
Hedges' g SMD								0.16 (-0.13	1, 0.452)	0.2798
Non-tBRCAm	84	78.67 (20.540)	-1.91 (1.384)	50	76.44 (18.833)	-0.43	1 (1.919)	-1.50 (-6.18	5, 3.187)	0.5277
Hedges' g SMD								-0.11 (-0.46	5, 0.236)	0.5219
Int. p-value										0.7033
Status somatic E	RCA m	utations								
sBRCAm		77.78 (21.390)	-6.15 (3.585)	6	87.50 (10.206)	-13.06	5 (6.359)	6.91 (-8.46	0. 22.281)	0.3598
Hedges' g SMD		(22,370)	1.15 (3.505)	3	220 (20.200)			0.44 (-0.49		0.3598
gBRCAm	56 '	72.52 (25.529)	-4.97 (2.016)	28	77.68 (19.311)	-3.63	1 (3.098)	-1.36 (-8.73		0.7151
Hedges' g SMD		, ,	,,		- , /		,,	-0.09 (-0.54		0.7083
Non-BRCAm	37 '	77.03 (23.357)	-0.60 (2.259)	21	78.57 (21.176)	-0.05	5 (3.243)	-0.56 (-8.52		0.8888
Hedges' g SMD		(======================================	., (,					-0.04 (-0.57		0.8872
Int. p-value								(3.37	,,	0.5545

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.6 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Social (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=255		Placebo + be (N=13		Difference between groups			
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value		
First line treat	ment outcome (IVRS)							
NED [PDS]	81 75.31 (25.428)	6.89 (1.618)	43 70.54 (24.623)	7.14 (2.350)	-0.24 (-5.897, 5.413)	0.9326		
Hedges' g SMD					-0.02 (-0.386, 0.354)	0.9316		
NED/CR [IDS]	61 74.04 (28.791)	1.49 (2.250)	33 78.79 (23.670)	7.04 (3.231)	-5.55 (-13.394, 2.300)	0.1632		
Hedges' g SMD					-0.31 (-0.733, 0.119)	0.1578		
NED/CR [Chemo]	37 65.32 (30.525)	6.77 (2.639)	20 70.83 (28.549)	10.06 (3.974)	-3.29 (-12.870, 6.287)	0.4938		
Hedges' g SMD					-0.19 (-0.740, 0.350)	0.4839		
PR	41 76.42 (29.809)	4.40 (2.479)	22 74.24 (25.054)	4.60 (3.700)	-0.20 (-9.127, 8.737)	0.9652		
Hedges' g SMD					-0.01 (-0.530, 0.506)	0.9645		
Int. p-value						0.3273		
Screening labora	itory tBRCA status (I	VRS)						
tBRCAm	130 69.23 (29.162)	5.81 (1.418)	59 72.88 (26.785)	8.62 (2.227)	-2.81 (-8.024, 2.402)	0.2887		
Hedges' g SMD					-0.17 (-0.478, 0.138)	0.2793		
non-tBRCAm	90 79.63 (25.574)	3.44 (1.565)	59 74.29 (23.432)	6.28 (2.055)	-2.85 (-7.969, 2.273)	0.2732		
Hedges' g SMD					-0.19 (-0.515, 0.143)	0.2675		
Int. p-value						0.4299		
First line treat	ment outcome (eCRF)							
NED [PDS]	79 74.26 (27.965)	6.15 (1.640)	42 73.02 (23.557)	7.56 (2.340)	-1.42 (-7.078, 4.242)	0.6206		
Hedges' g SMD		.,== (=.310)		(-0.10 (-0.470, 0.279)	0.6178		
NED/CR [IDS]	60 75.83 (29.976)	1.80 (2.124)	28 80.36 (24.867)	8.30 (3.378)		0.1072		
Hedges' g SMD	(=====,0,0,	, , , , , , , , , , , , , , , , , , , ,	(=/		-0.38 (-0.834, 0.071)	0.0984		
NED/CR [Chemo]	34 65.20 (28.239)	10.08 (2.860)	17 71.57 (28.726)	10.75 (4.733)	-0.67 (-11.798, 10.466)	0.9047		
Hedges' g SMD	(20.20)		_: /1.5/ (20./20)		-0.04 (-0.619, 0.545)	0.9005		
PR	44 76.52 (25.749)	3.38 (2.498)	30 70.00 (25.295)	4.40 (3.256)	-1.01 (-9.224, 7.195)	0.8057		
Hedges' g SMD	(===, 12)	-, (1,0)	(==:255)		-0.06 (-0.523, 0.405)	0.8040		
Int. p-value					3.100,	0.5350		

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

SMD = standardised mean difference. * p<0.05.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.6 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Social (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Ola	aparib + bev N=255)			Placebo + bev (N=132		Difference between groups		
Subgroup		n (SD) at eline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
Screening labor	atory tBRC	A status (e0	CRF)						
tBRCAm	127 68.50	(29.113)	5.89 (1.442)	60	73.33 (26.786)	8.94 (2.221)	-3.05 (-8.288,	2.178)	0.2509
Hedges' g SMD							-0.18 (-0.491,	0.124)	0.2418
non-tBRCAm	93 80.29	9 (25.413)	3.37 (1.527)	58	73.85 (23.386)	6.08 (2.063)	-2.71 (-7.801,	2.383)	0.2946
Hedges' g SMD							-0.18 (-0.507,	0.150)	0.2874
Int. p-value									0.4036
Age group									
<65 years	161 71.84	1 (29.002)	5.18 (1.274)	89	71.35 (25.375)	9.29 (1.815)	-4.11 (-8.478,	0.257)	0.0649
Hedges' g SMD							-0.25 (-0.508,	0.012)	0.0611
>=65 years	59 77.97	7 (25.416)	3.07 (1.768)	29	80.46 (23.175)	2.86 (2.723)	0.21 (-6.261,	6.688)	0.9478
Hedges' g SMD							0.02 (-0.429,	0.460)	0.9467
Int. p-value									0.3810
FIGO Stage (Dis	ease state)							
III	157 74.10	(28.143)	3.35 (1.243)	81	73.46 (26.716)	8.97 (1.818)	-5.62 (-9.957,	-1.279)	0.0114*
Hedges' g SMD							-0.35 (-0.623,		0.0103*
IV	63 71.96	5 (28.371)	9.26 (1.864)	37	73.87 (21.351)	3.96 (2.796)	5.30 (-1.388,		0.1189
Hedges' g SMD							0.34 (-0.073,	0.745)	0.1070
Int. p-value									0.0265*
Region									
Europe	211 72.99	9 (28.491)	4.94 (1.075)	112	73.66 (25.275)	6.98 (1.582)	-2.04 (-5.809,	1.720)	0.2861
Hedges' g SMD							-0.13 (-0.357,	0.102)	0.2765
Japan	9	NC	NC	6	NC	NC	NC		NC
Int. p-value									NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.6 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Social (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bev			Placebo + bev N=132		Difference between groups			
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value	
ECOG performance	e statı	us at Baseline								
(0) Normal activity	163 7	73.72 (27.273)	4.79 (1.172)	90	75.37 (24.381)	6.56 (1.643)	-1.78 (-5.754,	2.200)	0.3795	
Hedges' g SMD							-0.12 (-0.374,	0.141)	0.3747	
(1) Restricted activity	54 7	73.77 (31.323)	4.56 (2.367)	28	67.86 (26.808)	9.95 (3.694)	-5.39 (-14.134,	3.352)	0.2234	
Hedges' g SMD Int. p-value							-0.29 (-0.753,	0.165)	0.2088 0.1783	
Baseline CA-125										
<=ULN	195 7	74.02 (28.670)	5.35 (1.078)	105	74.13 (25.422)	8.18 (1.565)	, ,		0.1375	
Hedges' g SMD >ULN Int. p-value	25	NC	NC	13	NC	NC	-0.18 (-0.421, NC	0.054)	0.1307 NC NC	
Histological gra	ade									
High grade	220 7	3.48 (28.160)	4.82 (1.055)	118	73.59 (25.067)	7.60 (1.529)	-2.78 (-6.433,	0.877)	0.1358	
Hedges' g SMD Int. p-value							-0.17 (-0.397,	0.051)	0.1293 ID	
Cytoreductive su	urgery	outcome								
No residue		74.53 (28.675)	4.39 (1.293)	71	75.59 (24.207)	7.85 (1.921)	-3.46 (-8.029,	1.103)	0.1363	
Hedges' g SMD							-0.22 (-0.506,	0.065)	0.1307	
Residue	71 6	59.72 (27.647)	6.37 (1.977)	39	68.80 (27.620)	7.00 (2.868)	-0.63 (-7.553,		0.8561	
Hedges' g SMD							-0.04 (-0.427,	0.354)	0.8535	
Int. p-value									0.6062	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.6 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Functional scale: Social (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + ber		Difference between groups			
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (S Change f baseli	Erom	Estimated difference (99		p-value
Timing of cytore	oduati	no dirgori								
Upfront Hedges' g SMD		72.05 (28.621)	7.33 (1.298)	70	70.48 (25.248)	8.08 (1		-0.75 (-5.215, -0.05 (-0.340,		0.7421 0.7391
Interval Hedges' g SMD Int. p-value	83	74.30 (28.070)	1.23 (1.931)	40	77.92 (25.706)	5.75 (2		-4.52 (-11.562, -0.25 (-0.628,		0.2059 0.1963 0.2143
Myriad tumour B	RCA mu	tation status								
tBRCAm Hedges' g SMD		68.87 (29.593)	6.32 (1.395)	68	73.53 (26.113)	9.76 (2		-3.44 (-8.405, -0.21 (-0.498,	,	0.1734 0.1656
Non-tBRCAm Hedges' g SMD Int. p-value	84	80.95 (24.012)	2.17 (1.598)	50	73.67 (23.831)	4.88 (2		-2.71 (-8.144, -0.18 (-0.530,		0.3247 0.3161 0.4667
Status somatic E	RCA m	utations								
sBRCAm Hedges' g SMD		65.74 (21.747)	14.05 (2.879)	6	72.22 (25.092)	15.82 (5	,	-1.77 (-14.288, -0.14 (-1.063,	,	0.7716 0.7696
gBRCAm Hedges' g SMD	56 '	75.00 (27.707)	-0.90 (2.315)	28	71.43 (23.941)	11.10 (3	3.519) -	-12.00 (-20.401, -0.67 (-1.135,	-3.607)	0.0057
Non-BRCAm Hedges' g SMD Int. p-value	37	85.14 (19.557)	0.89 (1.894)	21	80.95 (21.269)	1.91 (2		-1.01 (-7.863, -0.08 (-0.619,	•	0.7677 0.7616 0.1285

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.7 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Loss of appetite (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bev (N=255			Placebo + bev (N=132		Difference between groups		
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
First line treat		, ,							
NED [PDS]	81	10.29 (19.466)	3.24 (1.553)	43	8.53 (19.372)	-0.36 (2.225)	3.59 (-1.785, 8.971)	0.1884	
Hedges' g SMD							0.25 (-0.120, 0.623)	0.1840	
NED/CR [IDS]	61	6.01 (12.922)	6.09 (2.011)	34	7.84 (14.352)	1.62 (2.795)	4.46 (-2.388, 11.317)	0.1986	
Hedges' g SMD							0.28 (-0.143, 0.700)	0.1957	
NED/CR [Chemo]	36	9.26 (20.488)	2.95 (2.278)	20	5.00 (12.212)	-2.89 (3.354)	5.84 (-2.316, 13.998)	0.1563	
Hedges' g SMD							0.41 (-0.145, 0.959)	0.1486	
PR	41	5.69 (14.724)	5.00 (2.658)	23	14.49 (22.079)	7.19 (4.108)	-2.19 (-12.181, 7.801)	0.6587	
Hedges' g SMD							-0.12 (-0.631, 0.391)	0.6451	
Int. p-value								0.2388	
Screening labora	atory	tBRCA status (IV	RS)						
tBRCAm	130	8.46 (17.269)	4.23 (1.351)	59	6.21 (13.093)	2.36 (2.111)	1.88 (-3.074, 6.826)	0.4554	
Hedges' g SMD							0.12 (-0.189, 0.427)	0.4475	
non-tBRCAm	89	7.49 (17.228)	4.04 (1.284)	61	11.48 (20.981)	-0.87 (1.646)	4.91 (0.785, 9.042)	0.0200	
Hedges' g SMD							0.39 (0.065, 0.723)	0.0190	
Int. p-value								0.6099	
First line treat	ment	outcome (eCRF)							
NED [PDS]		9.70 (19.357)	3.41 (1.573)	42	7.94 (19.211)	0.33 (2.226)	3.09 (-2.319, 8.494)	0.2602	
Hedges' g SMD		,			•	,	0.22 (-0.158, 0.592)	0.2571	
NED/CR [IDS]	60	5.56 (12.527)	6.84 (1.870)	29	5.75 (12.814)	1.12 (2.782)	5.72 (-0.951, 12.391)	0.0918	
Hedges' g SMD		,			•	,	0.39 (-0.060, 0.834)	0.0896	
NED/CR [Chemo]	33	9.09 (19.135)	1.58 (2.331)	17	7.84 (14.575)	-4.73 (3.764)	6.31 (-2.674, 15.293)	0.1628	
Hedges' g SMD		,	, - ,			, , ,	0.44 (-0.153, 1.031)	0.1458	
PR	44	8.33 (17.791)	5.09 (2.519)	31	12.90 (20.507)	5.81 (3.203)	-0.72 (-8.903, 7.458)	0.8606	
Hedges' g SMD							-0.04 (-0.501, 0.418)	0.8591	
Int. p-value							,	0.2541	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.7 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Loss of appetite (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bev			Placebo + be		Difference between groups		
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95	∛ CI)	p-value
Screening labora	atory	tBRCA status (eC	RF)						
tBRCAm	127	8.66 (17.423)	4.31 (1.371)	60	6.11 (13.007)	2.16 (2.103)	2.15 (-2.810,	7.109)	0.3935
Hedges' g SMD							0.14 (-0.171,	0.444)	0.3853
non-tBRCAm	92	7.25 (16.994)	3.92 (1.254)	60	11.67 (21.104)	-0.74 (1.650)	4.66 (0.558,	8.757)	0.0263
Hedges' g SMD							0.38 (0.048,	0.704)	0.0247
Int. p-value									0.7130
Age group									
<65 years	161	6.83 (14.962)	5.51 (1.109)	90	5.93 (13.755)	1.91 (1.565)	3.60 (-0.179,	7.381)	0.0617
Hedges' g SMD							0.25 (-0.009,	0.509)	0.0583
>=65 years	58	11.49 (22.121)	0.18 (2.058)	30	17.78 (24.343)	-2.05 (3.039)	2.22 (-5.103,	9.548)	0.5473
Hedges' g SMD							0.14 (-0.304,	0.579)	0.5411
Int. p-value									0.8547
FIGO Stage (Dis	ease s	state)							
III	156	9.40 (18.854)	4.70 (1.192)	83	8.43 (18.656)	-0.26 (1.700)	4.95 (0.861,	9.047)	0.0179
Hedges' g SMD							0.33 (0.059,	0.595)	0.0168
IV	63	4.76 (11.758)	2.86 (1.445)	37	9.91 (15.446)	3.68 (2.125)	-0.82 (-5.943,	4.308)	0.7519
Hedges' g SMD							-0.07 (-0.474,	0.339)	0.7447
Int. p-value									0.0416
Region									
Europe	210	7.94 (17.269)	4.23 (0.987)	114	9.36 (18.018)	0.81 (1.428)	3.42 (0.004,	6.839)	0.0497
Hedges' g SMD							0.23 (0.004,	0.462)	0.0458
Japan	9	NC	NC	6	NC	NC	NC		NC
Int. p-value									NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.7 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Loss of appetite (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

Subgroup		Olaparib + bev (N=255			Placebo + bev (N=132		Difference between groups		
	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
ECOG performance	e stat	us at Baseline							
(0) Normal	162	5.97 (14.816)	4.49 (1.005)	92	7.61 (17.888)	1.58 (1.392)	2.91 (-0.471, 6.297)	0.0913	
activity Hedges' g SMD							0.22 (-0.033, 0.480)	0.0879	
(1) Restricted	53	14.47 (22.178)	1.82 (2.270)	28	13.10 (16.578)	-1.48 (3.425)	3.30 (-4.878, 11.484)	0.4241	
activity									
Hedges' g SMD							0.19 (-0.267, 0.650)	0.4136	
Int. p-value								0.9856	
Baseline CA-125	value	2							
<=ULN	194	8.08 (17.238)	3.87 (0.999)	107	8.72 (17.337)	1.59 (1.422)	2.28 (-1.142, 5.698)	0.1910	
Hedges' g SMD							0.16 (-0.076, 0.396)	0.1846	
>ULN Int. p-value	25	NC	NC	13	NC	NC	NC	NC NC	
Histological gra	ade								
High grade		8.07 (17.219)	4.15 (0.958)	120	8.89 (17.677)	0.83 (1.370)	3.31 (0.026, 6.603)	0.0483	
Hedges' g SMD Int. p-value							0.23 (0.005, 0.452)	0.0450° ID	
Cytoreductive su	ıraerv	z outcome							
No residue		7.75 (16.686)	4.64 (1.168)	72	7.41 (16.990)	0.52 (1.699)	4.12 (0.057, 8.190)	0.0469	
Hedges' g SMD							0.29 (0.007, 0.577)	0.0448	
Residue	70	8.57 (18.551)	2.61 (1.627)	40	10.00 (17.213)	-1.82 (2.314)	4.43 (-1.183, 10.042)	0.1206	
Hedges' g SMD							0.31 (-0.076, 0.705)	0.1148	
Int. p-value								0.8556	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.7 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Loss of appetite (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bev			Placebo + be		Difference between groups		
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Timing of cytore	educti	ve surgery							
Upfront Hedges' g SMD Interval Hedges' g SMD Int. p-value	129	9.56 (19.182) 5.62 (13.594)	2.00 (1.139) 7.07 (1.667)		8.92 (18.651) 7.32 (13.969)	-0.90 (1.607) 0.62 (2.512)	2.90 (-0.981, 6.790) 0.22 (-0.070, 0.510) 6.45 (0.477, 12.429) 0.41 (0.036, 0.792)	0.1420 0.1376 0.0346* 0.0317* 0.2435	
Myriad tumour Bl	RCA mu	Itation status							
tBRCAm Hedges' g SMD		8.33 (17.093)	4.45 (1.307)	68	6.86 (14.749)	1.49 (1.949)	2.96 (-1.674, 7.591) 0.19 (-0.102, 0.482)	0.2093 0.2020	
Non-tBRCAm Hedges' g SMD Int. p-value	83	7.63 (17.519)	3.89 (1.357)	52	11.54 (20.753)	-0.34 (1.821)	4.23 (-0.275, 8.726) 0.33 (-0.017, 0.681)	0.0655 0.0624 0.9501	
Status somatic 1	BRCA m	nutations							
sBRCAm Hedges' g SMD		12.96 (23.260)	1.00 (3.286)	6	5.56 (13.608)	-4.03 (5.985)	5.03 (-9.343, 19.398) 0.34 (-0.586, 1.274)	0.4724	
gBRCAm Hedges' g SMD	57	8.77 (16.093)	8.53 (2.335)	28	7.14 (13.929)	1.09 (3.534)	7.43 (-1.009, 15.875) 0.41 (-0.047, 0.866)	0.0835	
Non-BRCAm Hedges' g SMD Int. p-value	36	4.63 (11.691)	5.90 (1.637)	22	9.09 (18.349)	3.02 (2.191)	2.88 (-2.590, 8.348) 0.28 (-0.249, 0.817)	0.2956 0.2961 0.6887	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.8 PAOLAl: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale:

Constipation (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olap	parib + bev (N=255			Placebo + bev		Difference between o	Difference between groups		
Subgroup		(SD) at line [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value		
First line treat	ment outcor	me (IVRS)								
NED [PDS]	80 21.25	(32.367)	0.07 (1.843)	43 1	17.05 (30.318)	0.83 (2.656)	-0.76 (-7.171, 5.654)	0.8152		
Hedges' g SMD							-0.04 (-0.415, 0.326)	0.8127		
NED/CR [IDS]	61 18.03	(24.776)	0.03 (1.732)	34 2	10.78 (17.829)	0.52 (2.538)	-0.49 (-6.653, 5.672)	0.8743		
Hedges' g SMD							-0.03 (-0.454, 0.385)	0.8710		
NED/CR [Chemo]	36 14.81	(25.751)	5.28 (3.409)	20 1	10.00 (19.041)	6.15 (5.398)		0.8929		
Hedges' g SMD							-0.04 (-0.586, 0.507)	0.8877		
PR	39 N	1C	NC	21	NC	NC	NC	NC		
Int. p-value								0.9412		
Screening labora	-	•	,		4 60 (00 505)	1.50 / 0.000		0 5505		
tBRCAm	128 17.97	(28.648)	0.78 (1.497)	5./ [14.62 (23.585)	1.59 (2.383)		0.7735		
Hedges' g SMD	00 16 68	(06 842)	0 10 / 1 612)	<i>c</i> 1 -	2 66 (05 260)	1 15 / 0 150)	-0.05 (-0.359, 0.265)	0.7687		
non-tBRCAm	88 16.67	(26.743)	2.18 (1.613)	6I -	13.66 (25.369)	1.15 (2.158)		0.7025		
Hedges' g SMD Int. p-value							0.06 (-0.262, 0.392)	0.6969 0.9630		
int. p-value								0.9630		
First line treat	ment outcor	me (eCRF)								
NED [PDS]	78 23.08	(33.681)	-0.37 (1.900)	42 1	17.46 (30.567)	0.39 (2.695)	-0.76 (-7.308, 5.791)	0.8189		
Hedges' g SMD							-0.04 (-0.419, 0.331)	0.8171		
NED/CR [IDS]	60 16.11	(24.156)	0.80 (1.683)	29	10.34 (18.046)	3.02 (2.702)	-2.22 (-8.589, 4.158)	0.4903		
Hedges' g SMD							-0.16 (-0.606, 0.282)	0.4740		
NED/CR [Chemo]	33 11.11	(19.837)	6.12 (3.623)	17	9.80 (19.596)	10.71 (6.238)	-4.59 (-19.191, 10.016)	0.5289		
Hedges' g SMD							-0.20 (-0.787, 0.386)	0.5037		
PR	42 15.08	(25.717)	3.77 (2.508)	29	16.09 (22.923)	-3.55 (3.389)		0.0887		
Hedges' g SMD							0.42 (-0.055, 0.902)	0.0828		
Int. p-value								0.3119		

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.8 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale:

Constipation (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be			Placebo + bev		Difference between groups			
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value	
Screening labor	atory tE	BRCA status (e	CRF)							
tBRCAm	125 18	.13 (28.869)	0.81 (1.522)	58	14.94 (23.506)	1.57 (2.374)	-0.76 (-6.327,	4.815)	0.7890	
Hedges' g SMD							-0.04 (-0.355,	0.268)	0.7848	
non-tBRCAm	91 16	.48 (26.469)	2.13 (1.575)	60	13.33 (25.453)	1.21 (2.160)	0.93 (-4.378,	6.235)	0.7297	
Hedges' g SMD							0.06 (-0.267,	0.385)	0.7238	
Int. p-value									0.8861	
Age group										
<65 years	159 16	.35 (28.032)	2.42 (1.345)	90	14.81 (23.497)	2.12 (1.935)	0.30 (-4.343,	4.951)	0.8975	
Hedges' g SMD							0.02 (-0.241,	0.276)	0.8954	
>=65 years	57 20	.47 (27.280)	-1.68 (1.794)	28	11.90 (27.539)	0.87 (2.805)	-2.55 (-9.224,	4.126)	0.4493	
Hedges' g SMD							-0.18 (-0.634,	0.272)	0.4345	
Int. p-value									0.2744	
FIGO Stage (Dis	ease sta	ate)								
III	154 18	.40 (28.275)	2.68 (1.313)	81	14.81 (26.874)	3.22 (1.921)	-0.55 (-5.144,	4.047)	0.8141	
Hedges' g SMD							-0.03 (-0.302,	0.236)	0.8108	
IV	62 15	.05 (26.774)	-1.96 (1.868)	37	12.61 (18.175)	-3.22 (2.830)	1.26 (-5.463,	7.985)	0.7106	
Hedges' g SMD							0.08 (-0.328,	0.487)	0.7011	
Int. p-value									0.2629	
Region										
Europe	207 18	.20 (28.188)	1.25 (1.139)	112	14.29 (24.797)	0.75 (1.689)	0.50 (-3.518,	4.511)	0.8080	
Hedges' g SMD							0.03 (-0.201,	0.259)	0.8030	
Japan Int. p-value	9	NC	NC	6	NC	NC	NC		NC NC	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.8 PAOLAl: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale:

Constipation (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be			Placebo + bev (N=132		Difference between groups			
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value	
ECOG performance	e stat	us at Baseline								
(0) Normal activity	161	15.32 (27.637)	2.08 (1.230)	91	15.38 (25.967)	1.53 (1.731)	0.55 (-3.638,	4.733)	0.7968	
Hedges' g SMD							0.03 (-0.223,	0.291)	0.7940	
(1) Restricted activity	52	24.36 (28.095)	-1.18 (2.668)	27	9.88 (18.057)	1.99 (4.320)	-3.17 (-13.464,	7.121)	0.5409	
Hedges' g SMD Int. p-value							-0.15 (-0.620,	0.311)	0.5165 0.8726	
Baseline CA-125		=								
<=ULN Hedges' g SMD	191	18.85 (28.708)	0.79 (1.175)	105	14.60 (25.286)	1.54 (1.716)	-0.75 (-4.854, -0.04 (-0.283,	,	0.7187 0.7124	
>ULN Int. p-value	25	NC	NC	13	NC	NC	NC	0.193)	NC NC	
Histological gra	ade									
High grade Hedges' g SMD Int. p-value	216	17.44 (27.832)	1.41 (1.100)	118	14.12 (24.424)	1.49 (1.611)	-0.09 (-3.931, -0.01 (-0.230,	•	0.9643 0.9635 ID	
Cytoreductive s	ırgerv	outcome								
No residue		19.62 (29.833)	0.15 (1.312)	72	14.35 (26.137)	1.33 (1.940)	-1.18 (-5.811,	3.450)	0.6157	
Hedges' g SMD							-0.07 (-0.358,	0.210)	0.6093	
Residue	68	13.73 (23.909)	3.99 (2.113)	40	10.83 (17.521)	3.48 (3.113)	0.51 (-6.967,	7.982)	0.8930	
Hedges' g SMD Int. p-value							0.03 (-0.363,	0.418)	0.8900 0.3505	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.8 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale:

Constipation (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be			Placebo + bev		Difference between groups			
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value		
Timing of cytore	oduati:	uo gurgoru								
Upfront Hedges' g SMD	126	18.78 (30.839)	2.22 (1.479)		13.62 (26.172)	1.88 (2.105)	0.33 (-4.751, 5.420) 0.02 (-0.271, 0.310)	0.8969 0.8951		
Interval Hedges' g SMD Int. p-value	83 1	16.06 (23.490)	-0.04 (1.584)	41	12.20 (17.882)	1.95 (2.507)	-1.99 (-7.881, 3.909) -0.13 (-0.506, 0.243)	0.5055 0.4905 0.9776		
Myriad tumour BI	RCA mu	tation status								
tBRCAm Hedges' g SMD	134	18.41 (28.192)	0.85 (1.455)	66	13.64 (22.628)	0.36 (2.221)	0.48 (-4.765, 5.728) 0.03 (-0.267, 0.323)	0.8565 0.8532		
Non-tBRCAm Hedges' g SMD Int. p-value	82 1	15.85 (27.330)	2.25 (1.668)	52	14.74 (26.743)	2.77 (2.328)	-0.53 (-6.214, 5.159) -0.03 (-0.381, 0.314)	0.8545 0.8512 0.5504		
Status somatic H	SRCA m	utations								
sBRCAm Hedges' g SMD		17.65 (20.809)	-2.19 (3.769)	6	16.67 (18.257)	6.03 (6.792)	-8.22 (-24.582, 8.144) -0.50 (-1.445, 0.443)	0.3048 0.2982		
gBRCAm Hedges' g SMD	56 3	19.64 (27.544)	3.25 (2.683)	28	16.67 (23.130)	2.23 (4.025)	1.01 (-8.633, 10.658) 0.05 (-0.405, 0.503)	0.8348 0.8324		
Non-BRCAm Hedges' g SMD Int. p-value	36 3	16.67 (25.820)	0.08 (2.497)	22	16.67 (30.429)	5.43 (3.496)	-5.35 (-13.961, 3.258) -0.34 (-0.874, 0.194)	0.2181 0.2123 0.4902		

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.9 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Diarrhoea (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + bev		Difference between groups			
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value	
First line treat	ment	outcome (IVRS)								
NED [PDS]	79	7.17 (15.717)	1.32 (1.416)	43	12.40 (20.604)	2.90 (2.009)	-1.58 (-6.462,	3.308)	0.5237	
Hedges' g SMD							-0.12 (-0.494,	0.249)	0.5184	
NED/CR [IDS]	61	12.57 (21.225)	-0.46 (1.966)	34	8.82 (23.654)	0.84 (2.830)	-1.30 (-8.194,	5.592)	0.7075	
Hedges' g SMD							-0.08 (-0.501,	0.338)	0.7023	
NED/CR [Chemo]	37	17.12 (25.606)	-3.66 (1.585)	20	23.33 (37.619)	-3.88 (2.526)	0.22 (-5.824,	6.257)	0.9428	
Hedges' g SMD							0.02 (-0.523,	0.565)	0.9402	
PR	40	NC	NC	22	NC	NC	NC		NC	
Int. p-value									0.9160	
tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value		9.56 (20.067) 9.09 (17.307)	-0.05 (1.104) 2.38 (1.609)		11.30 (23.660) 15.00 (26.343)	1.45 (1.729) 0.79 (2.095)	-1.50 (-5.553, -0.12 (-0.425, 1.59 (-3.665, 0.10 (-0.227,	0.191)	0.4649 0.4563 0.5498 0.5439 0.3174	
First line treat		, ,								
NED [PDS]	77	7.36 (15.879)	1.23 (1.454)	42	11.90 (20.589)	2.13 (2.040)	-0.90 (-5.881,		0.7213	
Hedges' g SMD							-0.07 (-0.445,		0.7188	
NED/CR [IDS]	60	9.44 (19.496)	2.49 (1.908)	29	6.90 (18.643)	3.48 (2.995)	-0.99 (-8.072,		0.7816	
Hedges' g SMD	2.4	10 62 /06 165	0.05 / 1.050	1.5	18 65 (22 552)	2 01 / 2 251	-0.06 (-0.508,		0.7760	
NED/CR [Chemo]	34	18.63 (26.197)	-2.87 (1.870)	Τ./	17.65 (33.578)	-3.81 (3.361)	0.94 (-6.922,		0.8084	
Hedges' g SMD PR	42	5.43 (14.420)	0.20 (2.112)	20	18.89 (29.921)	0.25 (2.833)	0.08 (-0.505, -0.05 (-7.316,		0.7944	
Hedges' g SMD	43	5.43 (14.420)	0.20 (2.112)	30	10.09 (29.921)	0.25 (2.833)	0.00 (-0.470,		0.9889	
Int. p-value							0.00 (-0.470,	0.403)	0.9886	
TITC. P-Value									0.910	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.9 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Diarrhoea (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be	vacizumab		cebo + bev (N=132		Difference k	oetween g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline		(SD) at Line [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
Screening labor	atorv	tBRCA status (e	CRF)						
tBRCAm	_		-0.09 (1.117)	60 11.11	(23.504)	1.46 (1.712)	-1.55 (-5.584,	2.490)	0.4505
Hedges' g SMD							-0.12 (-0.428,	0.187)	0.4424
non-tBRCAm	91	9.16 (17.260)	2.28 (1.563)	59 15.25	(26.495)	0.79 (2.094)	1.49 (-3.707,	6.690)	0.5706
Hedges' g SMD							0.10 (-0.231,	0.424)	0.5640
Int. p-value									0.3367
Age group									
<65 years	160	7.92 (16.503)	-0.22 (1.012)	90 15.56	(26.061)	0.70 (1.444)	-0.93 (-4.425,	2.568)	0.6014
Hedges' g SMD							-0.07 (-0.329,	0.188)	0.5928
>=65 years	57	13.45 (24.283)	3.58 (2.026)	29 5.75	(20.057)	3.32 (3.162)	0.26 (-7.319,	7.836)	0.9457
Hedges' g SMD							0.02 (-0.431,	0.463)	0.9437
Int. p-value									0.6227
FIGO Stage (Dis	ease s	state)							
III	155	10.32 (19.580)	2.06 (1.121)	82 8.94	(19.632)	1.44 (1.629)	0.62 (-3.285,	4.521)	0.7552
Hedges' g SMD							0.04 (-0.224,	0.311)	0.7514
IV	62	6.99 (17.217)	-1.57 (1.659)	37 22.52	(32.446)	2.50 (2.424)	-4.06 (-10.084,	1.955)	0.1829
Hedges' g SMD							-0.29 (-0.703,	0.115)	0.1589
Int. p-value									0.1430
Region									
Europe	208	9.46 (19.142)	0.90 (0.954)	113 13.27	(25.409)	1.41 (1.397)	-0.50 (-3.839,	2.834)	0.7671
Hedges' g SMD							-0.04 (-0.264,	0.194)	0.7619
Japan Int. p-value	9	NC	NC	6 N	С	NC	NC		NC NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.

SMD = standardised mean difference. * p<0.05.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.9 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Diarrhoea (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + bev		Difference l	oetween g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
ECOG performance	stat	us at Baseline							
(0) Normal activity	161	9.32 (19.441)	0.75 (1.099)	91	13.19 (25.278)	1.84 (1.520)	-1.10 (-4.802,	2.611)	0.5607
Hedges' g SMD							-0.08 (-0.334,		0.5563
(1) Restricted activity	53	10.06 (18.004)	2.71 (1.901)	28	13.10 (24.578)	0.36 (3.181)	2.36 (-5.046,	9.760)	0.5270
Hedges' g SMD Int. p-value							0.16 (-0.302,	0.615)	0.5039 0.2251
Baseline CA-125									
<=ULN Hedges' g SMD	192	9.72 (19.545)	0.82 (0.977)	106	11.95 (22.626)	1.41 (1.410)	-0.59 (-3.972, -0.04 (-0.280,		0.7303 0.7253
>ULN Int. p-value	25	NC	NC	13	NC	NC	0.04 (0.200, NC	0.193)	NC NC
Histological gra	ıde								
High grade	217	9.37 (18.956)	1.00 (0.927)	119	13.17 (25.012)	1.50 (1.341)	-0.50 (-3.715,	2.714)	0.7593
Hedges' g SMD Int. p-value							-0.04 (-0.259,	0.188)	0.7546 ID
Cytoreductive su	ırgery	outcome							
No residue		8.33 (17.486)	1.64 (1.160)	72	9.72 (19.730)	3.03 (1.702)	-1.39 (-5.456,	2.675)	0.5006
Hedges' g SMD							-0.10 (-0.384,		0.4944
Residue	70	12.38 (22.105)	-1.30 (1.484)	39	20.51 (32.994)	-3.22 (2.168)	1.92 (-3.347,		0.4703
Hedges' g SMD Int. p-value							0.15 (-0.244,	0.540)	0.4587

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.9 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Diarrhoea (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + be		Difference between o	groups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Timing of cytore	educti	ve surgery						
Upfront	127	9.45 (19.198)	0.63 (1.124)	70	15.71 (26.449)	0.31 (1.586)	0.31 (-3.540, 4.169)	0.8721
Hedges' g SMD							0.02 (-0.267, 0.316)	0.8702
Interval	83	10.04 (19.293)	1.10 (1.549)	41	9.76 (23.856)	1.97 (2.435)	-0.88 (-6.608, 4.854)	0.7620
Hedges' g SMD							-0.06 (-0.434, 0.315)	0.7547
Int. p-value								0.8974
Myriad tumour BI	RCA mu	itation status						
tBRCAm	135	9.63 (20.307)	-0.82 (1.100)	68	13.73 (27.157)	2.06 (1.648)	-2.88 (-6.804, 1.037)	0.1484
Hedges' g SMD							-0.22 (-0.512, 0.072)	0.1400
Non-tBRCAm	82	8.94 (16.604)	3.84 (1.631)	51	12.42 (22.071)	0.29 (2.238)	3.55 (-1.956, 9.056)	0.2034
Hedges' g SMD							0.23 (-0.119, 0.582)	0.1955
Int. p-value								0.0281
Status somatic E	BRCA m	nutations						
sBRCAm	17	9.80 (22.866)	-4.92 (2.953)	6 :	22.22 (34.427)	-1.76 (5.106)	-3.16 (-15.563, 9.242)	0.6008
Hedges' g SMD							-0.25 (-1.183, 0.685)	0.6019
gBRCAm	56	9.52 (22.665)	2.60 (1.791)	28	7.14 (16.623)	1.14 (2.782)	1.46 (-5.141, 8.054)	0.6614
Hedges' g SMD							0.10 (-0.350, 0.558)	0.6529
Non-BRCAm	36	9.26 (15.142)	5.80 (2.491)	22	9.09 (23.417)	3.73 (3.655)	2.07 (-6.896, 11.042)	0.6430
Hedges' g SMD							0.13 (-0.402, 0.660)	0.6328
Int. p-value								0.6544

SMD = standardised mean difference. * p<0.05.

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.10 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Dyspnoea (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bo (N=25		Placebo + be (N=13		Difference between g	roups
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
First line treat	ment outcome (IVRS)					
NED [PDS]	80 26.67 (28.758)	-2.40 (1.899)	43 20.16 (24.277)	-4.28 (2.716)	1.88 (-4.703, 8.454)	0.5735
Hedges' g SMD					0.11 (-0.263, 0.479)	0.5686
NED/CR [IDS]	61 24.59 (25.749)	-3.49 (2.051)	34 28.43 (32.960)	-6.98 (2.928)	3.49 (-3.632, 10.612)	0.3327
Hedges' g SMD					0.21 (-0.210, 0.632)	0.3253
NED/CR [Chemo]	36 24.07 (27.152)	-1.62 (2.962)	20 28.33 (24.839)	-10.32 (4.300)	8.70 (-1.798, 19.191)	0.1023
Hedges' g SMD					0.47 (-0.085, 1.023)	0.0973
PR	39 18.80 (26.263)	3.87 (3.490)	23 14.49 (19.659)	9.97 (4.893)	-6.10 (-18.223, 6.027)	0.3167
Hedges' g SMD					-0.27 (-0.786, 0.249)	0.3092
Int. p-value						0.2560
Screening labora	atory tBRCA status (1 129 27.65 (28.603)		59 16.95 (24.269)	-3.19 (2.473)	-0.07 (-5.901, 5.762)	0.9813
Hedges' g SMD	125 27.03 (20.003)	3.20 (1.301)	33 10.33 (21.203)	3.13 (2.173)	0.00 (-0.312, 0.304)	0.9809
non-tBRCAm	87 19.16 (24.183)	1.10 (1.837)	61 28.42 (27.780)	-4.12 (2.371)	,	0.0871
Hedges' g SMD	0, 19:10 (21:103)	1.10 (1.057)	01 20.12 (27.700)	1.12 (2.3/1)	0.29 (-0.036, 0.622)	0.0809
Int. p-value					0.25 (0.050, 0.022,	0.2079
First line treat	ment outcome (eCRF)					
NED [PDS]	78 27.78 (30.111)	-2.12 (1.945)	42 20.63 (24.364)	-4.22 (2.733)	2.10 (-4.565, 8.765)	0.5338
Hedges' g SMD					0.12 (-0.255, 0.496)	0.5304
NED/CR [IDS]	59 21.47 (24.575)	-2.93 (2.089)	29 29.89 (33.741)	-5.95 (3.246)	3.02 (-4.716, 10.759)	0.4394
Hedges' g SMD					0.18 (-0.264, 0.627)	0.4251
NED/CR [Chemo]	33 23.23 (25.665)	-1.06 (3.097)	17 27.45 (26.965)	-8.85 (4.851)	7.79 (-3.820, 19.397)	0.1834
Hedges' g SMD					0.41 (-0.178, 1.004)	0.1708
PR	43 20.93 (25.222)	2.20 (3.096)	31 17.20 (20.854)	2.55 (3.846)	-0.35 (-10.236, 9.540)	0.9442
Hedges' g SMD					-0.02 (-0.478, 0.445)	0.9439
Int. p-value						0.7092

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.10 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Dyspnoea (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	_	b + bevacizumab (N=255)	Placebo + be (N=13	– –	Difference between o	groups
Subgroup	n Mean (SD [a] Baseline		n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Screening labor	atory tBRCA sta	tus (eCRF)				
tBRCAm	126 28.04 (28	.725) -3.74 (1.605) 60 16.67 (24.162)	-3.03 (2.460)	-0.71 (-6.548, 5.129)	0.8109
Hedges' g SMD					-0.04 (-0.346, 0.269)	0.8063
non-tBRCAm	90 18.89 (23	.990) 1.58 (1.803) 60 28.89 (27.765)	-4.17 (2.389)	5.75 (-0.229, 11.732)	0.0593
Hedges' g SMD					0.32 (-0.005, 0.653)	0.0536
Int. p-value						0.1256
Age group						
<65 years	159 24.32 (27	.221) -0.59 (1.383	90 23.70 (26.082)	-3.74 (1.946)	3.15 (-1.549, 7.856)	0.1877
Hedges' g SMD					0.18 (-0.083, 0.436)	0.1817
>=65 years	57 23.98 (27	.280) -3.47 (2.320) 30 20.00 (28.500)	-2.98 (3.380)	-0.49 (-8.653, 7.670)	0.9049
Hedges' g SMD					-0.03 (-0.469, 0.415)	0.9037
Int. p-value						0.4536
FIGO Stage (Dis	ease state)					
III	154 25.97 (27	.810) -1.11 (1.398) 83 24.50 (27.588)	-4.31 (1.989)	3.20 (-1.589, 7.995)	0.1891
Hedges' g SMD					0.18 (-0.086, 0.449)	0.1843
IV	62 19.89 (25	.220) -2.17 (2.318) 37 18.92 (24.268)	-0.37 (3.492)	-1.80 (-10.116, 6.526)	0.6695
Hedges' g SMD					-0.09 (-0.499, 0.316)	0.6589
Int. p-value						0.3058
Region						
Europe	207 24.80 (27	.424) -1.72 (1.237) 114 23.68 (26.874)	-3.60 (1.778)	1.89 (-2.377, 6.149)	0.3848
Hedges' g SMD					0.10 (-0.126, 0.332)	0.3766
Japan Int. p-value	9 NC	NC	6 NC	NC	NC	NC NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.10 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Dyspnoea (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be			Placebo + bev		Difference between o	groups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
ECOG performance	statı	us at Baseline						
(0) Normal activity	159 2	21.17 (23.844)	0.01 (1.317)	92	21.74 (25.407)	-1.48 (1.800)	1.49 (-2.910, 5.880)	0.5062
Hedges' g SMD							0.09 (-0.169, 0.345)	0.5026
(1) Restricted activity	53 3	33.33 (33.968)	-6.20 (2.834)	28	26.19 (30.574)	-8.82 (4.429)	2.62 (-7.851, 13.091)	0.6200
Hedges' g SMD							0.12 (-0.338, 0.578)	0.6079
Int. p-value								0.4166
Baseline CA-125	value							
<=ULN	193 2	23.32 (27.064)	-1.74 (1.233)	107	23.99 (27.008)	-3.21 (1.752)	1.47 (-2.751, 5.681)	0.4945
Hedges' g SMD >ULN	23	NC	NC	13	NC	NC	0.08 (-0.153, 0.320) NC	0.4882 NC
Int. p-value	23	IVC	INC	13	IVC	INC	IVC	NC
Histological gra	ıde							
High grade	216 2	24.23 (27.174)	-1.48 (1.194)	120	22.78 (26.633)	-3.40 (1.697)	1.92 (-2.167, 5.997)	0.3567
Hedges' g SMD Int. p-value							0.11 (-0.117, 0.330)	0.3495 ID
Cytoreductive su	ırgery	outcome						
No residue	140 2	25.48 (28.163)	-2.36 (1.408)	72	24.07 (28.648)	-4.84 (2.053)	2.48 (-2.432, 7.384)	0.3211
Hedges' g SMD							0.15 (-0.139, 0.430)	0.3154
Residue	69 2	21.74 (24.136)	0.03 (2.259)	40	25.00 (23.570)	-2.58 (3.200)	2.61 (-5.165, 10.378)	0.5073
Hedges' g SMD Int. p-value							0.13 (-0.256, 0.524)	0.5005 0.7691

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.10 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Dyspnoea (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be N=255			Placebo + be N=13		ab	Difference between g	groups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Chan	n (SE) ge from seline	Estimated difference (95% CI)	p-value
Timing of cytore									
Upfront	127 2	24.41 (28.312)	-0.93 (1.542)	71	22.07 (23.867)	-3.40	(2.143)	2.48 (-2.734, 7.688)	0.3496
Hedges' g SMD								0.14 (-0.151, 0.431)	0.3453
Interval	82 2	23.98 (24.720)	-2.91 (1.889)	41	28.46 (31.235)	-6.24	(2.911)	3.33 (-3.565, 10.215)	0.3411
Hedges' g SMD								0.19 (-0.188, 0.563)	0.3280
Int. p-value									0.6551
Myriad tumour BF	RCA mut	tation status							
tBRCAm		26.67 (28.448)	-2.26 (1.552)	68	17.65 (24.751)	-3.32	(2.310)	1.05 (-4.462, 6.564)	0.7074
Hedges' g SMD								0.06 (-0.235, 0.348)	0.7017
Non-tBRCAm	81 2	20.16 (24.540)	0.13 (1.897)	52	29.49 (27.735)	-4.50	(2.554)	4.63 (-1.719, 10.984)	0.1514
Hedges' g SMD								0.26 (-0.088, 0.611)	0.1427
Int. p-value									0.5037
Status somatic E	BRCA mu	ıtations							
sBRCAm	18 2	24.07 (25.063)	-1.12 (0.000)	6	27.78 (32.773)	-11.87	(181.80)	10.76 (-396.81,418.319)	0.9567
Hedges' g SMD		, ,	, ,		, , , , , , , , , , , , , , , , , , , ,		,	0.05 (-0.875, 0.973)	0.9174
gBRCAm	56 3	31.55 (28.723)	-2.39 (2.560)	28	17.86 (26.422)	1.71	(3.894)	-4.11 (-13.493, 5.282)	0.3866
Hedges' g SMD								-0.21 (-0.662, 0.248)	0.3722
Non-BRCAm	35 1	7.14 (18.737)	-2.12 (2.569)	22	25.76 (27.084)	-4.43	(3.524)	2.31 (-6.604, 11.222)	0.6049
Hedges' g SMD								0.14 (-0.389, 0.679)	0.5948
Int. p-value								, , , , , , , , , , , , , , , , , , , ,	0.2598

SMD = standardised mean difference. * p<0.05.

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.11 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Fatigue (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=25		Placebo + ber		Difference between g	roups
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
First line treat	ment outcome (IVRS)					
NED [PDS]	81 32.44 (22.762)	0.59 (1.677)	43 35.92 (21.254)	-1.87 (2.412)	2.46 (-3.365, 8.277)	0.4052
Hedges' g SMD					0.16 (-0.211, 0.529)	0.4000
NED/CR [IDS]	61 32.88 (22.551)	1.46 (2.009)	34 33.82 (23.064)	-4.34 (2.866)	5.80 (-1.166, 12.759)	0.1015
Hedges' g SMD					0.36 (-0.065, 0.781)	0.0969
NED/CR [Chemo]	36 36.73 (26.537)	-1.89 (2.377)	20 32.22 (21.898)	-3.25 (3.526)	1.36 (-7.179, 9.896)	0.7506
Hedges' g SMD					0.09 (-0.456, 0.637)	0.7459
PR	40 29.72 (20.346)	1.08 (2.278)	22 34.85 (25.898)	6.82 (3.237)	-5.74 (-13.678, 2.204)	0.1533
Hedges' g SMD					-0.39 (-0.911, 0.139)	0.1497
Int. p-value						0.1486
Screening labora	atory tBRCA status (I	VRS)				
tBRCAm	130 34.83 (23.578)	-0.05 (1.370)	58 31.42 (22.526)	1.39 (2.155)	-1.44 (-6.489, 3.600)	0.5727
Hedges' g SMD					-0.09 (-0.400, 0.219)	0.5660
non-tBRCAm	88 29.73 (21.608)	1.09 (1.489)	61 37.43 (22.335)	-4.07 (1.891)	5.17 (0.379, 9.952)	0.0346
Hedges' g SMD					0.36 (0.030, 0.689)	0.0324
Int. p-value						0.1028
First line treat	ment outcome (eCRF)					
NED [PDS]	79 34.11 (24.520)	0.55 (1.691)	42 34.13 (20.443)	-1.57 (2.396)	2.12 (-3.686, 7.928)	0.4709
Hedges' g SMD			()	(=.350)	0.14 (-0.236, 0.513)	0.4684
NED/CR [IDS]	59 30.41 (20.435)	1.87 (2.083)	29 34.29 (24.581)	-3.69 (3.220)	5.56 (-2.092, 13.208)	0.1521
Hedges' g SMD	(23, 23, 23,	(= . 300)			0.33 (-0.112, 0.782)	0.1423
NED/CR [Chemo]	33 34.34 (25.210)	-0.53 (2.424)	17 31.37 (21.601)	-3.85 (3.973)	3.32 (-6.028, 12.669)	0.4777
Hedges' g SMD			((2.3737	0.22 (-0.366, 0.808)	0.4605
PR	44 32.32 (21.939)	0.27 (2.277)	30 35.93 (24.314)	2.96 (2.892)	-2.69 (-10.037, 4.665)	0.4685
Hedges' g SMD			,,	,,	-0.17 (-0.637, 0.292)	0.4670
Int. p-value					, , , , , , , , , , , , , , , , , , , ,	0.4257

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

SMD = standardised mean difference. * p<0.05.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.11 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Fatigue (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + bev		Difference l	oetween g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
Screening labor	atory t	BRCA status (e	CRF)						
tBRCAm	127 3	5.21 (23.707)	-0.14 (1.395)	59	31.26 (22.363)	1.14 (2.149)	-1.28 (-6.346,	3.781)	0.6179
Hedges' g SMD							-0.08 (-0.389,	0.229)	0.6119
non-tBRCAm	91 2	9.37 (21.361)	1.18 (1.448)	60	37.69 (22.435)	-4.07 (1.895)	5.24 (0.494,	9.991)	0.0307
Hedges' g SMD							0.37 (0.040,	0.697)	0.0281
Int. p-value									0.1113
Age group									
<65 years	161 3	4.51 (23.492)	0.36 (1.203)	90	34.44 (23.083)	-1.32 (1.697)	1.68 (-2.421,	5.776)	0.4209
Hedges' g SMD							0.11 (-0.151,	0.366)	0.4147
>=65 years	57 2	7.88 (20.506)	1.28 (1.916)	29	34.67 (21.137)	-1.22 (2.851)	2.50 (-4.352,	9.352)	0.4699
Hedges' g SMD							0.17 (-0.280,	0.616)	0.4628
Int. p-value									0.5973
FIGO Stage (Dis	ease st	cate)							
III	155 3	4.70 (24.144)	1.04 (1.239)	82	34.21 (23.432)	-2.53 (1.773)	3.57 (-0.692,	7.831)	0.1002
Hedges' g SMD							0.23 (-0.041,	0.496)	0.0967
IV	63 2	8.04 (18.811)	-0.12 (1.854)	37	35.14 (20.707)	-0.70 (2.822)	0.58 (-6.155,	7.310)	0.8651
Hedges' g SMD							0.04 (-0.369,	0.443)	0.8598
Int. p-value									0.1870
Region									
Europe	209 3	2.80 (23.241)	0.88 (1.053)	113	34.76 (22.686)	-1.13 (1.520)	2.01 (-1.631,	5.645)	0.2786
Hedges' g SMD							0.13 (-0.100,	0.358)	0.2707
Japan	9	NC	NC	6	NC	NC	NC		NC
Int. p-value									NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.11 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Fatigue (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be			Placebo + bev		Difference 1	between g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
ECOG performance	e stat	us at Baseline							
(0) Normal	161 2	29.81 (20.541)	2.95 (1.137)	92	32.49 (22.313)	0.73 (1.566)	2.22 (-1.592,	6.038)	0.2522
activity									
Hedges' g SMD	5 2	41 00 (06 068)	0 15 / 0 205)	0.5	41 26 (00 245)	E 01 (2 506)	0.15 (-0.105,	•	0.2475
(1) Restricted activity	53 4	11.82 (26.967)	-8.17 (2.307)	27	41.36 (22.347)	-7.91 (3.586)	-0.26 (-8.749,	8.237)	0.9522
Hedges' g SMD							-0.01 (-0.478,	0.449)	0.9509
Int. p-value							(,	,	0.4966
Baseline CA-125 <=ULN Hedges' g SMD >ULN Int. p-value		33.05 (23.302) NC	0.42 (1.077) NC	106 13	34.64 (22.150) NC	-1.35 (1.539) NC	1.78 (-1.923, 0.12 (-0.121, NC		0.3455 0.3385 NC NC
Histological gra	ade								
High grade		32.77 (22.890)	0.59 (1.022)	119	34.50 (22.536)	-1.36 (1.459)	1.95 (-1.554,	5.457)	0.2743
Hedges' g SMD Int. p-value							0.13 (-0.097,	0.350)	0.2675 ID
Cytoreductive s	urgery	outcome							
No residue	141 3	32.55 (22.767)	0.87 (1.287)	72	34.65 (22.232)	-2.62 (1.885)	3.49 (-1.017,	7.988)	0.1285
Hedges' g SMD							0.22 (-0.061,	0.508)	0.1236
Residue	70 3	33.97 (23.694)	-0.25 (1.711)	40	34.03 (24.531)	0.06 (2.431)	-0.32 (-6.216,	5.579)	0.9148
Hedges' g SMD							-0.02 (-0.410,	0.367)	0.9135
Int. p-value									0.7677

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.11 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Fatigue (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + bev		Difference between g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
m! !								
Timing of cytore			0 10 / 1 005)	71	24 25 (22 200)	1 00 / 1 002	\ 0.0F / 2.417 F 210\	0 ((00
Upfront	149 3	33.20 (24.121)	-0.13 (1.285)	/ 1	34.35 (22.209)	-1.08 (1.803		0.6682 0.6653
Hedges' g SMD Interval	02 2	22 72 /21 240\	1 44 / 1 750\	41	34.55 (24.517)	-3.66 (2.679	0.06 (-0.226, 0.354)	
Hedges' g SMD	8∠ 3	32.72 (21.348)	1.44 (1.750)	41	34.33 (24.31/)	-3.00 (2.0/9) 5.11 (-1.236, 11.447) 0.31 (-0.066, 0.688)	0.1135 0.1054
Int. p-value							0.31 (-0.000, 0.088)	0.1054
Myriad tumour BF	RCA mut	ation status						
tBRCAm	136 3	4.84 (23.419)	-0.32 (1.359)	67	32.84 (23.165)	-0.90 (2.034) 0.58 (-4.247, 5.405)	0.8133
Hedges' g SMD							0.04 (-0.257, 0.328)	0.8106
Non-tBRCAm	82 2	19.34 (21.689)	1.71 (1.501)	52	36.65 (21.734)	-2.07 (1.999) 3.78 (-1.196, 8.754)	0.1352
Hedges' g SMD							0.27 (-0.079, 0.619)	0.1294
Int. p-value								0.4241
Status somatic E	BRCA mu	ıtations						
sBRCAm	18 3	3.95 (22.698)	-3.20 (3.938)	6	33.33 (32.961)	-7.14 (7.080) 3.95 (-12.932, 20.827)	0.6313
Hedges' g SMD							0.23 (-0.700, 1.153)	0.6324
gBRCAm	57 3	8.69 (22.809)	3.29 (2.265)	28	30.16 (21.566)	0.99 (3.477) 2.31 (-6.020, 10.633)	0.5831
Hedges' g SMD							0.13 (-0.322, 0.583)	0.5726
Non-BRCAm	35 2	8.25 (19.866)	1.58 (2.236)	22	32.07 (18.139)	0.54 (3.029) 1.04 (-6.550, 8.623)	0.7847
Hedges' g SMD							0.08 (-0.458, 0.609)	0.7826
Int. p-value								0.9312

SMD = standardised mean difference. * p<0.05.

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.12 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Financial difficulties (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255	vacizumab		Placebo + be (N=13		Difference between g	roups
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-valu
First line treat	ment o	utcome (IVRS)						
NED [PDS]	81 1	7.70 (28.425)	-3.35 (1.824)	43 2	24.81 (34.190)	-1.19 (2.624)	-2.16 (-8.499, 4.176)	0.5009
Hedges' g SMD							-0.13 (-0.499, 0.241)	0.495
NED/CR [IDS]	60 1	7.78 (30.971)	-1.39 (2.365)	32 3	15.63 (29.310)	-10.72 (3.456)	9.33 (1.015, 17.649)	0.0283
Hedges' g SMD							0.49 (0.058, 0.929)	0.0263
NED/CR [Chemo]	36 1	9.44 (32.244)	1.01 (2.778)	20 1	18.33 (27.519)	-0.11 (3.823)	1.12 (-8.382, 10.624)	0.813
Hedges' g SMD							0.07 (-0.481, 0.612)	0.8138
PR	41	NC	NC	22	NC	NC	NC	NC
Int. p-value								0.0240
						2 01 / 0 565	0 00 / 5 150 / 005\	0 000
tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value		1.09 (29.540) 4.81 (30.440)	, ,		24.29 (31.459) 13.22 (27.173)	-3.91 (2.565) 0.03 (2.310)	0.87 (-5.157, 6.905) 0.05 (-0.263, 0.354) -0.89 (-6.639, 4.856) -0.05 (-0.382, 0.278)	0.771 0.759 0.757
Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value	90 1	4.81 (30.440)	-0.87 (1.763)	58 3	13.22 (27.173)	0.03 (2.310)	0.05 (-0.263, 0.354) -0.89 (-6.639, 4.856) -0.05 (-0.382, 0.278)	0.775 0.771 0.759 0.757 0.472
Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value First line treat NED [PDS]	90 1	4.81 (30.440)	, ,	58 3	13.22 (27.173)	0.03 (2.310)	0.05 (-0.263, 0.354) -0.89 (-6.639, 4.856) -0.05 (-0.382, 0.278) -1.40 (-8.060, 5.262)	0.771 0.759 0.757 0.472
Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value First line treat NED [PDS] Hedges' g SMD	90 1 ment o 79 1	4.81 (30.440) utcome (eCRF) 7.72 (28.662)	-0.87 (1.763) -2.55 (1.943)	58 £	23.81 (33.966)	0.03 (2.310)	0.05 (-0.263, 0.354) -0.89 (-6.639, 4.856) -0.05 (-0.382, 0.278) -1.40 (-8.060, 5.262) -0.08 (-0.454, 0.295)	0.771 0.759 0.757 0.472 0.678
Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value First line treat NED [PDS] Hedges' g SMD NED/CR [IDS]	90 1 ment o 79 1	4.81 (30.440)	-0.87 (1.763)	58 £	23.81 (33.966)	0.03 (2.310)	0.05 (-0.263, 0.354) -0.89 (-6.639, 4.856) -0.05 (-0.382, 0.278) -1.40 (-8.060, 5.262) -0.08 (-0.454, 0.295) 11.98 (3.249, 20.720)	0.771 0.759 0.757 0.472 0.678 0.676 0.007
Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value First line treat NED [PDS] Hedges' g SMD NED/CR [IDS] Hedges' g SMD	90 1 ment o 79 1 59 1	4.81 (30.440) utcome (eCRF) 7.72 (28.662) 8.64 (31.117)	-0.87 (1.763) -2.55 (1.943) 0.24 (2.351)	42 2	23.81 (33.966) 13.58 (29.612)	0.03 (2.310) -1.15 (2.740) -11.74 (3.713)	0.05 (-0.263, 0.354) -0.89 (-6.639, 4.856) -0.05 (-0.382, 0.278) -1.40 (-8.060, 5.262) -0.08 (-0.454, 0.295) 11.98 (3.249, 20.720) 0.64 (0.178, 1.110)	0.771 0.759 0.757 0.472 0.678 0.676 0.007 0.006
Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value First line treat NED [PDS] Hedges' g SMD	90 1 ment o 79 1	4.81 (30.440) utcome (eCRF) 7.72 (28.662)	-0.87 (1.763) -2.55 (1.943)	58 £	23.81 (33.966)	0.03 (2.310)	0.05 (-0.263, 0.354) -0.89 (-6.639, 4.856) -0.05 (-0.382, 0.278) -1.40 (-8.060, 5.262) -0.08 (-0.454, 0.295) 11.98 (3.249, 20.720)	0.771 0.759 0.757 0.472 0.678 0.676

Screening laboratory tBRCA status (eCRF)

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.12 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Financial difficulties (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=25		Placebo + bev		Difference between groups		
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value	125 21.33 (29.754) 93 14.70 (30.081)	-2.81 (1.679) -1.19 (1.735)	60 23.89 (31.349) 57 13.45 (27.357)	-4.08 (2.543) 0.16 (2.331)	1.27 (-4.744, 7.287) 0.07 (-0.242, 0.374) -1.36 (-7.104, 4.391) -0.08 (-0.409, 0.251)	0.6771 0.6726 0.6413 0.6383 0.3512	
Age group <65 years Hedges' g SMD >=65 years Hedges' g SMD Int. p-value	160 23.13 (31.941) 58 5.75 (18.875)	, ,	, ,	-4.43 (2.118) 5.02 (2.498)	2.54 (-2.562, 7.636) 0.13 (-0.129, 0.392) -7.22 (-13.274, -1.158) -0.54 (-0.996, -0.090)	0.3280 0.3215 0.0203* 0.0189* 0.0430*	
FIGO Stage (Dise	•						
III Hedges' g SMD	155 18.28 (29.716)	-1.47 (1.456)	81 19.34 (31.120)	-3.17 (2.100)	1.70 (-3.338, 6.731) 0.09 (-0.177, 0.361)	0.5074 0.5027	
IV Hedges' g SMD Int. p-value	63 19.05 (30.944)	-2.95 (2.393)	36 17.59 (27.005)	1.31 (3.541)	-4.26 (-12.758, 4.242) -0.21 (-0.624, 0.198)	0.3217 0.3093 0.6252	
Region							
Europe Hedges' g SMD	209 18.66 (30.268)	-2.19 (1.242)	111 18.62 (30.045)	-1.83 (1.804)	-0.36 (-4.665, 3.953) -0.02 (-0.250, 0.211)	0.8709 0.8689	
Japan Int. p-value	9 NC	NC	6 NC	NC	NC	NC NC	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

SMD = standardised mean difference. * p<0.05.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Table 2.5.4.12 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Financial difficulties (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=255		Placebo + bev		Difference between	groups
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
(0) Normal activity Hedges' g SMD	161 18.63 (30.010)	-2.38 (1.414)	89 17.60 (28.020)	-2.47 (1.964)	0.09 (-4.674, 4.861) 0.01 (-0.254, 0.264)	
(1) Restricted activity Hedges' g SMD	54 18.52 (30.828)	-0.44 (2.303)	28 22.62 (35.199)	-4.97 (3.542)	4.53 (-3.887, 12.942) 0.26 (-0.203, 0.714)	
Int. p-value					0.20 (0.203, 0.714)	0.6449
Baseline CA-125		0.50 (1.000)	104.10.01.(20.055)	0.75 / 1.040		0.0156
<=ULN	193 19.00 (30.172)	-2.52 (1.290)	104 18.91 (30.375)	-2.76 (1.848)		
Hedges' g SMD >ULN Int. p-value	25 NC	NC	13 NC	NC	0.01 (-0.225, 0.251) NC	0.9145 NC NC
Histological gra	ade					
High grade Hedges' g SMD Int. p-value	218 18.50 (30.006)	-2.08 (1.206)	117 18.80 (29.809)	-2.47 (1.730)	0.40 (-3.752, 4.544) 0.02 (-0.203, 0.246)	
Cytoreductive s						
No residue Hedges' g SMD	141 17.73 (29.427)	-1.40 (1.497)	70 20.48 (32.745)	-5.09 (2.210)	3.69 (-1.571, 8.955) 0.20 (-0.083, 0.491)	
Residue Hedges' g SMD Int. p-value	70 20.48 (30.715)	-3.62 (2.109)	39 15.38 (25.185)	1.49 (3.040)	-5.11 (-12.454, 2.240) -0.28 (-0.673, 0.114)	
Timing of cytore	eductive surgery					
Upfront	130 18.72 (29.931)	-4.06 (1.495)	70 21.90 (31.539)	0.22 (2.104)	-4.28 (-9.372, 0.815)	0.0991

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.

SMD = standardised mean difference. * p<0.05.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.12 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Financial difficulties (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be N=255			Placebo + be (N=13		Differ	ence between g	roups
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline		mated ce (95% CI)	p-value
Hedges' g SMD	[]	sasciine (z)	Dabetine	[]	Dagetine (2)	202011110		.539, 0.044)	0.0965
Interval	81 18	.52 (29.814)	0.56 (2.049)	39 1	.2.82 (27.161)	-9.23 (3.205)	9.79 (2	.264, 17.316)	0.0112*
Hedges' g SMD Int. p-value							0.51 (0	.125, 0.901)	0.0095* 0.0013*
Myriad tumour BI	RCA muta	ation status							
tBRCAm	134 21	.14 (29.629)	-2.84 (1.641)	68 2	21.57 (29.233)	-3.47 (2.416)	•	.134, 6.384)	0.8307
Hedges' g SMD							0.03 (-0	.260, 0.324)	0.8285
Non-tBRCAm	84 14	.29 (30.298)	-1.15 (1.707)	49 1	.4.97 (30.476)	-0.88 (2.349)	-0.28 (-6	.032, 5.481)	0.9247
Hedges' g SMD							-0.02 (-0	.369, 0.335)	0.9239
Int. p-value									0.6095
Status somatic I	BRCA mut	ations							
sBRCAm	18	NC	NC	6	NC	NC		NC	NC
gBRCAm	55 24	.85 (33.468)	-2.05 (2.478)	28 2	23.81 (29.893)	-10.83 (3.751)	8.78 (-0	.173, 17.738)	0.0545
Hedges' g SMD							0.46 (0	.000, 0.922)	0.0499*
Non-BRCAm	37	NC	NC	21	NC	NC		NC	NC
Int. p-value									NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.13 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Nausea and vomiting (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bev (N=255			Placebo + be		Difference between g	roups
Subgroup	n [a]		Mean (SE) Change from baseline	n [a]		Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
First line treat	ment	outcome (IVRS)						
NED [PDS] Hedges' g SMD		4.53 (10.210)	7.18 (1.159)	43	3.10 (6.562)	1.20 (1.667)	5.97 (1.947, 9.995) 0.56 (0.183, 0.936)	0.0040* 0.0036*
NED/CR [IDS]	61	6.01 (16.951)	6.64 (1.610)	34	0.98 (3.981)	0.02 (2.285)	6.62 (1.029, 12.208)	0.0209*
Hedges' g SMD NED/CR [Chemo]	36	3.24 (8.746)	3.44 (1.030)	20	4.17 (13.107)	-0.55 (1.575)	0.51 (0.085, 0.937) 3.99 (0.189, 7.797)	0.0187* 0.0401*
Hedges' g SMD PR	41	NC	NC	23	NC	NC	0.61 (0.047, 1.165) NC	0.0336* NC
Int. p-value								0.6571
Screening labora	atory	tBRCA status (IV	7RS)					
tBRCAm Hedges' g SMD	130	5.00 (13.553)	6.01 (1.026)	59	4.24 (10.538)	0.81 (1.592)	5.20 (1.464, 8.940) 0.44 (0.126, 0.748)	0.0066* 0.0059*
non-tBRCAm	89	3.56 (8.873)	6.47 (0.910)	61	2.46 (7.351)	0.50 (1.172)	5.97 (3.033, 8.907)	<0.0001*
Hedges' g SMD Int. p-value							0.67 (0.339, 1.008)	<0.0001* 0.9133
First line treat	ment	Outcome (eCRF)						
NED [PDS]		4.43 (10.235)	7.06 (1.184)	42	3.17 (6.624)	1.20 (1.686)	5.86 (1.775, 9.944)	0.0053*
Hedges' g SMD							0.55 (0.165, 0.927)	0.0049*
NED/CR [IDS] Hedges' g SMD	60	3.06 (9.939)	7.01 (1.098)	29	1.15 (4.298)	1.27 (1.700)	5.74 (1.696, 9.780) 0.65 (0.198, 1.106)	0.0060* 0.0049*
NED/CR [Chemo]	33	3.54 (9.088)	3.29 (1.083)	17	5.88 (14.363)	-2.66 (1.889)	5.95 (1.536, 10.363)	0.0049*
Hedges' g SMD		•	,		,		0.86 (0.252, 1.474)	0.0056*
PR	44	NC	NC	31	NC	NC	NC	NC
Int. p-value								0.9174

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.13 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Nausea and vomiting (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bev (N=255			Placebo + bev		Difference between	groups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Screening labor	atory	tBRCA status (e0	'유무')					
tBRCAm	_	5.12 (13.691)	6.00 (1.042)	60	4.17 (10.462)	0.69 (1.585)	5.31 (1.562, 9.053)	0.0057
Hedges' g SMD	12,	3.12 (13.0)1)	0.00 (1.012)	00	1.17 (10.102)	0.05 (1.505)	0.44 (0.133, 0.754)	0.0051
non-tBRCAm	92	3.44 (8.749)	6.50 (0.892)	60	2.50 (7.406)	0.61 (1.178)	5.89 (2.963, 8.812)	0.0001
Hedges' g SMD	22	3.44 (0.74)	0.30 (0.032)	00	2.30 (7.400)	0.01 (1.170)	0.67 (0.333, 1.001)	
Int. p-value							0.07 (0.333, 1.001)	0.9597
Inc. p-varue								0.9597
Age group								
<65 years	161	4.45 (12.466)	6.76 (0.880)	90	3.52 (9.512)	0.56 (1.233)	6.20 (3.213, 9.184)	<0.0001
Hedges' g SMD							0.54 (0.282, 0.806)	<0.0001
>=65 years	58	4.31 (10.145)	4.87 (1.111)	30	2.78 (7.686)	2.12 (1.718)	2.75 (-1.344, 6.845)	0.1847
Hedges' g SMD							0.31 (-0.133, 0.753)	0.1704
Int. p-value								0.4620
FIGO Stage (Dis	ease s	state)						
III		4.17 (10.129)	6.75 (0.829)	83	3.41 (8.926)	0.48 (1.187)	6.27 (3.418, 9.128)	<0.0001
Hedges' g SMD							0.59 (0.323, 0.866)	<0.0001
IV	63	5.03 (15.449)	4.82 (1.309)	37	3.15 (9.492)	1.24 (1.940)	3.58 (-1.076, 8.238)	0.1302
Hedges' g SMD		(=====,	(,		((,	0.33 (-0.083, 0.734)	0.1188
Int. p-value							0.55 (0.055, 0.751,	0.2738
ine. p varue								0.2750
Region								
Europe	210	4.60 (12.090)	6.33 (0.736)	114	3.36 (9.190)	0.62 (1.063)	5.71 (3.161, 8.252)	
Hedges' g SMD							0.52 (0.290, 0.754)	<0.0001
Japan Int. p-value	9	NC	NC	6	NC	NC	NC	NC NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.13 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Nausea and vomiting (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bev (N=255			Placebo + bev		Difference b	etween g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95	% CI)	p-value
ECOG performance	e stat	tus at Baseline							
(0) Normal activity	162	2.67 (8.079)	7.40 (0.761)	92	3.08 (9.222)	2.03 (1.058)	5.37 (2.802,	7.942)	<0.0001*
Hedges' g SMD							0.54 (0.283,	0.804)	<0.0001*
(1) Restricted activity	53	10.06 (18.588)	2.80 (1.739)	28	4.17 (8.636)	-3.97 (2.625)	6.77 (0.496,	13.047)	0.0348*
Hedges' g SMD							0.51 (0.047,	0.977)	0.0308*
Int. p-value									0.5611
Baseline CA-125	value	9							
<=ULN	194	4.47 (12.355)	5.95 (0.727)	107	3.74 (9.524)	0.90 (1.038)		7.552)	<0.0001*
Hedges' g SMD							,	0.727)	<0.0001*
>ULN Int. p-value	25	NC	NC	13	NC	NC	NC		NC NC
Histological gra	ade								
High grade	219	4.41 (11.873)	6.22 (0.709)	120	3.33 (9.065)	0.79 (1.013)	5.43 (2.995,	7.863)	<0.0001*
Hedges' g SMD Int. p-value							0.51 (0.280,	0.732)	<0.0001* ID
Cytoreductive s	urgery	outcome							
No residue		3.99 (10.108)	6.98 (0.822)	72	2.31 (5.804)	1.02 (1.210)	5.96 (3.071,	8.850)	<0.0001*
Hedges' g SMD							0.60 (0.308,	0.886)	<0.0001*
Residue	70	4.76 (14.783)	4.72 (1.414)	40	4.17 (12.375)	0.60 (1.960)	4.12 (-0.672,	8.908)	0.0913
Hedges' g SMD							0.34 (-0.051,	0.731)	0.0887
Int. p-value									0.5296

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.13 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Nausea and vomiting (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bev			Placebo + bev		Difference between g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Timing of cytor	0d11a+i	avvaavv						
Upfront Hedges' g SMD		3.88 (9.435)	6.26 (0.850)		2.82 (7.961)	1.35 (1.206)	4.91 (1.995, 7.824) 0.50 (0.203, 0.791)	0.0011* 0.0009*
Interval Hedges' g SMD Int. p-value	83	4.82 (14.846)	6.45 (1.360)	41	3.25 (10.014)	0.23 (2.052)	6.22 (1.340, 11.097) 0.49 (0.110, 0.868)	0.0129* 0.0115* 0.4423
Myriad tumour B	RCA mu	utation status						
tBRCAm Hedges' g SMD	136	4.90 (13.321)	6.32 (0.994)	68	3.68 (9.911)	0.86 (1.476)	5.46 (1.948, 8.978) 0.46 (0.167, 0.756)	0.0025* 0.0021*
Non-tBRCAm Hedges' g SMD Int. p-value	83	3.61 (9.035)	6.06 (0.937)	52	2.88 (7.894)	0.42 (1.260)	5.64 (2.532, 8.749) 0.64 (0.286, 0.996)	0.0005* 0.0004* 0.9362
Status somatic	BRCA m	nutations						
sBRCAm	18	NC	NC	6	NC	NC	NC	NC
gBRCAm Hedges' g SMD	5/	7.02 (17.522)	6.37 (1.693)	28	4.76 (10.978)	-1.08 (2.536)	7.45 (1.371, 13.522) 0.57 (0.107, 1.029)	0.0169* 0.0157*
Non-BRCAm	36	1.85 (5.312)	7.35 (1.530)	22	4.55 (10.518)	1.92 (2.102)	5.42 (0.178, 10.671)	0.0430*
Hedges' g SMD Int. p-value							0.57 (0.026, 1.108)	0.0400* 0.4422

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.5.4.14 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Pain (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be		Placebo + bev		Difference between groups		
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
First line treat	ment outcome (IVRS)						
NED [PDS]	82 23.37 (22.810)	0.28 (1.707)	43 30.62 (26.460)	-0.11 (2.470)	0.39 (-5.577, 6.354)	0.8976	
Hedges' g SMD					0.02 (-0.344, 0.394)	0.8962	
NED/CR [IDS]	61 19.95 (21.042)	8.57 (2.203)	34 16.67 (19.678)	6.27 (3.136)	2.30 (-5.322, 9.923)	0.5500	
Hedges' g SMD					0.13 (-0.290, 0.550)	0.5450	
NED/CR [Chemo]	37 23.42 (26.194)	3.69 (3.437)	20 23.33 (19.041)	7.94 (5.120)	-4.25 (-16.667, 8.167)	0.4944	
Hedges' g SMD					-0.19 (-0.739, 0.351)	0.4857	
PR	41 25.20 (24.473)	0.61 (2.569)	23 20.29 (20.693)	9.78 (3.600)	-9.17 (-18.048, -0.302)	0.0429	
Hedges' g SMD					-0.54 (-1.061, -0.022)	0.0412	
Int. p-value						0.5973	
Screening labora tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value	atory tBRCA status (I 131 25.32 (24.227) 90 19.07 (21.135)	7RS) 3.69 (1.651) 3.53 (1.617)		4.65 (2.576) 4.36 (2.087)	-0.96 (-7.003, 5.080) -0.05 (-0.357, 0.257) -0.83 (-6.063, 4.401) -0.05 (-0.378, 0.273)	0.7538 0.7501 0.7539 0.7513 0.7251	
	tment outcome (eCRF)						
NED [PDS]	80 23.54 (23.375)	0.96 (1.759)	42 28.17 (24.827)	1.20 (2.504)	, , , , , , , , , , , , , , , , , , , ,	0.9378	
Hedges' g SMD					-0.01 (-0.388, 0.358)	0.9373	
NED/CR [IDS]	60 17.50 (19.750)	8.75 (2.251)	29 17.82 (20.379)	6.35 (3.503)	, , , , , , , , , , , , , , , , , , , ,	0.5658	
Hedges' g SMD					0.13 (-0.311, 0.577)	0.5565	
NED/CR [Chemo]	34 23.04 (25.955)	2.42 (3.763)	17 23.53 (17.735)	7.70 (5.946)	, , , , , , , , , , , , , , , , , , , ,	0.4570	
Hedges' g SMD					-0.23 (-0.812, 0.356)	0.4438	
PR	44 27.27 (24.136)	2.42 (2.774)	31 22.04 (24.865)	8.34 (3.503)	-5.92 (-14.904, 3.059)	0.1925	
Hedges' g SMD					-0.31 (-0.773, 0.152)	0.1878	
Int. p-value						0.7951	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.14 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Pain (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bev			Placebo + bev (N=132		Difference k	oetween g	roups
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
Screening labor	atory tE	BRCA status (e0	CRF)						
tBRCAm	128 25	.91 (24.194)	3.55 (1.684)	60	23.33 (24.967)	4.72 (2.576)	-1.16 (-7.245,	4.916)	0.7060
Hedges' g SMD							-0.06 (-0.367,	0.247)	0.7017
non-tBRCAm	93 18	.46 (21.062)	3.77 (1.567)	60	23.61 (20.878)	4.19 (2.078)	-0.42 (-5.581,	4.743)	0.8727
Hedges' g SMD							-0.03 (-0.351,	0.298)	0.8709
Int. p-value									0.8341
Age group									
<65 years	162 23	.87 (23.561)	4.04 (1.416)	90	24.63 (23.468)	4.24 (1.999)	-0.20 (-5.029,	4.624)	0.9341
Hedges' g SMD							-0.01 (-0.269,	0.247)	0.9333
>=65 years	59 19	.77 (21.988)	2.71 (1.977)	30	20.00 (21.173)	6.28 (2.967)	-3.56 (-10.654,	3.527)	0.3203
Hedges' g SMD							-0.23 (-0.668,	0.214)	0.3123
Int. p-value									0.7317
FIGO Stage (Dis	ease sta	ate)							
III	158 24	.05 (23.306)	3.11 (1.348)	83	23.49 (23.431)	3.55 (1.940)	-0.44 (-5.095,	4.217)	0.8527
Hedges' g SMD							-0.03 (-0.291,	0.240)	0.8510
IV	63 19	.58 (22.707)	5.03 (2.329)	37	23.42 (22.034)	9.02 (3.365)	-3.99 (-12.127,	4.151)	0.3329
Hedges' g SMD							-0.21 (-0.613,	0.201)	0.3219
Int. p-value									0.8785
Region									
Europe	212 22	.88 (23.333)	3.76 (1.195)	114	23.83 (23.252)	5.47 (1.732)	-1.72 (-5.857,	2.424)	0.4153
Hedges' g SMD							-0.10 (-0.324,	0.131)	0.4074
Japan	9	NC	NC	6	NC	NC	NC		NC
Int. p-value									NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.14 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Pain (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be			Placebo + bev N=132		Difference between groups		
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
ECOG performance	e stat	us at Baseline							
(0) Normal activity	163	20.86 (22.016)	4.97 (1.296)	92	20.47 (22.114)	5.14 (1.789)	-0.16 (-4.517,	4.190)	0.9411
Hedges' g SMD							-0.01 (-0.265,	0.246)	0.9406
(1) Restricted activity	54	27.47 (25.730)	-1.27 (2.727)	28	33.33 (23.130)	3.72 (4.212)	-4.98 (-14.978,	5.011)	0.3242
Hedges' g SMD Int. p-value							-0.24 (-0.695,	0.221)	0.3103 0.4342
Baseline CA-125									
<=ULN	196	22.45 (23.340)	3.04 (1.206)	107	23.36 (22.297)	5.41 (1.724)	-2.37 (-6.512,		0.2604
Hedges' g SMD >ULN Int. p-value	25	NC	NC	13	NC	NC	-0.14 (-0.373, NC	0.098)	0.2535 NC NC
Histological gra	ade								
High grade	221	22.78 (23.173)	3.60 (1.166)	120	23.47 (22.917)	4.70 (1.669)	-1.10 (-5.103,	2.908)	0.5901
Hedges' g SMD Int. p-value							-0.06 (-0.284,	0.160)	0.5847 ID
Cytoreductive su	ırgery	outcome							
No residue	143	21.33 (22.150)	3.89 (1.356)	72	24.07 (23.385)	3.47 (2.000)	0.41 (-4.355,	5.181)	0.8646
Hedges' g SMD							0.02 (-0.258,	0.308)	0.8628
Residue	71	26.76 (25.585)	2.79 (2.406)	40	21.67 (23.020)	6.85 (3.466)	-4.05 (-12.479,	4.370)	0.3418
Hedges' g SMD							-0.19 (-0.581,	0.195)	0.3302
Int. p-value									0.5893

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.14 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Symptom scale: Pain (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=25		Placebo + bev		Difference between groups		
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Timing of cytore	eductive surgery						
Upfront	131 24.30 (24.131)	1.33 (1.499)	71 27.46 (24.250)	2.05 (2.113)	-0.72 (-5.834, 4.390)	0.7808	
Hedges' g SMD					-0.04 (-0.330, 0.247)	0.7789	
Interval	83 21.29 (22.287)	7.28 (1.916)	41 15.85 (19.347)	6.96 (2.929)	0.32 (-6.624, 7.270)	0.9268	
Hedges' g SMD					0.02 (-0.356, 0.392)	0.9251	
Int. p-value						0.8805	
Myriad tumour BI	RCA mutation status						
tBRCAm	137 24.70 (24.178)	3.68 (1.584)	68 23.28 (24.458)	3.16 (2.363)	0.53 (-5.085, 6.140)	0.8531	
Hedges' g SMD					0.03 (-0.263, 0.319)	0.8510	
Non-tBRCAm	84 19.64 (21.202)	3.76 (1.716)	52 23.72 (20.964)	6.00 (2.311)	-2.24 (-7.951, 3.464)	0.4381	
Hedges' g SMD					-0.14 (-0.485, 0.208)	0.4325	
Int. p-value						0.2513	
Status somatic H	BRCA mutations						
sBRCAm	18 24.07 (22.304)	-2.55 (3.514)	6 11.11 (17.213)	1.80 (6.664)	-4.35 (-20.181, 11.482)	0.5749	
Hedges' g SMD					-0.28 (-1.203, 0.652)	0.5606	
gBRCAm	57 24.56 (23.168)	9.32 (2.326)	28 25.60 (26.637)	-0.29 (3.522)	9.60 (1.206, 17.999)	0.0255	
Hedges' g SMD					0.53 (0.071, 0.991)	0.0235	
Non-BRCAm	37 21.17 (21.748)	3.17 (2.130)	22 18.18 (20.515)	7.24 (3.054)	-4.08 (-11.543, 3.393)	0.2791	
Hedges' g SMD					-0.30 (-0.829, 0.232)	0.2704	
Int. p-value						0.0130	

SMD = standardised mean difference. * p<0.05.

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.5.4.15 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Insomnia (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bev (N=255		Placebo + bev (N=132		Difference between groups		
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
First line treat	ment outcome (IVRS)						
NED [PDS]	80 29.17 (32.424)	4.94 (2.451)	43 23.26 (26.761)	6.56 (3.497)	-1.61 (-10.089, 6.867)	0.7073	
Hedges' g SMD					-0.07 (-0.443, 0.299)	0.7041	
NED/CR [IDS]	60 27.78 (28.235)	4.26 (2.715)	34 16.67 (20.515)	4.94 (3.889)	-0.67 (-10.163, 8.816)	0.8880	
Hedges' g SMD					-0.03 (-0.452, 0.390)	0.8856	
NED/CR [Chemo]	36 30.56 (32.244)	6.85 (3.765)	20 26.67 (29.814)	10.35 (5.677)	-3.50 (-17.213, 10.208)	0.6099	
Hedges' g SMD					-0.15 (-0.694, 0.401)	0.6004	
PR	41 25.20 (27.669)	3.49 (3.063)	23 26.09 (24.529)	7.03 (4.455)	-3.54 (-14.395, 7.311)	0.5155	
Hedges' g SMD					-0.17 (-0.684, 0.339)	0.5081	
Int. p-value						0.9411	
Screening labora tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value	atory tBRCA status (IV 128 31.25 (31.233) 89 23.97 (28.422)	7RS) 3.90 (1.868) 6.24 (2.247)	, ,	6.61 (2.911) 5.76 (2.899)	-2.71 (-9.565, 4.144) -0.13 (-0.434, 0.183) 0.48 (-6.773, 7.743) 0.02 (-0.304, 0.348)	0.4362 0.4260 0.8951 0.8941 0.9013	
	tment outcome (eCRF)						
NED [PDS]	78 32.05 (34.584)	4.52 (2.524)	42 23.02 (27.038)	7.48 (3.544)		0.5010	
Hedges' g SMD					-0.13 (-0.506, 0.245)	0.4969	
NED/CR [IDS]	59 26.55 (27.529)	3.61 (2.707)	29 16.09 (21.121)	6.17 (4.231)	-2.56 (-12.593, 7.471)	0.6128	
Hedges' g SMD					-0.12 (-0.563, 0.327)	0.6023	
NED/CR [Chemo]	33 27.27 (30.567)	7.02 (4.030)	17 23.53 (30.652)	9.52 (6.678)	-2.50 (-18.219, 13.216)	0.7497	
Hedges' g SMD					-0.10 (-0.685, 0.486)	0.7386	
PR	44 25.76 (25.783)	4.85 (2.897)	31 26.88 (23.443)	2.77 (3.818)		0.6662	
Hedges' g SMD					0.10 (-0.358, 0.562)	0.6628	
Int. p-value						0.7512	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.15 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Insomnia (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + (N=2		Placebo + bev		Difference between g	roups
Subgroup	n Mean (SD) at [a] Baseline [b]		n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Screening labor	atory tBRCA status	(eCRF)				
tBRCAm	125 32.00 (31.224) 3.72 (1.897)	60 21.67 (24.414)	6.42 (2.901)	-2.70 (-9.577, 4.176)	0.4393
Hedges' g SMD					-0.12 (-0.432, 0.184)	0.4289
non-tBRCAm	92 23.19 (28.275) 6.45 (2.191)	60 23.33 (26.254)	5.67 (2.897)	0.78 (-6.402, 7.966)	0.8299
Hedges' g SMD					0.04 (-0.289, 0.361)	0.8281
Int. p-value						0.9120
Age group						
<65 years	159 29.77 (30.141) 5.75 (1.685)	90 23.70 (25.106)	6.85 (2.387)	-1.10 (-6.872, 4.664)	0.7064
Hedges' g SMD					-0.05 (-0.309, 0.208)	0.7014
>=65 years	58 24.14 (30.457) 2.84 (2.479)	30 18.89 (25.795)	1.54 (3.671)	1.29 (-7.562, 10.149)	0.7717
Hedges' g SMD					0.07 (-0.374, 0.507)	0.7677
Int. p-value						0.4887
FIGO Stage (Dis	ease state)					
III	155 29.03 (30.320) 4.83 (1.741)	83 23.29 (27.906)	3.98 (2.494)	0.85 (-5.155, 6.861)	0.7799
Hedges' g SMD					0.04 (-0.228, 0.305)	0.7767
IV	62 26.34 (30.261) 4.59 (2.463)	37 20.72 (18.175)	9.19 (3.596)	-4.61 (-13.284, 4.072)	0.2944
Hedges' g SMD					-0.22 (-0.633, 0.184)	0.2811
Int. p-value						0.1953
Region						
Europe	208 28.85 (30.581) 5.10 (1.458)	114 22.81 (25.596)	5.84 (2.114)	-0.74 (-5.804, 4.324)	0.7739
Hedges' g SMD					-0.03 (-0.263, 0.194)	0.7691
Japan Int. p-value	9 NC	NC	6 NC	NC	NC	NC NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.15 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Insomnia (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olapa	arib + bev (N=255			Placebo + bev (N=132		Difference between groups			
Subgroup	n Mean [a] Basel	(SD) at ine [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimate difference (9		p-value	
ECOG performance	e status at I	Baseline								
(0) Normal activity	160 24.79 ((26.763)	6.04 (1.591)	92 1	9.93 (24.239)	7.82 (2.183)	-1.77 (-7.105,	3.559)	0.5129	
Hedges' g SMD							-0.09 (-0.343,		0.5084	
(1) Restricted activity	53 40.88 ((36.772)	0.40 (3.271)	28 3	0.95 (27.108)	-1.39 (5.097)	1.78 (-10.329,	13.895)	0.7701	
Hedges' g SMD							0.07 (-0.387,	0.529)	0.7619	
Int. p-value									0.7896	
Baseline CA-125	value									
<=ULN	193 28.50 ((30.613)	4.88 (1.463)	107 2	3.36 (25.988)	5.80 (2.095)	,		0.7183	
Hedges' g SMD >ULN	24 NO	7	NC	13	NC	NC	-0.04 (-0.281,	0.192)	0.7133 NC	
Int. p-value	24 100	-	NC	13	INC	INC	INC		NC	
Histological gra	ade									
High grade	217 28.26 ((30.258)	4.81 (1.421)	120 2	2.50 (25.258)	5.74 (2.031)	-0.93 (-5.818,	3.957)	0.7083	
Hedges' g SMD Int. p-value							-0.04 (-0.266,	0.180)	0.7031 ID	
Cytoreductive s	urgery outcom	me								
No residue	140 29.29 ((31.602)	3.96 (1.808)	72 2	(24.740)	6.58 (2.644)	-2.62 (-8.960,	3.719)	0.4159	
Hedges' g SMD							-0.12 (-0.405,		0.4082	
Residue	70 25.71 ((27.318)	6.34 (2.518)	40 2	5.00 (26.954)	6.43 (3.646)	,		0.9837	
Hedges' g SMD							0.00 (-0.393,	0.384)	0.9833	
Int. p-value									0.3963	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.5.4.15 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-C30 Single item symptom scale: Insomnia (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=25		Placebo + bev		Difference between g	roups
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
	eductive surgery	4 10 (1 000)	E1 00 4E (04 010)	0.00 (0.605)	4 00 / 10 450 0 000)	
Upfront	128 29.69 (32.733)	4.13 (1.876)	71 23.47 (24.813)	8.22 (2.625)	, , , , , , , , , , , , , , , , , , , ,	0.2077
Hedges' g SMD	00 05 61 (05 010)	6 01 (0 310)	41 10 51 (26 242)	0.00 / 0.556	-0.19 (-0.480, 0.102)	0.2022
Interval	82 25.61 (25.819)	6.01 (2.310)	41 19.51 (26.849)	2.99 (3.556)	, , , , , , , , , , , , , , , , , , , ,	0.4783
Hedges' g SMD					0.14 (-0.236, 0.515)	0.4667 0.1306
Int. p-value						0.1306
Myriad tumour BR	RCA mutation status					
tBRCAm	134 30.35 (30.715)	4.20 (1.807)	68 21.08 (23.679)	5.71 (2.696)	-1.52 (-7.944, 4.910)	0.6421
Hedges' g SMD					-0.07 (-0.363, 0.221)	0.6348
Non-tBRCAm	83 24.90 (29.376)	6.16 (2.347)	52 24.36 (27.309)	5.92 (3.148)	0.23 (-7.547, 8.013)	0.9527
Hedges' g SMD					0.01 (-0.336, 0.357)	0.9522
Int. p-value						0.8349
Status somatic E	BRCA mutations					
sBRCAm	18 25.93 (33.442)	5.28 (4.239)	6 11.11 (17.213)	18.59 (7.583)	-13.31 (-31.785, 5.163)	0.1467
Hedges' g SMD	(112)		()		-0.71 (-1.659, 0.240)	0.1432
gBRCAm	55 30.91 (29.989)	10.67 (2.874)	28 22.62 (20.394)	4.56 (4.363)	, , , , , , , , , , , , , , , , , , , ,	0.2476
Hedges' q SMD		,	, , , , , , , , , , , , , , , , , , , ,		0.28 (-0.181, 0.733)	0.2369
Non-BRCAm	36 25.00 (26.874)	0.60 (3.336)	22 21.21 (24.224)	3.99 (4.500)	, , , , , , , , , , , , , , , , , , , ,	0.5483
Hedges' g SMD			, , , , , , , , , , , , , , , , , , , ,		-0.16 (-0.695, 0.368)	0.5462
Int. p-value					21.7 (21.11.1, 31.500)	0.0431

SMD = standardised mean difference. * p<0.05.

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.6.1 PAOLA1: Summary of EORTC QLQ-OV28 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI	Olaparib + bevacizumab (N=255)	Baseline [a]	247	20.61	18.074	0.0	16.67	77.8
		Wk 12 (Day 85)	226	22.44	17.765	0.0	22.22	77.
		Wk 24 (Day 169)	201	22.29	18.533	0.0	16.67	77.
		Wk 36 (Day 253)	174	22.12	17.566	0.0	18.33	83.
		Wk 48 (Day 337)	175	22.05	18.992	0.0	16.67	94.
		Wk 60 (Day 421)	164	22.80	19.423	0.0	16.67	86.
		Wk 72 (Day 505)	158	21.30	17.913	0.0	16.67	77.
		Wk 84 (Day 589)	139	22.07	17.795	0.0	16.67	77.
		Wk 96 (Day 673)	136	23.95	19.575	0.0	22.22	77.
		Wk 108 (Day 757)	108	21.12	20.290	0.0	16.67	88.
		Wk 120 (Day 841)	1	11.11	NC	11.1	11.11	11.
		Wk 132 (Day 925)	1	11.11	NC	11.1	11.11	11.
		Wk 144 (Day 1009)	1	11.11	NC	11.1	11.11	11.
		Wk 156 (Day 1093)	1	11.11	NC	11.1	11.11	11.
		End of Treatment	129	25.10	20.922	0.0	22.22	94.
		30 day Follow-up	61	26.94	23.029	0.0	22.22	88.
	Placebo + bevacizumab (N=132)	Baseline [a]	125	20.04	16.776	0.0	16.67	77.
		Wk 12 (Day 85)	116	21.75	18.549	0.0	16.67	83.
		Wk 24 (Day 169)	104	24.05	19.420	0.0	16.67	72.
		Wk 36 (Day 253)	97	23.03	17.553	0.0	22.22	66.
		Wk 48 (Day 337)	86	23.71	18.746	0.0	22.22	88.
		Wk 60 (Day 421)	71	22.86	17.634	0.0	22.22	66.
		Wk 72 (Day 505)	68	21.26	17.253	0.0	16.67	61.
		Wk 84 (Day 589)	51	22.05	15.973	0.0	22.22	55.
		Wk 96 (Day 673)	41	25.20	18.092	0.0	22.22	72.
		Wk 108 (Day 757)	31	28.32	21.534	0.0	22.22	88.
		End of Treatment	69	26.80	20.441	0.0	22.22	72.
		30 day Follow-up	24	31.02	22.338	0.0	25.00	88.

Table 2.6.1 PAOLA1: Summary of EORTC QLQ-OV28 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-OV28 Symptom scale/items: Body image	Olaparib + bevacizumab (N=255)	Baseline [a]	245	38.23	31.667	0.0	33.33	100.
		Wk 12 (Day 85)	224	32.66	30.127	0.0	33.33	100.
		Wk 24 (Day 169)	198	30.81	26.802	0.0	33.33	100.
		Wk 36 (Day 253)	175	29.90	29.594	0.0	33.33	100.
		Wk 48 (Day 337)	174	29.02	26.681	0.0	33.33	100.
		Wk 60 (Day 421)	162	27.06	26.206	0.0	16.67	100.
		Wk 72 (Day 505)	156	28.10	27.747	0.0	33.33	100.
		Wk 84 (Day 589)	136	27.21	27.801	0.0	16.67	100.
		Wk 96 (Day 673)	134	28.61	28.658	0.0	33.33	100.
		Wk 108 (Day 757)	108	23.61	26.877	0.0	16.67	100.
		Wk 120 (Day 841)	1	0.00	NC	0.0	0.00	0.0
		Wk 132 (Day 925)	1	16.67	NC	16.7	16.67	16.
		Wk 144 (Day 1009)	1	0.00	NC	0.0	0.00	0.0
		Wk 156 (Day 1093)	1	0.00	NC	0.0	0.00	0.0
		End of Treatment	129	27.26	27.238	0.0	33.33	100
		30 day Follow-up	61	30.33	26.962	0.0	33.33	100
	Placebo + bevacizumab (N=132)	Baseline [a]	124	39.52	33.635	0.0	33.33	100
		Wk 12 (Day 85)	116	31.90	30.659	0.0	33.33	100
		Wk 24 (Day 169)	103	33.66	27.710	0.0	33.33	100
		Wk 36 (Day 253)	95	32.98	27.395	0.0	33.33	100
		Wk 48 (Day 337)	86	30.81	29.319	0.0	33.33	100
		Wk 60 (Day 421)	71	27.70	29.269	0.0	16.67	100
		Wk 72 (Day 505)	67	27.61	26.992	0.0	33.33	100
		Wk 84 (Day 589)	51	27.45	27.853	0.0	16.67	100
		Wk 96 (Day 673)	40	31.67	28.445	0.0	33.33	100
		Wk 108 (Day 757)	31	31.72	28.660	0.0	33.33	83.
		End of Treatment	69	34.06	28.787	0.0	33.33	100
		30 day Follow-up	23	31.16	29.432	0.0	33.33	100

Table 2.6.1 PAOLA1: Summary of EORTC QLQ-OV28 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects	Olaparib + bevacizumab (N=255)	Baseline [a]	247	25.75	18.713	0.0	26.67	88.9
		Wk 12 (Day 85)	226	25.16	18.157	0.0	20.00	93.3
		Wk 24 (Day 169)	200	24.84	17.435	0.0	20.00	86.7
		Wk 36 (Day 253)	173	23.54	15.963	0.0	20.00	100.
		Wk 48 (Day 337)	172	22.61	15.850	0.0	20.00	80.0
		Wk 60 (Day 421)	164	21.67	17.780	0.0	20.00	86.7
		Wk 72 (Day 505)	158	21.07	18.845	0.0	13.33	80.0
		Wk 84 (Day 589)	138	20.23	16.633	0.0	13.33	93.3
		Wk 96 (Day 673)	135	21.63	18.349	0.0	20.00	93.3
		Wk 108 (Day 757)	108	19.65	18.593	0.0	13.33	93.3
		Wk 120 (Day 841)	1	20.00	NC	20.0	20.00	20.
		Wk 132 (Day 925)	1	26.67	NC	26.7	26.67	26.
		Wk 144 (Day 1009)	1	20.00	NC	20.0	20.00	20.
		Wk 156 (Day 1093)	1	26.67	NC	26.7	26.67	26.
		End of Treatment	129	20.96	17.050	0.0	20.00	93.
		30 day Follow-up	61	20.33	16.619	0.0	20.00	66.
	Placebo + bevacizumab (N=132)	Baseline [a]	124	24.38	16.810	0.0	20.00	93.3
		Wk 12 (Day 85)	116	26.03	17.598	0.0	26.67	100
		Wk 24 (Day 169)	104	26.33	15.675	0.0	26.67	60.
		Wk 36 (Day 253)	96	24.53	16.073	0.0	23.33	66.
		Wk 48 (Day 337)	84	24.15	17.368	0.0	20.00	66.
		Wk 60 (Day 421)	71	21.22	16.169	0.0	20.00	100
		Wk 72 (Day 505)	68	18.70	16.207	0.0	20.00	66.
		Wk 84 (Day 589)	51	17.88	16.971	0.0	13.33	66.
		Wk 96 (Day 673)	41	20.61	16.439	0.0	20.00	60.
		Wk 108 (Day 757)	31	24.73	22.522	0.0	13.33	100
		End of Treatment	69	23.21	16.973	0.0	20.00	66.
		30 day Follow-up	23	24.93	22.761	0.0	20.00	100

Table 2.6.1 PAOLA1: Summary of EORTC QLQ-OV28 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment	Olaparib + bevacizumab (N=255)	Baseline [a]	243	56.88	27.626	0.0	55.56	100.
		Wk 12 (Day 85)	223	51.57	26.168	0.0	55.56	100.
		Wk 24 (Day 169)	197	48.67	24.111	0.0	44.44	100.
		Wk 36 (Day 253)	175	45.21	25.793	0.0	44.44	100.
		Wk 48 (Day 337)	173	43.93	25.468	0.0	44.44	100.
		Wk 60 (Day 421)	164	40.14	25.083	0.0	33.33	100.
		Wk 72 (Day 505)	155	38.85	24.336	0.0	33.33	100.
		Wk 84 (Day 589)	136	40.11	24.611	0.0	33.33	100.
		Wk 96 (Day 673)	136	38.11	25.945	0.0	33.33	100.
		Wk 108 (Day 757)	106	35.01	22.670	0.0	33.33	100.
		Wk 120 (Day 841)	1	11.11	NC	11.1	11.11	11.1
		Wk 132 (Day 925)	1	11.11	NC	11.1	11.11	11.1
		Wk 144 (Day 1009)	1	11.11	NC	11.1	11.11	11.
		Wk 156 (Day 1093)	1	22.22	NC	22.2	22.22	22.
		End of Treatment	128	43.79	25.355	0.0	33.33	100
		30 day Follow-up	59	40.87	24.711	0.0	33.33	88.9
	Placebo + bevacizumab (N=132)	Baseline [a]	124	56.94	26.926	0.0	55.56	100
		Wk 12 (Day 85)	114	48.20	28.188	0.0	44.44	100
		Wk 24 (Day 169)	103	47.57	26.503	0.0	44.44	100
		Wk 36 (Day 253)	94	45.33	26.516	0.0	44.44	100
		Wk 48 (Day 337)	84	39.81	25.884	0.0	33.33	100
		Wk 60 (Day 421)	69	40.66	25.786	0.0	33.33	100
		Wk 72 (Day 505)	67	36.98	28.795	0.0	33.33	100
		Wk 84 (Day 589)	51	37.69	28.465	0.0	33.33	100
		Wk 96 (Day 673)	40	42.50	29.115	0.0	38.89	100
		Wk 108 (Day 757)	31	43.73	29.805	0.0	44.44	100
		End of Treatment	67	47.35	23.676	0.0	44.44	88.
		30 day Follow-up	23	49.28	29.555	0.0	55.56	100

Table 2.6.1 PAOLA1: Summary of EORTC QLQ-OV28 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-OV28 Symptom scale/items: Hormonal	Olaparib + bevacizumab (N=255)	Baseline [a]	246	26.36	29.812	0.0	16.67	100.
		Wk 12 (Day 85)	225	20.52	26.586	0.0	16.67	100.
		Wk 24 (Day 169)	198	20.12	26.192	0.0	16.67	100.
		Wk 36 (Day 253)	175	22.67	27.806	0.0	16.67	100.
		Wk 48 (Day 337)	173	21.19	26.237	0.0	16.67	100.
		Wk 60 (Day 421)	164	22.76	27.878	0.0	16.67	100.
		Wk 72 (Day 505)	158	25.00	30.200	0.0	16.67	100.
		Wk 84 (Day 589)	137	27.49	30.154	0.0	16.67	100.
		Wk 96 (Day 673)	136	25.49	28.899	0.0	16.67	100.
		Wk 108 (Day 757)	108	25.15	27.953	0.0	16.67	100
		Wk 120 (Day 841)	1	16.67	NC	16.7	16.67	16.7
		Wk 132 (Day 925)	1	16.67	NC	16.7	16.67	16.
		Wk 144 (Day 1009)	1	33.33	NC	33.3	33.33	33.
		Wk 156 (Day 1093)	1	16.67	NC	16.7	16.67	16.
		End of Treatment	129	23.26	26.551	0.0	16.67	100
		30 day Follow-up	61	21.04	25.802	0.0	16.67	100
	Placebo + bevacizumab (N=132)	Baseline [a]	123	26.42	31.502	0.0	16.67	100
		Wk 12 (Day 85)	115	28.99	30.033	0.0	16.67	100
		Wk 24 (Day 169)	103	24.76	27.600	0.0	16.67	100
		Wk 36 (Day 253)	97	22.34	27.731	0.0	16.67	100
		Wk 48 (Day 337)	84	25.00	26.447	0.0	16.67	100
		Wk 60 (Day 421)	71	23.94	26.537	0.0	16.67	100
		Wk 72 (Day 505)	68	25.49	28.863	0.0	16.67	100
		Wk 84 (Day 589)	51	22.88	26.449	0.0	16.67	100
		Wk 96 (Day 673)	41	24.39	26.899	0.0	16.67	100
		Wk 108 (Day 757)	31	28.49	26.941	0.0	33.33	100
		End of Treatment	69	26.09	27.338	0.0	16.67	100
		30 day Follow-up	23	19.57	21.113	0.0	16.67	66.

Table 2.6.1 PAOLA1: Summary of EORTC QLQ-OV28 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy	Olaparib + bevacizumab (N=255)	Baseline [a]	247	42.58	34.808	0.0	33.33	100.
		Wk 12 (Day 85)	227	34.51	31.217	0.0	33.33	100.
		Wk 24 (Day 169)	200	28.08	28.920	0.0	16.67	100.
		Wk 36 (Day 253)	173	28.90	30.599	0.0	33.33	100.
		Wk 48 (Day 337)	173	26.69	28.001	0.0	16.67	100
		Wk 60 (Day 421)	164	27.74	27.954	0.0	33.33	100
		Wk 72 (Day 505)	158	25.32	27.629	0.0	16.67	100
		Wk 84 (Day 589)	139	24.82	29.717	0.0	16.67	100
		Wk 96 (Day 673)	135	26.05	29.829	0.0	16.67	100
		Wk 108 (Day 757)	108	25.31	28.229	0.0	16.67	100
		Wk 120 (Day 841)	1	33.33	NC	33.3	33.33	33.
		Wk 132 (Day 925)	1	33.33	NC	33.3	33.33	33.
		Wk 144 (Day 1009)	1	33.33	NC	33.3	33.33	33.
		Wk 156 (Day 1093)	1	50.00	NC	50.0	50.00	50.
		End of Treatment	129	29.59	29.773	0.0	16.67	100
		30 day Follow-up	61	29.78	30.141	0.0	33.33	100
	Placebo + bevacizumab (N=132)	Baseline [a]	124	45.43	35.201	0.0	33.33	100
		Wk 12 (Day 85)	116	35.06	34.219	0.0	33.33	100
		Wk 24 (Day 169)	104	31.41	30.130	0.0	33.33	100
		Wk 36 (Day 253)	96	26.04	28.899	0.0	16.67	100
		Wk 48 (Day 337)	84	25.99	28.146	0.0	16.67	100
		Wk 60 (Day 421)	71	25.82	28.283	0.0	16.67	100
		Wk 72 (Day 505)	68	23.77	28.260	0.0	16.67	100
		Wk 84 (Day 589)	51	25.82	28.346	0.0	16.67	100
		Wk 96 (Day 673)	41	26.42	30.726	0.0	16.67	100
		Wk 108 (Day 757)	31	25.27	28.826	0.0	16.67	100
		End of Treatment	69	24.40	25.176	0.0	16.67	100
		30 day Follow-up	23	19.57	22.837	0.0	16.67	66.

Table 2.6.1 PAOLA1: Summary of EORTC QLQ-OV28 results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EORTC QLQ-OV28 Symptom scale/items: Other single items	Olaparib + bevacizumab (N=255)	Baseline [a]	246	24.33	23.007	0.0	22.22	100.
		Wk 12 (Day 85)	226	17.15	18.534	0.0	11.11	100.
		Wk 24 (Day 169)	200	14.36	15.377	0.0	11.11	66.7
		Wk 36 (Day 253)	174	14.94	16.385	0.0	11.11	83.3
		Wk 48 (Day 337)	175	13.90	14.002	0.0	11.11	58.3
		Wk 60 (Day 421)	164	15.55	16.269	0.0	11.11	88.9
		Wk 72 (Day 505)	158	15.72	16.148	0.0	11.11	66.7
		Wk 84 (Day 589)	138	14.88	15.645	0.0	11.11	58.3
		Wk 96 (Day 673)	136	15.48	14.894	0.0	11.11	66.7
		Wk 108 (Day 757)	108	13.45	14.822	0.0	11.11	66.7
		Wk 120 (Day 841)	1	0.00	NC	0.0	0.00	0.0
		Wk 132 (Day 925)	1	0.00	NC	0.0	0.00	0.0
		Wk 144 (Day 1009)	1	0.00	NC	0.0	0.00	0.0
		Wk 156 (Day 1093)	1	0.00	NC	0.0	0.00	0.0
		End of Treatment	129	16.04	16.530	0.0	11.11	83.3
		30 day Follow-up	61	17.30	19.625	0.0	11.11	66.7
	Placebo + bevacizumab (N=132)	Baseline [a]	125	28.07	25.375	0.0	25.00	75.0
		Wk 12 (Day 85)	116	11.35	17.090	0.0	0.00	91.7
		Wk 24 (Day 169)	104	10.31	16.163	0.0	4.17	83.3
		Wk 36 (Day 253)	97	8.33	10.261	0.0	0.00	33.3
		Wk 48 (Day 337)	86	8.33	11.770	0.0	0.00	50.0
		Wk 60 (Day 421)	71	10.52	14.077	0.0	11.11	66.7
		Wk 72 (Day 505)	68	12.17	15.305	0.0	11.11	55.6
		Wk 84 (Day 589)	51	12.58	15.834	0.0	8.33	58.3
		Wk 96 (Day 673)	41	11.38	16.135	0.0	0.00	58.3
		Wk 108 (Day 757)	31	12.90	15.901	0.0	8.33	50.0
		End of Treatment	69	15.30	19.196	0.0	11.11	83.3
		30 day Follow-up	24	13.54	18.558	0.0	11.11	75.0

Table 2.6.2.1 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bev (N=255		Placebo + bev		Difference between	n groups
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Wk 12 (Day 85)	206 19.79 (17.735)	2.97 (0.995)	111 20.41 (16.944)	1.34 (1.354)	1.63 (-1.676, 4.933	0.3332
Wk 24 (Day 169)	187 19.66 (17.793)	2.28 (1.053)	100 19.50 (16.047)	4.71 (1.446)	-2.42 (-5.944, 1.096	0.1763
Wk 36 (Day 253)	162 18.50 (16.559)	3.23 (1.082)	92 20.23 (16.329)	4.10 (1.462)	-0.87 (-4.454, 2.70	0.6315
Wk 48 (Day 337)	165 19.07 (17.642)	2.96 (1.200)	77 20.13 (16.235)	4.74 (1.727)	-1.78 (-5.919, 2.358	0.3979
Wk 60 (Day 421)	155 19.54 (17.879)	3.01 (1.168)	67 20.15 (17.200)	6.26 (1.730)	-3.25 (-7.356, 0.869	0.1212
Wk 72 (Day 505)	149 18.81 (17.292)	1.95 (1.226)	61 19.03 (15.514)	2.72 (1.840)	-0.76 (-5.118, 3.588	0.7297
Wk 84 (Day 589)	133 18.92 (17.241)	2.48 (1.152)	47 19.05 (16.476)	3.30 (1.824)	-0.82 (-5.073, 3.425	0.7027
Wk 96 (Day 673)	129 19.12 (17.888)	4.96 (1.363)	36 21.02 (17.243)	4.97 (2.303)	-0.01 (-5.282, 5.264	0.9973
Average over all visits	221 20.32 (17.983)	2.98 (0.856)	119 19.88 (16.791)	4.02 (1.228)	-1.04 (-3.981, 1.90	0.4889
Hedges' g SMD					-0.08 (-0.303, 0.143	0.4828

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used model within-subject error.

SMD = standardised mean difference. * p<0.05.

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.2.2 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Body image (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=25		Placebo + be		Difference between	groups
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Wk 12 (Day 85)	202 37.95 (31.686)	-5.58 (1.551)	110 38.64 (33.708)	-7.39 (2.100)	1.82 (-3.317, 6.952)	0.4867
Wk 24 (Day 169)	183 38.98 (32.097)	-7.53 (1.606)	98 40.48 (33.675)	-5.49 (2.196)	-2.04 (-7.392, 3.316)	0.4544
Wk 36 (Day 253)	161 37.89 (31.677)	-8.26 (1.852)	89 41.76 (34.645)	-6.37 (2.528)	-1.89 (-8.065, 4.278)	0.5464
Wk 48 (Day 337)	163 38.65 (32.326)	-8.65 (1.798)	76 41.67 (34.801)	-8.37 (2.600)	-0.28 (-6.505, 5.945)	0.9295
Wk 60 (Day 421)	151 39.18 (32.444)	-11.64 (1.785)	66 41.92 (35.233)	-10.17 (2.635)	-1.47 (-7.738, 4.798)	0.6447
Wk 72 (Day 505)	145 39.43 (32.620)	-10.31 (1.865)	60 44.17 (35.498)	-9.64 (2.830)	-0.66 (-7.338, 6.015)	0.8454
Wk 84 (Day 589)	129 38.11 (32.422)	-12.23 (1.961)	46 39.86 (34.868)	-12.05 (3.062)	-0.18 (-7.345, 6.981)	0.9602
Wk 96 (Day 673)	126 39.81 (32.453)	-8.76 (2.145)	35 44.29 (35.226)	-7.51 (3.635)	-1.25 (-9.563, 7.070)	0.7680
Average over all visits	219 38.96 (31.546)	-9.12 (1.363)	118 38.98 (33.633)	-8.38 (1.953)	-0.74 (-5.430, 3.941)	0.7549
Hedges' g SMD					-0.04 (-0.260, 0.188)	0.7516

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.2.3 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=255		Placebo + bev		Difference between	groups
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Wk 12 (Day 85)	206 24.82 (18.058)	0.57 (1.041)	110 23.97 (17.213)	1.72 (1.423)	-1.15 (-4.614, 2.324)	0.5164
Wk 24 (Day 169)	186 25.09 (18.589)	-0.13 (1.009)	99 23.86 (17.215)	1.72 (1.385)	-1.85 (-5.222, 1.522)	0.2812
Wk 36 (Day 253)	161 24.68 (17.579)	-1.28 (1.063)	91 23.68 (17.346)	0.66 (1.435)	-1.94 (-5.455, 1.575)	0.2783
Wk 48 (Day 337)	162 25.63 (18.900)	-2.38 (1.070)	75 23.78 (16.944)	-1.08 (1.549)	-1.30 (-5.004, 2.411)	0.4918
Wk 60 (Day 421)	155 25.30 (18.437)	-3.76 (1.086)	66 23.16 (17.052)	-1.82 (1.631)	-1.94 (-5.806, 1.917)	0.3223
Wk 72 (Day 505)	149 25.63 (18.868)	-4.25 (1.205)	60 24.14 (17.975)	-5.93 (1.833)	1.68 (-2.644, 5.998)	0.4455
Wk 84 (Day 589)	132 24.96 (18.853)	-4.70 (1.137)	46 24.38 (18.676)	-6.38 (1.835)	1.68 (-2.569, 5.937)	0.4362
Wk 96 (Day 673)	128 25.08 (18.329)	-3.64 (1.225)	36 26.53 (18.569)	-4.59 (2.085)	0.95 (-3.812, 5.714)	0.6943
Average over all visits	221 25.55 (18.833)	-2.44 (0.794)	118 24.15 (16.945)	-1.96 (1.147)	-0.48 (-3.229, 2.263)	0.7295
Hedges' g SMD					-0.04 (-0.264, 0.183)	0.7255

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.2.4 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=25		Placebo + be		Difference b	etween g	roups
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE)	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from	Estimated difference (95	% CI)	p-value
Wk 12 (Day 85)	199 56.03 (27.443)	-5.30 (1.520)	108 56.22 (26.979)	-9.12 (2.059)	3.82 (-1.218,	8.851)	0.1368
Wk 24 (Day 169)	182 55.74 (27.608)	-7.98 (1.491)	98 56.52 (27.264)	-9.22 (2.029)	1.24 (-3.713,	6.198)	0.6220
Wk 36 (Day 253)	160 55.35 (26.864)	-10.21 (1.702)	88 56.31 (26.006)	-10.55 (2.311)	0.34 (-5.312,	5.987)	0.9065
Wk 48 (Day 337)	160 57.19 (27.638)	-12.04 (1.588)	74 55.03 (25.800)	-16.04 (2.295)	4.00 (-1.491,	9.497)	0.1526
Wk 60 (Day 421)	153 55.63 (28.003)	-15.19 (1.693)	64 54.60 (26.349)	-13.61 (2.512)	-1.58 (-7.547,	4.379)	0.6015
Wk 72 (Day 505)	143 56.02 (28.378)	-17.46 (1.730)	60 55.37 (27.522)	-17.36 (2.627)	-0.10 (-6.296,	6.093)	0.9743
Wk 84 (Day 589)	128 57.20 (27.654)	-15.99 (1.784)	46 53.38 (27.999)	-17.61 (2.874)	1.63 (-5.044,	8.296)	0.6314
Wk 96 (Day 673)	128 57.38 (27.146)	-17.11 (2.029)	35 57.62 (29.058)	-12.96 (3.523)	-4.15 (-12.161,	3.859)	0.3082
Average over all visits	217 56.37 (27.356)	-12.66 (1.203)	118 56.64 (26.927)	-13.31 (1.736)	0.65 (-3.506,	4.803)	0.7589
Hedges' g SMD					0.04 (-0.188,	0.260)	0.7549

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.2.5 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Hormonal (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		vacizumab	Placebo + bev		Difference between groups		
Timepoint		Mean (SE) Change from	n Mean (SD) at	Mean (SE)	Estimated	p-value	
Wk 12 (Day 85)	204 24.92 (29.150)	-4.41 (1.387)	108 26.39 (30.240)	3.81 (1.904)	-8.21 (-12.849, -3.581)	0.0006*	
Wk 24 (Day 169)	184 26.00 (30.462)	-5.32 (1.348)	97 24.74 (30.153)	-0.48 (1.866)	-4.84 (-9.374, -0.315)	0.0361*	
Wk 36 (Day 253)	162 26.13 (30.118)	-3.39 (1.745)	91 24.54 (29.953)	-2.05 (2.380)	-1.35 (-7.153, 4.461)	0.6486	
Wk 48 (Day 337)	163 26.69 (29.834)	-4.15 (1.562)	75 26.22 (30.777)	-1.16 (2.257)	-2.99 (-8.392, 2.413)	0.2770	
Wk 60 (Day 421)	154 27.16 (30.247)	-3.10 (1.701)	66 25.00 (32.324)	-1.06 (2.521)	-2.04 (-8.030, 3.947)	0.5027	
Wk 72 (Day 505)	148 26.80 (29.587)	-1.17 (1.892)	60 23.33 (31.173)	1.23 (2.858)	-2.40 (-9.152, 4.348)	0.4842	
Wk 84 (Day 589)	131 27.99 (30.119)	0.70 (1.843)	46 25.72 (33.093)	-2.01 (2.929)	2.71 (-4.109, 9.524)	0.4348	
Wk 96 (Day 673)	128 27.21 (29.937)	-1.42 (1.872)	35 30.00 (34.490)	-1.49 (3.213)	0.08 (-7.246, 7.397)	0.9838	
Average over all visits Hedges' g SMD	220 25.68 (29.768)	-2.78 (1.234)	117 26.50 (31.115)	-0.40 (1.777)	-2.38 (-6.637, 1.873) -0.13 (-0.352, 0.097)	0.2716 0.2652	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.2.6 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=25		Placebo + be		Difference between groups		
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Wk 12 (Day 85)	207 42.19 (34.282)	-7.34 (1.733)	110 43.03 (34.341)	-6.60 (2.373)	-0.74 (-6.521, 5.039)	0.8010	
Wk 24 (Day 169)	186 40.77 (34.658)	-13.12 (1.699)	99 42.93 (34.511)	-10.48 (2.336)	-2.64 (-8.321, 3.048)	0.3621	
Wk 36 (Day 253)	161 39.03 (34.666)	-12.50 (1.801)	91 44.14 (34.728)	-16.19 (2.447)	3.69 (-2.296, 9.670)	0.2262	
Wk 48 (Day 337)	163 40.49 (34.988)	-14.26 (1.672)	75 46.44 (35.227)	-17.74 (2.399)	3.48 (-2.277, 9.238)	0.2352	
Wk 60 (Day 421)	155 41.08 (35.446)	-14.18 (1.782)	66 45.96 (35.211)	-15.84 (2.664)	1.66 (-4.653, 7.973)	0.6051	
Wk 72 (Day 505)	149 41.05 (34.141)	-16.23 (1.794)	60 43.33 (34.608)	-18.94 (2.733)	2.71 (-3.730, 9.149)	0.4081	
Wk 84 (Day 589)	133 39.60 (33.685)	-15.88 (1.978)	46 43.12 (33.440)	-15.96 (3.170)	0.08 (-7.280, 7.444)	0.9825	
Wk 96 (Day 673)	128 38.93 (34.001)	-13.55 (1.990)	36 46.76 (35.146)	-15.92 (3.382)	2.37 (-5.369, 10.108)	0.5469	
Average over all visits	221 41.78 (34.474)	-13.38 (1.345)	118 44.63 (35.263)	-14.71 (1.933)	1.33 (-3.308, 5.961)	0.5738	
Hedges' g SMD					0.07 (-0.158, 0.289)	0.5686	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.6.2.7 PAOLA1: Summary of analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Other single items (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

				Difference between groups		
n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
205 23.50 (22.428)	-6.88 (1.148)	111 27.15 (25.702)	-13.56 (1.562)	6.68 (2.866, 10.500)	0.0006*	
185 23.06 (22.219)	-10.18 (1.048)	100 26.94 (25.518)	-14.61 (1.428)	4.43 (0.938, 7.917)	0.0131*	
161 23.24 (22.189)	-10.11 (0.991)	92 26.12 (24.297)	-15.61 (1.327)	5.50 (2.239, 8.766)	0.0010*	
164 23.51 (22.465)	-10.27 (0.968)	77 25.94 (24.560)	-15.56 (1.390)	5.29 (1.955, 8.626)	0.0020*	
154 22.40 (22.108)	-8.65 (1.141)	67 25.70 (24.558)	-12.72 (1.697)	4.07 (0.038, 8.100)	0.0479*	
148 24.27 (22.621)	-9.07 (1.192)	61 26.41 (25.868)	-11.49 (1.835)	2.42 (-1.891, 6.732)	0.2699	
131 23.09 (22.664)	-9.84 (1.273)	47 25.18 (25.983)	-10.20 (2.063)	0.37 (-4.416, 5.147)	0.8804	
128 21.40 (20.881)	-8.87 (1.194)	36 30.40 (27.021)	-13.14 (2.147)	4.27 (-0.602, 9.133)	0.0855	
220 24.04 (22.468)	-9.23 (0.745)	119 28.01 (25.527)	-13.36 (1.084)	, , ,	0.0019*	
	n Mean (SD) at [a] Baseline [b] 205 23.50 (22.428) 185 23.06 (22.219) 161 23.24 (22.189) 164 23.51 (22.465) 154 22.40 (22.108) 148 24.27 (22.621) 131 23.09 (22.664) 128 21.40 (20.881)	n Mean (SD) at Change from baseline 205 23.50 (22.428) -6.88 (1.148) 185 23.06 (22.219) -10.18 (1.048) 161 23.24 (22.189) -10.11 (0.991) 164 23.51 (22.465) -10.27 (0.968) 154 22.40 (22.108) -8.65 (1.141) 148 24.27 (22.621) -9.07 (1.192) 131 23.09 (22.664) -9.84 (1.273) 128 21.40 (20.881) -8.87 (1.194)	N=255 Mean (SE) N Mean (SD) at Easeline [b] Change from baseline [a] Baseline [b] N Mean (SE) N Mean (SD) at Easeline [b] N Mean (SE) N Mean (SD) at Easeline [b] N Me	Nean (SD) at Change from baseline Nean (SE) Change from baseline Nean (SD) at baseline Nean (SE) Change from baseline Nean (SD) at baseline Nean (SE) Change from baseline Nean (SD) at baseline Nean (SE) Change from baseline Nean (SD) at baseline Nean (SE) Nean (N=132 Mean (SD) at Mean (SE) Change from Nean (SE) Nean (SE) Change from Nean (SD) at Nean (SE) Nean (SE)	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Figure 2.6.3.1 PAOLA1: Mean (+/- SD) score for EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

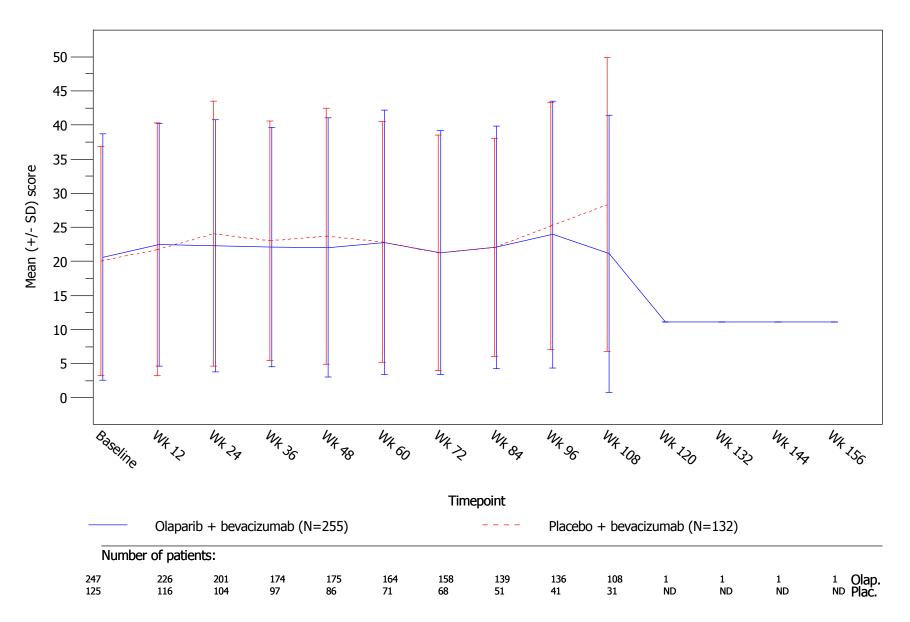


Figure 2.6.3.2 PAOLA1: Mean (+/- SD) score for EORTC QLQ-OV28 Symptom scale/items: Body image across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

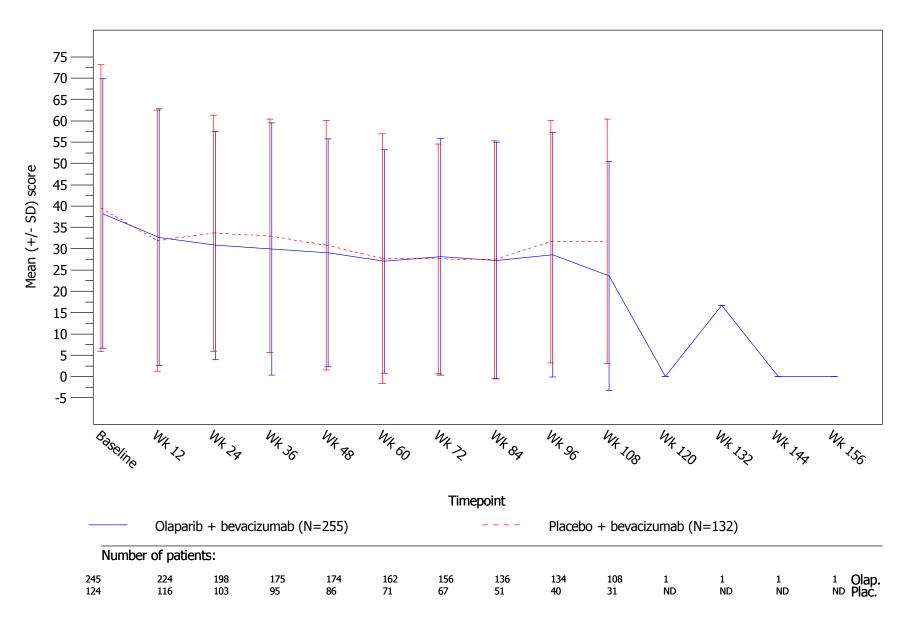


Figure 2.6.3.3 PAOLA1: Mean (+/- SD) score for EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

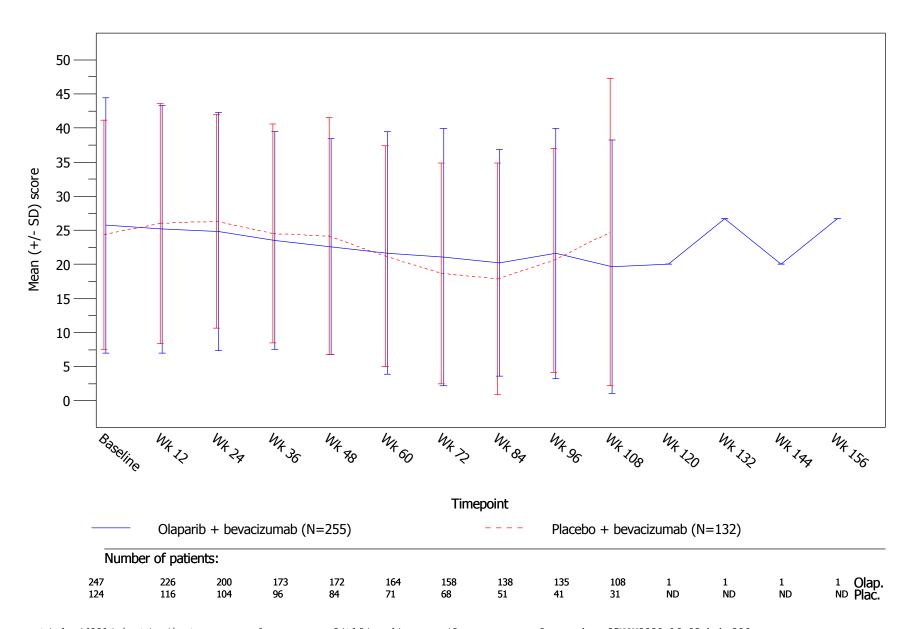


Figure 2.6.3.4 PAOLA1: Mean (+/- SD) score for EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

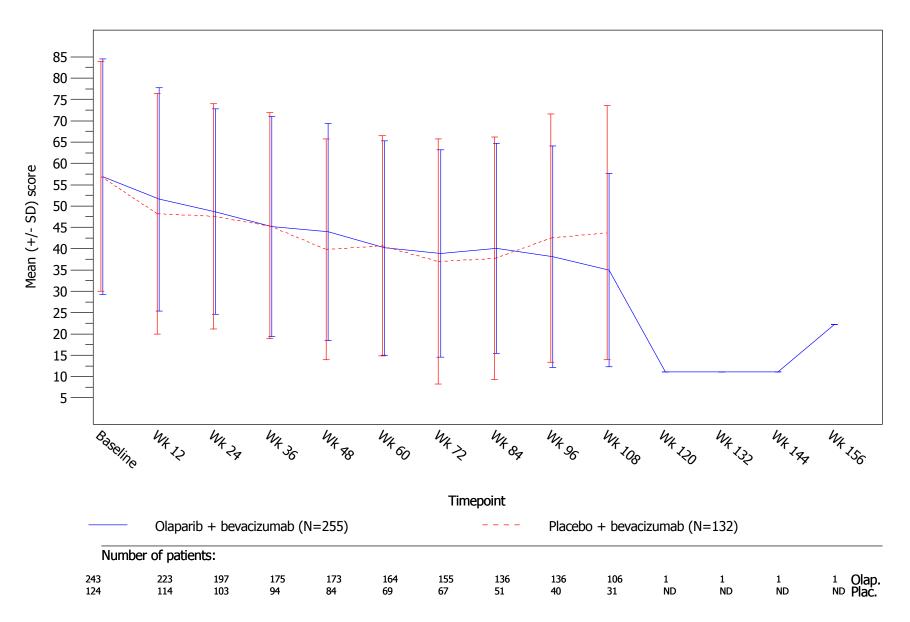


Figure 2.6.3.5 PAOLA1: Mean (+/- SD) score for EORTC QLQ-OV28 Symptom scale/items: Hormonal across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

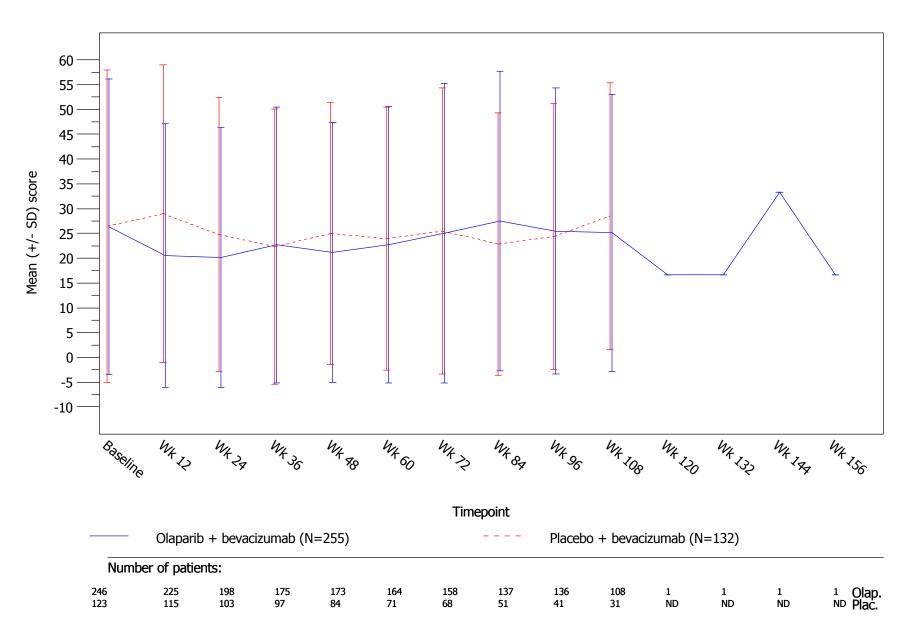


Figure 2.6.3.6 PAOLA1: Mean (+/- SD) score for EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

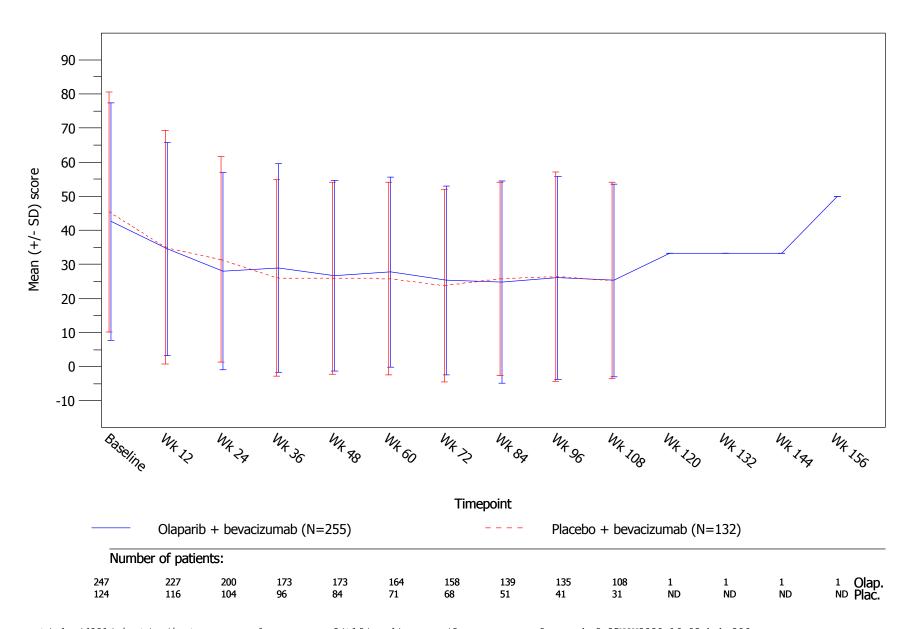


Figure 2.6.3.7 PAOLA1: Mean (+/- SD) score for EORTC QLQ-OV28 Symptom scale/items: Other single items across timepoints, by treatment group

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

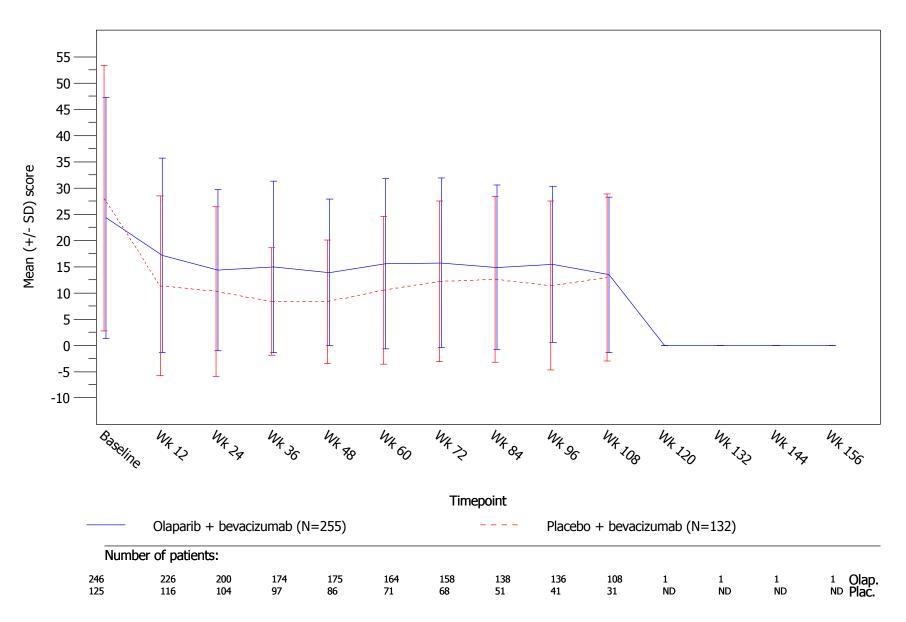


Table 2.6.4.1 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bev		Placebo + bev (N=132		Difference between groups		
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
First line treat	ment outcome (IVRS)						
NED [PDS]	82 19.00 (17.543)	3.54 (1.303)	42 22.30 (15.047)	6.19 (1.909)	-2.65 (-7.233, 1.932)	0.2545	
Hedges' g SMD					-0.22 (-0.593, 0.153)	0.2483	
NED/CR [IDS]	61 22.19 (18.596)	1.08 (1.718)	34 16.34 (16.193)	1.50 (2.438)	-0.41 (-6.376, 5.547)	0.8903	
Hedges' g SMD					-0.03 (-0.449, 0.389)	0.8885	
NED/CR [Chemo]	37 25.78 (21.435)	0.65 (2.351)	20 21.94 (16.664)	3.56 (3.503)	-2.92 (-11.436, 5.600)	0.4932	
Hedges' g SMD					-0.19 (-0.740, 0.351)	0.4841	
PR	41 15.28 (12.678)	6.82 (1.960)	23 18.89 (20.534)	1.89 (2.789)	4.93 (-1.914, 11.769)	0.1543	
Hedges' g SMD					0.38 (-0.136, 0.894)	0.1494	
Int. p-value						0.2893	
Screening labora	tory tBRCA status (IV	TRS)					
tBRCAm	131 22.21 (18.743)	2.61 (1.130)	58 17.95 (15.334)	2.42 (1.777)	0.19 (-3.973, 4.354)	0.9282	
Hedges' g SMD					0.01 (-0.295, 0.324)	0.9270	
non-tBRCAm	90 17.57 (16.532)	3.54 (1.331)	61 21.71 (18.003)	5.39 (1.710)	-1.85 (-6.149, 2.447)	0.3957	
Hedges' g SMD					-0.14 (-0.468, 0.183)	0.3908	
Int. p-value						0.8051	
First line treat	ment outcome (eCRF)						
NED [PDS]	80 20.83 (19.242)	2.83 (1.376)	41 22.38 (15.040)	4.88 (1.979)	-2.05 (-6.827, 2.724)	0.3966	
Hedges' g SMD	(======,		,	, , , , , , , , , , , , , , , , , , , ,	-0.16 (-0.541, 0.213)	0.3939	
NED/CR [IDS]	60 18.76 (15.845)	2.89 (1.643)	29 15.86 (16.354)	1.66 (2.524)	1.22 (-4.778, 7.225)	0.6860	
Hedges' g SMD	(2000)		(()	,	0.09 (-0.350, 0.537)	0.6803	
NED/CR [Chemo]	34 23.97 (19.742)	1.83 (2.486)	17 23.20 (22.069)	4.07 (4.060)	-2.24 (-11.911, 7.425)	0.6413	
Hedges' g SMD	(, 12)			, , , , , , , , , , , , , , , , , , , ,	-0.14 (-0.728, 0.438)	0.6267	
PR	44 18.66 (17.551)	3.74 (1.773)	31 17.89 (15.855)	3.90 (2.271)	-0.17 (-5.924, 5.591)	0.9541	
Hedges' g SMD		(7)		- / (/-)	-0.01 (-0.473, 0.446)	0.9539	
Int. p-value					212, 01110,	0.5023	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.4.1 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Ola	parib + bev (N=255			Placebo + bev (N=132		Difference between groups		
Subgroup		(SD) at line [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
Screening labor	atory tBRCA	status (e0	CRF)						
tBRCAm Hedges' g SMD	128 22.01	(18.878)	2.79 (1.140)	59	17.93 (15.202)	2.45 (1.757)	0.34 (-3.797, 0.03 (-0.282,	•	0.8698 0.8677
non-tBRCAm Hedges' g SMD Int. p-value	93 18.00	(16.490)	3.20 (1.309)	60 :	21.80 (18.143)	5.39 (1.723)	-2.19 (-6.474, -0.17 (-0.494,	,	0.3154 0.3099 0.6610
Age group	162 20.65	(18.012)	3.00 (1.020)	89 :	21.70 (15.707)	3.89 (1.449)	-0.89 (-4.380,	2.605)	0.6172
Hedges' g SMD >=65 years		(10.012)	NC	30	NC	NC	-0.07 (-0.326,	•	0.6124 NC
Int. p-value	39 1	NC .	INC	30	NC	NC .	NC NC		NC
FIGO Stage (Dis	ease state)								
III Hedges' g SMD	158 21.50	(18.996)	2.41 (1.051)	83	21.61 (18.123)	3.32 (1.512)	-0.92 (-4.543, -0.07 (-0.334,	•	0.6197 0.6158
IV	63 17.38	(14.881)	4.51 (1.477)	36	15.90 (12.556)	6.16 (2.219)	-1.65 (-6.946,	3.647)	0.5377
Hedges' g SMD Int. p-value							-0.13 (-0.543,	0.277)	0.5253 0.9716
Region									
Europe Hedges' g SMD	212 20.69	(18.108)	2.90 (0.885)	113	20.34 (17.028)	4.24 (1.284)	-1.33 (-4.402, -0.10 (-0.330,	•	0.3928 0.3851
Japan Int. p-value	9 1	NC	NC	6	NC	NC	NC	,	NC NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.

SMD = standardised mean difference. * p<0.05.

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.1 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (mixed model for repeated measures) - average over all visits Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be			Placebo + bev (N=132		Difference between groups		
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
ECOG performance	e stat	us at Baseline							
(0) Normal	163	19.53 (17.461)	4.28 (1.007)	91	19.39 (17.005)	4.62 (1.388)	-0.34 (-3.716,	3.037)	0.8432
activity									
Hedges' g SMD	5 4	02 10 (10 600)	1 10 / 1 654)	0.0	01 45 (16 054)	1 05 / 0 600)	-0.03 (-0.283,	,	0.8422
(1) Restricted activity	54	23.10 (19.692)	-1.17 (1.654)	28	21.47 (16.274)	1.87 (2.608)	-3.04 (-9.184,	3.113)	0.3286
Hedges' g SMD							-0.24 (-0.694,	0.222)	0.3122
Int. p-value								**,	0.9793
Baseline CA-125 <=ULN Hedges' g SMD >ULN Int. p-value		20.01 (18.202) NC	2.72 (0.897) NC	106 13	19.99 (17.054) NC	4.01 (1.288) NC	-1.29 (-4.381, -0.10 (-0.337, NC		0.4113 0.4045 NC NC
Histological gra	ade								
High grade		20.32 (17.983)	2.98 (0.856)	119	19.88 (16.791)	4.02 (1.228)	-1.04 (-3.981,	1.907)	0.4889
Hedges' g SMD Int. p-value							-0.08 (-0.303,	0.143)	0.4828 ID
Cytoreductive s	urgery	outcome							
No residue		19.97 (17.691)	2.83 (1.057)	71	19.95 (15.867)	4.20 (1.565)	-1.37 (-5.097,	2.351)	0.4680
Hedges' g SMD							-0.11 (-0.391,	0.178)	0.4627
Residue	71	21.44 (19.298)	2.85 (1.570)	40	19.94 (17.918)	3.24 (2.230)	-0.38 (-5.799,	5.037)	0.8892
Hedges' g SMD							-0.03 (-0.415,	0.360)	0.8876
Int. p-value									0.4787

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

> model within-subject error. SMD = standardised mean difference. * p<0.05.

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.1 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be			Placebo + bev		Difference between groups		
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
Timing of cytore	educti	ve surgery							
Upfront	131	20.54 (18.855)	3.63 (1.082)	70	21.16 (14.958)	4.86 (1.531)	-1.23 (-4.932,	2.467)	0.5120
Hedges' g SMD							-0.10 (-0.388,	0.192)	0.5085
Interval	83	20.32 (17.250)	1.80 (1.480)	41	17.89 (18.993)	1.89 (2.257)	-0.09 (-5.441,	5.267)	0.9743
Hedges' g SMD							-0.01 (-0.380,	0.368)	0.9738
Int. p-value									0.2218
Myriad tumour Bl	RCA mu	tation status							
tBRCAm	137	22.03 (18.847)	1.89 (1.080)	67	19.10 (16.246)	2.05 (1.621)	-0.16 (-4.006,	3.686)	0.9346
Hedges' g SMD							-0.01 (-0.305,	0.280)	0.9336
Non-tBRCAm	84	17.55 (16.204)	4.88 (1.403)	52	20.88 (17.577)	6.28 (1.873)	-1.40 (-6.041,	3.239)	0.5511
Hedges' g SMD							-0.11 (-0.452,	0.240)	0.5473
Int. p-value									0.7304
Status somatic 1	BRCA m	utations							
sBRCAm	18	17.10 (18.290)	-3.13 (2.384)	6	25.74 (13.433)	3.08 (4.181)	-6.20 (-16.338,	3.931)	0.2161
Hedges' g SMD							-0.59 (-1.532,	0.351)	0.2191
gBRCAm	57	21.07 (16.314)	4.73 (1.542)	27	18.44 (16.447)	1.74 (2.455)	3.00 (-2.778,	8.771)	0.3048
Hedges' g SMD							0.25 (-0.212,	0.707)	0.2917
Non-BRCAm	37	15.98 (13.350)	6.08 (1.783)	22	17.93 (17.563)	10.63 (2.485)	-4.55 (-10.695,	1.586)	0.1429
Hedges' g SMD							-0.40 (-0.936,	0.130)	0.1384
Int. p-value									0.1879

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.6.4.2 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Body image (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		bevacizumab 255)	Placebo + be (N=1)		Difference between groups		
Subgroup	n Mean (SD) at		n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
First line treat	ment outcome (IVRS)					
NED [PDS]	81 40.33 (30.942) -12.32 (2.181)	42 42.46 (32.552)	-5.14 (3.140)	-7.18 (-14.757, 0.400)	0.0632	
Hedges' g SMD					-0.36 (-0.734, 0.017)	0.0610	
NED/CR [IDS]	61 41.53 (29.443) -8.13 (2.784)	34 37.75 (37.674)	-11.64 (3.987)	3.51 (-6.163, 13.184)	0.4724	
Hedges' g SMD					0.16 (-0.264, 0.576)	0.4661	
NED/CR [Chemo]	36 42.59 (35.061) -9.29 (3.170)	20 28.33 (30.156)	-7.31 (4.627)	-1.98 (-13.387, 9.430)	0.7281	
Hedges' g SMD					-0.10 (-0.646, 0.448)	0.7216	
PR	41 29.27 (31.794) -4.40 (3.646)	22 43.94 (31.933)	-4.92 (5.637)	0.51 (-13.223, 14.248)	0.9404	
Hedges' g SMD					0.02 (-0.497, 0.539)	0.9376	
Int. p-value						0.0834	
Screening labora	tory tBRCA status	(IVRS)					
tBRCAm	131 41.35 (31.170) -8.93 (1.763)	57 35.67 (31.409)	-7.72 (2.825)	-1.21 (-7.782, 5.368)	0.7176	
Hedges' g SMD					-0.06 (-0.370, 0.253)	0.7122	
non-tBRCAm	88 35.42 (31.945) -9.82 (2.093)	61 42.08 (35.565)	-8.75 (2.616)	-1.06 (-7.714, 5.590)	0.7527	
Hedges' g SMD					-0.05 (-0.380, 0.274)	0.7508	
Int. p-value						0.9023	
First line treat	ment outcome (eCRF	')					
NED [PDS]	79 41.98 (32.671	•	41 42.28 (32.935)	-6.62 (3.123)	-5.20 (-12.769, 2.364)	0.1758	
Hedges' g SMD	,	,	(,,	-0.26 (-0.641, 0.116)	0.1741	
NED/CR [IDS]	60 39.72 (29.287) -9.06 (2.710)	29 31.61 (35.731)	-10.77 (4.225)	1.71 (-8.293, 11.713)	0.7345	
Hedges' g SMD	, , ,		, , , ,	, -,	0.08 (-0.365, 0.522)	0.7284	
NED/CR [Chemo]	33 36.87 (34.550) -7.58 (3.316)	17 31.37 (35.786)	-4.44 (5.118)	-3.14 (-15.592, 9.308)	0.6113	
Hedges' g SMD	, , , , , , , , , , , , , , , , , , , ,		, , ,	/	-0.16 (-0.743, 0.429)	0.6006	
PR	44 32.58 (30.701) -5.55 (3.335)	30 45.56 (31.237)	-8.64 (4.393)	3.10 (-8.106, 14.298)	0.5824	
Hedges' g SMD	,		, , , ,	, , ,	0.13 (-0.331, 0.598)	0.5728	
Int. p-value					, , , , , , , , , , , , , , , , , , , ,	0.3305	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.2 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Body image (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + bev (N=132		Difference l	oetween g	roups
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
Screening labor	atory tE	BRCA status (e	CRF)						
tBRCAm	128 41	.93 (31.232)	-9.01 (1.796)	58	35.06 (31.482)	-7.42 (2.822)	-1.59 (-8.199,	5.019)	0.6356
Hedges' g SMD							-0.08 (-0.387,	0.234)	0.6289
non-tBRCAm	91 34	.80 (31.686)	-9.67 (2.041)	60	42.78 (35.437)	-8.86 (2.621)	-0.81 (-7.414,	5.791)	0.8083
Hedges' g SMD							-0.04 (-0.367,	0.285)	0.8064
Int. p-value									0.9983
Age group									
<65 years	161 40	.79 (31.397)	-8.49 (1.637)	89	39.51 (32.893)	-7.90 (2.317)	-0.59 (-6.183,	4.996)	0.8345
Hedges' g SMD							-0.03 (-0.287,	0.231)	0.8324
>=65 years	58 33	.91 (31.679)	-10.47 (2.194)	29	37.36 (36.367)	-9.92 (3.337)	-0.54 (-8.493,	7.405)	0.8920
Hedges' g SMD							-0.03 (-0.477,	0.414)	0.8900
Int. p-value									0.5069
FIGO Stage (Dis	ease sta	ate)							
III	156 41	.56 (31.820)	-9.50 (1.586)	82	38.01 (34.174)	-9.40 (2.276)	-0.10 (-5.572,	5.364)	0.9700
Hedges' g SMD							-0.01 (-0.273,	0.262)	0.9697
IV	63 32	2.54 (30.147)	-7.08 (2.804)	36	41.20 (32.729)	-6.34 (4.146)	-0.74 (-10.726,	9.247)	0.8833
Hedges' g SMD							-0.03 (-0.441,	0.378)	0.8799
Int. p-value									0.9241
Region									
Europe	211 39	.89 (31.630)	-9.71 (1.389)	112	39.29 (33.992)	-9.11 (2.023)	-0.60 (-5.427,	4.230)	0.8074
Hedges' g SMD							-0.03 (-0.258,	0.200)	0.8042
Japan	8	NC	NC	6	NC	NC	NC		NC
Int. p-value									NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

SMD = standardised mean difference. * p<0.05.

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.2 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Body image (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=25			Placebo + be		Difference between groups		
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
ECOG performance	e stat	us at Baseline							
(0) Normal activity	162	37.55 (30.086)	-7.83 (1.607)	90	37.22 (33.244)	-6.62 (2.229)	-1.21 (-6.625,	4.205)	0.6600
Hedges' g SMD							-0.06 (-0.316,		0.6576
(1) Restricted activity	53	43.40 (35.707)	-11.23 (2.636)	28	44.64 (34.858)	-15.13 (3.981)	3.89 (-5.608,	13.398)	0.4172
Hedges' g SMD Int. p-value							0.19 (-0.265,	0.653)	0.4066 0.5597
Baseline CA-125 <=ULN Hedges' g SMD >ULN Int. p-value			-9.65 (1.407) NC	105	39.84 (34.398) NC	-8.07 (2.015)	-1.59 (-6.422, -0.08 (-0.317, NC		0.5192 0.5133 NC NC
Histological gra	ade								
High grade	219	38.96 (31.546)	-9.12 (1.363)	118	38.98 (33.633)	-8.38 (1.953)	-0.74 (-5.430,	,	0.7549
Hedges' g SMD Int. p-value							-0.04 (-0.260,	0.188)	0.7516 ID
Cytoreductive su	ırgerv	outcome							
No residue			-10.61 (1.712)	71	38.03 (34.061)	-7.57 (2.520)	-3.04 (-9.053,	2.963)	0.3188
Hedges' g SMD							-0.15 (-0.432,		0.3134
Residue	70	36.19 (32.473)	-7.12 (2.375)	40	40.83 (33.536)	-8.24 (3.330)	1.11 (-7.016,		0.7862
Hedges' g SMD							0.05 (-0.334,	0.443)	0.7835
Int. p-value									0.5050

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

SMD = standardised mean difference. * p<0.05.

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.2 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Body image (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=25			Placebo + be		Difference between groups		
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Timing of cytor	educti	ve surgery							
Upfront	130	39.62 (32.633)	-10.79 (1.724)	70	40.24 (32.039)	-6.40 (2.416)	-4.38 (-10.238, 1.475)	0.1417	
Hedges' g SMD							-0.22 (-0.511, 0.072)	0.1392	
Interval	82	39.84 (29.832)	-7.49 (2.340)	41	36.99 (36.801)	-10.58 (3.587)	3.09 (-5.404, 11.581)	0.4726	
Hedges' g SMD							0.14 (-0.234, 0.516)	0.4621	
Int. p-value								0.0746	
Myriad tumour B	RCA mu	itation status							
tBRCAm	137	40.88 (31.634)	-9.39 (1.755)	66	36.87 (32.491)	-8.40 (2.675)	-0.99 (-7.298, 5.323)	0.7579	
Hedges' g SMD							-0.05 (-0.341, 0.247)	0.7538	
Non-tBRCAm	82	35.77 (31.331)	-8.88 (2.124)	52	41.67 (35.162)	-8.04 (2.779)	-0.84 (-7.787, 6.112)	0.8118	
Hedges' g SMD							-0.04 (-0.390, 0.305)	0.8101	
Int. p-value								0.8718	
Status somatic	BRCA m	nutations							
sBRCAm	18	NC	NC	6	NC	NC	NC	NC	
gBRCAm	57	37.72 (31.580)	-1.59 (2.742)	27	38.89 (31.009)	-9.26 (4.314)	7.66 (-2.505, 17.829)	0.1376	
Hedges' g SMD							0.36 (-0.104, 0.818)	0.1291	
Non-BRCAm	35	41.43 (31.408)	-10.72 (3.334)	22	40.15 (35.134)	-9.59 (4.449)	-1.13 (-12.298, 10.041)	0.8400	
Hedges' g SMD							-0.06 (-0.589, 0.478)	0.8393	
Int. p-value								0.6300	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.6.4.3 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + b		Placebo + bev		Difference between g	roups
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
First line treat	ment outcome (IVRS)					
NED [PDS]	82 26.42 (16.176)	-4.56 (1.167)	42 26.55 (21.224)	-3.02 (1.717)	-1.55 (-5.657, 2.563)	0.4576
Hedges' g SMD					-0.14 (-0.515, 0.229)	0.4520
NED/CR [IDS]	61 25.25 (20.408)	-1.00 (1.561)	34 23.14 (14.609)	-2.17 (2.202)	1.16 (-4.204, 6.529)	0.6679
Hedges' g SMD					0.09 (-0.327, 0.512)	0.6647
NED/CR [Chemo]	37 26.82 (21.783)	-1.80 (2.431)	20 20.67 (13.663)	-2.46 (3.500)	0.66 (-7.940, 9.257)	0.8785
Hedges' g SMD					0.04 (-0.501, 0.587)	0.8768
PR	41 23.10 (18.968)	-0.02 (1.610)	22 24.32 (13.827)	1.42 (2.402)	-1.44 (-7.254, 4.370)	0.6202
Hedges' g SMD					-0.13 (-0.652, 0.385)	0.6131
Int. p-value						0.8218
Screening labora tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value	atory tBRCA status (131 27.11 (18.425) 90 23.28 (19.289)	-3.02 (1.105)	, ,		-1.25 (-5.376, 2.877) -0.10 (-0.408, 0.214) 0.21 (-3.394, 3.821) 0.02 (-0.305, 0.345)	0.5509 0.5418 0.9070 0.9060 0.2883
	ement outcome (eCRF)					
NED [PDS]	80 28.61 (18.254)	-5.45 (1.217)	41 26.22 (21.689)	-3.49 (1.759)	-1.95 (-6.194, 2.287)	0.3636
Hedges' g SMD					-0.18 (-0.553, 0.201)	0.3597
NED/CR [IDS]	60 23.22 (19.810)	-1.68 (1.522)	29 22.53 (14.520)	-1.34 (2.353)	-0.33 (-5.906, 5.241)	0.9059
Hedges' g SMD					-0.03 (-0.471, 0.416)	0.9040
NED/CR [Chemo]	34 25.00 (19.249)	-2.21 (2.508)	17 25.10 (13.023)	-3.45 (3.852)	1.24 (-8.021, 10.503)	0.7884
Hedges' g SMD					0.08 (-0.501, 0.664)	0.7847
PR	44 24.10 (18.580)	2.51 (1.573)	30 22.28 (14.118)	1.02 (2.081)	1.49 (-3.725, 6.706)	0.5700
Hedges' g SMD					0.14 (-0.328, 0.601)	0.5657
Int. p-value						0.5529

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.3 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + bev (N=132		Difference b	oetween g	roups
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
Screening labor	atory t	BRCA status (e	CRF)						
tBRCAm	128 2	7.27 (18.533)	-2.97 (1.119)	58	21.81 (14.157)	-1.70 (1.749)	-1.27 (-5.385,	2.850)	0.5444
Hedges' g SMD							-0.10 (-0.408,	0.212)	0.5358
non-tBRCAm	93 23	3.17 (19.082)	-1.97 (1.096)	60	26.42 (19.109)	-2.03 (1.450)	0.06 (-3.543,	3.659)	0.9747
Hedges' g SMD							0.01 (-0.319,	0.330)	0.9744
Int. p-value									0.2910
Age group									
<65 years	162 26	6.47 (18.922)	-3.09 (0.975)	88	24.89 (17.461)	-3.05 (1.392)	-0.04 (-3.388,	3.310)	0.9816
Hedges' g SMD							0.00 (-0.263,	0.256)	0.9814
>=65 years	59 23	3.00 (18.506)	-0.64 (1.296)	30	22.00 (15.403)	0.70 (1.960)	-1.35 (-6.028,	3.336)	0.5686
Hedges' g SMD							-0.13 (-0.570,	0.309)	0.5609
Int. p-value									0.8819
FIGO Stage (Dis	ease st	ate)							
III	158 2	7.27 (19.953)	-2.58 (0.967)	83	25.34 (17.842)	-3.09 (1.391)	0.51 (-2.835,	3.848)	0.7657
Hedges' g SMD							0.04 (-0.225,	0.307)	0.7630
IV	63 23	1.22 (14.959)	-1.79 (1.428)	35	21.33 (14.444)	0.27 (2.159)	-2.05 (-7.193,	3.090)	0.4301
Hedges' g SMD							-0.17 (-0.585,	0.242)	0.4166
Int. p-value									0.1306
Region									
Europe	212 20	6.10 (18.908)	-2.66 (0.819)	112	24.73 (17.017)	-1.80 (1.198)	-0.86 (-3.718,	1.994)	0.5532
Hedges' g SMD							-0.07 (-0.300,	0.158)	0.5461
Japan	9	NC	NC	6	NC	NC	NC		NC
Int. p-value									NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/mmrmsubpr_v3.sas emmrmsubpr_v3bc 25NOV2020:12:36 kvbv306

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.3 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be			Placebo + bev		Difference between groups		
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
ECOG performance	e stat	us at Baseline							
(0) Normal activity	163	23.92 (16.657)	-1.62 (0.840)	90	22.96 (17.275)	-1.41 (1.172)	-0.21 (-3.045,	2.635)	0.8868
Hedges' g SMD (1) Restricted	54	30.27 (24.020)	-5.55 (2.008)	28	27.98 (15.512)	-3.51 (3.061)	-0.02 (-0.276, -2.04 (-9.325,		0.8859 0.5797
activity Hedges' g SMD Int. p-value							-0.13 (-0.589,	0.325)	0.5704 0.8108
Baseline CA-125			2 02 / 0 020)	105	OF 11 (17 201)	1 06 / 1 010)	1 00 / 2 077	1 005)	0.4660
<=ULN Hedges' g SMD	196	25.55 (18.846)	-3.03 (0.839)	105	25.11 (17.381)	-1.96 (1.212)	-1.08 (-3.977, -0.09 (-0.327,		0.4592
>ULN Int. p-value	25	NC	NC	13	NC	NC	NC	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	NC NC
Histological gra	ade								
High grade	221	25.55 (18.833)	-2.44 (0.794)	118	24.15 (16.945)	-1.96 (1.147)	-0.48 (-3.229,	2.263)	0.7295
Hedges' g SMD Int. p-value							-0.04 (-0.264,	0.183)	0.7255 ID
Cytoreductive su	ırgery	outcome							
No residue	143	26.12 (18.916)	-3.92 (0.934)	71	24.72 (18.881)	-2.71 (1.386)	-1.21 (-4.509,	2.084)	0.4693
Hedges' g SMD							-0.11 (-0.391,	•	0.4635
Residue	71	25.45 (19.233)	0.35 (1.514)	40	22.38 (13.418)	-0.77 (2.149)	1.11 (-4.105,	,	0.6729
Hedges' g SMD							0.08 (-0.303,	0.472)	0.6684
Int. p-value									0.2232

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.4.3 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + bev		Difference between	groups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Timing of cytore	educti	ve surgery						
Upfront	131 2	26.48 (18.379)	-3.87 (1.005)	70	24.90 (18.706)	-2.35 (1.426	5) -1.52 (-4.967, 1.919	0.3837
Hedges' g SMD							-0.13 (-0.421, 0.160	0.3792
Interval	83 2	24.98 (19.969)	-0.29 (1.371)	41	22.11 (13.939)	-1.60 (2.094	1) 1.31 (-3.649, 6.270	0.6019
Hedges' g SMD							0.10 (-0.273, 0.476	0.5941
Int. p-value								0.3886
Myriad tumour BI	RCA mu	tation status						
tBRCAm	137 2	26.60 (18.200)	-3.21 (1.059)	66	23.41 (15.690)	-2.22 (1.602	2) -0.99 (-4.782, 2.806	0.6081
Hedges' g SMD							-0.08 (-0.372, 0.216	0.6021
Non-tBRCAm	84 2	23.83 (19.813)	-1.43 (1.181)	52	25.10 (18.528)	-1.50 (1.592	2) 0.06 (-3.863, 3.986	0.9752
Hedges' g SMD							0.01 (-0.340, 0.351	0.9750
Int. p-value								0.4032
Status somatic H	BRCA m	utations						
sBRCAm	18 2	20.74 (16.151)	0.52 (2.352)	6	12.22 (7.794)	-1.93 (4.229	9) 2.45 (-7.777, 12.681	0.6224
Hedges' g SMD							0.24 (-0.692, 1.162	0.6190
gBRCAm	57 2	27.37 (18.969)	0.32 (1.907)	26	21.03 (12.464)	-1.70 (3.001	l) 2.01 (-5.095, 9.121	0.5747
Hedges' g SMD							0.14 (-0.328, 0.600	0.5664
Non-BRCAm	37 2	25.00 (22.845)	-2.16 (1.614)	22	24.09 (21.527)	-0.06 (2.332	2) -2.10 (-7.792, 3.599	0.4635
Hedges' g SMD							-0.20 (-0.731, 0.327	0.4543
Int. p-value								0.8042

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.6.4.4 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		bevacizumab =255)	Placebo + be (N=13		Difference between g	roups
Subgroup	n Mean (SD) a [a] Baseline [b		n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
First line treat	ment outcome (IVR	S)				
NED [PDS]	81 59.60 (26.10)	2) -16.58 (1.796)	42 59.26 (26.307)	-12.70 (2.616)	-3.88 (-10.163, 2.400)	0.2236
Hedges' g SMD					-0.23 (-0.609, 0.139)	0.2186
NED/CR [IDS]	60 51.76 (29.79)	5) -8.45 (2.510)	34 52.78 (29.659)	-14.46 (3.510)	6.01 (-2.560, 14.588)	0.1668
Hedges' g SMD					0.30 (-0.122, 0.724)	0.1631
NED/CR [Chemo]	36 58.02 (28.49)	7) -9.22 (3.182)	20 50.00 (24.845)	-13.63 (4.690)	4.41 (-7.073, 15.893)	0.4434
Hedges' g SMD					0.22 (-0.328, 0.768)	0.4318
PR	40 55.28 (24.84	4) -14.10 (3.094)	22 63.64 (24.829)	-6.25 (4.720)	-7.84 (-19.274, 3.584)	0.1735
Hedges' g SMD					-0.38 (-0.903, 0.147)	0.1582
Int. p-value						0.0489
tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value	·	(IVRS) 9) -12.57 (1.590) 3) -13.10 (1.850)	, ,		1.97 (-3.960, 7.904) 0.11 (-0.205, 0.417) -1.91 (-7.844, 4.022) -0.11 (-0.436, 0.221)	0.5127 0.5042 0.5251 0.5210 0.6449
	ement outcome (eCR					
NED [PDS]	79 61.11 (26.99)	3) -15.63 (1.890)	41 57.99 (26.030)	-11.94 (2.706)	-3.69 (-10.235, 2.847)	0.2657
Hedges' g SMD	50 50 05 (00 15	2) 2 5 7 7 2	00.40.01.700.555	10 00 / 0 000	-0.22 (-0.594, 0.162)	0.2628
NED/CR [IDS]	59 52.07 (29.45)	0) -9.57 (2.557)	29 49.81 (29.639)	-13.38 (3.882)	3.81 (-5.430, 13.053)	0.4144
Hedges' g SMD	00.54.55.705.00		15 51 60 (04 55 1)	0 44 (5 242)	0.19 (-0.257, 0.634)	0.4072
NED/CR [Chemo]	33 54.55 (27.82)	3) -10.25 (3.349)	17 51.63 (24.514)	-9.11 (5.241)	, , , , , , , , , , , , , , , , , , , ,	0.8560
Hedges' g SMD	40.55.56.701.10		20 50 50 (05 555)	10 00 (0 = :=:	-0.06 (-0.641, 0.529)	0.8513
PR	43 55.56 (24.48)	3) -11.09 (2.816)	30 62.78 (25.360)	-13.93 (3.747)	2.83 (-6.588, 12.253)	0.5498
Hedges' g SMD					0.14 (-0.322, 0.612)	0.5428
Int. p-value						0.3316

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.4 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be	evacizumab 5)		Placebo + be (N=13		Difference l	Difference between groups		
Subgroup	n [a]	Mean (SD) at	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value	
Screening labor	atory t	BRCA status (e	eCRF)							
tBRCAm	128 5	8.51 (27.100)	-12.09 (1.604)	58	54.89 (26.777)	-14.90 (2.523)	2.81 (-3.093,	8.715)	0.3488	
Hedges' g SMD							0.15 (-0.159,	0.462)	0.3394	
non-tBRCAm	89 5	3.31 (27.584)	-13.80 (1.811)	60	58.33 (27.188)	-10.83 (2.373)	-2.97 (-8.883,	2.935)	0.3214	
Hedges' g SMD							-0.17 (-0.496,	0.160)	0.3155	
Int. p-value									0.3589	
Age group										
<65 years	160 5	7.99 (25.869)	-12.30 (1.417)	89	58.05 (26.232)	-14.16 (2.019)	1.85 (-3.008,	6.712)	0.4536	
Hedges' g SMD							0.10 (-0.159,	0.360)	0.4466	
>=65 years	57 5	1.85 (30.956)	-13.26 (2.270)	29	52.30 (29.002)	-11.40 (3.452)	-1.86 (-10.087,	6.375)	0.6548	
Hedges' g SMD							-0.10 (-0.552,	0.343)	0.6476	
Int. p-value									0.7172	
FIGO Stage (Dis	ease st	cate)								
III	154 5	66.85 (28.355)	-13.15 (1.469)	82	56.23 (28.306)	-14.04 (2.102)	0.89 (-4.166,	5.941)	0.7296	
Hedges' g SMD							0.05 (-0.220,	0.316)	0.7267	
IV	63 5	5.20 (24.923)	-11.83 (1.913)	36	57.56 (23.836)	-13.26 (3.109)	1.43 (-5.830,	8.680)	0.6971	
Hedges' g SMD							0.09 (-0.324,	0.495)	0.6822	
Int. p-value									0.7150	
Region										
Europe	209 5	7.10 (27.406)	-12.89 (1.239)	112	57.39 (26.974)	-13.14 (1.815)	0.25 (-4.077,	4.570)	0.9107	
Hedges' g SMD							0.01 (-0.216,	0.243)	0.9089	
Japan Int. p-value	8	NC	NC	6	NC	NC	NC		NC NC	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.4.4 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=25			Placebo + be		Difference b	etween g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
ECOG performance	stat	us at Baseline							
(0) Normal activity	161	54.76 (26.534)	-11.64 (1.420)	90	56.05 (27.979)	-11.29 (1.975)	-0.35 (-5.136,		0.8872
Hedges' g SMD (1) Restricted	52	61.86 (28.626)	-16.06 (2.351)	28	58.53 (23.594)	-21.30 (3.745)	-0.02 (-0.277, 5.25 (-3.547,		0.8862 0.2388
activity Hedges' g SMD Int. p-value							0.29 (-0.173,	0.750)	0.2204 0.2736
Baseline CA-125			10 77 / 1 070)	105	FF 02 /26 460\	10 20 / 1 000)	0.20 / 4.760	4 001)	0.0640
K=ULN Hedges' g SMD	192	56.48 (27.537)	-12.77 (1.270)	105	55.93 (26.468)	-12.39 (1.828)	-0.38 (-4.760, -0.02 (-0.259,	,	0.8648 0.8626
>ULN Int. p-value	25	NC	NC	13	NC	NC	NC		NC NC
Histological gra	.de								
High grade		56.37 (27.356)	-12.66 (1.203)	118	56.64 (26.927)	-13.31 (1.736)	0.65 (-3.506,	,	0.7589
Hedges' g SMD Int. p-value							0.04 (-0.188,	0.260)	0.7549 ID
Cytoreductive su	rgery	outcome							
No residue	141	57.05 (28.218)	-13.49 (1.492)	71	55.24 (27.997)	-13.01 (2.205)	-0.48 (-5.731,	4.769)	0.8569
Hedges' g SMD							-0.03 (-0.312,	•	0.8550
Residue	69	57.33 (25.892)	-11.76 (2.269)	40	59.44 (25.387)	-13.28 (3.229)	,		0.7016
Hedges' g SMD Int. p-value							0.08 (-0.312,	0.467)	0.6965 0.4759

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

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Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.4.4 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be			Placebo + be N=13		Difference between g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Timing of cytore	educti	ve surgery						
Upfront			-15.18 (1.531)	70	58.65 (24.712)	-12.68 (2.162)	-2.50 (-7.721, 2.729)	0.3473
Hedges' g SMD			, ,		,	,	-0.14 (-0.432, 0.150)	0.3428
Interval	81	54.12 (28.161)	-9.44 (2.090)	41	53.52 (30.675)	-13.91 (3.153)	4.47 (-3.023, 11.961)	0.2399
Hedges' g SMD							0.23 (-0.146, 0.607)	0.2309
Int. p-value								0.1525
Myriad tumour Bi tBRCAm Hedges' g SMD Non-tBRCAm Hedges' g SMD Int. p-value	137	57.50 (26.683)	-12.11 (1.539) -13.90 (1.946)		, ,	-14.58 (2.353) -11.26 (2.584)	0.13 (-0.160, 0.428)	0.3810 0.3720 0.4173 0.4119 0.3788
Status somatic I	BRCA m	utations						
sBRCAm	18	55.56 (24.403)	-12.35 (3.061)	6	53.70 (30.157)	-18.77 (5.576)	6.43 (-6.871, 19.724)	0.3268
Hedges' g SMD							0.47 (-0.463, 1.408)	0.3225
gBRCAm	57	56.14 (27.752)	-8.94 (2.342)	27	59.47 (27.598)	-15.71 (3.702)	6.77 (-1.967, 15.502)	0.1270
Hedges' g SMD							0.37 (-0.093, 0.830)	0.1174
Non-BRCAm	33	55.89 (31.240)	-16.58 (2.611)	22	52.53 (30.895)	-10.64 (3.570)	-5.94 (-14.845, 2.964)	0.1861
Hedges' g SMD							-0.37 (-0.917, 0.172)	0.1797
Int. p-value								0.132

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Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.6.4.5 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Hormonal (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=25!			Placebo + bev		Difference between o	groups
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-valu
First line treat	tment o	utcome (IVRS)						
NED [PDS]	81 31	1.89 (32.829)	-6.22 (2.087)	42 33	3.33 (31.666)	-0.66 (3.037)	-5.55 (-12.849, 1.741)	0.134
Hedges' g SMD							-0.29 (-0.664, 0.085)	0.130
NED/CR [IDS]	61 25	5.41 (30.823)	1.52 (2.374)	33 17	7.68 (30.029)	-1.84 (3.400)	3.35 (-4.923, 11.630)	0.423
Hedges' g SMD							0.18 (-0.248, 0.600)	0.416
NED/CR [Chemo]	37 17	7.57 (25.137)	-2.78 (3.015)	20 25	5.83 (26.752)	1.54 (4.231)	-4.32 (-14.778, 6.130)	0.410
Hedges' g SMD							-0.23 (-0.776, 0.316)	0.408
PR	41	NC	NC	22	NC	NC	NC	NC
Int. p-value								0.516
tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value		4.53 (30.679)	-2.21 (1.637) -3.53 (1.903)		9.17 (31.342) 4.04 (30.961)	0.27 (2.405)	-1.68 (-7.787, 4.418) -0.09 (-0.401, 0.225) -3.80 (-9.857, 2.263) -0.21 (-0.533, 0.120)	0.586 0.580 0.217 0.215 0.998
First line treat NED [PDS]		utcome (eCRF) 2.28 (32.395)	-6.86 (2.144)	41 33	3.33 (32.059)	-0.20 (3.064)	-6.66 (-14.066, 0.749)	0.077
Hedges' g SMD	10 02	2.20 (32.333)	0.00 (2.111)	11 3.	3.33 (32.03)	0.20 (5.001)	-0.34 (-0.724, 0.036)	0.076
NED/CR [IDS]	60 23	3.33 (30.097)	2.35 (2.322)	28 13	2.50 (25.909)	-0.04 (3.631)	, , , , , , , , , , , , , , , , , , , ,	0.583
	00 23	3.33 (30.037)	2.33 (2.322)	20 12	2.00 (20.00)	0.01 (0.001)	0.13 (-0.320, 0.578)	0.573
Hedges' a SMD	34	NC	NC	17	NC	NC	NC	NC
Hedges' g SMD NED/CR [Chemo]						NC	NC	NC
NED/CR [Chemo] PR	44	NC	NC	30	NC	INC	NC	NC

Screening laboratory tBRCA status (eCRF)

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

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[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.4.5 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Hormonal (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=255		Placebo + bev		Difference between groups		
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
tBRCAm Hedges' g SMD non-tBRCAm Hedges' g SMD Int. p-value	128 26.95 (29.365) 92 23.91 (30.393)	-2.25 (1.670) -3.43 (1.844)	57 30.12 (31.879) 60 23.06 (30.236)	-0.34 (2.622) 0.09 (2.393)	-1.91 (-8.046, 4.225) -0.10 (-0.411, 0.213) -3.53 (-9.495, 2.445) -0.19 (-0.521, 0.131)	0.5398 0.5335 0.2451 0.2421 0.8439	
Age group <65 years Hedges' g SMD >=65 years Hedges' g SMD Int. p-value	162 29.84 (31.218) 58 14.08 (21.585)			, ,	-1.81 (-7.110, 3.500) -0.09 (-0.350, 0.170) -2.56 (-7.972, 2.860) -0.22 (-0.663, 0.231)	0.5034 0.4975 0.3506 0.3430 0.9803	
FIGO Stage (Disc III Hedges' g SMD IV Hedges' g SMD Int. p-value	ease state) 157 25.69 (28.836) 63 25.66 (32.216)		, ,		-3.48 (-8.701, 1.743) -0.18 (-0.450, 0.087) 0.96 (-6.245, 8.168) 0.06 (-0.353, 0.466)	0.1907 0.1857 0.7918 0.7878 0.5197	
Region Europe Hedges' g SMD Japan Int. p-value	211 26.30 (30.155) 9 NC	-2.88 (1.281) NC	111 27.78 (31.409) 6 NC	-0.41 (1.866)	-2.46 (-6.915, 1.993) -0.13 (-0.359, 0.101) NC	0.2778 0.2704 NC NC	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.

SMD = standardised mean difference. * p<0.05.

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.4.5 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Hormonal (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=255		Placebo + bev		Difference betw	een groups
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% C	I) p-value
(0) Normal activity Hedges' g SMD (1) Restricted activity Hedges' g SMD Int. p-value	163 28.22 (31.824) 53 18.55 (21.844)	-4.63 (1.468) 2.89 (2.284)	90 25.74 (30.309) 27 29.01 (34.154)	-1.49 (2.034) 1.46 (3.587)	-3.15 (-8.091, 1. -0.17 (-0.423, 0. 1.43 (-7.050, 9. 0.08 (-0.382, 0.	092) 0.2081 917) 0.7377
Baseline CA-125 <=ULN Hedges' g SMD >ULN Int. p-value		-2.74 (1.312) NC	104 27.40 (31.410) 13 NC	-0.54 (1.891) NC	-2.20 (-6.731, 2. -0.12 (-0.356, 0. NC	
Histological gra High grade Hedges' g SMD Int. p-value		-2.78 (1.234)	117 26.50 (31.115)	-0.40 (1.777)	-2.38 (-6.637, 1. -0.13 (-0.352, 0.	
Cytoreductive so No residue Hedges' g SMD Residue Hedges' g SMD Int. p-value	rgery outcome 142 27.82 (31.556) 71 22.30 (27.013)		, ,	,	-2.89 (-8.372, 2. -0.15 (-0.440, 0. -1.69 (-9.145, 5. -0.09 (-0.477, 0.	133) 0.2937 765) 0.6541
Timing of cytore	eductive surgery 130 26.54 (30.689)	-5.05 (1.647)	70 30.48 (30.026)	0.53 (2.306)	-5.59 (-11.177, 0.	003) 0.0501

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.5 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Hormonal (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be (N=255		Placebo + bev		Difference between groups		
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Hedges' g SMD Interval Hedges' g SMD Int. p-value	83 25.10 (29.491)	0.68 (1.965)	40 17.92 (29.811)	-0.61 (3.022)	-0.29 (-0.586, -0.002) 1.30 (-5.871, 8.462) 0.07 (-0.307, 0.448)	0.0488* 0.7211 0.7149 0.3429	
Myriad tumour B	RCA mutation status						
tBRCAm Hedges' g SMD	137 26.16 (29.275)	-2.91 (1.588)	65 29.74 (32.741)	-1.08 (2.419)	-1.84 (-7.547, 3.870) -0.10 (-0.392, 0.198)	0.5262 0.5199	
Non-tBRCAm Hedges' g SMD Int. p-value	83 24.90 (30.729)	-2.40 (1.977)	52 22.44 (28.751)	0.88 (2.608)	-3.28 (-9.753, 3.195) -0.18 (-0.525, 0.169)	0.3183 0.3152 0.9017	
Status somatic I	BRCA mutations						
sBRCAm	18 NC	NC	5 NC	NC	NC	NC	
gBRCAm	57 27.78 (27.877)	5.23 (2.655)	27 24.69 (30.791)	2.74 (4.196)	2.49 (-7.431, 12.410)	0.6188	
Hedges' g SMD					0.12 (-0.339, 0.578)	0.6089	
Non-BRCAm	36 25.00 (28.031)	-1.65 (2.614)	22 18.94 (24.825)	-0.25 (3.660)	-1.41 (-10.454, 7.639)	0.7565	
Hedges' g SMD Int. p-value					-0.09 (-0.616, 0.445)	0.7524 0.8212	

model within-subject error.

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.4.6 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

Subgroup		+ bevacizumab (=255)	Placebo + b (N=1		Difference between groups		
	n Mean (SD) [a] Baseline [n Mean (SD) at [a] Baseline [b]		Estimated difference (95% CI)	p-value	
First line treat	ment outcome (IV)	RS)					
NED [PDS]	82 50.00 (35.33	31) -20.55 (2.544)	42 50.79 (31.441)	-21.63 (3.694)	1.08 (-7.800, 9.955)	0.8105	
Hedges' g SMD					0.05 (-0.326, 0.418)	0.8089	
NED/CR [IDS]	61 36.61 (35.33	32) -12.50 (2.170)	34 39.22 (37.128)	-12.62 (3.038)	0.11 (-7.345, 7.570)	0.9761	
Hedges' g SMD					0.01 (-0.413, 0.426)	0.9758	
NED/CR [Chemo]	37 38.74 (31.44	10) -3.02 (3.066)	20 41.67 (38.805)	-11.63 (4.531)	8.61 (-2.320, 19.530)	0.1200	
Hedges' g SMD					0.44 (-0.109, 0.992)	0.1159	
PR	41 35.77 (31.96	54) -7.53 (2.697)	22 43.94 (36.567)	7.21 (4.358)	-0.32 (-10.658, 10.021)	0.9509	
Hedges' g SMD					-0.02 (-0.535, 0.501)	0.9485	
Int. p-value						0.9243	
Screening labora	itory tBRCA statu:	s (IVRS)					
tBRCAm	131 40.59 (34.53	36) -11.13 (1.671)	57 36.26 (30.719)	9.58 (2.668)	-1.55 (-7.764, 4.667)	0.6235	
Hedges' g SMD					-0.08 (-0.390, 0.232)	0.6171	
non-tBRCAm	90 43.52 (34.50	01) -17.01 (2.222)	61 52.46 (37.618)	-20.48 (2.852)	3.47 (-3.696, 10.635)	0.3402	
Hedges' g SMD					0.16 (-0.166, 0.486)	0.3353	
Int. p-value						0.2407	
First line treat	ment outcome (eCl	RF)					
NED [PDS]	,	90) -20.40 (2.630)	41 47.97 (31.666)) -22.76 (3.763)	2.36 (-6.736, 11.459)	0.6083	
Hedges' g SMD	,	,	(0.10 (-0.278, 0.476)	0.6066	
NED/CR [IDS]	60 35.00 (35.48	32) -12.40 (2.214)	29 40.23 (37.932)) -10.73 (3.404)		0.6826	
Hedges' g SMD	,	,	,	,	-0.09 (-0.538, 0.349)	0.6755	
NED/CR [Chemo]	34 37.25 (30.16	58) -4.07 (3.094)	17 52.94 (38.295)) -13.88 (5.067)		0.1060	
Hedges' g SMD	,	•	, , , , , , , , , , , , , , , , , , , ,	,	0.51 (-0.083, 1.099)	0.0921	
PR	44 35.98 (31.93	36) -6.74 (2.587)	30 37.78 (34.722)	9.14 (3.543)		0.5867	
Hedges' g SMD	,	, , , ,		, ,	0.13 (-0.334, 0.596)	0.5804	
Int. p-value					, , , , , , , , , , , , , , , , , , , ,	0.9337	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.6 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)				Placebo + be		Difference between groups		
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
Screening labor	atory tH	BRCA status (eCRF)						
tBRCAm	128 40	.89 (34.801)	-10.87 (1.682)	58	36.21 (30.451)	-9.67 (2.631)	-1.20 (-7.368, 4.966)	0.7011	
Hedges' g SMD							-0.06 (-0.372, 0.248)	0.6961	
non-tBRCAm	93 43	.01 (34.167)	-17.25 (2.191)	60	52.78 (37.853)	-20.43 (2.882)	3.18 (-3.999, 10.356)	0.3830	
Hedges' g SMD							0.15 (-0.179, 0.471)	0.3773	
Int. p-value								0.3086	
Age group									
<65 years	162 38	3.48 (34.417)	-13.62 (1.509)	88	40.53 (34.072)	-14.21 (2.160)	0.60 (-4.596, 5.789)	0.8212	
Hedges' g SMD							0.03 (-0.229, 0.290)	0.8187	
>=65 years	59 50	.85 (33.250)	-12.54 (2.739)	30	56.67 (36.515)	-15.89 (4.020)	3.35 (-6.319, 13.020)	0.4930	
Hedges' g SMD							0.16 (-0.285, 0.596)	0.4888	
Int. p-value								0.5458	
FIGO Stage (Dis	ease sta	ate)							
III	158 44	.62 (35.556)	-13.93 (1.683)	83	47.99 (34.472)	-17.06 (2.407)	3.13 (-2.662, 8.915)	0.2884	
Hedges' g SMD							0.15 (-0.121, 0.411)	0.2838	
IV	63 34	.66 (30.717)	-11.64 (2.132)	35	36.67 (36.335)	-9.52 (3.339)	-2.12 (-9.988, 5.745)	0.5935	
Hedges' g SMD							-0.12 (-0.531, 0.296)	0.5790	
Int. p-value								0.1740	
Region									
Europe	212 42	.30 (34.815)	-13.55 (1.384)	112	44.64 (36.019)	-14.18 (2.010)	0.63 (-4.173, 5.433)	0.7965	
Hedges' g SMD							0.03 (-0.198, 0.260)	0.7934	
Japan Int. p-value	9	NC	NC	6	NC	NC	NC	NC NC	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.6 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)				Placebo + be (N=13		Difference between groups		
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value	
ECOG performance	e stat	us at Baseline							
(0) Normal activity	163	41.31 (33.709)	-12.72 (1.579)	90	39.81 (34.682)	-14.07 (2.182)	1.36 (-3.949, 6.661	0.6150	
Hedges' g SMD (1) Restricted	54	42.59 (36.725)	-16.10 (2.673)	28	60.12 (33.129)	-15.71 (4.353)	0.07 (-0.191, 0.324 -0.38 (-10.669, 9.905		
activity Hedges' g SMD Int. p-value							-0.02 (-0.475, 0.438	0.9380 0.9244	
Baseline CA-125			-13.96 (1.428)	105	46.03 (34.863)	-14.53 (2.049)	0.58 (-4.342, 5.493	0.8181	
Hedges' g SMD >ULN Int. p-value	25	NC	NC	13	NC	NC	0.03 (-0.209, 0.265 NC		
Histological gra	ade								
High grade Hedges' g SMD Int. p-value	221	41.78 (34.474)	-13.38 (1.345)	118	44.63 (35.263)	-14.71 (1.933)	1.33 (-3.308, 5.961 0.07 (-0.158, 0.289		
Cytoreductive su	ırgery	outcome							
No residue	143	44.64 (36.019)	-17.49 (1.758)	71	45.54 (34.728)	-18.06 (2.589)	0.57 (-5.602, 6.740		
Hedges' g SMD	71	20 50 (21 212)	F 30 / 0 000	4.0	40 50 (27 544)	10 40 / 0 050	0.03 (-0.258, 0.311		
Residue Hedges' g SMD Int. p-value	/1	38.50 (31.319)	-5.32 (2.096)	40	42.50 (37.544)	-10.48 (2.978)	5.16 (-2.052, 12.381 0.28 (-0.106, 0.673		

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.6 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)			Placebo + bevacizumab (N=132)				Difference between groups		
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (S Change i baseli	from	Estimated difference (95% CI)	p-value	
Timing of cytore	educti	ve surgery								
Upfront	131	48.85 (33.791)	-15.91 (1.928)	70	48.10 (35.160)	-17.02 (2	2.705)	1.11 (-5.437, 7.663)	0.7380	
Hedges' g SMD								0.05 (-0.240, 0.340)	0.7367	
Interval	83	32.73 (33.682)	-9.14 (1.874)	41	38.21 (35.987)	-10.44 (2	2.858)	1.30 (-5.503, 8.099)	0.7059	
Hedges' g SMD								0.07 (-0.301, 0.448)	0.6993	
Int. p-value									0.7796	
Myriad tumour Bl	RCA mu	tation status								
tBRCAm	137	39.17 (35.274)	-11.04 (1.624)	66	41.41 (33.492)	-11.17 (2	2.457)	0.13 (-5.673, 5.942)	0.9637	
Hedges' g SMD								0.01 (-0.287, 0.301)	0.9631	
Non-tBRCAm	84	46.03 (32.891)	-17.76 (2.356)	52	48.72 (37.318)	-20.63 (3	3.168)	2.87 (-4.943, 10.673)	0.4693	
Hedges' g SMD								0.13 (-0.217, 0.475)	0.4650	
Int. p-value									0.7921	
Status somatic 1	BRCA m	utations								
sBRCAm	18	25.93 (30.903)	-10.22 (3.393)	6	30.56 (37.143)	-0.72 (6	5.089)	-9.49 (-24.008, 5.022)	0.1881	
Hedges' g SMD								-0.63 (-1.576, 0.313)	0.1900	
gBRCAm	57	36.26 (34.521)	-11.84 (2.270)	26	35.90 (25.251)	-13.38 (3	3.681)	1.54 (-7.095, 10.166)	0.7236	
Hedges' g SMD								0.09 (-0.378, 0.550)	0.7159	
Non-BRCAm	37	41.44 (32.778)	-18.30 (2.936)	22	43.18 (33.594)	-20.40 (4	4.183)	2.10 (-8.162, 12.362)	0.6831	
Hedges' g SMD								0.11 (-0.416, 0.640)	0.6781	
Int. p-value									0.6334	

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.6.4.7 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Other single items (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + 1 (N=2		Placebo + b		Difference between g	roups
Subgroup	n Mean (SD) at [a] Baseline [b]		n Mean (SD) at [a] Baseline [b]		Estimated difference (95% CI)	p-value
First line treat	ment outcome (IVRS)				
NED [PDS]	81 25.99 (25.985)	-10.97 (1.068)	42 28.57 (25.021)	-14.26 (1.593)	3.29 (-0.510, 7.088)	0.0891
Hedges' g SMD					0.33 (-0.043, 0.707)	0.0831
NED/CR [IDS]	61 19.40 (18.782)	-4.92 (1.516)	34 24.51 (25.093)	-12.74 (2.150)	7.82 (2.567, 13.075)	0.0040*
Hedges' g SMD					0.64 (0.212, 1.071)	0.0034*
NED/CR [Chemo]	37 27.33 (21.769)	-13.29 (1.742)	20 32.50 (24.830)	-18.35 (2.626)	5.06 (-1.284, 11.397)	0.1154
Hedges' g SMD					0.45 (-0.098, 1.004)	0.1069
PR	41 24.12 (20.179)	-7.58 (1.935)	23 28.26 (28.521)	-9.39 (2.874)	1.80 (-5.132, 8.740)	0.6033
Hedges' g SMD					0.14 (-0.373, 0.649)	0.5965
Int. p-value						0.2171
Screening labora	itory tBRCA status	(IVRS)				
tBRCAm	130 24.55 (22.582)	-8.32 (0.993)	58 26.39 (24.655)	-13.16 (1.585)	4.84 (1.150, 8.529)	0.0104
Hedges' g SMD					0.42 (0.105, 0.730)	0.0089
non-tBRCAm	90 23.30 (22.408)	-10.47 (1.154)	61 29.55 (26.440)	-13.83 (1.510)	3.36 (-0.414, 7.139)	0.0806
Hedges' g SMD					0.30 (-0.031, 0.623)	0.0757
Int. p-value						0.6830
First line treat	ment outcome (eCRF)				
NED [PDS]	79 27.92 (26.509)	,	41 27.37 (25.020)	-15.07 (1.577)	3.21 (-0.574, 6.990)	0.0956
Hedges' g SMD	()		()	2.2. (2.3///	0.33 (-0.052, 0.707)	0.0912
NED/CR [IDS]	60 18.33 (18.184)	-4.47 (1.568)	29 22.70 (22.665)	-12.20 (2.396)	7.73 (2.024, 13.440)	0.0085
Hedges' g SMD				, ,	0.62 (0.166, 1.072)	0.0074
NED/CR [Chemo]	34 22.55 (17.986)	-9.89 (1.904)	17 35.78 (23.051)	-12.72 (3.216)	2.83 (-4.805, 10.460)	0.4579
Hedges' g SMD			,	. , ,	0.23 (-0.349, 0.819)	0.4305
PR	44 25.88 (21.364)	-8.06 (1.969)	31 28.85 (29.777)	-14.25 (2.549)	6.19 (-0.258, 12.646)	0.0596
Hedges' g SMD	, ,	, , ,	, ,	, ,	0.45 (-0.013, 0.918)	0.0566
Int. p-value					, , , , , , , , , , , , , , , , , , , ,	0.5887

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

SMD = standardised mean difference. * p<0.05.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.6.4.7 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Other single items (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=25			Placebo + be (N=13		Difference k	oetween g	roups
Subgroup		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
Screening labor	atory t	BRCA status (e	eCRF)						
tBRCAm	127 2	4.61 (22.804)	-8.12 (1.006)	59	26.13 (24.522)	-13.18 (1.575)	5.06 (1.370,	8.746)	0.0075*
Hedges' g SMD							0.43 (0.123,	0.747)	0.0063*
non-tBRCAm	93 2	3.27 (22.101)	-10.70 (1.130)	60	29.86 (26.553)	-13.79 (1.514)	3.09 (-0.665,	6.844)	0.1060
Hedges' g SMD							0.27 (-0.052,	0.600)	0.0998
Int. p-value									0.5265
Age group									
<65 years	162 2	2.36 (21.052)	-8.02 (0.829)	89	25.91 (24.489)	-13.52 (1.204)	5.50 (2.617,	8.386)	0.0002*
Hedges' g SMD							0.51 (0.243,	0.768)	0.0002*
>=65 years	58 2	8.74 (25.637)	-12.96 (1.675)	30	34.26 (27.886)	-13.02 (2.482)	0.07 (-5.905,	6.041)	0.9820
Hedges' g SMD							0.01 (-0.436,	0.446)	0.9817
Int. p-value									0.0742
FIGO Stage (Dis	ease st	ate)							
III	157 2	5.65 (23.972)	-11.04 (0.875)	83	31.16 (24.583)	-14.64 (1.263)	3.59 (0.562,	6.627)	0.0204*
Hedges' g SMD							0.32 (0.054,	0.589)	0.0186*
IV	63 2	0.02 (17.724)	-4.35 (1.573)	36	20.76 (26.523)	-10.24 (2.388)	5.89 (0.199,	11.577)	0.0427*
Hedges' g SMD							0.44 (0.029,	0.858)	0.0360*
Int. p-value									0.6977
Region									
Europe	211 2	4.28 (22.715)	-9.53 (0.768)	113	28.49 (25.864)	-13.51 (1.131)	3.98 (1.290,	6.677)	0.0039*
Hedges' g SMD		,	,				0.35 (0.116,	0.577)	0.0032*
Japan	9	NC	NC	6	NC	NC	NC		NC
Int. p-value									NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.

SMD = standardised mean difference. * p<0.05.

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.6.4.7 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Other single items (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=25			Placebo + be		Difference	between g	groups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimate difference (9		p-value
ECOG performance	e stat	us at Baseline							
(0) Normal activity	163	22.43 (21.480)	-8.10 (0.871)	91	26.62 (26.366)	-12.61 (1.216)	4.51 (1.560	, 7.460)	0.0029*
Hedges' g SMD							0.40 (0.139	, 0.657)	0.0026*
(1) Restricted activity	53	28.88 (24.859)	-12.60 (1.531)	28	32.54 (22.425)	-13.34 (2.492)	0.74 (-5.080	, 6.556)	0.8014
Hedges' g SMD Int. p-value							0.06 (-0.396	, 0.520)	0.7924 0.7608
Baseline CA-125									
<=ULN	195	23.23 (22.221)	-9.12 (0.771)	106	28.51 (25.647)	-13.12 (1.117)	4.00 (1.328		0.0035*
Hedges' g SMD >ULN Int. p-value	25	NC	NC	13	NC	NC	0.36 (0.124 NC	, 0.600)	0.0029* NC NC
Histological gra	ade								
High grade		24.04 (22.468)	-9.23 (0.745)	119	28.01 (25.527)	-13.36 (1.084)	4.13 (1.538	, 6.718)	0.0019*
Hedges' g SMD Int. p-value							0.36 (0.139	, 0.588)	0.0015* ID
Cytoreductive su	ırgery	outcome							
No residue		23.83 (23.823)	-9.35 (0.879)	71	25.78 (24.020)	-13.80 (1.314)	4.45 (1.330	, 7.565)	0.0054*
Hedges' g SMD							0.42 (0.128	, 0.703)	0.0047*
Residue	71	24.88 (20.138)	-9.10 (1.384)	40	30.76 (27.766)	-14.48 (2.029)	•	, 10.264)	0.0313*
Hedges' g SMD Int. p-value							0.44 (0.050	, 0.834)	0.0272* 0.8901

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis. Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to

Table 2.6.4.7 PAOLA1: Summary of subgroup analysis of change from baseline in EORTC QLQ-OV28 Symptom scale/items: Other single items (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=25			Placebo + be (N=13		Difference between g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Timing of cytore	educti	ve surgery						
Upfront	130	26.71 (24.274)	-11.70 (0.875)	70	29.01 (25.174)	-15.05 (1.265)	3.36 (0.323, 6.391)	0.0303*
Hedges' g SMD							0.33 (0.036, 0.620)	0.0278*
Interval	83	20.21 (19.227)	-4.94 (1.325)	41	25.14 (25.969)	-12.71 (2.026)	7.77 (2.964, 12.571)	0.0018*
Hedges' g SMD							0.62 (0.241, 1.006)	0.0014*
Int. p-value								0.1624
Myriad tumour B	RCA mu	tation status						
tBRCAm	136	24.65 (22.502)	-9.41 (0.957)	67	28.73 (25.309)	-14.53 (1.458)	5.12 (1.678, 8.557)	0.0037*
Hedges' g SMD							0.45 (0.151, 0.742)	0.0031*
Non-tBRCAm	84	23.05 (22.512)	-8.92 (1.199)	52	27.08 (26.022)	-11.85 (1.636)	2.92 (-1.098, 6.948)	0.1527
Hedges' g SMD							0.26 (-0.090, 0.604)	0.1465
Int. p-value								0.8001
Status somatic 1	BRCA m	utations						
sBRCAm	17	14.71 (13.313)	-3.07 (1.967)	6	24.07 (22.884)	-7.72 (3.690)	4.66 (-4.046, 13.360)	0.2789
Hedges' g SMD							0.54 (-0.408, 1.484)	0.2650
gBRCAm	57	21.44 (22.529)	-5.12 (1.487)	27	27.98 (24.325)	-14.27 (2.361)	9.14 (3.575, 14.711)	0.0016*
Hedges' g SMD							0.78 (0.309, 1.257)	0.0012*
Non-BRCAm	37	21.85 (22.578)	-5.28 (1.413)	22	20.83 (24.782)	-6.37 (2.020)	1.10 (-3.872, 6.063)	0.6588
Hedges' g SMD							0.12 (-0.407, 0.649)	0.6533
Int. p-value								0.1525

SMD = standardised mean difference. * p<0.05.

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.7.1 PAOLA1: Summary of EQ-5D-5L results across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

					Res	ult		
Parameter	Group	Time Point	n	Mean	SD	Min	Median	Max
EQ-5D-5L Visual analogue scale	Olaparib + bevacizumab (N=255)	Baseline [a]	244	72.3	16.40	10	75.0	100
		Wk 12 (Day 85)	225	71.3	15.75	20	70.0	100
		Wk 24 (Day 169)	202	72.4	15.74	1	75.0	100
		Wk 36 (Day 253)	179	73.1	15.77	25	75.0	100
		Wk 48 (Day 337)	174	73.7	15.76	20	75.0	100
		Wk 60 (Day 421)	164	75.5	15.29	30	80.0	100
		Wk 72 (Day 505)	160	75.0	16.17	7	80.0	100
		Wk 84 (Day 589)	137	75.7	15.09	40	80.0	100
		Wk 96 (Day 673)	135	76.7	15.26	30	80.0	100
		Wk 108 (Day 757)	109	78.7	15.36	40	80.0	10
		Wk 120 (Day 841)	1	70.0	NC	70	70.0	70
		Wk 132 (Day 925)	1	75.0	NC	75	75.0	75
		Wk 144 (Day 1009)	1	75.0	NC	75	75.0	75
		Wk 156 (Day 1093)	1	75.0	NC	75	75.0	75
		End of Treatment	130	73.3	18.88	1	75.0	10
		30 day Follow-up	60	75.8	16.68	20	80.0	100
	Placebo + bevacizumab (N=132)	Baseline [a]	127	72.1	14.76	35	70.0	100
		Wk 12 (Day 85)	117	71.6	15.92	25	75.0	10
		Wk 24 (Day 169)	104	71.6	15.53	30	72.5	98
		Wk 36 (Day 253)	97	73.3	13.80	40	75.0	10
		Wk 48 (Day 337)	84	73.4	16.09	30	75.0	10
		Wk 60 (Day 421)	68	77.9	13.27	35	80.0	10
		Wk 72 (Day 505)	67	77.8	14.88	40	80.0	10
		Wk 84 (Day 589)	51	79.0	13.30	50	80.0	10
		Wk 96 (Day 673)	41	79.7	13.30	40	80.0	100
		Wk 108 (Day 757)	31	78.9	16.32	30	80.0	10
		End of Treatment	69	71.2	16.07	35	73.0	10
		30 day Follow-up	21	75.4	21.45	30	80.0	100

Table 2.7.2 PAOLA1: Summary of analysis of change from baseline in EQ-5D-5L VAS (mixed model for repeated measures)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + be		Placebo + bev		Difference be	etween g	roups
Timepoint	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95%	CI)	p-value
Wk 12 (Day 85)	204 72.45 (16.434)	-2.14 (0.889)	114 72.83 (14.510)	-1.27 (1.189)	-0.87 (-3.787,	2.054)	0.5599
Wk 24 (Day 169)	188 72.60 (16.816)	-0.43 (0.943)	103 71.53 (14.766)	-0.68 (1.273)	0.25 (-2.868,	3.368)	0.8750
Wk 36 (Day 253)	168 72.40 (16.964)	0.03 (1.013)	94 72.00 (14.538)	-0.10 (1.362)	0.13 (-3.213,	3.468)	0.9402
Wk 48 (Day 337)	163 72.35 (17.091)	0.70 (1.066)	78 73.50 (14.371)	0.25 (1.501)	0.45 (-3.176,	4.071)	0.8080
Wk 60 (Day 421)	154 71.66 (17.141)	3.12 (0.994)	66 74.68 (13.373)	3.00 (1.448)	0.12 (-3.343,	3.577)	0.9470
Wk 72 (Day 505)	149 72.14 (17.471)	3.40 (1.091)	64 74.58 (13.520)	3.32 (1.614)	0.08 (-3.759,	3.914)	0.9684
Wk 84 (Day 589)	130 70.90 (17.617)	3.56 (1.014)	49 75.78 (12.435)	3.07 (1.532)	0.49 (-3.140,	4.111)	0.7923
Wk 96 (Day 673)	127 71.65 (17.883)	3.51 (1.128)	39 75.33 (12.848)	3.59 (1.808)	-0.08 (-4.284,	4.122)	0.9699
Average over all	217 72.62 (16.458)	1.47 (0.791)	121 72.34 (14.661)	1.40 (1.104)	0.07 (-2.604,	2.743)	0.9592
Hedges' g SMD					0.01 (-0.217,	0.228)	0.9587

SMD = standardised mean difference. * p<0.05.

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Figure 2.7.3.1 PAOLA1: Mean (+/- SD) score for EQ-5D-5L Visual analogue scale across timepoints, by treatment group Full Analysis Set, HRD[42] positive, DCO 22MAR2020

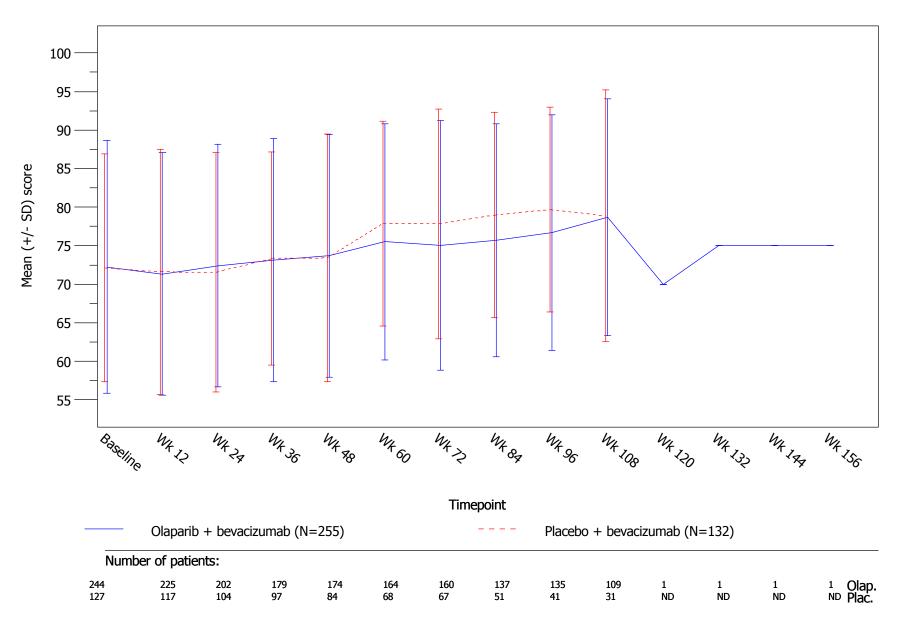


Table 2.7.4.1 PAOLA1: Summary of subgroup analysis of change from baseline in EQ-5D-5L Visual analogue scale (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be (N=255			Placebo + bev		Difference between g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% CI)	p-value
Rivet line tweet		out some (TIMC)						
First line treat NED [PDS]		71.85 (18.295)	4.05 (1.151)	1 E	72 42 /12 014\	2 04 / 1 E06\	0 21 / 2 670 4 002)	0.9144
Hedges' g SMD	19	/1.85 (18.295)	4.05 (1.151)	45	72.42 (13.814)	3.84 (1.586)	0.21 (-3.670, 4.092) 0.02 (-0.346, 0.386)	0.9144
NED/CR [IDS]	60.5	72.87 (14.426)	-1.22 (1.597)	2.2	72.73 (15.314)	2.61 (2.245)	, , , , , , , , , , , , , , , , , , , ,	0.1685
Hedges' q SMD	62	/2.8/ (14.426)	-1.22 (1.597)	33	/2./3 (15.314)	2.01 (2.245)	-3.83 (-9.311, 1.854) -0.30 (-0.724, 0.125)	0.1685
NED/CR [Chemo]	25 (50 46 (15 042)	2 07 / 1 065\	20	71 70 (10 740)	0 00 / 0 766)	, , , , , , , , , , , , , , , , , , , ,	
	35 (59.46 (15.942)	3.07 (1.965)	20	71.70 (12.749)	-0.09 (2.766)	3.16 (-3.664, 9.978)	0.3568
Hedges' g SMD	41 5	76 44 (15 064)	1 27 / 2 220)	2.2	70 17 (17 570)	F 00 / 2 020)	0.26 (-0.290, 0.813)	0.3527
PR	41	76.44 (15.864)	-1.37 (2.220)	23	72.17 (17.570)	-5.28 (3.238)	, , , , , , , , , , , , , , , , , , , ,	0.3242
Hedges' g SMD Int. p-value							0.26 (-0.250, 0.775)	0.3154
inc. p-value								0.2296
Screening labora	atory	tBRCA status (I	VRS)					
tBRCAm	128 7	70.83 (16.684)	2.00 (0.978)	60	71.92 (14.960)	2.50 (1.503)	-0.50 (-4.042, 3.037)	0.7796
Hedges' g SMD							-0.04 (-0.351, 0.262)	0.7761
non-tBRCAm	89 7	75.20 (15.866)	0.93 (1.303)	61	72.75 (14.474)	-0.35 (1.624)	1.28 (-2.847, 5.404)	0.5408
Hedges' g SMD							0.10 (-0.224, 0.428)	0.5392
Int. p-value								0.6352
First line treat	ment.	outcome (eCRF)						
NED [PDS]		70.86 (18.645)	4.21 (1.218)	44	72.48 (13.589)	3.58 (1.646)	0.63 (-3.430, 4.688)	0.7595
Hedges' g SMD		(======)	()		(==:==)/		0.06 (-0.312, 0.428)	0.7590
NED/CR [IDS]	61 5	74.48 (15.443)	-0.91 (1.598)	28	72.68 (16.414)	1.90 (2.439)	-2.82 (-8.627, 2.989)	0.3367
Hedges' g SMD		(=====)			(/	(= . 10)	-0.22 (-0.670, 0.227)	0.3331
NED/CR [Chemo]	32 5	70.81 (16.119)	2.57 (2.115)	17	71.41 (13.491)	-1.79 (3.151)	4.36 (-3.296, 12.010)	0.2569
Hedges' g SMD	32 .	(=0.11)	,,	_,	(_ 0 , _ , _)	(3.131)	0.35 (-0.245, 0.940)	0.2500
PR	44 7	74.86 (14.245)	-1.38 (1.947)	31	72.90 (15.640)	-2.27 (2.495)	0.89 (-5.431, 7.212)	0.7794
Hedges' q SMD			(1, /	5 4	(13 . 3 10)	(2.199)	0.07 (-0.394, 0.526)	0.7781
Int. p-value							3.320	0.4238

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.7.4.1 PAOLA1: Summary of subgroup analysis of change from baseline in EQ-5D-5L Visual analogue scale (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Ol	aparib + bev N=255)			Placebo + bev (N=132		Difference	between g	roups
Subgroup		n (SD) at eline [b]	Mean (SE) Change from baseline		Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (99		p-value
Screening labor	atory tBRC	A status (e0	CRF)						
tBRCAm	125 70.69	9 (16.783)	1.92 (0.994)	61	71.72 (14.913)	2.46 (1.497)	-0.54 (-4.088,	3.008)	0.7643
Hedges' g SMD							-0.05 (-0.354,	0.259)	0.7608
non-tBRCAm	92 75.2	5 (15.716)	1.01 (1.268)	60	72.97 (14.500)	-0.21 (1.622)	1.22 (-2.863,	5.293)	0.5565
Hedges' g SMD							0.10 (-0.227,	0.424)	0.5545
Int. p-value									0.5441
Age group									
<65 years	159 71.9	4 (17.201)	1.37 (0.923)	92	72.38 (14.446)	2.41 (1.269)	-1.04 (-4.134,	2.053)	0.5081
Hedges' g SMD							-0.09 (-0.345,	0.169)	0.5036
>=65 years	58 74.48	3 (14.194)	1.77 (1.561)	29	72.21 (15.589)	-1.69 (2.288)	3.46 (-2.055,	8.968)	0.2156
Hedges' g SMD							0.28 (-0.163,	0.733)	0.2127
Int. p-value									0.3047
FIGO Stage (Dis	ease state)							
III	156 71.9	7 (16.804)	1.93 (0.920)	84	73.49 (14.576)	2.64 (1.295)	-0.71 (-3.837,	2.426)	0.6575
Hedges' g SMD							-0.06 (-0.326,	0.205)	0.6548
IV	61 74.28	3 (15.548)	0.40 (1.508)	37	69.73 (14.717)	-0.87 (2.117)	1.27 (-3.909,	6.446)	0.6275
Hedges' g SMD							0.10 (-0.306,	0.512)	0.6209
Int. p-value									0.4558
Region									
Europe	208 72.3	5 (16.609)	1.38 (0.820)	115	72.33 (14.602)	1.25 (1.155)	0.14 (-2.652,	2.924)	0.9236
Hedges' g SMD							0.01 (-0.216,	0.239)	0.9227
Japan	9	NC	NC	6	NC	NC	NC		NC
Int. p-value									NC

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.7.4.1 PAOLA1: Summary of subgroup analysis of change from baseline in EQ-5D-5L Visual analogue scale (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bo (N=25		Placebo + bev (N=132		Difference between groups			
Subgroup	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	n Mean (SD) at [a] Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95% C	CI) p-value		
ECOG performance	e status at Baseline							
(0) Normal	161 73.37 (16.688)	1.16 (0.909)	92 75.03 (13.996)	0.09 (1.234)	1.08 (-1.948, 4	.099) 0.4841		
activity								
Hedges' g SMD	50 50 00 (15 556)	0.01 (1.606)	00 50 55 (10 001)	5 04 (0 551)	0.09 (-0.164, 0	*		
(1) Restricted activity	52 70.90 (15.576)	2.91 (1.626)	28 63.57 (13.801)	7.24 (2.551)	-4.33 (-10.415, 1	.750) 0.1602		
Hedges' g SMD					-0.35 (-0.810, 0	.116) 0.1417		
Int. p-value					0.55 (0.010)	0.1574		
Baseline CA-125	value							
<=ULN	192 72.34 (16.405)	2.06 (0.816)	108 72.20 (14.791)	1.64 (1.134)	0.42 (-2.327, 3	.174) 0.7622		
Hedges' g SMD					0.04 (-0.199, 0	,		
>ULN Int. p-value	25 NC	NC	13 NC	NC	NC	NC NC		
Histological gra	ade							
High grade	217 72.62 (16.458)	1.47 (0.791)	121 72.34 (14.661)	1.40 (1.104)	0.07 (-2.604, 2	.743) 0.9592		
Hedges' g SMD					0.01 (-0.217, 0	.228) 0.9587		
Int. p-value						ID		
Cytoreductive s	urgery outcome							
No residue	141 72.33 (17.194)	2.12 (0.944)	73 72.32 (14.682)	3.06 (1.347)	-0.94 (-4.184, 2	.304) 0.5683		
Hedges' g SMD					-0.08 (-0.366, 0	.200) 0.5659		
Residue	68 72.65 (15.070)	0.48 (1.507)	40 73.10 (14.510)	0.64 (2.082)	-0.17 (-5.274, 4	.940) 0.9483		
Hedges' g SMD					-0.01 (-0.404, 0	.377) 0.9477		
Int. p-value						0.9815		

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

model within-subject error.
SMD = standardised mean difference. * p<0.05.</pre>

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[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Table 2.7.4.1 PAOLA1: Summary of subgroup analysis of change from baseline in EQ-5D-5L Visual analogue scale (mixed model for repeated measures) - average over all visits

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + be			Placebo + bev		Difference b	oetween g	roups
Subgroup	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	n [a]	Mean (SD) at Baseline [b]	Mean (SE) Change from baseline	Estimated difference (95		p-value
mining of autom									
Timing of cytore Upfront Hedges' g SMD		71.40 (17.408)	3.85 (0.980)	73	72.11 (14.391)	2.73 (1.320)	1.12 (-2.122, 0.10 (-0.188,		0.4958 0.4937
Interval Hedges' g SMD Int. p-value	83 '	74.00 (14.977)	-2.16 (1.365)	40	73.48 (15.011)	1.48 (2.064)	-3.65 (-8.549, -0.29 (-0.666,	•	0.1433 0.1381 0.1546
Myriad tumour BI	RCA mu	tation status							
tBRCAm Hedges' g SMD		71.28 (16.508)	1.71 (0.964)	68	71.76 (14.397)	2.78 (1.418)	-1.07 (-4.449, -0.09 (-0.386,	2.319) 0.198)	0.5354 0.5299
Non-tBRCAm Hedges' g SMD Int. p-value	83	74.80 (16.240)	1.07 (1.355)	53	73.08 (15.099)	-0.91 (1.753)	1.98 (-2.410, 0.16 (-0.187,		0.3729 0.3705 0.2959
Status somatic H	SRCA m	utations							
sBRCAm Hedges' g SMD		69.35 (16.871)	3.00 (2.521)	6	70.00 (24.290)	8.55 (4.315)	-5.54 (-15.957, -0.51 (-1.457,	4.870) 0.433)	0.2802 0.2881
gBRCAm Hedges' g SMD	56 '	72.36 (16.124)	-0.71 (1.401)	29	73.24 (11.544)	3.79 (2.070)	-4.50 (-9.483, -0.42 (-0.869,	0.481)	0.0759 0.0716
Non-BRCAm Hedges' g SMD Int. p-value	37	72.65 (15.469)	1.06 (1.915)	22	72.45 (13.996)	0.06 (2.643)	1.00 (-5.618, 0.08 (-0.445,	7.616)	0.7611 0.7590 0.0973

SMD = standardised mean difference. * p<0.05.

[[]a] = number of patients with change from baseline at each timepoint in the MMRM analysis.

[[]b] = mean baseline for patients with change from baseline at each timepoint in the MMRM analysis.

Calculated from a mixed model for repeated measures (MMRM) analysis of the change from baseline score with treatment, visit and treatment by visit interaction included as explanatory variables and the baseline score included as a covariate along with the baseline score by visit interaction. The study discontinuation visit and the safety follow-up visit are excluded and only visits where change from baseline is available for at least 25% of patients in both treatment arms are included in the analysis.

Treatment, visit and treatment by visit interaction are fixed effects in the model, an unstructured covariance matrix is used to model within-subject error.

Table 2.8.1 PAOLA1: Summary of compliance with EORTC QLQ-C30 questionnaire by planned visit Full Analysis Set, HRD[42] positive, DCO 22MAR2020

Group	Time Point [a]	Expected [b]	Received [c]	Evaluable [d]	Compliance rate (%) [e]	Evaluability rate (%) [f]
Olaparib + bevacizumab (N=255)	VISIT 1 (Day 1)	255	251	251	98.4	100.0
	VISIT 7 (Day 85)	232	210	210	90.5	100.0
	VISIT 11 (Day 169)	218	192	192	88.1	100.0
	VISIT 13 (Day 253)	203	174	174	85.7	100.0
	VISIT 15 (Day 337)	193	169	169	87.6	100.0
	VISIT 17 (Day 421)	179	155	155	86.6	100.0
	VISIT 19 (Day 505)	169	148	148	87.6	100.0
	VISIT 21 (Day 589)	156	131	131	84.0	100.0
	VISIT 23 (Day 673)	141	124	124	87.9	100.0
	VISIT 25 (Day 757)	3	1	1	33.3	100.0
	END OF TREATMENT [a]	250	198	198	79.2	100.0
	30 DAY FOLLOW-UP	224	48	48	21.4	100.0
Placebo + bevacizumab (N=132)	VISIT 1 (Day 1)	132	127	127	96.2	100.0
	VISIT 7 (Day 85)	127	110	110	86.6	100.0
	VISIT 11 (Day 169)	113	98	98	86.7	100.0
	VISIT 13 (Day 253)	102	86	86	84.3	100.0
	VISIT 15 (Day 337)	90	80	80	88.9	100.0
	VISIT 17 (Day 421)	74	65	65	87.8	100.0
	VISIT 19 (Day 505)	65	60	60	92.3	100.0
	VISIT 21 (Day 589)	52	44	44	84.6	100.0
	VISIT 23 (Day 673)	42	37	37	88.1	100.0
	VISIT 25 (Day 757)	2	2	2	100.0	100.0
	END OF TREATMENT [a]	127	98	98	77.2	100.0
	30 DAY FOLLOW-UP	118	19	19	16.1	100.0

[[]a] End of Treatment refers to discontinuation from treatment with olaparib or placebo. Date of study discontinuation is mapped to the nearest visit date to define the number of expected forms.

[[]b] Expected = number of patients still on study. Number of patients expected at baseline equals to full analysis set.
[c] Received = forms received back plus those recorded as: Subject too heavily affected by symptoms of disease under investigation.

[[]d] Evaluable = forms where at least one subscale that can be determined or where a reason for not completing the form is:

Subject too heavily affected by symptoms of disease under investigation.

[[]e] Compliance Rate = Evaluable/Expected * 100.

[[]f] Evaluability Rate = Evaluable/Received * 100.

Table 2.8.2 PAOLA1: Summary of compliance with EORTC QLQ-OV28 questionnaire by planned visit Full Analysis Set, HRD[42] positive, DCO 22MAR2020

Group	Time Point [a]	Expected [b]	Received [c]	Evaluable [d]	Compliance rate (%) [e]	Evaluability rate (%) [f]
Olaparib + bevacizumab (N=255)	VISIT 1 (Day 1)	255	251	251	98.4	100.0
	VISIT 7 (Day 85)	232	211	211	90.9	100.0
	VISIT 11 (Day 169)	218	193	193	88.5	100.0
	VISIT 13 (Day 253)	203	173	173	85.2	100.0
	VISIT 15 (Day 337)	193	169	169	87.6	100.0
	VISIT 17 (Day 421)	179	155	155	86.6	100.0
	VISIT 19 (Day 505)	169	147	147	87.0	100.0
	VISIT 21 (Day 589)	156	131	131	84.0	100.0
	VISIT 23 (Day 673)	141	125	125	88.7	100.0
	VISIT 25 (Day 757)	3	1	1	33.3	100.0
	END OF TREATMENT [a]	250	197	197	78.8	100.0
	30 DAY FOLLOW-UP	224	48	48	21.4	100.0
Placebo + bevacizumab (N=132)	VISIT 1 (Day 1)	132	126	126	95.5	100.0
	VISIT 7 (Day 85)	127	109	109	85.8	100.0
	VISIT 11 (Day 169)	113	98	98	86.7	100.0
	VISIT 13 (Day 253)	102	86	86	84.3	100.0
	VISIT 15 (Day 337)	90	80	80	88.9	100.0
	VISIT 17 (Day 421)	74	65	65	87.8	100.0
	VISIT 19 (Day 505)	65	60	60	92.3	100.0
	VISIT 21 (Day 589)	52	44	44	84.6	100.0
	VISIT 23 (Day 673)	42	37	37	88.1	100.0
	VISIT 25 (Day 757)	2	2	2	100.0	100.0
	END OF TREATMENT [a]	127	98	98	77.2	100.0
	30 DAY FOLLOW-UP	118	19	19	16.1	100.0

[[]a] End of Treatment refers to discontinuation from treatment with olaparib or placebo. Date of study discontinuation is mapped to the nearest visit date to define the number of expected forms.

[[]b] Expected = number of patients still on study. Number of patients expected at baseline equals to full analysis set.
[c] Received = forms received back plus those recorded as: Subject too heavily affected by symptoms of disease under investigation.

[[]d] Evaluable = forms where at least one subscale that can be determined or where a reason for not completing the form is:

Subject too heavily affected by symptoms of disease under investigation.

[[]e] Compliance Rate = Evaluable/Expected * 100.

[[]f] Evaluability Rate = Evaluable/Received * 100.

Table 2.8.3 PAOLA1: Summary of compliance with EQ-5D questionnaire by planned visit Full Analysis Set, HRD[42] positive, DCO 22MAR2020

Group	Time Point [a]	Expected [b]	Received [c]	Evaluable [d]	Compliance rate (%) [e]	Evaluability rate (%) [f]
Olaparib + bevacizumab (N=255)	VISIT 1 (Day 1)	255	247	247	96.9	100.0
	VISIT 7 (Day 85)	232	208	208	89.7	100.0
	VISIT 11 (Day 169)	218	192	192	88.1	100.0
	VISIT 13 (Day 253)	203	172	172	84.7	100.0
	VISIT 15 (Day 337)	193	167	167	86.5	100.0
	VISIT 17 (Day 421)	179	154	154	86.0	100.0
	VISIT 19 (Day 505)	169	148	148	87.6	100.0
	VISIT 21 (Day 589)	156	129	129	82.7	100.0
	VISIT 23 (Day 673)	141	123	123	87.2	100.0
	VISIT 25 (Day 757)	3	1	1	33.3	100.0
	END OF TREATMENT [a]	250	197	197	78.8	100.0
	30 DAY FOLLOW-UP	224	48	48	21.4	100.0
Placebo + bevacizumab (N=132)	VISIT 1 (Day 1)	132	127	127	96.2	100.0
	VISIT 7 (Day 85)	127	109	109	85.8	100.0
	VISIT 11 (Day 169)	113	97	97	85.8	100.0
	VISIT 13 (Day 253)	102	85	85	83.3	100.0
	VISIT 15 (Day 337)	90	80	80	88.9	100.0
	VISIT 17 (Day 421)	74	64	64	86.5	100.0
	VISIT 19 (Day 505)	65	59	59	90.8	100.0
	VISIT 21 (Day 589)	52	44	44	84.6	100.0
	VISIT 23 (Day 673)	42	37	37	88.1	100.0
	VISIT 25 (Day 757)	2	2	2	100.0	100.0
	END OF TREATMENT [a]	127	95	95	74.8	100.0
	30 DAY FOLLOW-UP	118	16	16	13.6	100.0

[[]a] End of Treatment refers to discontinuation from treatment with olaparib or placebo. Date of study discontinuation is mapped to the nearest visit date to define the number of expected forms.

[[]b] Expected = number of patients still on study. Number of patients expected at baseline equals to full analysis set.
[c] Received = forms received back plus those recorded as: Subject too heavily affected by symptoms of disease under investigation.

[[]d] Evaluable = forms where at least one subscale that can be determined or where a reason for not completing the form is:

Subject too heavily affected by symptoms of disease under investigation.

[[]e] Compliance Rate = Evaluable/Expected * 100.

[[]f] Evaluability Rate = Evaluable/Received * 100.

Table 3.1 PAOLA1: Summary of observation period (months) for adverse events Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=131)
All AE endpoints	n	255	131
_	Median	24.80	17.77
	Min	1.2	1.1
	Max	36.8	26.3
All AESI endpoints	n	255	131
	Median	38.51	36.76
	Min	8.9	5.3
	Max	55.6	53.8

Table 3.2.1 PAOLA1: Summary of analysis of time to first occurrence of adverse events Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	-	bevacizumab 255)		pevacizumab 131)			
	Number (%) of patient n with event	(95% CI)	Number (%) of patients n with events		Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AE	255 255 (100	0.2 (0.2, 0.3)	131 127 (96.9)	0.3 (0.2, 0.7)	1.43	1.15, 1.80	0.0023*
AE SOC: General disorders and administration site conditions	255 156 (61.2	5.6 (2.8,11.0)	131 57 (43.5)	NE (NE, NE)	1.70	1.26, 2.33	0.0006*
AE PT: Fatigue	255 141 (55.3	8.5 (3.5,15.3)	131 44 (33.6)	NE (NE, NE)	2.01	1.44, 2.86	<0.0001*
AE PT: Pyrexia	255 16 (6.3	NE (NE, NE)	131 4 (3.1)	NE (NE, NE)	2.12	0.77, 7.46	0.1717
AE PT: Oedema	255 8 (3.1	NE (NE, NE)	131 4 (3.1)	NE (NE, NE)	0.89	0.28, 3.39	0.8579
AE PT: Oedema peripheral	255 15 (5.9	NE (NE, NE)	131 7 (5.3)	NE (NE, NE)	1.01	0.42, 2.67	0.9846
AE PT: Mucosal inflammation	255 15 (5.9	NE (NE, NE)	131 4 (3.1)	NE (NE, NE)	1.95	0.70, 6.89	0.2285
AE PT: Pain	255 7 (2.7	NE (NE, NE)	131 4 (3.1)	NE (NE, NE)	0.79	0.24, 3.04	0.7052
AE SOC: Eye disorders	255 9 (3.5	NE (NE, NE)	131 4 (3.1)	NE (NE, NE)	0.89	0.28, 3.32	0.8478
AE SOC: Surgical and medical procedures	255 10 (3.9	NE (NE, NE)	131 3 (2.3)	NE (NE, NE)	1.69	0.51, 7.56	0.4248
AE SOC: Endocrine disorders	255 4 (1.6	NE (NE, NE)	131 9 (6.9)	NE (NE, NE)	0.20	0.05, 0.62	0.0033*

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.1 PAOLA1: Summary of analysis of time to first occurrence of adverse events Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2				Placebo + be (N=1)				
		Number (%) of patients with events	Median ti (95% CI) (months) [)	(Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AE SOC: Respiratory, thoracic and mediastinal disorders	255	62 (24.3)	NE (NE,	NE)	131	28 (21.4)	NE (NE, NE)	1.04	0.67, 1.65	0.8671
AE PT: Dyspnoea	255	22 (8.6)	NE (NE,	NE)	131	3 (2.3)	NE (NE, NE)	4.05	1.40, 17.15	0.0142*
AE PT: Epistaxis	255	18 (7.1)	NE (NE,	NE)	131	7 (5.3)	NE (NE, NE)	1.28	0.56, 3.31	0.5776
AE PT: Cough	255	11 (4.3)	NE (NE,	NE)	131	6 (4.6)	NE (NE, NE)	0.87	0.32, 2.58	0.7899
AE PT: Rhinorrhoea	255	5 (2.0)	NE (NE,	NE)	131	3 (2.3)	NE (NE, NE)	0.82	0.20, 4.00	0.7811
AE PT: Oropharyngeal pain	255	4 (1.6)	NE (NE,	NE)	131	4 (3.1)	NE (NE, NE)	0.47	0.11, 2.01	0.2793
AE SOC: Skin and subcutaneous tissue disorders	255	44 (17.3)	NE (NE,	NE)	131	19 (14.5)	NE (NE, NE)	1.16	0.68, 2.04	0.5923
AE PT: Alopecia	255	8 (3.1)	NE (NE,	NE)	131	2 (1.5)	NE (NE, NE)	2.02	0.50, 13.47	0.3661
AE PT: Rash	255	10 (3.9)	NE (NE,	NE)	131	7 (5.3)	NE (NE, NE)	0.76	0.29, 2.11	0.5832
AE PT: Erythema	255	8 (3.1)	NE (NE,	NE)	131	0	NE (NE, NE)	NC	NC	0.0569
AE PT: Pruritus	255	5 (2.0)	NE (NE,	NE)	131	3 (2.3)	NE (NE, NE)	0.81	0.20, 3.98	0.7736
AE SOC: Renal and urinary disorders	255	27 (10.6)	NE (NE,	NE)	131	24 (18.3)	NE (NE, NE)	0.50	0.29, 0.88	0.0133*
AE PT: Proteinuria	255	19 (7.5)	NE (NE,	NE)	131	19 (14.5)	NE (NE, NE)	0.46	0.24, 0.89	0.0163*

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.
[c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.1 PAOLA1: Summary of analysis of time to first occurrence of adverse events Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	_	bevacizumab =255)		pevacizumab 131)			
	Number (%) of patient n with event	(95% CI)	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AE SOC: Blood and lymphatic system disorders	255 148 (58.0	4.8 (2.8,12.3)	131 41 (31.3)	NE (NE, NE)	2.43	1.73, 3.49	<0.0001*
AE PT: Anaemia	255 102 (40.0	NE (NE, NE)	131 12 (9.2)	NE (NE, NE)	5.43	3.10, 10.43	<0.0001*
AE PT: Leukopenia	255 46 (18.0	NE (NE, NE)	131 11 (8.4)	NE (NE, NE)	2.34	1.26, 4.78	0.0097*
AE PT: Lymphopenia	255 60 (23.5	NE (NE, NE)	131 10 (7.6)	NE (NE, NE)	3.36	1.79, 6.99	0.0002*
AE PT: Neutropenia	255 30 (11.8	NE (NE, NE)	131 15 (11.5)	NE (NE, NE)	0.96	0.52, 1.84	0.8855
AE PT: Thrombocytopenia	255 12 (4.7	NE (NE, NE)	131 3 (2.3)	NE (NE, NE)	1.88	0.59, 8.31	0.3264
AE SOC: Gastrointestinal disorders	255 189 (74.1	0.8 (0.5, 1.9)	131 83 (63.4)	6.9 (3.4,10.6)	1.56	1.20, 2.04	0.0012*
AE PT: Abdominal pain	255 56 (22.0	NE (NE, NE)	131 32 (24.4)	NE (NE, NE)	0.84	0.55, 1.32	0.4401
AE PT: Intestinal obstruction	255 5 (2.0	NE (NE, NE)	131 1 (0.8)	NE (NE, NE)	1.83	0.29, 35.00	0.5771
AE PT: Diarrhoea	255 50 (19.6	NE (NE, NE)	131 25 (19.1)	NE (NE, NE)	1.01	0.63, 1.67	0.9506
AE PT: Dyspepsia	255 12 (4.7	NE (NE, NE)	131 3 (2.3)	NE (NE, NE)	1.78	0.56, 7.87	0.3692
AE PT: Vomiting	255 54 (21.2	NE (NE, NE)	131 16 (12.2)	NE (NE, NE)	1.77	1.03, 3.20	0.0438*
AE PT: Haemorrhoids	255 7 (2.7	NE (NE, NE)	131 4 (3.1)	NE (NE, NE)	0.84	0.25, 3.24	0.7851

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.1 PAOLA1: Summary of analysis of time to first occurrence of adverse events Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1				Placebo + be (N=13				
	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		(Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AE PT: Constipation	255	28 (11.0)	NE (NE, N	IE)	131	15 (11.5)	NE (NE, NE)	0.87	0.47, 1.69	0.6718
AE PT: Abdominal pain upper	255	10 (3.9)	NE (NE, N	1E)	131	4 (3.1)	NE (NE, NE)	1.17	0.39, 4.26	0.7958
AE PT: Stomatitis	255	12 (4.7)	NE (NE, N	IE)	131	2 (1.5)	NE (NE, NE)	3.35	0.91, 21.59	0.0951
AE PT: Subileus	255	2 (0.8)	NE (NE, N	IE)	131	2 (1.5)	NE (NE, NE)	0.60	0.07, 5.12	0.6125
AE PT: Nausea	255	144 (56.5)	2.9 (0.8,14.	5)	131	30 (22.9)	NE (NE, NE)	3.38	2.30, 5.13	<0.0001*
AE PT: Gingival bleeding	255	9 (3.5)	NE (NE, N	IE)	131	2 (1.5)	NE (NE, NE)	2.38	0.60, 15.78	0.2596
AE PT: Toothache	255	6 (2.4)	NE (NE, N	IE)	131	3 (2.3)	NE (NE, NE)	1.04	0.27, 5.02	0.9527
AE SOC: Immune system disorders	255	7 (2.7)	NE (NE, N	1E)	131	1 (0.8)	NE (NE, NE)	3.67	0.64, 69.24	0.1965
AE SOC: Nervous system disorders	255	87 (34.1)	NE (NE, N	1E)	131	32 (24.4)	NE (NE, NE)	1.48	0.99, 2.25	0.0607
AE PT: Dysgeusia	255	23 (9.0)	NE (NE, N	IE)	131	2 (1.5)	NE (NE, NE)	6.19	1.82, 38.62	0.0049*
AE PT: Headache	255	39 (15.3)	NE (NE, N	IE)	131	22 (16.8)	NE (NE, NE)	0.89	0.53, 1.53	0.6675
AE PT: Neuropathy peripheral	255	22 (8.6)	NE (NE, N	1E)	131	5 (3.8)	NE (NE, NE)	2.45	0.997, 7.33	0.0637
AE PT: Polyneuropathy	255	8 (3.1)	NE (NE, N	IE)	131	2 (1.5)	NE (NE, NE)	2.10	0.52, 13.95	0.3386

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.1 PAOLA1: Summary of analysis of time to first occurrence of adverse events Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1				bevacizumab 131)			
	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AE PT: Dizziness	255	8 (3.1)	NE (NE, NE)	131	3 (2.3)	NE (NE, NE)	1.30	0.37, 5.97	0.7019
AE SOC: Ear and labyrinth disorders	255	10 (3.9)	NE (NE, NE)	131	8 (6.1)	NE (NE, NE)	0.59	0.23, 1.56	0.2663
AE PT: Vertigo	255	6 (2.4)	NE (NE, NE)	131	6 (4.6)	NE (NE, NE)	0.48	0.15, 1.56	0.2022
AE SOC: Vascular disorders	255	137 (53.7)	8.3 (5.8,14.9)	131	82 (62.6)	4.9 (3.4, 8.4)	0.81	0.61, 1.07	0.1305
AE PT: Hot flush	255	7 (2.7)	NE (NE, NE)	131	3 (2.3)	NE (NE, NE)	1.08	0.30, 5.04	0.9058
AE PT: Hypertension	255	122 (47.8)	14.1 (8.3, NE)	131	78 (59.5)	5.4 (3.3,11.0)	0.72	0.54, 0.97	0.0301*
AE SOC: Cardiac disorders	255	7 (2.7)	NE (NE, NE)	131	7 (5.3)	NE (NE, NE)	0.43	0.14, 1.29	0.1140
AE SOC: Infections and infestations	255	128 (50.2)	15.0 (11.0,21.5)	131	63 (48.1)	16.4 (9.0, NE)	0.98	0.72, 1.34	0.8979
AE PT: Bronchitis	255	13 (5.1)	NE (NE, NE)	131	3 (2.3)	NE (NE, NE)	2.27	0.73, 9.92	0.1915
AE PT: Gastroenteritis	255	13 (5.1)	NE (NE, NE)	131	0	NE (NE, NE)	NC	NC	0.0119*
AE PT: Gingivitis	255	7 (2.7)	NE (NE, NE)	131	2 (1.5)	NE (NE, NE)	1.87	0.44, 12.68	0.4343
AE PT: Influenza	255	8 (3.1)	NE (NE, NE)	131	1 (0.8)	NE (NE, NE)	3.38	0.62, 62.93	0.2231
AE PT: Urinary tract infection	255	41 (16.1)	NE (NE, NE)	131	12 (9.2)	NE (NE, NE)	1.77	0.95, 3.55	0.0811

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.1 PAOLA1: Summary of analysis of time to first occurrence of adverse events Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	-	bevacizumab 255)	Placebo + b (N=1				
	Number (%) of patients n with events		Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AE PT: Infection	255 9 (3.5)	NE (NE, NE)	131 1 (0.8)	NE (NE, NE)	4.68	0.87, 86.59	0.1085
AE PT: Nasopharyngitis	255 15 (5.9)	NE (NE, NE)	131 10 (7.6)	NE (NE, NE)	0.66	0.30, 1.53	0.3109
AE PT: Pharyngitis	255 8 (3.1)	NE (NE, NE)	131 4 (3.1)	NE (NE, NE)	0.88	0.28, 3.35	0.8428
AE PT: Rhinitis	255 10 (3.9)	NE (NE, NE)	131 4 (3.1)	NE (NE, NE)	1.17	0.38, 4.30	0.7973
AE PT: Tooth abscess	255 8 (3.1)	NE (NE, NE)	131 4 (3.1)	NE (NE, NE)	1.08	0.34, 4.07	0.9033
AE PT: Tooth infection	255 7 (2.7)	NE (NE, NE)	131 2 (1.5)	NE (NE, NE)	1.57	0.38, 10.55	0.5740
AE PT: Cystitis	255 11 (4.3)	NE (NE, NE)	131 9 (6.9)	NE (NE, NE)	0.62	0.25, 1.55	0.2851
AE SOC: Psychiatric disorders	255 21 (8.2)	NE (NE, NE)	131 13 (9.9)	NE (NE, NE)	0.77	0.39, 1.59	0.4585
AE PT: Anxiety	255 7 (2.7)	NE (NE, NE)	131 5 (3.8)	NE (NE, NE)	0.67	0.21, 2.29	0.4979
AE PT: Depression	255 6 (2.4)	NE (NE, NE)	131 1 (0.8)	NE (NE, NE)	2.62	0.44, 49.87	0.3601
AE PT: Sleep disorder	255 7 (2.7)	NE (NE, NE)	131 7 (5.3)	NE (NE, NE)	0.50	0.17, 1.49	0.1968
AE SOC: Musculoskeletal and connective tissue disorders	255 106 (41.6)	NE (NE, NE)	131 57 (43.5)	NE (NE, NE)	0.88	0.64, 1.22	0.4318
AE PT: Arthralgia	255 64 (25.1)	NE (NE, NE)	131 30 (22.9)	NE (NE, NE)	1.08	0.70, 1.70	0.7283

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.1 PAOLA1: Summary of analysis of time to first occurrence of adverse events Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b				Placebo + be				
		Number (%) of patients with events	Median tir (95% CI) (months) [Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AE PT: Muscle spasms	255	11 (4.3)	NE (NE,	NE)	131	8 (6.1)	NE (NE, NE)	0.62	0.25, 1.62	0.3068
AE PT: Myalgia	255	21 (8.2)	NE (NE,	NE)	131	7 (5.3)	NE (NE, NE)	1.60	0.71, 4.09	0.2795
AE PT: Neck pain	255	4 (1.6)	NE (NE,	NE)	131	5 (3.8)	NE (NE, NE)	0.42	0.10, 1.60	0.1864
AE PT: Back pain	255	17 (6.7)	NE (NE,	NE)	131	8 (6.1)	NE (NE, NE)	0.92	0.40, 2.28	0.8429
AE PT: Pain in extremity	255	17 (6.7)	NE (NE,	NE)	131	8 (6.1)	NE (NE, NE)	1.16	0.51, 2.87	0.7297
AE PT: Musculoskeletal pain	255	9 (3.5)	NE (NE,	NE)	131	9 (6.9)	NE (NE, NE)	0.53	0.21, 1.37	0.1739
AE SOC: Metabolism and nutrition disorders	255	30 (11.8)	NE (NE,	NE)	131	9 (6.9)	NE (NE, NE)	1.51	0.74, 3.39	0.2790
AE PT: Decreased appetite	255	23 (9.0)	NE (NE,	NE)	131	4 (3.1)	NE (NE, NE)	2.74	1.05, 9.36	0.0543
AE SOC: Investigations	255	73 (28.6)	NE (NE,	NE)	131	29 (22.1)	NE (NE, NE)	1.31	0.86, 2.06	0.2213
AE PT: Weight increased	255	11 (4.3)	NE (NE,	NE)	131	10 (7.6)	NE (NE, NE)	0.53	0.22, 1.30	0.1504
AE PT: Blood creatinine increased	255	13 (5.1)	NE (NE,	NE)	131	1 (0.8)	NE (NE, NE)	6.54	1.28,119.35	0.0386*
AE PT: White blood cell count decreased	255	8 (3.1)	NE (NE,	NE)	131	2 (1.5)	NE (NE, NE)	1.64	0.41, 10.93	0.5261

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.1 PAOLA1: Summary of analysis of time to first occurrence of adverse events Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	_	bevacizumab 255)	Placebo + b (N=1				2-sided [b] p-value[c]
	Number (%) of patients n with events	,	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	
AE PT: Lymphocyte count decreased	255 7 (2.7)	NE (NE, NE)	131 1 (0.8)	NE (NE, NE)	3.50	0.62, 65.62	0.2131
AE PT: Neutrophil count decreased	255 20 (7.8)	NE (NE, NE)	131 6 (4.6)	NE (NE, NE)	1.72	0.73, 4.75	0.2477
AE PT: Platelet count decreased	255 5 (2.0)	NE (NE, NE)	131 1 (0.8)	NE (NE, NE)	3.06	0.49, 58.98	0.2847
AE SOC: Injury, poisoning and procedural complications	255 15 (5.9)	NE (NE, NE)	131 9 (6.9)	NE (NE, NE)	0.72	0.31, 1.73	0.4350

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.2 PAOLA1: Summary of analysis of time to first occurrence of serious adverse events Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b				Placebo + be (N=1)				
		Number (%) of patients with events	Median to (95% CI (months))		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
SAE	255	73 (28.6)	NE (NE,	NE)	131	45 (34.4)	NE (NE, NE)	0.75	0.52, 1.10	0.1332
SAE SOC: Respiratory, thoracic and mediastinal disorders	255	7 (2.7)	NE (NE,	NE)	131	1 (0.8)	NE (NE, NE)	4.27	0.75, 80.04	0.1411
SAE SOC: Blood and lymphatic system disorders	255	17 (6.7)	NE (NE,	NE)	131	1 (0.8)	NE (NE, NE)	8.49	1.74,153.16	0.0127*
SAE PT: Anaemia	255	13 (5.1)	NE (NE,	NE)	131	1 (0.8)	NE (NE, NE)	6.96	1.38,126.51	0.0300*
SAE SOC: Gastrointestinal disorders	255	11 (4.3)	NE (NE,	NE)	131	10 (7.6)	NE (NE, NE)	0.48	0.20, 1.17	0.0936
SAE SOC: Vascular disorders	255	24 (9.4)	NE (NE,	NE)	131	16 (12.2)	NE (NE, NE)	0.72	0.38, 1.38	0.3039
SAE PT: Hypertension	255	20 (7.8)	NE (NE,	NE)	131	16 (12.2)	NE (NE, NE)	0.59	0.30, 1.16	0.1151
SAE SOC: Infections and infestations	255	12 (4.7)	NE (NE,	NE)	131	9 (6.9)	NE (NE, NE)	0.64	0.27, 1.58	0.3126

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.3 PAOLA1: Summary of analysis of time to first occurrence of adverse event leading to treatment discontinuation

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)		Placebo + b (N=1				
	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95	95% CI [b]	2-sided p-value [c]
AE leading to discontinuation of treatment	255 50 (19.6)	NE (NE, NE)	131 8 (6.1)	NE (NE, NE)	3.14	1.57, 7.18	0.0017*

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 23.0.

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.4 PAOLA1: Summary of analysis of time to first occurrence of severe adverse events with max. CTCAE grade >=3 including grade 5
Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)				Placebo + bevacizumab (N=131)						
		Number (%) of patients with events	Median t (95% CI (months)	.)		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AE max CTCAE grade>=3	255	147 (57.6)	8.6 (5.6,	15.3)	131	65 (49.6)	16.7 (6.6,	NE)	1.20	0.90, 1.63	0.2205
AE G>=3 SOC: General disorders and administration site conditions	255	15 (5.9)	NE (NE,	NE)	131	1 (0.8)	NE (NE,	NE)	7.83	1.58,141.83	0.0185*
AE G>=3 PT: Fatigue	255	14 (5.5)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	0.0065*
AE G>=3 SOC: Respiratory, thoracic and mediastinal disorders	255	6 (2.4)	NE (NE,	NE)	131	2 (1.5)	NE (NE,	NE)	1.78	0.41, 12.18	0.4769
AE G>=3 SOC: Renal and urinary disorders	255	3 (1.2)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	0.2780
AE G>=3 SOC: Blood and lymphatic system disorders	255	71 (27.8)	NE (NE,	NE)	131	5 (3.8)	NE (NE,	NE)	8.16	3.64, 23.32	<0.0001*
AE G>=3 PT: Anaemia	255	47 (18.4)	NE (NE,	NE)	131	1 (0.8)	NE (NE,	NE)	27.85	6.08,493.74	<0.0001*
AE G>=3 PT: Leukopenia	255	6 (2.4)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	0.0945
AE G>=3 PT: Lymphopenia	255	19 (7.5)	NE (NE,	NE)	131	3 (2.3)	NE (NE,	NE)	2.97	1.01, 12.69	0.0670
AE G>=3 PT: Neutropenia	255	12 (4.7)	NE (NE,	NE)	131	1 (0.8)	NE (NE,	NE)	5.91	1.15,107.89	0.0537

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.4 PAOLAl: Summary of analysis of time to first occurrence of severe adverse events with max. CTCAE grade >=3 including grade 5

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)					Placebo + be (N=1				_
		Number (%) of patients with events	Median t (95% CI (months))		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AE G>=3 SOC: Gastrointestinal disorders	255	30 (11.8)	NE (NE	NE)	131	12 (9.2)	NE (NE, NE) 1.20	0.63, 2.45	0.5931
AE G>=3 PT: Diarrhoea	255	7 (2.7)	NE (NE	NE)	131	3 (2.3)	NE (NE, NE) 1.18	0.33, 5.54	0.8073
AE G>=3 PT: Nausea	255	8 (3.1)	NE (NE	NE)	131	0	NE (NE, NE) NC	NC	0.0590
AE G>=3 SOC: Nervous system disorders	255	7 (2.7)	NE (NE	NE)	131	4 (3.1)	NE (NE, NE	0.74	0.22, 2.91	0.6454
AE G>=3 SOC: Vascular disorders	255	48 (18.8)	NE (NE	NE)	131	42 (32.1)	NE (NE, NE	0.51	0.33, 0.78	0.0013*
AE G>=3 PT: Hypertension	255	45 (17.6)	NE (NE	NE)	131	42 (32.1)	NE (NE, NE	0.47	0.30, 0.72	0.0004*
AE G>=3 SOC: Infections and infestations	255	14 (5.5)	NE (NE	NE)	131	10 (7.6)	NE (NE, NE	0.62	0.28, 1.46	0.2559
AE G>=3 SOC: Investigations	255	16 (6.3)	NE (NE	NE)	131	5 (3.8)	NE (NE, NE) 1.72	0.67, 5.30	0.2888
AE G>=3 PT: Neutrophil count decreased	255	6 (2.4)	NE (NE	NE)	131	1 (0.8)	NE (NE, NE) 3.12	0.53, 59.15	0.2692

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.
[c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.5 PAOLA1: Summary of analysis of time to first occurrence of non-severe adverse events with max. CTCAE grade 1 or 2

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + b (N=2		Placebo + b (N=1				
	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AE max CTCAE grade=1 or 2	255 108 (42.4)	NE (NE, NE)	131 62 (47.3)	NE (NE, NE)	0.93	0.68, 1.28	0.6317

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 23.0.

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Figure 3.3.1 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

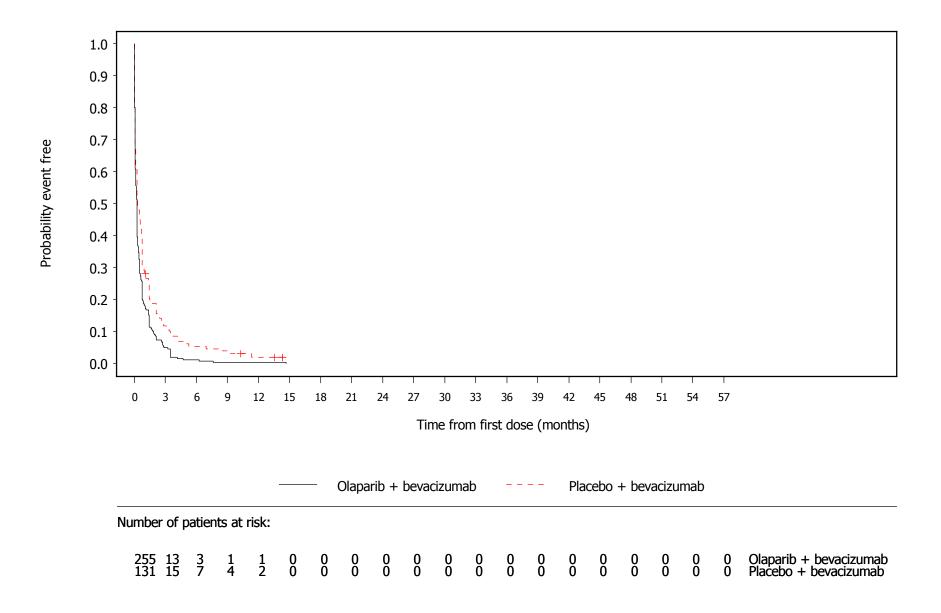


Figure 3.3.2 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: General disorders and administration site conditions Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

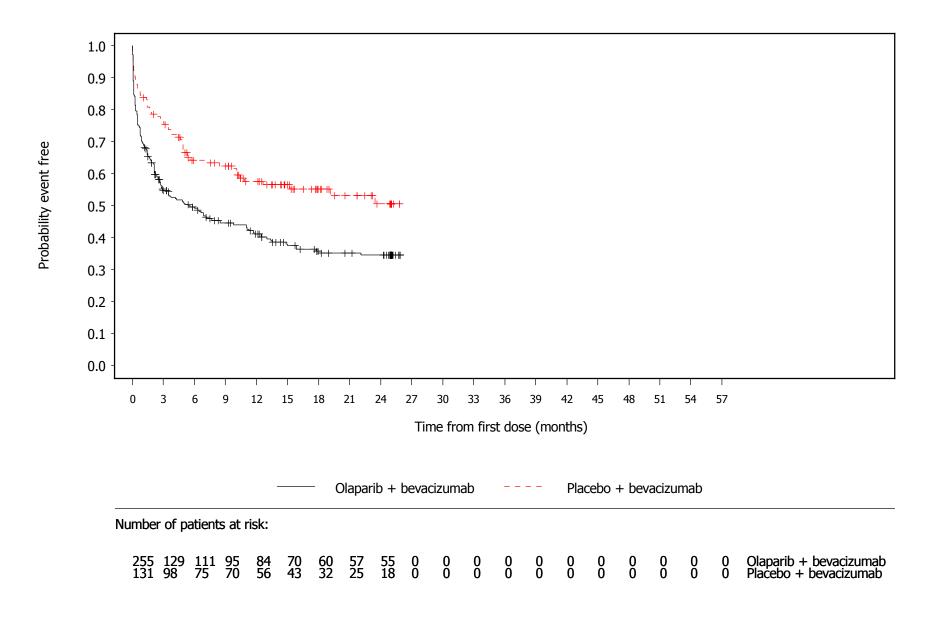


Figure 3.3.3 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Fatigue Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

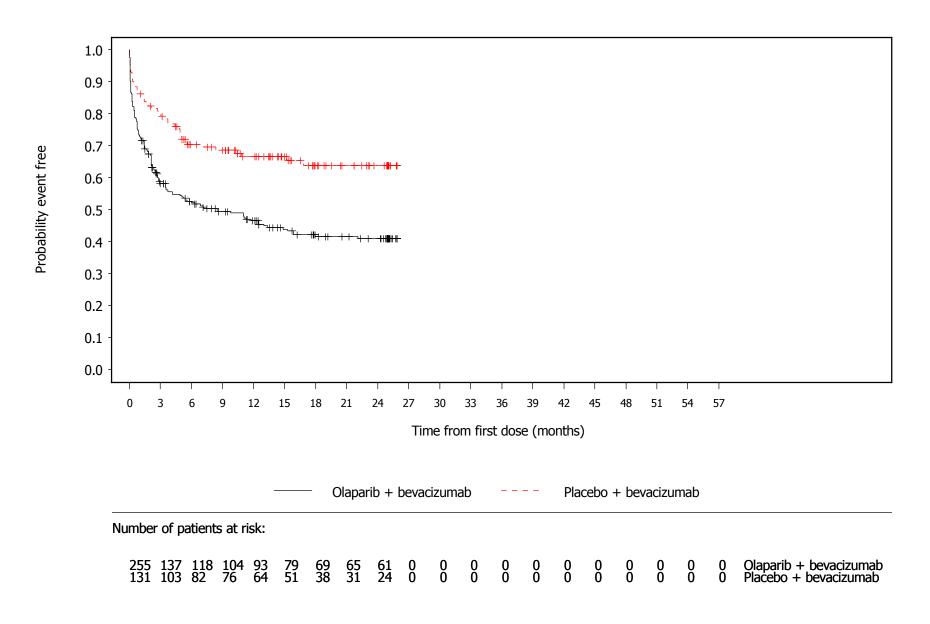


Figure 3.3.4 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Pyrexia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

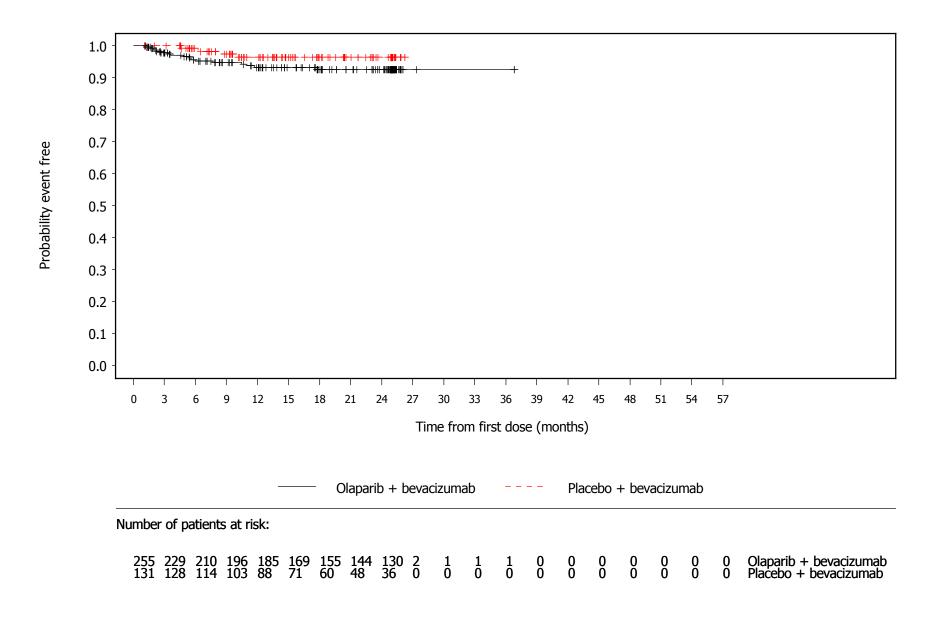


Figure 3.3.5 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Oedema Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

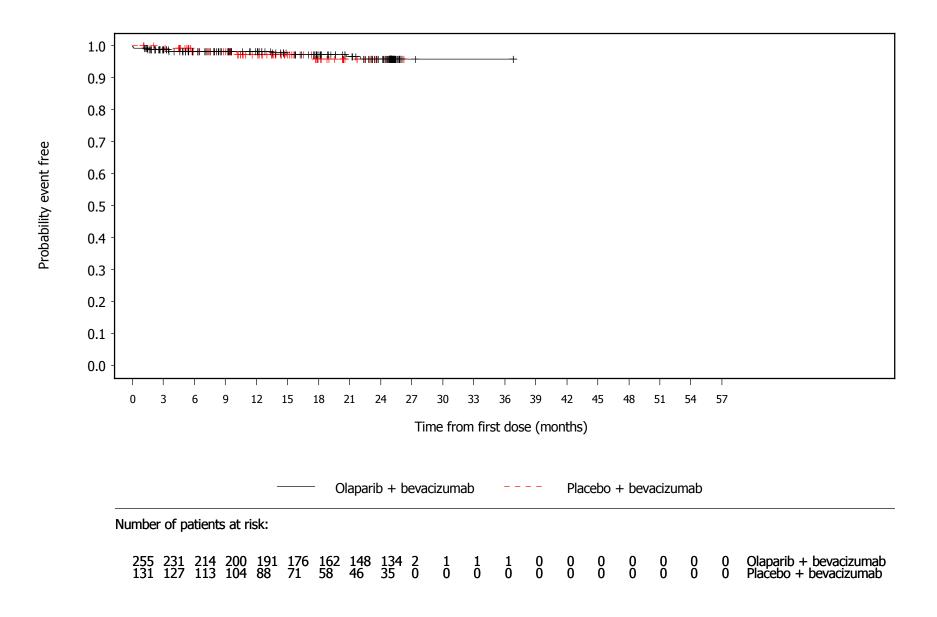


Figure 3.3.6 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Oedema peripheral Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

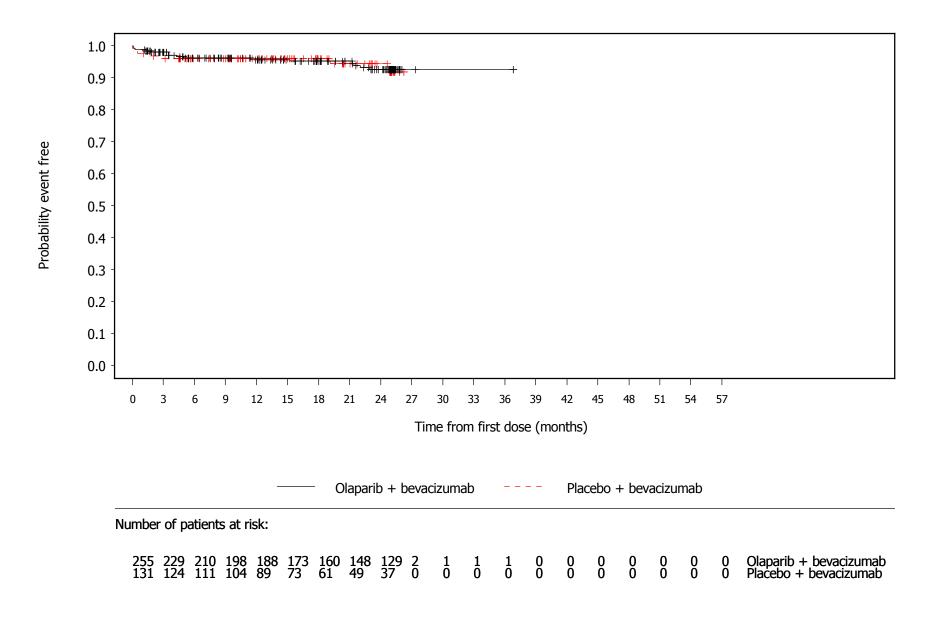


Figure 3.3.7 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Mucosal inflammation Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

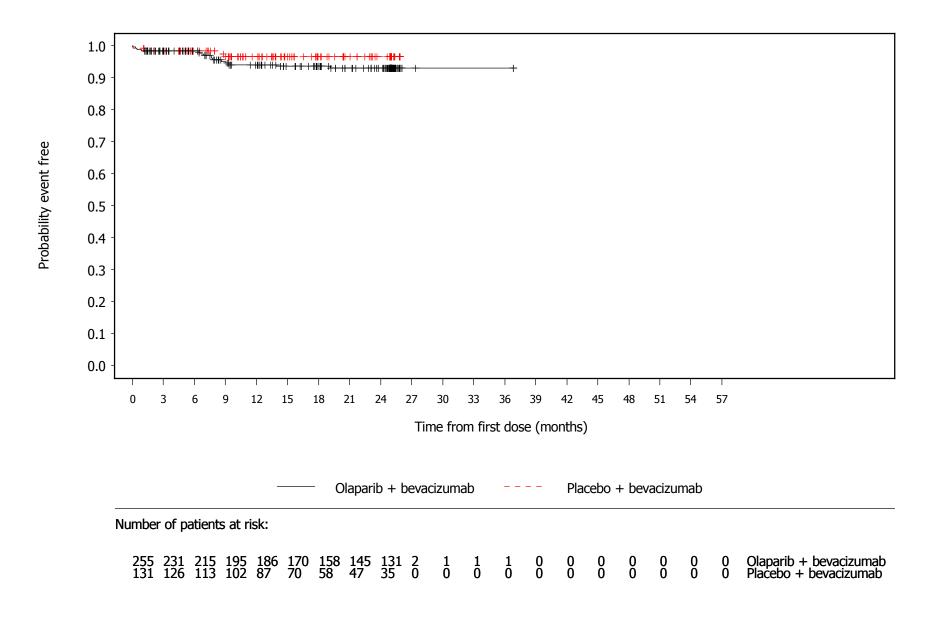


Figure 3.3.8 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Pain Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

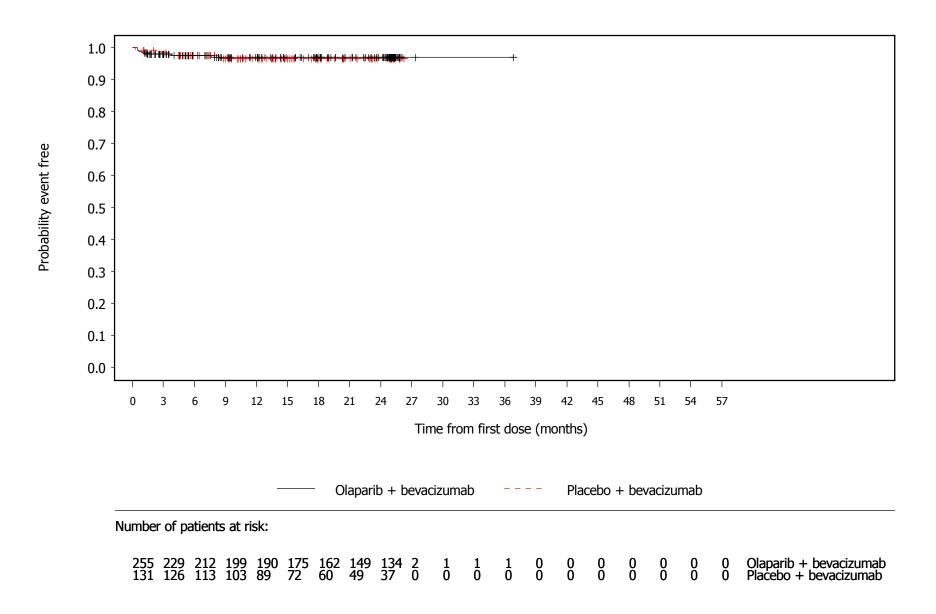


Figure 3.3.9 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Eye disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

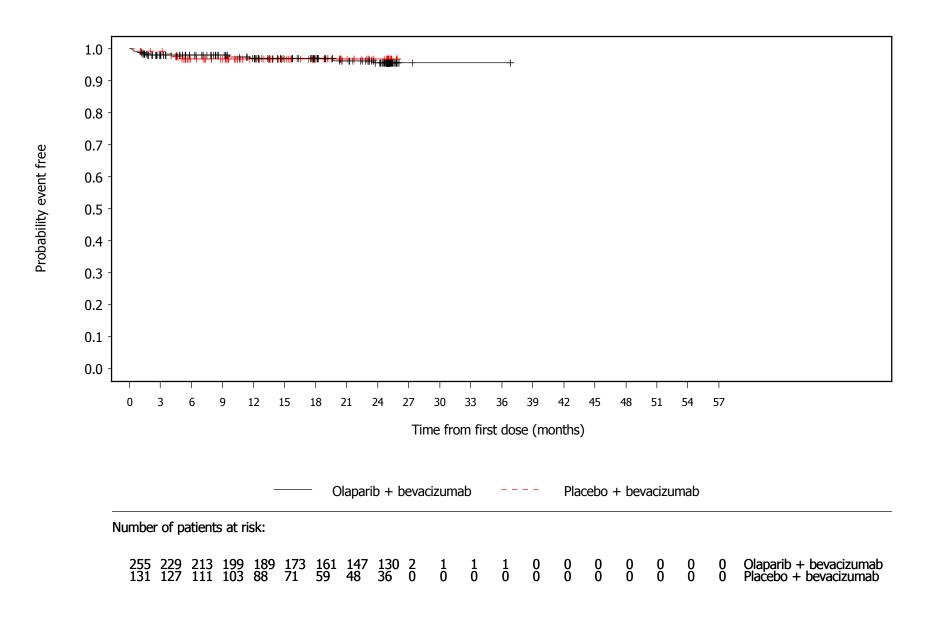


Figure 3.3.10 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Surgical and medical procedures Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

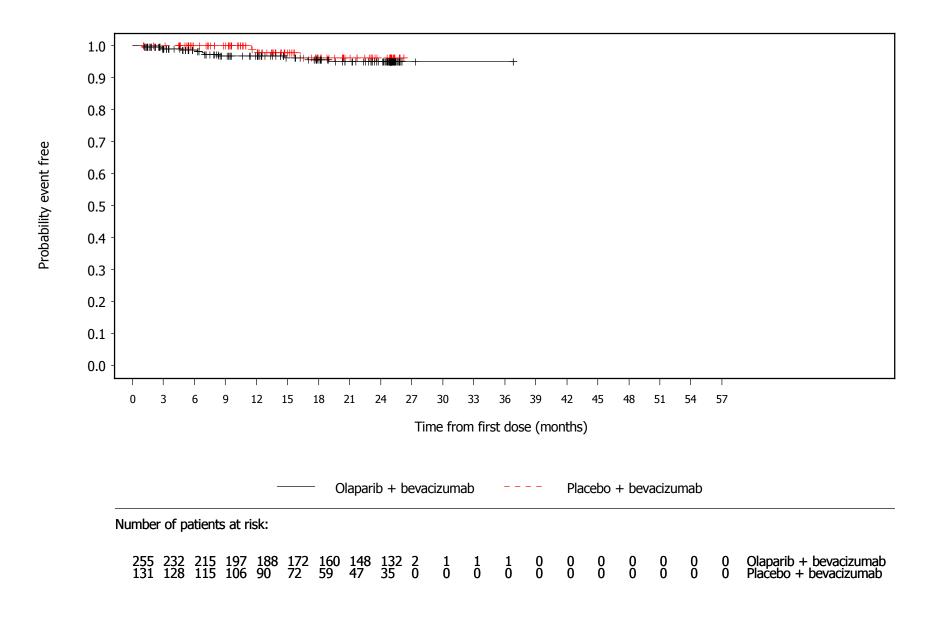


Figure 3.3.11 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Endocrine disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

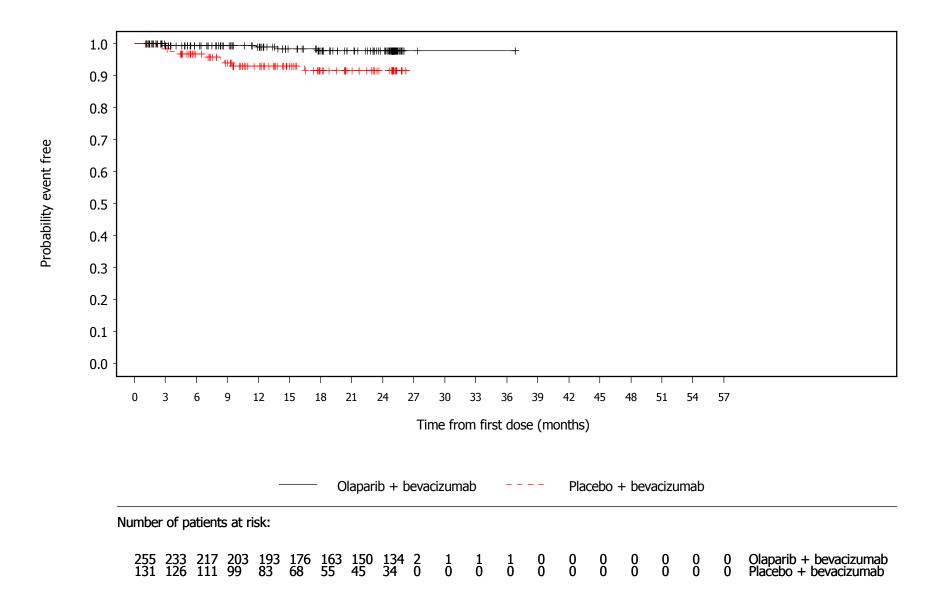


Figure 3.3.12 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Respiratory, thoracic and mediastinal disorders
Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

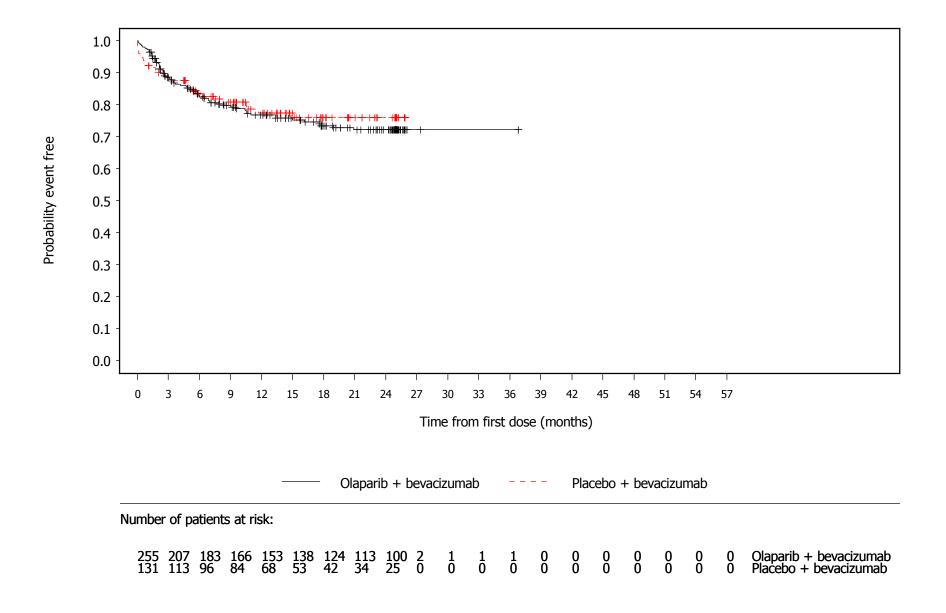


Figure 3.3.13 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Dyspnoea Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

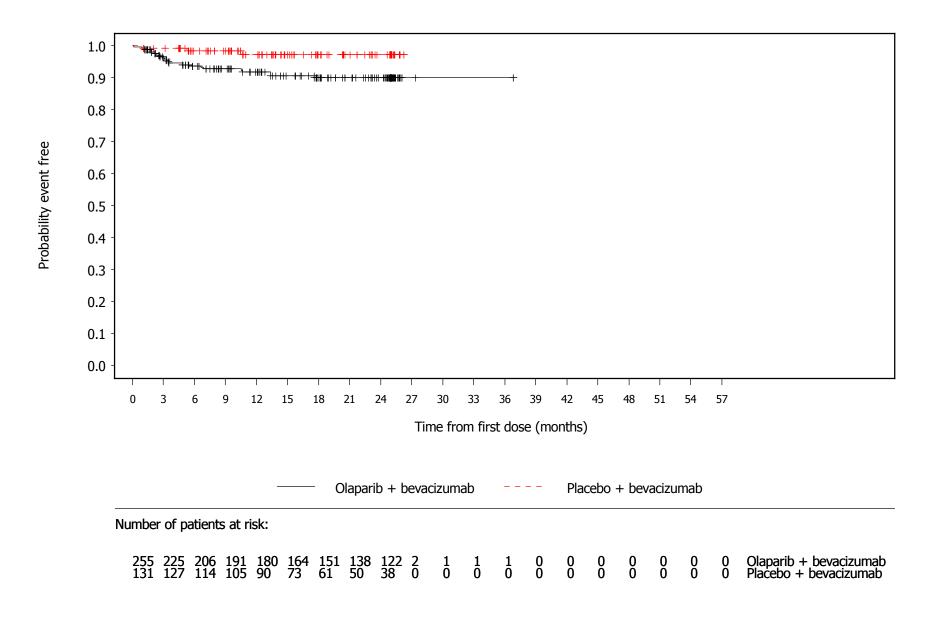


Figure 3.3.14 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Epistaxis Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

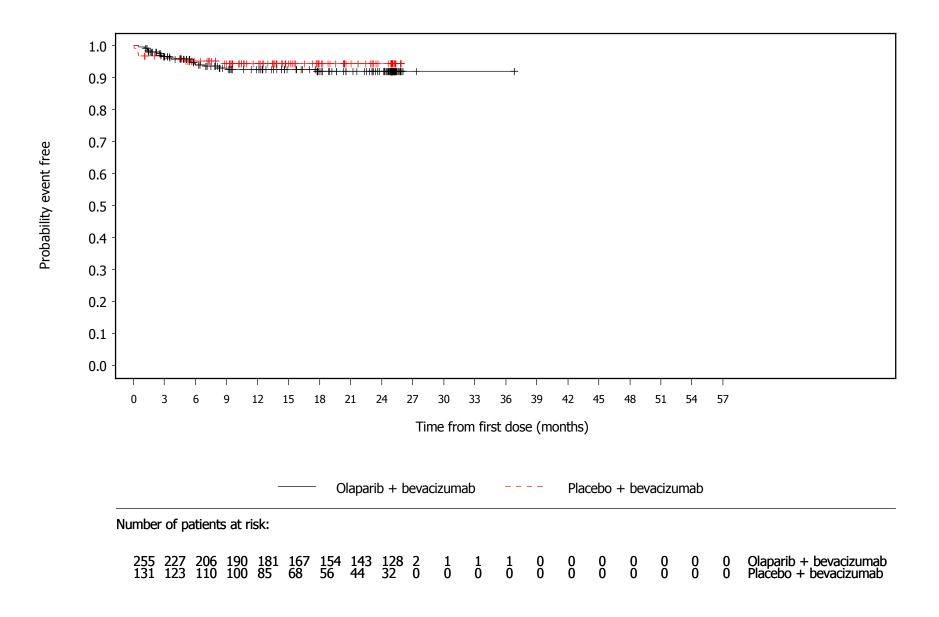


Figure 3.3.15 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Cough Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

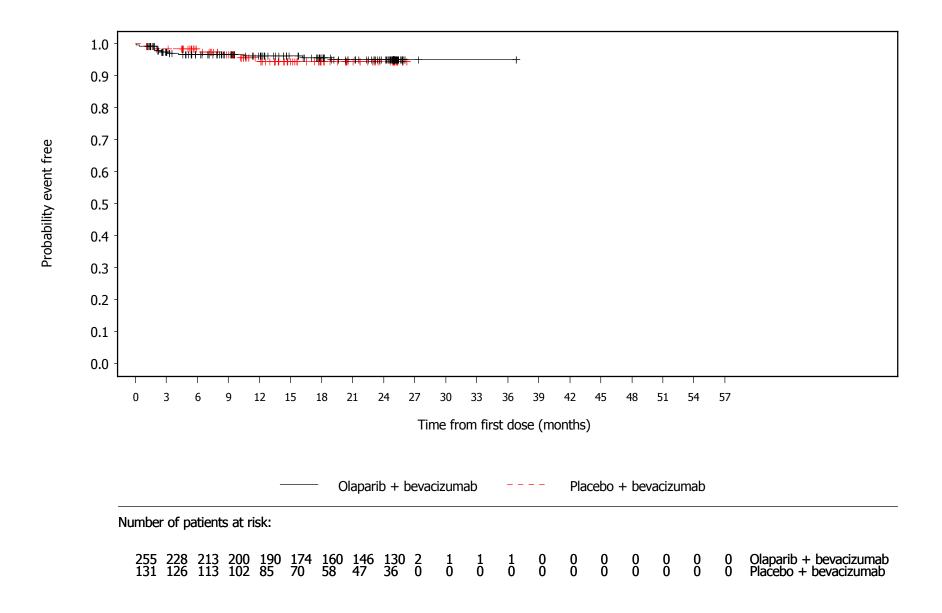


Figure 3.3.16 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Rhinorrhoea Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

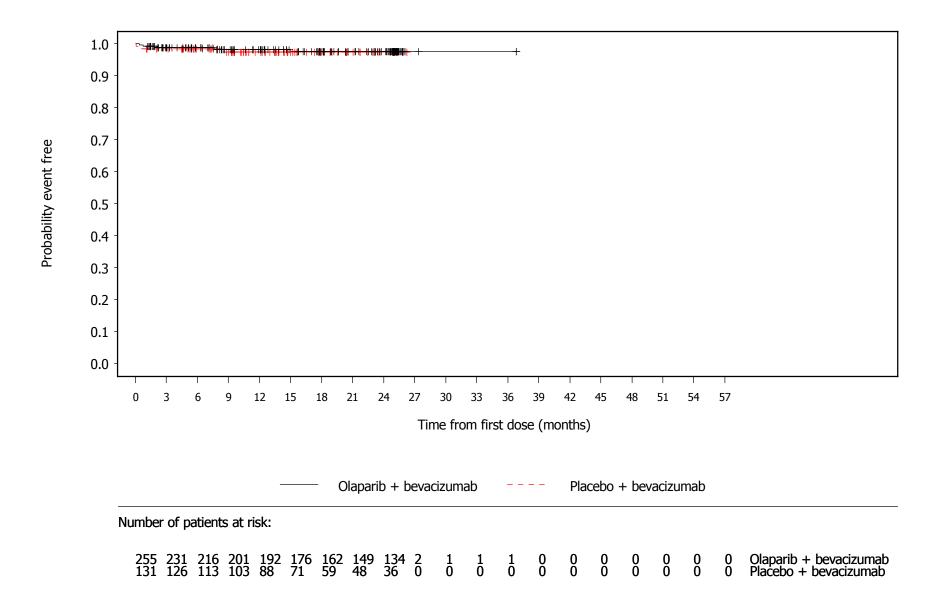


Figure 3.3.17 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Oropharyngeal pain Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

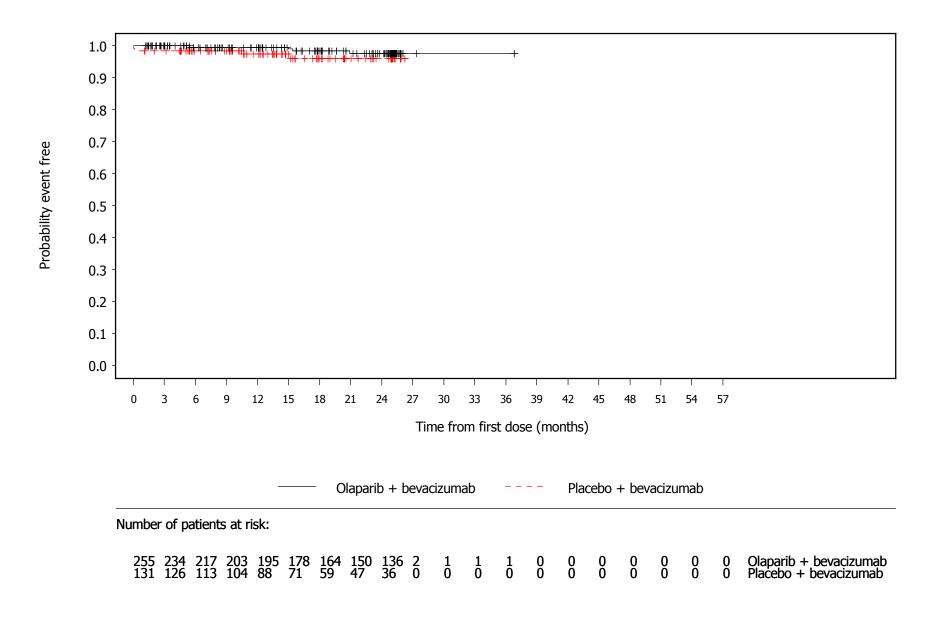


Figure 3.3.18 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Skin and subcutaneous tissue disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

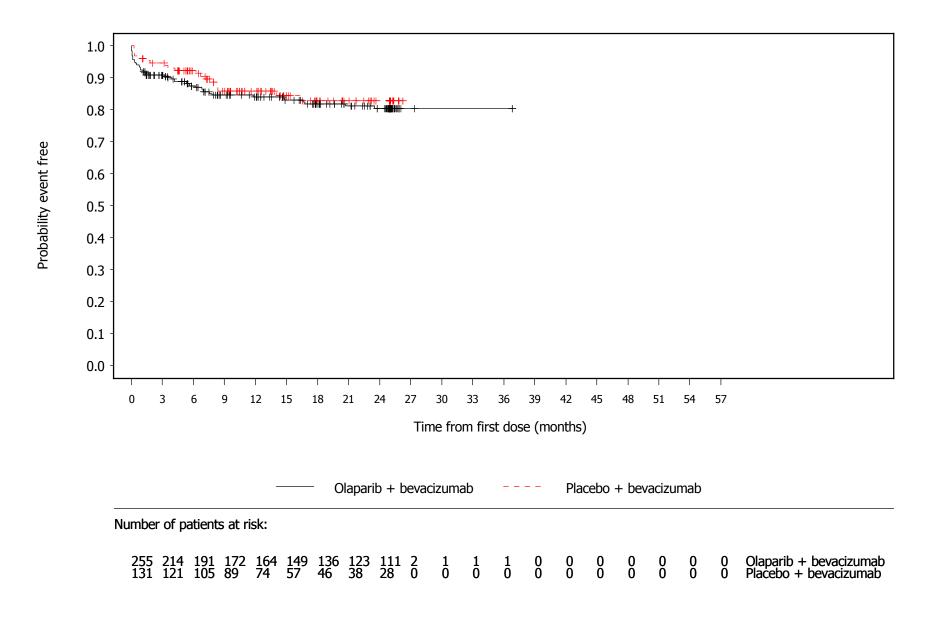


Figure 3.3.19 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Alopecia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

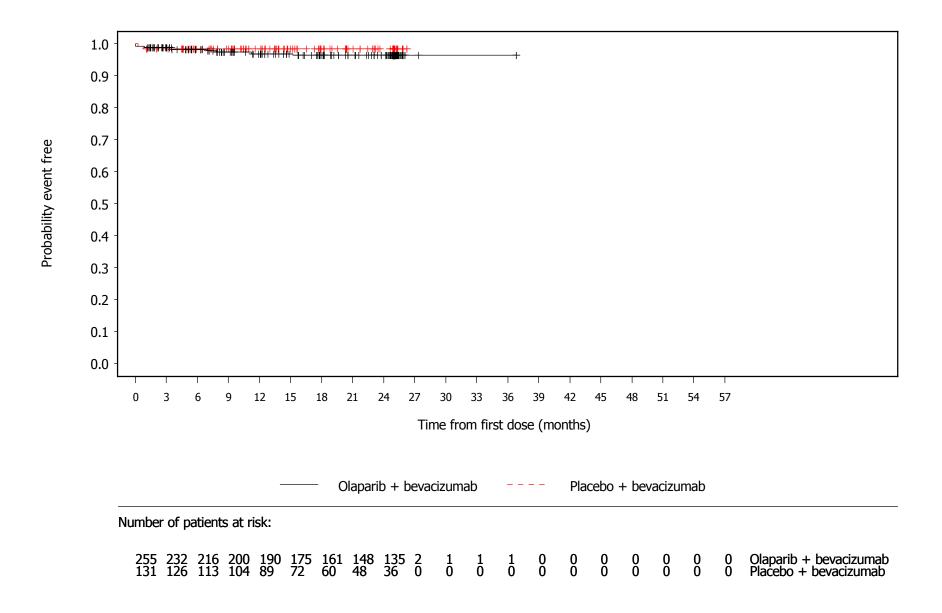


Figure 3.3.20 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Rash Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

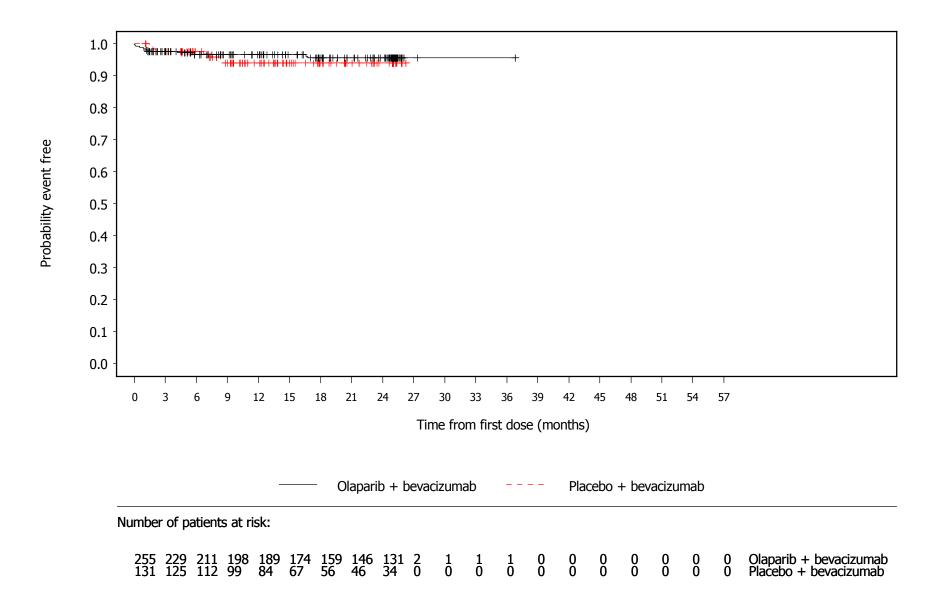


Figure 3.3.21 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Erythema Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

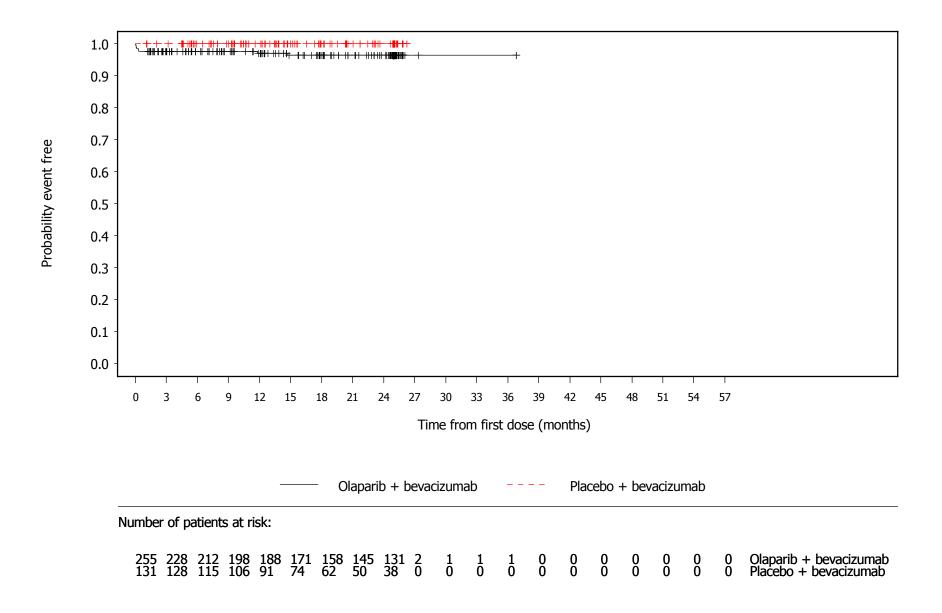


Figure 3.3.22 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Pruritus Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

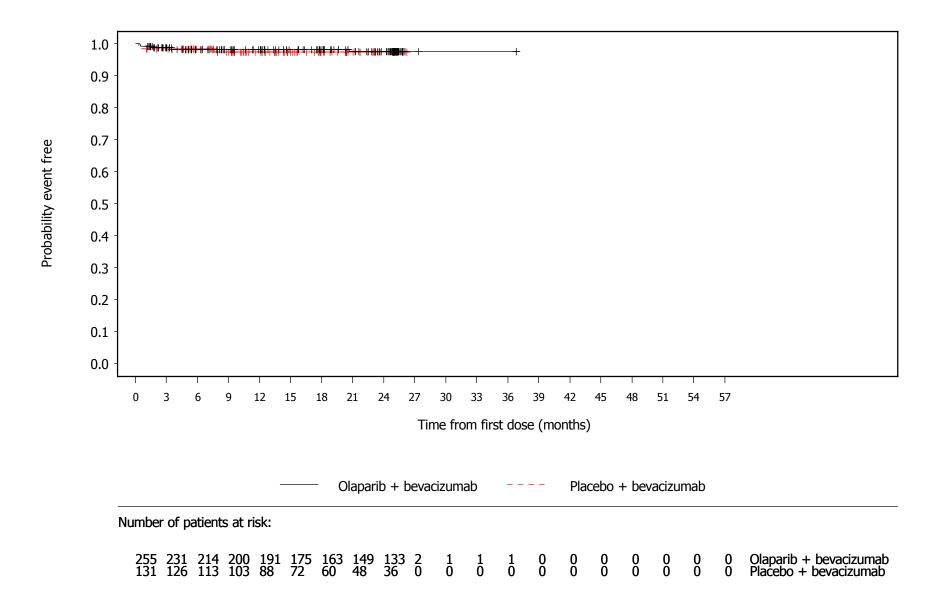


Figure 3.3.23 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Renal and urinary disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

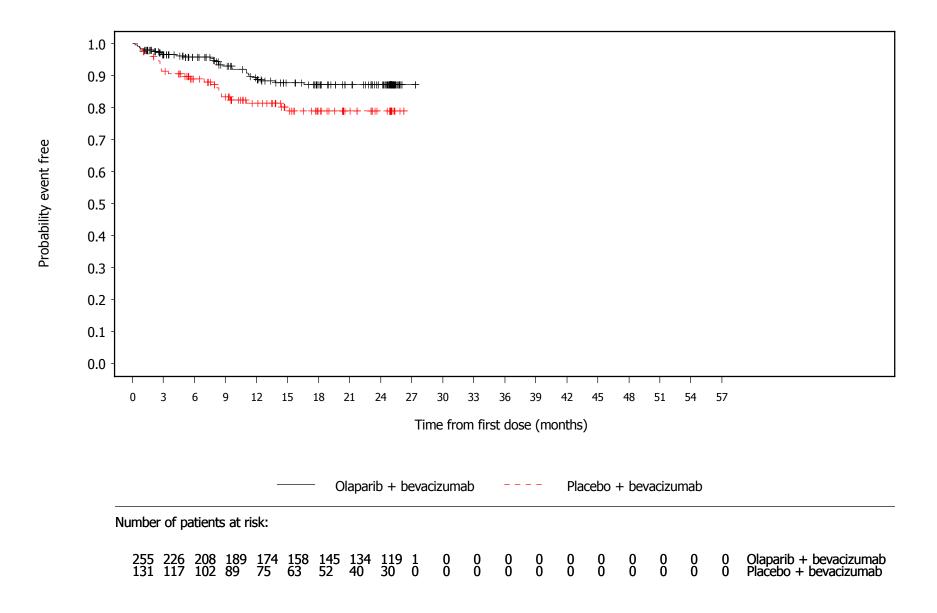


Figure 3.3.24 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

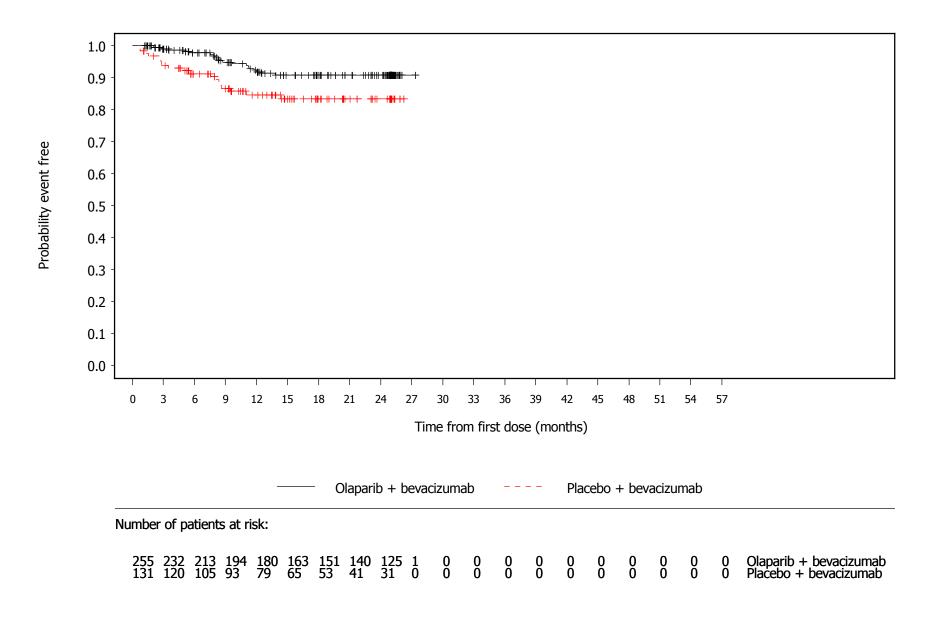


Figure 3.3.25 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Blood and lymphatic system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

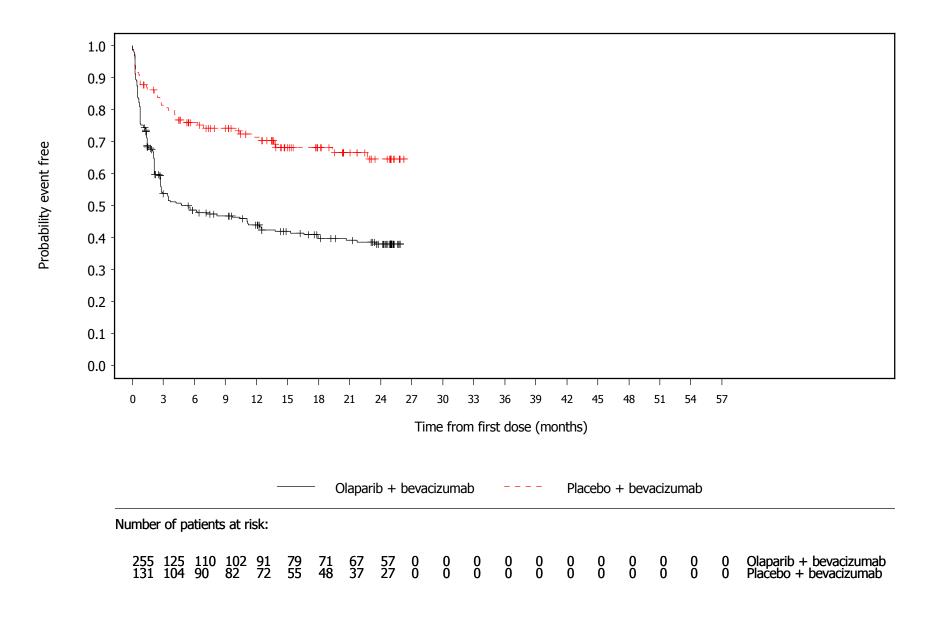


Figure 3.3.26 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

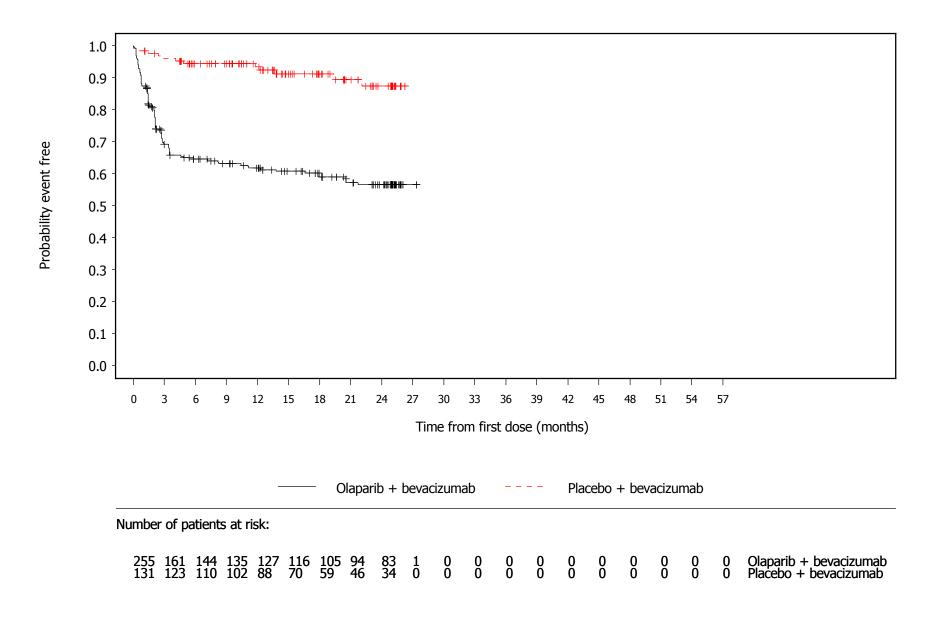


Figure 3.3.27 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Leukopenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

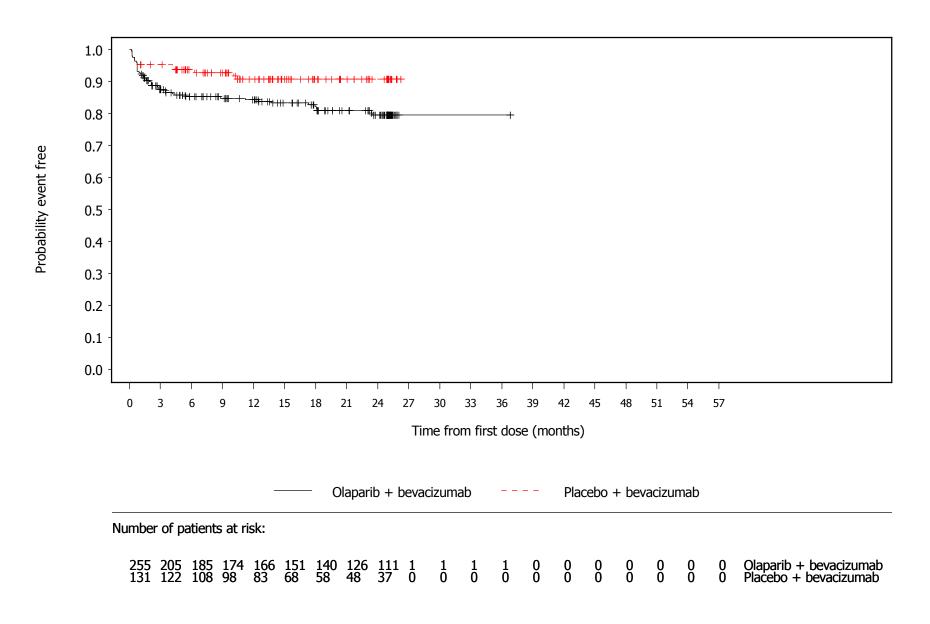


Figure 3.3.28 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Lymphopenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

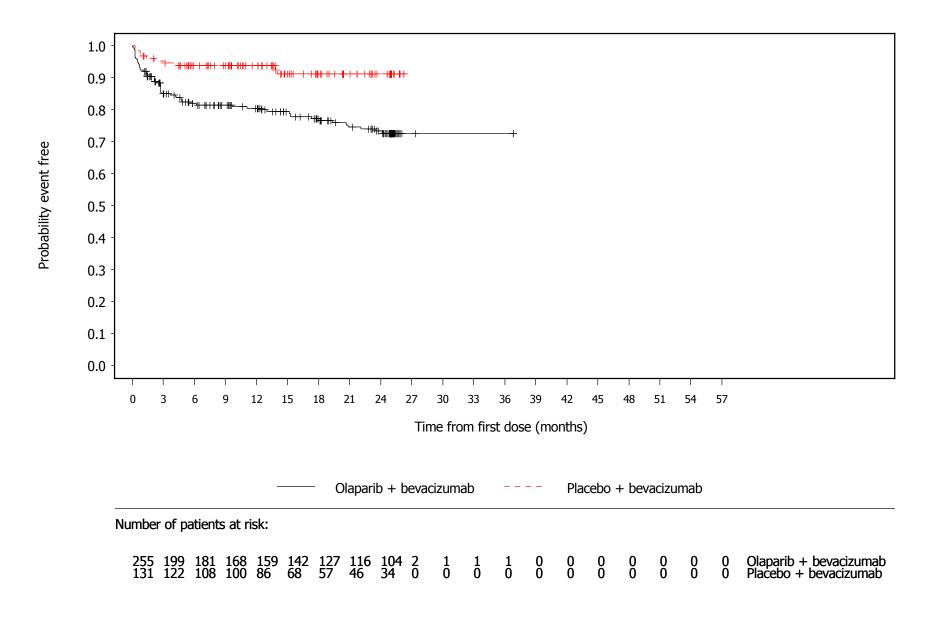


Figure 3.3.29 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

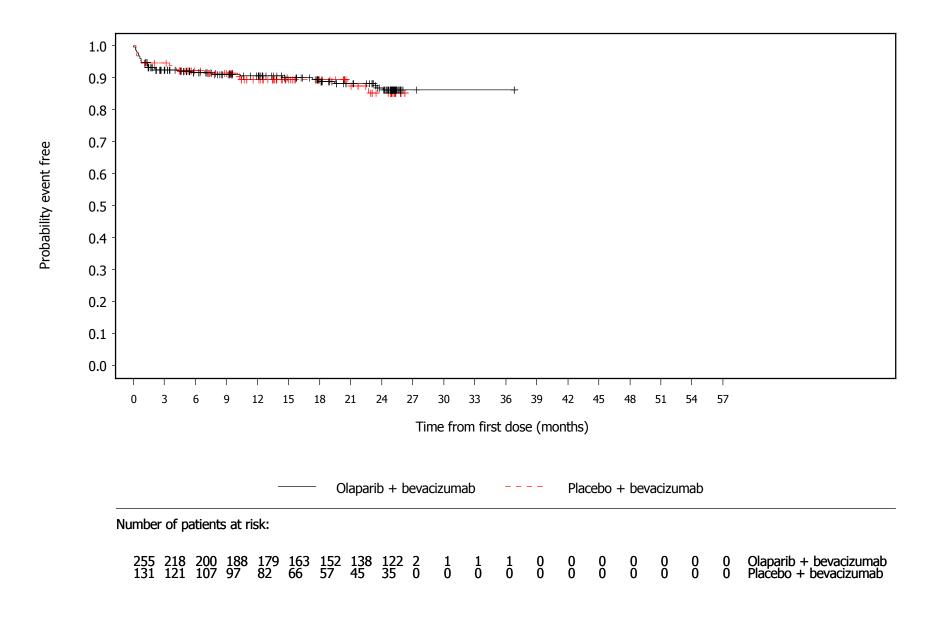


Figure 3.3.30 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

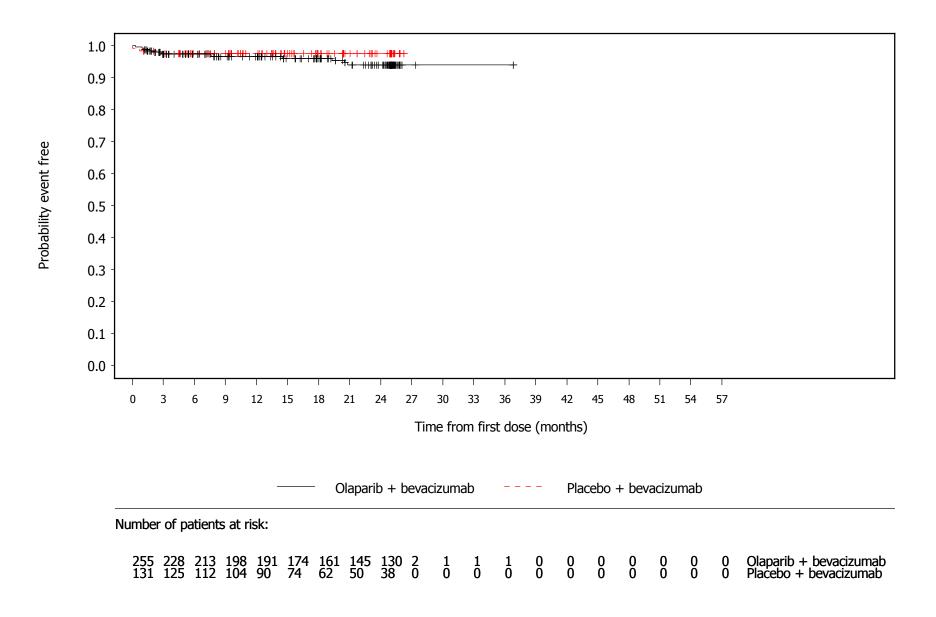


Figure 3.3.31 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Gastrointestinal disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

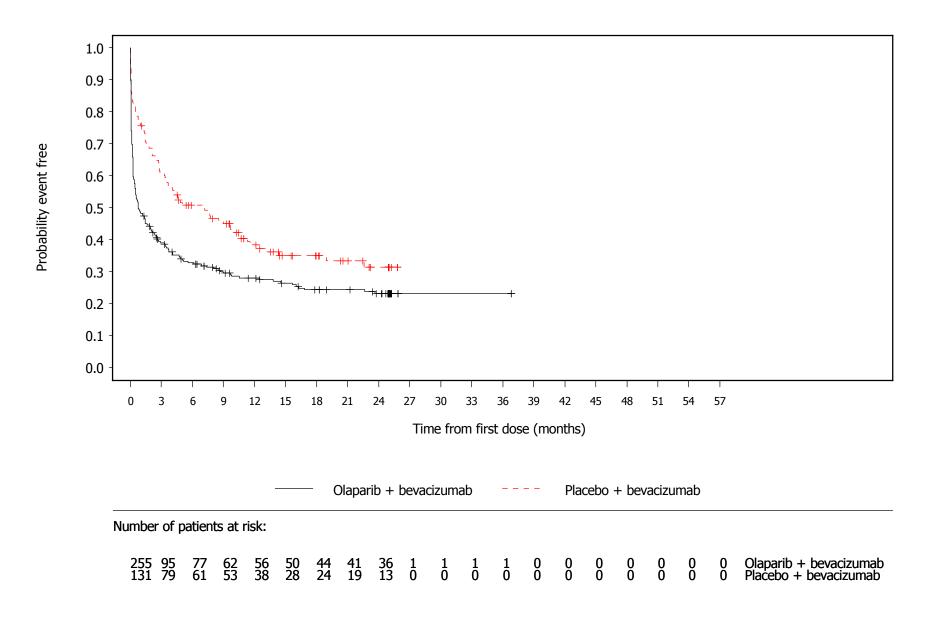


Figure 3.3.32 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Abdominal pain Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

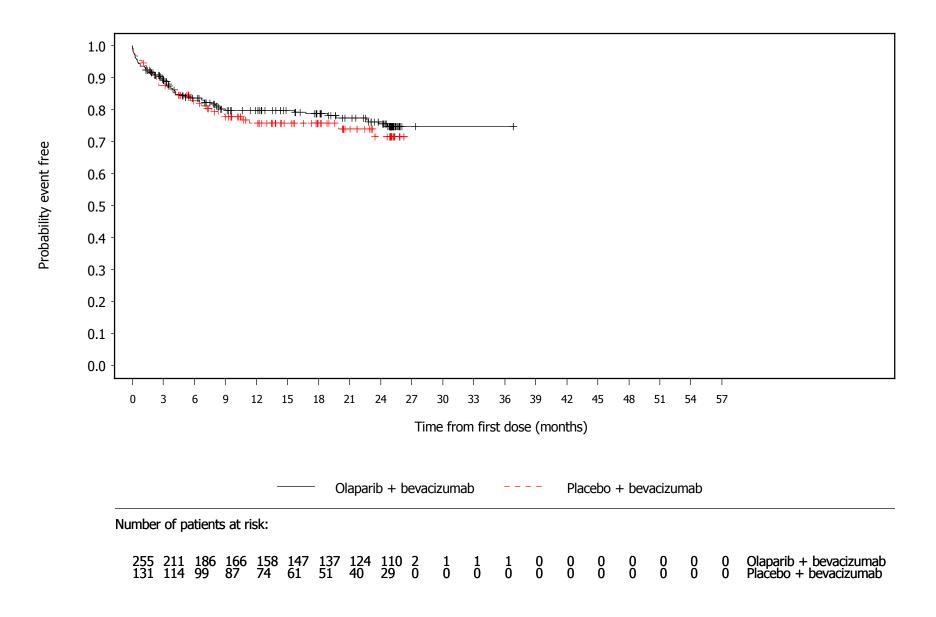


Figure 3.3.33 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Intestinal obstruction Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

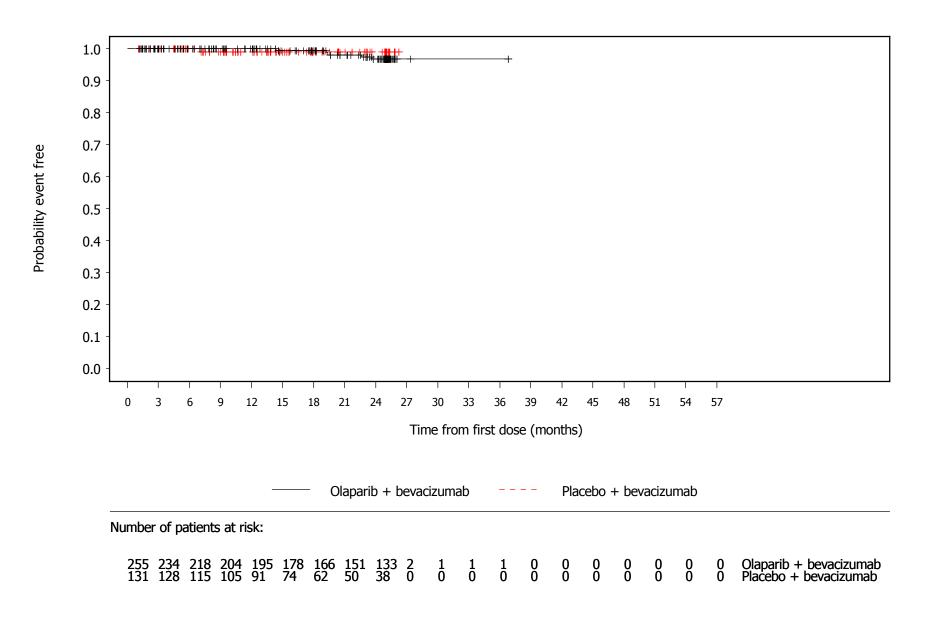


Figure 3.3.34 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Diarrhoea Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

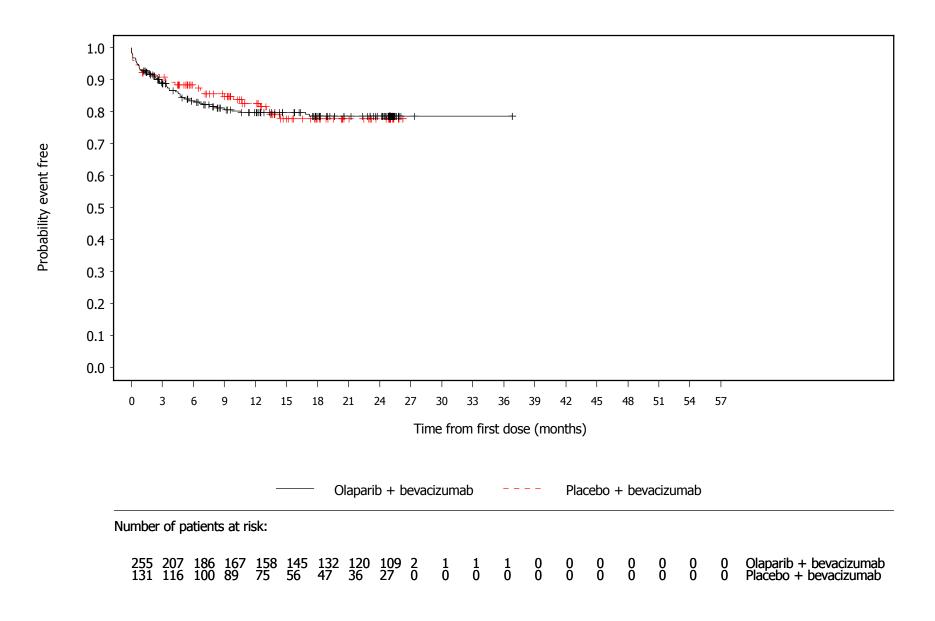


Figure 3.3.35 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Dyspepsia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

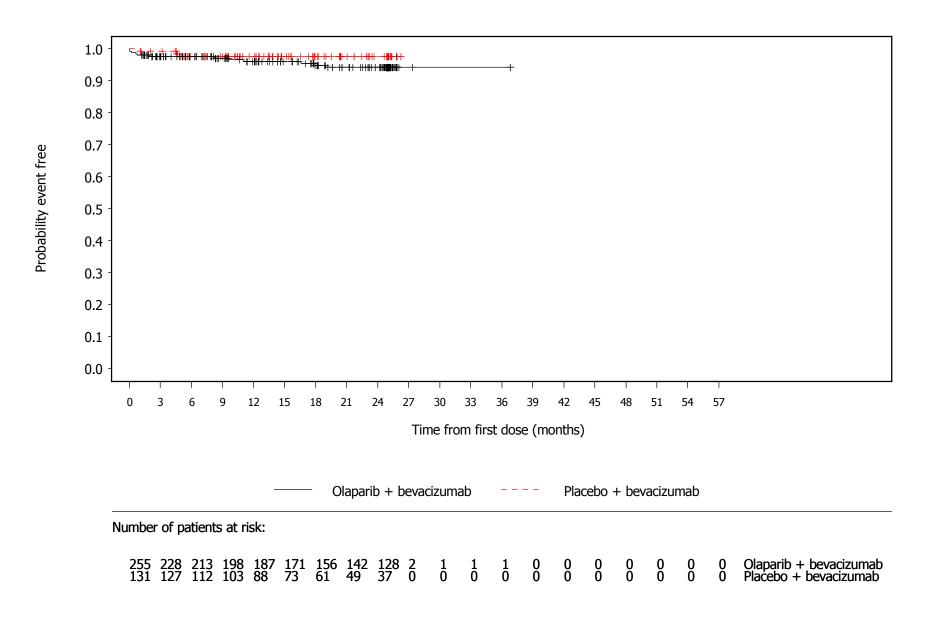


Figure 3.3.36 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

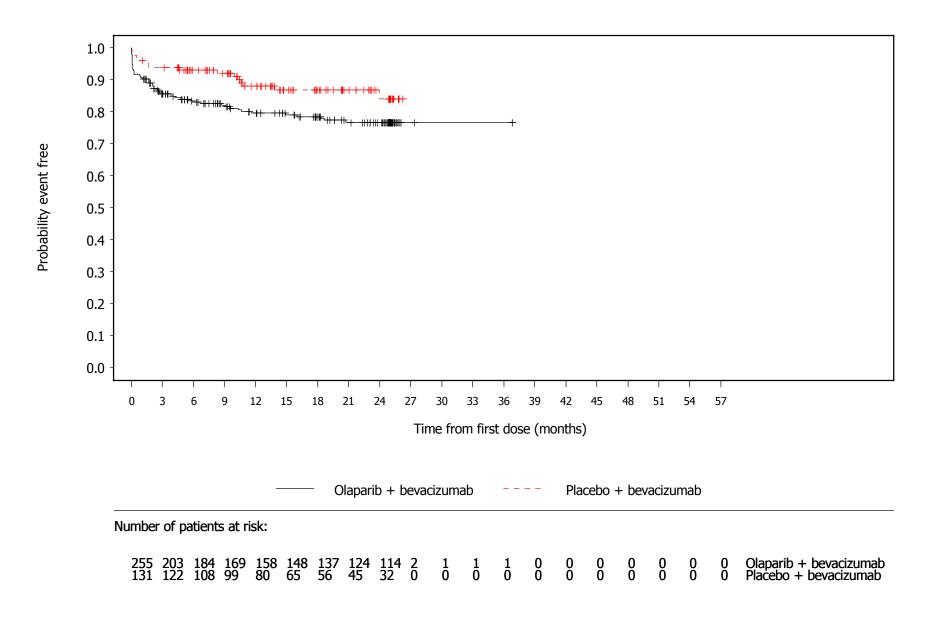


Figure 3.3.37 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Haemorrhoids Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

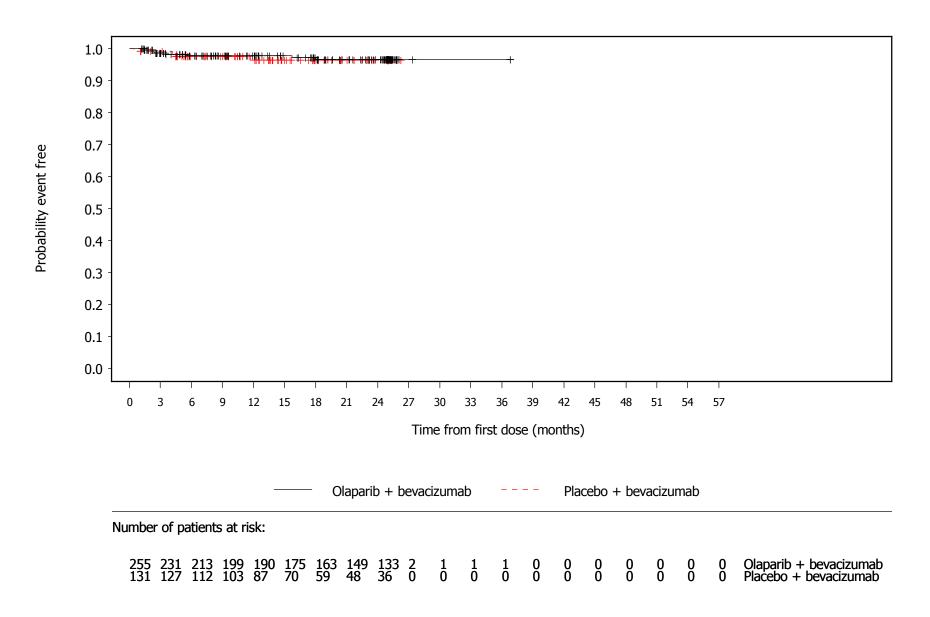


Figure 3.3.38 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Constipation Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

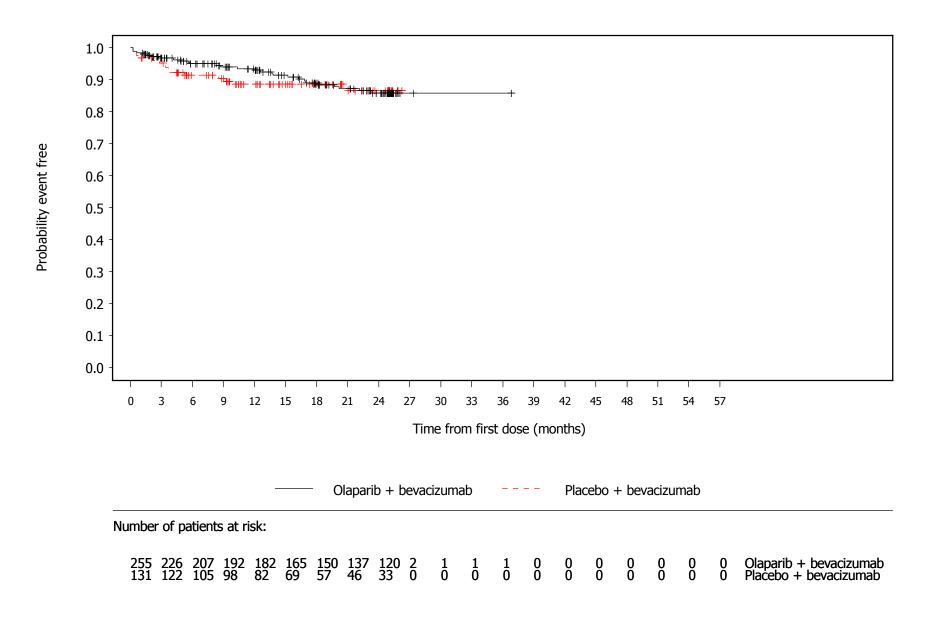


Figure 3.3.39 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Abdominal pain upper Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

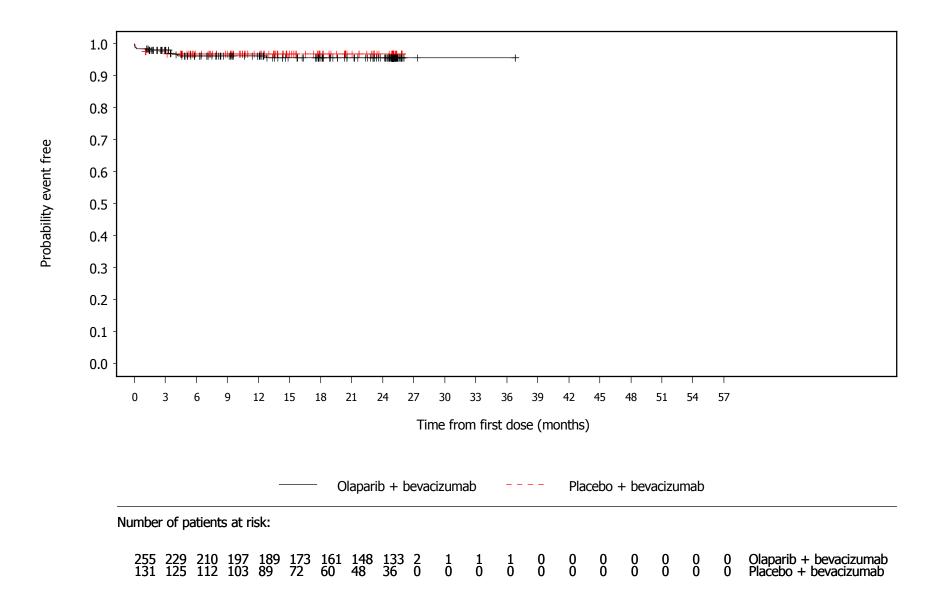


Figure 3.3.40 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Stomatitis Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

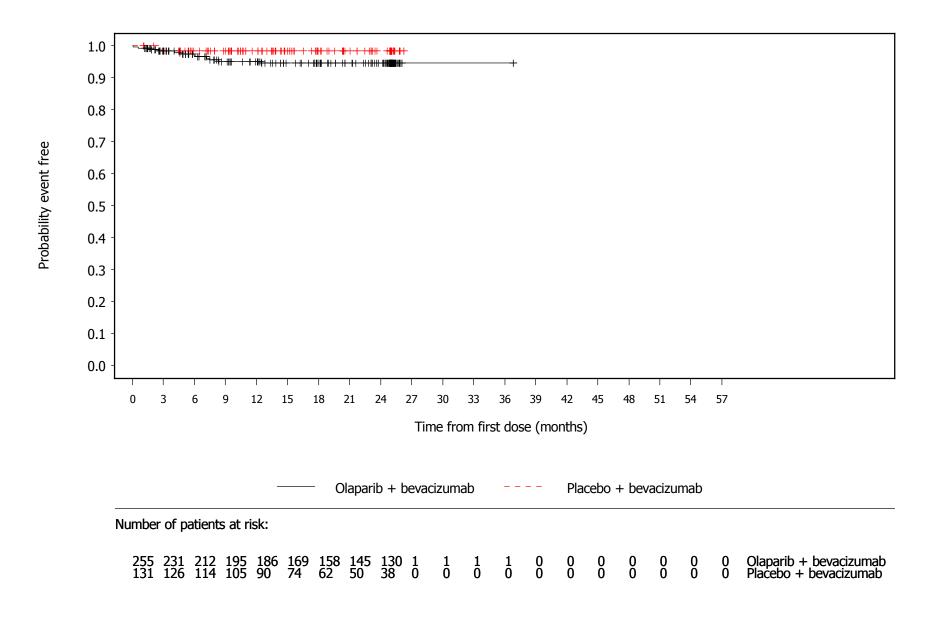


Figure 3.3.41 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Subileus Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

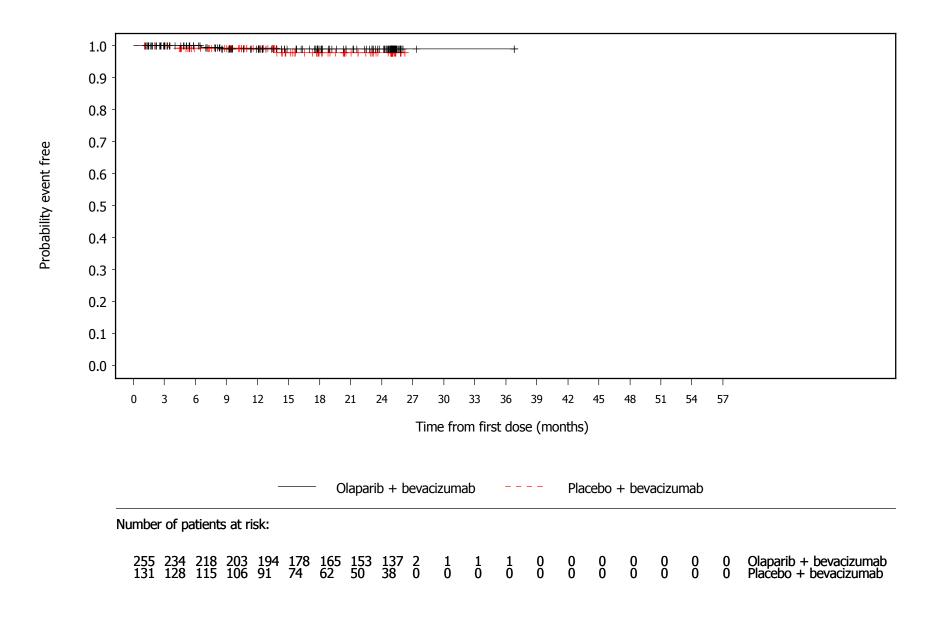


Figure 3.3.42 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Nausea Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

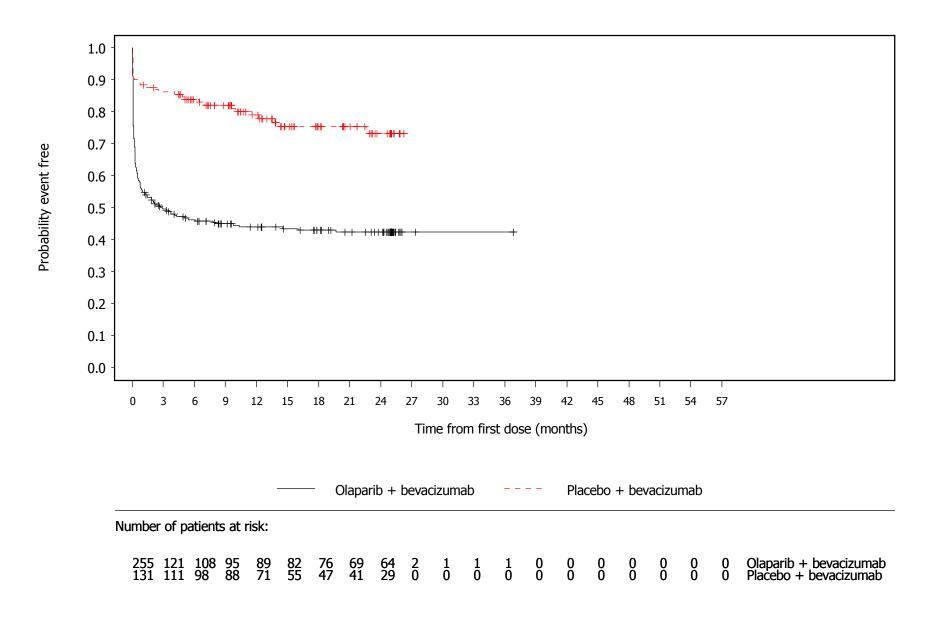


Figure 3.3.43 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Gingival bleeding Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

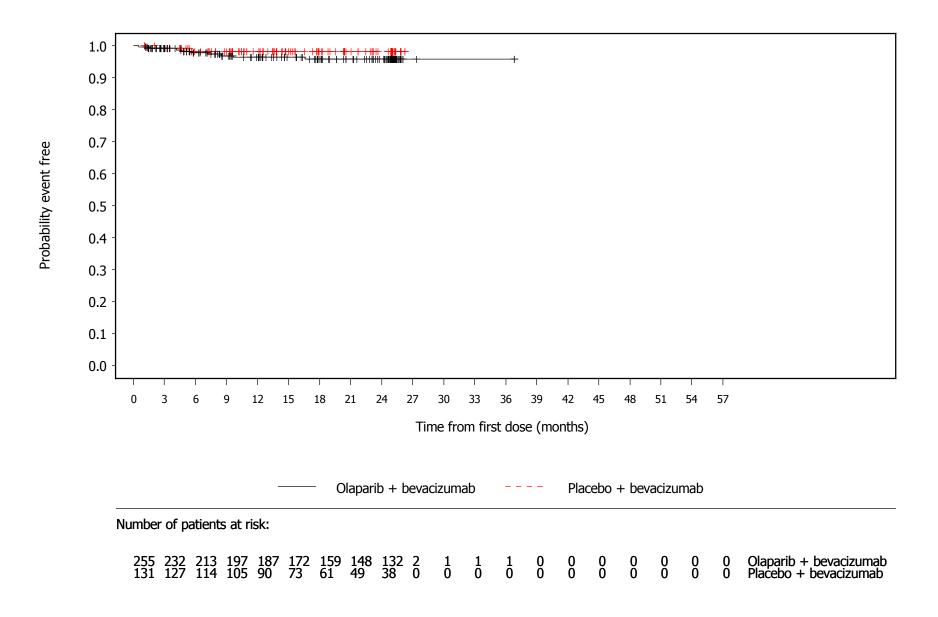


Figure 3.3.44 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Toothache Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

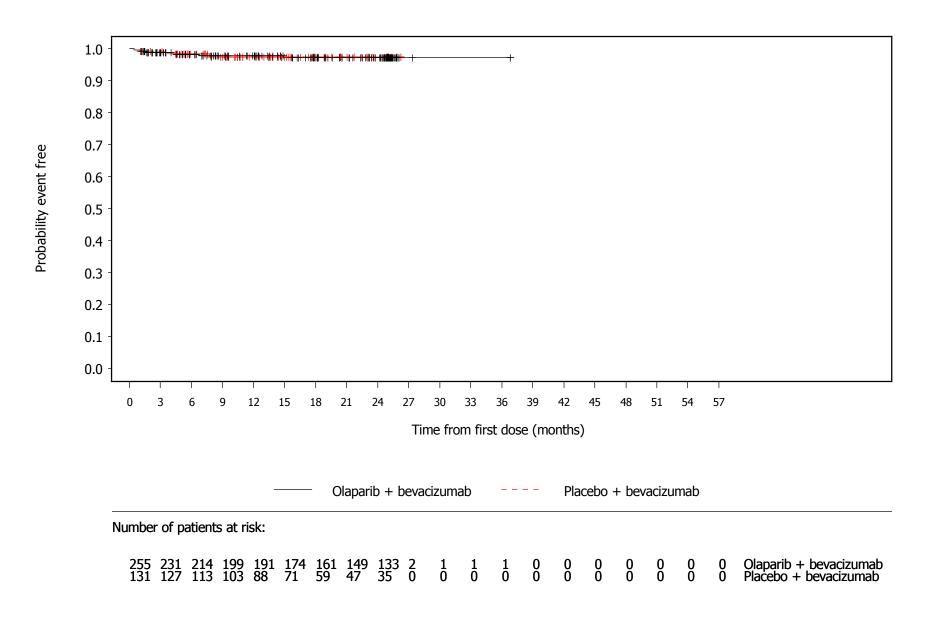


Figure 3.3.45 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Immune system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

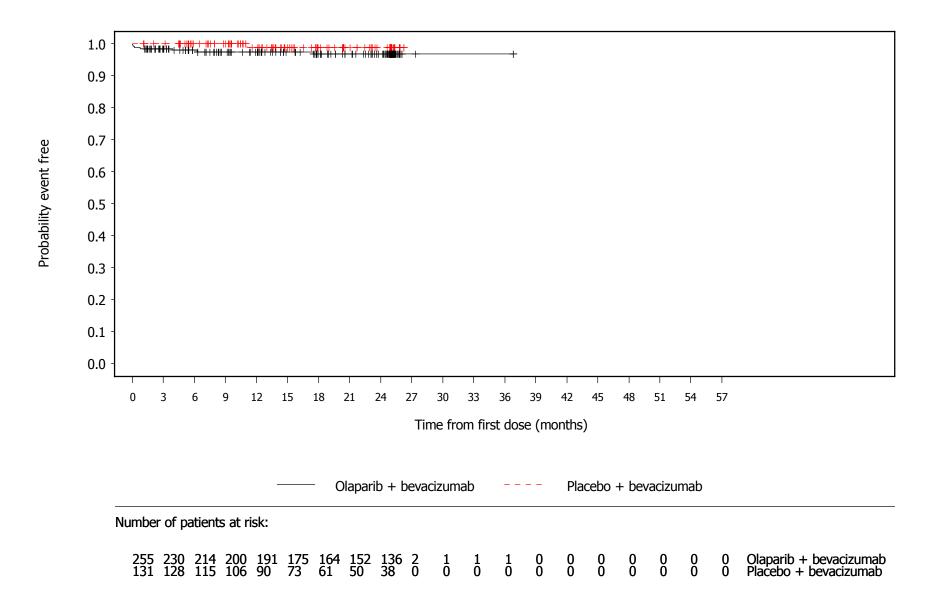


Figure 3.3.46 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Nervous system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

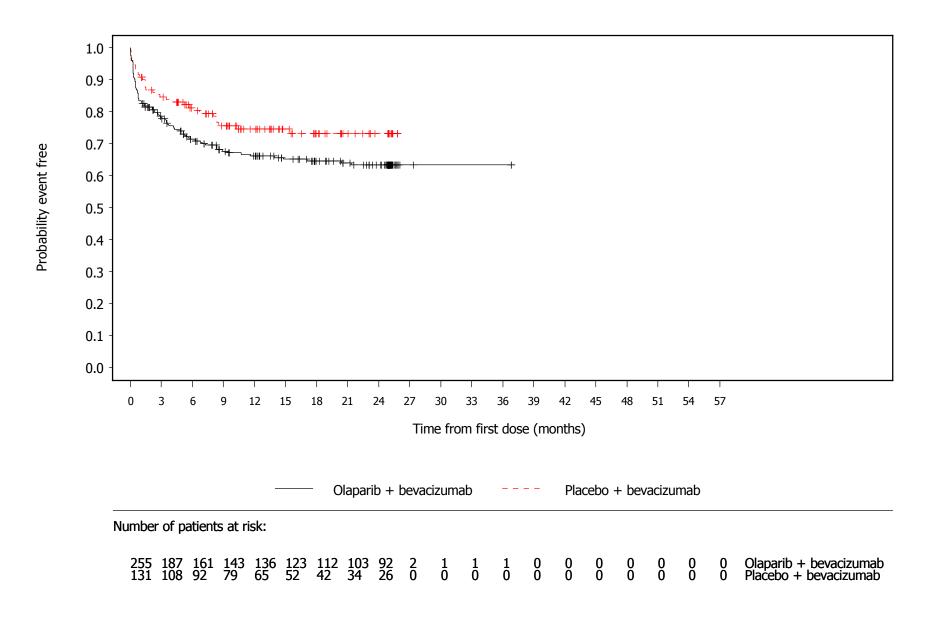


Figure 3.3.47 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Dysgeusia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

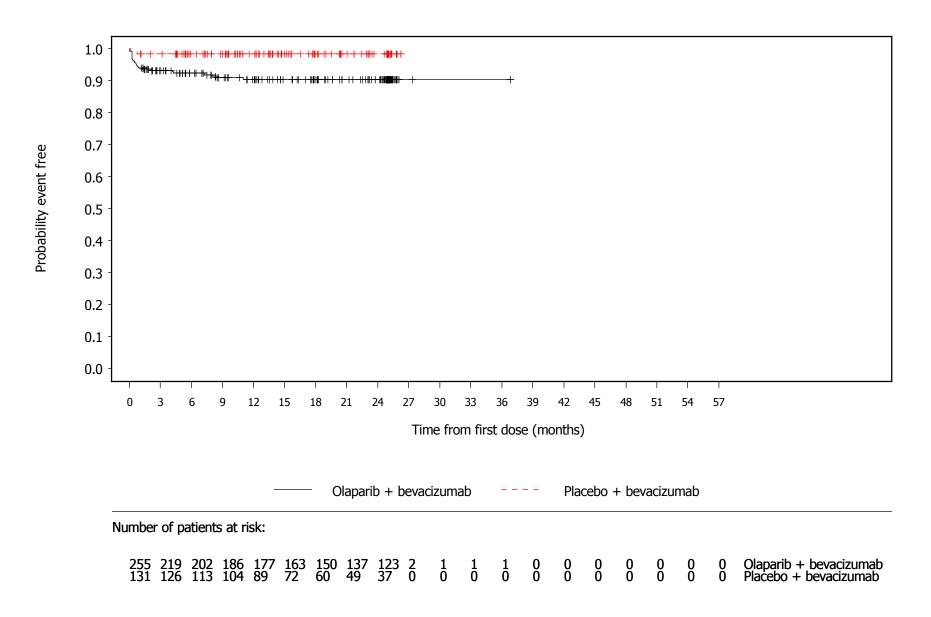


Figure 3.3.48 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Headache Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

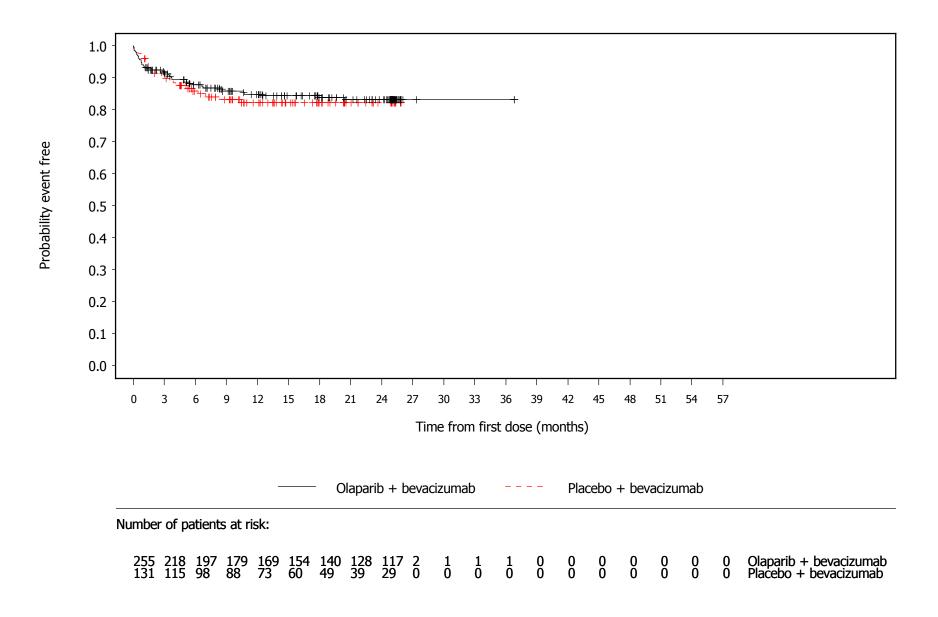


Figure 3.3.49 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Neuropathy peripheral Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

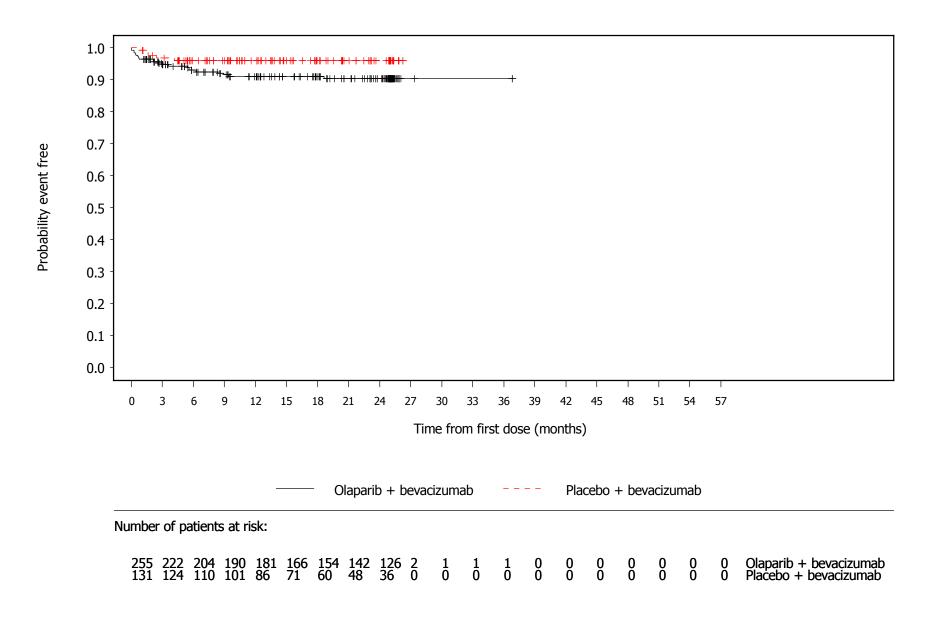


Figure 3.3.50 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Polyneuropathy Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

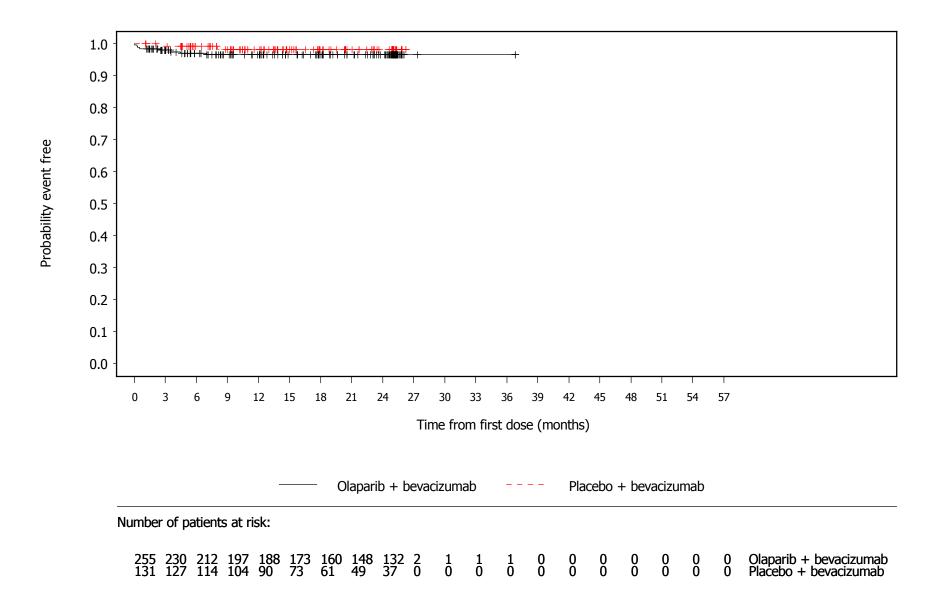


Figure 3.3.51 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Dizziness Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

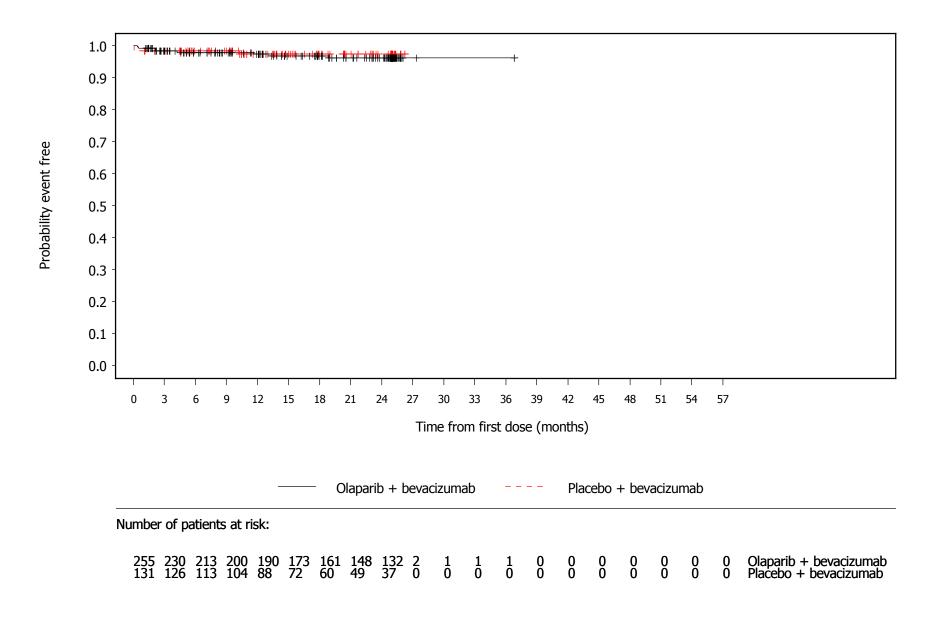


Figure 3.3.52 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Ear and labyrinth disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

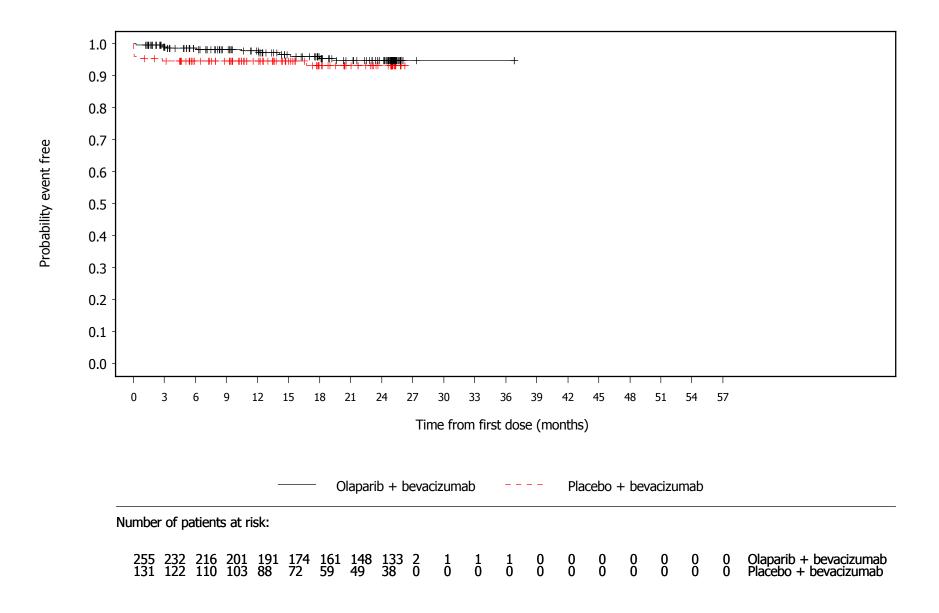


Figure 3.3.53 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Vertigo Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

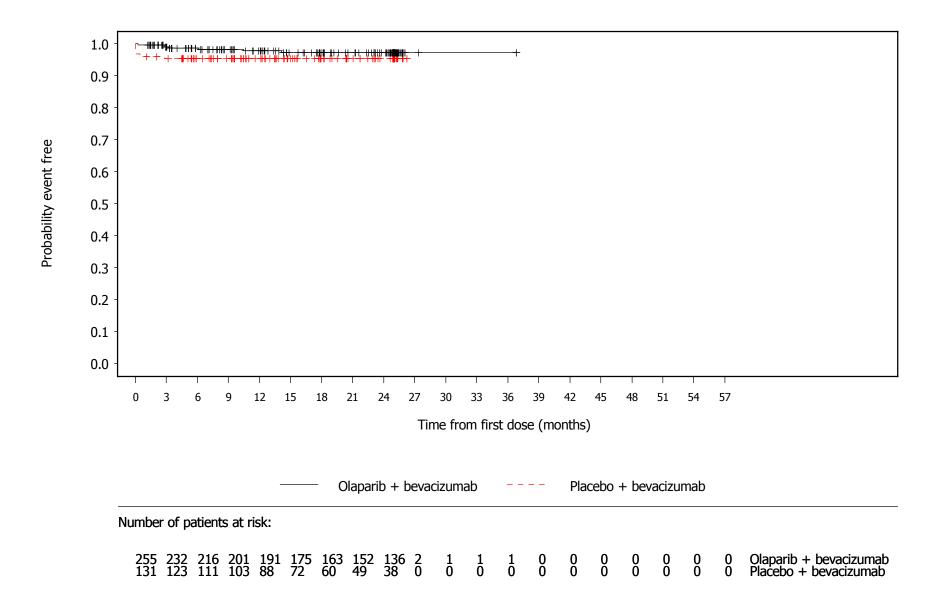


Figure 3.3.54 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Vascular disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

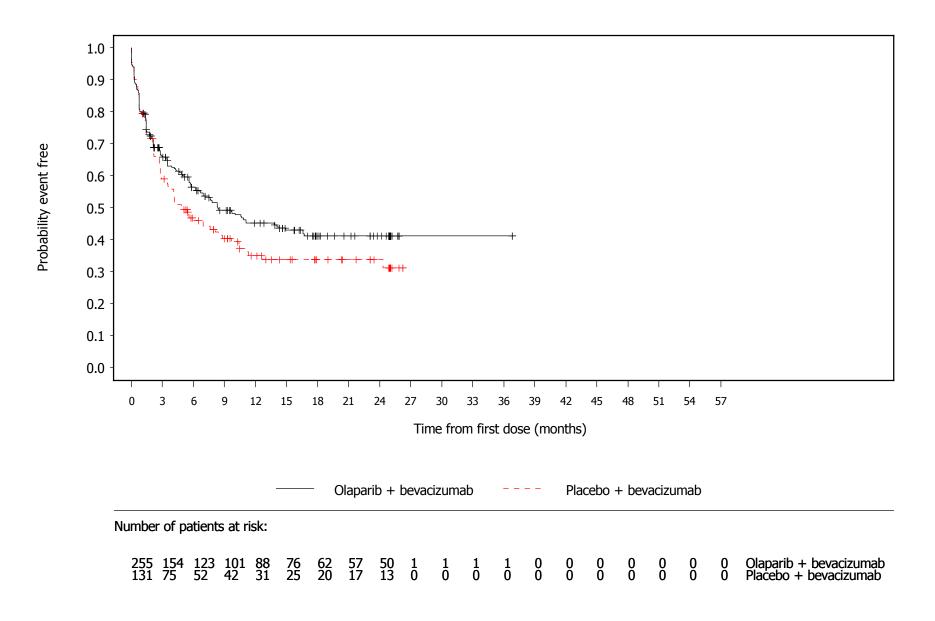


Figure 3.3.55 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Hot flush Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

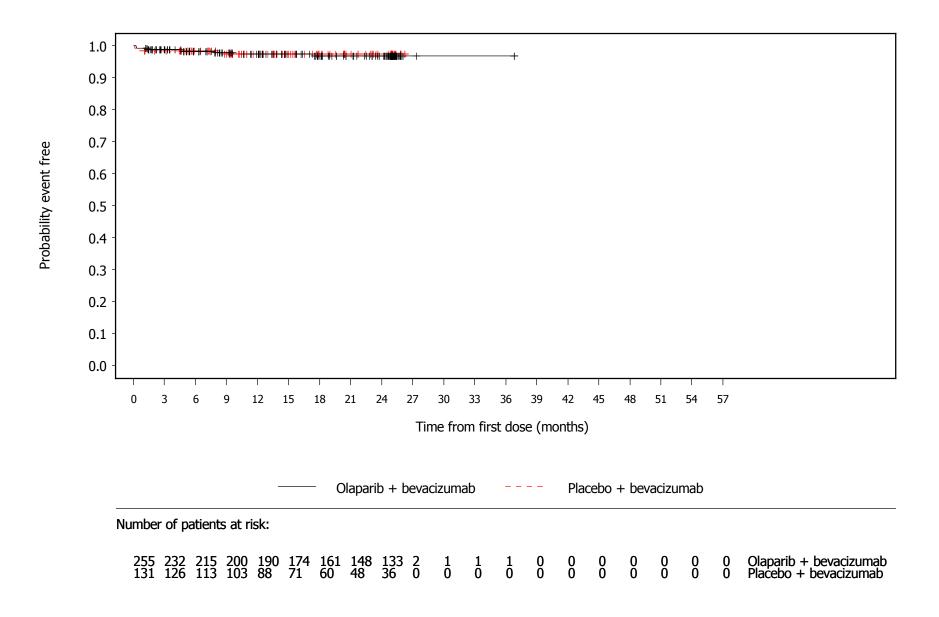


Figure 3.3.56 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

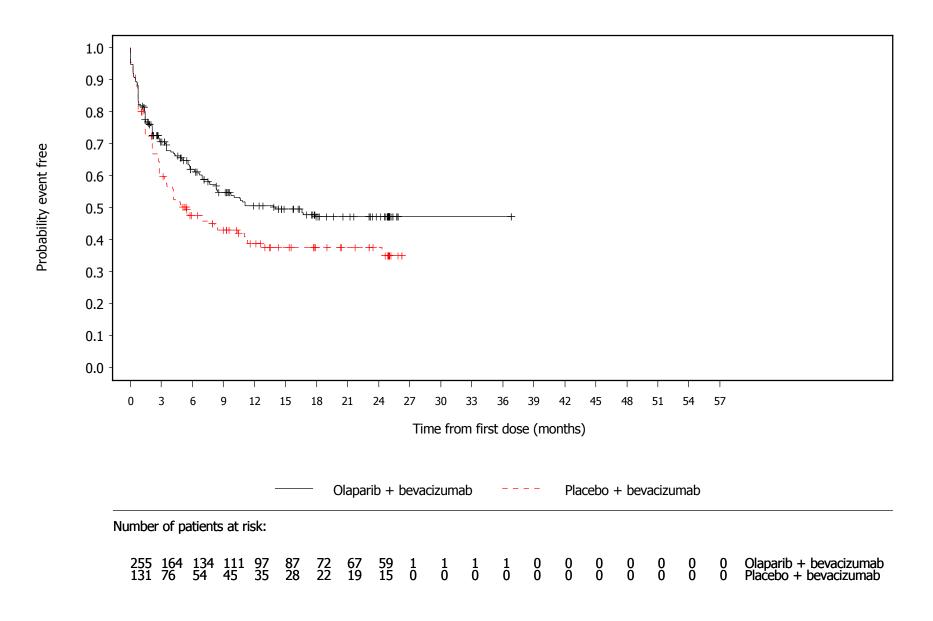


Figure 3.3.57 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Cardiac disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

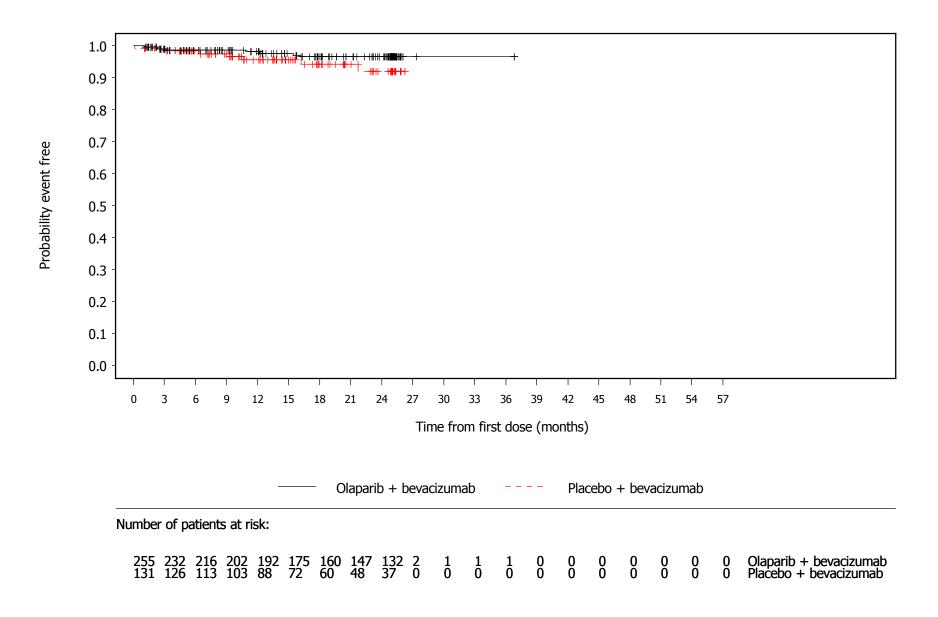


Figure 3.3.58 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Infections and infestations Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

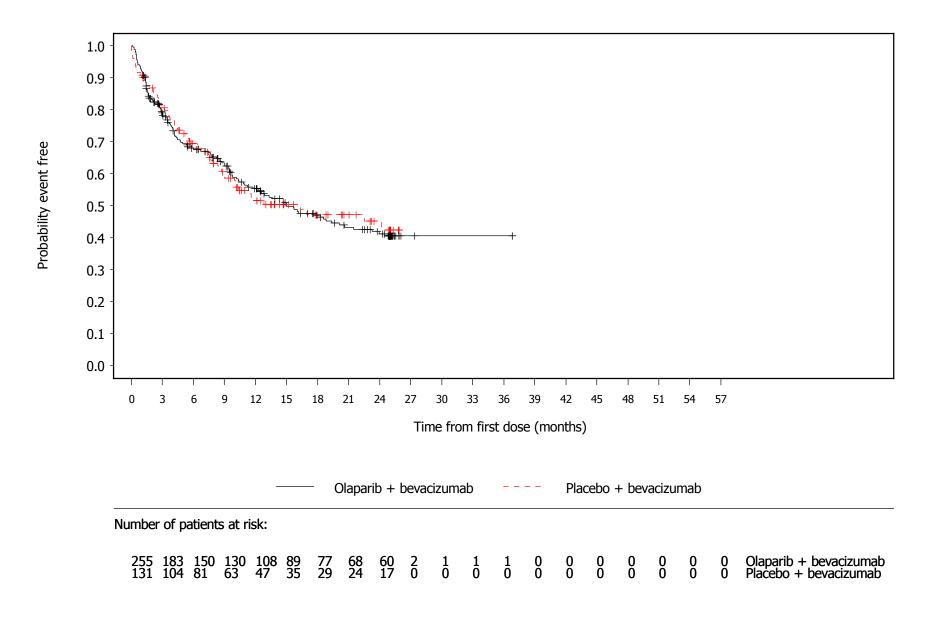


Figure 3.3.59 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Bronchitis Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

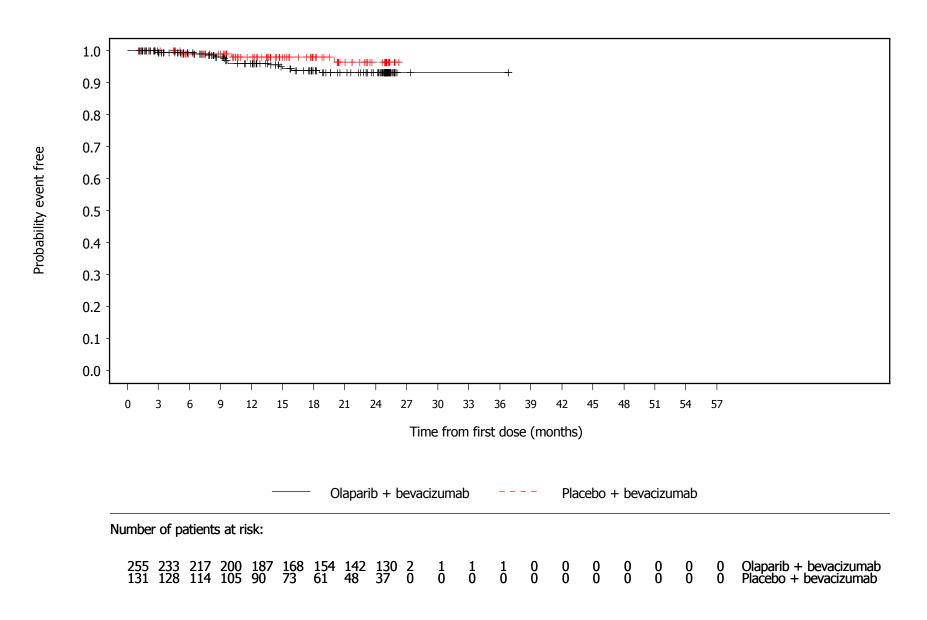


Figure 3.3.60 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Gastroenteritis Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

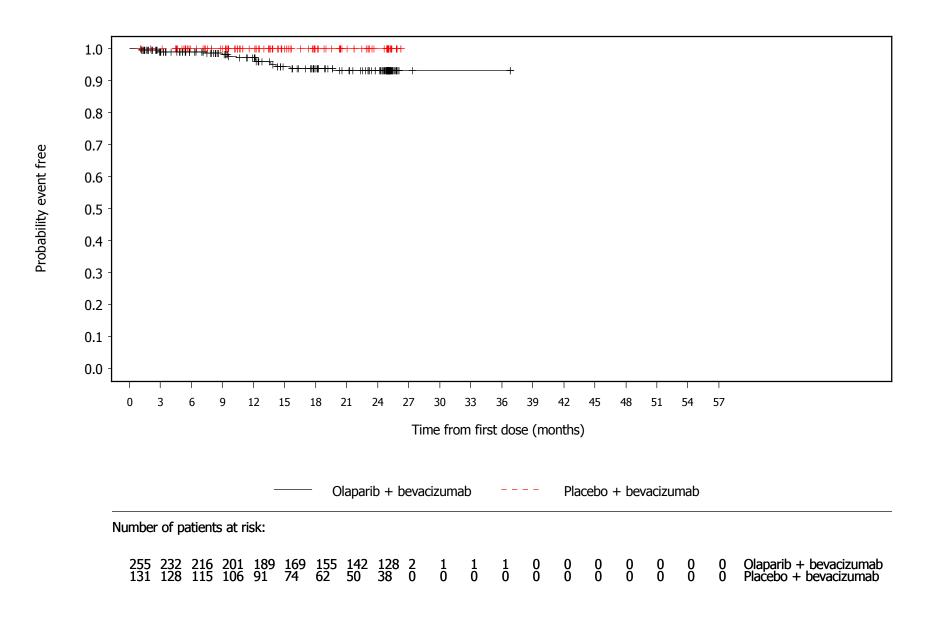


Figure 3.3.61 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Gingivitis Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

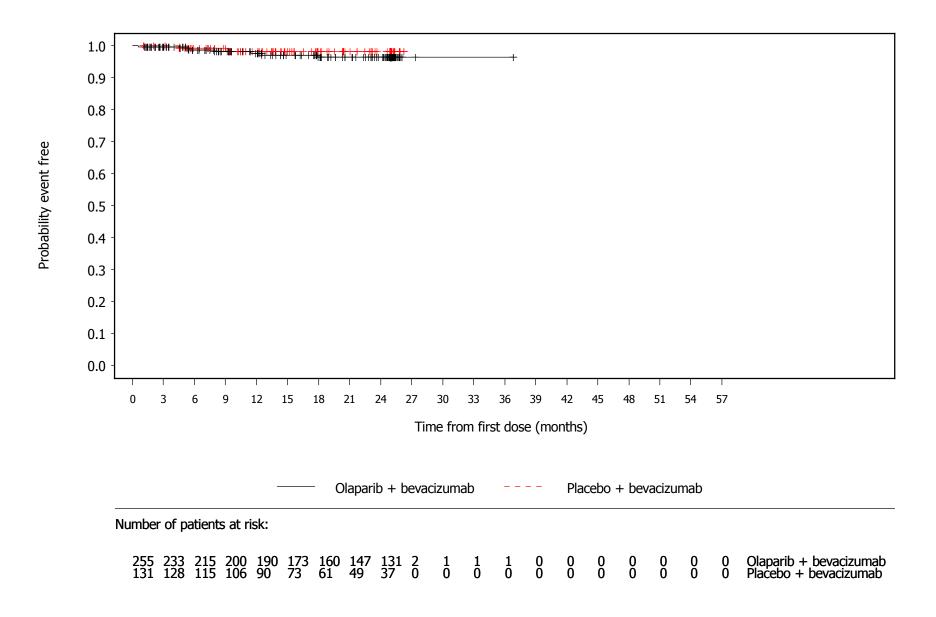


Figure 3.3.62 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Influenza Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

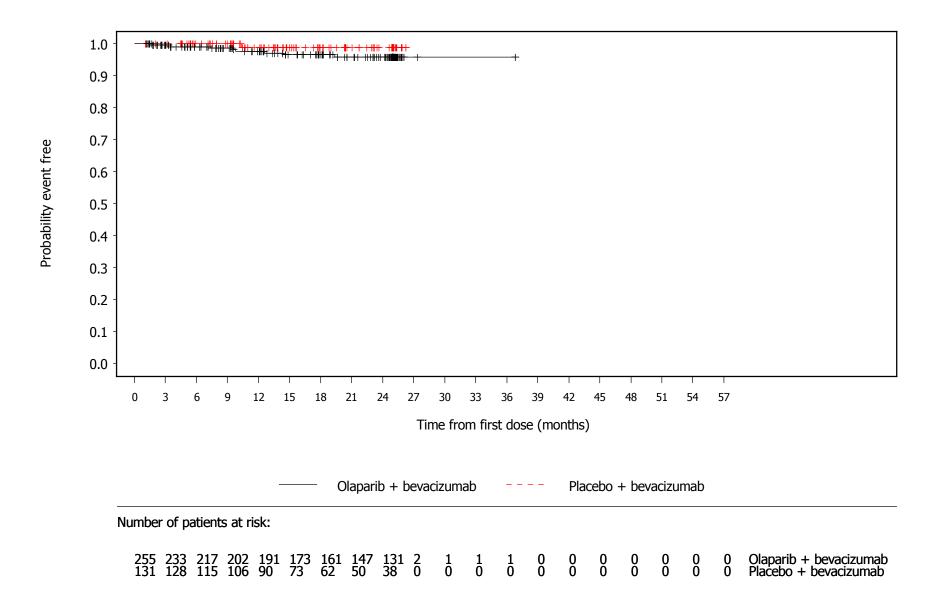


Figure 3.3.63 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Urinary tract infection Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

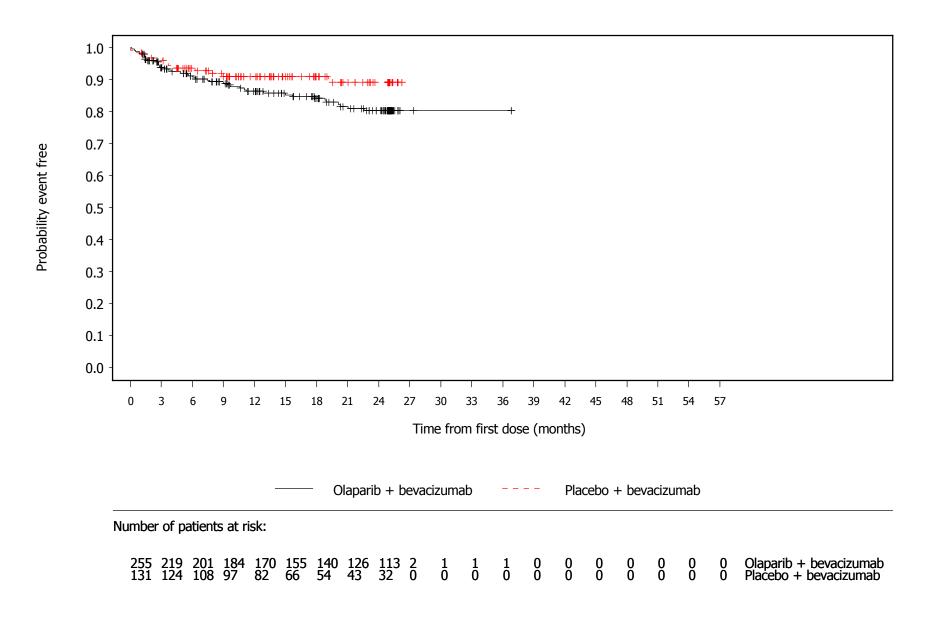


Figure 3.3.64 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Infection Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

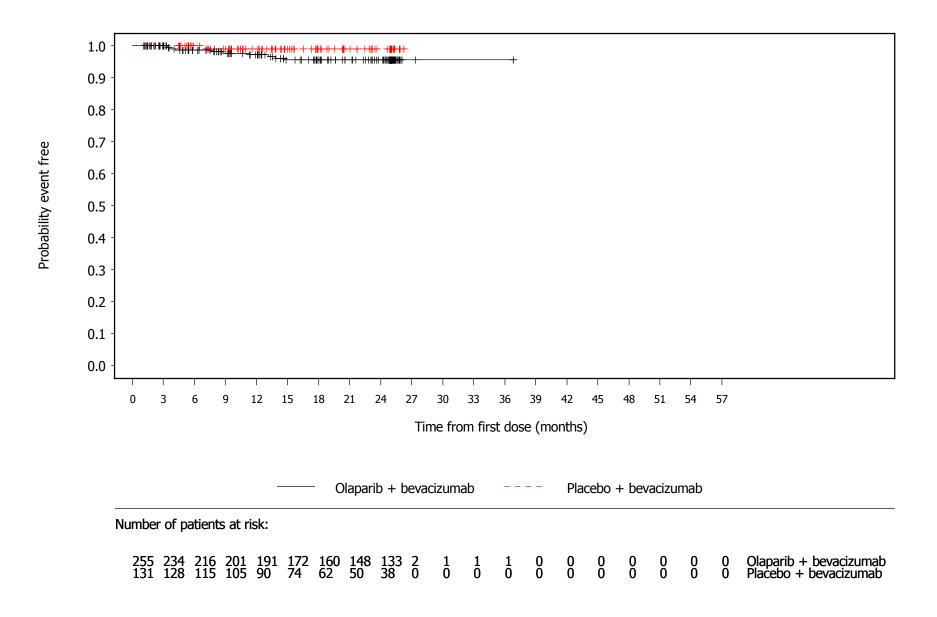


Figure 3.3.65 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Nasopharyngitis Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

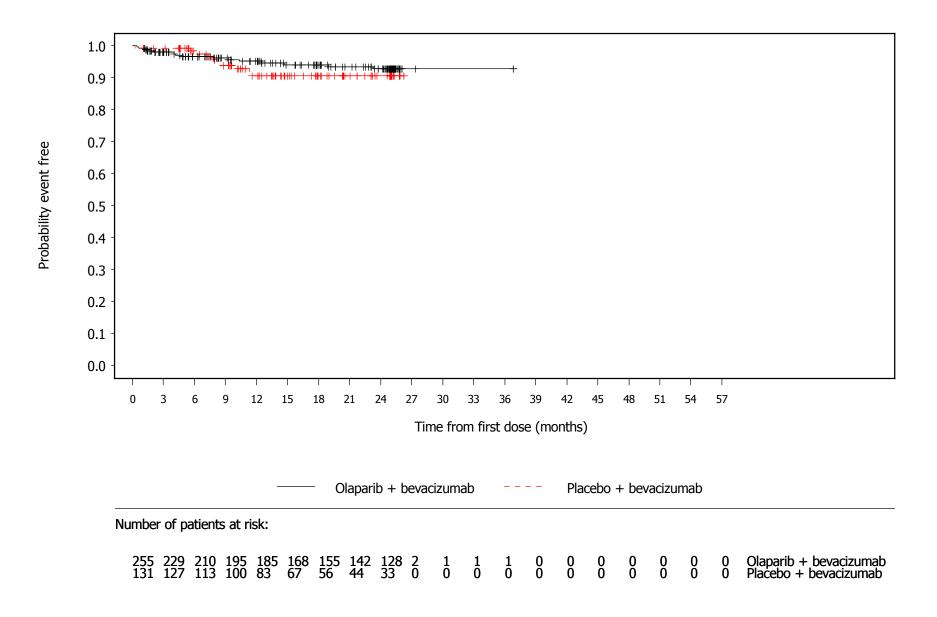


Figure 3.3.66 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Pharyngitis Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

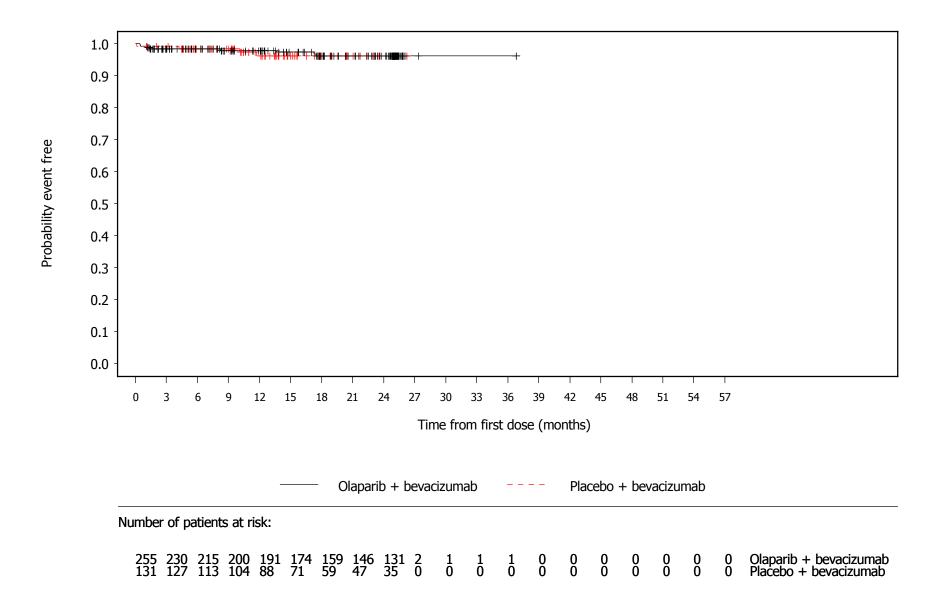


Figure 3.3.67 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Rhinitis Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

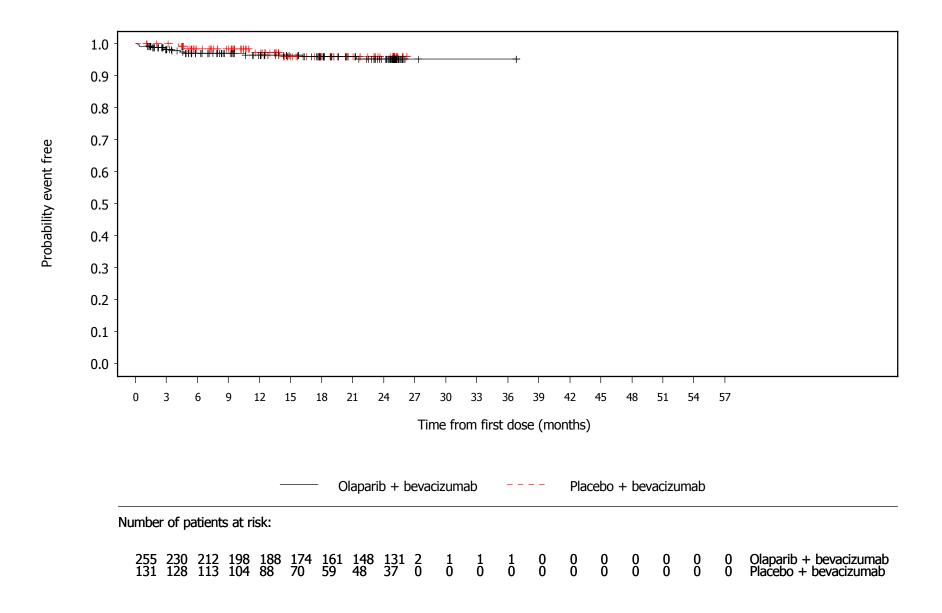


Figure 3.3.68 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Tooth abscess Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

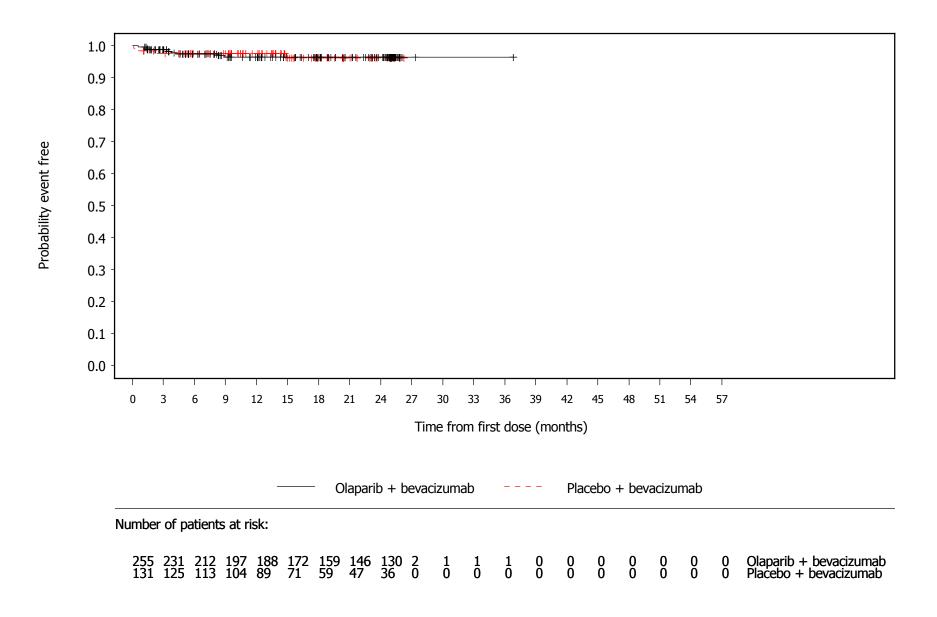


Figure 3.3.69 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Tooth infection Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

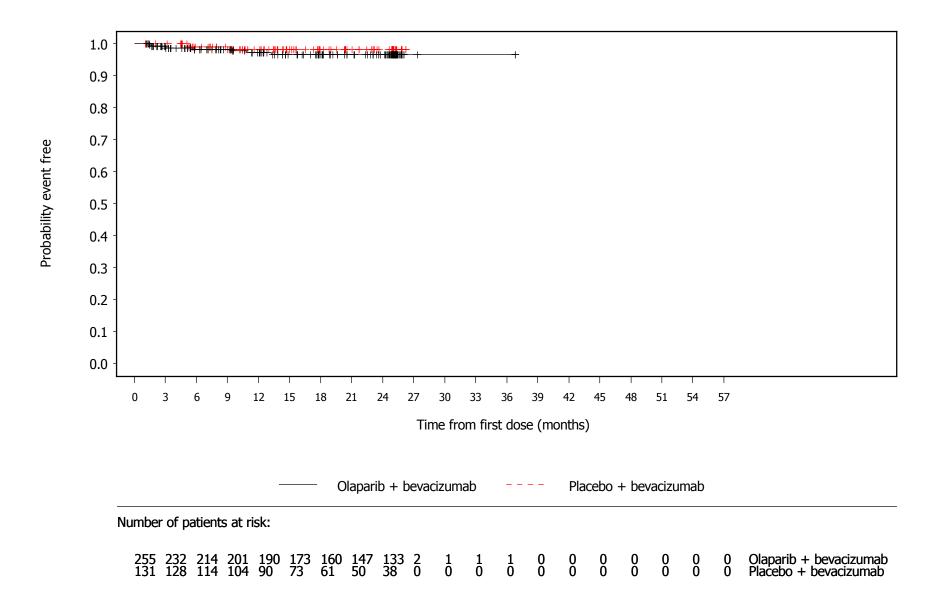


Figure 3.3.70 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Cystitis Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

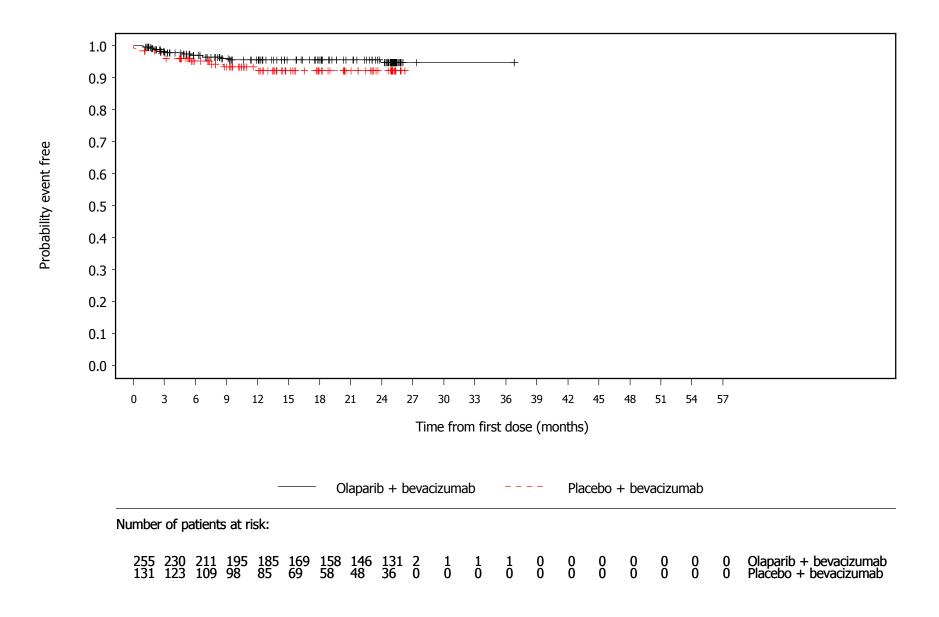


Figure 3.3.71 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Psychiatric disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

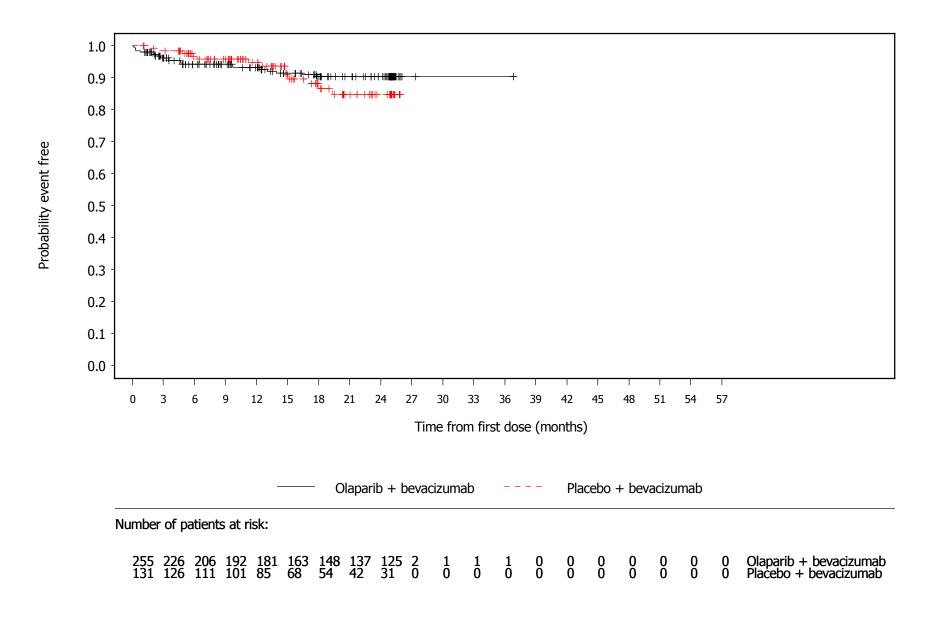


Figure 3.3.72 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Anxiety Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

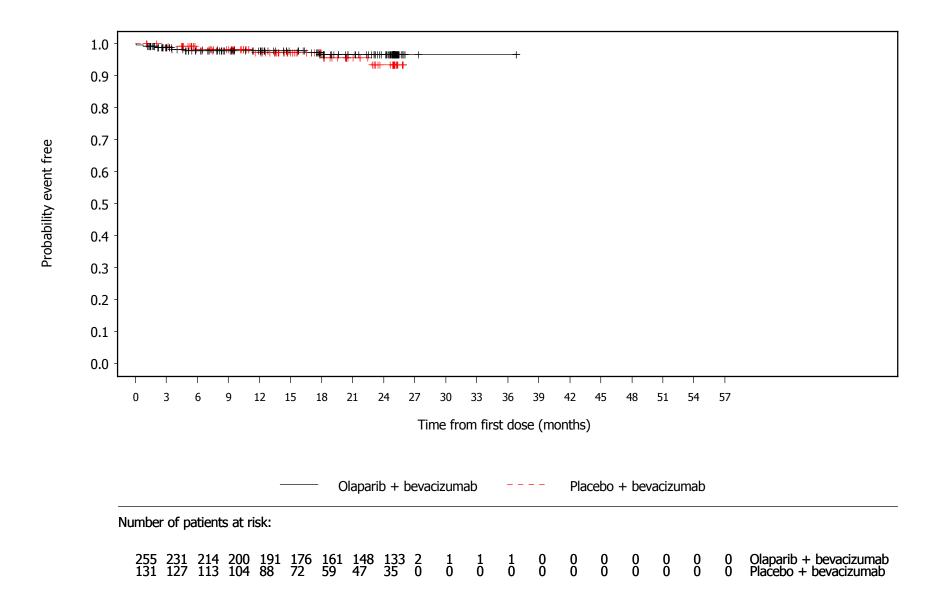


Figure 3.3.73 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Depression Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

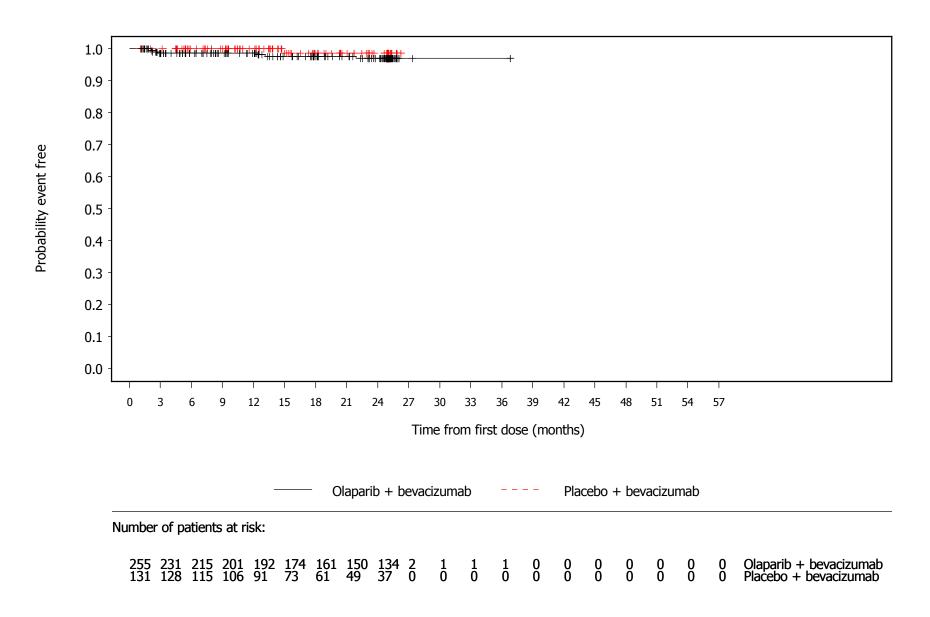


Figure 3.3.74 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Sleep disorder Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

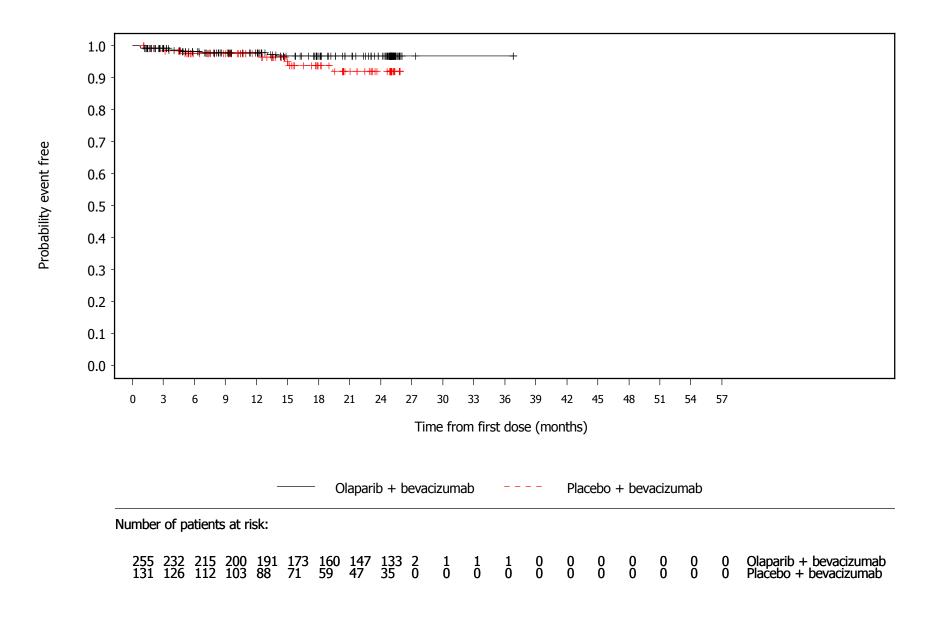


Figure 3.3.75 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Musculoskeletal and connective tissue disorders

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

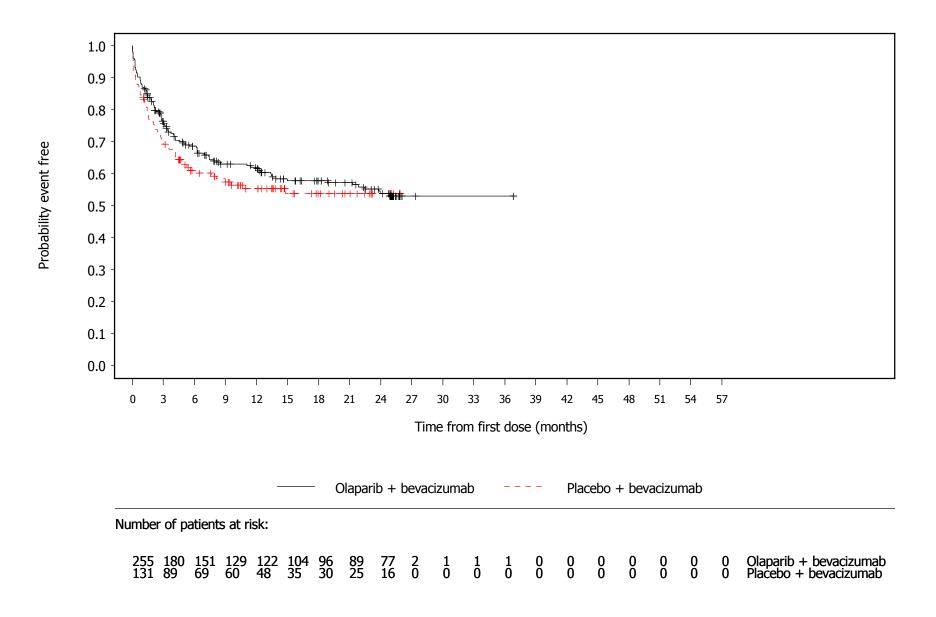


Figure 3.3.76 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Arthralgia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

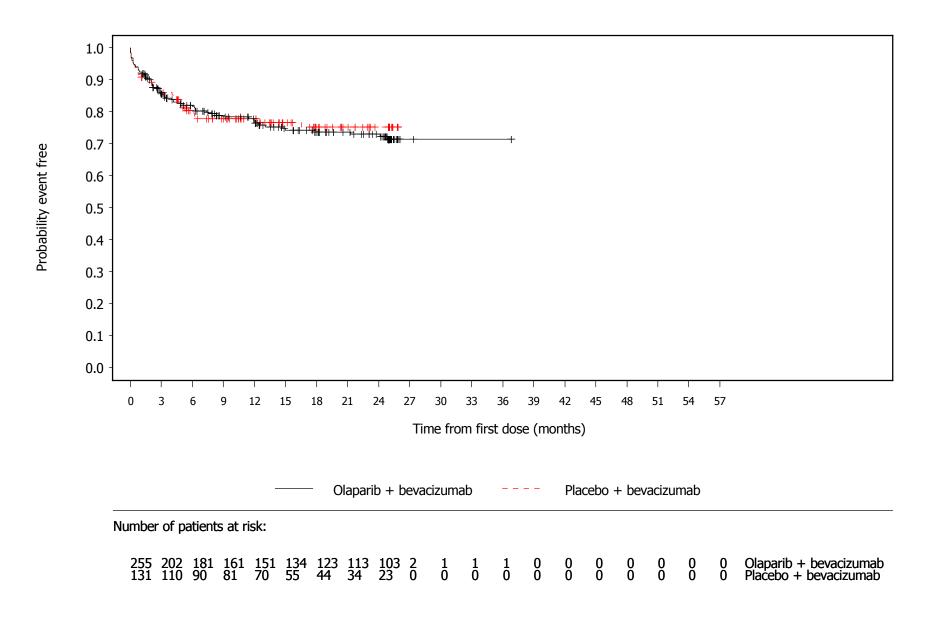


Figure 3.3.77 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Muscle spasms Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

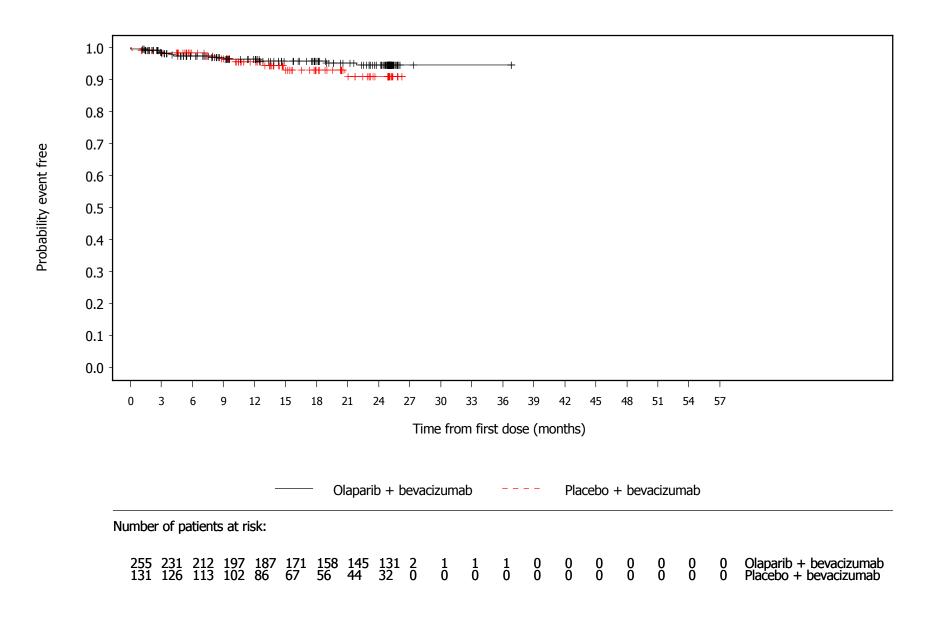


Figure 3.3.78 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Myalgia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

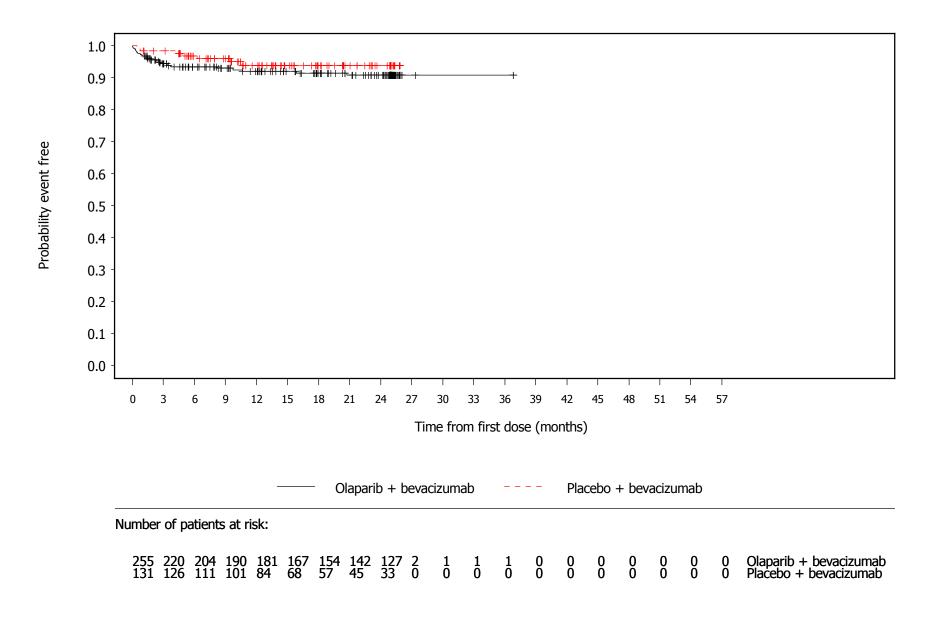


Figure 3.3.79 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Neck pain Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

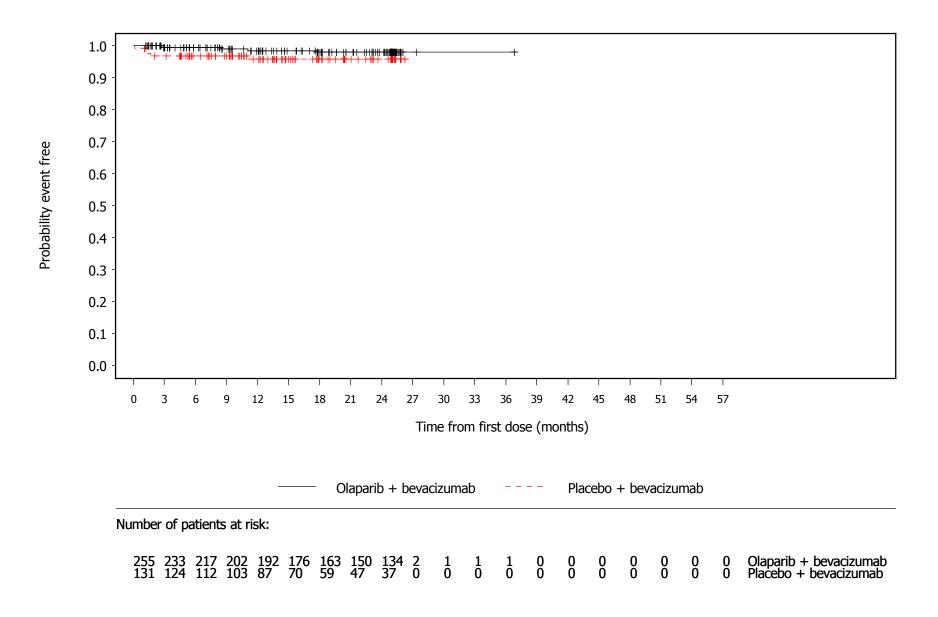


Figure 3.3.80 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Back pain Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

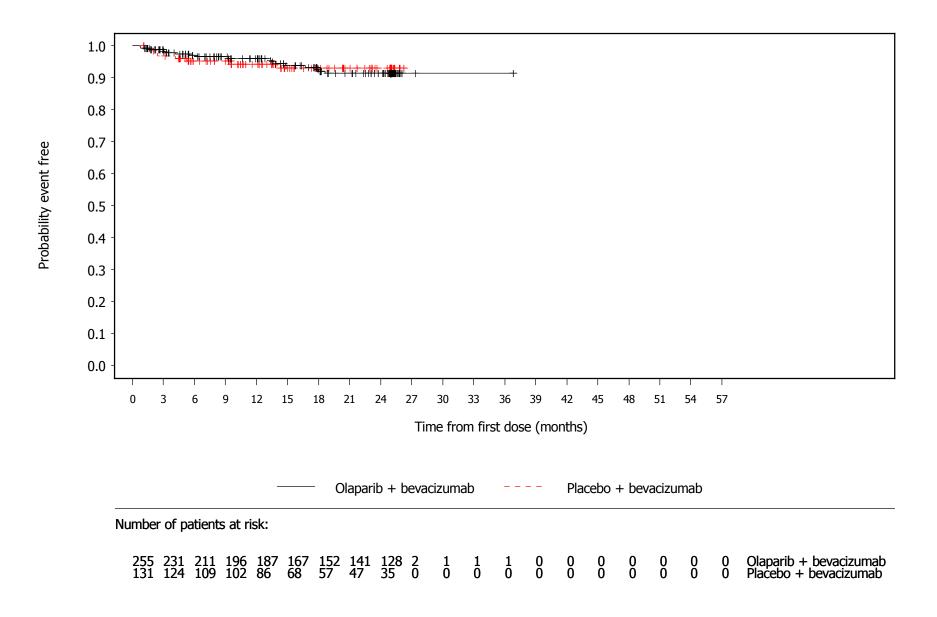


Figure 3.3.81 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Pain in extremity Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

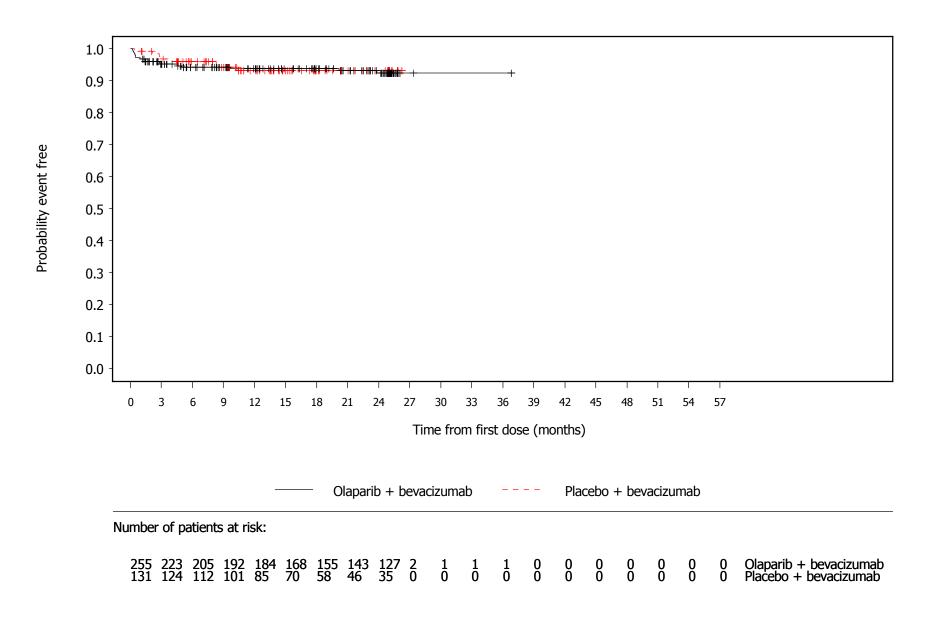


Figure 3.3.82 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Musculoskeletal pain Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

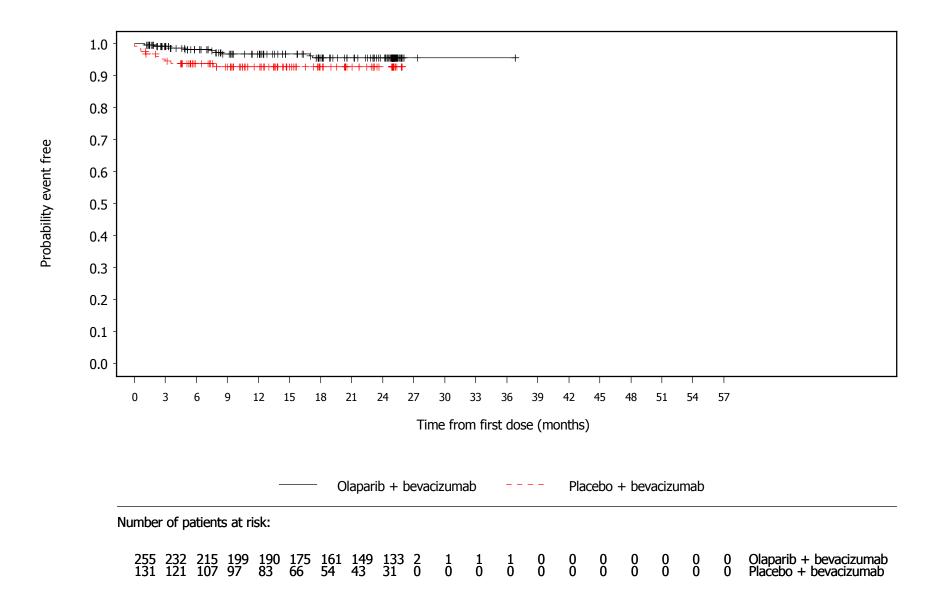


Figure 3.3.83 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Metabolism and nutrition disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

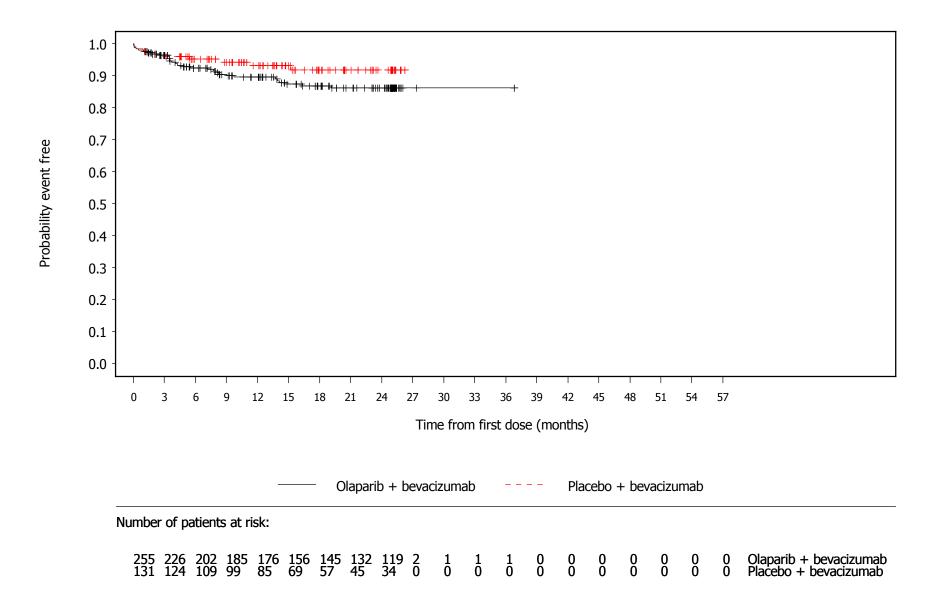


Figure 3.3.84 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Decreased appetite Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

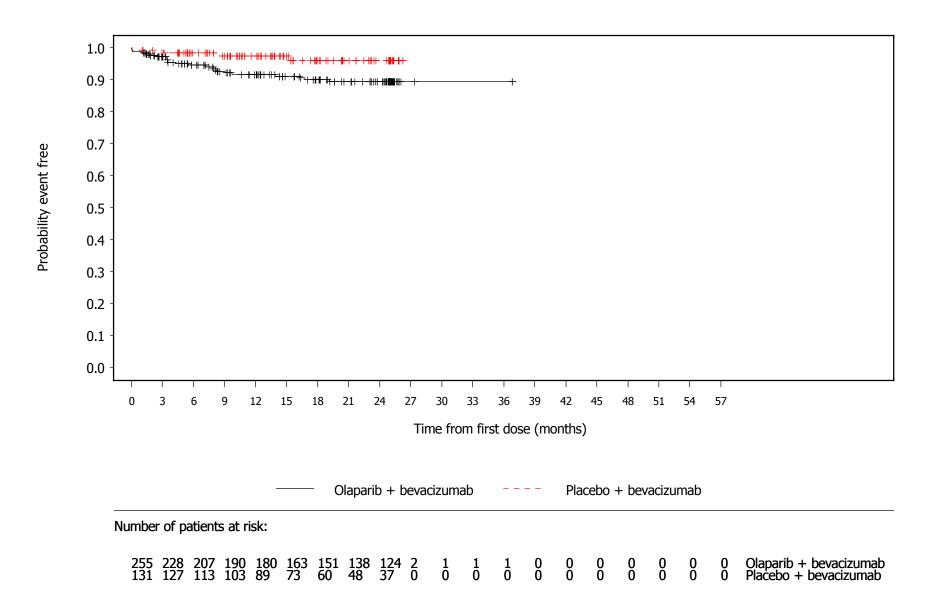


Figure 3.3.85 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Investigations Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

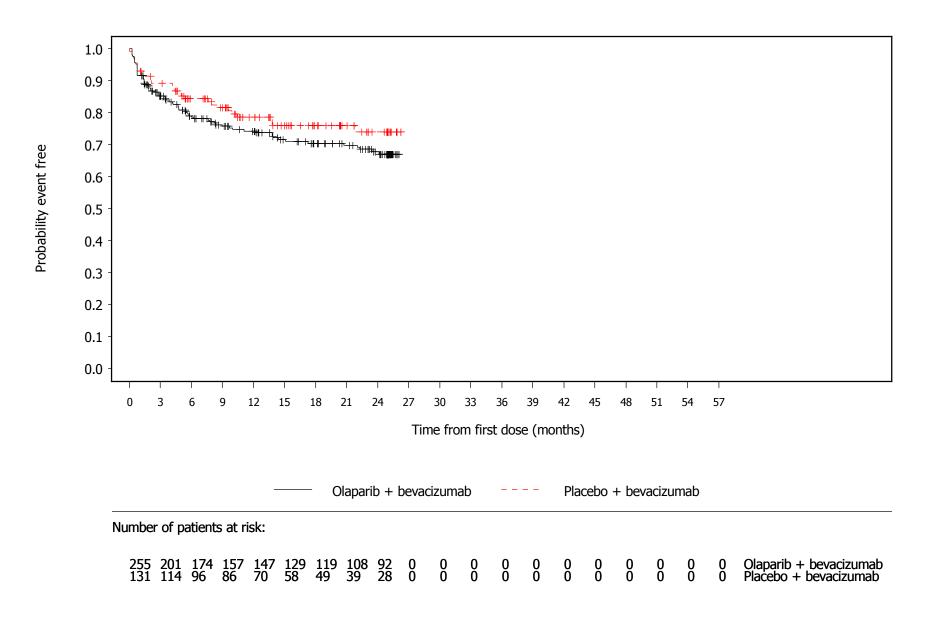


Figure 3.3.86 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Weight increased Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

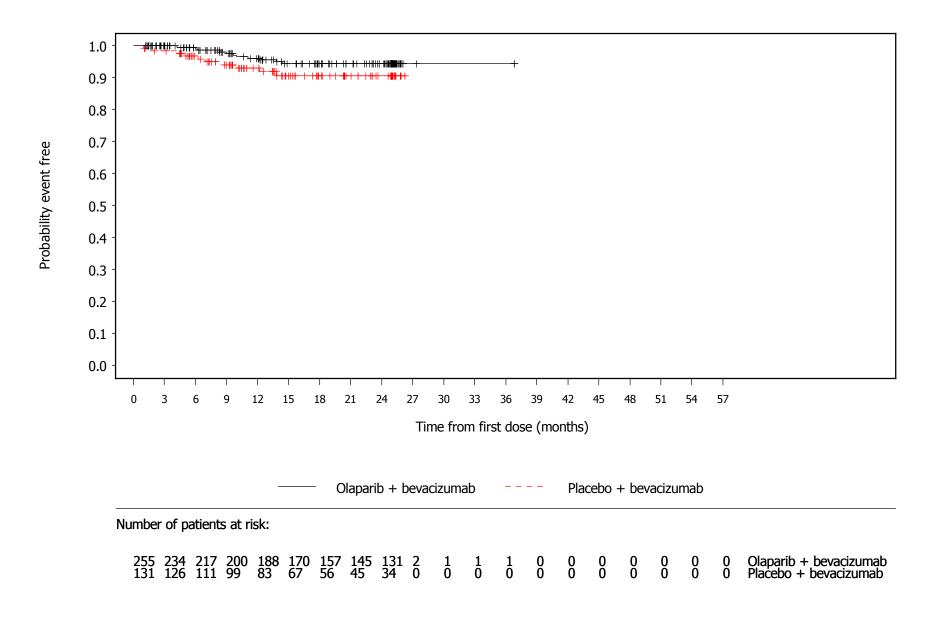


Figure 3.3.87 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Blood creatinine increased Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

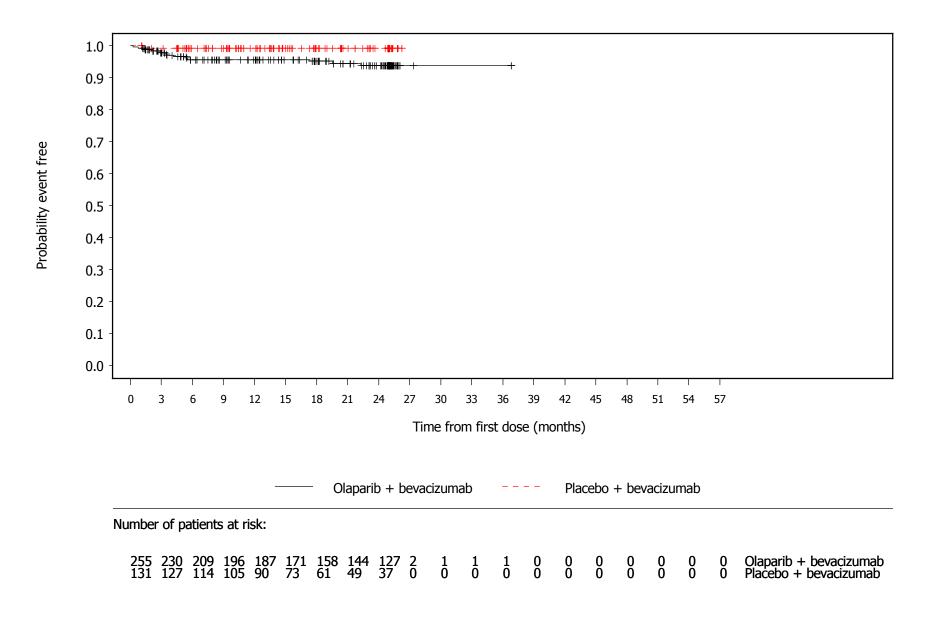


Figure 3.3.88 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: White blood cell count decreased Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

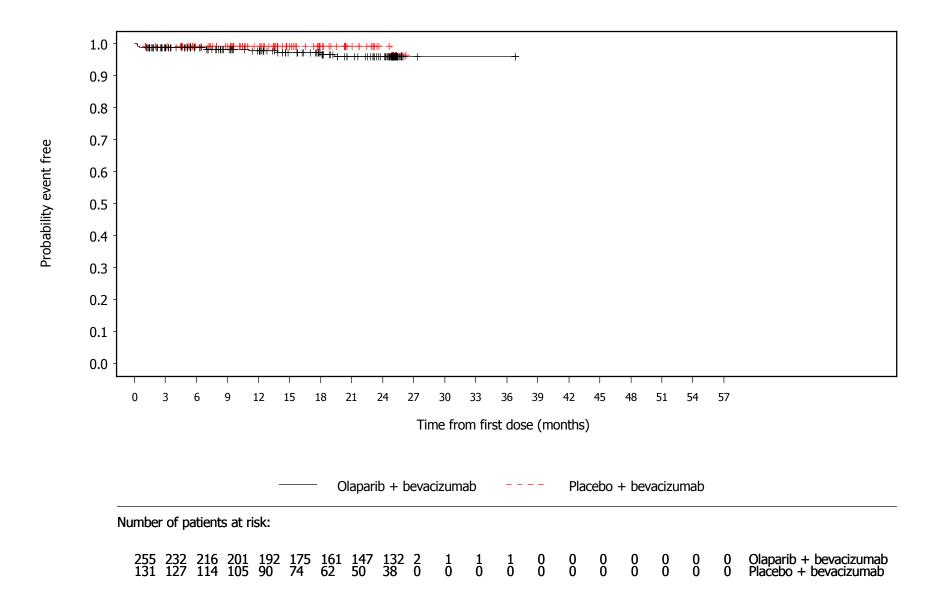


Figure 3.3.89 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Lymphocyte count decreased Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

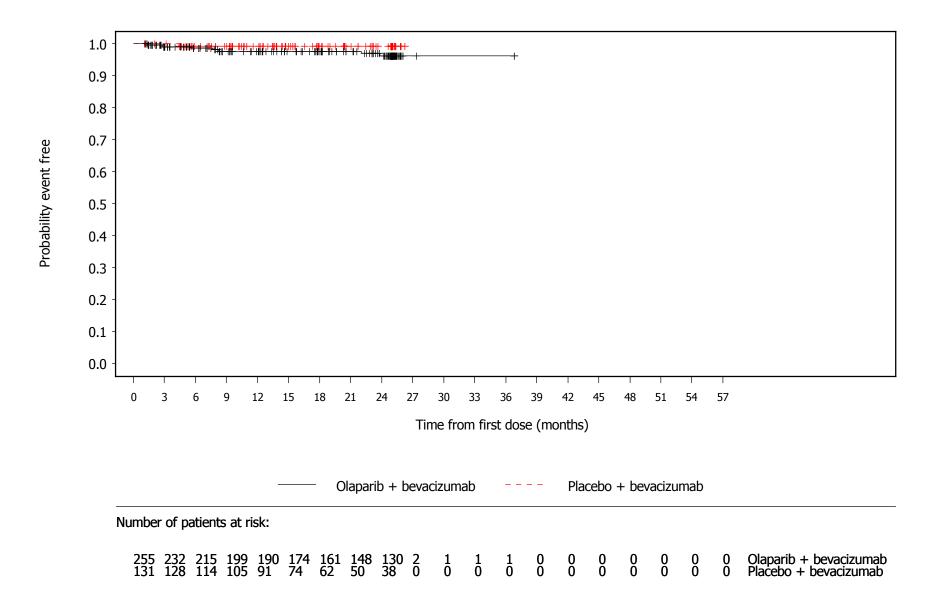


Figure 3.3.90 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Neutrophil count decreased Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

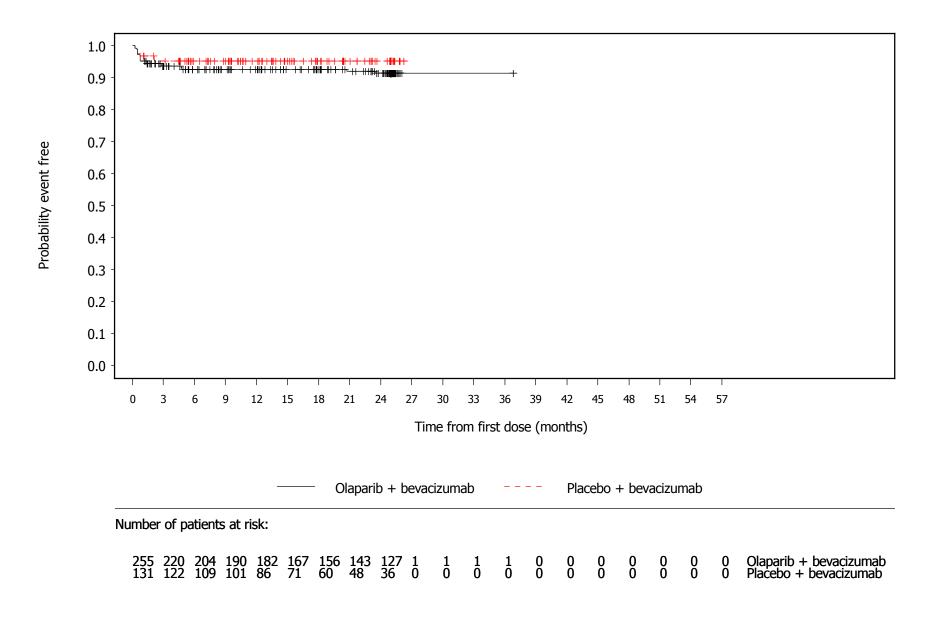


Figure 3.3.91 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE PT: Platelet count decreased Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

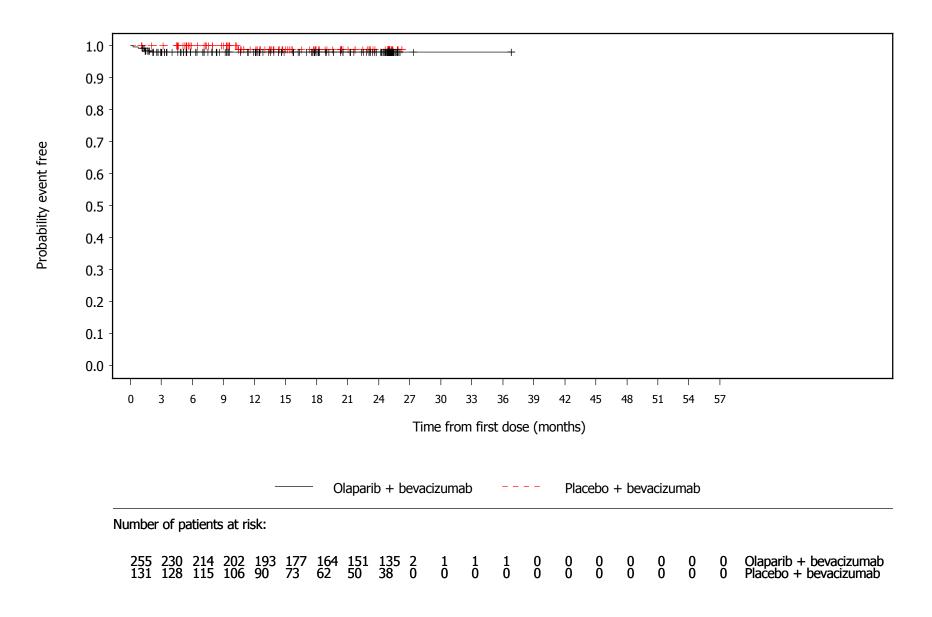


Figure 3.3.92 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE SOC: Injury, poisoning and procedural complications Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

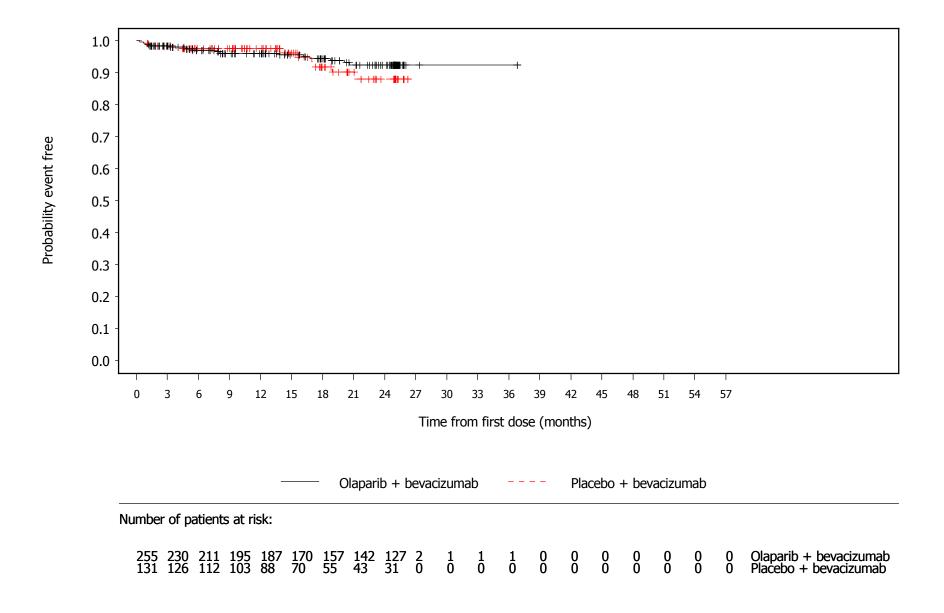


Figure 3.3.93 PAOLA1: Kaplan-Meier plot of time to first occurrence of SAE Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

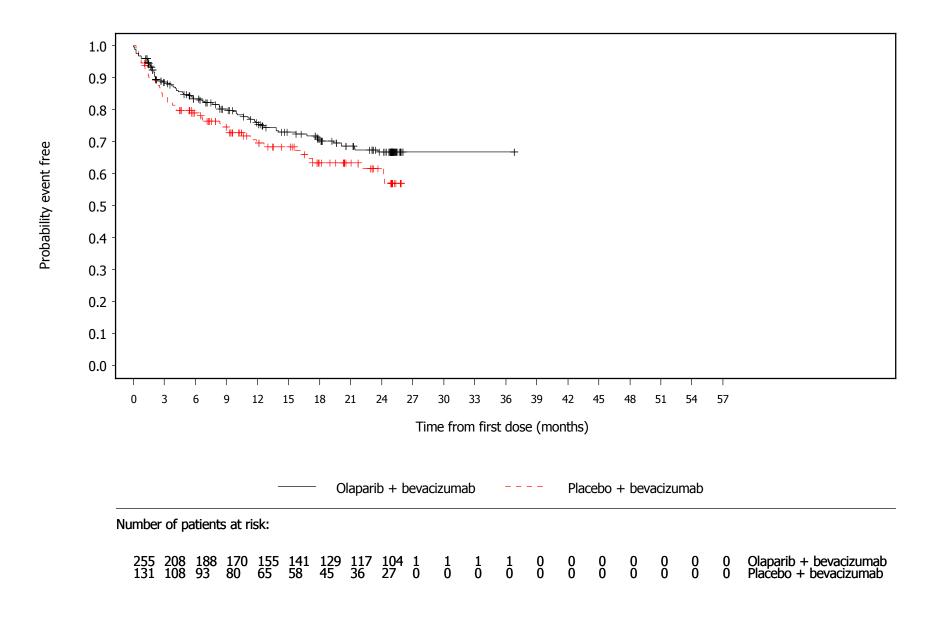


Figure 3.3.94 PAOLA1: Kaplan-Meier plot of time to first occurrence of SAE SOC: Respiratory, thoracic and mediastinal disorders

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

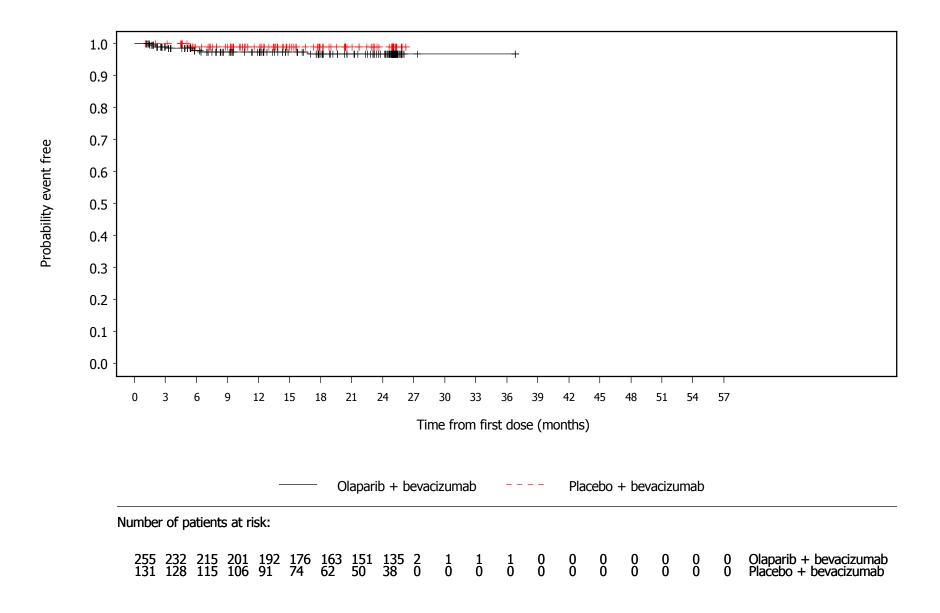


Figure 3.3.95 PAOLA1: Kaplan-Meier plot of time to first occurrence of SAE SOC: Blood and lymphatic system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

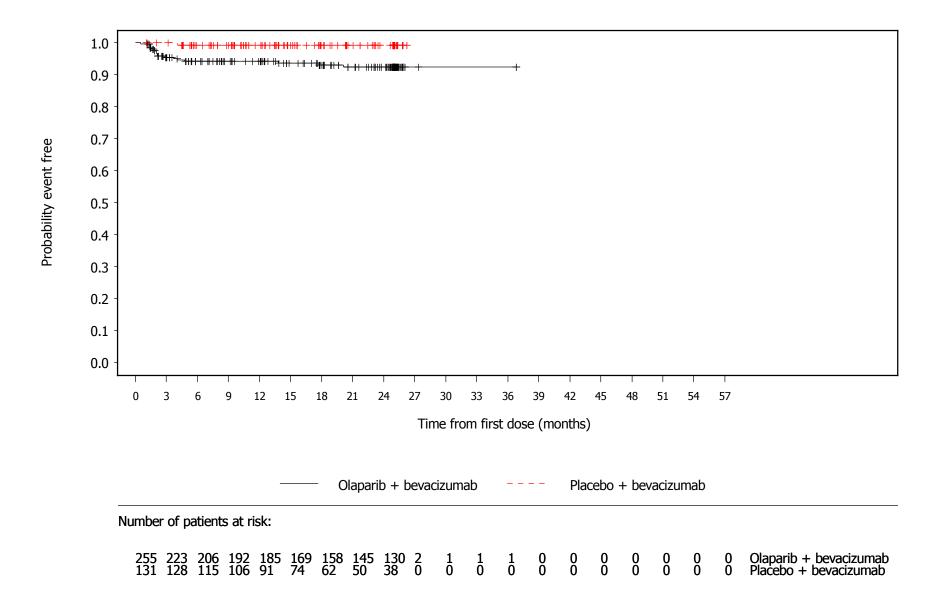


Figure 3.3.96 PAOLA1: Kaplan-Meier plot of time to first occurrence of SAE PT: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

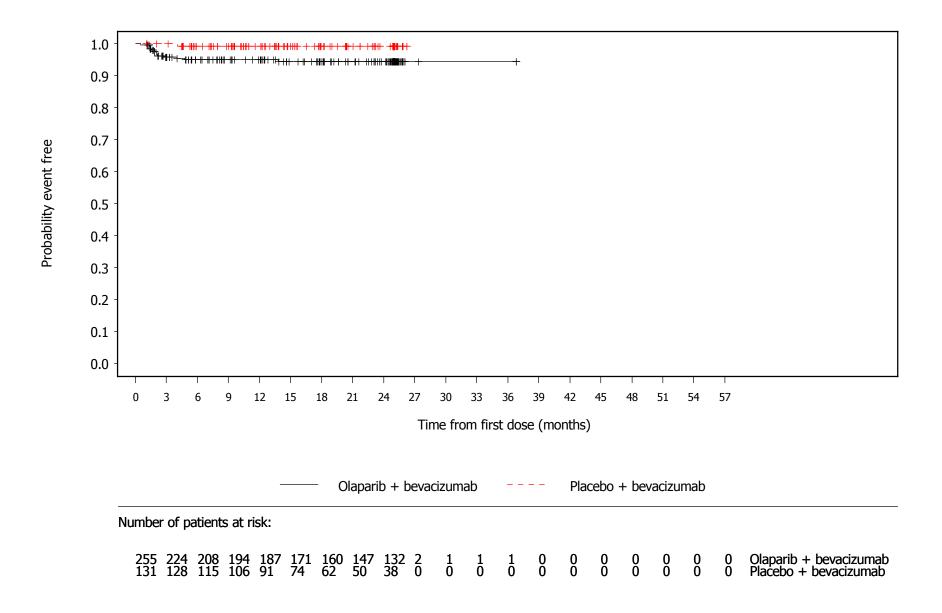


Figure 3.3.97 PAOLA1: Kaplan-Meier plot of time to first occurrence of SAE SOC: Gastrointestinal disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

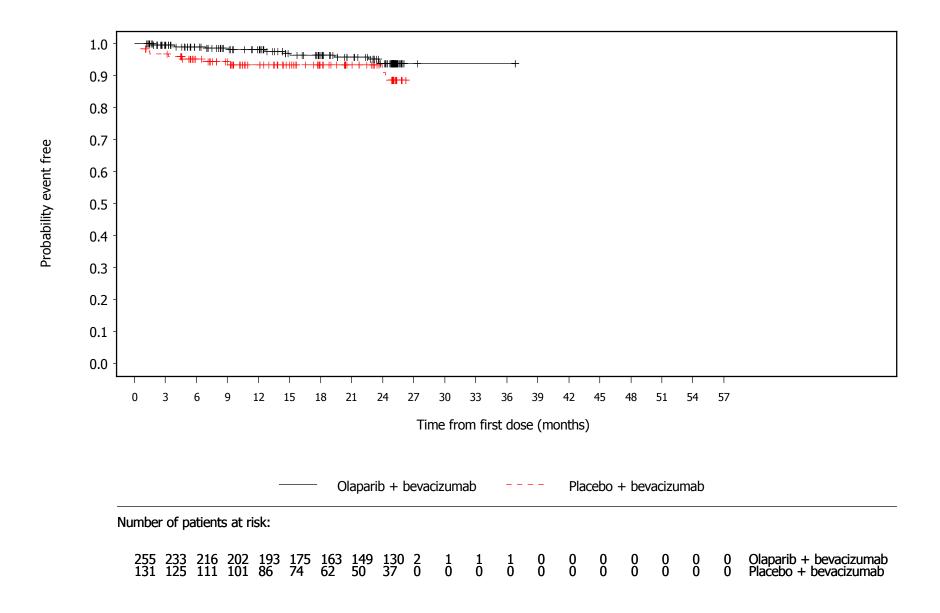


Figure 3.3.98 PAOLA1: Kaplan-Meier plot of time to first occurrence of SAE SOC: Vascular disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

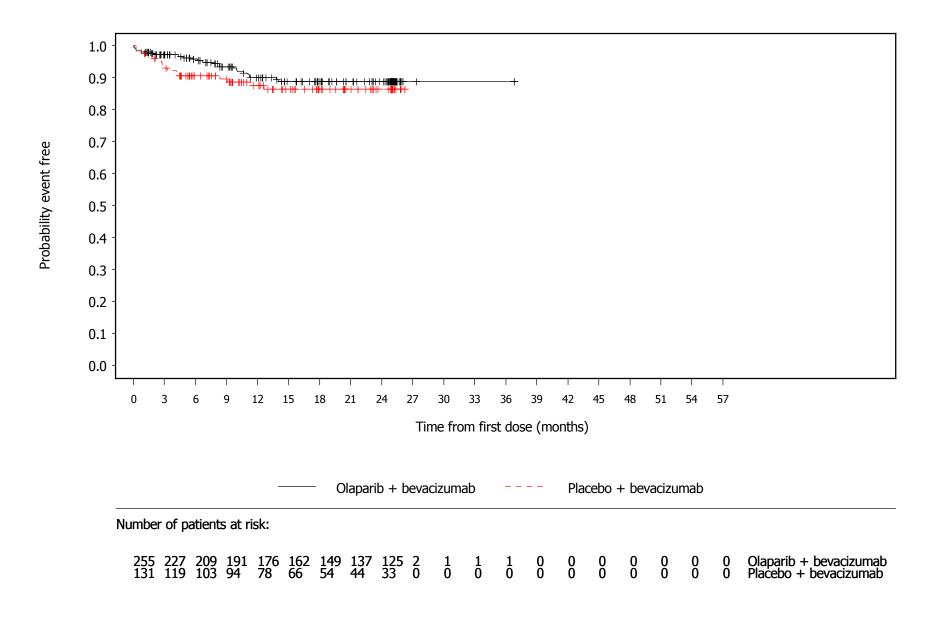


Figure 3.3.99 PAOLA1: Kaplan-Meier plot of time to first occurrence of SAE PT: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

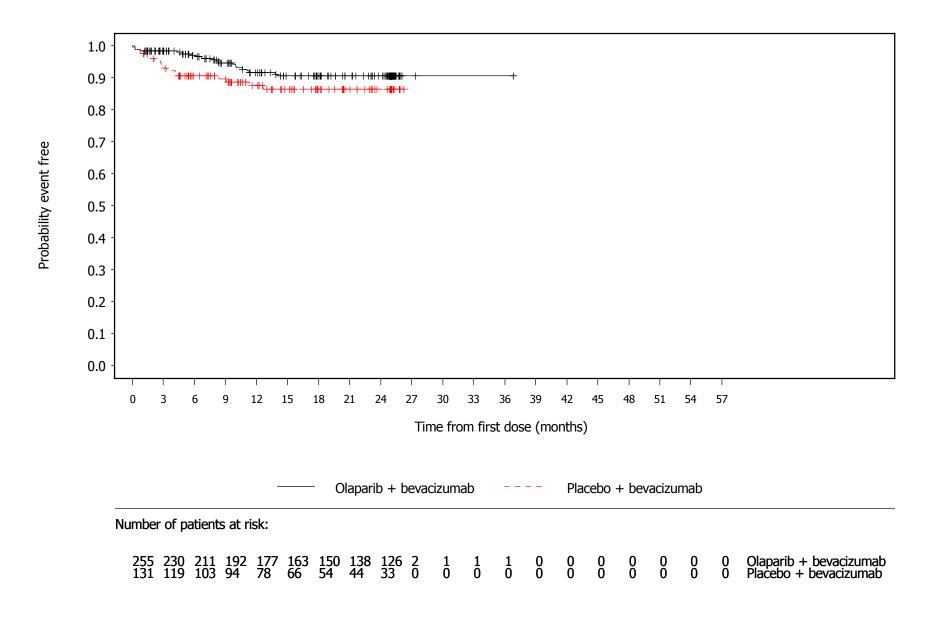


Figure 3.3.100 PAOLA1: Kaplan-Meier plot of time to first occurrence of SAE SOC: Infections and infestations Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

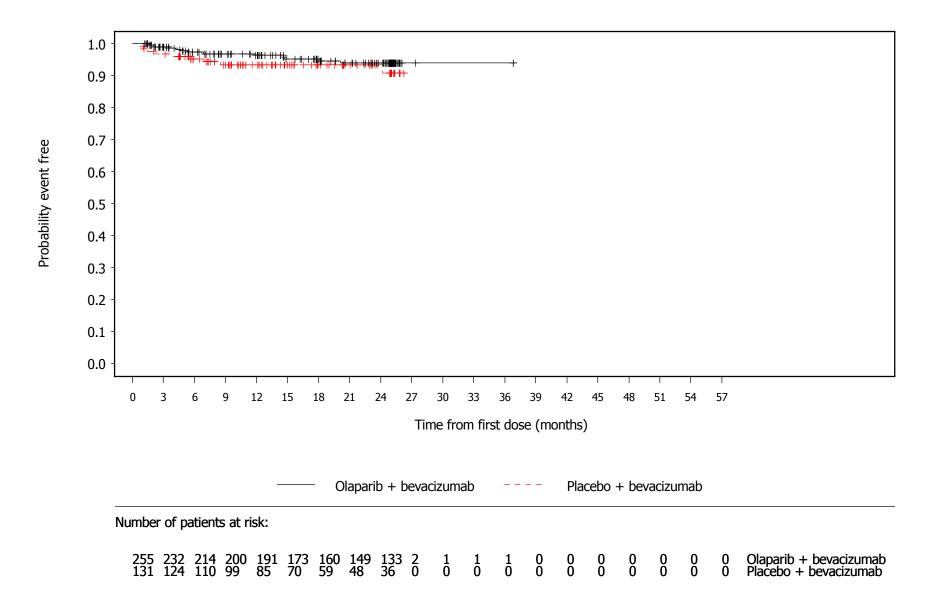


Figure 3.3.101 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE leading to discontinuation of treatment Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

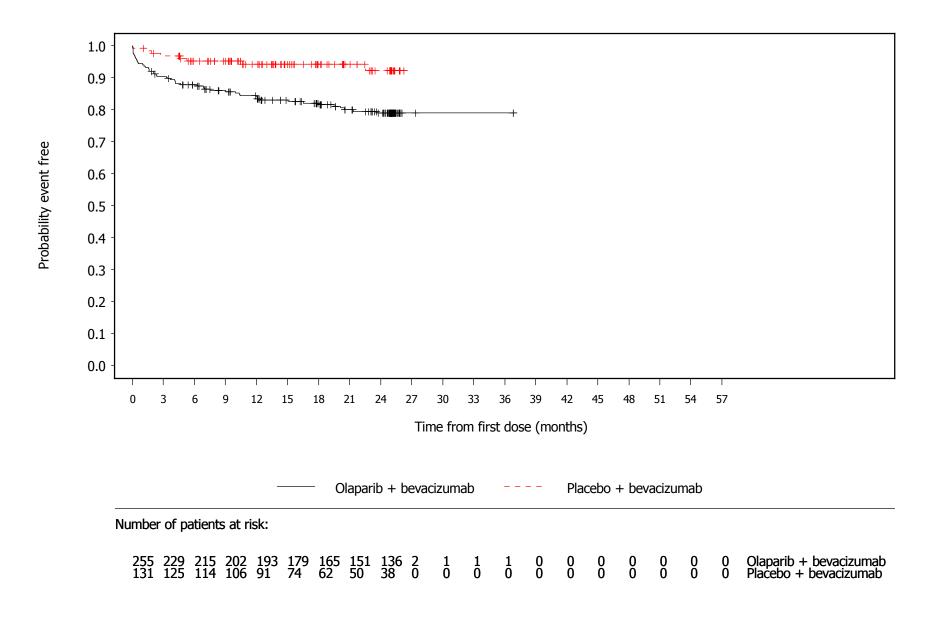


Figure 3.3.102 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE max CTCAE grade>=3 Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

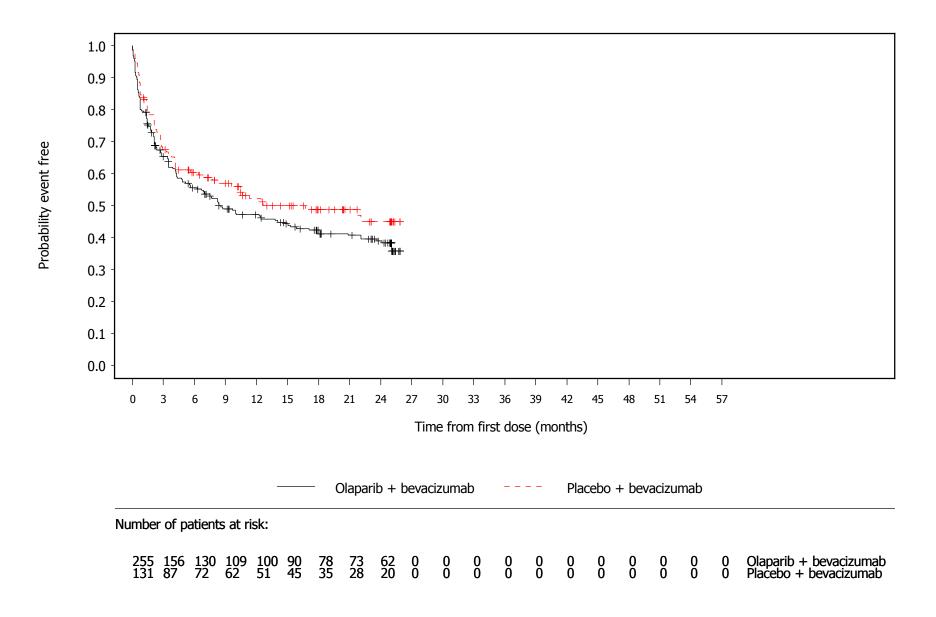


Figure 3.3.103 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 SOC: General disorders and administration site conditions

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

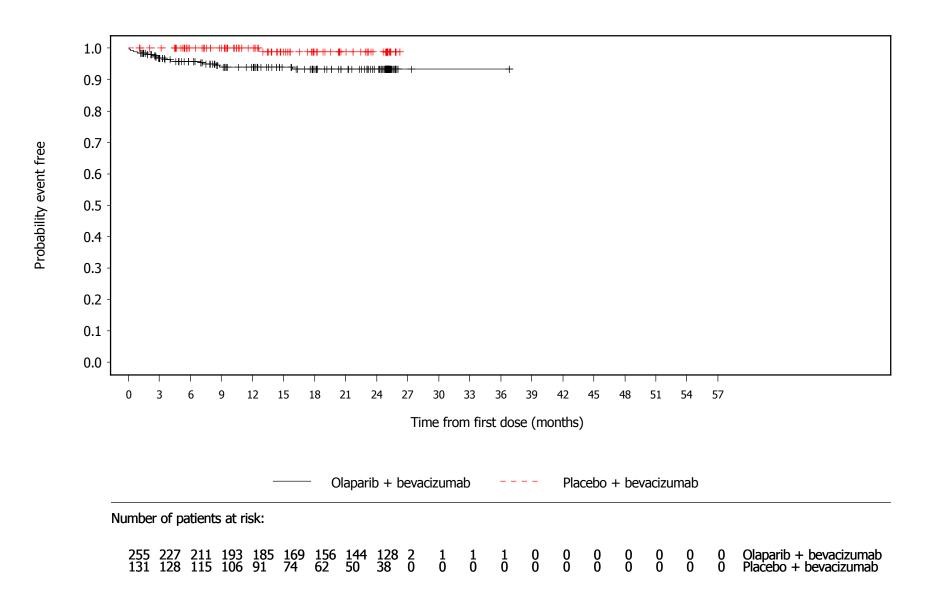


Figure 3.3.104 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 PT: Fatigue Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

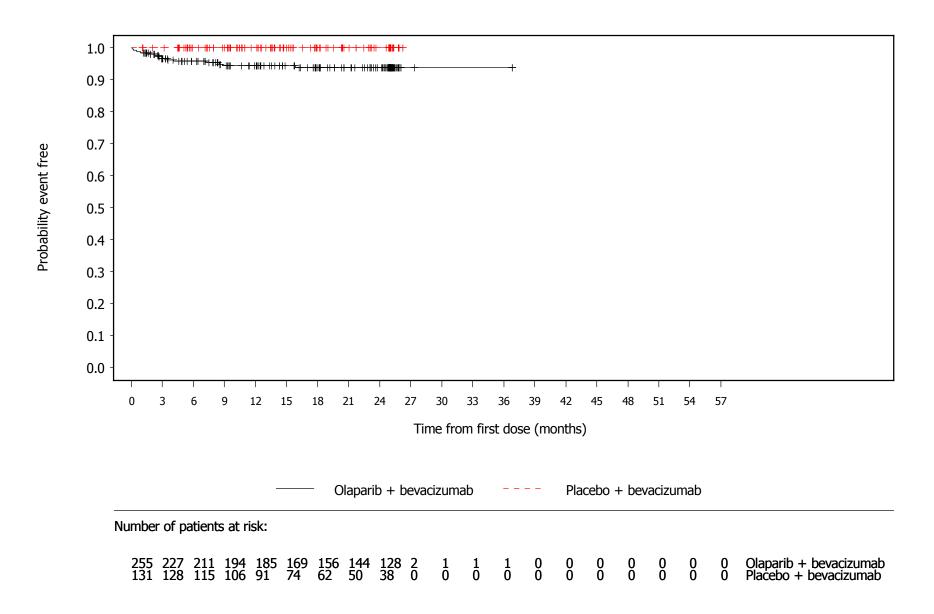


Figure 3.3.105 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 SOC: Respiratory, thoracic and mediastinal disorders

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

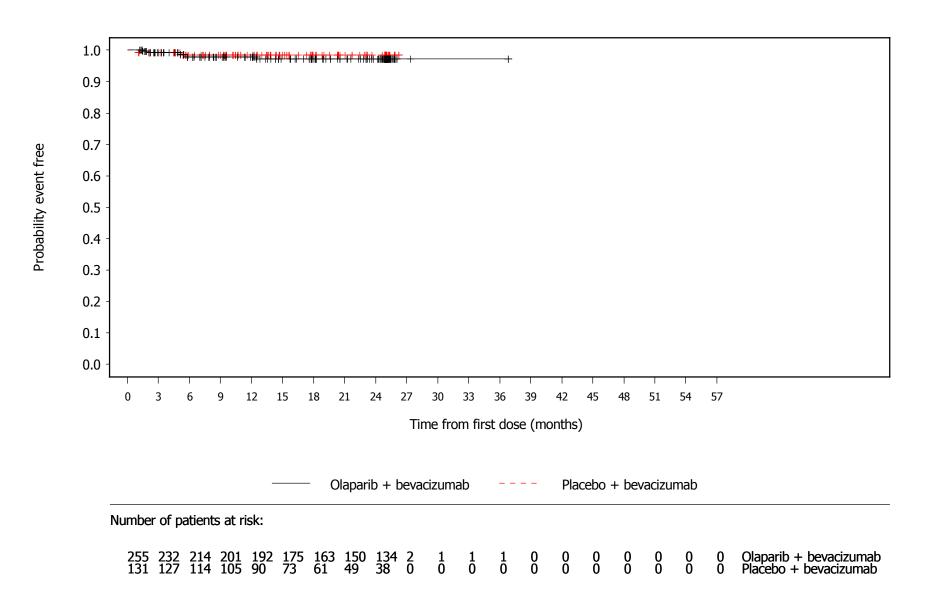


Figure 3.3.106 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 SOC: Renal and urinary disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

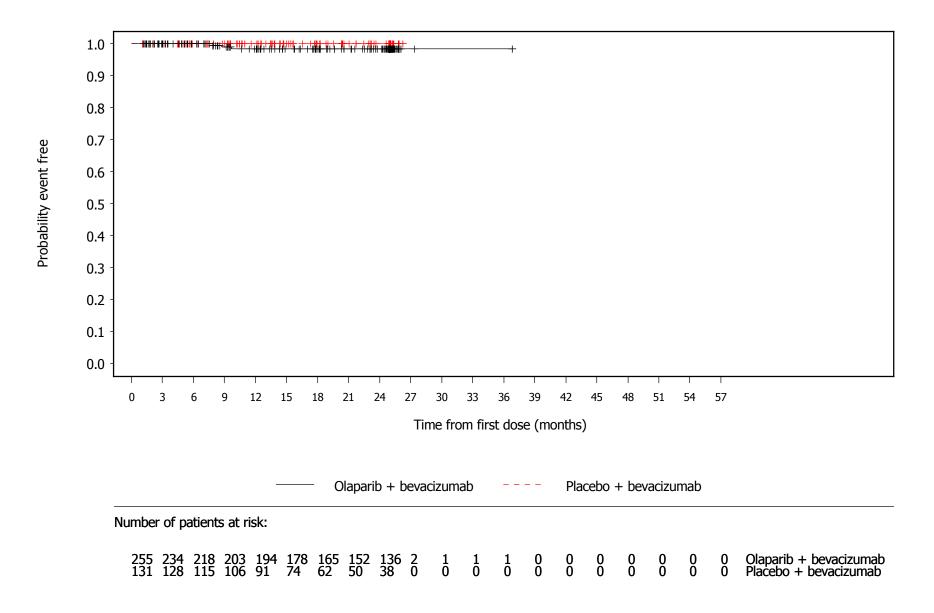


Figure 3.3.107 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 SOC: Blood and lymphatic system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

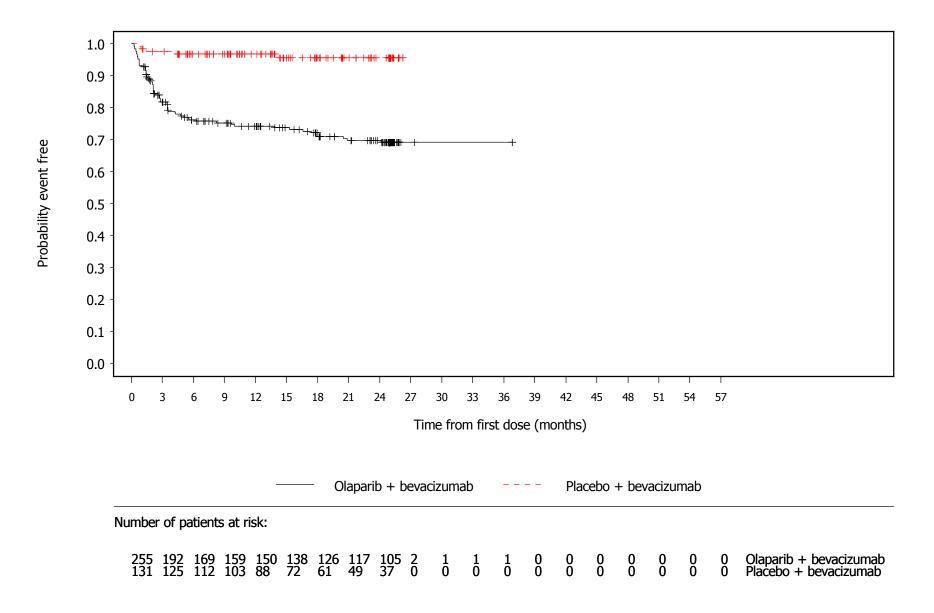


Figure 3.3.108 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 PT: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

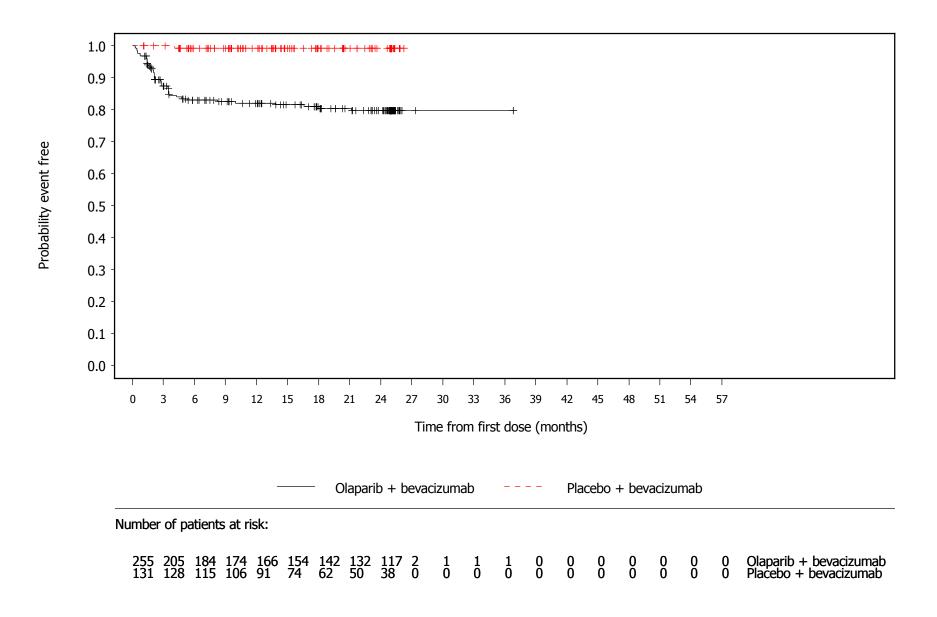


Figure 3.3.109 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 PT: Leukopenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

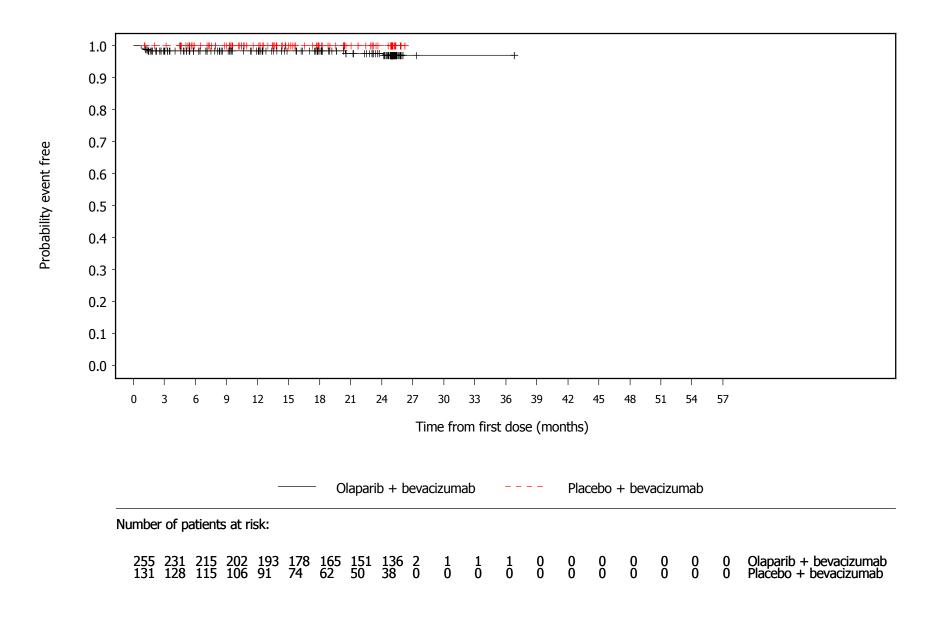


Figure 3.3.110 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 PT: Lymphopenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

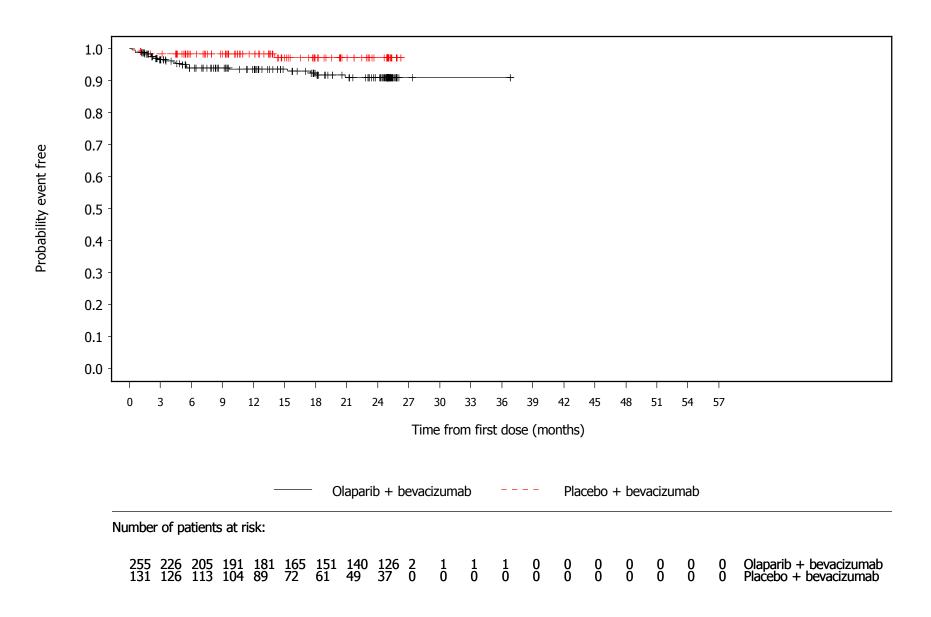


Figure 3.3.111 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 PT: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

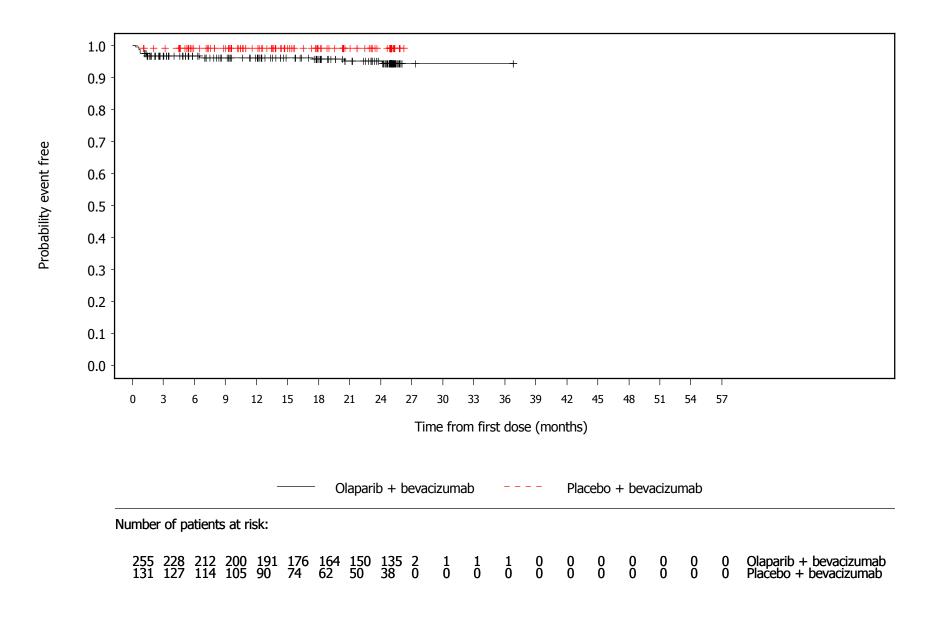


Figure 3.3.112 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 SOC: Gastrointestinal disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

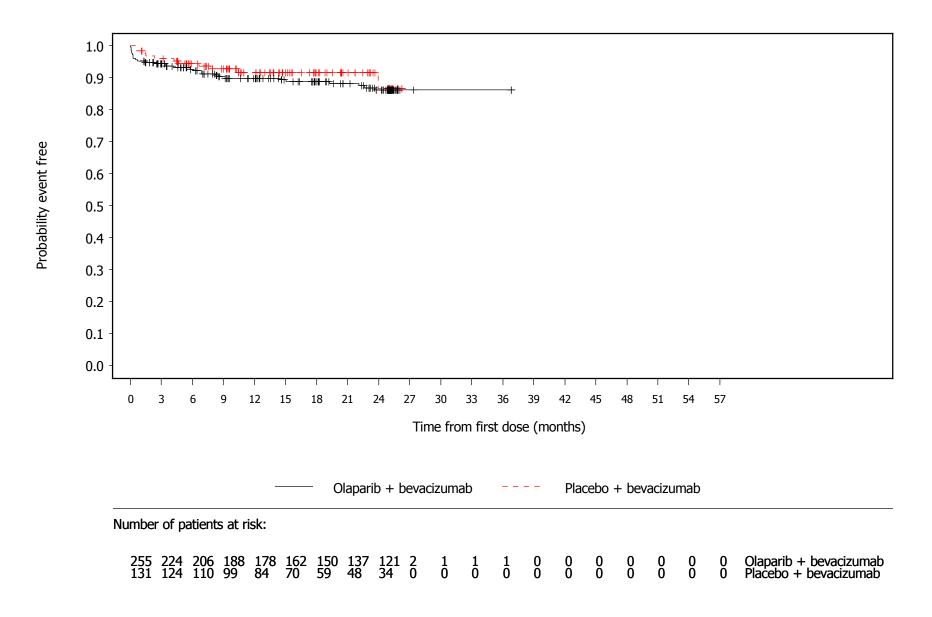


Figure 3.3.113 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 PT: Diarrhoea Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

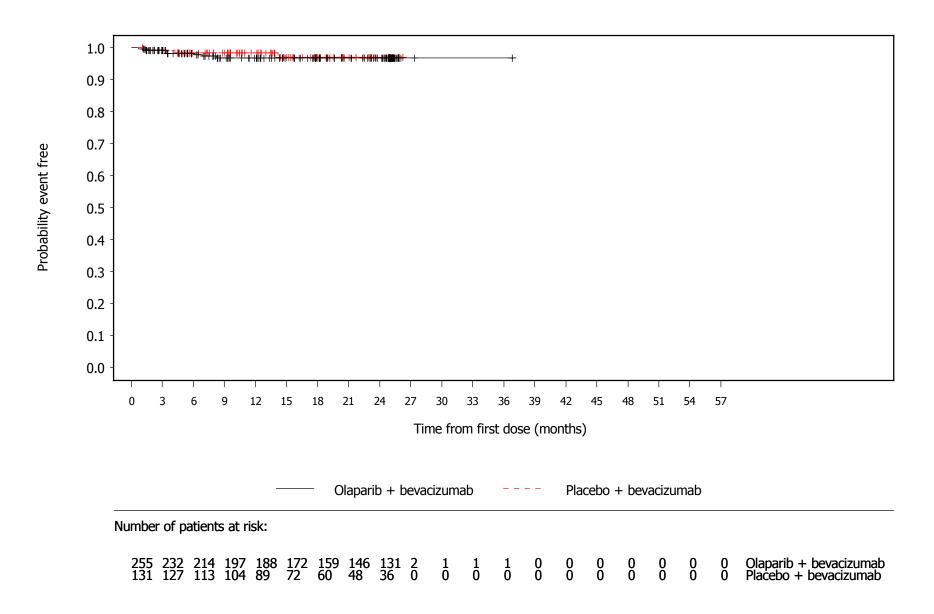


Figure 3.3.114 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 PT: Nausea Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

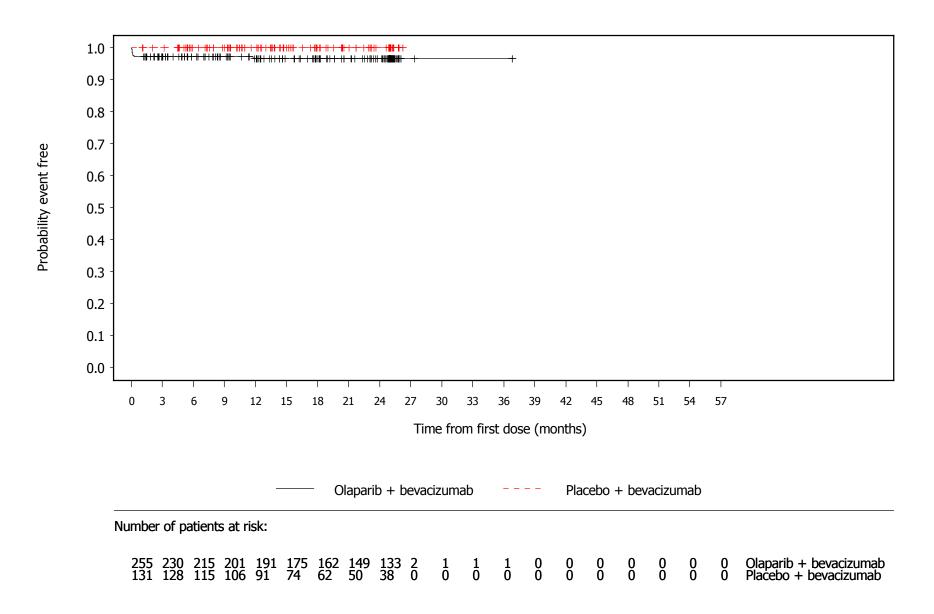


Figure 3.3.115 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 SOC: Nervous system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

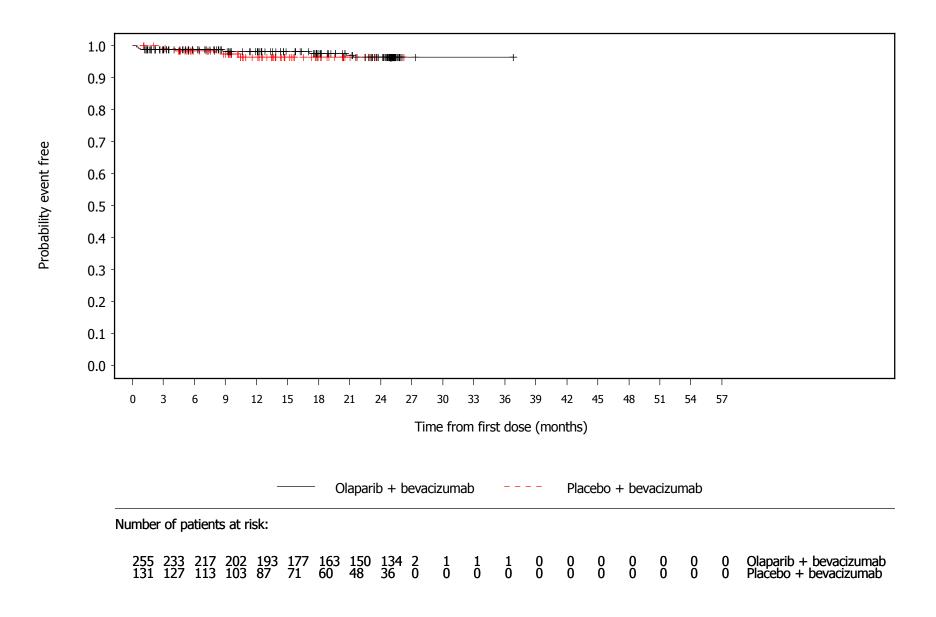


Figure 3.3.116 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 SOC: Vascular disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

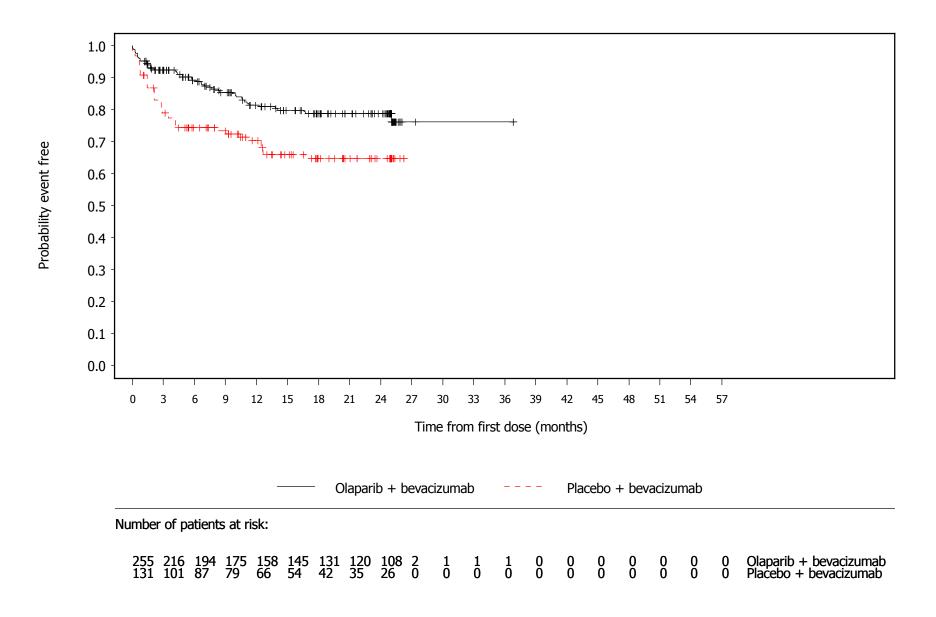


Figure 3.3.117 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 PT: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

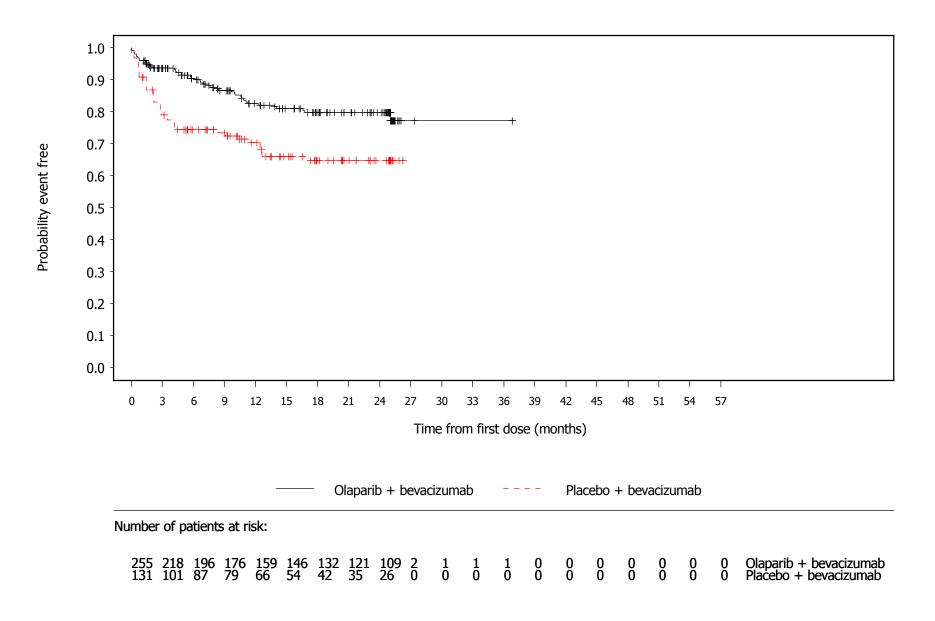


Figure 3.3.118 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 SOC: Infections and infestations Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

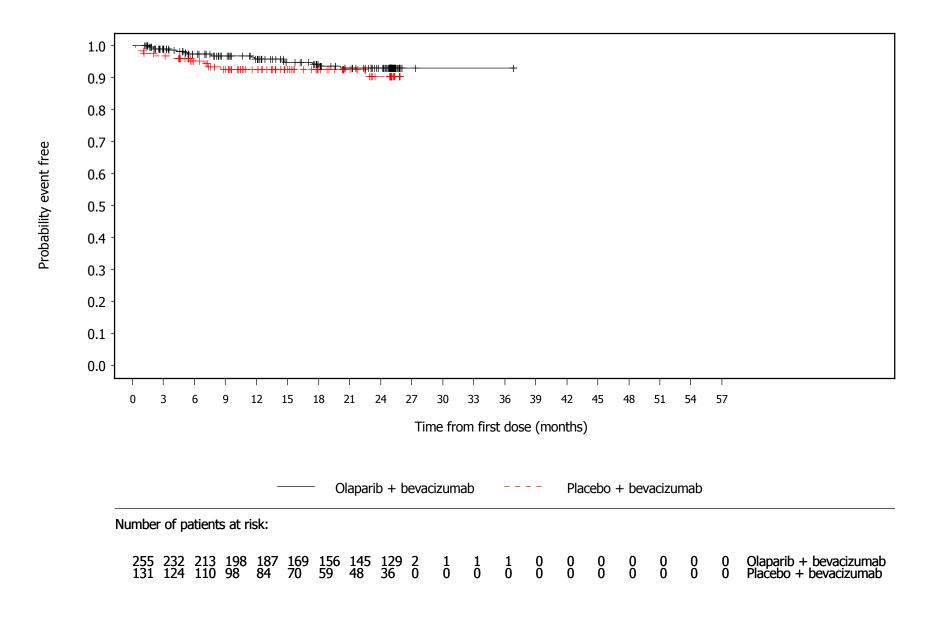


Figure 3.3.119 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 SOC: Investigations Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

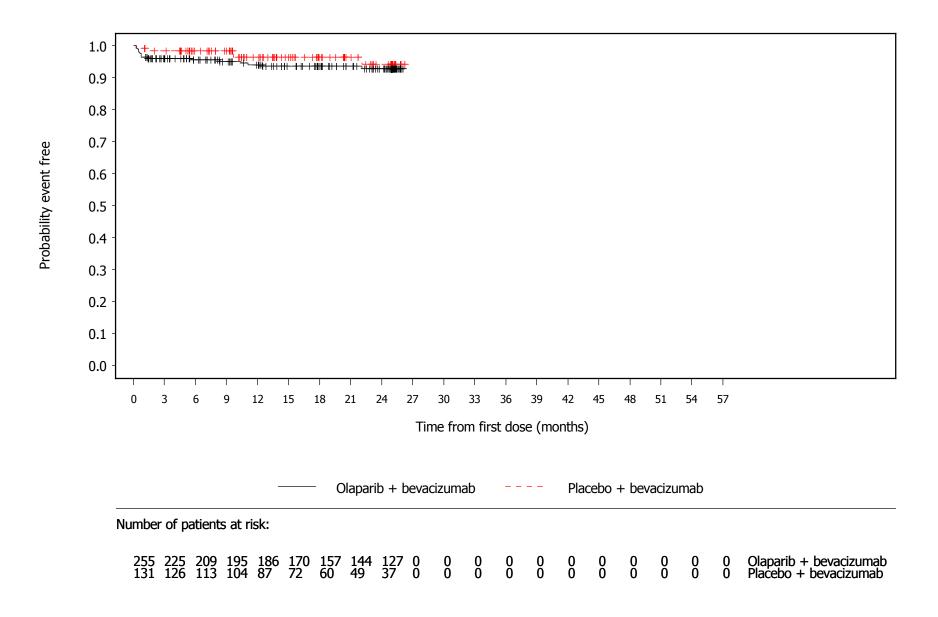


Figure 3.3.120 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE G>=3 PT: Neutrophil count decreased Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

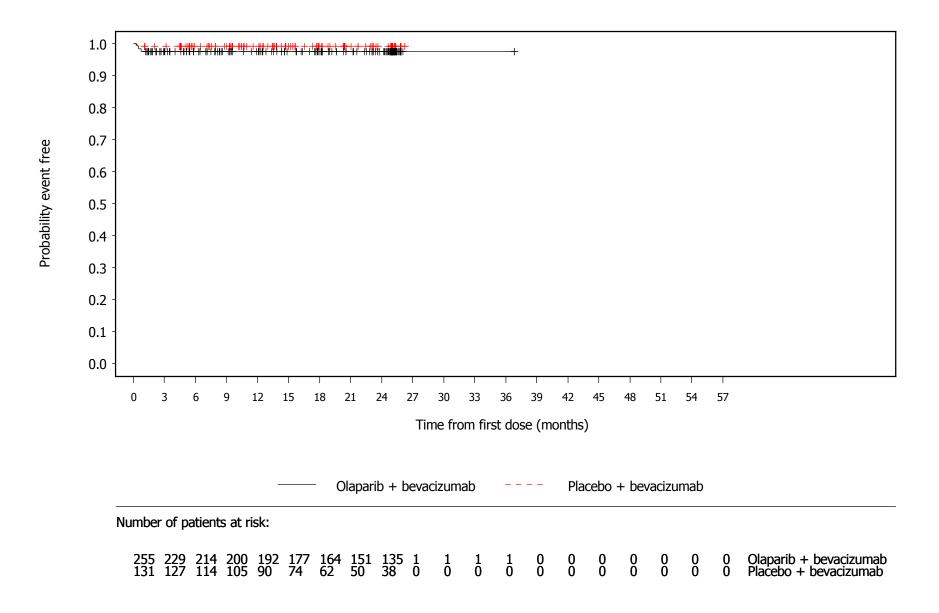


Figure 3.3.121 PAOLA1: Kaplan-Meier plot of time to first occurrence of AE max CTCAE grade=1 or 2 Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

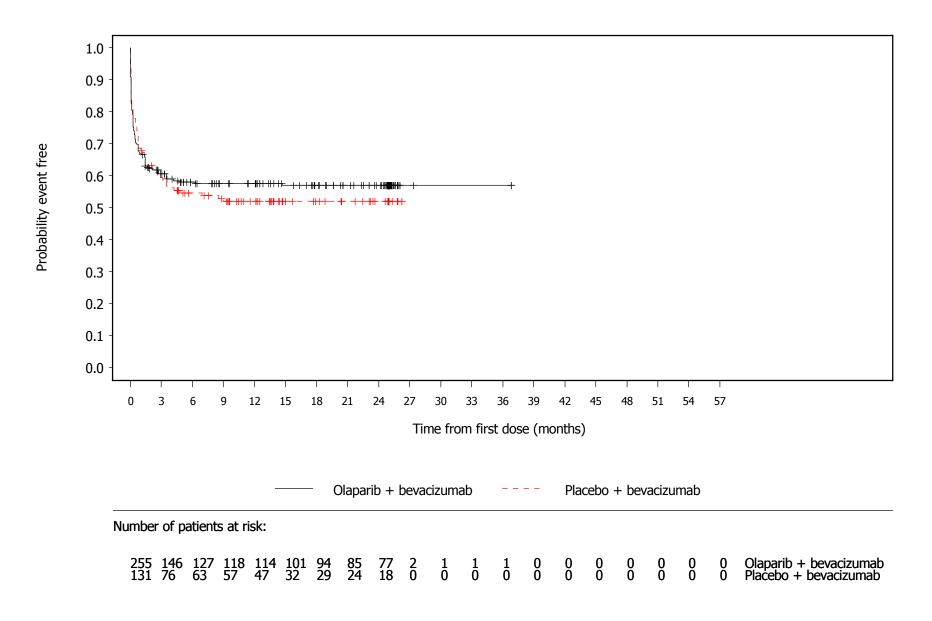


Table 3.4.1 PAOLA1: Summary of subgroup analysis of AE Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + N=2			Placebo + h (N=1					
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]			2-sided p-value [b]
First line treatment out	come (IV	RS)								
NED [PDS]	92	92 (100)	0.2 (0.1, 0.3)	48	47 (97.9)	0.6 (0.3, 0.9)	1.51	1.07,	2.16	0.0196*
NED/CR [IDS]	74	74 (100)	0.2 (0.1, 0.3)	38	37 (97.4)	0.2 (0.1, 0.7)	1.42	0.96,	2.14	0.0769
NED/CR [Chemo]	40	40 (100)	0.2 (0.1, 0.3)	20	19 (95.0)	0.6 (0.0, 0.7)	1.34	0.79,	2.38	0.2843
PR	49	49 (100)	0.3 (0.2, 0.5)	25	24 (96.0)	0.3 (0.1, 0.7)	1.30	0.80,	2.16	0.2906
Interaction p-value										0.9636
Screening laboratory tBR0	CA statu	ıs (IVRS)								
tBRCAm	150	150 (100)	0.2 (0.2, 0.3)	65	63 (96.9)	0.3 (0.2, 0.7)	1.29	0.97,	1.75	0.0848
non-tBRCAm	105	105 (100)	0.2 (0.1, 0.3)	66	64 (97.0)	0.3 (0.1, 0.7)	1.60	1.17,	2.21	0.0030*
Interaction p-value										0.3256
First line treatment out	come (eC	PRF)								
NED [PDS]	89	89 (100)	0.2 (0.1, 0.3)	47	46 (97.9)	0.6 (0.3, 0.9)	1.49	1.05,	2.15	0.0248*
NED/CR [IDS]	74	74 (100)	0.2 (0.1, 0.3)	32	31 (96.9)	0.3 (0.1, 1.0)	1.60	1.06,	2.48	0.0255*
NED/CR [Chemo]	39	39 (100)	0.2 (0.1, 0.5)	17	16 (94.1)	0.2 (0.0, 0.7)	1.12	0.64,	2.07	0.6972
PR	50	50 (100)	0.2 (0.2, 0.4)	34	33 (97.1)	0.3 (0.1, 0.7)	1.22	0.79,	1.91	0.3777
Interaction p-value										0.6987
Screening laboratory tBR0	CA statu	ıs (eCRF)								
tBRCAm	147	147 (100)	0.2 (0.2, 0.3)	67	65 (97.0)	0.3 (0.2, 0.7)	1.29	0.97,	1.74	0.0851
non-tBRCAm	108	108 (100)	0.2 (0.1, 0.3)	64	62 (96.9)	0.3 (0.1, 0.7)	1.61	1.18,	2.22	0.0029*
Interaction p-value										0.3115
Age group										
<65 years	185	185 (100)	0.2 (0.1, 0.3)	98	94 (95.9)	0.6 (0.2, 0.7)	1.60	1.25,	2.07	0.0002*
>=65 years	70	70 (100)	0.3 (0.2, 0.4)	33	33 (100)	0.3 (0.1, 0.5)	0.95	0.63,	1.46	0.8099
Interaction p-value										0.0380*

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaa 25NOV2020:11:06 kvbv306

Olaparib PAOLA1, German benefit assessment

Table 3.4.1 PAOLA1: Summary of subgroup analysis of AE Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + h			Placebo + b (N=1					
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]	n	Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	182 (100)	0.2 (0.2, 0.3)	89	87 (97.8)	0.3 (0.2, 0.7)	1.38	1.07,	1.79	0.0131*
IV	73	73 (100)	0.2 (0.1, 0.3)	42	40 (95.2)	0.4 (0.1, 0.7)	1.49	1.01,	2.21	0.0416*
Interaction p-value										0.7507
Region										
Europe	245	245 (100)	0.2 (0.2, 0.3)	125	121 (96.8)	0.3 (0.2, 0.7)	1.43	1.15,	1.79	0.0012*
Japan	10	10 (100)	0.5 (0.2, 0.7)	6	6 (100)	0.7 (0.3, 1.4)	1.05	0.39,	3.09	0.9237
Interaction p-value										0.5636
ECOG performance status at	Basel	ine								
(0) Normal activity	190	190 (100)	0.2 (0.2, 0.3)	100	96 (96.0)	0.3 (0.2, 0.7)	1.42	1.11,	1.83	0.0050*
(1) Restricted activity	61	61 (100)	0.2 (0.1, 0.3)	30	30 (100)	0.6 (0.1, 0.7)	1.32	0.86,	2.08	0.2042
Interaction p-value										0.7846
Baseline CA-125 value										
<=ULN	228	228 (100)	0.2 (0.2, 0.3)	117		0.3 (0.2, 0.6)	1.38	1.10,		0.0057*
>ULN	27	27 (100)	0.2 (0.0, 0.4)	14	14 (100)	0.7 (0.0, 2.4)	1.71	0.91,	3.36	0.0959
Interaction p-value										0.5302
Histological grade										
High grade	255	255 (100)	0.2 (0.2, 0.3)	131	127 (96.9)	0.3 (0.2, 0.7)	1.41	1.14,	1.76	0.0015*
Interaction p-value										NC
Cytoreductive surgery outco	me									
No residue	166	166 (100)	0.2 (0.1, 0.3)	80	78 (97.5)	0.5 (0.2, 0.7)	1.53	1.17,	2.02	0.0017*
Residue	79	79 (100)	0.2 (0.2, 0.3)	43	41 (95.3)	0.2 (0.1, 0.5)	1.22	0.84,	1.80	0.3032
Interaction p-value										0.3412

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaa 25NOV2020:11:06 kvbv306

Table 3.4.1 PAOLA1: Summary of subgroup analysis of AE Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2			Placebo + b (N=1					
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	urgery									
Upfront	146	146 (100)	0.2 (0.1, 0.3)	78	76 (97.4)	0.5 (0.2, 0.7)	1.44	1.09,	1.91	0.0096*
Interval	99	99 (100)	0.2 (0.2, 0.3)	45	43 (95.6)	0.2 (0.1, 0.3)	1.39	0.97,	2.02	0.0698
Interaction p-value										0.8800
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	158 (100)	0.2 (0.1, 0.3)	77	75 (97.4)	0.3 (0.1, 0.6)	1.28	0.98,	1.70	0.0732
Non-tBRCAm	97	97 (100)	0.2 (0.2, 0.3)	54	52 (96.3)	0.6 (0.2, 0.7)	1.65	1.18,	2.33	0.0035*
Interaction p-value										0.2614
Status somatic BRCA mutat	ions									
sBRCAm	22	22 (100)	0.2 (0.1, 0.7)	7	7 (100)	0.3 (0.1, 2.8)	1.32	0.59,	3.37	0.5142
gBRCAm	66	66 (100)	0.2 (0.1, 0.3)	31	31 (100)	0.5 (0.1, 0.7)	1.39	0.91,	2.17	0.1301
Non-BRCAm	41	41 (100)	0.2 (0.1, 0.5)	22	22 (100)	0.3 (0.1, 0.7)	1.28	0.77,	2.19	0.3535
Interaction p-value										0.9701

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaa 25NOV2020:11:06 kvbv306

Table 3.4.2 PAOLA1: Summary of subgroup analysis of AE SOC: General disorders and administration site conditions Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		-	bevacizumab 255)			bevacizumab 131)					
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)									
NED [PDS]	92	55 (59.8)	6.3 (2.8,15.0)	48	21 (43.8)	23.5 (9.8,	NE)	1.61	0.99,	2.73	0.0540
NED/CR [IDS]	74	53 (71.6)	2.4 (0.9, 6.9)	38	18 (47.4)	19.1 (4.4,	NE)	1.98	1.18,	3.47	0.0087*
NED/CR [Chemo]	40	23 (57.5)	2.8 (0.9, NE)	20	10 (50.0)	NE (NE,	NE)	1.30	0.63,	2.86	0.4855
PR	49	25 (51.0)	11.3 (2.9, NE)	25	8 (32.0)	NE (NE,	NE)	1.88	0.89,	4.45	0.1030
Interaction p-value											0.8238
Screening laboratory tBR0	CA statu	s (IVRS)									
tBRCAm	150	92 (61.3)	5.4 (2.5,11.7)	65	28 (43.1)	23.5 (10.1,	NE)	1.71	1.14,	2.66	0.0093*
non-tBRCAm	105	64 (61.0)	6.0 (2.2,13.0)	66	29 (43.9)	NE (NE,	NE)	1.66	1.08,	2.62	0.0193*
Interaction p-value											0.9275
First line treatment out	come (eC	RF)									
NED [PDS]	89	53 (59.6)	6.5 (2.8,15.9)	47	21 (44.7)	23.5 (6.9,	NE)	1.56	0.96,	2.65	0.0750
NED/CR [IDS]	74	53 (71.6)	2.2 (0.9, 6.6)	32	12 (37.5)	NE (NE,	NE)	2.89	1.60,	5.67	0.0003*
NED/CR [Chemo]	39	21 (53.8)	11.7 (0.5, NE)	17	8 (47.1)	NE (NE,	NE)	1.25	0.58,	3.01	0.5825
PR	50	27 (54.0)	11.1 (2.7, NE)	34	15 (44.1)	NE (NE,	NE)	1.22	0.66,	2.36	0.5259
Interaction p-value											0.1983
Screening laboratory tBR0	CA statu	s (eCRF)									
tBRCAm	147	90 (61.2)	5.4 (2.5,11.7)	67	29 (43.3)	23.5 (10.1,	NE)	1.70	1.14,	2.64	0.0093*
non-tBRCAm	108	66 (61.1)	6.0 (2.2,13.0)	64	28 (43.8)	NE (NE,	NE)	1.68	1.09,	2.65	0.0182*
Interaction p-value											0.9567
Age group											
<65 years	185	114 (61.6)	5.4 (2.4,11.0)	98	39 (39.8)	NE (NE,	NE)	1.91	1.34,	2.79	0.0002*
>=65 years	70	42 (60.0)	6.9 (2.8,17.8)	33	18 (54.5)	19.1 (4.6,	NE)	1.23	0.72,	2.20	0.4504
Interaction p-value											0.1982

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aab 25NOV2020:11:06 kvbv306

Table 3.4.2 PAOLA1: Summary of subgroup analysis of AE SOC: General disorders and administration site conditions Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		-	bevacizumab =255)			bevacizumab 131)				
Subgroup	n	Number (%) of patient with event	s (95% CI)		Number (%) of patients with events	Median tir (95% CI) (months) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	110 (60.4) 6.0 (2.8,11.7)	89	41 (46.1)	23.5 (9.8,	NE)	1.55	1.09, 2.24	0.0142*
IV	73	46 (63.0) 4.2 (1.4,13.4)	42	16 (38.1)	NE (NE,	NE)	2.10	1.22, 3.83	0.0071*
Interaction p-value										0.3651
Region										
Europe	245	155 (63.3) 4.8 (2.7, 7.7)	125	56 (44.8)	23.5 (10.1,	NE)	1.71	1.27, 2.34	0.0004*
Japan	10	1 (10.0) NE (NE, NE)	6	1 (16.7)	NE (NE,	NE)	0.67	0.03, 16.77	0.7728
Interaction p-value										0.5140
ECOG performance status at	Basel	ine								
(0) Normal activity	190	117 (61.6) 5.4 (2.5, 9.8)	100	43 (43.0)	NE (NE,	NE)	1.76	1.25, 2.53	0.0010*
(1) Restricted activity	61	35 (57.4) 11.7 (2.5, NE)	30	13 (43.3)	NE (NE,	NE)	1.48	0.80, 2.90	0.2166
Interaction p-value										0.6372
Baseline CA-125 value										
<=ULN	228	136 (59.6) 6.6 (3.4,11.7)	117	51 (43.6)	NE (NE,	NE)	1.62	1.18, 2.26	0.0023*
>ULN	27	20 (74.1) 1.4 (0.5, 6.9)	14	6 (42.9)	12.6 (0.8,	NE)	2.45	1.04, 6.71	0.0392*
Interaction p-value										0.3910
Histological grade										
High grade	255	156 (61.2) 5.6 (2.8,11.0)	131	57 (43.5)	NE (NE,	NE)	1.69	1.26, 2.31	0.0004*
Interaction p-value										NC
Cytoreductive surgery outco	ome									
No residue	166	108 (65.1) 4.3 (2.2, 7.1)	80	34 (42.5)	NE (NE,	NE)	2.00	1.38, 2.99	0.0002*
Residue	79	42 (53.2) 11.7 (2.7, NE)	43	19 (44.2)	NE (NE,	NE)	1.22	0.72, 2.14	0.4743
Interaction p-value										0.1469

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aab 25NOV2020:11:06 kvbv306

Table 3.4.2 PAOLA1: Summary of subgroup analysis of AE SOC: General disorders and administration site conditions Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2	bevacizumab 255)			oevacizumab 131)					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% C	I [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	83 (56.8)	7.7 (2.9,15.9)	78	32 (41.0)	NE (NE,	NE)	1.62	1.09,	2.46	0.0175*
Interval	99	67 (67.7)	2.7 (1.2, 6.6)	45	21 (46.7)	19.1 (4.6,	NE)	1.82	1.13,	3.04	0.0125*
Interaction p-value											0.7187
Myriad tumour BRCA mutati	ion stat	us									
tBRCAm	158	98 (62.0)	5.4 (2.5, 9.8)	77	32 (41.6)	NE (NE,	NE)	1.80	1.22,	2.73	0.0025*
Non-tBRCAm	97	58 (59.8)	6.0 (2.2,15.0)	54	25 (46.3)	NE (NE,	NE)	1.55	0.98,	2.52	0.0614
Interaction p-value											0.6282
Status somatic BRCA mutat	cions										
sBRCAm	22	12 (54.5)	12.5 (2.5, NE)	7	1 (14.3)	NE (NE,	NE)	4.78	0.94,	87.10	0.0610
gBRCAm	66	49 (74.2)	1.8 (0.7, 4.8)	31	14 (45.2)	NE (NE,	NE)	2.52	1.43,	4.75	0.0011*
Non-BRCAm	41	24 (58.5)	3.5 (1.4, NE)	22	14 (63.6)	4.9 (1.4,	NE)	0.94	0.49,	1.87	0.8552
Interaction p-value											0.0522

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aab 25NOV2020:11:06 kvbv306

Table 3.4.3 PAOLA1: Summary of subgroup analysis of AE PT: Fatigue Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2			Placebo + be (N=13		b				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)									
NED [PDS]	92	49 (53.3)	7.2 (2.8, NE)	48	17 (35.4)	NE (N	E, NE)	1.82	1.07,	3.26	0.0260*
NED/CR [IDS]	74	49 (66.2)	3.5 (1.8,11.0)	38	15 (39.5)	NE (N	E, NE)	2.09	1.20,	3.87	0.0080*
NED/CR [Chemo]	40	19 (47.5)	NE (NE, NE)	20	5 (25.0)	NE (N	E, NE)	2.27	0.91,	6.86	0.0796
PR	49	24 (49.0)	11.3 (3.5, NE)	25	7 (28.0)	NE (N	E, NE)	2.00	0.91,	5.03	0.0873
Interaction p-value											0.9781
Screening laboratory tBR0	CA statu	s (IVRS)									
tBRCAm	150	82 (54.7)	7.2 (3.5,18.0)	65	23 (35.4)	NE (N	E, NE)	1.80	1.15,	2.92	0.0088*
non-tBRCAm	105	59 (56.2)	11.0 (2.7,22.1)	66	21 (31.8)	NE (N	E, NE)	2.19	1.36,	3.69	0.0011*
Interaction p-value											0.5679
First line treatment out	come (eC	RF)									
NED [PDS]	89	48 (53.9)	7.2 (2.8, NE)	47	18 (38.3)	NE (N	E, NE)	1.70	1.01,	3.01	0.0458*
NED/CR [IDS]	74	48 (64.9)	3.5 (1.4, 9.8)	32	9 (28.1)	NE (N	E, NE)	3.36	1.73,	7.33	0.0002*
NED/CR [Chemo]	39	18 (46.2)	NE (NE, NE)	17	4 (23.5)	NE (N	E, NE)	2.29	0.85,	7.93	0.1045
PR	50	26 (52.0)	14.9 (3.5, NE)	34	13 (38.2)	NE (N	E, NE)	1.34	0.70,	2.69	0.3833
Interaction p-value											0.2620
Screening laboratory tBR0	CA statu	s (eCRF)									
tBRCAm	147	80 (54.4)	7.2 (3.5, NE)	67	23 (34.3)	NE (N	E, NE)	1.86	1.19,	3.03	0.0057*
non-tBRCAm	108	61 (56.5)	8.5 (2.7,22.1)	64	21 (32.8)	NE (N	E, NE)	2.13	1.32,	3.57	0.0017*
Interaction p-value											0.7009
Age group											
<65 years	185	104 (56.2)	7.2 (2.8,15.3)	98	30 (30.6)	NE (N	E, NE)	2.26	1.53,	3.46	<0.0001*
>=65 years	70	37 (52.9)	11.3 (2.9, NE)	33	14 (42.4)	NE (N	E, NE)	1.40	0.78,	2.69	0.2688
Interaction p-value											0.2108

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aac 25NOV2020:11:06 kvbv306

Table 3.4.3 PAOLA1: Summary of subgroup analysis of AE PT: Fatigue Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olapar	rib + 1 (N=2	bevacizumab 255)		Placebo + be (N=1)		mab					
Subgroup	n	Number of pat with e	ients	Median time (95% CI) (months) [a]		Number (%) of patients with events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	99 (54.4)	8.5 (3.5,18.0)	89	31 (34.8)	NE (NE,	NE)	1.86	1.26,	2.83	0.0015*
IV	73	42 (57.5)	11.0 (2.1, NE)	42	13 (31.0)	NE (NE,	NE)	2.29	1.27,	4.45	0.0053*
Interaction p-value													0.5794
Region													
Europe	245	141 (57.6)	6.3 (3.5,12.5)	125	44 (35.2)	NE (NE,	NE)	1.98	1.42,	2.81	<0.0001*
Japan	10	0		NE (NE, NE)	6	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value													NC
ECOG performance status at	Basel	ine											
(0) Normal activity	190	105 (55.3)	7.2 (3.5,13.4)	100	32 (32.0)	NE (NE,	NE)	2.11	1.43,	3.18	<0.0001*
(1) Restricted activity	61	32 (52.5)	15.3 (2.7, NE)	30	11 (36.7)	NE (NE,	NE)	1.63	0.85,	3.39	0.1471
Interaction p-value													0.5309
Baseline CA-125 value													
<=ULN	228			11.0 (4.1,15.8)	117	39 (33.3)	NE (NE,	NE)	1.97	1.39,		<0.0001*
>ULN	27	16 (59.3)	2.1 (0.5, NE)	14	5 (35.7)	NE (NE,	NE)	2.07	0.81,	6.32	0.1338
Interaction p-value													0.9320
Histological grade													
High grade	255	141 (55.3)	8.5 (3.5,15.3)	131	44 (33.6)	NE (NE,	NE)	1.98	1.43,	2.81	<0.0001*
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	97 (58.4)	5.6 (2.5,12.5)	80	27 (33.8)	NE (NE,	NE)	2.25	1.49,	3.52	<0.0001*
Residue	79	39 (49.4)	15.8 (3.5, NE)	43	13 (30.2)	NE (NE,	NE)	1.69	0.92,	3.28	0.0902
Interaction p-value													0.4570

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aac 25NOV2020:11:06 kvbv306

Table 3.4.3 PAOLA1: Summary of subgroup analysis of AE PT: Fatigue Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2			Placebo + b (N=1						
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	75 (51.4)	13.0 (4.9, NE)	78	23 (29.5)	NE (NE,	NE)	2.07	1.32,	3.38	0.0012*
Interval	99	61 (61.6)	3.7 (2.1,11.8)	45	17 (37.8)	NE (NE,	NE)	1.98	1.19,	3.50	0.0082*
Interaction p-value											0.9015
Myriad tumour BRCA mutati	ion stat	us									
tBRCAm	158	87 (55.1)	7.2 (3.5,18.0)	77	26 (33.8)	NE (NE,	NE)	1.90	1.24,	3.00	0.0025*
Non-tBRCAm	97	54 (55.7)	11.0 (2.7, NE)	54	18 (33.3)	NE (NE,	NE)	2.10	1.26,	3.69	0.0040*
Interaction p-value											0.7720
Status somatic BRCA mutat	cions										
sBRCAm	22	10 (45.5)	NE (NE, NE)	7	0	NE (NE,	NE)	NC	NC		NC
gBRCAm	66	45 (68.2)	2.4 (1.1, 6.9)	31	13 (41.9)	NE (NE,	NE)	2.29	1.27,	4.43	0.0050*
Non-BRCAm	41	24 (58.5)	4.1 (1.4, NE)	22	12 (54.5)	8.3 (2.8,	NE)	1.24	0.63,	2.56	0.5439
Interaction p-value											0.1942

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aac 25NOV2020:11:06 kvbv306

Table 3.4.4 PAOLA1: Summary of subgroup analysis of AE SOC: Endocrine disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	. (Olaparib + b (N=2		ab		Placebo + be (N=1)		b				
Subgroup	C	Number (%) of patients n with events		time CI)) [a]	(Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IVR	S)										
NED [PDS]	92	1 (1.1)	NE (N	E, NE)	48	5 (10.4)	NE (IE, NE)	NC	NC		NC
NED/CR [IDS]	74	2 (2.7)	NE (N	E, NE)	38	3 (7.9)	NE (E, NE)	NC	NC		NC
NED/CR [Chemo]	40	0	NE (N	E, NE)	20	1 (5.0)	NE (E, NE)	NC	NC		NC
PR	49	1 (2.0)	NE (N	E, NE)	25	0	NE (E, NE)	NC	NC		NC
Interaction p-value												NC
Screening laboratory tBR0	CA status	(IVRS)										
tBRCAm	150	4 (2.7)	NE (N	E, NE)	65	5 (7.7)	NE (E, NE)	NC	NC		NC
non-tBRCAm	105	0	NE (N	E, NE)	66	4 (6.1)	NE (IE, NE)	NC	NC		NC
Interaction p-value												NC
First line treatment out	come (eCR	F)										
NED [PDS]	89	1 (1.1)	NE (N	E, NE)	47	5 (10.6)	NE (N	E, NE)	NC	NC		NC
NED/CR [IDS]	74	2 (2.7)	NE (N	E, NE)	32	3 (9.4)	NE (N	E, NE)	NC	NC		NC
NED/CR [Chemo]	39	0	NE (N	E, NE)	17	1 (5.9)	NE (N	E, NE)	NC	NC		NC
PR	50	1 (2.0)	NE (N	E, NE)	34	0	NE (N	E, NE)	NC	NC		NC
Interaction p-value												NC
Screening laboratory tBR0	CA status	(eCRF)										
tBRCAm	147	3 (2.0)	NE (N	E, NE)	67	5 (7.5)	NE (IE, NE)	NC	NC		NC
non-tBRCAm	108	1 (0.9)	NE (N	E, NE)	64	4 (6.3)	NE (E, NE)	NC	NC		NC
Interaction p-value												NC
Age group												
<65 years	185	3 (1.6)	NE (N	E, NE)	98	9 (9.2)	NE (IE, NE)	0.17	0.04,	0.55	0.0031*
>=65 years	70	1 (1.4)	NE (N	E, NE)	33	0	NE (IE, NE)	NC	NC		NC
Interaction p-value												NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aad 25NOV2020:11:06 kvbv306

Table 3.4.4 PAOLA1: Summary of subgroup analysis of AE SOC: Endocrine disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	I	Olaparib + b (N=2!		ımab			Placebo + be (N=13		mab					
Subgroup	Number (%) of patients n with events		Media (95 (mont	% CI)	C	Number (%) of patients with events	(95	an ti % CI) hs) [)	Hazard ratio [b]	95% C	I [b]	2-sided p-value [b]
FIGO Stage (Disease state)														
III	182	4 (2.2)	NE (NE,	NE)	89	9 (10.1)	NE (NE,	NE)	0.20	0.06,	0.63	0.0053*
IV	73	0	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
Region														
Europe	245	4 (1.6)	NE (NE,	NE)	125	9 (7.2)	NE (NE,	NE)	0.20	0.06,	0.63	0.0055*
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
ECOG performance status at	Baseli	.ne												
(0) Normal activity	190	4 (2.1)	NE (NE,	NE)	100	8 (8.0)	NE (NE,	NE)	0.25	0.07,	0.80	0.0190*
(1) Restricted activity	61	0	NE (NE,	NE)	30	1 (3.3)	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
Baseline CA-125 value														
<=ULN	228	3 (1.3)	NE (NE,	NE)	117	9 (7.7)	NE (NE,	NE)	0.16	0.03,	0.53	0.0023*
>ULN	27	1 (3.7)	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
Histological grade														
High grade	255	4 (1.6)	NE (NE,	NE)	131	9 (6.9)	NE (NE,	NE)	0.21	0.06,	0.65	0.0063*
Interaction p-value														NC
Cytoreductive surgery outco	ome													
No residue	166	3 (1.8)	NE (NE,	NE)	80	8 (10.0)	NE (NE,	NE)	0.17	0.04,	0.59	0.0047*
Residue	79	1 (1.3)	NE (NE,	NE)	43	1 (2.3)	NE (NE,	NE)	0.48	0.02,	12.22	0.6108
Interaction p-value														0.5092

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aad 25NOV2020:11:06 kvbv306

Table 3.4.4 PAOLA1: Summary of subgroup analysis of AE SOC: Endocrine disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	1	Olaparib + b (N=2		•		Placebo + be (N=1)		b			
Subgroup	C	Number (%) of patients with events	Median t (95% CI (months)	.)	C	Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	1 (0.7)	NE (NE,	NE)	78	6 (7.7)	NE (1	IE, NE)	NC	NC	NC
Interval	99	3 (3.0)	NE (NE,	NE)	45	3 (6.7)	NE (IE, NE)	NC	NC	NC
Interaction p-value											NC
Myriad tumour BRCA mutati	on statu	ıs									
tBRCAm	158	4 (2.5)	NE (NE,	NE)	77	6 (7.8)	NE (1	IE, NE)	0.28	0.07, 0.998	0.0496*
Non-tBRCAm	97	0	NE (NE,	NE)	54	3 (5.6)	NE (1	IE, NE)	NC	NC	NC
Interaction p-value											NC
Status somatic BRCA mutat	ions										
sBRCAm	22	0	NE (NE,	NE)	7	2 (28.6)	NE (IE, NE)	NC	NC	NC
gBRCAm	66	2 (3.0)	NE (NE,	NE)	31	3 (9.7)	NE (IE, NE)	NC	NC	NC
Non-BRCAm	41	0	NE (NE,	NE)	22	1 (4.5)	NE (1	IE, NE)	NC	NC	NC
Interaction p-value											NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aad 25NOV2020:11:06 kvbv306

Table 3.4.5 PAOLA1: Summary of subgroup analysis of AE PT: Dyspnoea Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olaparib + k (N=2		umab	:		Placebo + be (N=1)		ımab				
Subgroup		Number (%) of patients with events		an ti 5% CI :hs))	C	Number (%) of patients with events	(95	an ti 5% CI :hs))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IV	RS)											
NED [PDS]	92	15 (16.3)	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	8.42	1.71,152.21	0.0049*
NED/CR [IDS]	74	4 (5.4)	NE (NE,	NE)	38	1 (2.6)	NE (NE,	NE)	2.13	0.31, 41.60	0.4692
NED/CR [Chemo]	40	1 (2.5)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	2 (4.1)	NE (NE,	NE)	25	1 (4.0)	NE (NE,	NE)	0.98	0.09, 20.97	0.9836
Interaction p-value													0.3674
Screening laboratory tBR0	CA statu	s (IVRS)											
tBRCAm	150	11 (7.3)	NE (NE,	NE)	65	2 (3.1)	NE (NE,	NE)	2.41	0.65, 15.55	0.2087
non-tBRCAm	105	11 (10.5)	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	7.43	1.45,135.83	0.0122*
Interaction p-value													0.3692
First line treatment out	come (eCl	RF)											
NED [PDS]	89	13 (14.6)	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	7.33	1.46,133.04	0.0111*
NED/CR [IDS]	74	5 (6.8)	NE (NE,	NE)	32	1 (3.1)	NE (NE,	NE)	2.35	0.38, 44.94	0.3950
NED/CR [Chemo]	39	2 (5.1)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	1 (2.0)	NE (NE,	NE)	34	1 (2.9)	NE (NE,	NE)	0.63	0.02, 15.90	0.7445
Interaction p-value													0.3554
Screening laboratory tBR0	CA statu	s (eCRF)											
tBRCAm	147	10 (6.8)	NE (NE,	NE)	67	2 (3.0)	NE (NE,	NE)	2.30	0.61, 14.97	0.2407
non-tBRCAm	108	12 (11.1)	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	7.62	1.50,138.72	0.0100*
Interaction p-value													0.3402
Age group													
<65 years	185	15 (8.1)	NE (NE,	NE)	98	2 (2.0)	NE (NE,	NE)	4.11	1.16, 26.06	0.0262*
>=65 years	70	7 (10.0)	NE (NE,	NE)	33	1 (3.0)	NE (NE,	NE)	3.40	0.60, 63.50	0.1863
Interaction p-value													0.8855

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aae 25NOV2020:11:06 kvbv306

Table 3.4.5 PAOLA1: Summary of subgroup analysis of AE PT: Dyspnoea Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		ıb		Placebo + be (N=1)					
Subgroup		Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	Median to (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	15 (8.2)	NE (N	E, NE)	89	2 (2.2)	NE (NE,	NE)	3.83	1.08, 24.30	0.0359*
IV	73	7 (9.6)	NE (N	E, NE)	42	1 (2.4)	NE (NE,	NE)	4.08	0.73, 76.27	0.1217
Interaction p-value											0.9617
Region											
Europe	245	22 (9.0)	NE (N	E, NE)	125	3 (2.4)	NE (NE,	NE)	3.85	1.33, 16.26	0.0101*
Japan	10	0	NE (N	E, NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	15 (7.9)	NE (N	E, NE)	100	2 (2.0)	NE (NE,	NE)	4.14	1.17, 26.30	0.0251*
(1) Restricted activity	61	6 (9.8)	NE (N	E, NE)	30	1 (3.3)	NE (NE,	NE)	2.86	0.49, 54.11	0.2734
Interaction p-value											0.7826
Baseline CA-125 value											
<=ULN	228	18 (7.9)	NE (N	E, NE)	117	3 (2.6)	NE (NE,	NE)	3.15	1.07, 13.47	0.0365*
>ULN	27	4 (14.8)	NE (N	E, NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	22 (8.6)	NE (N	E, NE)	131	3 (2.3)	NE (NE,	NE)	3.89	1.35, 16.46	0.0094*
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	19 (11.4)	NE (N	E, NE)	80	2 (2.5)	NE (NE,	NE)	4.95	1.44, 31.08	0.0081*
Residue	79	2 (2.5)	NE (N	E, NE)	43	1 (2.3)	NE (NE,	NE)	1.02	0.10, 21.90	0.9880
Interaction p-value											0.2972

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aae 25NOV2020:11:06 kvbv306

Table 3.4.5 PAOLA1: Summary of subgroup analysis of AE PT: Dyspnoea Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		Place	ebo + bo (N=1	evacizumab 31)	•			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Numbe of pat n with e	ients	Median tim (95% CI) (months) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	15 (10.3)	NE (NE, NE)	78 2 (2.6)	NE (NE,	NE)	4.15	1.17, 26.32	0.0250*
Interval	99	6 (6.1)	NE (NE, NE)	45 1 (2.2)	NE (NE,	NE)	2.78	0.47, 52.52	0.2884
Interaction p-value										0.7654
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	11 (7.0)	NE (NE, NE)	77 2 (2.6)	NE (NE,	NE)	2.70	0.72, 17.42	0.1512
Non-tBRCAm	97	11 (11.3)	NE (NE, NE)	54 1 (1.9)	NE (NE,	NE)	6.63	1.29,121.23	0.0195*
Interaction p-value										0.4749
Status somatic BRCA mutat	ions									
sBRCAm	22	1 (4.5)	NE (NE, NE)	7 0		NE (NE,	NE)	NC	NC	NC
gBRCAm	66	4 (6.1)	NE (NE, NE)	31 1 (3.2)	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	41	3 (7.3)	NE (NE, NE)	22 0		NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aae 25NOV2020:11:06 kvbv306

Table 3.4.6 PAOLA1: Summary of subgroup analysis of AE SOC: Renal and urinary disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olaparib + b (N=2		umab			Placebo + b		ımab					
Subgroup		Number (%) of patients with events	(95	an ti 5% CI :hs))		Number (%) of patients with events	(9!	an ti 5% CI ths))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment outo	come (IV	RS)												
NED [PDS]	92	10 (10.9)	NE (NE,	NE)	48	11 (22.9)	NE (NE,	NE)	0.43	0.18,	1.01	0.0525
NED/CR [IDS]	74	8 (10.8)	NE (NE,	NE)	38	4 (10.5)	NE (NE,	NE)	1.04	0.33,	3.90	0.9486
NED/CR [Chemo]	40	4 (10.0)	NE (NE,	NE)	20	3 (15.0)	NE (NE,	NE)	0.62	0.14,	3.13	0.5322
PR	49	5 (10.2)	NE (NE,	NE)	25	6 (24.0)	NE (NE,	NE)	0.38	0.11,	1.26	0.1102
Interaction p-value														0.5961
Screening laboratory tBRO	CA statu	s (IVRS)												
tBRCAm	150	21 (14.0)	NE (NE,	NE)	65	11 (16.9)	NE (NE,	NE)	0.78	0.39,	1.69	0.5197
non-tBRCAm	105	6 (5.7)	NE (NE,	NE)	66	13 (19.7)	NE (NE,	NE)	0.27	0.09,	0.68	0.0050*
Interaction p-value														0.0730
First line treatment outo	come (eC	RF)												
NED [PDS]	89	7 (7.9)	NE (NE,	NE)	47	10 (21.3)	NE (NE,	NE)	0.34	0.12,	0.87	0.0255*
NED/CR [IDS]	74	10 (13.5)	NE (NE,	NE)	32	3 (9.4)	NE (NE,	NE)	1.53	0.47,	6.82	0.5041
NED/CR [Chemo]	39	3 (7.7)	NE (NE,	NE)	17	3 (17.6)	NE (NE,	NE)	0.39	0.07,	2.12	0.2590
PR	50	6 (12.0)	NE (NE,	NE)	34	7 (20.6)	NE (NE,	NE)	0.51	0.16,	1.53	0.2224
Interaction p-value														0.2583
Screening laboratory tBRO	CA statu	s (eCRF)												
tBRCAm	147	20 (13.6)	NE (NE,	NE)	67	11 (16.4)	NE (NE,	NE)	0.78	0.38,	1.69	0.5149
non-tBRCAm	108	7 (6.5)	NE (NE,	NE)	64	13 (20.3)	NE (NE,	NE)	0.30	0.11,	0.72	0.0073*
Interaction p-value														0.0987
Age group														
<65 years	185	20 (10.8)	NE (NE,	NE)	98	20 (20.4)	NE (NE,	NE)	0.50	0.27,	0.93	0.0282*
>=65 years	70	7 (10.0)	NE (NE,	NE)	33	4 (12.1)	NE (NE,	NE)	0.76	0.23,	2.89	0.6618
Interaction p-value														0.5424

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaf 25NOV2020:11:06 kvbv306

Table 3.4.6 PAOLA1: Summary of subgroup analysis of AE SOC: Renal and urinary disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		umab			Placebo + be (N=1)		nab	 :				
Subgroup		Number (%) of patients with events	(95	an ti 5% CI :hs))		Number (%) of patients with events	(95	n time % CI) hs) [a		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)														
III	182	19 (10.4)	NE (NE,	NE)	89	15 (16.9)	NE (NE,	NE)	0.58	0.30,	1.17	0.1264
IV	73	8 (11.0)	NE (NE,	NE)	42	9 (21.4)	NE (NE,	NE)	0.47	0.17,	1.22	0.1184
Interaction p-value														0.7042
Region														
Europe	245	27 (11.0)	NE (NE,	NE)	125	21 (16.8)	NE (NE,	NE)	0.61	0.35,	1.10	0.0977
Japan	10	0	NE (NE,	NE)	6	3 (50.0)	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
ECOG performance status at	Basel	ine												
(0) Normal activity	190	21 (11.1)	NE (NE,	NE)	100	17 (17.0)	NE (NE,	NE)	0.63	0.33,	1.21	0.1633
(1) Restricted activity	61	6 (9.8)	NE (NE,	NE)	30	7 (23.3)	NE (NE,	NE)	0.35	0.11,	1.04	0.0594
Interaction p-value														0.3520
Baseline CA-125 value														
<=ULN	228	23 (10.1)	NE (NE,	NE)	117	22 (18.8)	NE (NE,	NE)	0.50	0.28,		0.0229*
>ULN	27	4 (14.8)	NE (NE,	NE)	14	2 (14.3)	NE (NE,	NE)	0.91	0.18,	6.57	0.9149
Interaction p-value														0.5078
Histological grade														
High grade	255	27 (10.6)	NE (NE,	NE)	131	24 (18.3)	NE (NE,	NE)	0.54	0.31,	0.94	0.0299*
Interaction p-value														NC
Cytoreductive surgery outco	ome													
No residue	166	18 (10.8)	NE (NE,	NE)	80	14 (17.5)	NE (NE,	NE)	0.60	0.30,	1.23	0.1565
Residue	79	8 (10.1)	NE (NE,	NE)	43	9 (20.9)	NE (NE,	NE)	0.42	0.16,	1.09	0.0730
Interaction p-value														0.5440

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaf 25NOV2020:11:06 kvbv306

Table 3.4.6 PAOLA1: Summary of subgroup analysis of AE SOC: Renal and urinary disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2				Placebo + be (N=1)		mab	·			
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events	(95	an time % CI) hs) [a]	Hazard ratio [b]	95% C	I [b]	2-sided p-value [b]
Timing of cytoreductive :	surgery											
Upfront	146	12 (8.2)	NE (NE,	NE)	78	17 (21.8)	NE (NE, NE)	0.34	0.16,	0.70	0.0038*
Interval	99	14 (14.1)	NE (NE,	NE)	45	6 (13.3)	NE (NE, NE)	1.04	0.42,	2.94	0.9350
Interaction p-value												0.0611
Myriad tumour BRCA mutat:	ion stat	us										
tBRCAm	158	21 (13.3)	NE (NE,	NE)	77	12 (15.6)	NE (NE, NE)	0.81	0.40,	1.69	0.5574
Non-tBRCAm	97	6 (6.2)	NE (NE,	NE)	54	12 (22.2)	NE (NE, NE)	0.25	0.09,	0.66	0.0044*
Interaction p-value												0.0550
Status somatic BRCA muta	cions											
sBRCAm	22	2 (9.1)	NE (NE,	NE)	7	1 (14.3)	NE (NE, NE)	0.52	0.05,	11.27	0.6130
gBRCAm	66	9 (13.6)	NE (NE,	NE)	31	5 (16.1)	NE (NE, NE)	0.89	0.31,	2.89	0.8311
Non-BRCAm	41	3 (7.3)	NE (NE,	NE)	22	3 (13.6)	NE (NE, NE)	0.49	0.09,	2.66	0.3900
Interaction p-value												0.8095

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaf 25NOV2020:11:06 kvbv306

Table 3.4.7 PAOLA1: Summary of subgroup analysis of AE PT: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	;	Olaparib + b (N=2		ımab	:		Placebo + be (N=1)		nab	•				
Subgroup		Number (%) of patients with events	Media (95 (mont)	% CI)			Number (%) of patients with events	(95	an time % CI) hs) [a		Hazard ratio [b]	95% C	I [b]	2-sided p-value [b]
First line treatment out	come (IVI	RS)												
NED [PDS]	92	6 (6.5)	NE (NE,	NE)	48	9 (18.8)	NE (NE,	NE)	0.31	0.10,	0.85	0.0239*
NED/CR [IDS]	74	7 (9.5)	NE (NE,	NE)	38	3 (7.9)	NE (NE,	NE)	1.22	0.34,	5.68	0.7674
NED/CR [Chemo]	40	3 (7.5)	NE (NE,	NE)	20	3 (15.0)	NE (NE,	NE)	0.47	0.09,	2.51	0.3543
PR	49	3 (6.1)	NE (NE,	NE)	25	4 (16.0)	NE (NE,	NE)	0.36	0.07,	1.64	0.1818
Interaction p-value														0.4102
Screening laboratory tBR0	CA status	s (IVRS)												
tBRCAm	150	15 (10.0)	NE (NE,	NE)	65	9 (13.8)	NE (NE,	NE)	0.67	0.30,	1.60	0.3569
non-tBRCAm	105	4 (3.8)	NE (NE,	NE)	66	10 (15.2)	NE (NE,	NE)	0.24	0.07,	0.72	0.0103*
Interaction p-value														0.1452
First line treatment out	come (eCI	RF)												
NED [PDS]	89	5 (5.6)	NE (NE,	NE)	47	8 (17.0)	NE (NE,	NE)	0.30	0.09,	0.90	0.0320*
NED/CR [IDS]	74	9 (12.2)	NE (NE,	NE)	32	2 (6.3)	NE (NE,	NE)	2.12	0.55,	13.93	0.2993
NED/CR [Chemo]	39	1 (2.6)	NE (NE,	NE)	17	2 (11.8)	NE (NE,	NE)	0.20	0.01,	2.09	0.1724
PR	50	3 (6.0)	NE (NE,	NE)	34	6 (17.6)	NE (NE,	NE)	0.29	0.06,	1.10	0.0695
Interaction p-value														0.0915
Screening laboratory tBR0	CA status	s (eCRF)												
tBRCAm	147	14 (9.5)	NE (NE,	NE)	67	9 (13.4)	NE (NE,	NE)	0.66	0.29,	1.58	0.3349
non-tBRCAm	108	5 (4.6)	NE (NE,	NE)	64	10 (15.6)	NE (NE,	NE)	0.29	0.09,	0.80	0.0173*
Interaction p-value														0.2213
Age group														
<65 years	185	14 (7.6)	NE (NE,	NE)	98	17 (17.3)	NE (NE,	NE)	0.41	0.20,	0.83	0.0135*
>=65 years	70	5 (7.1)	NE (NE,	NE)	33	2 (6.1)	NE (NE,	NE)	1.12	0.24,	7.80	0.8939
Interaction p-value														0.2508

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aag 25NOV2020:11:06 kvbv306

Table 3.4.7 PAOLA1: Summary of subgroup analysis of AE PT: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b)		Placebo + be (N=1)						
Subgroup		Number (%) of patients with events	Median (95% C (months)	I)		Number (%) of patients with events	Median t (95% C (months)	[)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	14 (7.7)	NE (NE	, NE)	89	12 (13.5)	NE (NE	NE)	0.54	0.25,	1.18	0.1187
IV	73	5 (6.8)	NE (NE	, NE)	42	7 (16.7)	NE (NE	NE)	0.39	0.12,	1.22	0.1053
Interaction p-value												0.6513
Region												
Europe	245	19 (7.8)	NE (NE	, NE)	125	16 (12.8)	NE (NE	NE)	0.57	0.29,	1.13	0.1043
Japan	10	0	NE (NE	, NE)	6	3 (50.0)	NE (NE	NE)	NC	NC		NC
Interaction p-value												NC
ECOG performance status at	Basel	ine										
(0) Normal activity	190	15 (7.9)	NE (NE	, NE)	100	13 (13.0)	NE (NE	NE)	0.61	0.29,	1.29	0.1900
(1) Restricted activity	61	4 (6.6)	NE (NE	, NE)	30	6 (20.0)	NE (NE	NE)	0.26	0.07,	0.90	0.0335*
Interaction p-value												0.2452
Baseline CA-125 value												
<=ULN	228	17 (7.5)	NE (NE	, NE)	117	17 (14.5)	NE (NE	NE)	0.49	0.25,	0.96	0.0388*
>ULN	27	2 (7.4)	NE (NE	, NE)	14	2 (14.3)	NE (NE	NE)	0.43	0.05,	3.61	0.4093
Interaction p-value												0.9090
Histological grade												
High grade	255	19 (7.5)	NE (NE	, NE)	131	19 (14.5)	NE (NE	NE)	0.48	0.25,	0.92	0.0267*
Interaction p-value												NC
Cytoreductive surgery outc	ome											
No residue	166	15 (9.0)	NE (NE	, NE)	80	11 (13.8)	NE (NE	NE)	0.65	0.30,	1.44	0.2772
Residue	79	4 (5.1)	NE (NE	, NE)	43	7 (16.3)	NE (NE	NE)	0.27	0.07,	0.90	0.0328*
Interaction p-value												0.2349

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aag 25NOV2020:11:06 kvbv306

Table 3.4.7 PAOLA1: Summary of subgroup analysis of AE PT: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		Plac	ebo + b (N=1	evacizumab 31)	•		
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		er (%) atients events	Median time (95% CI) (months) [a]	- Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery								
Upfront	146	8 (5.5)	NE (NE, NE)	78 14	(17.9)	NE (NE, N	E) 0.28	0.11, 0.65	0.0030*
Interval	99	11 (11.1)	NE (NE, NE)	45 4	(8.9)	NE (NE, N	E) 1.25	0.43, 4.53	0.6931
Interaction p-value									0.0313*
Myriad tumour BRCA mutati	on stat	us							
tBRCAm	158	16 (10.1)	NE (NE, NE)	77 10	(13.0)	NE (NE, N	E) 0.72	0.33, 1.65	0.4312
Non-tBRCAm	97	3 (3.1)	NE (NE, NE)	54 9	(16.7)	NE (NE, N	E) 0.18	0.04, 0.60	0.0046*
Interaction p-value									0.0581
Status somatic BRCA mutat	ions								
sBRCAm	22	2 (9.1)	NE (NE, NE)	7 1	(14.3)	NE (NE, N	E) 0.51	0.05, 10.99	0.5998
gBRCAm	66	7 (10.6)	NE (NE, NE)	31 5	(16.1)	NE (NE, N	E) 0.66	0.21, 2.24	0.4909
Non-BRCAm	41	2 (4.9)	NE (NE, NE)	22 2	(9.1)	NE (NE, N	E) 0.51	0.06, 4.24	0.5034
Interaction p-value									0.9636

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aag 25NOV2020:11:06 kvbv306

Table 3.4.8 PAOLA1: Summary of subgroup analysis of AE SOC: Blood and lymphatic system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1			Placebo + be (N=1			:			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median (95% ((months)	I)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)									
NED [PDS]	92	53 (57.6)	4.8 (2.8,23.5)	48	15 (31.3)	NE (NE	, NE)	2.35	1.36,	4.32	0.0017*
NED/CR [IDS]	74	49 (66.2)	2.1 (1.4, 6.2)	38	13 (34.2)	NE (NE	, NE)	2.56	1.43,	4.92	0.0011*
NED/CR [Chemo]	40	20 (50.0)	12.5 (2.8, NE)	20	7 (35.0)	NE (NE	, NE)	1.64	0.72,	4.17	0.2451
PR	49	26 (53.1)	11.1 (2.1, NE)	25	6 (24.0)	NE (NE	, NE)	2.87	1.26,	7.72	0.0102*
Interaction p-value											0.8208
Screening laboratory tBRG	CA statu	s (IVRS)									
tBRCAm	150	84 (56.0)	7.2 (2.8,18.0)	65	22 (33.8)	NE (NE	, NE)	2.01	1.28,	3.30	0.0019*
non-tBRCAm	105	64 (61.0)	2.8 (2.1,12.5)	66	19 (28.8)	NE (NE	, NE)	2.85	1.74,	4.90	<0.0001*
Interaction p-value											0.3244
First line treatment out	come (eC	RF)									
NED [PDS]	89	52 (58.4)	4.8 (2.8,23.5)	47	14 (29.8)	NE (NE	, NE)	2.56	1.46,	4.80	0.0007*
NED/CR [IDS]	74	48 (64.9)	3.4 (1.9, 8.2)	32	10 (31.3)	NE (NE	, NE)	2.80	1.48,	5.88	0.0010*
NED/CR [Chemo]	39	17 (43.6)	NE (NE, NE)	17	7 (41.2)	NE (NE	, NE)	1.07	0.46,	2.76	0.8827
PR	50	29 (58.0)	2.8 (2.0, NE)	34	10 (29.4)	NE (NE	, NE)	2.56	1.29,	5.54	0.0063*
Interaction p-value											0.3726
Screening laboratory tBRG	CA statu	s (eCRF)									
tBRCAm	147	83 (56.5)	7.2 (2.8,18.0)	67	23 (34.3)	NE (NE	, NE)	2.00	1.28,	3.25	0.0018*
non-tBRCAm	108	65 (60.2)	2.8 (2.1,13.8)	64	18 (28.1)	NE (NE	, NE)	2.87	1.74,	4.99	<0.0001*
Interaction p-value											0.3089
Age group											
<65 years	185	103 (55.7)	7.2 (2.8,18.0)	98	31 (31.6)	NE (NE	, NE)	2.18	1.48,	3.32	<0.0001*
>=65 years	70	45 (64.3)	2.7 (1.4,12.3)	33	10 (30.3)	NE (NE	, NE)	2.87	1.51,	6.03	0.0009*
Interaction p-value											0.4966

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aah 25NOV2020:11:06 kvbv306

Table 3.4.8 PAOLA1: Summary of subgroup analysis of AE SOC: Blood and lymphatic system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		-	bevacizumab 255)		Placebo + b						
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median tim (95% CI) (months) [a		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	107 (58.8)	3.7 (2.6,15.3)	89	26 (29.2)	NE (NE,	NE)	2.69	1.78,	4.21	<0.0001*
IV	73	41 (56.2)	11.1 (2.8, NE)	42	15 (35.7)	NE (NE,	NE)	1.78	1.01,	3.31	0.0478*
Interaction p-value											0.2718
Region											
Europe	245	139 (56.7)	5.6 (2.8,13.8)	125	37 (29.6)	NE (NE,	NE)	2.46	1.73,	3.58	<0.0001*
Japan	10	9 (90.0)	2.1 (0.3, 2.8)	6	4 (66.7)	0.7 (0.3,	NE)	1.40	0.45,	5.16	0.5717
Interaction p-value											0.3857
ECOG performance status at	Basel	ine									
(0) Normal activity	190	109 (57.4)	4.7 (2.8,12.5)	100	29 (29.0)	NE (NE,	NE)	2.59	1.74,	3.97	<0.0001*
(1) Restricted activity	61	37 (60.7)	5.4 (2.0,21.7)	30	12 (40.0)	NE (NE,	NE)	1.71	0.92,	3.42	0.0934
Interaction p-value											0.2979
Baseline CA-125 value											
<=ULN	228	131 (57.5)	, , , ,	117	35 (29.9)	` ,	NE)	2.45	1.70,		<0.0001*
>ULN	27	17 (63.0)	2.7 (1.1, NE)	14	6 (42.9)	NE (NE,	NE)	1.82	0.76,	5.04	0.1889
Interaction p-value											0.5701
Histological grade											
High grade	255	148 (58.0)	4.8 (2.8,12.3)	131	41 (31.3)	NE (NE,	NE)	2.35	1.68,	3.37	<0.0001*
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	102 (61.4)	3.5 (2.7,10.4)	80	24 (30.0)	NE (NE,	NE)	2.74	1.78,	4.37	<0.0001*
Residue	79	41 (51.9)	11.2 (2.7, NE)	43	14 (32.6)	NE (NE,	NE)	1.82	1.02,	3.47	0.0425*
Interaction p-value											0.2967

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aah 25NOV2020:11:06 kvbv306

Table 3.4.8 PAOLA1: Summary of subgroup analysis of AE SOC: Blood and lymphatic system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2			Placebo + be (N=1)		mab					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	(95	an tir % CI) hs) [Hazard ratio [b]	95% C	I [b]	2-sided p-value [b]
Timing of cytoreductive :	surgery											
Upfront	146	77 (52.7)	12.5 (3.4, NE)	78	21 (26.9)	NE (NE,	NE)	2.45	1.54,	4.07	<0.0001*
Interval	99	66 (66.7)	2.7 (2.0, 5.6)	45	17 (37.8)	NE (NE,	NE)	2.26	1.36,	3.98	0.0013*
Interaction p-value												0.8264
Myriad tumour BRCA mutat:	ion stat	us										
tBRCAm	158	93 (58.9)	4.7 (2.7,12.3)	77	23 (29.9)	NE (NE,	NE)	2.53	1.63,	4.10	<0.0001*
Non-tBRCAm	97	55 (56.7)	5.4 (2.1,23.5)	54	18 (33.3)	NE (NE,	NE)	2.12	1.27,	3.71	0.0036*
Interaction p-value												0.6168
Status somatic BRCA muta	cions											
sBRCAm	22	14 (63.6)	2.8 (0.5, NE)	7	2 (28.6)	NE (NE,	NE)	2.95	0.82,	18.82	0.1031
gBRCAm	66	41 (62.1)	2.8 (2.1, 9.7)	31	14 (45.2)	NE (NE,	NE)	1.82	1.02,	3.47	0.0431*
Non-BRCAm	41	27 (65.9)	2.8 (1.1,18.0)	22	7 (31.8)	NE (NE,	NE)	2.55	1.17,	6.35	0.0169*
Interaction p-value												0.7298

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aah 25NOV2020:11:06 kvbv306

Table 3.4.9 PAOLA1: Summary of subgroup analysis of AE PT: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

Subgroup	·	Olaparib + (N=2	bevacizum 255)	zumab Placebo + bevacizumab (N=131)									
		Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]		
First line treatment outcome (IVRS)													
NED [PDS]	92	37 (40.2)	NE (I	NE, NE)	48	8 (16.7)	NE (1	NE, NE)	2.98	1.46, 6.90	0.0019*		
NED/CR [IDS]	74	34 (45.9)	21.7 (4	.6, NE)	38	0	NE (NE, NE)	NC	NC	NC		
NED/CR [Chemo]	40	13 (32.5)	NE (I	NE, NE)	20	1 (5.0)	NE (NE, NE)	7.15	1.42,129.87	0.0123*		
PR	49	18 (36.7)	NE (I	NE, NE)	25	3 (12.0)	NE (1	NE, NE)	3.78	1.28, 16.12	0.0139*		
Interaction p-value											0.6834		
Screening laboratory tBR0	CA statu	s (IVRS)											
tBRCAm	150	55 (36.7)	NE (I	NE, NE)	65	7 (10.8)	NE (1	NE, NE)	4.00	1.95, 9.64	<0.0001*		
non-tBRCAm	105	47 (44.8)	NE (I	NE, NE)	66	5 (7.6)	NE (1	NE, NE)	7.66	3.35, 22.08	<0.0001*		
Interaction p-value											0.2896		
First line treatment out	come (eC	RF)											
NED [PDS]	89	35 (39.3)	NE (I	NE, NE)	47	8 (17.0)	NE (1	NE, NE)	2.82	1.38, 6.54	0.0037*		
NED/CR [IDS]	74	32 (43.2)	NE (I	NE, NE)	32	0	NE (1	NE, NE)	NC	NC	NC		
NED/CR [Chemo]	39	11 (28.2)	NE (I	NE, NE)	17	1 (5.9)	NE (1	NE, NE)	5.06	0.98, 92.48	0.0528		
PR	50	22 (44.0)	NE (I	NE, NE)	34	3 (8.8)	NE (1	NE, NE)	6.03	2.09, 25.49	0.0003*		
Interaction p-value											0.5317		
Screening laboratory tBR0	CA statu	s (eCRF)											
tBRCAm	147	55 (37.4)	NE (NE, NE)	67	7 (10.4)	NE (1	NE, NE)	4.23	2.06, 10.20	<0.0001*		
non-tBRCAm	108	47 (43.5)	NE (I	NE, NE)	64	5 (7.8)	NE (1	NE, NE)	7.15	3.13, 20.62	<0.0001*		
Interaction p-value											0.3921		
Age group													
<65 years	185	70 (37.8)	NE (I	NE, NE)	98	9 (9.2)	NE (1	NE, NE)	4.88	2.57, 10.49	<0.0001*		
>=65 years	70	32 (45.7)	NE (I	NE, NE)	33	3 (9.1)	NE (1	NE, NE)	6.80	2.43, 28.30	<0.0001*		
Interaction p-value											0.6278		

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aai 25NOV2020:11:06 kvbv306

Table 3.4.9 PAOLA1: Summary of subgroup analysis of AE PT: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bevacizumab Placebo + bevacizumab (N=255) (N=131)									
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	73 (40.1)	NE (NE,	NE)	89	9 (10.1)	NE (NE,	NE)	4.94	2.61, 10.61	<0.0001*
IV	73	29 (39.7)	NE (NE,	NE)	42	3 (7.1)	NE (NE,	NE)	6.57	2.33, 27.44	<0.0001*
Interaction p-value											0.6793
Region											
Europe	245	96 (39.2)	NE (NE,	NE)	125	12 (9.6)	NE (NE,	NE)	4.93	2.82, 9.48	<0.0001*
Japan	10	6 (60.0)	2.8 (0.3,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	74 (38.9)	NE (NE,	NE)	100	8 (8.0)	NE (NE,	NE)	6.14	3.15, 13.84	<0.0001*
(1) Restricted activity	61	28 (45.9)	NE (NE,	NE)	30	4 (13.3)	NE (NE,	NE)	3.89	1.53, 13.16	0.0029*
Interaction p-value											0.4936
Baseline CA-125 value											
<=ULN	228	89 (39.0)	NE (NE,	NE)	117	11 (9.4)	NE (NE,	NE)	5.06	2.83, 10.04	<0.0001*
>ULN	27	13 (48.1)	NE (NE,	NE)	14	1 (7.1)	NE (NE,	NE)	8.66	1.73,157.31	0.0049*
Interaction p-value											0.5983
Histological grade											
High grade	255	102 (40.0)	NE (NE,	NE)	131	12 (9.2)	NE (NE,	NE)	5.35	3.07, 10.26	<0.0001*
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	69 (41.6)	NE (NE,	NE)	80	8 (10.0)	NE (NE,	NE)	5.32	2.72, 12.01	<0.0001*
Residue	79	28 (35.4)	NE (NE,	NE)	43	2 (4.7)	NE (NE,	NE)	8.56	2.58, 53.08	<0.0001*
Interaction p-value											0.5468

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aai 25NOV2020:11:06 kvbv306

Table 3.4.9 PAOLA1: Summary of subgroup analysis of AE PT: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		_	bevacizumab 255)		Placebo + b (N=1				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) Median ti of patients (95% CI) n with events (months) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery								
Upfront	146	52 (35.6)	NE (NE,	NE)	8 10 (12.8)	NE (NE, NE)	3.27	1.74, 6.83	0.0001*
Interval	99	45 (45.5)	NE (NE,	NE) 4	.5 0	NE (NE, NE)	NC	NC	NC
Interaction p-value									NC
Myriad tumour BRCA mutat	ion stat	us							
tBRCAm	158	63 (39.9)	NE (NE,	NE)	7 6 (7.8)	NE (NE, NE)	6.25	2.94, 16.17	<0.0001*
Non-tBRCAm	97	39 (40.2)	NE (NE,	NE) 5	6 (11.1)	NE (NE, NE)	4.46	2.04, 11.73	<0.0001*
Interaction p-value									0.5823
Status somatic BRCA mutat	cions								
sBRCAm	22	10 (45.5)	NE (NE,	NE)	7 0	NE (NE, NE)	NC	NC	NC
gBRCAm	66	27 (40.9)	NE (NE,	NE) 3	1 4 (12.9)	NE (NE, NE)	4.11	1.61, 13.93	0.0020*
Non-BRCAm	41	20 (48.8)	21.7 (2.8,	NE) 2	12 0	NE (NE, NE)	NC	NC	NC
Interaction p-value									NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aai 25NOV2020:11:06 kvbv306

Table 3.4.10 PAOLA1: Summary of subgroup analysis of AE PT: Leukopenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olaparib + h (N=2		ımab	:		Placebo + be (N=13		mab				
Subgroup		Number (%) of patients n with events		Median time (95% CI) (months) [a]		Number (%) of patients n with events		Median time (95% CI) (months) [a]		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]	
First line treatment out	come (IV	RS)											
NED [PDS]	92	16 (17.4)	NE (NE,	NE)	48	3 (6.3)	NE (NE,	NE)	2.78	0.93, 11.96	0.0701
NED/CR [IDS]	74	17 (23.0)	NE (NE,	NE)	38	5 (13.2)	NE (NE,	NE)	1.79	0.71, 5.44	0.2303
NED/CR [Chemo]	40	7 (17.5)	NE (NE,	NE)	20	2 (10.0)	NE (NE,	NE)	1.86	0.45, 12.50	0.4140
PR	49	6 (12.2)	NE (NE,	NE)	25	1 (4.0)	NE (NE,	NE)	3.07	0.52, 58.04	0.2390
Interaction p-value													0.9294
Screening laboratory tBR0	CA statu	s (IVRS)											
tBRCAm	150	26 (17.3)	NE (NE,	NE)	65	4 (6.2)	NE (NE,	NE)	2.89	1.13, 9.80	0.0254*
non-tBRCAm	105	20 (19.0)	NE (NE,	NE)	66	7 (10.6)	NE (NE,	NE)	1.82	0.81, 4.65	0.1534
Interaction p-value													0.5012
First line treatment out	come (eCl	RF)											
NED [PDS]	89	18 (20.2)	NE (NE,	NE)	47	3 (6.4)	NE (NE,	NE)	3.23	1.09, 13.81	0.0322*
NED/CR [IDS]	74	16 (21.6)	NE (NE,	NE)	32	5 (15.6)	NE (NE,	NE)	1.41	0.55, 4.31	0.4921
NED/CR [Chemo]	39	3 (7.7)	NE (NE,	NE)	17	1 (5.9)	NE (NE,	NE)	1.32	0.17, 26.64	0.8071
PR	50	9 (18.0)	NE (NE,	NE)	34	2 (5.9)	NE (NE,	NE)	3.11	0.80, 20.41	0.1063
Interaction p-value													0.6807
Screening laboratory tBR0	CA statu	s (eCRF)											
tBRCAm	147	26 (17.7)	NE (NE,	NE)	67	5 (7.5)	NE (NE,	NE)	2.43	1.02, 7.19	0.0458*
non-tBRCAm	108	20 (18.5)	NE (NE,	NE)	64	6 (9.4)	NE (NE,	NE)	2.00	0.85, 5.47	0.1153
Interaction p-value													0.7722
Age group													
<65 years	185	29 (15.7)	NE (NE,	NE)	98	8 (8.2)	NE (NE,	NE)	1.96	0.94, 4.60	0.0747
>=65 years	70	17 (24.3)	NE (NE,	NE)	33	3 (9.1)	NE (NE,	NE)	2.69	0.90, 11.53	0.0784
Interaction p-value													0.6642

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaj 25NOV2020:11:06 kvbv306

Table 3.4.10 PAOLA1: Summary of subgroup analysis of AE PT: Leukopenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)					Placebo + b						
Subgroup		Number (%) of patients with events	Median t (95% C (months)	I)		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]		[b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	38 (20.9)	NE (NE	, NE)	89	7 (7.9)	NE (NE,	NE)	2.79	1.33,	6.83	0.0054*
IV	73	8 (11.0)	NE (NE	, NE)	42	4 (9.5)	NE (NE,	NE)	1.09	0.34,	4.08	0.8907
Interaction p-value												0.2117
Region												
Europe	245	39 (15.9)	NE (NE	, NE)	125	7 (5.6)	NE (NE,	NE)	2.89	1.38,	7.07	0.0038*
Japan	10	7 (70.0)	2.4 (0.7	,18.0)	6	4 (66.7)	2.6 (0.5,	NE)	1.11	0.33,	4.26	0.8664
Interaction p-value												0.2123
ECOG performance status at	Basel	ine										
(0) Normal activity	190	34 (17.9)	NE (NE	, NE)	100	8 (8.0)	NE (NE,	NE)	2.33	1.14,	5.42	0.0196*
(1) Restricted activity	61	11 (18.0)	NE (NE	, NE)	30	3 (10.0)	NE (NE,	NE)	1.67	0.52,	7.38	0.4113
Interaction p-value												0.6646
Baseline CA-125 value												
<=ULN	228	41 (18.0)	NE (NE		117	11 (9.4)	NE (NE,	NE)	1.97	1.05,	4.04	0.0337*
>ULN	27	5 (18.5)	NE (NE	, NE)	14	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value												NC
Histological grade												
High grade	255	46 (18.0)	NE (NE	, NE)	131	11 (8.4)	NE (NE,	NE)	2.19	1.18,	4.45	0.0123*
Interaction p-value												NC
Cytoreductive surgery outco	me											
No residue	166	34 (20.5)	NE (NE	, NE)	80	8 (10.0)	NE (NE,	NE)	2.09	1.02,	4.86	0.0440*
Residue	79	9 (11.4)	NE (NE	, NE)	43	3 (7.0)	NE (NE,	NE)	1.61	0.48,	7.26	0.4590
Interaction p-value												0.7374

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaj 25NOV2020:11:06 kvbv306

Table 3.4.10 PAOLA1: Summary of subgroup analysis of AE PT: Leukopenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2			Placebo + be (N=1						
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	C	Number (%) of patients with events	Median time (95% CI) (months) [a]		Hazard ratio [b] 95% CI [b]		[b]	2-sided p-value [b]
Timing of cytoreductive :	surgery										
Upfront	146	22 (15.1)	NE (NE, NE)	78	5 (6.4)	NE (NE,	NE)	2.38	0.97,	7.11	0.0574
Interval	99	21 (21.2)	NE (NE, NE)	45	6 (13.3)	NE (NE,	NE)	1.58	0.68,	4.31	0.3043
Interaction p-value											0.5450
Myriad tumour BRCA mutat:	ion stat	us									
tBRCAm	158	28 (17.7)	NE (NE, NE)	77	6 (7.8)	NE (NE,	NE)	2.31	1.02,	6.18	0.0431*
Non-tBRCAm	97	18 (18.6)	NE (NE, NE)	54	5 (9.3)	NE (NE,	NE)	2.05	0.82,	6.21	0.1318
Interaction p-value											0.8588
Status somatic BRCA muta	cions										
sBRCAm	22	7 (31.8)	NE (NE, NE)	7	0	NE (NE,	NE)	NC	NC		NC
gBRCAm	66	13 (19.7)	NE (NE, NE)	31	3 (9.7)	NE (NE,	NE)	2.18	0.70,	9.50	0.1905
Non-BRCAm	41	10 (24.4)	NE (NE, NE)	22	3 (13.6)	NE (NE,	NE)	1.87	0.57,	8.35	0.3171
Interaction p-value											0.8685

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaj 25NOV2020:11:06 kvbv306

Table 3.4.11 PAOLA1: Summary of subgroup analysis of AE PT: Lymphopenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		mab		Placebo + be (N=1)					
Subgroup		Number (%) of patients n with events		Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IV	RS)									
NED [PDS]	92	23 (25.0)	NE (NE, NE)	48	2 (4.2)	NE (NE,	NE)	5.95	1.76, 37.07	0.0020*
NED/CR [IDS]	74	22 (29.7)	NE (NE, NE)	38	3 (7.9)	NE (NE,	NE)	4.41	1.53, 18.65	0.0041*
NED/CR [Chemo]	40	7 (17.5)	NE (NE, NE)	20	3 (15.0)	NE (NE,	NE)	1.11	0.31, 5.14	0.8832
PR	49	8 (16.3)	NE (NE, NE)	25	2 (8.0)	NE (NE,	NE)	2.12	0.53, 14.05	0.3086
Interaction p-value											0.3356
Screening laboratory tBR0	CA statu	s (IVRS)									
tBRCAm	150	36 (24.0)	NE (NE, NE)	65	4 (6.2)	NE (NE,	NE)	4.13	1.65, 13.82	0.0012*
non-tBRCAm	105	24 (22.9)	NE (NE, NE)	66	6 (9.1)	NE (NE,	NE)	2.58	1.13, 6.97	0.0239*
Interaction p-value											0.4957
First line treatment out	come (eCl	RF)									
NED [PDS]	89	23 (25.8)	NE (NE, NE)	47	2 (4.3)	NE (NE,	NE)	6.14	1.82, 38.23	0.0016*
NED/CR [IDS]	74	18 (24.3)	NE (NE, NE)	32	1 (3.1)	NE (NE,	NE)	8.90	1.84,160.16	0.0030*
NED/CR [Chemo]	39	7 (17.9)	NE (NE, NE)	17	4 (23.5)	NE (NE,	NE)	0.67	0.20, 2.56	0.5319
PR	50	11 (22.0)	NE (NE, NE)	34	3 (8.8)	NE (NE,	NE)	2.65	0.83, 11.71	0.1053
Interaction p-value											0.0596
Screening laboratory tBR0	CA statu	s (eCRF)									
tBRCAm	147	35 (23.8)	NE (NE, NE)	67	4 (6.0)	NE (NE,	NE)	4.20	1.68, 14.05	0.0011*
non-tBRCAm	108	25 (23.1)	NE (NE, NE)	64	6 (9.4)	NE (NE,	NE)	2.55	1.12, 6.87	0.0248*
Interaction p-value											0.4708
Age group											
<65 years	185	34 (18.4)	NE (NE, NE)	98	7 (7.1)	NE (NE,	NE)	2.61	1.23, 6.44	0.0108*
>=65 years	70	26 (37.1)	NE (NE, NE)	33	3 (9.1)	NE (NE,	NE)	4.55	1.60, 19.10	0.0026*
Interaction p-value											0.4413

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aak 25NOV2020:11:06 kvbv306

Table 3.4.11 PAOLA1: Summary of subgroup analysis of AE PT: Lymphopenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2				Placebo + be (N=1		·		
Subgroup		Number (%) of patients with events	Median t (95% C (months)	[)		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	46 (25.3)	NE (NE	, NE)	89	4 (4.5)	NE (NE, N	6.10	2.48, 20.22	<0.0001*
IV	73	14 (19.2)	NE (NE	, NE)	42	6 (14.3)	NE (NE, N	1.27	0.51, 3.60	0.6153
Interaction p-value										0.0266*
Region										
Europe	245	56 (22.9)	NE (NE	, NE)	125	7 (5.6)	NE (NE, N	4.29	2.09, 10.34	<0.0001*
Japan	10	4 (40.0)	NE (NE	, NE)	6	3 (50.0)	NE (NE, N	0.67	0.15, 3.42	0.6072
Interaction p-value										0.0418*
ECOG performance status at	Basel	ine								
(0) Normal activity	190	39 (20.5)	NE (NE	NE)	100	6 (6.0)	NE (NE, N	3.61	1.65, 9.49	0.0007*
(1) Restricted activity	61	20 (32.8)	NE (NE	, NE)	30	4 (13.3)	NE (NE, N	2.43	0.92, 8.35	0.0766
Interaction p-value										0.5764
Baseline CA-125 value										
<=ULN	228	52 (22.8)	NE (NE	, NE)	117	8 (6.8)	NE (NE, N	3.49	1.76, 7.96	0.0002*
>ULN	27	8 (29.6)	NE (NE	, NE)	14	2 (14.3)	NE (NE, N	2.04	0.51, 13.49	0.3370
Interaction p-value										0.5531
Histological grade										
High grade	255	60 (23.5)	NE (NE	NE)	131	10 (7.6)	NE (NE, N	3.21	1.72, 6.67	0.0001*
Interaction p-value										NC
Cytoreductive surgery outco	ome									
No residue	166	42 (25.3)	NE (NE	, NE)	80	3 (3.8)	NE (NE, N	7.23	2.63, 29.84	<0.0001*
Residue	79	16 (20.3)	NE (NE	, NE)	43	6 (14.0)	NE (NE, N	1.40	0.58, 3.92	0.4676
Interaction p-value										0.0266*

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aak 25NOV2020:11:06 kvbv306

Table 3.4.11 PAOLA1: Summary of subgroup analysis of AE PT: Lymphopenia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2			Placebo + b (N=1		·		
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	- Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	urgery								
Upfront	146	30 (20.5)	NE (NE, NE) 78	3 (3.8)	NE (NE, NI	E) 5.35	1.91, 22.32	0.0006*
Interval	99	28 (28.3)	NE (NE, NE) 45	6 (13.3)	NE (NE, NI	E) 2.30	1.02, 6.15	0.0446*
Interaction p-value									0.2499
Myriad tumour BRCA mutati	on stat	us							
tBRCAm	158	38 (24.1)	NE (NE, NE) 77	4 (5.2)	NE (NE, NI	E) 4.91	1.97, 16.39	0.0002*
Non-tBRCAm	97	22 (22.7)	NE (NE, NE) 54	6 (11.1)	NE (NE, NI	E) 2.07	0.89, 5.62	0.0932
Interaction p-value									0.2101
Status somatic BRCA mutat	ions								
sBRCAm	22	8 (36.4)	NE (NE, NE) 7	0	NE (NE, NI	E) NC	NC	NC
gBRCAm	66	20 (30.3)	NE (NE, NE) 31	3 (9.7)	NE (NE, N	3.73	1.28, 15.82	0.0135*
Non-BRCAm	41	14 (34.1)	NE (NE, NE) 22	5 (22.7)	NE (NE, NI	E) 1.43	0.55, 4.43	0.4798
Interaction p-value									0.2291

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aak 25NOV2020:11:06 kvbv306

Table 3.4.12 PAOLA1: Summary of subgroup analysis of AE SOC: Gastrointestinal disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + N			Placebo + b					
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment outo	come (IV	RS)								
NED [PDS]	92	64 (69.6)	0.9 (0.3, 2.5)	48	32 (66.7)	4.2 (1.8,12.5)	1.38	0.91,	2.13	0.1338
NED/CR [IDS]	74	64 (86.5)	0.5 (0.2, 1.4)	38	28 (73.7)	4.2 (1.4,10.0)	1.65	1.07,	2.61	0.0237*
NED/CR [Chemo]	40	29 (72.5)	0.6 (0.2, 8.7)	20	13 (65.0)	7.2 (0.9, NE)	1.39	0.74,	2.77	0.3150
PR	49	32 (65.3)	4.1 (0.3,15.7)	25	10 (40.0)	NE (NE, NE)	2.13	1.08,	4.56	0.0274*
Interaction p-value										0.7353
Screening laboratory tBRO	CA statu	ıs (IVRS)								
tBRCAm	150	115 (76.7)	0.7 (0.4, 1.9)	65	39 (60.0)	7.7 (3.4,22.6)	1.75	1.23,	2.56	0.0016*
non-tBRCAm	105	74 (70.5)	1.0 (0.3, 3.3)	66	44 (66.7)	4.6 (2.1,10.6)	1.34	0.93,	1.96	0.1221
Interaction p-value										0.3071
First line treatment outo	come (eC	CRF)								
NED [PDS]	89	62 (69.7)	0.9 (0.3, 2.7)	47	30 (63.8)	6.9 (2.1,13.0)	1.48	0.96,	2.31	0.0739
NED/CR [IDS]	74	61 (82.4)	0.6 (0.2, 1.6)	32	25 (78.1)	4.2 (1.4, 9.7)	1.50	0.95,	2.44	0.0797
NED/CR [Chemo]	39	28 (71.8)	0.6 (0.2, 9.7)	17	10 (58.8)	7.8 (0.5,19.0)	1.37	0.69,	2.97	0.3790
PR	50	36 (72.0)	2.0 (0.3, 9.0)	34	17 (50.0)	9.7 (3.4, NE)	1.84	1.05,	3.36	0.0325*
Interaction p-value										0.9145
Screening laboratory tBRO	CA statu	ıs (eCRF)								
tBRCAm	147	113 (76.9)	0.8 (0.5, 1.9)	67	41 (61.2)	7.7 (2.9,14.4)	1.70	1.20,	2.46	0.0026*
non-tBRCAm	108	76 (70.4)	1.0 (0.3, 3.3)	64	42 (65.6)	4.8 (2.1,10.7)	1.38	0.95,	2.03	0.0899
Interaction p-value										0.4332
Age group										
<65 years	185	136 (73.5)	0.7 (0.3, 1.6)	98	59 (60.2)	7.8 (3.6,13.0)	1.69	1.25,	2.31	0.0006*
>=65 years	70	53 (75.7)	1.5 (0.4, 4.7)	33	24 (72.7)	3.5 (0.8,10.6)	1.21	0.76,	2.00	0.4297
Interaction p-value										0.2613

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aal 25NOV2020:11:06 kvbv306

Table 3.4.12 PAOLA1: Summary of subgroup analysis of AE SOC: Gastrointestinal disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k				bevacizumab 131)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	136 (74.7)	0.8 (0.4, 1.9)	89	61 (68.5)	4.6 (2.1, 9.7)	1.41	1.05,	1.92	0.0234*
IV	73	53 (72.6)	0.8 (0.3, 2.6)	42	22 (52.4)	10.7 (3.7, NE)	1.91	1.17,	3.20	0.0085*
Interaction p-value										0.3053
Region										
Europe	245	185 (75.5)	0.8 (0.4, 1.6)	125	79 (63.2)	6.9 (3.3,10.6)	1.59	1.22,	2.08	0.0004*
Japan	10	4 (40.0)	NE (NE, NE)	6	4 (66.7)	14.3 (1.4, NE)	0.69	0.16,	2.93	0.6036
Interaction p-value										0.2541
ECOG performance status at	Basel	ine								
(0) Normal activity	190	139 (73.2)	0.8 (0.4, 2.0)	100	60 (60.0)	9.0 (3.3,14.4)	1.64	1.22,	2.24	0.0010*
(1) Restricted activity	61	47 (77.0)	1.4 (0.2, 4.7)	30	22 (73.3)	4.5 (2.7, 7.8)	1.30	0.79,	2.20	0.2988
Interaction p-value										0.4466
Baseline CA-125 value										
<=ULN	228	168 (73.7)	0.8 (0.4, 2.0)	117	74 (63.2)	, , , ,	1.52	1.16,		0.0023*
>ULN	27	21 (77.8)	0.7 (0.1, 4.8)	14	9 (64.3)	8.3 (0.6, NE)	1.81	0.85,	4.15	0.1256
Interaction p-value										0.6786
Histological grade										
High grade	255	189 (74.1)	0.8 (0.5, 1.9)	131	83 (63.4)	6.9 (3.4,10.6)	1.55	1.20,	2.01	0.0007*
Interaction p-value										NC
Cytoreductive surgery outco	ome									
No residue	166	125 (75.3)	0.8 (0.3, 1.4)	80	56 (70.0)	4.2 (2.6,10.0)	1.47	1.08,	2.04	0.0140*
Residue	79	56 (70.9)	1.2 (0.3, 6.1)	43	24 (55.8)	9.0 (2.9, NE)	1.51	0.95,	2.48	0.0851
Interaction p-value										0.9384

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aal 25NOV2020:11:06 kvbv306

Table 3.4.12 PAOLA1: Summary of subgroup analysis of AE SOC: Gastrointestinal disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olaparib + N=2			Placebo + b (N=1					
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	99 (67.8)	1.1 (0.4, 3.7)	78	48 (61.5)	7.2 (2.8,13.0)	1.41	1.01,	2.01	0.0457*
Interval	99	82 (82.8)	0.6 (0.3, 1.4)	45	32 (71.1)	4.6 (1.4, 9.7)	1.55	1.04,	2.36	0.0321*
Interaction p-value										0.7413
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	119 (75.3)	0.7 (0.4, 1.9)	77	48 (62.3)	5.0 (2.8,13.0)	1.60	1.15,	2.26	0.0049*
Non-tBRCAm	97	70 (72.2)	1.3 (0.3, 3.4)	54	35 (64.8)	7.7 (2.6,11.3)	1.46	0.98,	2.22	0.0611
Interaction p-value										0.7445
Status somatic BRCA mutat	ions									
sBRCAm	22	16 (72.7)	0.6 (0.1, 9.0)	7	3 (42.9)	NE (NE, NE)	2.10	0.70,	9.03	0.2016
gBRCAm	66	56 (84.8)	0.6 (0.2, 1.6)	31	21 (67.7)	4.6 (2.8,22.6)	1.92	1.18,	3.26	0.0081*
Non-BRCAm	41	30 (73.2)	1.4 (0.2, 3.4)	22	15 (68.2)	4.6 (0.9,10.7)	1.39	0.76,	2.65	0.2943
Interaction p-value										0.6885

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aal 25NOV2020:11:06 kvbv306

Table 3.4.13 PAOLA1: Summary of subgroup analysis of AE PT: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		b		Placebo + be (N=1		b	:			
Subgroup		Number (%) of patients with events	Median (95% ((months	CI)		Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI	I [b]	2-sided p-value [b]
First line treatment outo	come (IV	RS)										
NED [PDS]	92	23 (25.0)	NE (N	E, NE)	48	7 (14.6)	NE (N	E, NE)	1.80	0.81,	4.54	0.1539
NED/CR [IDS]	74	18 (24.3)	NE (N	E, NE)	38	6 (15.8)	NE (N	E, NE)	1.58	0.66,	4.37	0.3133
NED/CR [Chemo]	40	4 (10.0)	NE (N	E, NE)	20	1 (5.0)	NE (N	E, NE)	2.00	0.30,	39.05	0.5097
PR	49	9 (18.4)	NE (N	E, NE)	25	2 (8.0)	NE (N	E, NE)	2.50	0.64,	16.40	0.2001
Interaction p-value												0.9655
Screening laboratory tBR0	CA statu	s (IVRS)										
tBRCAm	150	34 (22.7)	NE (N	E, NE)	65	7 (10.8)	NE (N	E, NE)	2.21	1.04,	5.43	0.0389*
non-tBRCAm	105	20 (19.0)	NE (N	E, NE)	66	9 (13.6)	NE (N	E, NE)	1.45	0.68,	3.36	0.3420
Interaction p-value												0.4680
First line treatment out	come (eC	RF)										
NED [PDS]	89	21 (23.6)	NE (N	E, NE)	47	7 (14.9)	NE (N	E, NE)	1.68	0.75,	4.25	0.2189
NED/CR [IDS]	74	17 (23.0)	NE (N	E, NE)	32	4 (12.5)	NE (N	E, NE)	1.94	0.72,	6.75	0.2033
NED/CR [Chemo]	39	5 (12.8)	NE (N	E, NE)	17	2 (11.8)	NE (N	E, NE)	1.08	0.23,	7.56	0.9235
PR	50	9 (18.0)	NE (N	E, NE)	34	3 (8.8)	NE (N	E, NE)	2.07	0.62,	9.34	0.2495
Interaction p-value												0.9370
Screening laboratory tBR0	CA statu	s (eCRF)										
tBRCAm	147	33 (22.4)	NE (N	E, NE)	67	7 (10.4)	NE (N	E, NE)	2.25	1.06,	5.54	0.0348*
non-tBRCAm	108	21 (19.4)	NE (N	E, NE)	64	9 (14.1)	NE (N	E, NE)	1.44	0.68,	3.31	0.3516
Interaction p-value												0.4360
Age group												
<65 years	185	43 (23.2)	NE (N	E, NE)	98	10 (10.2)	NE (N	E, NE)	2.44	1.28,	5.13	0.0059*
>=65 years	70	11 (15.7)	NE (N	E, NE)	33	6 (18.2)	NE (N	E, NE)	0.84	0.32,	2.43	0.7286
Interaction p-value												0.0909

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aam 25NOV2020:11:06 kvbv306

Table 3.4.13 PAOLA1: Summary of subgroup analysis of AE PT: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		umab				pevacizumab 131)					
Subgroup		Number (%) of patients with events		an ti % CI; hs) [)		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	41 (22.5)	NE (NE,	NE)	89	12 (13.5)	NE (NE,	NE)	1.79	0.97,	3.57	0.0621
IV	73	13 (17.8)	NE (NE,	NE)	42	4 (9.5)	NE (NE,	NE)	1.83	0.65,	6.49	0.2693
Interaction p-value													0.9770
Region													
Europe	245	54 (22.0)	NE (NE,	NE)	125	14 (11.2)	NE (NE,	NE)	2.07	1.18,	3.87	0.0099*
Japan	10	0	NE (NE,	NE)	6	2 (33.3)	24.0 (1.4,	NE)	NC	NC		NC
Interaction p-value													NC
ECOG performance status at	Basel	ine											
(0) Normal activity	190	41 (21.6)	NE (NE,	NE)	100	11 (11.0)	NE (NE,	NE)	2.07	1.10,	4.23	0.0225*
(1) Restricted activity	61	13 (21.3)	NE (NE,	NE)	30	5 (16.7)	NE (NE,	NE)	1.29	0.48,	4.02	0.6254
Interaction p-value													0.4559
Baseline CA-125 value													
<=ULN	228	43 (18.9)	NE (NE,	NE)	117	15 (12.8)	NE (NE,	NE)	1.51	0.86,	2.81	0.1553
>ULN	27	11 (40.7)	NE (NE,	NE)	14	1 (7.1)	NE (NE,	NE)	6.64	1.29,1	21.39	0.0195*
Interaction p-value													0.1091
Histological grade													
High grade	255	54 (21.2)	NE (NE,	NE)	131	16 (12.2)	NE (NE,	NE)	1.81	1.06,	3.26	0.0293*
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	40 (24.1)	NE (NE,	NE)	80	11 (13.8)	NE (NE,	NE)	1.87	0.99,	3.82	0.0533
Residue	79	12 (15.2)	NE (NE,	NE)	43	5 (11.6)	NE (NE,	NE)	1.30	0.48,	4.09	0.6166
Interaction p-value													0.5715

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aam 25NOV2020:11:06 kvbv306

Table 3.4.13 PAOLA1: Summary of subgroup analysis of AE PT: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		Placebo + b (N=1			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	29 (19.9)	NE (NE, NE)	78 9 (11.5)	NE (NE, NE)	1.82 0.89, 4.07	0.1014
Interval	99	23 (23.2)	NE (NE, NE)	45 7 (15.6)	NE (NE, NE)	1.51 0.68, 3.81	0.3230
Interaction p-value							0.7498
Myriad tumour BRCA mutati	on stat	us					
tBRCAm	158	36 (22.8)	NE (NE, NE)	77 8 (10.4)	NE (NE, NE)	2.31 1.13, 5.36	0.0199*
Non-tBRCAm	97	18 (18.6)	NE (NE, NE)	54 8 (14.8)	NE (NE, NE)	1.28 0.58, 3.12	0.5544
Interaction p-value							0.3061
Status somatic BRCA mutat	ions						
sBRCAm	22	4 (18.2)	NE (NE, NE)	7 1 (14.3)	NE (NE, NE)	1.13 0.17, 22.09	0.9129
gBRCAm	66	15 (22.7)	NE (NE, NE)	31 4 (12.9)	NE (NE, NE)	1.93 0.70, 6.78	0.2154
Non-BRCAm	41	5 (12.2)	NE (NE, NE)	22 4 (18.2)	NE (NE, NE)	0.61 0.16, 2.47	0.4698
Interaction p-value							0.4193

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aam 25NOV2020:11:06 kvbv306

Table 3.4.14 PAOLA1: Summary of subgroup analysis of AE PT: Nausea Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olaparib + 1 (N=2			Placebo + be (N=1)			:		
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median t (95% C (months)	[)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IV	RS)								
NED [PDS]	92	52 (56.5)	3.7 (0.6, NE)	48	11 (22.9)	NE (NE	, NE)	3.34	1.81, 6.76	<0.0001*
NED/CR [IDS]	74	48 (64.9)	1.4 (0.3, 4.3)	38	13 (34.2)	NE (NE	, NE)	2.44	1.36, 4.69	0.0022*
NED/CR [Chemo]	40	21 (52.5)	0.6 (0.2, NE)	20	2 (10.0)	NE (NE	, NE)	7.71	2.26, 48.22	0.0003*
PR	49	23 (46.9)	NE (NE, NE)	25	4 (16.0)	NE (NE	, NE)	3.63	1.39, 12.37	0.0064*
Interaction p-value										0.4609
Screening laboratory tBR0	CA statu	ıs (IVRS)								
tBRCAm	150	87 (58.0)	2.6 (0.7,16.0)	65	12 (18.5)	NE (NE	, NE)	4.23	2.41, 8.15	<0.0001*
non-tBRCAm	105	57 (54.3)	3.3 (0.3, NE)	66	18 (27.3)	NE (NE	, NE)	2.60	1.56, 4.55	0.0002*
Interaction p-value										0.2328
First line treatment out	come (eC	RF)								
NED [PDS]	89	51 (57.3)	2.7 (0.5, NE)	47	10 (21.3)	NE (NE	, NE)	3.69	1.95, 7.71	<0.0001*
NED/CR [IDS]	74	49 (66.2)	1.0 (0.3, 3.4)	32	10 (31.3)	NE (NE	, NE)	2.89	1.53, 6.05	0.0007*
NED/CR [Chemo]	39	19 (48.7)	NE (NE, NE)	17	2 (11.8)	NE (NE	, NE)	5.91	1.71, 37.10	0.0027*
PR	50	24 (48.0)	14.5 (2.1, NE)	34	7 (20.6)	NE (NE	, NE)	2.71	1.23, 6.82	0.0121*
Interaction p-value										0.7610
Screening laboratory tBR0	CA statu	ıs (eCRF)								
tBRCAm	147	85 (57.8)	2.9 (0.8,16.0)	67	13 (19.4)	NE (NE	, NE)	3.97	2.30, 7.47	<0.0001*
non-tBRCAm	108	59 (54.6)	2.7 (0.3, NE)	64	17 (26.6)	NE (NE	, NE)	2.72	1.62, 4.81	<0.0001*
Interaction p-value										0.3475
Age group										
<65 years	185	109 (58.9)	2.1 (0.6, 8.0)	98	21 (21.4)	NE (NE	, NE)	3.74	2.39, 6.13	<0.0001*
>=65 years	70	35 (50.0)	19.7 (0.8, NE)	33	9 (27.3)	NE (NE	, NE)	2.28	1.15, 5.06	0.0175*
Interaction p-value										0.2774

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aan 25NOV2020:11:06 kvbv306

Table 3.4.14 PAOLA1: Summary of subgroup analysis of AE PT: Nausea Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1				bevacizumab 131)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	100 (54.9)	3.7 (0.8, NE)	89	25 (28.1)	NE (NE,	NE)	2.54	1.67, 4.03	<0.0001*
IV	73	44 (60.3)	2.1 (0.3, NE)	42	5 (11.9)	NE (NE,	NE)	6.98	3.04, 20.18	<0.0001*
Interaction p-value										0.0370*
Region										
Europe	245	141 (57.6)	2.6 (0.8, 9.8)	125	30 (24.0)	NE (NE,	NE)	3.19	2.18, 4.82	<0.0001*
Japan	10	3 (30.0)	NE (NE, NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
ECOG performance status at	Basel	ine								
(0) Normal activity	190	113 (59.5)	2.0 (0.5, 5.4)	100	19 (19.0)	NE (NE,	NE)	4.31	2.72, 7.24	<0.0001*
(1) Restricted activity	61	30 (49.2)	19.7 (0.8, NE)	30	11 (36.7)	14.1 (10.0,	NE)	1.61	0.83, 3.35	0.1650
Interaction p-value										0.0266*
Baseline CA-125 value										
<=ULN	228	129 (56.6)	2.7 (0.8,14.5)	117	28 (23.9)	NE (NE,	NE)	3.13	2.11, 4.81	<0.0001*
>ULN	27	15 (55.6)	4.1 (0.1, NE)	14	2 (14.3)	NE (NE,	NE)	5.24	1.48, 33.26	0.0076*
Interaction p-value										0.4854
Histological grade										
High grade	255	144 (56.5)	2.9 (0.8,14.5)	131	30 (22.9)	NE (NE,	NE)	3.27	2.24, 4.94	<0.0001*
Interaction p-value										NC
Cytoreductive surgery outco	ome									
No residue	166	101 (60.8)	1.4 (0.4, 5.0)	80	21 (26.3)	NE (NE,	NE)	3.17	2.02, 5.22	<0.0001*
Residue	79	39 (49.4)	14.5 (0.6, NE)	43	9 (20.9)	NE (NE,	NE)	2.95	1.50, 6.50	0.0012*
Interaction p-value										0.8703

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aan 25NOV2020:11:06 kvbv306

Table 3.4.14 PAOLA1: Summary of subgroup analysis of AE PT: Nausea Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + N=2			Placebo + be (N=1)		mab				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patient: n with event:		Median time (95% CI) (months) [a]		Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
Timing of cytoreductive s	urgery										
Upfront	146	77 (52.7)	5.4 (0.6, NE)	78	14 (17.9)	NE (NE, NE	4.00	2.34,	7.37	<0.0001*
Interval	99	63 (63.6)	1.9 (0.4, 5.0)	45	16 (35.6)	NE (NE, NE	2.26	1.34,	4.05	0.0018*
Interaction p-value											0.1570
Myriad tumour BRCA mutati	on stat	us									
tBRCAm	158	92 (58.2)	2.3 (0.7,14.5)	77	15 (19.5)	NE (NE, NE	4.00	2.39,	7.20	<0.0001*
Non-tBRCAm	97	52 (53.6)	3.4 (0.4, NE)	54	15 (27.8)	NE (NE, NE	2.52	1.45,	4.64	0.0007*
Interaction p-value											0.2519
Status somatic BRCA mutat	ions										
sBRCAm	22	13 (59.1)	2.2 (0.1, NE)	7	2 (28.6)	NE (NE, NE	2.45	0.68,	15.66	0.1906
gBRCAm	66	42 (63.6)	2.1 (0.3, 6.1)	31	6 (19.4)	NE (NE, NE	4.61	2.11,	12.11	<0.0001*
Non-BRCAm	41	22 (53.7)	2.7 (0.2, NE)	22	6 (27.3)	NE (NE, NE	2.56	1.10,	6.96	0.0275*
Interaction p-value											0.5900

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aan 25NOV2020:11:06 kvbv306

Table 3.4.15 PAOLA1: Summary of subgroup analysis of AE PT: Dysgeusia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		ımab			Placebo + be (N=1)		ımab				
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI))	C	Number (%) of patients with events	(95	an ti 5% CI ths))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IVI	RS)											
NED [PDS]	92	11 (12.0)	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	6 (8.1)	NE (NE,	NE)	38	1 (2.6)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	4 (10.0)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	2 (4.1)	NE (NE,	NE)	25	1 (4.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA status	s (IVRS)											
tBRCAm	150	15 (10.0)	NE (NE,	NE)	65	1 (1.5)	NE (NE,	NE)	6.73	1.36,121.65	0.0144*
non-tBRCAm	105	8 (7.6)	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	5.26	0.96, 97.53	0.0560
Interaction p-value													0.8676
First line treatment out	come (eCI	RF)											
NED [PDS]	89	9 (10.1)	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	8 (10.8)	NE (NE,	NE)	32	1 (3.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	3 (7.7)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	2 (4.0)	NE (NE,	NE)	34	1 (2.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA status	s (eCRF)											
tBRCAm	147	15 (10.2)	NE (NE,	NE)	67	1 (1.5)	NE (NE,	NE)	7.08	1.44,128.03	0.0114*
non-tBRCAm	108	8 (7.4)	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	4.95	0.91, 91.82	0.0672
Interaction p-value													0.8091
Age group													
<65 years	185	15 (8.1)	NE (NE,	NE)	98	2 (2.0)	NE (NE,	NE)	4.11	1.16, 26.06	0.0262*
>=65 years	70	8 (11.4)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aao 25NOV2020:11:06 kvbv306

Table 3.4.15 PAOLA1: Summary of subgroup analysis of AE PT: Dysgeusia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2				Placebo + be (N=13					
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]		(Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	18 (9.9)	NE (NE	, NE)	89	2 (2.2)	NE (NE,	NE)	4.58	1.32, 28.83	0.0131*
IV	73	5 (6.8)	NE (NE	, NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	22 (9.0)	NE (NE	, NE)	125	2 (1.6)	NE (NE,	NE)	5.82	1.72, 36.33	0.0025*
Japan	10	1 (10.0)	NE (NE	, NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	19 (10.0)	NE (NE	, NE)	100	2 (2.0)	NE (NE,	NE)	5.27	1.53, 33.04	0.0056*
(1) Restricted activity	61	4 (6.6)	NE (NE	, NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	21 (9.2)	NE (NE	, NE)	117	2 (1.7)	NE (NE,	NE)	5.64	1.66, 35.27	0.0032*
>ULN	27	2 (7.4)	NE (NE	, NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	23 (9.0)	NE (NE	, NE)	131	2 (1.5)	NE (NE,	NE)	6.14	1.82, 38.28	0.0016*
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	18 (10.8)	NE (NE	, NE)	80	1 (1.3)	NE (NE,	NE)	9.23	1.91,165.96	0.0024*
Residue	79	4 (5.1)	NE (NE	, NE)	43	1 (2.3)	NE (NE,	NE)	2.17	0.32, 42.39	0.4576
Interaction p-value											0.3524

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aao 25NOV2020:11:06 kvbv306

Table 3.4.15 PAOLA1: Summary of subgroup analysis of AE PT: Dysgeusia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2			Placebo + bo (N=1					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	13 (8.9)	NE (NE, NE	78	1 (1.3)	NE (NE,	NE)	7.23	1.44,131.32	0.0117*
Interval	99	9 (9.1)	NE (NE, NE	2) 45	1 (2.2)	NE (NE,	NE)	4.24	0.80, 78.18	0.0988
Interaction p-value										0.7194
Myriad tumour BRCA mutati	ion stat	us								
tBRCAm	158	14 (8.9)	NE (NE, NE	3) 77	1 (1.3)	NE (NE,	NE)	7.05	1.42,127.64	0.0124*
Non-tBRCAm	97	9 (9.3)	NE (NE, NE	54	1 (1.9)	NE (NE,	NE)	5.27	0.99, 97.09	0.0518
Interaction p-value										0.8439
Status somatic BRCA mutat	cions									
sBRCAm	22	2 (9.1)	NE (NE, NE	2) 7	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	66	7 (10.6)	NE (NE, NE	31	1 (3.2)	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	41	5 (12.2)	NE (NE, NE	22	1 (4.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aao 25NOV2020:11:06 kvbv306

Table 3.4.16 PAOLA1: Summary of subgroup analysis of AE PT: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + : (N=2		:			bevacizumab 131)			
Subgroup		Number (%) of patients with events	Median ti (95% CI) (months) [Number (%) of patients with events		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IVI	RS)								
NED [PDS]	92	45 (48.9)	13.8 (6.7,	NE)	48	26 (54.2)	10.4 (4.2, NE)	0.93	0.58, 1.53	0.7847
NED/CR [IDS]	74	34 (45.9)	16.6 (6.9,	NE)	38	22 (57.9)	4.9 (2.1, NE)	0.67	0.39, 1.15	0.1447
NED/CR [Chemo]	40	20 (50.0)	10.1 (3.4,	NE)	20	13 (65.0)	4.1 (0.8, NE)	0.75	0.38, 1.54	0.4238
PR	49	23 (46.9)	16.6 (4.4,	NE)	25	17 (68.0)	2.8 (0.7,11.3)	0.49	0.26, 0.94	0.0310*
Interaction p-value										0.4534
Screening laboratory tBRO	CA status	s (IVRS)								
tBRCAm	150	68 (45.3)	NE (NE,	NE)	65	40 (61.5)	5.6 (2.9,11.1)	0.65	0.44, 0.97	0.0344*
non-tBRCAm	105	54 (51.4)	8.3 (5.7,	NE)	66	38 (57.6)	4.3 (2.2, NE)	0.84	0.56, 1.29	0.4278
Interaction p-value										0.3673
First line treatment outo	come (eCI	RF)								
NED [PDS]	89	43 (48.3)	11.1 (6.7,	NE)	47	25 (53.2)	11.0 (4.2, NE)	0.95	0.59, 1.58	0.8485
NED/CR [IDS]	74	31 (41.9)	NE (NE,	NE)	32	17 (53.1)	6.9 (2.8, NE)	0.71	0.40, 1.31	0.2679
NED/CR [Chemo]	39	22 (56.4)	3.9 (0.7,	NE)	17	12 (70.6)	4.1 (0.8, NE)	0.89	0.45, 1.86	0.7482
PR	50	25 (50.0)	16.6 (5.6,	NE)	34	23 (67.6)	2.1 (0.7, 5.6)	0.46	0.26, 0.82	0.0087*
Interaction p-value										0.2677
Screening laboratory tBRO	CA status	s (eCRF)								
tBRCAm	147	66 (44.9)	NE (NE,	NE)	67	40 (59.7)	6.9 (3.3,11.3)	0.67	0.45, 1.002	0.0509
non-tBRCAm	108	56 (51.9)	8.3 (6.2,	NE)	64	38 (59.4)	4.2 (2.1, NE)	0.81	0.54, 1.24	0.3299
Interaction p-value										0.5067
Age group										
<65 years	185	80 (43.2)	NE (NE,	NE)	98	54 (55.1)	8.1 (4.1, NE)	0.72	0.51, 1.02	0.0679
>=65 years	70	42 (60.0)	5.7 (2.8,1	4.1)	33	24 (72.7)	2.8 (1.4, 5.4)	0.69	0.42, 1.16	0.1567
Interaction p-value										0.8877

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aap 25NOV2020:11:06 kvbv306

Table 3.4.16 PAOLA1: Summary of subgroup analysis of AE PT: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		-		bevacizumab 255)			Placebo + k	pevacizumab 131)	;			
Subgroup	n	Number (of paties with ever	nts	Median ti (95% CI) (months) [)		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	85 (46	7)	16.6 (8.3,	NE)	89	53 (59.6)	6.9 (4.1,12.6)	0.74	0.53,	1.05	0.0879
IV	73	37 (50	7)	9.7 (5.6,	NE)	42	25 (59.5)	3.4 (2.1, NE)	0.70	0.43,	1.19	0.1841
Interaction p-value												0.8823
Region												
Europe	245	119 (48	6)	13.8 (8.3,	NE)	125	72 (57.6)	5.5 (3.5,11.1)	0.78	0.58,	1.04	0.0938
Japan	10	3 (30	0)	NE (NE,	NE)	6	6 (100)	2.1 (0.3,24.3)	0.19	0.04,	0.70	0.0135*
Interaction p-value												0.0399*
ECOG performance status at	Basel	ine										
(0) Normal activity	190	86 (45	3)	NE (NE,	NE)	100	59 (59.0)	4.2 (2.8, 7.6)	0.67	0.48,	0.94	0.0210*
(1) Restricted activity	61	33 (54	1)	8.3 (5.1,	NE)	30	19 (63.3)	10.4 (2.8,12.6)	0.87	0.50,	1.55	0.6215
Interaction p-value												0.4451
Baseline CA-125 value												
<=ULN	228			16.6 (8.3,		117	70 (59.8)	, , ,	0.73	0.54,		0.0414*
>ULN	27	14 (51	9)	6.9 (2.8,	NE)	14	8 (57.1)	4.3 (0.7, NE)	0.72	0.31,	1.79	0.4585
Interaction p-value												0.9714
Histological grade												
High grade	255	122 (47	. 8)	14.1 (8.3,	NE)	131	78 (59.5)	5.4 (3.3,11.0)	0.73	0.55,	0.97	0.0308*
Interaction p-value												NC
Cytoreductive surgery outco	me											
No residue	166	75 (45	2)	NE (NE,	NE)	80	43 (53.8)	10.4 (4.2, NE)	0.82	0.57,	1.21	0.3190
Residue	79	40 (50	6)	11.1 (3.5,	NE)	43	27 (62.8)	4.1 (2.1,11.3)	0.69	0.42,	1.13	0.1399
Interaction p-value												0.5643

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aap 25NOV2020:11:06 kvbv306

Table 3.4.16 PAOLA1: Summary of subgroup analysis of AE PT: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				Placebo + b (N=1		,			
Subgroup		Number (%) of patients with events	Median ti (95% CI) (months) [Number (%) Median time of patients (95% CI) n with events (months) [a]		Hazard ratio [b]	95% CI	2-sided p-value [b]		
Timing of cytoreductive s	urgery										
Upfront	146	69 (47.3)	13.8 (7.6,	NE)	78	45 (57.7)	7.6 (4.2,24.3	0.81	0.56,	1.19	0.2794
Interval	99	46 (46.5)	16.6 (7.5,	NE)	45	25 (55.6)	5.6 (2.8, NE) 0.72	0.45,	1.19	0.1998
Interaction p-value											0.7137
Myriad tumour BRCA mutati	on stat	us									
tBRCAm	158	75 (47.5)	16.6 (8.4,	NE)	77	45 (58.4)	6.9 (3.5,11.3	0.73	0.51,	1.07	0.1017
Non-tBRCAm	97	47 (48.5)	8.3 (6.6,	NE)	54	33 (61.1)	4.3 (2.1, NE) 0.73	0.47,	1.14	0.1664
Interaction p-value											0.9887
Status somatic BRCA mutat	ions										
sBRCAm	22	10 (45.5)	NE (NE,	NE)	7	3 (42.9)	NE (NE, NE	0.88	0.27,	3.94	0.8526
gBRCAm	66	29 (43.9)	NE (NE,	NE)	31	19 (61.3)	6.9 (3.3, NE	0.66	0.37,	1.19	0.1607
Non-BRCAm	41	20 (48.8)	10.6 (6.2,	NE)	22	14 (63.6)	3.1 (0.8, NE	0.60	0.31,	1.22	0.1552
Interaction p-value											0.8725

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aap 25NOV2020:11:06 kvbv306

Table 3.4.17 PAOLA1: Summary of subgroup analysis of AE PT: Gastroenteritis Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	;	Olaparib + k (N=2		ımab			Placebo + be (N=13		mab				
Subgroup	(Number (%) of patients with events	Media (95 (mont	% CI)	С	Number (%) of patients with events		an ti % CI) hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IVF	RS)											
NED [PDS]	92	3 (3.3)	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	5 (6.8)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	4 (10.0)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	1 (2.0)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA status	s (IVRS)											
tBRCAm	150	8 (5.3)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	5 (4.8)	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment outo	come (eCF	RF)											
NED [PDS]	89	3 (3.4)	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	6 (8.1)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	3 (7.7)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	1 (2.0)	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA status	s (eCRF)											
tBRCAm	147	7 (4.8)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	6 (5.6)	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	7 (3.8)	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	6 (8.6)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aag 25NOV2020:11:06 kvbv306

Table 3.4.17 PAOLA1: Summary of subgroup analysis of AE PT: Gastroenteritis Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		b		Placebo + be (N=1)					
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]		(Number (%) of patients with events	Median (95% ((months)	!I)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	10 (5.5)	NE (NI	E, NE)	89	0	NE (NE	, NE)	NC	NC	NC
IV	73	3 (4.1)	NE (NI	E, NE)	42	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	13 (5.3)	NE (NI	E, NE)	125	0	NE (NE	, NE)	NC	NC	NC
Japan	10	0	NE (NI	E, NE)	6	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	9 (4.7)	NE (NI	E, NE)	100	0	NE (NE	, NE)	NC	NC	NC
(1) Restricted activity	61	4 (6.6)	NE (NI	E, NE)	30	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	13 (5.7)	NE (NI	E, NE)	117	0	NE (NE	, NE)	NC	NC	NC
>ULN	27	0	NE (NI	E, NE)	14	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	13 (5.1)	NE (NI	E, NE)	131	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Cytoreductive surgery outc	ome										
No residue	166	9 (5.4)	NE (NI	E, NE)	80	0	NE (NE	, NE)	NC	NC	NC
Residue	79	3 (3.8)	NE (NI	E, NE)	43	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaq 25NOV2020:11:06 kvbv306

Table 3.4.17 PAOLA1: Summary of subgroup analysis of AE PT: Gastroenteritis Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	(Olaparib + b (N=2!		Placebo + b (N=1				
Subgroup	C	Number (%) of patients vith events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery							
Upfront	146	5 (3.4)	NE (NE, NE)	78 0	NE (NE, NE)	NC	NC	NC
Interval	99	7 (7.1)	NE (NE, NE)	45 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC
Myriad tumour BRCA mutat:	ion statu	ıs						
tBRCAm	158	8 (5.1)	NE (NE, NE)	77 0	NE (NE, NE)	NC	NC	NC
Non-tBRCAm	97	5 (5.2)	NE (NE, NE)	54 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC
Status somatic BRCA muta	tions							
sBRCAm	22	0	NE (NE, NE)	7 0	NE (NE, NE)	NC	NC	NC
gBRCAm	66	6 (9.1)	NE (NE, NE)	31 0	NE (NE, NE)	NC	NC	NC
Non-BRCAm	41	2 (4.9)	NE (NE, NE)	22 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aag 25NOV2020:11:06 kvbv306

Table 3.4.18 PAOLA1: Summary of subgroup analysis of AE PT: Blood creatinine increased Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	,	Olaparib + b (N=2!		mab	:		Placebo + be (N=1)			
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]		C	Number (%) of patients with events	Median (95% ((months)	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]	
First line treatment out	come (IVF	RS)										
NED [PDS]	92	6 (6.5)	NE (NE,	NE)	48	0	NE (NE	E, NE)	NC	NC	NC
NED/CR [IDS]	74	3 (4.1)	NE (NE,	NE)	38	1 (2.6)	NE (NE	E, NE)	NC	NC	NC
NED/CR [Chemo]	40	4 (10.0)	NE (NE,	NE)	20	0	NE (NE	E, NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NI	E, NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR0	CA status	s (IVRS)										
tBRCAm	150	7 (4.7)	NE (NE,	NE)	65	1 (1.5)	NE (NE	E, NE)	NC	NC	NC
non-tBRCAm	105	6 (5.7)	NE (NE,	NE)	66	0	NE (NE	E, NE)	NC	NC	NC
Interaction p-value												NC
First line treatment out	come (eCF	RF)										
NED [PDS]	89	6 (6.7)	NE (NE,	NE)	47	0	NE (NI	E, NE)	NC	NC	NC
NED/CR [IDS]	74	3 (4.1)	NE (NE,	NE)	32	1 (3.1)	NE (NE	E, NE)	NC	NC	NC
NED/CR [Chemo]	39	3 (7.7)	NE (NE,	NE)	17	0	NE (NE	E, NE)	NC	NC	NC
PR	50	1 (2.0)	NE (NE,	NE)	34	0	NE (NE	E, NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR0	CA status	(eCRF)										
tBRCAm	147	7 (4.8)	NE (NE,	NE)	67	1 (1.5)	NE (NI	E, NE)	NC	NC	NC
non-tBRCAm	108	6 (5.6)	NE (NE,	NE)	64	0	NE (NE	E, NE)	NC	NC	NC
Interaction p-value												NC
Age group												
<65 years	185	6 (3.2)	NE (NE,	NE)	98	1 (1.0)	NE (NI	E, NE)	NC	NC	NC
>=65 years	70	7 (10.0)	NE (NE,	NE)	33	0	NE (NE	E, NE)	NC	NC	NC
Interaction p-value												NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aar 25NOV2020:11:06 kvbv306

Table 3.4.18 PAOLA1: Summary of subgroup analysis of AE PT: Blood creatinine increased Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b				Placebo + be (N=1)					
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]		C	Number (%) of patients with events	Median to (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	11 (6.0)	NE (NE,	NE)	89	1 (1.1)	NE (NE,	NE)	5.38	1.05, 98.34	0.0430*
IV	73	2 (2.7)	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	13 (5.3)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	1 (16.7)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	10 (5.3)	NE (NE,	NE)	100	1 (1.0)	NE (NE,	NE)	5.37	1.03, 98.48	0.0458*
(1) Restricted activity	61	3 (4.9)	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	12 (5.3)	NE (NE,	NE)	117	1 (0.9)	NE (NE,	NE)	6.01	1.18,109.57	0.0274*
>ULN	27	1 (3.7)	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	13 (5.1)	NE (NE,	NE)	131	1 (0.8)	NE (NE,	NE)	6.53	1.30,118.62	0.0185*
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	9 (5.4)	NE (NE,	NE)	80	1 (1.3)	NE (NE,	NE)	4.48	0.84, 82.51	0.0850
Residue	79	4 (5.1)	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aar 25NOV2020:11:06 kvbv306

Table 3.4.18 PAOLA1: Summary of subgroup analysis of AE PT: Blood creatinine increased Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2!		Placebo + b (N=1			
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]
Timing of cytoreductive	surgery						
Upfront	146	9 (6.2)	NE (NE, NE)	78 0	NE (NE, NE)	NC NC	NC
Interval	99	4 (4.0)	NE (NE, NE)	45 1 (2.2)	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Myriad tumour BRCA mutat	ion statu	ıs					
tBRCAm	158	7 (4.4)	NE (NE, NE)	77 1 (1.3)	NE (NE, NE)	NC NC	NC
Non-tBRCAm	97	6 (6.2)	NE (NE, NE)	54 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Status somatic BRCA muta	tions						
sBRCAm	22	1 (4.5)	NE (NE, NE)	7 0	NE (NE, NE)	NC NC	NC
gBRCAm	66	2 (3.0)	NE (NE, NE)	31 1 (3.2)	NE (NE, NE)	NC NC	NC
Non-BRCAm	41	1 (2.4)	NE (NE, NE)	22 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aar 25NOV2020:11:06 kvbv306

Table 3.4.19 PAOLA1: Summary of subgroup analysis of SAE Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	·	Olaparib + b		ımab			Placeb		pevacizumab 131)					
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI))		Number of pati with ev	ents	Median ti (95% CI (months))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)												
NED [PDS]	92	23 (25.0)	NE (NE,	NE)	48	15 (3	31.3)	NE (NE,	NE)	0.69	0.36,	1.35	0.2723
NED/CR [IDS]	74	20 (27.0)	NE (NE,	NE)	38	9 (2	23.7)	NE (NE,	NE)	1.06	0.50,	2.46	0.8792
NED/CR [Chemo]	40	16 (40.0)	NE (NE,	NE)	20	10 (5	0.0)	15.6 (4.2,	NE)	0.71	0.33,	1.63	0.4065
PR	49	14 (28.6)	NE (NE,	NE)	25	11 (4	4.0)	17.3 (6.7,	NE)	0.61	0.28,	1.38	0.2283
Interaction p-value														0.7722
Screening laboratory tBR0	CA statu	s (IVRS)												
tBRCAm	150	45 (30.0)	NE (NE,	NE)	65	24 (3	6.9)	NE (NE,	NE)	0.71	0.43,	1.18	0.1777
non-tBRCAm	105	28 (26.7)	NE (NE,	NE)	66	21 (3	31.8)	NE (NE,	NE)	0.80	0.45,	1.42	0.4346
Interaction p-value														0.7532
First line treatment out	come (eC	RF)												
NED [PDS]	89	23 (25.8)	NE (NE,	NE)	47	14 (2	9.8)	NE (NE,	NE)	0.78	0.41,	1.56	0.4716
NED/CR [IDS]	74	21 (28.4)	NE (NE,	NE)	32	8 (2	25.0)	NE (NE,	NE)	1.12	0.52,	2.70	0.7790
NED/CR [Chemo]	39	13 (33.3)	NE (NE,	NE)	17	9 (5	2.9)	10.6 (4.2,	NE)	0.56	0.24,	1.36	0.1954
PR	50	16 (32.0)	NE (NE,	NE)	34	13 (3	88.2)	17.3 (8.5,	NE)	0.71	0.34,	1.51	0.3662
Interaction p-value														0.6997
Screening laboratory tBR0	CA statu	s (eCRF)												
tBRCAm	147	43 (29.3)	NE (NE,	NE)	67	24 (3	5.8)	NE (NE,	NE)	0.71	0.43,	1.18	0.1809
non-tBRCAm	108	30 (27.8)	NE (NE,	NE)	64	21 (3	32.8)	NE (NE,	NE)	0.81	0.47,	1.43	0.4630
Interaction p-value														0.7191
Age group														
<65 years	185	49 (26.5)	NE (NE,	NE)	98	35 (3	35.7)	NE (NE,	NE)	0.65	0.42,	1.02	0.0590
>=65 years	70	24 (34.3)	NE (NE,	NE)	33	10 (3	30.3)	NE (NE,	NE)	1.10	0.54,	2.41	0.8017
Interaction p-value														0.2270

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aas 25NOV2020:11:06 kvbv306

Table 3.4.19 PAOLA1: Summary of subgroup analysis of SAE Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		mab				oevacizumab 131)					
Subgroup		Number (%) of patients with events	Media (95° (mont)	% CI)			Number (%) of patients with events	Median ti (95% CI) (months) [Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	55 (30.2)	NE (NE,	NE)	89	28 (31.5)	NE (NE,	NE)	0.93	0.60,	1.49	0.7721
IV	73	18 (24.7)	NE (NE,	NE)	42	17 (40.5)	16.2 (8.3,	NE)	0.46	0.23,	0.90	0.0233*
Interaction p-value													0.0822
Region													
Europe	245	70 (28.6)	NE (NE,	NE)	125	44 (35.2)	NE (NE,	NE)	0.72	0.50,	1.06	0.0959
Japan	10	3 (30.0)	NE (NE,	NE)	6	1 (16.7)	NE (NE,	NE)	2.25	0.29,	45.51	0.4566
Interaction p-value													0.2973
ECOG performance status at	Basel	ine											
(0) Normal activity	190	46 (24.2)	NE (NE,	NE)	100	35 (35.0)	NE (NE,	NE)	0.63	0.41,	0.99	0.0457*
(1) Restricted activity	61	26 (42.6)	NE (NE,	NE)	30	10 (33.3)	NE (NE,	NE)	1.10	0.55,	2.39	0.8010
Interaction p-value													0.1985
Baseline CA-125 value													
<=ULN	228	64 (28.1)	NE (NE,	NE)	117	42 (35.9)	NE (NE,	NE)	0.71	0.48,	1.06	0.0914
>ULN	27	9 (33.3)	NE (NE,	NE)	14	3 (21.4)	NE (NE,	NE)	1.33	0.40,	5.98	0.6647
Interaction p-value													0.3516
Histological grade													
High grade	255	73 (28.6)	NE (NE,	NE)	131	45 (34.4)	NE (NE,	NE)	0.75	0.52,	1.10	0.1416
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	44 (26.5)	NE (NE,	NE)	80	23 (28.8)	NE (NE,	NE)	0.86	0.52,	1.44	0.5516
Residue	79	22 (27.8)	NE (NE,	NE)	43	19 (44.2)	15.6 (9.1,	NE)	0.52	0.28,	0.97	0.0406*
Interaction p-value													0.2183

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aas 25NOV2020:11:06 kvbv306

Table 3.4.19 PAOLA1: Summary of subgroup analysis of SAE Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		Placebo + b (N=1			
Subgroup		Number (%) Median time of patients (95% CI) n with events (months) [a		Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	39 (26.7)	NE (NE, NE)	78 30 (38.5)	NE (NE, NE)	0.61 0.38, 0.99	6 0.0482*
Interval	99	27 (27.3)	NE (NE, NE)	45 12 (26.7)	NE (NE, NE)	0.93 0.48, 1.9	1 0.8425
Interaction p-value							0.3182
Myriad tumour BRCA mutati	on stat	us					
tBRCAm	158	49 (31.0)	NE (NE, NE)	77 29 (37.7)	NE (NE, NE)	0.72 0.46, 1.1	6 0.1769
Non-tBRCAm	97	24 (24.7)	NE (NE, NE)	54 16 (29.6)	NE (NE, NE)	0.79 0.42, 1.5	1 0.4639
Interaction p-value							0.8343
Status somatic BRCA mutat	ions						
sBRCAm	22	7 (31.8)	NE (NE, NE)	7 0	NE (NE, NE)	NC NC	NC
gBRCAm	66	20 (30.3)	NE (NE, NE)	31 12 (38.7)	NE (NE, NE)	0.75 0.37, 1.5	8 0.4343
Non-BRCAm	41	9 (22.0)	NE (NE, NE)	22 5 (22.7)	NE (NE, NE)	0.90 0.31, 2.9	5 0.8579
Interaction p-value							0.7760

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aas 25NOV2020:11:06 kvbv306

Table 3.4.20 PAOLA1: Summary of subgroup analysis of SAE SOC: Blood and lymphatic system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2				Placebo + be (N=1)					
Subgroup	C	Number (%) of patients with events	Median ti (95% CI (months))	C	Number (%) of patients with events	Median t (95% C] (months)	_)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVF	RS)									
NED [PDS]	92	5 (5.4)	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	4 (5.4)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	3 (7.5)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	5 (10.2)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	s (IVRS)									
tBRCAm	150	9 (6.0)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	8 (7.6)	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
First line treatment out	come (eCF	RF)									
NED [PDS]	89	5 (5.6)	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	6 (8.1)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	2 (5.1)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	4 (8.0)	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	s (eCRF)									
tBRCAm	147	9 (6.1)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	8 (7.4)	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Age group											
<65 years	185	11 (5.9)	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	5.91	1.15,108.03	0.0307*
>=65 years	70	6 (8.6)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aat 25NOV2020:11:06 kvbv306

Table 3.4.20 PAOLA1: Summary of subgroup analysis of SAE SOC: Blood and lymphatic system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		nab		Placebo + be (N=1					
Subgroup		Number (%) of patients with events		n time (CI) (S) [a]	(Number (%) of patients with events	Median to (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	9 (4.9)	NE (NE, NE) 89	1 (1.1)	NE (NE,	NE)	4.68	0.88, 86.36	0.0744
IV	73	8 (11.0)	NE (NE, NE) 42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	15 (6.1)	NE (NE, NE) 125	1 (0.8)	NE (NE,	NE)	7.70	1.56,139.17	0.0078*
Japan	10	2 (20.0)	NE (NE, NE) 6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel:	ine									
(0) Normal activity	190	12 (6.3)	NE (NE, NE) 100	1 (1.0)	NE (NE,	NE)	6.46	1.27,117.59	0.0204*
(1) Restricted activity	61	5 (8.2)	NE (NE, NE) 30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	15 (6.6)	NE (NE, NE) 117	1 (0.9)	NE (NE,	NE)	8.04	1.63,145.46	0.0062*
>ULN	27	2 (7.4)	NE (NE, NE) 14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	17 (6.7)	NE (NE, NE) 131	1 (0.8)	NE (NE,	NE)	8.91	1.83,160.63	0.0031*
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	11 (6.6)	NE (NE, NE) 80	1 (1.3)	NE (NE,	NE)	5.60	1.09,102.30	0.0374*
Residue	79	3 (3.8)	NE (NE, NE) 43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aat 25NOV2020:11:06 kvbv306

Table 3.4.20 PAOLA1: Summary of subgroup analysis of SAE SOC: Blood and lymphatic system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2				Placebo + be (N=13		nab			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a		C	Number (%) of patients with events	(95	an time % CI) hs) [a]	Hazaro ratio [b]	-	2-sided p-value [b]
Timing of cytoreductive	surgery										
Upfront	146	8 (5.5)	NE (NE,	NE)	78	1 (1.3)	NE (NE, NE) NC	NC	NC
Interval	99	6 (6.1)	NE (NE,	NE)	45	0	NE (NE, NE) NC	NC	NC
Interaction p-value											NC
Myriad tumour BRCA mutat	ion stat	us									
tBRCAm	158	12 (7.6)	NE (NE,	NE)	77	0	NE (NE, NE) NC	NC	NC
Non-tBRCAm	97	5 (5.2)	NE (NE,	NE)	54	1 (1.9)	NE (NE, NE) NC	NC	NC
Interaction p-value											NC
Status somatic BRCA muta	cions										
sBRCAm	22	3 (13.6)	NE (NE,	NE)	7	0	NE (NE, NE) NC	NC	NC
gBRCAm	66	6 (9.1)	NE (NE,	NE)	31	0	NE (NE, NE) NC	NC	NC
Non-BRCAm	41	2 (4.9)	NE (NE,	NE)	22	0	NE (NE, NE) NC	NC	NC
Interaction p-value											NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aat 25NOV2020:11:06 kvbv306

Table 3.4.21 PAOLA1: Summary of subgroup analysis of SAE PT: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=	bevaciz 255)	umab	:		Placebo + be (N=1		b			
Subgroup	C	Number (%) of patients with events	(9	an ti 5% CI ths))	C	Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVF	RS)										
NED [PDS]	92	5 (5.4)	NE (NE,	NE)	48	1 (2.1)	NE (N	E, NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (NE,	NE)	38	0	NE (N	E, NE)	NC	NC	NC
NED/CR [Chemo]	40	2 (5.0)	NE (NE,	NE)	20	0	NE (N	E, NE)	NC	NC	NC
PR	49	4 (8.2)	NE (NE,	NE)	25	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR0	CA status	s (IVRS)										
tBRCAm	150	6 (4.0)	NE (NE,	NE)	65	0	NE (N	E, NE)	NC	NC	NC
non-tBRCAm	105	7 (6.7)	NE (NE,	NE)	66	1 (1.5)	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
First line treatment out	come (eCF	RF)										
NED [PDS]	89	5 (5.6)	NE (NE,	NE)	47	1 (2.1)	NE (N	E, NE)	NC	NC	NC
NED/CR [IDS]	74	3 (4.1)	NE (NE,	NE)	32	0	NE (N	E, NE)	NC	NC	NC
NED/CR [Chemo]	39	2 (5.1)	NE (NE,	NE)	17	0	NE (N	E, NE)	NC	NC	NC
PR	50	3 (6.0)	NE (NE,	NE)	34	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR0	CA status	s (eCRF)										
tBRCAm	147	6 (4.1)	NE (NE,	NE)	67	0	NE (N	E, NE)	NC	NC	NC
non-tBRCAm	108	7 (6.5)	NE (NE,	NE)	64	1 (1.6)	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Age group												
<65 years	185	9 (4.9)	NE (NE,	NE)	98	1 (1.0)	NE (N	E, NE)	4.92	0.92, 90.72	0.0642
>=65 years	70	4 (5.7)	NE (NE,	NE)	33	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aau 25NOV2020:11:06 kvbv306

Table 3.4.21 PAOLA1: Summary of subgroup analysis of SAE PT: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2)		Placebo + be (N=1)					
Subgroup		Number (%) of patients with events	Median (95% ((months)	I)	(Number (%) of patients with events	Median t (95% C] (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	8 (4.4)	NE (NE	, NE)	89	1 (1.1)	NE (NE,	NE)	NC	NC	NC
IV	73	5 (6.8)	NE (NE	, NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	11 (4.5)	NE (NE	, NE)	125	1 (0.8)	NE (NE,	NE)	5.79	1.13,105.87	0.0329*
Japan	10	2 (20.0)	NE (NE	, NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	9 (4.7)	NE (NE	, NE)	100	1 (1.0)	NE (NE,	NE)	4.98	0.93, 91.79	0.0619
(1) Restricted activity	61	4 (6.6)	NE (NE	, NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	12 (5.3)	NE (NE	, NE)	117	1 (0.9)	NE (NE,	NE)	6.49	1.28,118.23	0.0199*
>ULN	27	1 (3.7)	NE (NE	, NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	13 (5.1)	NE (NE	, NE)	131	1 (0.8)	NE (NE,	NE)	6.96	1.39,126.39	0.0139*
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	8 (4.8)	NE (NE	, NE)	80	1 (1.3)	NE (NE,	NE)	NC	NC	NC
Residue	79	3 (3.8)	NE (NE	, NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aau 25NOV2020:11:06 kvbv306

Table 3.4.21 PAOLA1: Summary of subgroup analysis of SAE PT: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	,	Olaparib + b (N=2		Placebo + b (N=1			
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI []	2-sided b] p-value[b]
Timing of cytoreductive s	surgery						
Upfront	146	8 (5.5)	NE (NE, NE)	78 1 (1.3)	NE (NE, NE)	NC NC	NC
Interval	99	3 (3.0)	NE (NE, NE)	45 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Myriad tumour BRCA mutati	ion statu	ıs					
tBRCAm	158	8 (5.1)	NE (NE, NE)	77 0	NE (NE, NE)	NC NC	NC
Non-tBRCAm	97	5 (5.2)	NE (NE, NE)	54 1 (1.9)	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Status somatic BRCA mutat	cions						
sBRCAm	22	2 (9.1)	NE (NE, NE)	7 0	NE (NE, NE)	NC NC	NC
gBRCAm	66	3 (4.5)	NE (NE, NE)	31 0	NE (NE, NE)	NC NC	NC
Non-BRCAm	41	2 (4.9)	NE (NE, NE)	22 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aau 25NOV2020:11:06 kvbv306

Table 3.4.22 PAOLA1: Summary of subgroup analysis of AE leading to discontinuation of treatment Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	·	Olaparib + b (N=2		mab			Placebo + be (N=1)		ımab				
Subgroup		Number (%) of patients with events	Media (95 (mont)	% CI)		C	Number (%) of patients with events	(95	an ti 5% CI :hs))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IV	RS)											
NED [PDS]	92	16 (17.4)	NE (NE,	NE)	48	3 (6.3)	NE (NE,	NE)	2.64	0.88, 11.36	0.0873
NED/CR [IDS]	74	21 (28.4)	NE (NE,	NE)	38	4 (10.5)	NE (NE,	NE)	2.65	1.01, 9.09	0.0479*
NED/CR [Chemo]	40	7 (17.5)	NE (NE,	NE)	20	1 (5.0)	NE (NE,	NE)	3.36	0.60, 62.84	0.1909
PR	49	6 (12.2)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													0.9775
Screening laboratory tBR0	CA statu	s (IVRS)											
tBRCAm	150	31 (20.7)	NE (NE,	NE)	65	3 (4.6)	NE (NE,	NE)	4.36	1.56, 18.16	0.0030*
non-tBRCAm	105	19 (18.1)	NE (NE,	NE)	66	5 (7.6)	NE (NE,	NE)	2.35	0.94, 7.08	0.0678
Interaction p-value													0.4244
First line treatment out	come (eC	RF)											
NED [PDS]	89	17 (19.1)	NE (NE,	NE)	47	3 (6.4)	NE (NE,	NE)	2.91	0.98, 12.47	0.0556
NED/CR [IDS]	74	21 (28.4)	NE (NE,	NE)	32	3 (9.4)	NE (NE,	NE)	3.06	1.05, 12.97	0.0384*
NED/CR [Chemo]	39	6 (15.4)	NE (NE,	NE)	17	1 (5.9)	NE (NE,	NE)	2.49	0.42, 46.99	0.3500
PR	50	6 (12.0)	NE (NE,	NE)	34	1 (2.9)	NE (NE,	NE)	3.88	0.66, 73.29	0.1461
Interaction p-value													0.9928
Screening laboratory tBR0	CA statu	s (eCRF)											
tBRCAm	147	31 (21.1)	NE (NE,	NE)	67	3 (4.5)	NE (NE,	NE)	4.57	1.63, 19.06	0.0021*
non-tBRCAm	108	19 (17.6)	NE (NE,	NE)	64	5 (7.8)	NE (NE,	NE)	2.21	0.89, 6.68	0.0905
Interaction p-value													0.3481
Age group													
<65 years	185	34 (18.4)	NE (NE,	NE)	98	4 (4.1)	NE (NE,	NE)	4.43	1.76, 14.83	0.0007*
>=65 years	70	16 (22.9)	NE (NE,	NE)	33	4 (12.1)	NE (NE,	NE)	1.78	0.65, 6.23	0.2750
Interaction p-value													0.2418

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aav 25NOV2020:11:06 kvbv306

Table 3.4.22 PAOLA1: Summary of subgroup analysis of AE leading to discontinuation of treatment Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		nab		Placebo + be (N=1					
Subgroup		Number (%) of patients with events	(95%	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	39 (21.4)	NE (NE, NE) 89	6 (6.7)	NE (NE,	NE)	3.16	1.44, 8.32	0.0027*
IV	73	11 (15.1)	NE (NE, NE) 42	2 (4.8)	NE (NE,	NE)	2.95	0.79, 19.06	0.1149
Interaction p-value											0.9367
Region											
Europe	245	47 (19.2)	NE (NE, NE) 125	8 (6.4)	NE (NE,	NE)	2.89	1.44, 6.61	0.0019*
Japan	10	3 (30.0)	NE (NE, NE) 6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel:	ine									
(0) Normal activity	190	35 (18.4)	NE (NE, NE) 100	5 (5.0)	NE (NE,	NE)	3.69	1.58, 10.76	0.0015*
(1) Restricted activity	61	15 (24.6)	NE (NE, NE) 30	3 (10.0)	NE (NE,	NE)	2.18	0.72, 9.42	0.1822
Interaction p-value											0.5140
Baseline CA-125 value											
<=ULN	228	44 (19.3)	NE (NE, NE) 117	8 (6.8)	NE (NE,	NE)	2.79	1.39, 6.40	0.0030*
>ULN	27	6 (22.2)	NE (NE, NE) 14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	50 (19.6)	NE (NE, NE) 131	8 (6.1)	NE (NE,	NE)	3.13	1.57, 7.14	0.0007*
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	38 (22.9)	NE (NE, NE) 80	6 (7.5)	NE (NE,	NE)	3.03	1.38, 7.97	0.0043*
Residue	79	11 (13.9)	NE (NE, NE) 43	2 (4.7)	NE (NE,	NE)	2.82	0.76, 18.26	0.1313
Interaction p-value											0.9385

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aav 25NOV2020:11:06 kvbv306

Table 3.4.22 PAOLA1: Summary of subgroup analysis of AE leading to discontinuation of treatment Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2)		Placebo + b (N=1				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery							
Upfront	146	26 (17.8)	NE (NE, NE)	78 4 (5.1)	NE (NE, NE)	3.39	1.32, 11.49	0.0091*
Interval	99	23 (23.2)	NE (NE, NE)	45 4 (8.9)	NE (NE, NE)	2.52	0.97, 8.58	0.0594
Interaction p-value								0.6969
Myriad tumour BRCA mutati	on stat	us						
tBRCAm	158	33 (20.9)	NE (NE, NE)	77 5 (6.5)	NE (NE, NE)	3.10	1.32, 9.06	0.0072*
Non-tBRCAm	97	17 (17.5)	NE (NE, NE)	54 3 (5.6)	NE (NE, NE)	3.13	1.05, 13.40	0.0396*
Interaction p-value								0.9909
Status somatic BRCA mutat	ions							
sBRCAm	22	3 (13.6)	NE (NE, NE)	7 0	NE (NE, NE)	NC	NC	NC
gBRCAm	66	14 (21.2)	NE (NE, NE)	31 0	NE (NE, NE)	NC	NC	NC
Non-BRCAm	41	6 (14.6)	NE (NE, NE)	22 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aav 25NOV2020:11:06 kvbv306

Table 3.4.23 PAOLA1: Summary of subgroup analysis of AE max CTCAE grade>=3
Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olapar	ib + bevacizum (N=255)	mab		pevacizumab 131)					
Subgroup	Number of pat n with e	ients (95%	n time s CI) ns) [a]	Number (%) of patients n with events	Median tir (95% CI) (months) [1	Mazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IVRS)									
NED [PDS]	92 47 (!	51.1) 18.0 (8	3.2, NE) 4	8 22 (45.8)	21.7 (4.2,	NE)	1.07	0.65,	1.81	0.7896
NED/CR [IDS]	74 49 (6	56.2) 4.2 (2	2.1,12.3) 3	8 17 (44.7)	22.1 (3.3,	NE)	1.79	1.05,	3.20	0.0307*
NED/CR [Chemo]	40 23 (57.5) 10.1 (2	2.3, NE) 2	0 12 (60.0)	9.6 (1.4,	NE)	1.02	0.52,	2.13	0.9470
PR	49 28 (57.1) 7.0 (4	1.2, NE) 2	5 14 (56.0)	11.3 (2.1,	NE)	1.00	0.53,	1.95	0.9917
Interaction p-value										0.4179
Screening laboratory tBR0	CA status (IVR	5)								
tBRCAm	150 90 (6	50.0) 8.7 (5	5.5,17.1) 6	5 32 (49.2)	11.3 (3.5,	NE)	1.23	0.83,	1.88	0.2977
non-tBRCAm	105 57 (54.3) 8.3 (3	3.4, NE) 6	6 33 (50.0)	16.7 (6.4,	NE)	1.17	0.77,	1.82	0.4603
Interaction p-value										0.8667
First line treatment out	come (eCRF)									
NED [PDS]	89 47 (!	52.8) 15.3 (5	7.7, NE) 4	7 21 (44.7)	NE (NE,	NE)	1.19	0.72,	2.02	0.5119
NED/CR [IDS]	74 49 (6	56.2) 4.2 (2	2.8, 8.6) 3	2 13 (40.6)	NE (NE,	NE)	2.11	1.18,	4.06	0.0105*
NED/CR [Chemo]	39 19 (4	18.7) 18.0 (2	2.3, NE) 1	7 12 (70.6)	5.7 (1.4,	NE)	0.68	0.34,	1.45	0.3092
PR	50 31 (52.0) 7.0 (2	2.3,22.1) 3	4 18 (52.9)	11.3 (2.1,	NE)	1.09	0.62,	1.99	0.7651
Interaction p-value										0.1172
Screening laboratory tBR0	CA status (eCRI	₹)								
tBRCAm	147 88 (59.9) 9.7 (5	5.2,18.0) 6	7 32 (47.8)	21.7 (4.2,	NE)	1.28	0.86,	1.95	0.2245
non-tBRCAm	108 59 (!	54.6) 8.3 (3	3.5, NE) 6	4 33 (51.6)	12.6 (4.2,	NE)	1.14	0.75,	1.76	0.5505
Interaction p-value										0.6960
Age group										
<65 years	185 101 (54.6) 10.1 (6	5.9,24.1) 9	8 46 (46.9)	21.7 (6.6,	NE)	1.18	0.84,	1.68	0.3543
>=65 years	70 46 (6	55.7) 3.5 (2	2.1,14.1) 3	3 19 (57.6)	8.3 (2.9,	NE)	1.30	0.77,	2.27	0.3259
Interaction p-value										0.7573

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaw 25NOV2020:11:06 kvbv306

Table 3.4.23 PAOLA1: Summary of subgroup analysis of AE max CTCAE grade>=3
Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		-	bevacizumab 255)			oevacizumab 131)					
Subgroup	Number (%) Med of patients (%) n with events (mon				Number (%) of patients with events	Median ti (95% CI) (months) [Hazard ratio [b]	95% C	I [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	110 (60.4)	8.6 (4.9,14.8)	89	40 (44.9)	NE (NE,	NE)	1.52	1.07,	2.20	0.0202*
IV	73	37 (50.7)	8.3 (4.2, NE)	42	25 (59.5)	6.6 (2.1,	NE)	0.74	0.45,	1.24	0.2514
Interaction p-value											0.0255*
Region											
Europe	245	141 (57.6)	8.6 (5.6,15.3)	125	63 (50.4)	12.6 (6.6,	NE)	1.19	0.89,	1.61	0.2450
Japan	10	6 (60.0)	4.9 (0.7, NE)	6	2 (33.3)	NE (NE,	NE)	1.93	0.45,	13.19	0.3970
Interaction p-value											0.5461
ECOG performance status at	Basel	ine									
(0) Normal activity	190	99 (52.1)	15.0 (6.9, NE)	100	50 (50.0)	16.7 (5.7,	NE)	1.06	0.76,	1.50	0.7381
(1) Restricted activity	61	44 (72.1)	4.3 (1.9, 8.4)	30	15 (50.0)	10.5 (1.4,	NE)	1.60	0.91,	2.98	0.1017
Interaction p-value											0.2232
Baseline CA-125 value											
<=ULN			8.7 (5.6,17.1)	117		16.7 (8.3,		1.25		1.72	0.1544
>ULN	27	17 (63.0)	6.2 (2.7,20.9)	14	8 (57.1)	2.4 (0.7,	NE)	0.92	0.41,	2.26	0.8518
Interaction p-value											0.5129
Histological grade											
High grade	255	147 (57.6)	8.6 (5.6,15.3)	131	65 (49.6)	16.7 (6.6,	NE)	1.21	0.91,	1.64	0.1886
Interaction p-value											NC
Cytoreductive surgery outco	me										
No residue	166	97 (58.4)	8.4 (4.8,15.3)	80	35 (43.8)	NE (NE,	NE)	1.47	1.01,	2.20	0.0437*
Residue	79	42 (53.2)	15.8 (4.2, NE)	43	24 (55.8)	10.5 (2.8,	NE)	0.94	0.57,	1.58	0.8130
Interaction p-value			<u> </u>								0.1681

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaw 25NOV2020:11:06 kvbv306

Table 3.4.23 PAOLA1: Summary of subgroup analysis of AE max CTCAE grade>=3
Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				bevacizumab 131)					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	of patients (95)	Hazard ratio [b] 95% CI		[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	78 (53.4)	15.0 (7.7, NE)	78	39 (50.0)	12.6 (6.4,	NE)	1.08	0.74,	1.60	0.6990
Interval	99	61 (61.6)	4.8 (3.5,14.1)	45	20 (44.4)	22.1 (4.2,	NE)	1.60	0.98,	2.71	0.0604
Interaction p-value											0.2230
Myriad tumour BRCA mutati	on stat	us									
tBRCAm	158	96 (60.8)	8.4 (4.8,15.3)	77	38 (49.4)	11.3 (4.2,	NE)	1.27	0.88,	1.87	0.2088
Non-tBRCAm	97	51 (52.6)	8.6 (3.9, NE)	54	27 (50.0)	16.7 (4.2,	NE)	1.12	0.71,	1.81	0.6312
Interaction p-value											0.6864
Status somatic BRCA mutat	ions										
sBRCAm	22	12 (54.5)	18.0 (5.2, NE)	7	2 (28.6)	NE (NE,	NE)	1.71	0.47,	11.02	0.4522
gBRCAm	66	45 (68.2)	5.5 (2.3, 8.7)	31	15 (48.4)	NE (NE,	NE)	1.69	0.97,	3.14	0.0667
Non-BRCAm	41	24 (58.5)	7.7 (2.8, NE)	22	11 (50.0)	16.7 (2.4,	NE)	1.27	0.64,	2.70	0.5041
Interaction p-value											0.8227

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaw 25NOV2020:11:06 kvbv306

Table 3.4.24 PAOLA1: Summary of subgroup analysis of AE G>=3 SOC: General disorders and administration site conditions Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2		ab		Placebo + b (N=1		nab	•			
Subgroup	C	Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVF	RS)										
NED [PDS]	92	4 (4.3)	NE (N	E, NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	6 (8.1)	NE (N	E, NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	3 (7.5)	NE (N	E, NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	2 (4.1)	NE (N	E, NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR	CA status	(IVRS)										
tBRCAm	150	9 (6.0)	NE (N	E, NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	6 (5.7)	NE (N	E, NE)	66	1 (1.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
First line treatment out	come (eCF	RF)										
NED [PDS]	89	4 (4.5)	NE (N	E, NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	5 (6.8)	NE (N	E, NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	3 (7.7)	NE (N	E, NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	3 (6.0)	NE (N	E, NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR	CA status	s (eCRF)										
tBRCAm	147	9 (6.1)	NE (N	E, NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	6 (5.6)	NE (N	E, NE)	64	1 (1.6)	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Age group												
<65 years	185	9 (4.9)	NE (N	E, NE)	98	1 (1.0)	NE (NE,	NE)	4.83	0.91, 89.00	0.0682
>=65 years	70	6 (8.6)	NE (N	E, NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aax 25NOV2020:11:06 kvbv306

Table 3.4.24 PAOLA1: Summary of subgroup analysis of AE G>=3 SOC: General disorders and administration site conditions Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b		ab		Placebo + be (N=1)					
Subgroup		Number (%) of patients with events	ients (95% CI) of patients (95% CI) ratio		95% CI [b]	2-sided p-value [b]					
FIGO Stage (Disease state)											
III	182	12 (6.6)	NE (N	E, NE)	89	1 (1.1)	NE (NE	, NE)	6.13	1.21,111.65	0.0250*
IV	73	3 (4.1)	NE (N	E, NE)	42	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	15 (6.1)	NE (N	E, NE)	125	1 (0.8)	NE (NE	, NE)	7.87	1.59,142.24	0.0069*
Japan	10	0	NE (N	E, NE)	6	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	8 (4.2)	NE (N	E, NE)	100	0	NE (NE	, NE)	NC	NC	NC
(1) Restricted activity	61	7 (11.5)	NE (N	E, NE)	30	1 (3.3)	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	14 (6.1)	NE (N	E, NE)	117	0	NE (NE	, NE)	NC	NC	NC
>ULN	27	1 (3.7)	NE (N	E, NE)	14	1 (7.1)	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	15 (5.9)	NE (N	E, NE)	131	1 (0.8)	NE (NE	, NE)	7.96	1.61,143.95	0.0065*
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	9 (5.4)	NE (N	E, NE)	80	1 (1.3)	NE (NE	, NE)	4.45	0.84, 82.09	0.0863
Residue	79	5 (6.3)	NE (N	E, NE)	43	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aax 25NOV2020:11:06 kvbv306

Table 3.4.24 PAOLA1: Summary of subgroup analysis of AE G>=3 SOC: General disorders and administration site conditions Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	(Olaparib + b (N=2!		Placebo + b (N=1				2-sided p-value [b]
Subgroup	C	Number (%) of patients vith events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	
Timing of cytoreductive s	surgery							
Upfront	146	7 (4.8)	NE (NE, NE)	78 1 (1.3)	NE (NE, NE)	NC	NC	NC
Interval	99	7 (7.1)	NE (NE, NE)	45 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC
Myriad tumour BRCA mutati	ion statu	ıs						
tBRCAm	158	9 (5.7)	NE (NE, NE)	77 0	NE (NE, NE)	NC	NC	NC
Non-tBRCAm	97	6 (6.2)	NE (NE, NE)	54 1 (1.9)	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC
Status somatic BRCA mutat	cions							
sBRCAm	22	1 (4.5)	NE (NE, NE)	7 0	NE (NE, NE)	NC	NC	NC
gBRCAm	66	7 (10.6)	NE (NE, NE)	31 0	NE (NE, NE)	NC	NC	NC
Non-BRCAm	41	2 (4.9)	NE (NE, NE)	22 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aax 25NOV2020:11:06 kvbv306

Table 3.4.25 PAOLA1: Summary of subgroup analysis of AE G>=3 PT: Fatigue Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	(Olaparib + b (N=2)				Placebo + be (N=13		mab				
Subgroup	C	Number (%) of patients with events	Median ti (95% CI (months))	C	Number (%) of patients with events		an ti % CI: hs))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)										
NED [PDS]	92	4 (4.3)	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	5 (6.8)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	3 (7.5)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	2 (4.1)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR0	CA status	(IVRS)										
tBRCAm	150	8 (5.3)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	6 (5.7)	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
First line treatment out	come (eCR	F)										
NED [PDS]	89	4 (4.5)	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	4 (5.4)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	3 (7.7)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	3 (6.0)	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR0	CA status	(eCRF)										
tBRCAm	147	8 (5.4)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	6 (5.6)	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Age group												
<65 years	185	8 (4.3)	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	6 (8.6)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aay 25NOV2020:11:06 kvbv306

Table 3.4.25 PAOLA1: Summary of subgroup analysis of AE G>=3 PT: Fatigue Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		ımab			Placebo + be (N=1)		mab	•			
Subgroup		Number (%) of patients with events	Medi (95 (mont	% CI)	C	Number (%) of patients with events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	12 (6.6)	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	2 (2.7)	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	14 (5.7)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Basel	ine											
(0) Normal activity	190	7 (3.7)	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	7 (11.5)	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	13 (5.7)	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	1 (3.7)	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	14 (5.5)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	8 (4.8)	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	5 (6.3)	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aay 25NOV2020:11:06 kvbv306

Table 3.4.25 PAOLA1: Summary of subgroup analysis of AE G>=3 PT: Fatigue Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		Placebo + b (N=1			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI	2-sided [b] p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	7 (4.8)	NE (NE, NE)	78 0	NE (NE, NE)	NC NC	NC
Interval	99	6 (6.1)	NE (NE, NE)	45 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Myriad tumour BRCA mutati	ion stati	ıs					
tBRCAm	158	8 (5.1)	NE (NE, NE)	77 0	NE (NE, NE)	NC NC	NC
Non-tBRCAm	97	6 (6.2)	NE (NE, NE)	54 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Status somatic BRCA mutat	ions						
sBRCAm	22	1 (4.5)	NE (NE, NE)	7 0	NE (NE, NE)	NC NC	NC
gBRCAm	66	6 (9.1)	NE (NE, NE)	31 0	NE (NE, NE)	NC NC	NC
Non-BRCAm	41	2 (4.9)	NE (NE, NE)	22 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aay 25NOV2020:11:06 kvbv306

Table 3.4.26 PAOLA1: Summary of subgroup analysis of AE G>=3 SOC: Blood and lymphatic system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olaparib + b (N=2		mab			Placebo + be (N=13		ımab				2-sided p-value [b]
Subgroup		Number (%) of patients with events	Media (95% (month	& CI)		C	Number (%) of patients with events	(95	an ti 5% CI ths))	Hazard ratio [b]	95% CI [b]	
First line treatment out	come (IV	RS)											
NED [PDS]	92	27 (29.3)	NE (NE,	NE)	48	4 (8.3)	NE (NE,	NE)	3.77	1.47, 12.75	0.0040*
NED/CR [IDS]	74	22 (29.7)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	10 (25.0)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	12 (24.5)	NE (NE,	NE)	25	1 (4.0)	NE (NE,	NE)	6.74	1.33,122.77	0.0171*
Interaction p-value													0.6032
Screening laboratory tBR0	CA statu	s (IVRS)											
tBRCAm	150	40 (26.7)	NE (NE,	NE)	65	2 (3.1)	NE (NE,	NE)	9.47	2.91, 58.20	<0.0001*
non-tBRCAm	105	31 (29.5)	NE (NE,	NE)	66	3 (4.5)	NE (NE,	NE)	7.57	2.70, 31.55	<0.0001*
Interaction p-value													0.8107
First line treatment out	come (eC	RF)											
NED [PDS]	89	28 (31.5)	NE (NE,	NE)	47	4 (8.5)	NE (NE,	NE)	4.00	1.57, 13.50	0.0023*
NED/CR [IDS]	74	24 (32.4)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	6 (15.4)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	12 (24.0)	NE (NE,	NE)	34	1 (2.9)	NE (NE,	NE)	8.79	1.73,160.30	0.0050*
Interaction p-value													0.4742
Screening laboratory tBR0	CA statu	s (eCRF)											
tBRCAm	147	40 (27.2)	NE (NE,	NE)	67	2 (3.0)	NE (NE,	NE)	9.99	3.07, 61.38	<0.0001*
non-tBRCAm	108	31 (28.7)	NE (NE,	NE)	64	3 (4.7)	NE (NE,	NE)	7.09	2.53, 29.53	<0.0001*
Interaction p-value													0.7133
Age group													
<65 years	185	47 (25.4)	NE (NE,	NE)	98	4 (4.1)	NE (NE,	NE)	6.77	2.75, 22.44	<0.0001*
>=65 years	70	24 (34.3)	NE (NE,	NE)	33	1 (3.0)	NE (NE,	NE)	13.80	2.92,246.76	<0.0001*
Interaction p-value													0.5092

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaz 25NOV2020:11:06 kvbv306

Table 3.4.26 PAOLA1: Summary of subgroup analysis of AE G>=3 SOC: Blood and lymphatic system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		-		Placebo + be (N=1)					
Subgroup		Number (%) of patients with events	ents (95% CI) of patients (95% CI) ratio		95% CI [b]	2-sided p-value [b]					
FIGO Stage (Disease state)											
III	182	55 (30.2)	NE (NE,	NE)	89	3 (3.4)	NE (NE,	NE)	10.37	3.83, 42.56	<0.0001*
IV	73	16 (21.9)	NE (NE,	NE)	42	2 (4.8)	NE (NE,	NE)	4.80	1.36, 30.33	0.0115*
Interaction p-value											0.4333
Region											
Europe	245	67 (27.3)	NE (NE,	NE)	125	3 (2.4)	NE (NE,	NE)	12.82	4.77, 52.42	<0.0001*
Japan	10	4 (40.0)	NE (NE,	NE)	6	2 (33.3)	NE (NE,	NE)	1.25	0.24, 9.03	0.7938
Interaction p-value											0.0406*
ECOG performance status at	Basel	ine									
(0) Normal activity	190	49 (25.8)	NE (NE,	NE)	100	3 (3.0)	NE (NE,	NE)	9.64	3.54, 39.70	<0.0001*
(1) Restricted activity	61	22 (36.1)	NE (NE,	NE)	30	2 (6.7)	NE (NE,	NE)	6.09	1.79, 38.03	0.0018*
Interaction p-value											0.6331
Baseline CA-125 value											
<=ULN	228	64 (28.1)	NE (NE,	NE)	117	5 (4.3)	NE (NE,	NE)	7.44	3.31, 21.27	<0.0001*
>ULN	27	7 (25.9)	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	71 (27.8)	NE (NE,	NE)	131	5 (3.8)	NE (NE,	NE)	8.17	3.65, 23.31	<0.0001*
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	53 (31.9)	NE (NE,	NE)	80	4 (5.0)	NE (NE,	NE)	7.39	3.03, 24.44	<0.0001*
Residue	79	13 (16.5)	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaz 25NOV2020:11:06 kvbv306

Table 3.4.26 PAOLA1: Summary of subgroup analysis of AE G>=3 SOC: Blood and lymphatic system disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		Placebo + b (N=1			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	39 (26.7)	NE (NE, NE)	78 4 (5.1)	NE (NE, NE)	5.58 2.25, 18.62	<0.0001*
Interval	99	27 (27.3)	NE (NE, NE)	45 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Myriad tumour BRCA mutat	ion stat	us					
tBRCAm	158	46 (29.1)	NE (NE, NE)	77 3 (3.9)	NE (NE, NE)	8.20 3.00, 33.78	<0.0001*
Non-tBRCAm	97	25 (25.8)	NE (NE, NE)	54 2 (3.7)	NE (NE, NE)	8.05 2.40, 50.05	0.0002*
Interaction p-value							0.9848
Status somatic BRCA mutat	cions						
sBRCAm	22	8 (36.4)	NE (NE, NE)	7 0	NE (NE, NE)	NC NC	NC
gBRCAm	66	17 (25.8)	NE (NE, NE)	31 2 (6.5)	NE (NE, NE)	4.73 1.35, 29.82	0.0117*
Non-BRCAm	41	14 (34.1)	NE (NE, NE)	22 1 (4.5)	NE (NE, NE)	8.92 1.79,161.67	0.0039*
Interaction p-value							0.6103

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aaz 25NOV2020:11:06 kvbv306

Table 3.4.27 PAOLA1: Summary of subgroup analysis of AE G>=3 PT: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olaparib + b (N=2		ımab			Placebo + be (N=13		ımab				
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI)	C	Number (%) of patients with events	(9	an ti 5% CI ths))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IV	RS)											
NED [PDS]	92	18 (19.6)	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	10.44	2.16,187.81	0.0011*
NED/CR [IDS]	74	14 (18.9)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	7 (17.5)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	8 (16.3)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA statu	s (IVRS)											
tBRCAm	150	23 (15.3)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	24 (22.9)	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	17.46	3.69,312.15	<0.0001*
Interaction p-value													NC
First line treatment outo	come (eCl	RF)											
NED [PDS]	89	19 (21.3)	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	11.35	2.36,203.99	0.0006*
NED/CR [IDS]	74	16 (21.6)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	5 (12.8)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	7 (14.0)	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA statu	s (eCRF)											
tBRCAm	147	23 (15.6)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	24 (22.2)	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	16.37	3.46,292.62	<0.0001*
Interaction p-value													NC
Age group													
<65 years	185	32 (17.3)	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	18.83	4.05,334.95	<0.0001*
>=65 years	70	15 (21.4)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aba 25NOV2020:11:06 kvbv306

Table 3.4.27 PAOLA1: Summary of subgroup analysis of AE G>=3 PT: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		ab		Placebo + be (N=13					
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]		(Number (%) of patients with events	Median t: (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	35 (19.2)	NE (N	IE, NE)	89	1 (1.1)	NE (NE,	NE)	19.08	4.13,339.02	<0.0001*
IV	73	12 (16.4)	NE (N	IE, NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	43 (17.6)	NE (N	IE, NE)	125	1 (0.8)	NE (NE,	NE)	23.96	5.23,424.77	<0.0001*
Japan	10	4 (40.0)	NE (N	IE, NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	33 (17.4)	NE (N	IE, NE)	100	1 (1.0)	NE (NE,	NE)	19.14	4.13,340.29	<0.0001*
(1) Restricted activity	61	14 (23.0)	NE (N	IE, NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	44 (19.3)	NE (N	IE, NE)	117	1 (0.9)	NE (NE,	NE)	25.57	5.59,453.20	<0.0001*
>ULN	27	3 (11.1)	NE (N	IE, NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	47 (18.4)	NE (N	IE, NE)	131	1 (0.8)	NE (NE,	NE)	26.71	5.85,473.03	<0.0001*
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	35 (21.1)	NE (N	IE, NE)	80	1 (1.3)	NE (NE,	NE)	19.54	4.23,347.11	<0.0001*
Residue	79	8 (10.1)	NE (N	IE, NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aba 25NOV2020:11:06 kvbv306

Table 3.4.27 PAOLA1: Summary of subgroup analysis of AE G>=3 PT: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2			oevacizumab 131)			
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery							
Upfront	146	26 (17.8)	NE (NE, NE)	78 1 (1.3)	NE (NE, NE)	15.29	3.25,272.91	<0.0001*
Interval	99	17 (17.2)	NE (NE, NE)	45 0	NE (NE, NE)	NC :	NC	NC
Interaction p-value								NC
Myriad tumour BRCA mutat:	ion stat	us						
tBRCAm	158	27 (17.1)	NE (NE, NE)	77 0	NE (NE, NE)	NC	NC	NC
Non-tBRCAm	97	20 (20.6)	NE (NE, NE)	54 1 (1.9)	NE (NE, NE)	12.75	2.66,228.74	0.0002*
Interaction p-value								NC
Status somatic BRCA muta	tions							
sBRCAm	22	3 (13.6)	NE (NE, NE)	7 0	NE (NE, NE)	NC :	NC	NC
gBRCAm	66	10 (15.2)	NE (NE, NE)	31 0	NE (NE, NE)	NC :	NC	NC
Non-BRCAm	41	10 (24.4)	NE (NE, NE)	22 0	NE (NE, NE)	NC :	NC	NC
Interaction p-value								NC

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3aba 25NOV2020:11:06 kvbv306

Table 3.4.28 PAOLA1: Summary of subgroup analysis of AE G>=3 SOC: Vascular disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olaparib + b (N=2		umab			Placebo + (N=	bevacizum 131)	ab	;			
Subgroup		Number (%) of patients with events		an ti % CI hs))		Number (%) of patients with events	(95%	n time (CI) (S) [a]	Hazard ratio [b]	95% CI	% CI [b]	2-sided p-value [b]
First line treatment outo	come (IV	RS)											
NED [PDS]	92	15 (16.3)	NE (NE,	NE)	48	11 (22.9)	NE (NE, NE	0.64	0.30,	1.44	0.2744
NED/CR [IDS]	74	13 (17.6)	NE (NE,	NE)	38	13 (34.2)	NE (NE, NE	0.45	0.21,	0.98	0.0447*
NED/CR [Chemo]	40	8 (20.0)	NE (NE,	NE)	20	7 (35.0)	NE (NE, NE	0.54	0.19,	1.53	0.2365
PR	49	12 (24.5)	NE (NE,	NE)	25	11 (44.0)	12.6 (2	2.1, NE	0.47	0.21,	1.09	0.0776
Interaction p-value													0.9229
Screening laboratory tBRO	CA statu	s (IVRS)											
tBRCAm	150	28 (18.7)	NE (NE,	NE)	65	23 (35.4)	NE (NE, NE	0.44	0.25,	0.77	0.0043*
non-tBRCAm	105	20 (19.0)	NE (NE,	NE)	66	19 (28.8)	NE (NE, NE	0.64	0.34,	1.21	0.1642
Interaction p-value													0.3782
First line treatment outo	come (eCl	RF)											
NED [PDS]	89	13 (14.6)	NE (NE,	NE)	47	10 (21.3)	NE (NE, NE	0.63	0.28,	1.47	0.2770
NED/CR [IDS]	74	12 (16.2)	NE (NE,	NE)	32	9 (28.1)	NE (NE, NE	0.55	0.23,	1.35	0.1833
NED/CR [Chemo]	39	8 (20.5)	NE (NE,	NE)	17	7 (41.2)	NE (NE, NE	0.49	0.18,	1.41	0.1789
PR	50	15 (30.0)	NE (NE,	NE)	34	15 (44.1)	12.6 (2	2.1, NE	0.52	0.25,	1.08	0.0805
Interaction p-value													0.9831
Screening laboratory tBRO	CA statu	s (eCRF)											
tBRCAm	147	26 (17.7)	NE (NE,	NE)	67	23 (34.3)	NE (NE, NE	0.43	0.24,	0.76	0.0038*
non-tBRCAm	108	22 (20.4)	NE (NE,	NE)	64	19 (29.7)	NE (NE, NE	0.67	0.36,	1.24	0.1979
Interaction p-value													0.2973
Age group													
<65 years	185	27 (14.6)	NE (NE,	NE)	98	29 (29.6)	NE (NE, NE	0.44	0.26,	0.74	0.0022*
>=65 years	70	21 (30.0)	NE (NE,	NE)	33	13 (39.4)	NE (NE, NE	0.68	0.35,	1.40	0.2899
Interaction p-value													0.3115

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3abb 25NOV2020:11:06 kvbv306

Table 3.4.28 PAOLA1: Summary of subgroup analysis of AE G>=3 SOC: Vascular disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		mab	,		Placebo + b						
Subgroup		Number (%) of patients with events	Media (95° (mont)	% CI)			Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	36 (19.8)	NE (NE,	NE)	89	25 (28.1)	NE (NE,	NE)	0.67	0.41,	1.13	0.1338
IV	73	12 (16.4)	NE (NE,	NE)	42	17 (40.5)	NE (NE,	NE)	0.31	0.14,	0.64	0.0017*
Interaction p-value													0.0861
Region													
Europe	245	48 (19.6)	NE (NE,	NE)	125	41 (32.8)	NE (NE,	NE)	0.53	0.35,	0.81	0.0033*
Japan	10	0	NE (NE,	NE)	6	1 (16.7)	NE (NE,	NE)	NC	NC		NC
Interaction p-value													NC
ECOG performance status at	Basel	ine											
(0) Normal activity	190	29 (15.3)	NE (NE,	NE)	100	32 (32.0)	NE (NE,	NE)	0.43	0.26,	0.72	0.0012*
(1) Restricted activity	61	17 (27.9)	NE (NE,	NE)	30	10 (33.3)	NE (NE,	NE)	0.72	0.34,	1.64	0.4272
Interaction p-value													0.2705
Baseline CA-125 value													
<=ULN	228	43 (18.9)	NE (NE,	NE)	117	35 (29.9)	NE (NE,	NE)	0.58	0.37,		0.0183*
>ULN	27	5 (18.5)	NE (NE,	NE)	14	7 (50.0)	4.3 (0.7,	NE)	0.25	0.07,	0.78	0.0175*
Interaction p-value													0.1738
Histological grade													
High grade	255	48 (18.8)	NE (NE,	NE)	131	42 (32.1)	NE (NE,	NE)	0.52	0.35,	0.80	0.0027*
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	25 (15.1)	NE (NE,	NE)	80	20 (25.0)	NE (NE,	NE)	0.56	0.31,	1.01	0.0557
Residue	79	16 (20.3)	NE (NE,	NE)	43	16 (37.2)	NE (NE,	NE)	0.48	0.24,	0.96	0.0391*
Interaction p-value													0.7391

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3abb 25NOV2020:11:06 kvbv306

Table 3.4.28 PAOLA1: Summary of subgroup analysis of AE G>=3 SOC: Vascular disorders Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		Placebo + b (N=1			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	23 (15.8)	NE (NE, NE)	78 21 (26.9)	NE (NE, NE)	0.54 0.30, 0.98	0.0432*
Interval	99	18 (18.2)	NE (NE, NE)	45 15 (33.3)	NE (NE, NE)	0.48 0.24, 0.96	0.0395*
Interaction p-value							0.8002
Myriad tumour BRCA mutat:	ion stat	us					
tBRCAm	158	30 (19.0)	NE (NE, NE)	77 25 (32.5)	NE (NE, NE)	0.49 0.29, 0.84	0.0105*
Non-tBRCAm	97	18 (18.6)	NE (NE, NE)	54 17 (31.5)	NE (NE, NE)	0.57 0.29, 1.12	0.1037
Interaction p-value							0.7244
Status somatic BRCA muta	cions						
sBRCAm	22	4 (18.2)	NE (NE, NE)	7 2 (28.6)	NE (NE, NE)	0.49 0.10, 3.57	0.4383
gBRCAm	66	13 (19.7)	NE (NE, NE)	31 11 (35.5)	NE (NE, NE)	0.50 0.22, 1.13	0.0930
Non-BRCAm	41	11 (26.8)	NE (NE, NE)	22 8 (36.4)	NE (NE, NE)	0.72 0.29, 1.85	0.4761
Interaction p-value							0.8257

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3abb 25NOV2020:11:06 kvbv306

Table 3.4.29 PAOLA1: Summary of subgroup analysis of AE G>=3 PT: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		umab			Placebo + (N=	bevacizuma :131)	lb	,			
Subgroup		Number (%) of patients with events		an ti % CI .hs))		Number (%) of patients with events		CI)	Hazard ratio [b]	95% C]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IV	RS)											
NED [PDS]	92	14 (15.2)	NE (NE,	NE)	48	11 (22.9)	NE (NE, NE)	0.59	0.27,	1.33	0.2006
NED/CR [IDS]	74	12 (16.2)	NE (NE,	NE)	38	13 (34.2)	NE (I	NE, NE)	0.41	0.19,	0.91	0.0288*
NED/CR [Chemo]	40	8 (20.0)	NE (NE,	NE)	20	7 (35.0)	NE (I	NE, NE)	0.53	0.19,	1.53	0.2335
PR	49	11 (22.4)	NE (NE,	NE)	25	11 (44.0)	12.6 (2	1, NE)	0.43	0.18,	0.996	0.0490*
Interaction p-value													0.9116
Screening laboratory tBRO	CA statu	s (IVRS)											
tBRCAm	150	27 (18.0)	NE (NE,	NE)	65	23 (35.4)	NE (NE, NE)	0.42	0.24,	0.73	0.0027*
non-tBRCAm	105	18 (17.1)	NE (NE,	NE)	66	19 (28.8)	NE (NE, NE)	0.57	0.30,	1.09	0.0887
Interaction p-value													0.4743
First line treatment outo	come (eC	RF)											
NED [PDS]	89	13 (14.6)	NE (NE,	NE)	47	10 (21.3)	NE (NE, NE)	0.63	0.28,	1.47	0.2759
NED/CR [IDS]	74	11 (14.9)	NE (NE,	NE)	32	9 (28.1)	NE (NE, NE)	0.50	0.21,	1.24	0.1322
NED/CR [Chemo]	39	8 (20.5)	NE (NE,	NE)	17	7 (41.2)	NE (NE, NE)	0.49	0.18,	1.40	0.1774
PR	50	13 (26.0)	NE (NE,	NE)	34	15 (44.1)	12.6 (2	1, NE)	0.44	0.20,	0.92	0.0291*
Interaction p-value													0.9344
Screening laboratory tBRO	CA statu	s (eCRF)											
tBRCAm	147	25 (17.0)	NE (NE,	NE)	67	23 (34.3)	NE (NE, NE)	0.41	0.23,	0.72	0.0023*
non-tBRCAm	108	20 (18.5)	NE (NE,	NE)	64	19 (29.7)	NE (NE, NE)	0.60	0.32,	1.13	0.1141
Interaction p-value													0.3654
Age group													
<65 years	185	25 (13.5)	NE (NE,	NE)	98	29 (29.6)	NE (NE, NE)	0.40	0.23,	0.68	0.0009*
>=65 years	70	20 (28.6)	NE (NE,	NE)	33	13 (39.4)	NE (NE, NE)	0.64	0.32,	1.33	0.2265
Interaction p-value													0.2876

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3abc 25NOV2020:11:06 kvbv306

Table 3.4.29 PAOLA1: Summary of subgroup analysis of AE G>=3 PT: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		mab	,		Placebo + b						
Subgroup		Number (%) of patients with events	Media (95° (mont)	% CI)			Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	33 (18.1)	NE (NE,	NE)	89	25 (28.1)	NE (NE,	NE)	0.61	0.36,	1.03	0.0662
IV	73	12 (16.4)	NE (NE,	NE)	42	17 (40.5)	NE (NE,	NE)	0.31	0.14,	0.64	0.0016*
Interaction p-value													0.1317
Region													
Europe	245	45 (18.4)	NE (NE,	NE)	125	41 (32.8)	NE (NE,	NE)	0.49	0.32,	0.75	0.0012*
Japan	10	0	NE (NE,	NE)	6	1 (16.7)	NE (NE,	NE)	NC	NC		NC
Interaction p-value													NC
ECOG performance status at	Basel	ine											
(0) Normal activity	190	27 (14.2)	NE (NE,	NE)	100	32 (32.0)	NE (NE,	NE)	0.40	0.24,	0.66	0.0005*
(1) Restricted activity	61	16 (26.2)	NE (NE,	NE)	30	10 (33.3)	NE (NE,	NE)	0.67	0.31,	1.54	0.3363
Interaction p-value													0.2696
Baseline CA-125 value													
<=ULN	228	40 (17.5)	NE (NE,	NE)	117	35 (29.9)	NE (NE,	NE)	0.53	0.34,	0.84	0.0075*
>ULN	27	5 (18.5)	NE (NE,	NE)	14	7 (50.0)	4.3 (0.7,	NE)	0.24	0.07,	0.77	0.0166*
Interaction p-value													0.2140
Histological grade													
High grade	255	45 (17.6)	NE (NE,	NE)	131	42 (32.1)	NE (NE,	NE)	0.49	0.32,	0.74	0.0010*
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	24 (14.5)	NE (NE,	NE)	80	20 (25.0)	NE (NE,	NE)	0.53	0.29,	0.98	0.0415*
Residue	79	15 (19.0)	NE (NE,	NE)	43	16 (37.2)	NE (NE,	NE)	0.44	0.21,	0.89	0.0233*
Interaction p-value													0.6755

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3abc 25NOV2020:11:06 kvbv306

Table 3.4.29 PAOLA1: Summary of subgroup analysis of AE G>=3 PT: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + b (N=2		Placebo + b (N=1			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	23 (15.8)	NE (NE, NE)	78 21 (26.9)	NE (NE, NE)	0.54 0.30, 0.98	0.0425*
Interval	99	16 (16.2)	NE (NE, NE)	45 15 (33.3)	NE (NE, NE)	0.42 0.21, 0.85	0.0174*
Interaction p-value							0.5926
Myriad tumour BRCA mutati	on stat	us					
tBRCAm	158	29 (18.4)	NE (NE, NE)	77 25 (32.5)	NE (NE, NE)	0.47 0.28, 0.81	0.0069*
Non-tBRCAm	97	16 (16.5)	NE (NE, NE)	54 17 (31.5)	NE (NE, NE)	0.51 0.25, 1.01	0.0517
Interaction p-value							0.8708
Status somatic BRCA mutat	ions						
sBRCAm	22	4 (18.2)	NE (NE, NE)	7 2 (28.6)	NE (NE, NE)	0.49 0.10, 3.55	0.4339
gBRCAm	66	12 (18.2)	NE (NE, NE)	31 11 (35.5)	NE (NE, NE)	0.45 0.20, 1.03	0.0590
Non-BRCAm	41	10 (24.4)	NE (NE, NE)	22 8 (36.4)	NE (NE, NE)	0.65 0.25, 1.69	0.3613
Interaction p-value							0.8440

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3abc 25NOV2020:11:06 kvbv306

Table 3.4.30 PAOLA1: Summary of subgroup analysis of AE max CTCAE grade=1 or 2 Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	-	bevacizumab 255)			Placebo + b						
Subgroup	(Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	l 95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IVI	RS)										
NED [PDS]	92	45 (48.9)	NE (NE,	NE)	48	25 (52.1)	7.0 (0.7,	NE)	0.97	0.60,	1.61	0.9150
NED/CR [IDS]	74	25 (33.8)	NE (NE,	NE)	38	20 (52.6)	4.8 (1.4,	NE)	0.63	0.35,	1.14	0.1219
NED/CR [Chemo]	40	17 (42.5)	NE (NE,	NE)	20	7 (35.0)	NE (NE,	NE)	1.40	0.61,	3.63	0.4398
PR	49	21 (42.9)	NE (NE,	NE)	25	10 (40.0)	NE (NE,	NE)	1.03	0.50,	2.29	0.9314
Interaction p-value												0.4389
Screening laboratory tBR0	CA status	s (IVRS)										
tBRCAm	150	60 (40.0)	NE (NE,	NE)	65	31 (47.7)	NE (NE,	NE)	0.81	0.53,	1.27	0.3598
non-tBRCAm	105	48 (45.7)	NE (NE,	NE)	66	31 (47.0)	NE (NE,	NE)	1.06	0.68,	1.67	0.8118
Interaction p-value												0.4168
First line treatment out	come (eCI	RF)										
NED [PDS]	89	42 (47.2)	NE (NE,	NE)	47	25 (53.2)	7.0 (1.1,	NE)	0.91	0.56,	1.52	0.7222
NED/CR [IDS]	74	25 (33.8)	NE (NE,	NE)	32	18 (56.3)	4.1 (1.0,	NE)	0.59	0.33,	1.11	0.0989
NED/CR [Chemo]	39	20 (51.3)	14.7 (0.5,	NE)	17	4 (23.5)	NE (NE,	NE)	2.75	1.04,	9.46	0.0406*
PR	50	19 (38.0)	NE (NE,	NE)	34	15 (44.1)	NE (NE,	NE)	0.78	0.40,	1.56	0.4771
Interaction p-value												0.0680
Screening laboratory tBR0	CA status	s (eCRF)										
tBRCAm	147	59 (40.1)	NE (NE,	NE)	67	33 (49.3)	NE (NE,	NE)	0.79	0.52,	1.22	0.2729
non-tBRCAm	108	49 (45.4)	NE (NE,	NE)	64	29 (45.3)	NE (NE,	NE)	1.10	0.70,	1.75	0.6957
Interaction p-value												0.2980
Age group												
<65 years	185	84 (45.4)	NE (NE,	NE)	98	48 (49.0)	NE (NE,	NE)	0.98	0.69,	1.41	0.9112
>=65 years	70	24 (34.3)	NE (NE,	NE)	33	14 (42.4)	NE (NE,	NE)	0.76	0.40,	1.50	0.4176
Interaction p-value												0.5054

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3abd 25NOV2020:11:06 kvbv306

Table 3.4.30 PAOLA1: Summary of subgroup analysis of AE max CTCAE grade=1 or 2 Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2		umab	•		Placebo + b						
Subgroup		Number (%) of patients with events	(95	an ti % CI .hs))		Number (%) of patients with events	Median tim (95% CI) (months) [a	-	Hazard ratio [b] 95% CI [[b]	2-sided b] p-value[b]	
FIGO Stage (Disease state)													
III	182	72 (39.6)	NE (NE,	NE)	89	47 (52.8)	7.0 (1.4,	NE)	0.71	0.50,	1.04	0.0776
IV	73	36 (49.3)	NE (NE,	NE)	42	15 (35.7)	NE (NE,	NE)	1.63	0.91,	3.06	0.1024
Interaction p-value													0.0195*
Region													
Europe	245	104 (42.4)	NE (NE,	NE)	125	58 (46.4)	NE (NE,	NE)	0.94	0.68,	1.30	0.7030
Japan	10	4 (40.0)	NE (NE,	NE)	6	4 (66.7)	1.1 (0.3,	NE)	0.54	0.13,	2.30	0.3929
Interaction p-value													0.4542
ECOG performance status at	Basel	ine											
(0) Normal activity	190	91 (47.9)	NE (NE,	NE)	100	46 (46.0)	NE (NE,	NE)	1.14	0.80,	1.63	0.4739
(1) Restricted activity	61	17 (27.9)	NE (NE,	NE)	30	15 (50.0)	NE (NE,	NE)	0.47	0.24,	0.96	0.0384*
Interaction p-value													0.0295*
Baseline CA-125 value													
<=ULN	228	98 (43.0)	NE (NE,	NE)	117	56 (47.9)	NE (NE,	NE)	0.91	0.66,		0.5835
>ULN	27	10 (37.0)	NE (NE,	NE)	14	6 (42.9)	NE (NE,	NE)	0.92	0.34,	2.71	0.8759
Interaction p-value													0.9834
Histological grade													
High grade	255	108 (42.4)	NE (NE,	NE)	131	62 (47.3)	NE (NE,	NE)	0.91	0.67,	1.26	0.5739
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	69 (41.6)	NE (NE,	NE)	80	43 (53.8)	4.8 (1.4,	NE)	0.79	0.54,	1.16	0.2201
Residue	79	37 (46.8)	NE (NE,	NE)	43	17 (39.5)	NE (NE,	NE)	1.24	0.71,	2.25	0.4658
Interaction p-value													0.1942

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. *p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3abd 25NOV2020:11:06 kvbv306

Table 3.4.30 PAOLA1: Summary of subgroup analysis of AE max CTCAE grade=1 or 2 Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olaparib + b			pevacizumab 131)		
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	68 (46.6)	NE (NE, NE)	78 37 (47.4)	NE (NE, NE)	1.02 0.69, 1.54	0.9129
Interval	99	38 (38.4)	NE (NE, NE)	45 23 (51.1)	8.6 (1.4, NE)	0.75 0.45, 1.28	0.2861
Interaction p-value							0.3582
Myriad tumour BRCA mutati	ion stat	us					
tBRCAm	158	62 (39.2)	NE (NE, NE)	77 37 (48.1)	NE (NE, NE)	0.80 0.54, 1.22	0.2955
Non-tBRCAm	97	46 (47.4)	NE (NE, NE)	54 25 (46.3)	NE (NE, NE)	1.11 0.69, 1.84	0.6672
Interaction p-value							0.3132
Status somatic BRCA mutat	cions						
sBRCAm	22	10 (45.5)	NE (NE, NE)	7 5 (71.4)	2.4 (0.1, NE)	0.53 0.19, 1.72	0.2731
gBRCAm	66	21 (31.8)	NE (NE, NE)	31 16 (51.6)	8.6 (0.6, NE)	0.56 0.29, 1.09	0.0879
Non-BRCAm	41	17 (41.5)	NE (NE, NE)	22 11 (50.0)	4.8 (0.7, NE)	0.84 0.40, 1.85	0.6514
Interaction p-value							0.6852

The time to event endpoint is the time to first AE or the time to censoring if the AE has not occurred by 30 days following the date of last dose of olaparib/placebo. Includes AEs with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 22.0. NC = not calculable.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubae_v3.sas ettesubae_v3abd 25NOV2020:11:06 kvbv306

Figure 3.5.1 PAOLA1: Kaplan-Meier plot of AE for Age group=<65 years Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

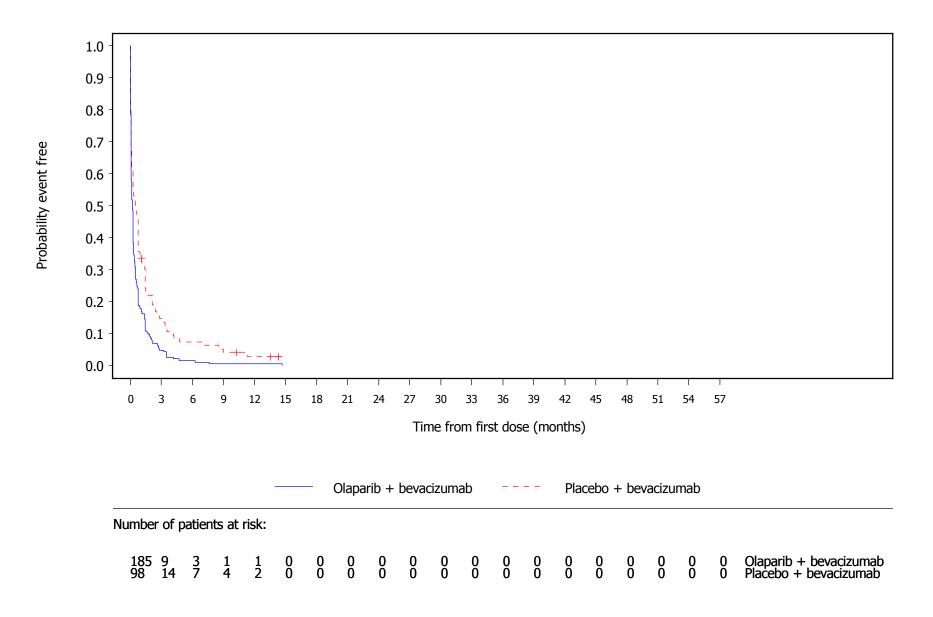


Figure 3.5.2 PAOLA1: Kaplan-Meier plot of AE for Age group=>=65 years Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

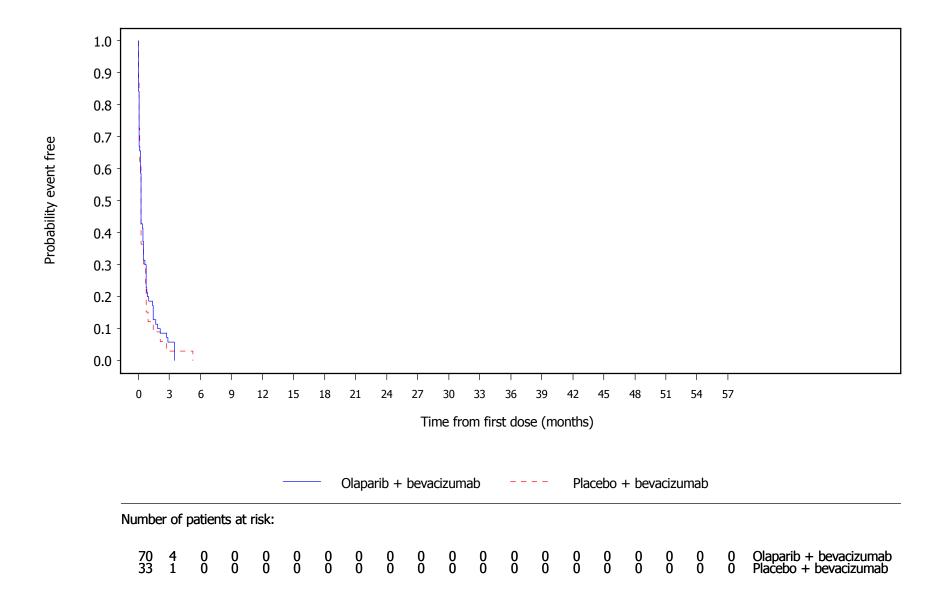


Figure 3.5.3 PAOLA1: Kaplan-Meier plot of AE PT: Proteinuria for Timing of cytoreductive surgery=Upfront Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

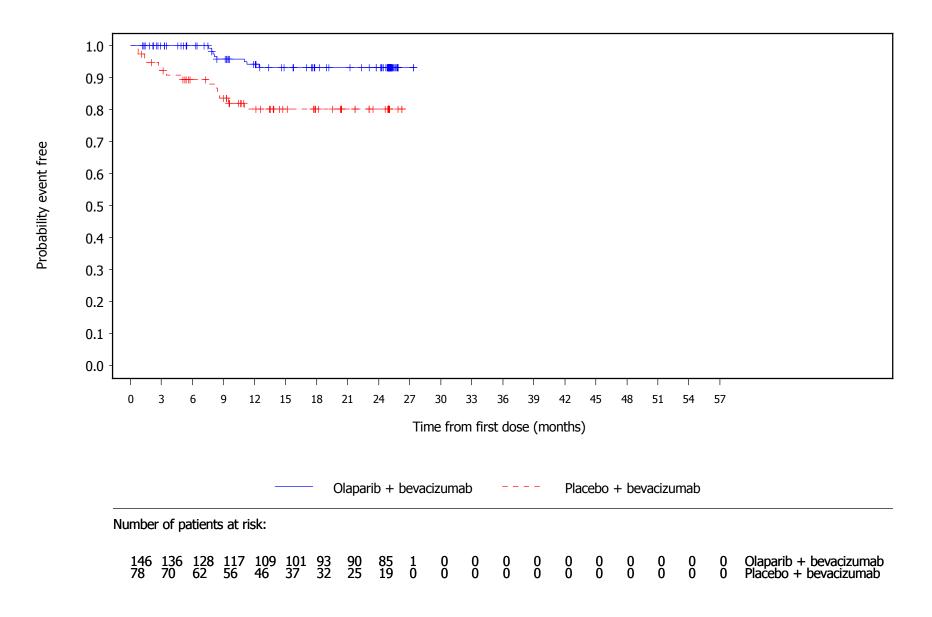


Figure 3.5.4 PAOLA1: Kaplan-Meier plot of AE PT: Proteinuria for Timing of cytoreductive surgery=Interval Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

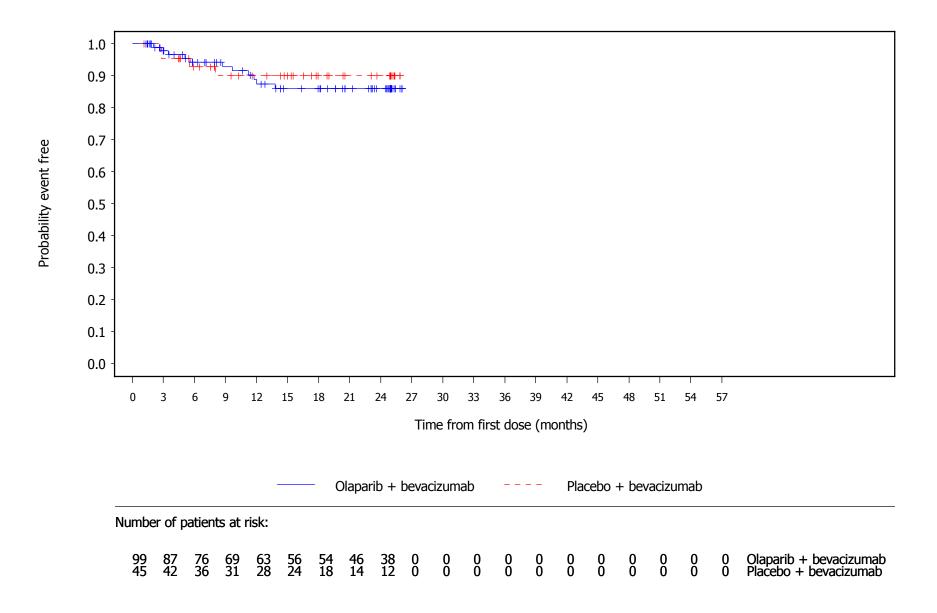


Figure 3.5.5 PAOLA1: Kaplan-Meier plot of AE PT: Lymphopenia for FIGO Stage (Disease state)=III Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

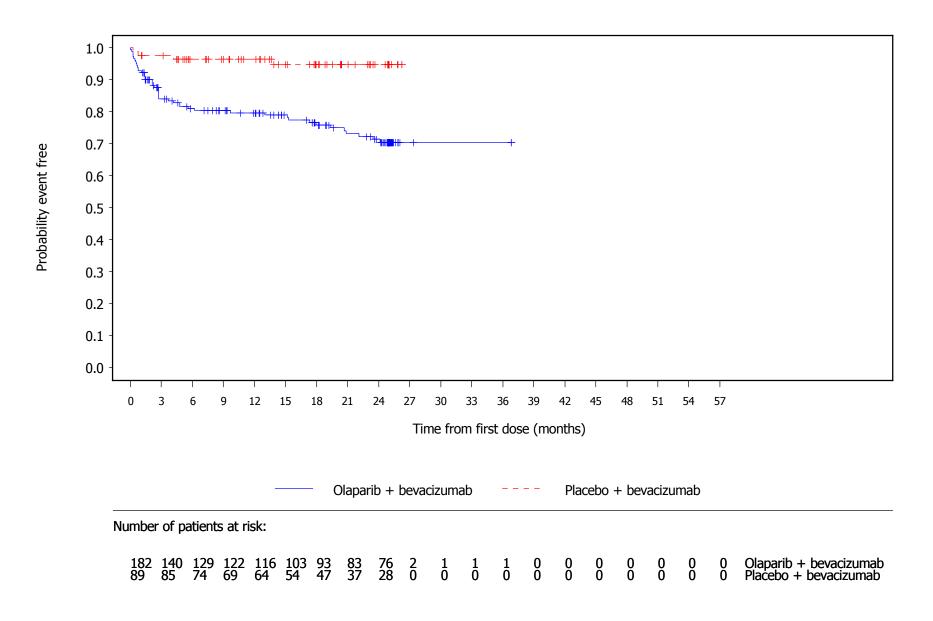


Figure 3.5.6 PAOLA1: Kaplan-Meier plot of AE PT: Lymphopenia for FIGO Stage (Disease state)=IV Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

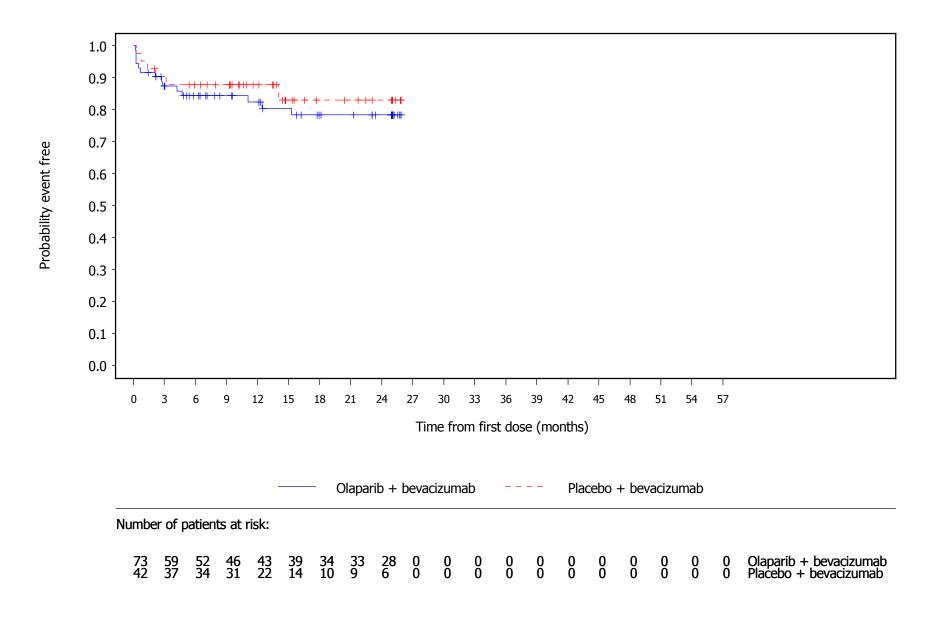


Figure 3.5.7 PAOLA1: Kaplan-Meier plot of AE PT: Lymphopenia for Region=Europe Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

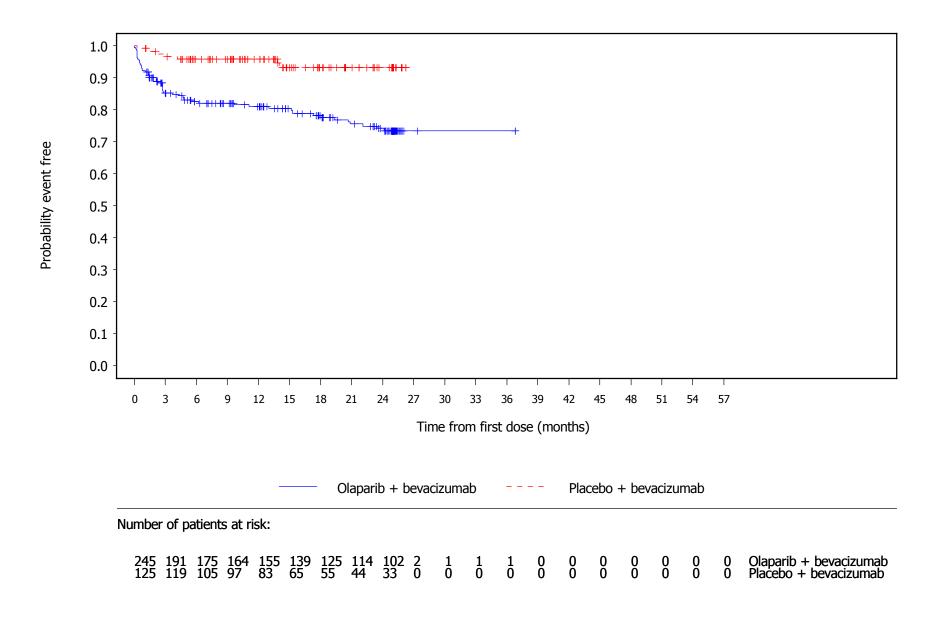


Figure 3.5.8 PAOLA1: Kaplan-Meier plot of AE PT: Lymphopenia for Region=Japan Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

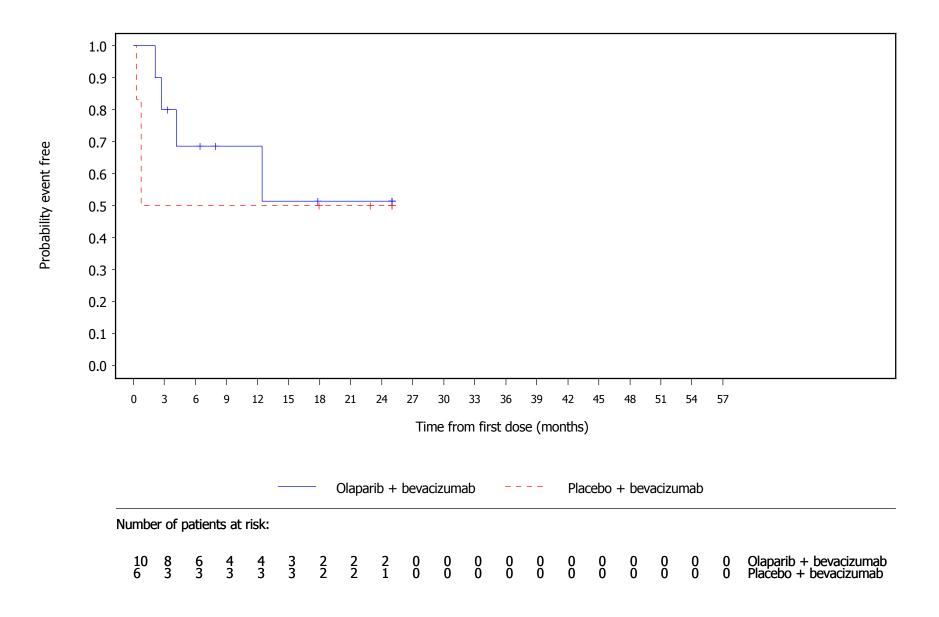


Figure 3.5.9 PAOLA1: Kaplan-Meier plot of AE PT: Lymphopenia for Cytoreductive surgery outcome=No residue Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

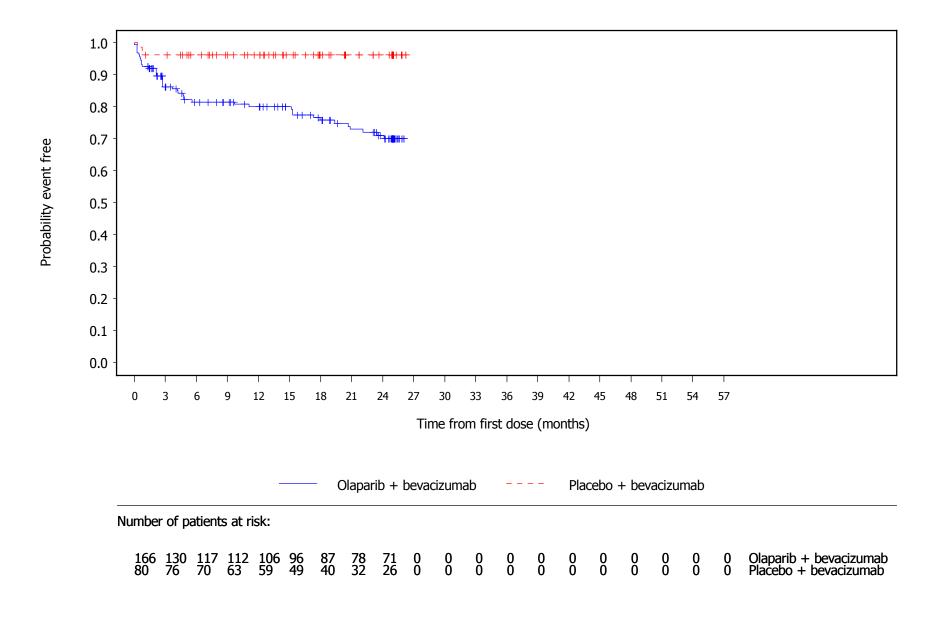


Figure 3.5.10 PAOLA1: Kaplan-Meier plot of AE PT: Lymphopenia for Cytoreductive surgery outcome=Residue Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

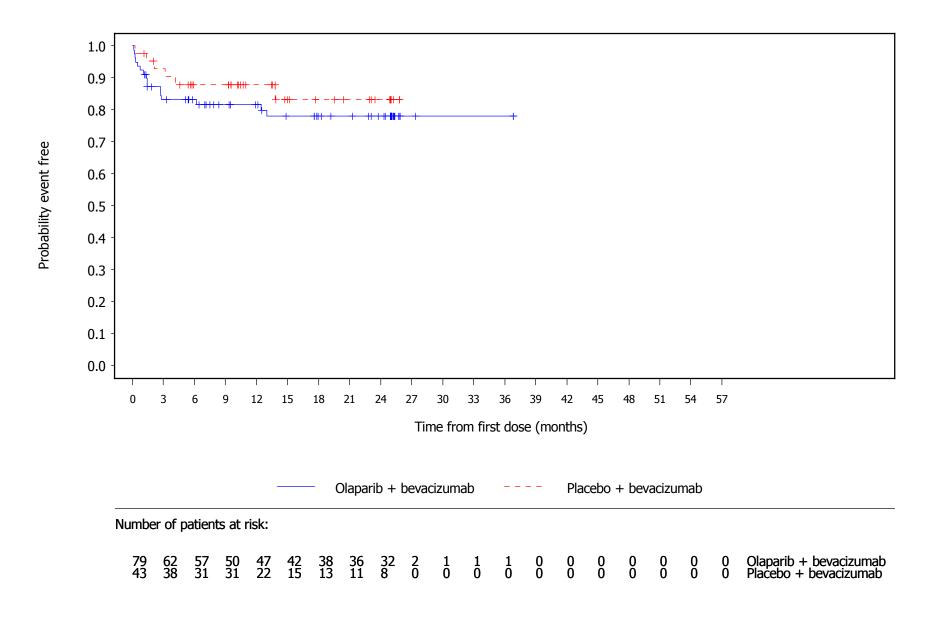


Figure 3.5.11 PAOLA1: Kaplan-Meier plot of AE PT: Nausea for FIGO Stage (Disease state)=III Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

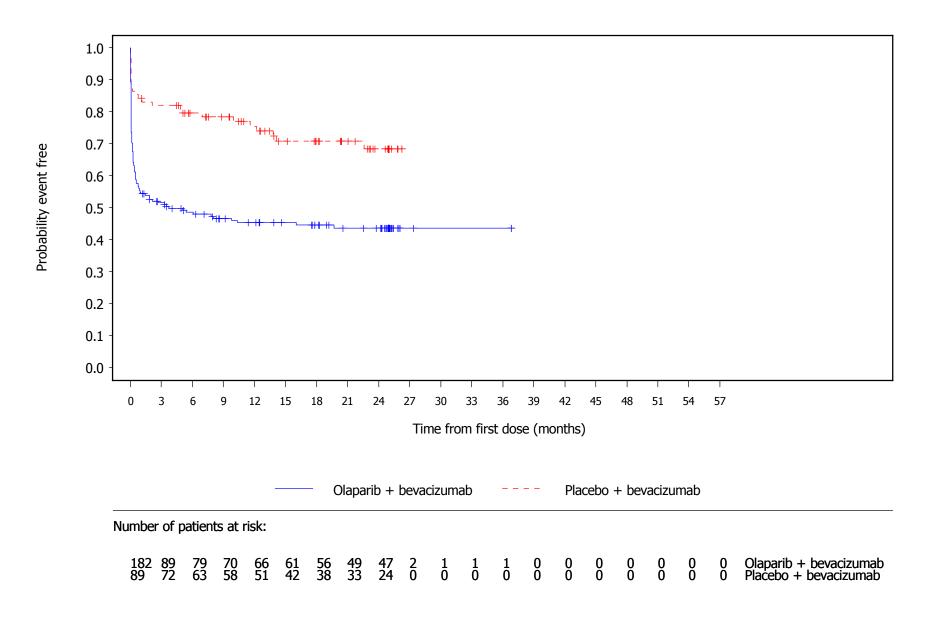


Figure 3.5.12 PAOLA1: Kaplan-Meier plot of AE PT: Nausea for FIGO Stage (Disease state)=IV Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

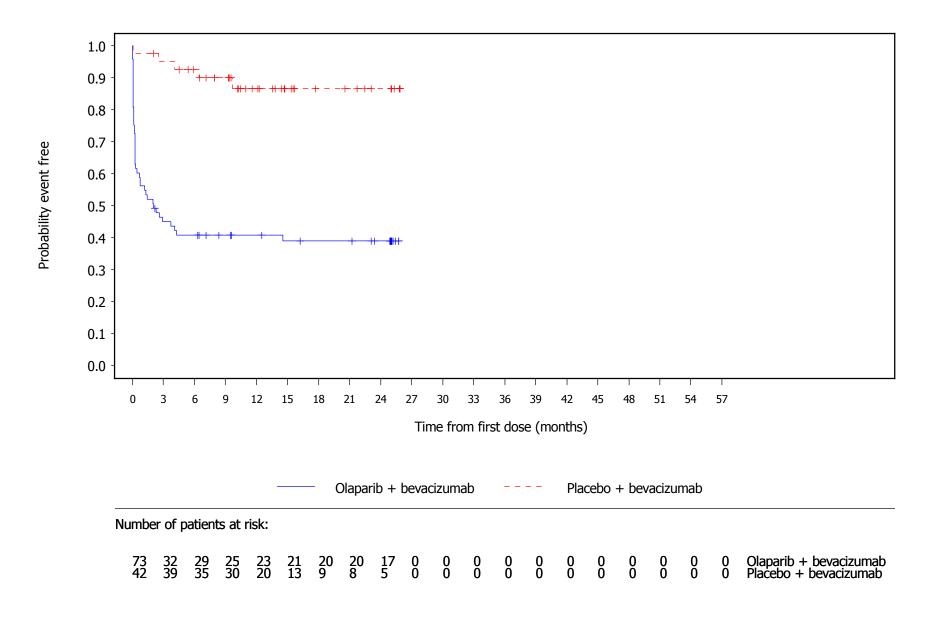


Figure 3.5.13 PAOLA1: Kaplan-Meier plot of AE PT: Nausea for ECOG performance status at Baseline=(0) Normal activity Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

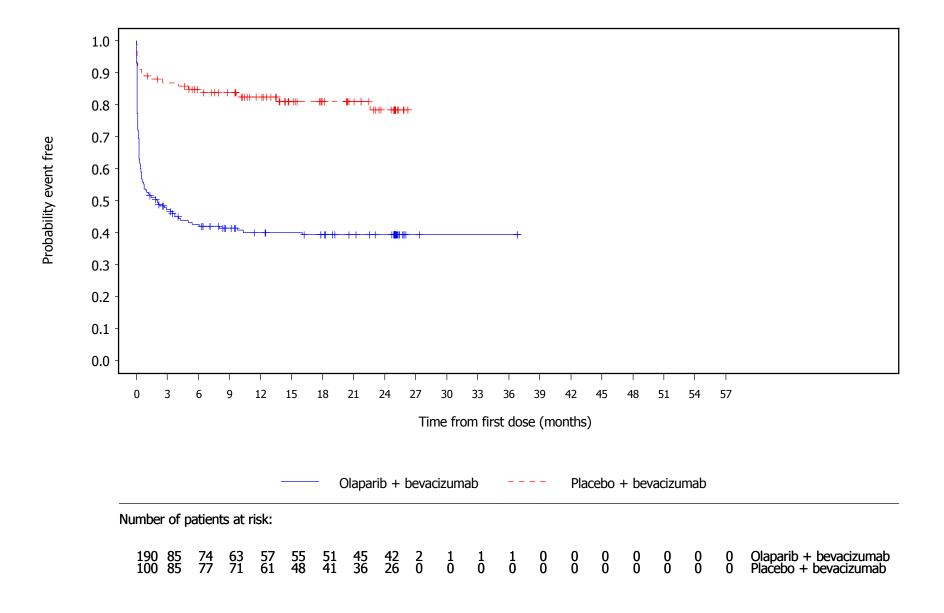


Figure 3.5.14 PAOLA1: Kaplan-Meier plot of AE PT: Nausea for ECOG performance status at Baseline=(1) Restricted activity Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

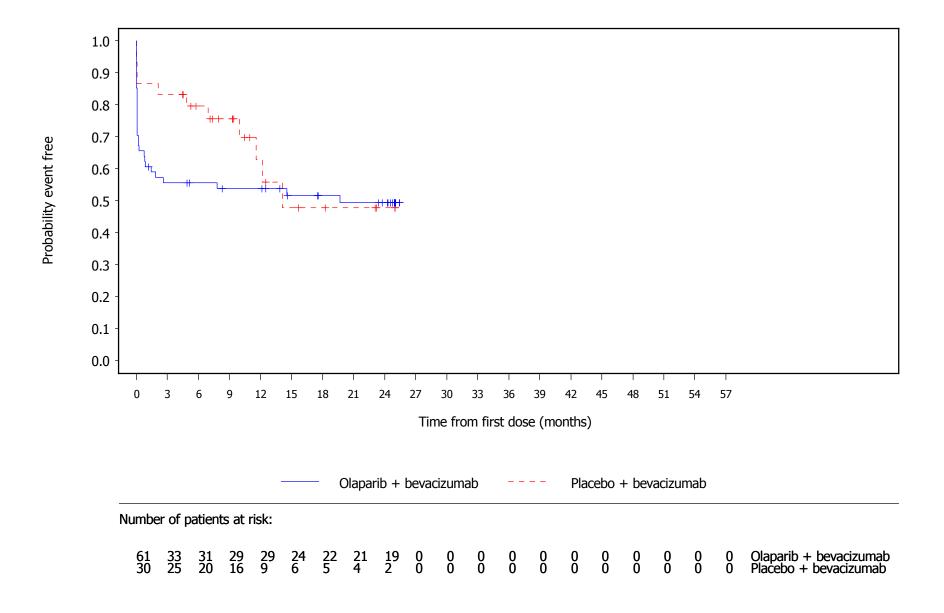


Figure 3.5.15 PAOLA1: Kaplan-Meier plot of AE PT: Hypertension for Region=Europe Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

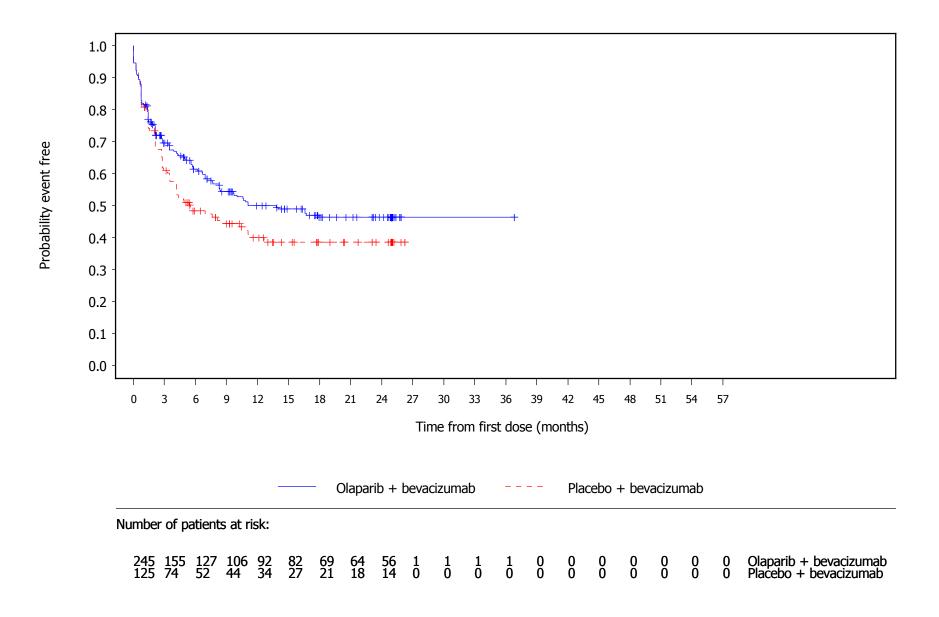


Figure 3.5.16 PAOLA1: Kaplan-Meier plot of AE PT: Hypertension for Region=Japan Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

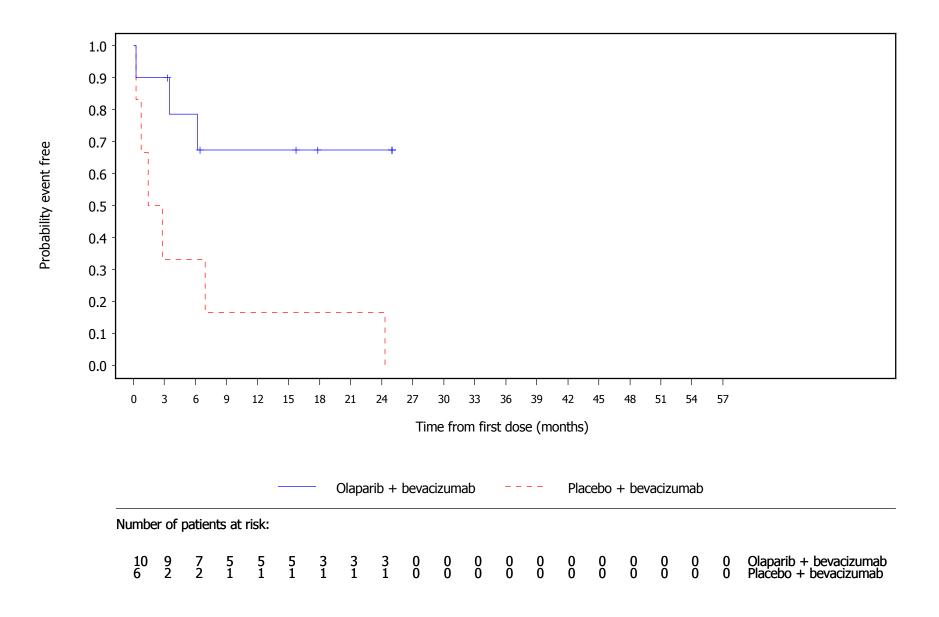


Figure 3.5.17 PAOLA1: Kaplan-Meier plot of AE max CTCAE grade>=3 for FIGO Stage (Disease state)=III Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

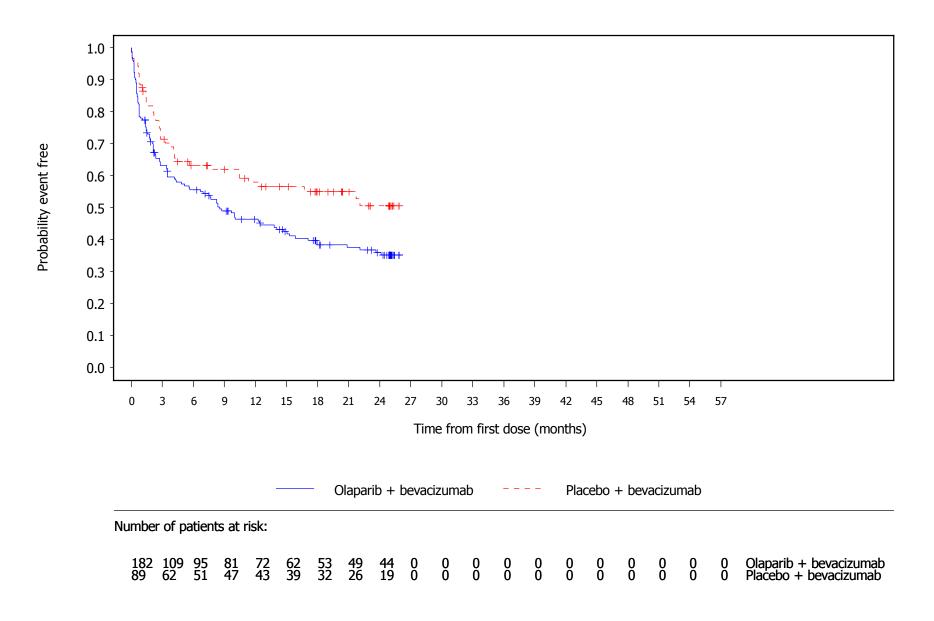


Figure 3.5.18 PAOLA1: Kaplan-Meier plot of AE max CTCAE grade>=3 for FIGO Stage (Disease state)=IV Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

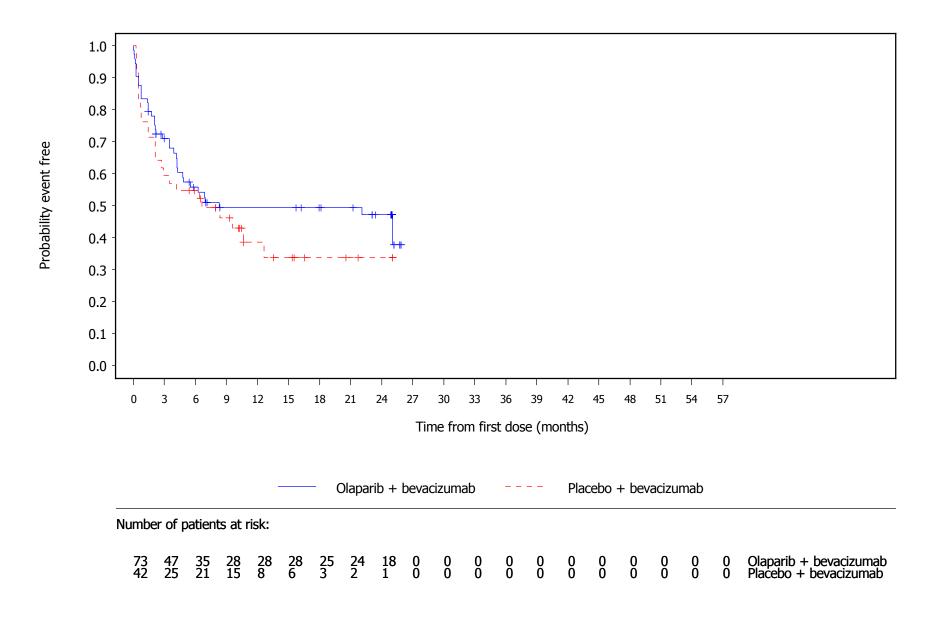


Figure 3.5.19 PAOLA1: Kaplan-Meier plot of AE G>=3 SOC: Blood and lymphatic system disorders for Region=Europe Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

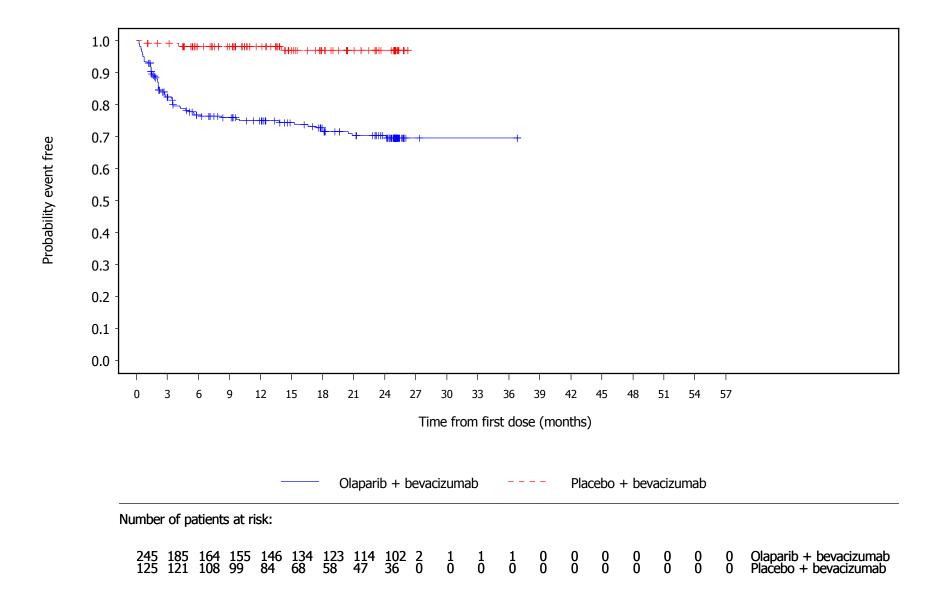


Figure 3.5.20 PAOLA1: Kaplan-Meier plot of AE G>=3 SOC: Blood and lymphatic system disorders for Region=Japan Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

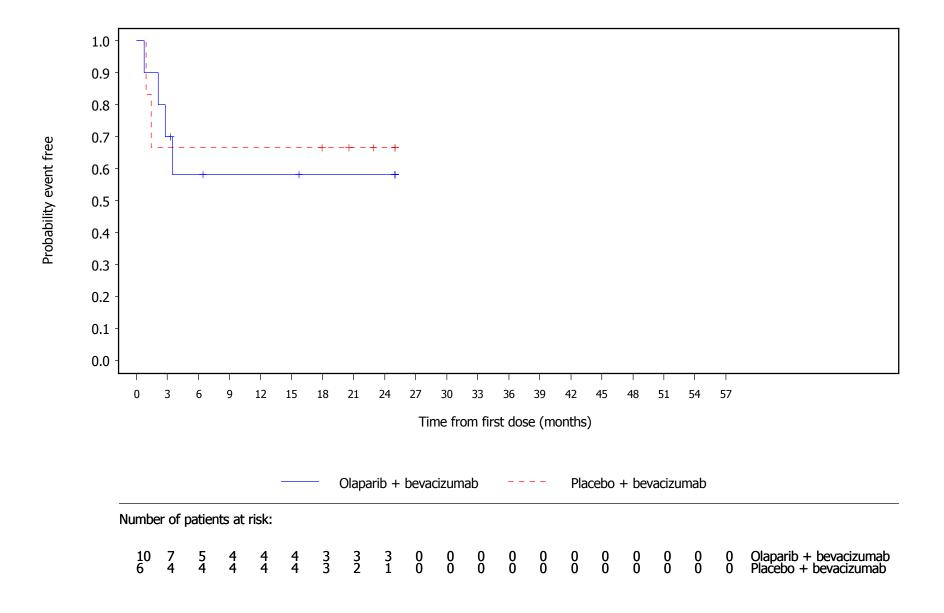


Figure 3.5.21 PAOLA1: Kaplan-Meier plot of AE max CTCAE grade=1 or 2 for FIGO Stage (Disease state)=III Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

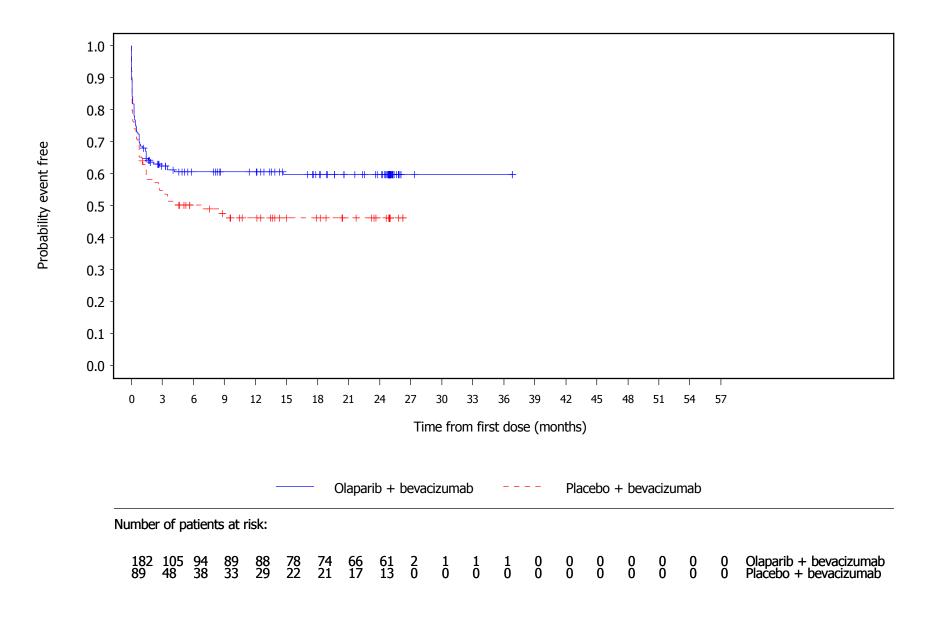


Figure 3.5.22 PAOLA1: Kaplan-Meier plot of AE max CTCAE grade=1 or 2 for FIGO Stage (Disease state)=IV Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

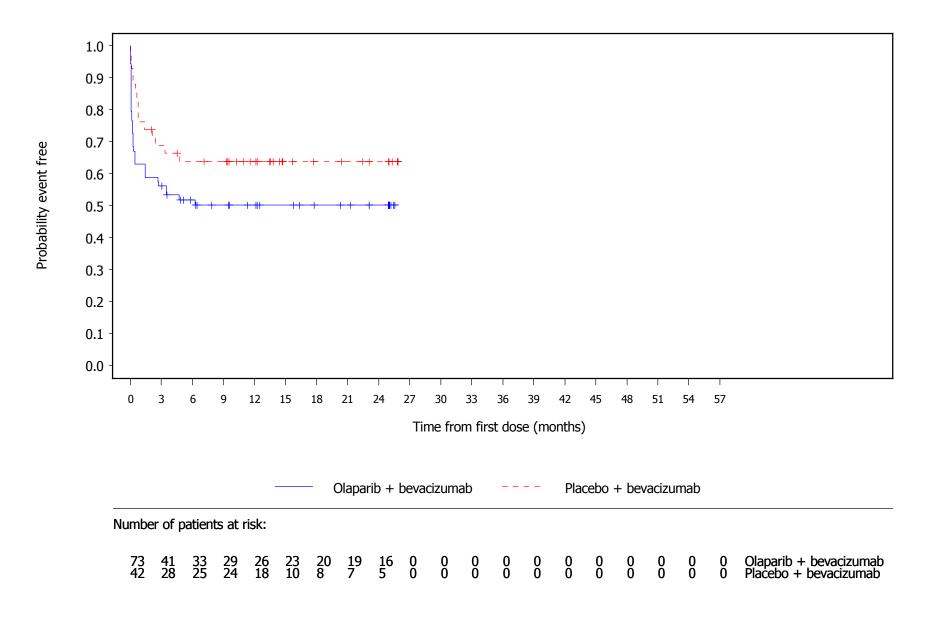


Figure 3.5.23 PAOLA1: Kaplan-Meier plot of AE max CTCAE grade=1 or 2 for ECOG performance status at Baseline=(0) Normal activity Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

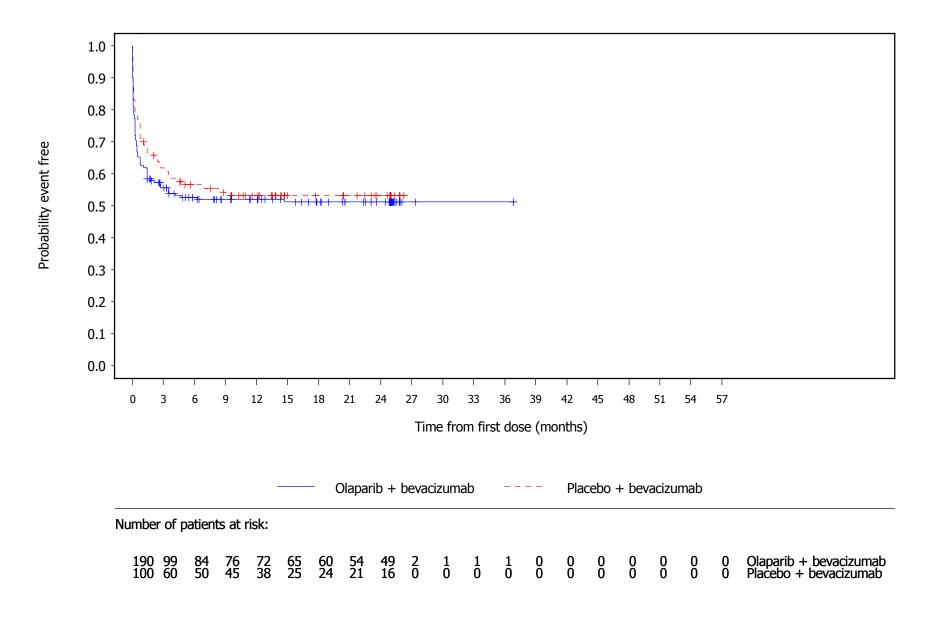
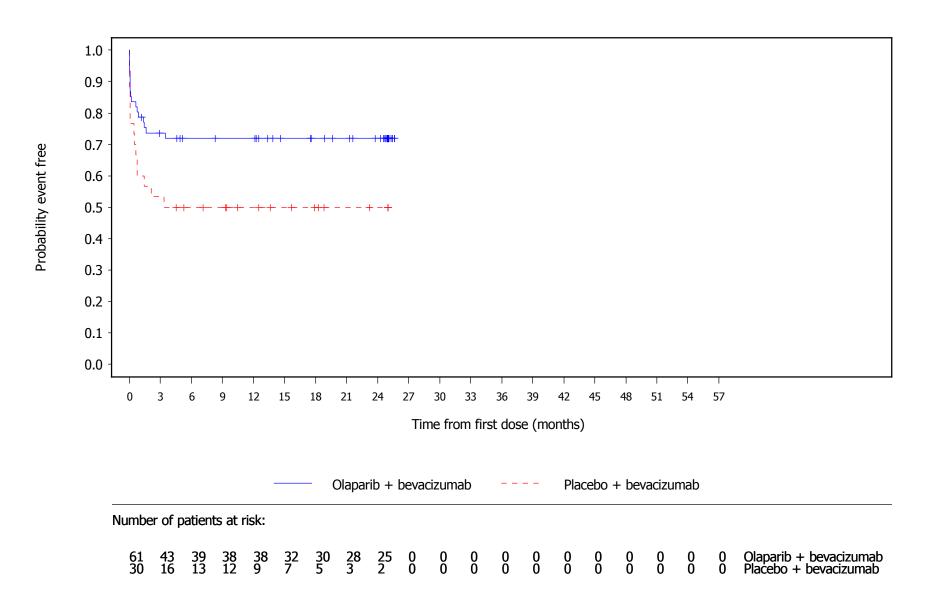


Figure 3.5.24 PAOLA1: Kaplan-Meier plot of AE max CTCAE grade=1 or 2 for ECOG performance status at Baseline=(1) Restricted activity

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020



					Treatment effect										
		laparib + vacizumab (N=255)		lacebo + vacizumab (N=131)		Odds	Ratio		Rela	tive Ris	k	Risk D	ifferen	ce	
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estima (95% C		2- sided p- value	Estim (95%)		2- sided p- value	Estimat (95% Cl		2- sided p- value	
AE [c][f][i]	255	255(100)	131	127(96.9)	18.04(1.90,	2396.90)	0.0083	1.03(1.001	, 1.06)	0.0438	1	IC		
AE SOC: General disorders and administration site conditions [b][e][h]	255	156(61.2)	131	57(43.5)	2.05(1.34,	3.15)	0.0010	1.41(1.14	, 1.77)	0.0010	0.18(0.07,	0.28)	0.0010	
AE PT: Fatigue [b][e][h]	255	141(55.3)	131	44(33.6)	2.45(1.58,	3.81)	<0.0001 *	1.65(1.28	, 2.18)	<0.0001 *	0.22(0.11,	0.32)	<0.0001 *	
AE PT: Pyrexia [b][e][h]	255	16(6.3)	131	4(3.1)	2.13(0.76,	7.54)	0.1584	2.05(0.77	, 7.06)	0.1584	0.03(-0.01,	0.07)	0.1584	
AE PT: Oedema [b][e][h]	255	8(3.1)	131	4(3.1)	1.03(0.32,	3.91)	0.9641	1.03(0.33	, 3.79)	0.9641	0.00(-0.04,	0.04)	0.9641	
AE PT: Oedema peripheral [b][e][h]	255	15(5.9)	131	7(5.3)	1.11(0.45,	2.96)	0.8280	1.10(0.48	, 2.82)	0.8280	0.01(-0.05,	0.05)	0.8280	
AE PT: Mucosal inflammation [b][e][h]	255	15(5.9)	131	4(3.1)	1.98(0.70,	7.07)	0.2062	1.93(0.72	, 6.65)	0.2062	0.03(-0.02,	0.07)	0.2062	
AE PT: Pain [b][e][h]	255	7(2.7)	131	4(3.1)	0.90(0.27,	3.47)	0.8639	0.90(0.28	, 3.38)	0.8639	-0.00(-0.05,	0.03)	0.8639	

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NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

									Treatm	ment eff	ect			
	bev	aparib + vacizumab (N=255)	bev	lacebo + vacizumab (N=131)		Odds	Ratio		Rela	tive Ris	k	Risk D	ifferen	ce
	n	Number (%) of patients with events [a]		Number (%) of patients with events [a]		Estimat (95% C	I)	2- sided p- value	Estim (95%)	CI)	2- sided p- value	Estimat (95% C	[)	2- sided p- value
AE SOC: Eye disorders [b][e][h]	255	9(3.5)	131	4(3.1)	1.16(0.37,	4.35)	0.8046	1.16(0.38	, 4.20)	0.8046	0.00(-0.04,	0.04)	0.8046
AE SOC: Surgical and medical procedures [b][e][h]	255	10(3.9)	131	3(2.3)	1.74(0.52,	7.87)	0.3853	1.71(0.53	, 7.54)	0.3853	0.02(-0.02,	0.05)	0.3853
AE SOC: Endocrine disorders [b][e][h]	255	4(1.6)	131	9(6.9)	0.22(0.06,	0.68)	0.0083	0.23(0.06	, 0.69)	0.0083	-0.05(-0.11,	-0.01)	0.0083
AE SOC: Respiratory, thoracic and mediastinal disorders [b][e][h]	255	62(24.3)	131	28(21.4)	1.18(0.72,	1.98)	0.5157	1.14(0.78	, 1.72)	0.5157	0.03(-0.06,	0.11)	0.5157
AE PT: Dyspnoea [b][e][h]	255	22(8.6)	131	3(2.3)	4.03(1.36,	17.24)	0.0094	3.77(1.34	, 15.71)	0.0094	0.06(0.02,	0.11)	0.0094
AE PT: Epistaxis [b][e][h]	255	18(7.1)	131	7(5.3)	1.35(0.57,	3.54)	0.5105	1.32(0.59	, 3.32)	0.5105	0.02(-0.04,	0.07)	0.5105
AE PT: Cough [b][e][h]	255	11(4.3)	131	6(4.6)	0.94(0.35,	2.78)	0.9041	0.94(0.37	, 2.68)	0.9041	-0.00(-0.05,	0.04)	0.9041

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NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

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[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c].

[e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h].

Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

									Т	reatme	ent effe	ect			
	ber	aparib + vacizumab (N=255)	ber	lacebo + vacizumab (N=131)		Odds	Ratio			Relat	ive Ris	k	Risk D	ifferen	ce
	n	<pre>Number (%) of patients with events [a]</pre>	n	<pre>Number (%) of patients with events [a]</pre>		Estimat (95% C		2- sided p- value		Estimat (95% C		2- sided p- value	Estima (95% C		2- sided p- value
AE PT: Rhinorrhoea [b][e][h]	255	5(2.0)	131	3(2.3)	0.85(0.21,	4.21)	0.8310	0.86(0.21,	4.12)	0.8310	-0.00(-0.04,	0.03)	0.8310
AE PT: Oropharyngeal pain [b][e][h]	255	4(1.6)	131	4(3.1)	0.51(0.12,	2.17)	0.3452	0.51(0.12,	2.14)	0.3452	-0.01(-0.06,	0.02)	0.3452
AE SOC: Skin and subcutaneous tissue disorders [b][e][h]	255	44(17.3)	131	19(14.5)	1.23(0.69,	2.25)	0.4852	1.19(0.74,	2.00)	0.4852	0.03(-0.05,	0.10)	0.4852
AE PT: Alopecia [b][e][h]	255	8(3.1)	131	2(1.5)	2.09(0.51,	13.97)	0.3245	2.05(0.52,	13.49)	0.3245	0.02(-0.02,	0.05)	0.3245
AE PT: Rash [b][e][h]	255	10(3.9)	131	7(5.3)	0.72(0.27,	2.03)	0.5251	0.73(0.29,	1.98)	0.5251	-0.01(-0.07,	0.03)	0.5251
AE PT: Erythema [c][g][i]	255	8(3.1)	131	0	9.03(1.11,1	171.24	0.0367		1	NC		1	NC	
AE PT: Pruritus [b][e][h]	255	5(2.0)	131	3(2.3)	0.85(0.21,	4.21)	0.8310	0.86(0.21,	4.12)	0.8310	-0.00(-0.04,	0.03)	0.8310
AE SOC: Renal and urinary disorders [b][e][h]	255	27(10.6)	131	24(18.3)	0.53(0.29,	0.96)	0.0373	0.58(0.35,	0.97)	0.0373	-0.08(-0.16,	-0.00)	0.0373

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[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

					Treatment eff	ect		
	Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=131)	Odds Ratio		Relative Ris	sk	Risk Differen	ce
	Number (%) of patients with n events [a			2- sided p- value	Estimate (95% CI)	2- sided p- value	Estimate (95% CI)	2- sided p- value
AE PT: Proteinuria [b][e][h]	255 19(7.5)	131 19(14.5)	0.47(0.24, 0.94)	0.0316	0.51(0.28, 0.94)	0.0316	-0.07(-0.14, -0.01)	0.0316
AE SOC: Blood and lymphatic system disorders [b][e][h]	255 148(58.0)	131 41(31.3)	3.04(1.96, 4.77)	<0.0001	1.85(1.43, 2.48)	<0.0001	0.27(0.17, 0.36)	<0.0001
AE PT: Anaemia [b][e][h]	255 102(40.0)	131 12(9.2)	6.61(3.60, 13.19)	<0.0001 *	4.37(2.62, 8.11)	<0.0001	0.31(0.23, 0.38)	<0.0001 *
AE PT: Leukopenia [b][e][h]	255 46(18.0)	131 11(8.4)	2.40(1.24, 5.04)	0.0085	2.15(1.20, 4.24)	0.0085	0.10(0.03, 0.16)	0.0085
AE PT: Lymphopenia [b][e][h]	255 60(23.5)	131 10(7.6)	3.72(1.91, 7.98)	<0.0001	3.08(1.72, 6.22)	<0.0001	0.16(0.09, 0.23)	<0.0001
AE PT: Neutropenia [b][e][h]	255 30(11.8)	131 15(11.5)	1.03(0.54, 2.04)	0.9273	1.03(0.58, 1.90)	0.9273	0.00(-0.07, 0.07)	0.9273
AE PT: Thrombocytopeni a [b][e][h]	255 12(4.7)	131 3(2.3)	2.11(0.66, 9.37)	0.2247	2.05(0.67, 8.90)	0.2247	0.02(-0.02, 0.06)	0.2247

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[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

									Treatr	ment eff	ect			
	bev	aparib + vacizumab (N=255)	bev	lacebo + vacizumab (N=131)		Odds	s Ratio		Rela	tive Ris	k	Risk D	ifferen	ce
	n	Number (%) of patients with events [a]		Number (%) of patients with events [a]		Estimat		2- sided p- value	Estim (95%)		2- sided p- value	Estimat (95% Cl		2- sided p- value
AE SOC: Gastrointestina l disorders [b][e][h]	255	189(74.1)	131	83(63.4)	1.66(1.05,	2.60)	0.0297	1.17(1.01	, 1.37)	0.0297	0.11(0.01,	0.21)	0.0297
AE PT: Abdominal pain [b][e][h]	255	56(22.0)	131	32(24.4)	0.87(0.53,	1.44)	0.5858	0.90(0.62	, 1.33)	0.5858	-0.02(-0.12,	0.06)	0.5858
AE PT: Intestinal obstruction [b][e][h]	255	5(2.0)	131	1(0.8)	2.60(0.41,	50.06)	0.3394	2.57(0.42	, 48.92)	0.3394	0.01(-0.02,	0.04)	0.3394
AE PT: Diarrhoea [b][e][h]	255	50(19.6)	131	25(19.1)	1.03(0.61,	1.79)	0.9019	1.03(0.68	, 1.61)	0.9019	0.01(-0.08,	0.09)	0.9019
AE PT: Dyspepsia [b][e][h]	255	12(4.7)	131	3(2.3)	2.11(0.66,	9.37)	0.2247	2.05(0.67	, 8.90)	0.2247	0.02(-0.02,	0.06)	0.2247
AE PT: Vomiting [b][e][h]	255	54(21.2)	131	16(12.2)	1.93(1.08,	3.63)	0.0263	1.73(1.06	, 3.02)	0.0263	0.09(0.01,	0.16)	0.0263
AE PT: Haemorrhoids [b][e][h]	255	7(2.7)	131	4(3.1)	0.90(0.27,	3.47)	0.8639	0.90(0.28	, 3.38)	0.8639	-0.00(-0.05,	0.03)	0.8639
AE PT: Constipation [b][e][h]	255	28(11.0)	131	15(11.5)	0.95(0.50,	1.90)	0.8897	0.96(0.54	, 1.78)	0.8897	-0.00(-0.08,	0.06)	0.8897

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[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

								Т	reatme	ent effe	ect			
	be ⁻	.aparib + vacizumab (N=255)	ber	lacebo + vacizumab (N=131)	Odo	ds Ratio			Relat	ive Ris	k	Risk Di	fferen	ce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]	Estim (95%		2- sided p- value		Estima (95% C		2- sided p- value	Estimat (95% CI		2- sided p- value
AE PT: Abdominal pain upper [b][e][h]	255	10(3.9)	131	4(3.1)	1.30(0.42	, 4.80)	0.6614	1.28(0.44,	4.61)	0.6614	0.01(-0.04,	0.05)	0.6614
AE PT: Stomatitis [b][e][h]	255	12(4.7)	131	2(1.5)	3.19(0.85	, 20.66)	0.0897	3.08(0.86,	19.62)	0.0897	0.03(-0.01,	0.07)	0.0897
AE PT: Subileus [c][e][h]	255	2(0.8)	131	2(1.5)	0.51(0.08	, 3.34)	0.4599	0.51(0.06,	4.24)	0.5069	-0.01(-0.04,	0.01)	0.5069
AE PT: Nausea [b][e][h]	255	144(56.5)	131	30(22.9)	4.37(2.74	, 7.12)	<0.0001	2.47(1.81,	3.52)	<0.0001	0.34(0.24,	0.43)	<0.0001
AE PT: Gingival bleeding [b][e][h]	255	9(3.5)	131	2(1.5)	2.36(0.60	, 15.63)	0.2385	2.31(0.61,	15.02)	0.2385	0.02(-0.02,	0.05)	0.2385
AE PT: Toothache [b][e][h]	255	6(2.4)	131	3(2.3)	1.03(0.27	, 4.93)	0.9690	1.03(0.28,	4.81)	0.9690	0.00(-0.04,	0.03)	0.9690
AE SOC: Immune system disorders [b][e][h]	255	7(2.7)	131	1(0.8)	3.67(0.64	, 68.96)	0.1603	3.60(0.65,	66.85)	0.1603	0.02(-0.01,	0.05)	0.1603
AE SOC: Nervous system disorders [b][e][h]	255	87(34.1)	131	32(24.4)	1.60(1.003	, 2.60)	0.0483	1.40(1	L.002,	2.01)	0.0483	0.10(0.001,	0.19)	0.0483

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[e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h].

Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

									ect						
	be	.aparib + vacizumab (N=255)	ber	lacebo + vacizumab (N=131)		Odd	s Ratio			Relat	ive Ris	k	Risk D	ifferen	ce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estima (95% C		2- sided p- value		Estima (95% C		2- sided p- value	Estimat (95% CI		2- sided p- value
AE PT: Dysgeusia [b][e][h]	255	23(9.0)	131	2(1.5)	6.39(1.85,	40.24)	0.0016	5.91(1.79,	36.45)	0.0016	0.07(0.03,	0.12)	0.0016
AE PT: Headache [b][e][h]	255	39(15.3)	131	22(16.8)	0.89(0.51,	1.60)	0.7032	0.91(0.57,	1.50)	0.7032	-0.01(-0.10,	0.06)	0.7032
AE PT: Neuropathy peripheral [b][e][h]	255	22(8.6)	131	5(3.8)	2.38(0.95,	7.24)	0.0657	2.26(0.95,	6.63)	0.0657	0.05(-0.00,	0.10)	0.0657
AE PT: Polyneuropathy [b][e][h]	255	8(3.1)	131	2(1.5)	2.09(0.51,	13.97)	0.3245	2.05(0.52,	13.49)	0.3245	0.02(-0.02,	0.05)	0.3245
AE PT: Dizziness [b][e][h]	255	8(3.1)	131	3(2.3)	1.38(0.39,	6.39)	0.6297	1.37(0.40,	6.18)	0.6297	0.01(-0.03,	0.04)	0.6297
AE SOC: Ear and labyrinth disorders [b][e][h]	255	10(3.9)	131	8(6.1)	0.63(0.24,	1.68)	0.3442	0.64(0.26,	1.65)	0.3442	-0.02(-0.08,	0.02)	0.3442
AE PT: Vertigo [b][e][h]	255	6(2.4)	131	6(4.6)	0.50(0.15,	1.63)	0.2450	0.51(0.16,	1.61)	0.2450	-0.02(-0.07,	0.01)	0.2450

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[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

					Treatment effe	ect		
	Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=131)	Odds Ratio		Relative Ris	k	Risk Differen	ce
	Number (%) of patients with n events [a]	Number (%) of patients with n events [a]	Estimate	2- sided p- value	Estimate (95% CI)	2- sided p- value	Estimate (95% CI)	2- sided p- value
AE SOC: Vascular disorders [b][e][h]	255 137(53.7)	131 82(62.6)	0.69(0.45, 1.06)	0.0946	0.86(0.72, 1.03)	0.0946	-0.09(-0.19, 0.02)	0.0946
AE PT: Hot flush [b][e][h]	255 7(2.7)	131 3(2.3)	1.20(0.33, 5.66)	0.7878	1.20(0.34, 5.49)	0.7878	0.00(-0.03, 0.04)	0.7878
AE PT: Hypertension [b][e][h]	255 122(47.8)	131 78(59.5)	0.62(0.41, 0.95)	0.0290	0.80(0.67, 0.98)	0.0290	-0.12(-0.22, -0.01)	0.0290
AE SOC: Cardiac disorders [b][e][h]	255 7(2.7)	131 7(5.3)	0.50(0.17, 1.49)	0.2079	0.51(0.18, 1.47)	0.2079	-0.03(-0.08, 0.01)	0.2079
AE SOC: Infections and infestations [b][e][h]	255 128(50.2)	131 63(48.1)	1.09(0.71, 1.66)	0.6953	1.04(0.85, 1.31)	0.6953	0.02(-0.08, 0.13)	0.6953
AE PT: Bronchitis [b][e][h]	255 13(5.1)	131 3(2.3)	2.29(0.72, 10.13)	0.1691	2.23(0.73, 9.58)	0.1691	0.03(-0.01, 0.07)	0.1691
AE PT: Gastroenteritis [c][g][i]	255 13(5.1)	131 0	14.64(1.92,1878.82)	0.0042	NC		NC	
AE PT: Gingivitis [b][e][h]	255 7(2.7)	131 2(1.5)	1.82(0.43, 12.33)	0.4367	1.80(0.44, 11.95)	0.4367	0.01(-0.02, 0.04)	0.4367

Includes adverse events with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 23.0.

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c].

[e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h].

Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

								Treatm	ent effe	ect			
	bev	aparib + vacizumab (N=255)	be	lacebo + vacizumab (N=131)	Odd	ls Ratio		Rela	tive Ris	k	Risk Di	fferen	ce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]	Estim (95% (2- sided p- value	Estima (95% (2- sided p- value	Estimat (95% CI		2- sided p- value
AE PT: Influenza [b][e][h]	255	8(3.1)	131	1(0.8)	4.21(0.76	, 78.53)	0.1092	4.11(0.77	75.81)	0.1092	0.02(-0.01,	0.05)	0.1092
AE PT: Urinary tract infection [b][e][h]	255	41(16.1)	131	12(9.2)	1.90(0.99	, 3.91)	0.0542	1.76(0.99	3.38)	0.0542	0.07(-0.00,	0.13)	0.0542
AE PT: Infection [b][e][h]	255	9(3.5)	131	1(0.8)	4.76(0.88	, 88.16)	0.0741	4.62(0.88	84.77)	0.0741	0.03(-0.00,	0.06)	0.0741
AE PT: Nasopharyngitis [b][e][h]	255	15(5.9)	131	10(7.6)	0.76(0.33	, 1.79)	0.5130	0.77(0.36	1.73)	0.5130	-0.02(-0.08,	0.03)	0.5130
AE PT: Pharyngitis [b][e][h]	255	8(3.1)	131	4(3.1)	1.03(0.32	, 3.91)	0.9641	1.03(0.33	3.79)	0.9641	0.00(-0.04,	0.04)	0.9641
AE PT: Rhinitis [b][e][h]	255	10(3.9)	131	4(3.1)	1.30(0.42	, 4.80)	0.6614	1.28(0.44	4.61)	0.6614	0.01(-0.04,	0.05)	0.6614
AE PT: Tooth abscess [b][e][h]	255	8(3.1)	131	4(3.1)	1.03(0.32	, 3.91)	0.9641	1.03(0.33	3.79)	0.9641	0.00(-0.04,	0.04)	0.9641
AE PT: Tooth infection [b][e][h]	255	7(2.7)	131	2(1.5)	1.82(0.43	, 12.33)	0.4367	1.80(0.44	11.95)	0.4367	0.01(-0.02,	0.04)	0.4367

Includes adverse events with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 23.0.

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c].

[e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h].

Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

					Treatment effo	ect		
	Olaparib + bevacizumab (N=255) _	Placebo + bevacizumab (N=131)	Odds Ratio		Relative Ris	k	Risk Differen	ce
AE PT: Cystitis	Number (%) of patients with n events [255 11(4.3)		Estimate (95% CI) 0.61(0.25, 1.55)	2- sided p- value	Estimate (95% CI) 0.63(0.27, 1.52)	2- sided p- value	Estimate (95% CI) -0.03(-0.08, 0.02)	2- sided p- value 0.2929
[b][e][h]	233 11(1.3	, 131)(0.),	0.01(0.23, 1.33)	0.2525	0.03(0.27, 1.32,	0.2525	0.03(0.00, 0.02)	0.2525
AE SOC: Psychiatric disorders [b][e][h]	255 21(8.2	131 13(9.9)	0.81(0.40, 1.72)	0.5827	0.83(0.43, 1.65)	0.5827	-0.02(-0.08, 0.04)	0.5827
AE PT: Anxiety [b][e][h]	255 7(2.7) 131 5(3.8)	0.71(0.22, 2.44)	0.5716	0.72(0.23, 2.39)	0.5716	-0.01(-0.06, 0.03)	0.5716
AE PT: Depression [b][e][h]	255 6(2.4	131 1(0.8)	3.13(0.53, 59.48)	0.2339	3.08(0.53, 57.89)	0.2339	0.02(-0.01, 0.04)	0.2339
AE PT: Sleep disorder [b][e][h]	255 7(2.7) 131 7(5.3)	0.50(0.17, 1.49)	0.2079	0.51(0.18, 1.47)	0.2079	-0.03(-0.08, 0.01)	0.2079
AE SOC: Musculoskeletal and connective tissue disorders [b][e][h]	255 106(41.6) 131 57(43.5)	0.92(0.60, 1.42)	0.7146	0.96(0.75, 1.23)	0.7146	-0.02(-0.12, 0.08)	0.7146
AE PT: Arthralgia [b][e][h]	255 64(25.1) 131 30(22.9)	1.13(0.69, 1.87)	0.6327	1.10(0.76, 1.63)	0.6327	0.02(-0.07, 0.11)	0.6327

Includes adverse events with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 23.0.

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c].

[e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h].

Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

									Trea	atment eff	ect			
	bev	aparib + vacizumab (N=255)	ber	lacebo + vacizumab (N=131)		Odds	s Ratio		Re	lative Ris	k	Risk Di	fferen	ce
	n	<pre>Number (%) of patients with events [a]</pre>	n	<pre>Number (%) of patients with events [a]</pre>		stima 95% C		2- sided p- value		imate % CI)	2- sided p- value	Estimat (95% CI		2- sided p- value
AE PT: Muscle spasms [b][e][h]	255	11(4.3)	131	8(6.1)	0.69(0.27,	1.83)	0.4478	0.71(0.	29, 1.78)	0.4478	-0.02(-0.07,	0.03)	0.4478
AE PT: Myalgia [b][e][h]	255	21(8.2)	131	7(5.3)	1.59(0.69,	4.13)	0.2882	1.54(0.	71, 3.83)	0.2882	0.03(-0.03,	0.08)	0.2882
AE PT: Neck pain [b][e][h]	255	4(1.6)	131	5(3.8)	0.40(0.10,	1.54)	0.1794	0.41(0.	10, 1.53)	0.1794	-0.02(-0.07,	0.01)	0.1794
AE PT: Back pain [b][e][h]	255	17(6.7)	131	8(6.1)	1.10(0.47,	2.76)	0.8317	1.09(0.	50, 2.61)	0.8317	0.01(-0.05,	0.05)	0.8317
AE PT: Pain in extremity [b][e][h]	255	17(6.7)	131	8(6.1)	1.10(0.47,	2.76)	0.8317	1.09(0.	50, 2.61)	0.8317	0.01(-0.05,	0.05)	0.8317
AE PT: Musculoskeletal pain [b][e][h]	255	9(3.5)	131	9(6.9)	0.50(0.19,	1.30)	0.1510	0.51(0.	21, 1.29)	0.1510	-0.03(-0.09,	0.01)	0.1510
AE SOC: Metabolism and nutrition disorders [b][e][h]	255	30(11.8)	131	9(6.9)	1.81(0.86,	4.15)	0.1198	1.71(0.	88, 3.73)	0.1198	0.05(-0.01,	0.11)	0.1198
AE PT: Decreased appetite [b][e][h]	255	23(9.0)	131	4(3.1)	3.15(1.18,	10.91)	0.0202	2.95(1.	17, 9.92)	0.0202	0.06(0.01,	0.11)	0.0202

Includes adverse events with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 23.0.

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

										Treatme	ent effe	ect			
	bev	aparib + vacizumab (N=255)	be	lacebo + vacizumab (N=131)		Odds	Ratio			Relat	ive Ris	k	Risk Di	fferen	ce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estimat		2- sided p- value		Estima (95% C		2- sided p- value	Estimat (95% CI		2- sided p- value
AE SOC: Investigations [b][e][h]	255	73(28.6)	131	29(22.1)	1.41(0.87,	2.34)	0.1668	1.29(0.90,	1.92)	0.1668	0.06(-0.03,	0.15)	0.1668
AE PT: Weight increased [b][e][h]	255	11(4.3)	131	10(7.6)	0.55(0.22,	1.34)	0.1834	0.57(0.24,	1.32)	0.1834	-0.03(-0.09,	0.01)	0.1834
AE PT: Blood creatinine increased [b][e][h]	255	13(5.1)	131	1(0.8)	6.98(1.37,1	L27.51)	0.0151	6.68(1.35,	120.60)	0.0151	0.04(0.01,	0.08)	0.0151
AE PT: White blood cell count decreased [b][e][h]	255	8(3.1)	131	2(1.5)	2.09(0.51,	13.97)	0.3245	2.05(0.52,	13.49)	0.3245	0.02(-0.02,	0.05)	0.3245
AE PT: Lymphocyte count decreased [b][e][h]	255	7(2.7)	131	1(0.8)	3.67(0.64,	68.96)	0.1603	3.60(0.65,	66.85)	0.1603	0.02(-0.01,	0.05)	0.1603
AE PT: Neutrophil count decreased [b][e][h]	255	20(7.8)	131	6(4.6)	1.77(0.73,	4.95)	0.2119	1.71(0.75,	4.59)	0.2119	0.03(-0.02,	0.08)	0.2119
AE PT: Platelet count decreased [b][e][h]	255	5(2.0)	131	1(0.8)	2.60(0.41,	50.06)	0.3394	2.57(0.42,	48.92)	0.3394	0.01(-0.02,	0.04)	0.3394

Includes adverse events with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 23.0.

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

										Treatme	ent effe	ect			
	beva	parib + acizumab N=255)	bev	lacebo + vacizumab (N=131)		Odds	Ratio			Relat	ive Ris	k	Risk D	ifferer	ice
	v V	Number (%) of patients with events [a]		Number (%) of patients with events [a]		Estimat (95% Ci		2- sided p- value		Estimat (95% CI		2- sided p- value	Estima (95% C		2- sided p- value
AE SOC: Injury, poisoning and procedural complications [b][e][h]	255	15(5.9)	131	9(6.9)	0.85(0.37,	2.07)	0.7056	0.86(0.39,	1.99)	0.7056	-0.01(-0.07,	0.04)	0.7056

Includes adverse events with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 23.0.

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c].

[e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h].

Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

Table 3.6.2 PAOLA1: Summary of analysis of serious adverse events (odds ratio, relative risk and risk difference)
Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

									T	reatme	ent effe	ect			
	bev	aparib + vacizumab (N=255)	bes	lacebo + vacizumab (N=131)		Odd	s Ratio			Relat	ive Ris	k	Risk Di	ifferen	ce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estima (95% C		2- sided p- value		stima 95% C		2- sided p- value	Estimat (95% CI		2- sided p- value
SAE [b][e][h]	255	73(28.6)	131	45(34.4)	0.77(0.49,	1.21)	0.2501	0.83(0.62,	1.14)	0.2501	-0.06(-0.16,	0.04)	0.2501
SAE SOC: Respiratory, thoracic and mediastinal disorders [b][e][h]	255	7(2.7)	131	1(0.8)	3.67(0.64,	68.96)	0.1603	3.60(0.65,	66.85)	0.1603	0.02(-0.01,	0.05)	0.1603
SAE SOC: Blood and lymphatic system disorders [b][e][h]	255	17(6.7)	131	1(0.8)	9.29(1.87,	168.17)	0.0029	8.73(1.83,	156.43)	0.0029	0.06(0.02,	0.10)	0.0029
SAE PT: Anaemia [b][e][h]	255	13(5.1)	131	1(0.8)	6.98(1.37,	127.51)	0.0151 *	6.68(1.35,	120.60)	0.0151 *	0.04(0.01,	0.08)	0.0151
SAE SOC: Gastrointestina l disorders [b][e][h]	255	11(4.3)	131	10(7.6)	0.55(0.22,	1.34)	0.1834	0.57(0.24,	1.32)	0.1834	-0.03(-0.09,	0.01)	0.1834
SAE SOC: Vascular disorders [b][e][h]	255	24(9.4)	131	16(12.2)	0.75(0.38,	1.48)	0.3977	0.77(0.43,	1.43)	0.3977	-0.03(-0.10,	0.04)	0.3977
SAE PT: Hypertension [b][e][h]	255	20(7.8)	131	16(12.2)	0.61(0.31,	1.24)	0.1697	0.64(0.35,	1.22)	0.1697	-0.04(-0.11,	0.02)	0.1697

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

								Treatr	ment eff	ect			
	be	laparib + vacizumab (N=255)		lacebo + vacizumab (N=131)	0dd	s Ratio		Rela	tive Ris	k	Risk D	ifferen	ıce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]	Estima (95% (2- sided p- value	Estim (95%		2- sided p- value	Estimat (95% Cl		2- sided p- value
SAE SOC: Infections and infestations [b][e][h]	255	12(4.7)	131	9(6.9)	0.67(0.28,	1.68)	0.3827	0.68(0.30	, 1.64)	0.3827	-0.02(-0.08,	0.03)	0.3827

Includes adverse events with an onset date on or after the date of first dose and up to and including 30 days following the date of last dose of olaparib/placebo. MedDRA version 23.0.

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

Table 3.6.3 PAOLA1: Summary of analysis of adverse events leading to discontinuation of study treatment (odds ratio, relative risk and risk difference)

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

									Tr	reatme:	nt effe	ect			
	be	laparib + vacizumab (N=255)		lacebo + vacizumab (N=131)		Odds	Ratio]	Relati	ive Ris	k	Risk D	ifferen	ice
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estimat		2- sided p- value		stimat 95% CI		2- sided p- value	Estima (95% C		2- sided p- value
AE leading to discontinuation of treatment [b][e][h]	255	50(19.6)	131	8(6.1)	3.75(1.81,	8.79)	0.0002	3.21(1.67,	7.15)	0.0002	0.14(0.07,	0.20)	0.0002

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

											Treatr	ment eff	ect			
	be	laparib + vacizumab (N=255)	be	lacebo + vacizumab (N=131)		Ode	ds Rat	tio			Rela	tive Ris	k	Risk	Differer	nce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estim (95%			2- sided p- value		Estim (95%		2- sided p- value	Esti (95%		2- sided p- value
AE max CTCAE grade>=3 [b][e][h]	255	147(57.6)	131	65(49.6)	1.38(0.91	, 2.	11)	0.1337	1.16(0.96	, 1.44)	0.1337	0.08(-0.0	2, 0.18)	0.1337
AE G>=3 SOC: General disorders and administration site conditions [b][e][h]	255	15(5.9)	131	1(0.8)	8.12(1.62	,147.	67)	0.0067	7.71(1.59	,138.51)	0.0067	0.05(0.0	2, 0.09)	0.0067
AE G>=3 PT: Fatigue [c][g][i]	255	14(5.5)	131	0	15.79(2.08	,2023	.85)	0.0027			NC			NC	
AE G>=3 SOC: Respiratory, thoracic and mediastinal disorders [b][e][h]	255	6(2.4)	131	2(1.5)	1.55(0.35	, 10.	71)	0.5801	1.54(0.36	, 10.42)	0.5801	0.01(-0.0	0.04)	0.5801
AE G>=3 SOC: Renal and urinary disorders [c][g][i]	255	3(1.2)	131	0	3.65(0.35	,491.	54)	0.3214			NC			NC	

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

										Treatment ef	fect				
	ber	aparib + vacizumab (N=255)	be ⁻	lacebo + vacizumab (N=131)		Odd	ls Ratio			Relative R	.sk		Risk Di	ifferen	ce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estima (95% (2- sided p- value		Estimate (95% CI)	2- sided p- value		Estimat (95% CI		2- sided p- value
AE G>=3 SOC: Blood and lymphatic system disorders [b][e][h]	255	71(27.8)	131	5(3.8)	9.72(4.20,	28.31)	<0.0001 *	7.29(3.37, 20.42) <0.0001 *	0.24(0.17,	0.30)	<0.0001 *
AE G>=3 PT: Anaemia [b][e][h]	255	47(18.4)	131	1(0.8)	29.37(6.31,	523.04)	<0.0001	24.15(5.39,425.07) <0.0001	0.18(0.13,	0.23)	<0.0001
AE G>=3 PT: Leukopenia [c][g][i]	255	6(2.4)	131	0	6.85(0.80,	896.13)	0.0873		NC			N	IC	
AE G>=3 PT: Lymphopenia [b][e][h]	255	19(7.5)	131	3(2.3)	3.44(1.14,	14.81)	0.0260	3.25(1.13, 13.67) 0.0260	0.05(0.01,	0.09)	0.0260
AE G>=3 PT: Neutropenia [b][e][h]	255	12(4.7)	131	1(0.8)	6.42(1.24,	117.55)	0.0226	6.16(1.23,111.64) 0.0226	0.04(0.01,	0.07)	0.0226
AE G>=3 SOC: Gastrointestina l disorders [b][e][h]	255	30(11.8)	131	12(9.2)	1.32(0.67,	2.77)	0.4307	1.28(0.70, 2.53) 0.4307	0.03(-0.04,	0.09)	0.4307
AE G>=3 PT: Diarrhoea [b][e][h]	255	7(2.7)	131	3(2.3)	1.20(0.33,	5.66)	0.7878	1.20(0.34, 5.49) 0.7878	0.00(-0.03,	0.04)	0.7878

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

									Tı	reatme	ent effe	ect			
	bev	aparib + vacizumab (N=255)	ber	lacebo + vacizumab (N=131)		Odd	s Ratio			Relat	ive Ris	k	Risk D	ifferen	.ce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estima (95% C		2- sided p- value		stima 95% C		2- sided p- value	Estima (95% C	I)	2- sided p- value
AE G>=3 PT: Nausea [c][g][i]	255	8(3.1)	131	0	9.03(1.11,	1171.24	*]	NC]	NC	
AE G>=3 SOC: Nervous system disorders [b][e][h]	255	7(2.7)	131	4(3.1)	0.90(0.27,	3.47)	0.8639	0.90(0.28,	3.38)	0.8639	-0.00(-0.05,	0.03)	0.8639
AE G>=3 SOC: Vascular disorders [b][e][h]	255	48(18.8)	131	42(32.1)	0.49(0.30,	0.80)	0.0041	0.59(0.41,	0.84)	0.0041	-0.13(-0.23,	-0.04)	0.0041
AE G>=3 PT: Hypertension [b][e][h]	255	45(17.6)	131	42(32.1)	0.45(0.28,	0.74)	0.0016	0.55(0.38,	0.79)	0.0016	-0.14(-0.24,	-0.05)	0.0016
AE G>=3 SOC: Infections and infestations [b][e][h]	255	14(5.5)	131	10(7.6)	0.70(0.31,	1.67)	0.4158	0.72(0.33,	1.63)	0.4158	-0.02(-0.08,	0.03)	0.4158
AE G>=3 SOC: Investigations [b][e][h]	255	16(6.3)	131	5(3.8)	1.69(0.64,	5.25)	0.2998	1.64(0.66,	4.94)	0.2998	0.02(-0.02,	0.07)	0.2998
AE G>=3 PT: Neutrophil count decreased [b][e][h]	255	6(2.4)	131	1(0.8)	3.13(0.53,	59.48)	0.2339	3.08(0.53,	57.89)	0.2339	0.02(-0.01,	0.04)	0.2339

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.</p>

Table 3.6.5 PAOLA1: Summary of analysis of non-severe adverse events with max. CTCAE grade 1 or 2 (odds ratio, relative risk and risk difference)

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

					Treatment ef:	ect		
	Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=131)	Odds Ratio		Relative Ri	sk	Risk Differer	nce
	Number (%) of patients with n events [a]	Number (%) of patients with n events [a]	Estimate (95% CI)	2- sided p- value	Estimate (95% CI)	2- sided p- value	Estimate (95% CI)	2- sided p- value
AE max CTCAE grade=1 or 2 [b][e][h]	255 108(42.4)	131 62(47.3)	0.82(0.54, 1.25)	0.3517	0.89(0.71, 1.14) 0.3517	-0.05(-0.15, 0.05)	0.3517

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

Olaparib PAOLA1, German benefit assessment

Table 3.8 PAOLA1: Duration of olaparib or placebo exposure (Safety analysis set) Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=131)
Total treatment duration [a]	n	255	131
	Mean	17.9	15.7
	SD	8.46	7.50
	Median	23.8	16.8
	Min	0	0
	Max	36	25
Actual treatment duration [b]	n	255	131
	Mean	17.3	15.5
	SD	8.45	7.46
	Median	22.1	16.8
	Min	0	0
	Max	36	25

[[]a] Total treatment duration (months) = (last dose date - first dose date + 1) / 30.4375.

[b] Actual treatment duration (months) = (total treatment duration - total duration of dose interruptions) / 30.4375.

Dose interruptions include those where the patient forgot to take all doses on a given day.

If patient is ongoing, data-cut-off has been used to calculate duration.

SD = standard deviation.

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Table 3.9 PAOLA1: Summary of analysis of adverse events leading to discontinuation of study treatment (total, and by SOC and PT)

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	Number (%)	of patients
System organ class / MedDRA Preferred term	Olaparib+ bevacizumab (N=255)	Placebo + bevacizumab (N=131)
Patients with any AE leading to discontinuation	50 (19.6)	8 (6.1)
Infections and infestations	2 (0.8)	1 (0.8)
Cellulitis	0 (0.0)	1 (0.8)
Cytomegalovirus infection	1 (0.4)	0 (0.0)
Infection	1 (0.4)	0 (0.0)
Neoplasms benign, malignant and unspecified (incl cysts and polyps)	3 (1.2)	1 (0.8)
Breast cancer	0 (0.0)	1 (0.8)
Bronchial carcinoma	1 (0.4)	0 (0.0)
Metastases to liver	1 (0.4)	0 (0.0)
Myelodysplastic syndrome	1 (0.4)	0 (0.0)
Pancreatic carcinoma	1 (0.4)	0 (0.0)
Blood and lymphatic system disorders	11 (4.3)	0 (0.0)
Anaemia	8 (3.1)	0 (0.0)
Aplastic anaemia	1 (0.4)	0 (0.0)
Haemolytic uraemic syndrome	1 (0.4)	0 (0.0)
Neutropenia	1 (0.4)	0 (0.0)
Psychiatric disorders	1 (0.4)	0 (0.0)
Mood altered	1 (0.4)	0 (0.0)
Nervous system disorders	2 (0.8)	1 (0.8)
Encephalitis autoimmune	1 (0.4)	0 (0.0)
Headache	1 (0.4)	1 (0.8)
Cardiac disorders	1 (0.4)	1 (0.8)

[[]a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

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Table 3.9 PAOLA1: Summary of analysis of adverse events leading to discontinuation of study treatment (total, and by SOC and PT)

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	Number (%)	of patients
System organ class / MedDRA Preferred term	Olaparib+ bevacizumab (N=255)	Placebo + bevacizumab (N=131)
Myocardial infarction	0 (0.0)	1 (0.8)
Prinzmetal angina	1 (0.4)	0 (0.0)
Respiratory, thoracic and mediastinal disorders	2 (0.8)	1 (0.8)
Dyspnoea	1 (0.4)	1 (0.8)
Pneumonitis	1 (0.4)	0 (0.0)
Gastrointestinal disorders	16 (6.3)	2 (1.5)
Abdominal pain	1 (0.4)	0 (0.0)
Gastric haemorrhage	0 (0.0)	1 (0.8)
Gastric perforation	1 (0.4)	0 (0.0)
Mechanical ileus	1 (0.4)	0 (0.0)
Nausea	10 (3.9)	1 (0.8)
Pancreatitis acute	1 (0.4)	0 (0.0)
Stomatitis	1 (0.4)	0 (0.0)
Vomiting	1 (0.4)	0 (0.0)
Skin and subcutaneous tissue disorders	1 (0.4)	0 (0.0)
Erythema nodosum	1 (0.4)	0 (0.0)
Musculoskeletal and connective tissue disorders	3 (1.2)	1 (0.8)
Arthralgia	2 (0.8)	1 (0.8)
Myalgia	1 (0.4)	0 (0.0)
General disorders and administration site conditions	6 (2.4)	0 (0.0)
Fatigue	4 (1.6)	0 (0.0)
Mucosal inflammation	2 (0.8)	0 (0.0)
Investigations	2 (0.8)	0 (0.0)

[[]a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

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Table 3.9 PAOLA1: Summary of analysis of adverse events leading to discontinuation of study treatment (total, and by SOC and PT)

Safety Analysis Set, HRD[42] positive, DCO 22MAR2020

	Number (%)	of patients
System organ class / MedDRA Preferred term	Olaparib+ bevacizumab (N=255)	Placebo + bevacizumab (N=131)
Neutrophil count decreased	1 (0.4)	0 (0.0)
Weight increased	1 (0.4)	0 (0.0)

Table 3.1 PAOLA1 Appendix: Summary of analysis of time to worsening in EORTC QLQ-C30 global QoL/health status, functional, symptom and single item scales (MID=15)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=132)		
	Number (%) Median time of patients (95% CI) n with events (months) [a]	Number (%) Median time of patients (95% CI) n with events (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [c]
EORTC QLQ-C30 Global QoL/health status (MID = 15)	255 146 (57.3) 16.6 (11.5,21.8)	132 81 (61.4) 13.8 (9.3,17.2)	0.85 0.65, 1.12	0.2343
EORTC QLQ-C30 Functional scale: Physical (MID = 15)	255 93 (36.5) 52.5 (26.3,52.5)	132 56 (42.4) 25.4 (19.9,30.7)	0.80 0.58, 1.13	0.1960
EORTC QLQ-C30 Functional scale: Role (MID = 15)	255 167 (65.5) 8.4 (5.8,11.2)	132 82 (62.1) 9.3 (6.1,16.2)	1.11 0.85, 1.46	0.4501
EORTC QLQ-C30 Functional scale: Cognitive (MID = 15)	255 174 (68.2) 11.1 (8.5,14.0)	132 85 (64.4) 8.5 (5.9,13.6)	0.91 0.70, 1.19	0.4835
EORTC QLQ-C30 Functional scale: Emotional (MID = 15)	255 157 (61.6) 13.8 (11.0,19.6)	132 85 (64.4) 11.2 (8.3,13.9)	0.93 0.71, 1.22	0.5616
EORTC QLQ-C30 Functional scale: Social (MID = 15)	255 148 (58.0) 13.5 (8.6,19.6)	132 81 (61.4) 11.3 (8.5,16.4)	0.91 0.69, 1.20	0.4710
EORTC QLQ-C30 Single item symptom scale: Loss of appetite (MID = 15)	255 146 (57.3) 13.6 (11.1,22.1)	132 65 (49.2) 22.3 (16.6,28.7)	1.42 1.06, 1.92	0.0227*
EORTC QLQ-C30 Single item symptom scale: Constipation (MID = 15)	255 133 (52.2) 19.9 (16.6,23.4)	132 69 (52.3) 19.7 (14.0,22.3)	1.03 0.77, 1.39	0.8313

Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

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[[]b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Table 3.1 PAOLA1 Appendix: Summary of analysis of time to worsening in EORTC QLQ-C30 global QoL/health status, functional, symptom and single item scales (MID=15)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	-	bevacizumab 255)		pevacizumab 132)			
	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	,	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
EORTC QLQ-C30 Single item symptom scale: Diarrhoea (MID = 15)	255 124 (48.6)	24.0 (16.6,25.9)	132 58 (43.9)	23.5 (19.9,35.0)	1.15	0.84, 1.58	0.4093
EORTC QLQ-C30 Single item symptom scale: Dyspnoea (MID = 15)	255 125 (49.0)	20.7 (16.0,52.5)	132 67 (50.8)	18.7 (12.3,24.9)	0.92	0.68, 1.25	0.5796
EORTC QLQ-C30 Symptom scale: Fatigue (MID = 15)	255 144 (56.5)	17.0 (12.5,22.0)	132 84 (63.6)	13.9 (9.6,17.5)	0.84	0.64, 1.11	0.2089
EORTC QLQ-C30 Single item symptom scale: Financial difficulties (MID = 15)	255 77 (30.2)	38.4 (38.4, NE)	132 48 (36.4)	NE (NE, NE)	0.72	0.50, 1.04	0.0709
EORTC QLQ-C30 Symptom scale: Nausea and vomiting (MID = 15)	255 178 (69.8)	5.8 (5.6, 8.7)	132 70 (53.0)	19.2 (12.7,23.5)	1.81	1.37, 2.42	<0.0001*
EORTC QLQ-C30 Symptom scale: Pain (MID = 15)	255 183 (71.8)	5.8 (5.6, 8.3)	132 95 (72.0)	5.6 (3.0, 8.1)	0.92	0.72, 1.19	0.5505
EORTC QLQ-C30 Single item symptom scale: Insomnia (MID = 15)	255 159 (62.4)	11.3 (8.4,14.0)	132 91 (68.9)	8.3 (5.6,11.1)	0.73	0.56, 0.95	0.0185*

Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation.

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

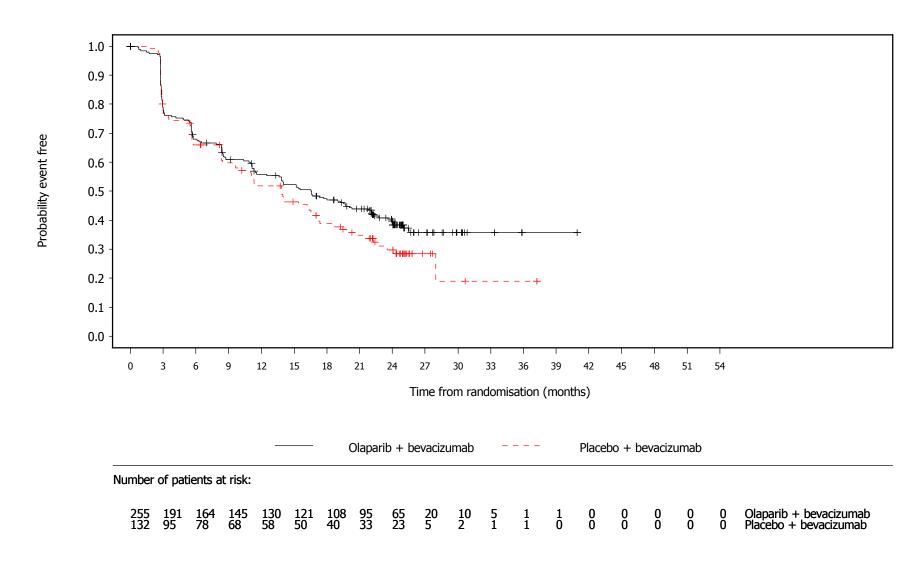
[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Figure 3.1.1 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Global QoL/health status (MID = 15) time to clinically meaningful worsening (first occurrence)

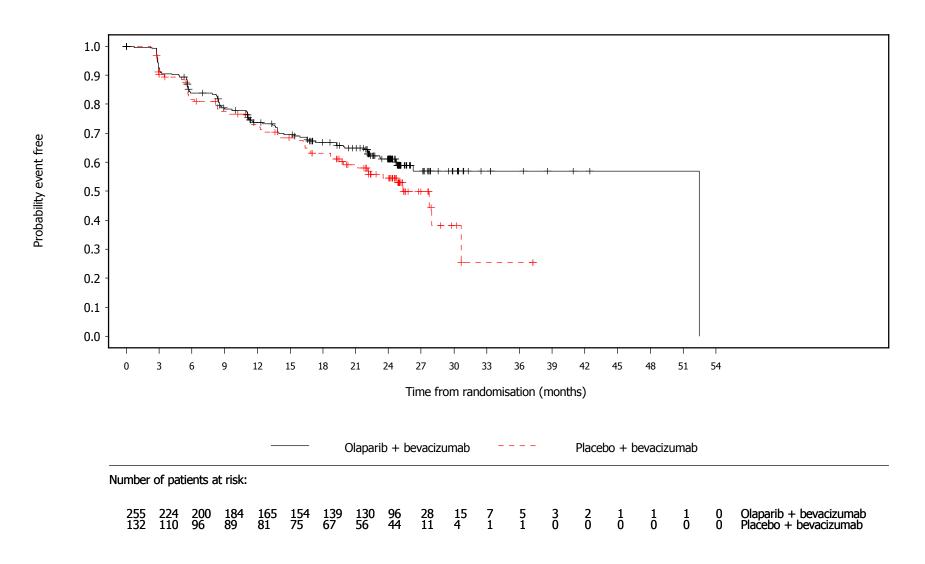
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2faa 15FEB2021:09:35 kvbv306

Figure 3.1.2 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Physical (MID = 15) time to clinically meaningful worsening (first occurrence)

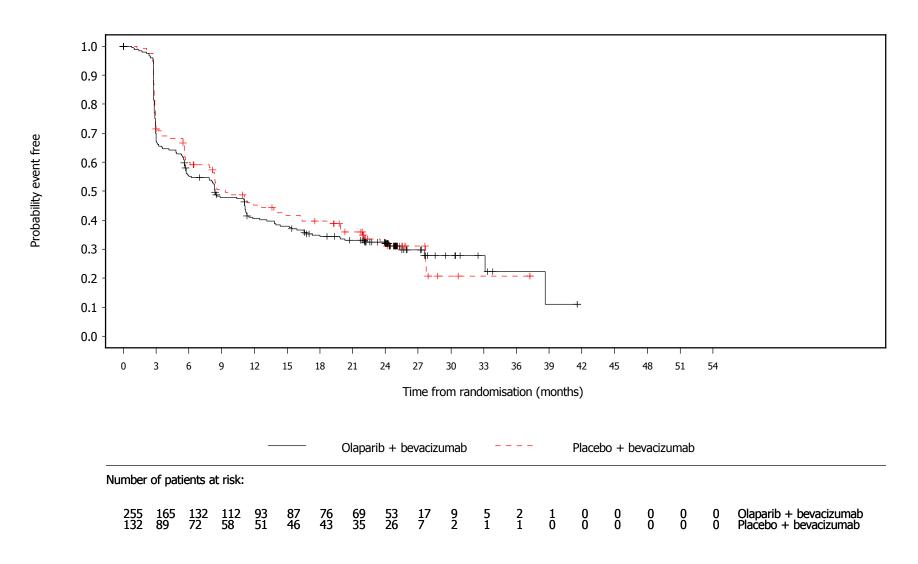
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2fab 15FEB2021:09:35 kvbv306

Figure 3.1.3 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Role (MID = 15) time to clinically meaningful worsening (first occurrence)

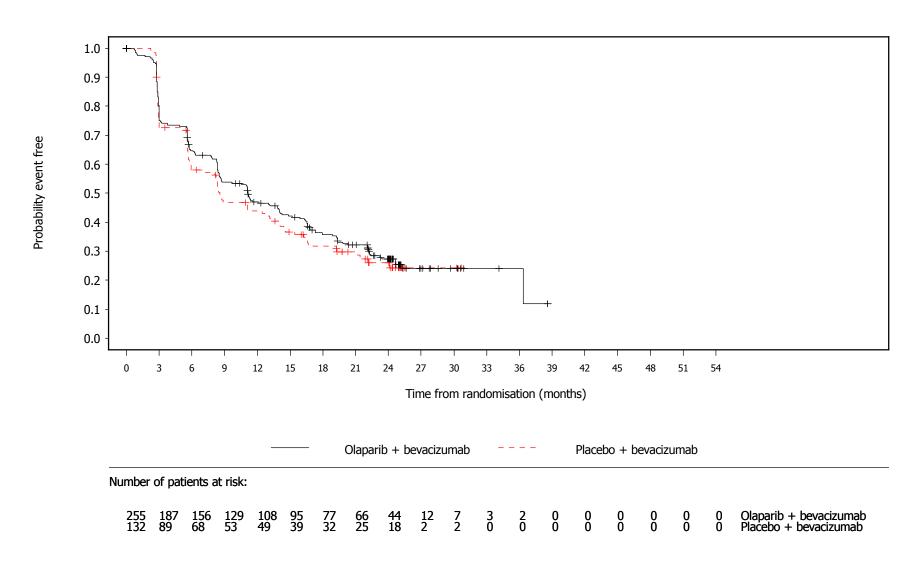
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2fac 15FEB2021:09:35 kvbv306

Figure 3.1.4 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Cognitive (MID = 15) time to clinically meaningful worsening (first occurrence)

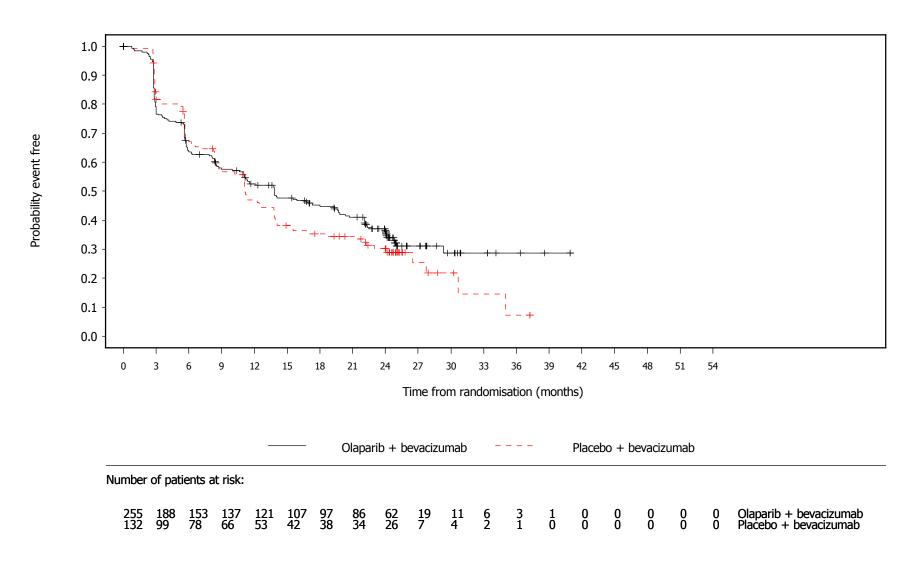
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2fad 15FEB2021:09:35 kvbv306

Figure 3.1.5 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Emotional (MID = 15) time to clinically meaningful worsening (first occurrence)

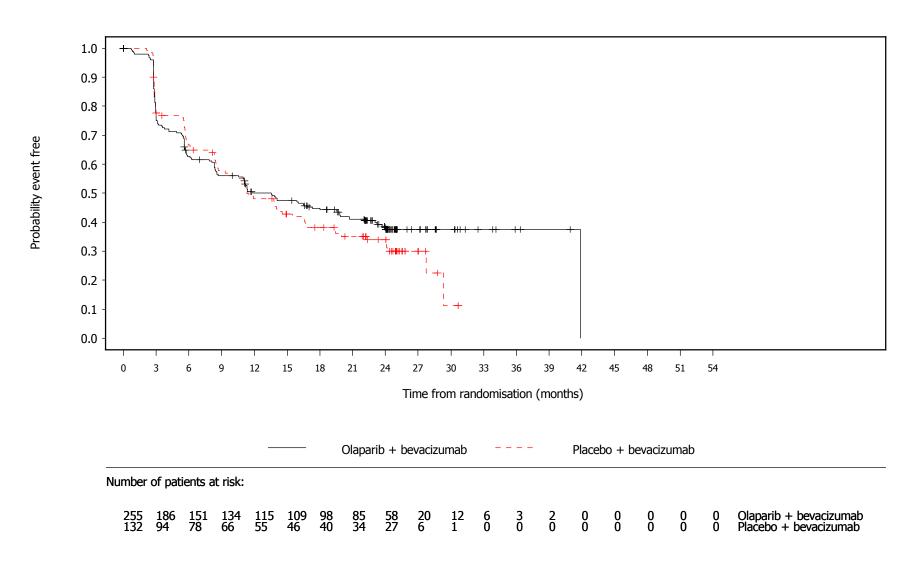
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2fae 15FEB2021:09:35 kvbv306

Figure 3.1.6 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Social (MID = 15) time to clinically meaningful worsening (first occurrence)

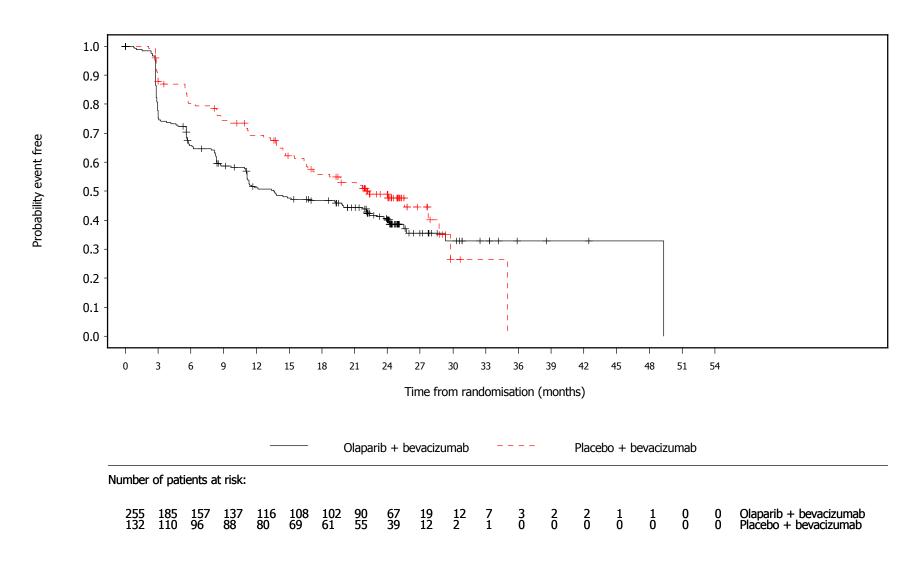
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2faf 15FEB2021:09:35 kvbv306

Figure 3.1.7 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Loss of appetite (MID = 15) time to clinically meaningful worsening (first occurrence)

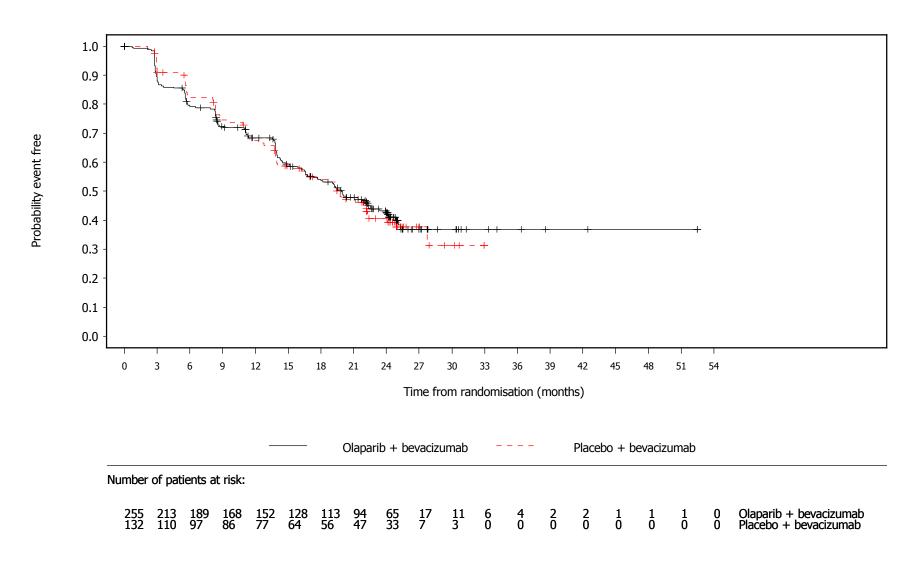
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2fag 15FEB2021:09:35 kvbv306

Figure 3.1.8 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Constipation (MID = 15) time to clinically meaningful worsening (first occurrence)

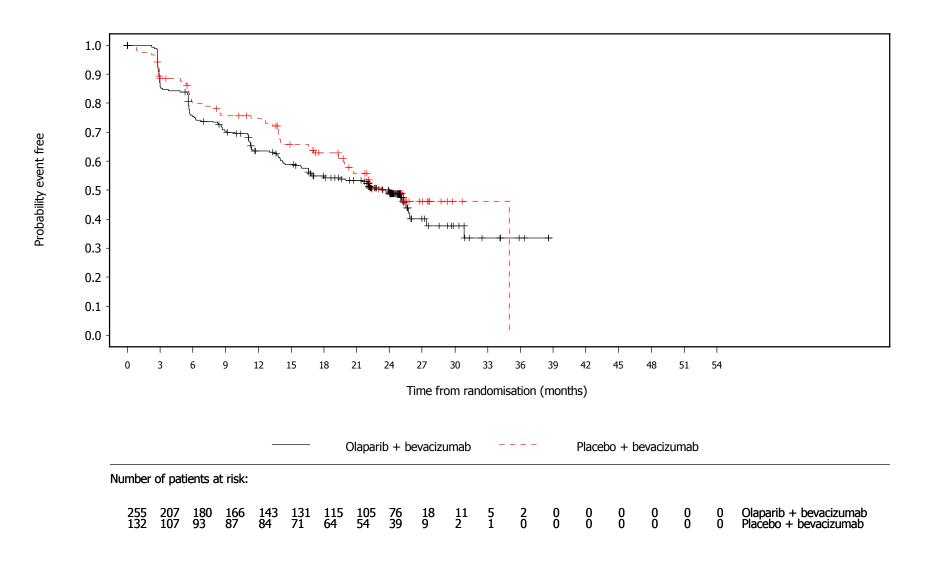
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2fah 15FEB2021:09:35 kvbv306

Figure 3.1.9 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Diarrhoea (MID = 15) time to clinically meaningful worsening (first occurrence)

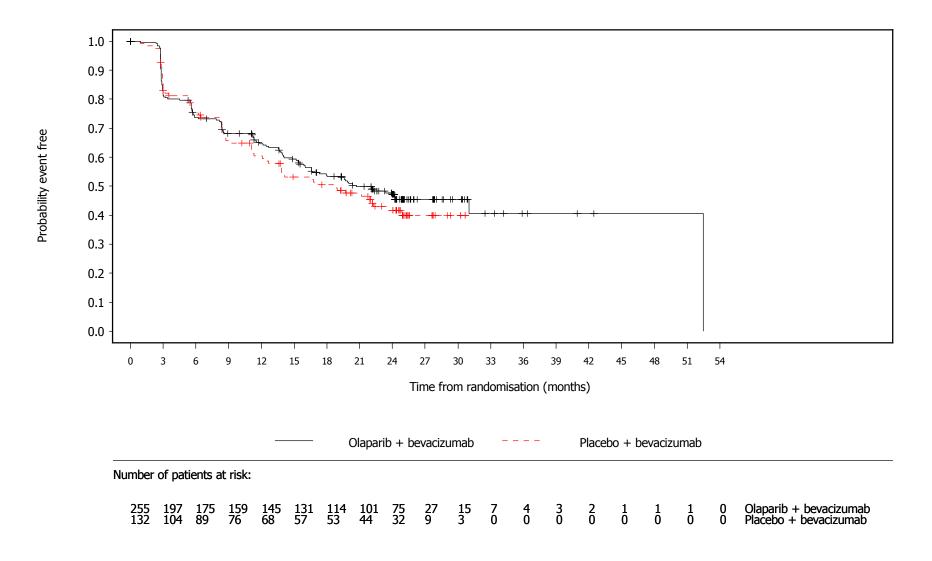
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2fai 15FEB2021:09:35 kvbv306

Figure 3.1.10 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Dyspnoea (MID = 15) time to clinically meaningful worsening (first occurrence)

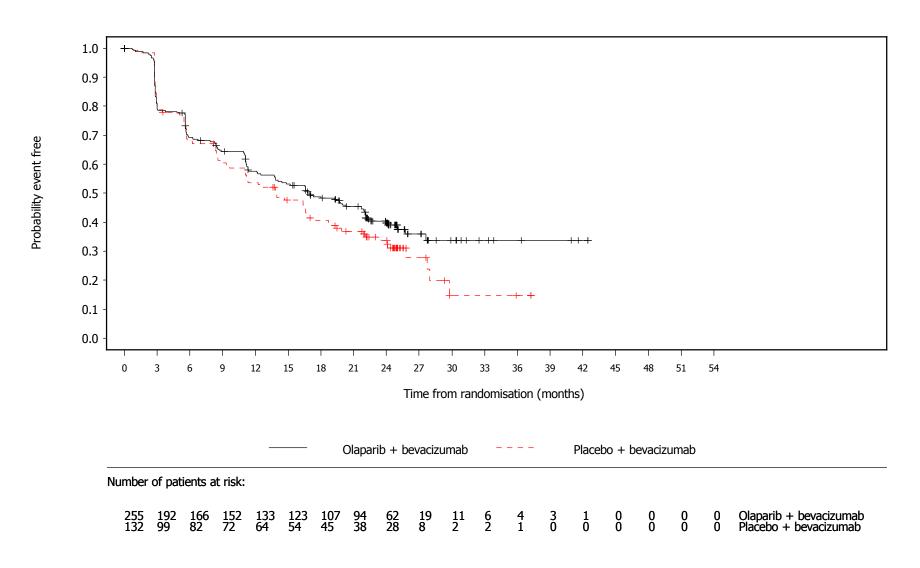
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2faj 15FEB2021:09:35 kvbv306

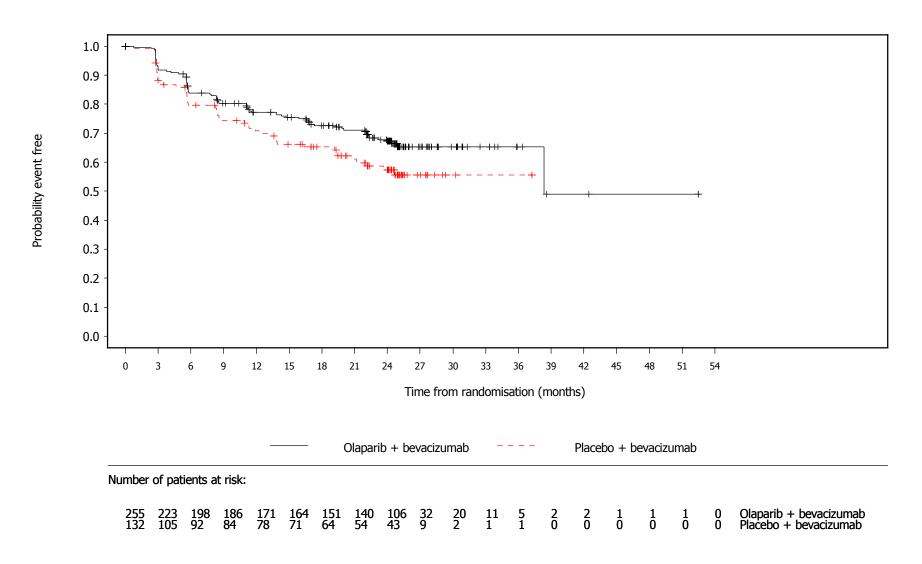
Figure 3.1.11 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Fatigue (MID = 15) time to clinically meaningful worsening (first occurrence)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2fak 15FEB2021:09:35 kvbv306

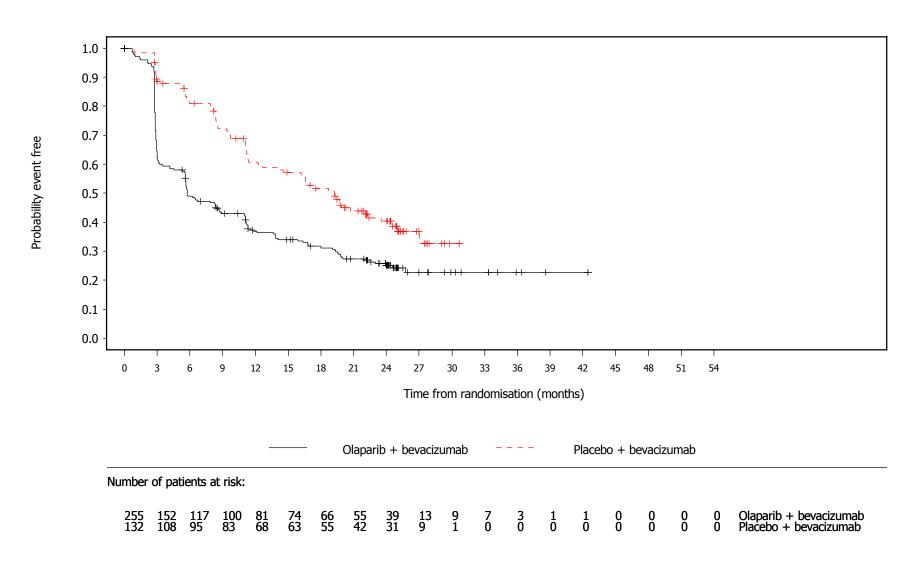
Figure 3.1.12 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Financial difficulties (MID = 15)
time to clinically meaningful worsening (first occurrence)
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2fal 15FEB2021:09:35 kvbv306

Figure 3.1.13 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Nausea and vomiting (MID = 15) time to clinically meaningful worsening (first occurrence)

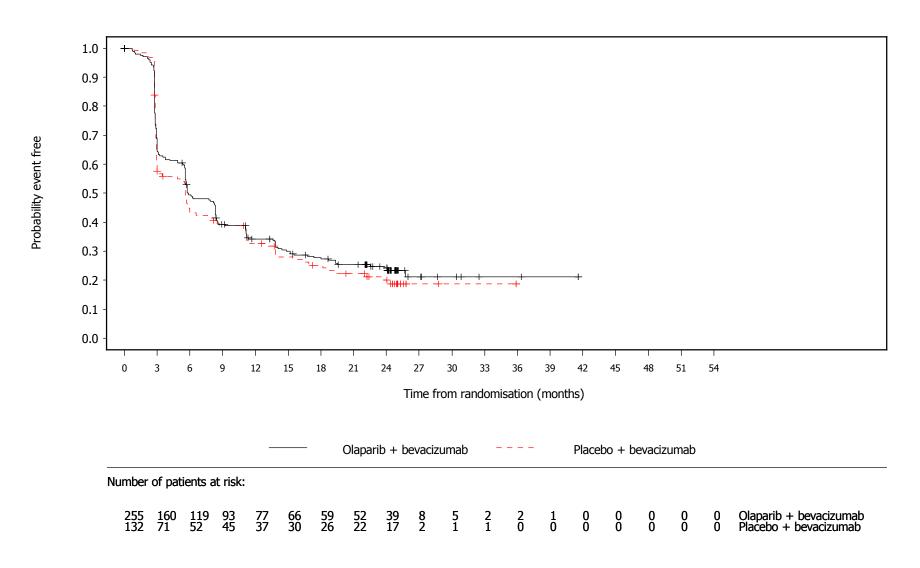
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2fam 15FEB2021:09:35 kvbv306

Figure 3.1.14 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Pain (MID = 15) time to clinically meaningful worsening (first occurrence)

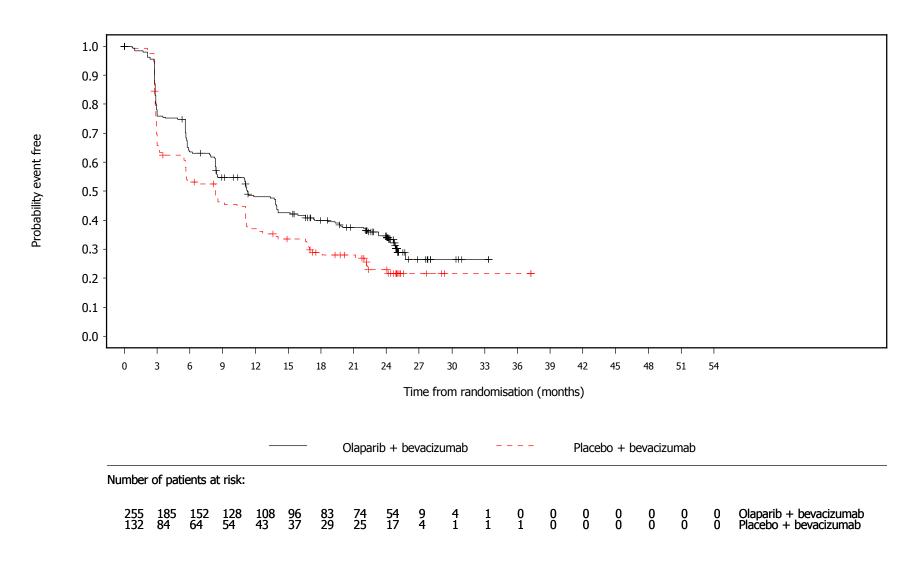
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2fan 15FEB2021:09:35 kvbv306

Figure 3.1.15 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Insomnia (MID = 15) time to clinically meaningful worsening (first occurrence)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2fao 15FEB2021:09:35 kvbv306

Table 3.2 PAOLA1 Appendix: Summary of analysis of time to worsening in EORTC QLQ-OV28 symptom and single item scales (MID=15)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevac (N=255)	zizumab	Placebo + b (N=1				
	of patients	edian time (95% CI) onths) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 15)	255 139 (54.5) 17.9	(13.8,23.1)	132 81 (61.4)	11.4 (8.3,18.7)	0.76	0.58, 1.01	0.0549
EORTC QLQ-OV28 Symptom scale/items: Body image (MID = 15)	255 126 (49.4) 21.9	(12.7, NE)	132 71 (53.8)	18.7 (11.5,25.1)	0.93	0.70, 1.26	0.6383
EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (MID = 15)	255 95 (37.3) NE	(NE, NE)	132 59 (44.7)	24.1 (17.0, NE)	0.76	0.55, 1.07	0.1095
EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (MID = 15)	255 95 (37.3) 52.	5 (NE, NE)	132 45 (34.1)	NE (NE, NE)	1.12	0.79, 1.62	0.5166
EORTC QLQ-OV28 Symptom scale/items: Hormonal (MID = 15)	255 135 (52.9) 19.1	(14.3,24.2)	132 76 (57.6)	11.3 (5.6,19.1)	0.75	0.56, 0.996	0.0462*
EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (MID = 15)	255 114 (44.7) 25.3	(18.6, NE)	132 58 (43.9)	23.0 (12.7, NE)	0.93	0.68, 1.29	0.6541
EORTC QLQ-OV28 Symptom scale/items: Other single items (MID = 15)	255 96 (37.6) 30.4	(25.3, NE)	132 45 (34.1)	28.0 (25.4, NE)	1.07	0.76, 1.55	0.7102

Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation.

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

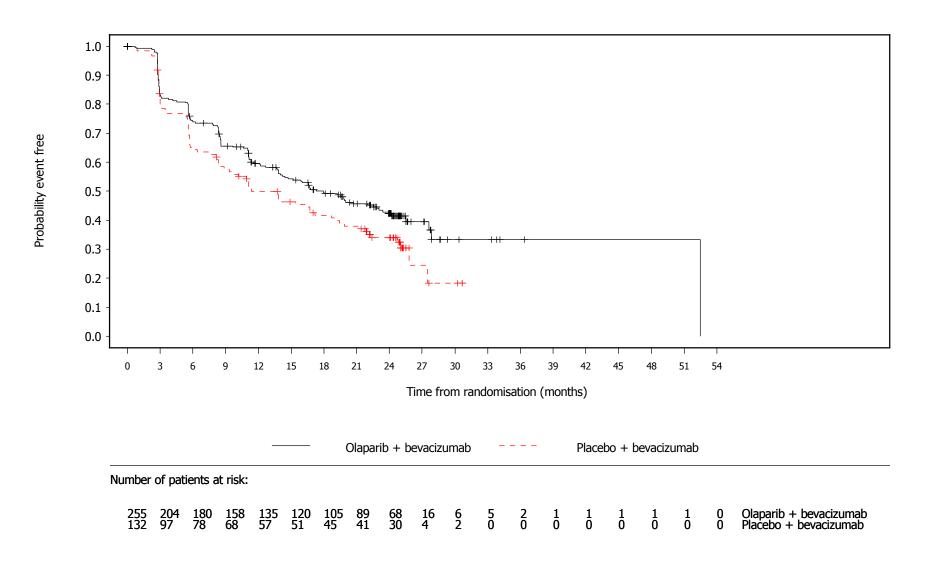
[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Figure 3.2.1 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 15) time to clinically meaningful worsening (first occurrence)

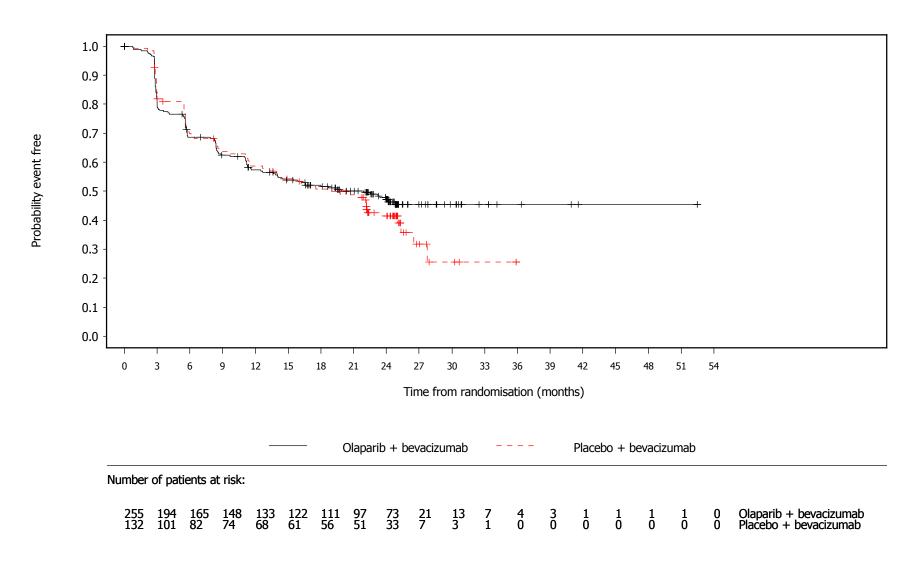
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2gaa 15FEB2021:09:35 kvbv306

Figure 3.2.2 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Body image (MID = 15) time to clinically meaningful worsening (first occurrence)

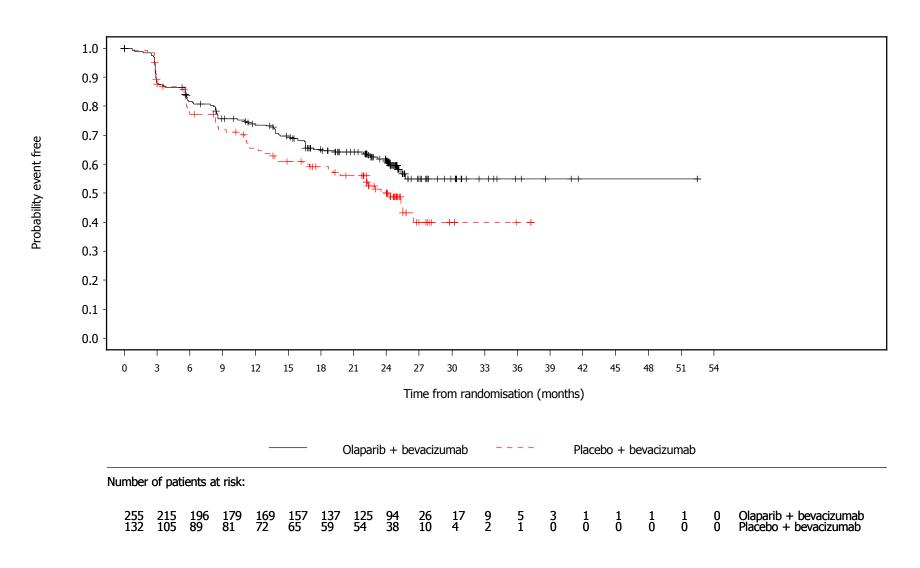
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2gab 15FEB2021:09:35 kvbv306

Figure 3.2.3 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (MID = 15) time to clinically meaningful worsening (first occurrence)

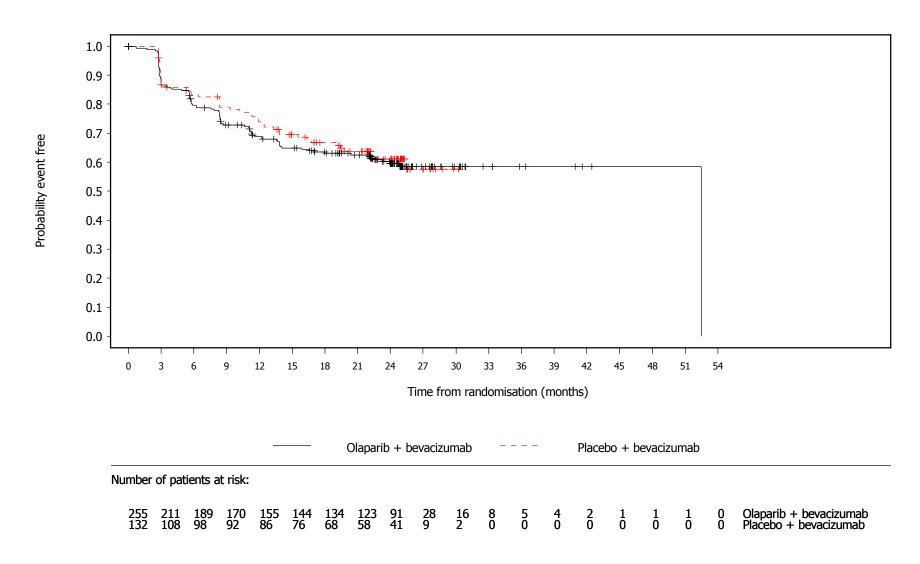
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2gac 15FEB2021:09:35 kvbv306

Figure 3.2.4 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (MID = 15) time to clinically meaningful worsening (first occurrence)

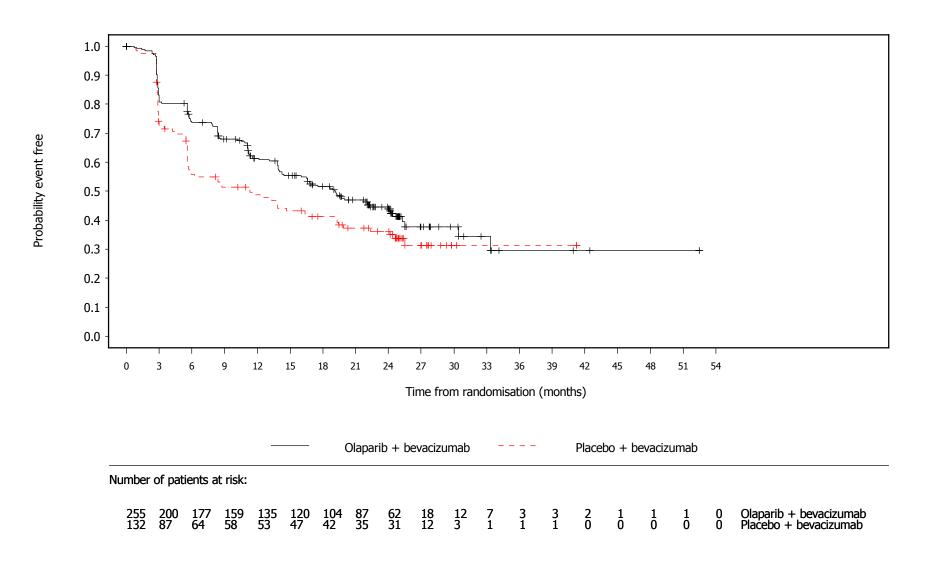
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2gad 15FEB2021:09:35 kvbv306

Figure 3.2.5 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Hormonal (MID = 15) time to clinically meaningful worsening (first occurrence)

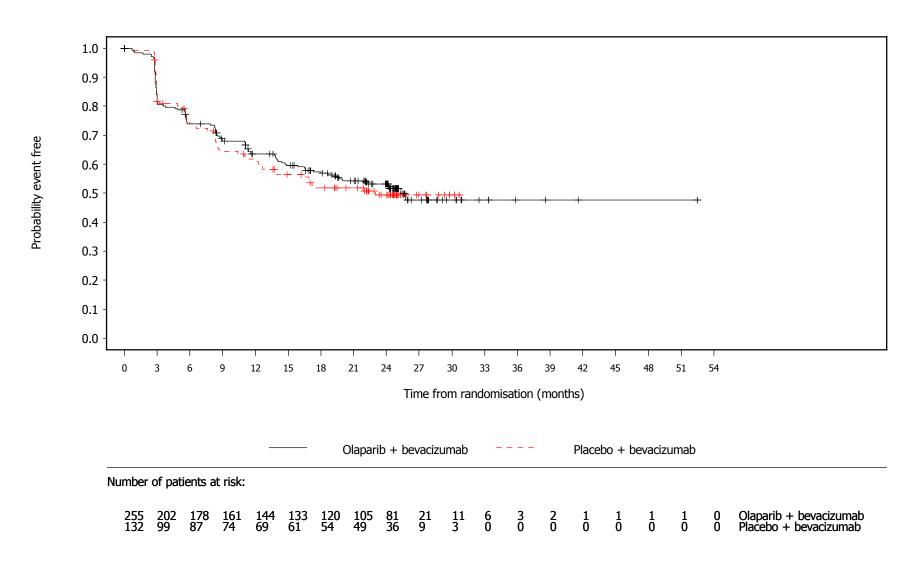
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2gae 15FEB2021:09:35 kvbv306

Figure 3.2.6 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (MID = 15) time to clinically meaningful worsening (first occurrence)

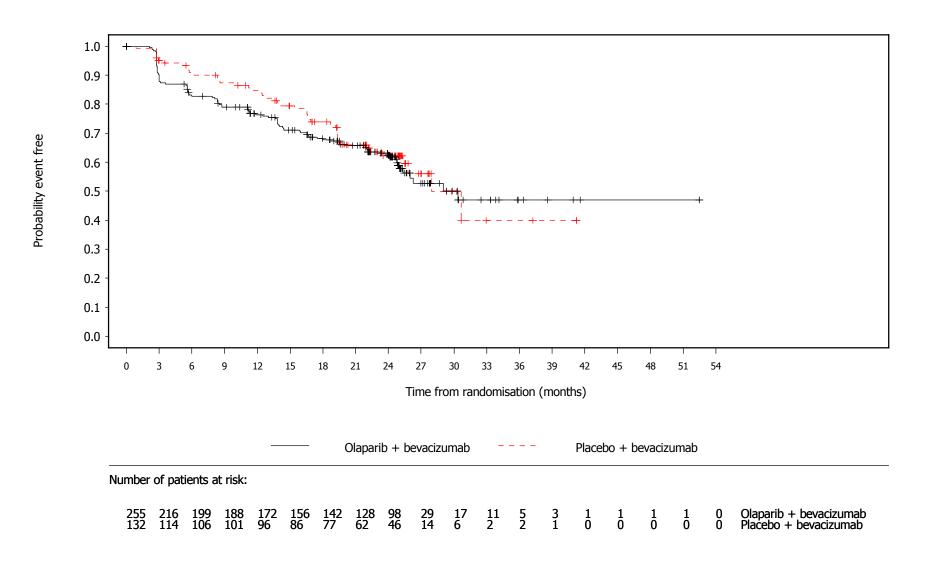
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2gaf 15FEB2021:09:35 kvbv306

Figure 3.2.7 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Other single items (MID = 15) time to clinically meaningful worsening (first occurrence)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2gag 15FEB2021:09:35 kvbv306

Table 3.3 PAOLA1 Appendix: Summary of analysis of time to worsening in EQ-5D VAS (MID = 15) Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + b (N=2		Placebo + b (N=1					
	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]	
EQ-5D-5L Visual analogue scale (MID = 15)	255 116 (45.5)	25.3 (17.5, NE)	132 58 (43.9)	26.7 (19.9, NE)	1.05	0.77, 1.46	0.7493	

Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation.

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status

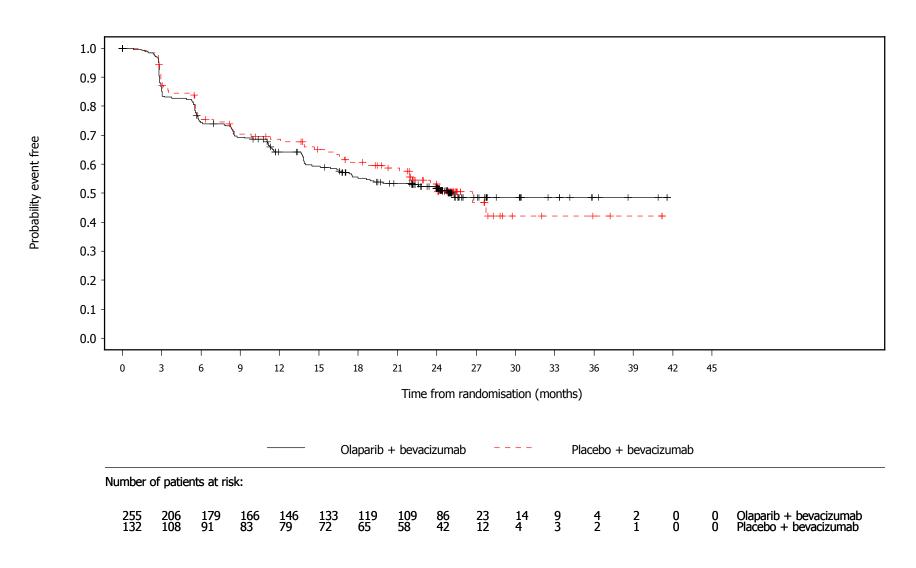
included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Figure 3.3.1 PAOLA1 Appendix: Kaplan-Meier plot of EQ-5D-5L Visual analogue scale (MID = 15) time to clinically meaningful worsening (first occurrence)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from date of randomisation until earliest date of a clinically meaningful worsening in the respective score. Patients whose score has not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at date of their last PRO assessment. Patients who died within 168 days of last PRO assessment, irrespective of baseline score, will be censored on their death date. Patients with no evaluable data at baseline will be censored at date of randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttemainpr2.sas ettemainpr2haa 15FEB2021:09:35 kvbv306

Table 2.2.3.1 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Global QoL/health status (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2				oevacizumab 132)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	TRS)								
NED [PDS]	92	49 (53.3)	20.0 (13.6, NE)	48	28 (58.3)	14.0 (5.6,22.7)	0.72	0.46,	1.16	0.1747
NED/CR [IDS]	74	46 (62.2)	11.1 (6.5,16.6)	38	20 (52.6)	13.8 (5.5, NE)	1.17	0.70,	2.02	0.5567
NED/CR [Chemo]	40	23 (57.5)	19.7 (8.3, NE)	20	15 (75.0)	9.8 (3.5,22.3)	0.68	0.36,	1.34	0.2578
PR	49	28 (57.1)	15.3 (5.6,24.0)	26	18 (69.2)	14.0 (5.6,19.9)	0.75	0.42,	1.38	0.3464
Interaction p-value										0.4751
Screening laboratory tBR0	CA statu	ıs (IVRS)								
tBRCAm	150	78 (52.0)	19.7 (14.0,25.5)	65	40 (61.5)	11.3 (8.3,21.4)	0.75	0.51,	1.11	0.1446
non-tBRCAm	105	68 (64.8)	11.4 (8.4,17.7)	67	41 (61.2)	13.9 (8.3,17.4)	0.94	0.64,	1.40	0.7568
Interaction p-value										0.4128
First line treatment out	come (eC	CRF)								
NED [PDS]	89	50 (56.2)	18.0 (11.0, NE)	47	28 (59.6)	13.8 (5.6,22.7)	0.77	0.49,	1.24	0.2767
NED/CR [IDS]	74	44 (59.5)	11.5 (6.5,22.1)	32	17 (53.1)	13.8 (5.6, NE)	1.04	0.61,	1.88	0.8796
NED/CR [Chemo]	39	19 (48.7)	22.1 (15.2, NE)	18	14 (77.8)	8.3 (3.0,17.2)	0.44	0.22,	0.90	0.0255*
PR	50	32 (64.0)	10.9 (4.9,23.6)	34	22 (64.7)	14.0 (8.3,23.5)	1.03	0.60,	1.80	0.9087
Interaction p-value										0.2199
Screening laboratory tBR0	CA statu	ıs (eCRF)								
tBRCAm	147	76 (51.7)	20.3 (13.8,25.5)	67	40 (59.7)	11.3 (8.3,22.3)	0.77	0.52,	1.13	0.1773
non-tBRCAm	108	70 (64.8)	13.6 (8.4,16.7)	65	41 (63.1)	13.9 (8.3,17.2)	0.91	0.62,	1.35	0.6255
Interaction p-value										0.5368
Age group										
<65 years	185	109 (58.9)	15.2 (11.0,19.7)	98	56 (57.1)	16.2 (9.3,20.8)	0.97	0.70,	1.34	0.8427
>=65 years	70	37 (52.9)	22.1 (11.3, NE)	34	25 (73.5)	9.9 (5.5,15.4)	0.51	0.31,	0.86	0.0126*
Interaction p-value										0.0411*

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aaa 24MAR2022:14:14 kvbv306

Table 2.2.3.1 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Global QoL/health status (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olapa	arib + (N=)	bevaci 255)	izumab			+ be	evacizumab 32)				
Subgroup	n	of pa	er (%) tients events	(dian time 95% CI) onths) [a]		Number (9 of patien with even	ts	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	105	(57.7)	16.6	(11.5,21.8)	90	54 (60.	0)	11.5 (8.3,17.4)	0.80	0.58,	1.12	0.1990
IV	73	41	(56.2)	15.6	(8.4,25.0)	42	27 (64.	3)	14.0 (8.3,21.4)	0.85	0.53,	1.40	0.5261
Interaction p-value													0.8411
Region													
Europe	245	141	(57.6)	16.6	(11.4,20.3)	126	77 (61.	1)	13.8 (9.3,17.2)	0.83	0.63,	1.10	0.1954
Japan	10	5	(50.0)	24.0	(2.8, NE)	6	4 (66.	7)	10.1 (2.8, NE)	0.60	0.16,	2.42	0.4524
Interaction p-value													0.6370
ECOG performance status at	Basel	ine											
(0) Normal activity	190	110	(57.9)	15.2	(11.1,19.7)	100	66 (66.	0)	11.2 (5.8,16.4)	0.77	0.57,	1.05	0.1012
(1) Restricted activity	61	33	(54.1)	22.1	(11.4, NE)	31	15 (48.	4)	18.6 (11.3, NE)	0.99	0.55,	1.88	0.9734
Interaction p-value													0.4725
Baseline CA-125 value													
<=ULN	228				(13.8,22.8)	118	•	,	13.9 (8.3,17.2)	0.74	0.56,		0.0440*
>ULN	27	21	(77.8)	5.9	(2.9,11.2)	14	8 (57.	1)	11.3 (5.5, NE)	1.90	0.88,	4.58	0.1065
Interaction p-value													0.0253*
Histological grade													
High grade	255	146	(57.3)	16.6	(11.5,21.8)	132	81 (61.	4)	13.8 (9.3,17.2)	0.82	0.63,	1.08	0.1557
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	95	(57.2)	15.2	(11.1,20.3)	80	45 (56.	3)	14.0 (8.3,19.3)	0.88	0.62,		0.5001
Residue	79	45	(57.0)	21.8	(11.3,25.0)	44	29 (65.	9)	12.6 (8.3,17.2)	0.76	0.48,	1.23	0.2577
Interaction p-value													0.6172

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aaa 24MAR2022:14:14 kvbv306

Table 2.2.3.1 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Global QoL/health status (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2				bevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	urgery									
Upfront	146	81 (55.5)	20.3 (15.3,25.0)	79	47 (59.5)	13.8 (8.3,19.3)	0.73	0.51,	1.06	0.0957
Interval	99	59 (59.6)	11.1 (6.1,16.6)	45	27 (60.0)	14.0 (5.6,23.5)	1.04	0.67,	1.67	0.8544
Interaction p-value										0.2306
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	81 (51.3)	21.8 (13.8, NE)	77	45 (58.4)	11.3 (8.3,21.4)	0.76	0.53,	1.10	0.1390
Non-tBRCAm	97	65 (67.0)	11.4 (8.4,17.7)	55	36 (65.5)	13.9 (8.3,17.2)	0.93	0.62,	1.41	0.7284
Interaction p-value										0.4575
Status somatic BRCA mutat	ions									
sBRCAm	22	9 (40.9)	25.5 (3.0, NE)	7	3 (42.9)	NE (NE, NE)	0.81	0.24,	3.64	0.7526
gBRCAm	66	37 (56.1)	13.8 (5.7, NE)	31	17 (54.8)	19.3 (8.3, NE)	1.10	0.63,	2.01	0.7389
Non-BRCAm	41	26 (63.4)	16.6 (8.3,22.1)	22	16 (72.7)	11.5 (3.5,16.6)	0.72	0.39,	1.38	0.3146
Interaction p-value										0.6127

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aaa 24MAR2022:14:14 kvbv306

Table 2.2.3.2 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Physical (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)				pevacizumab 132)					
Subgroup		Number (%) of patients with events	Median ti (95% CI) (months) [)		Number (%) of patients with events	Median t (95% C (months)	I)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV)	RS)										
NED [PDS]	92	29 (31.5)	52.5 (NE,	NE)	48	22 (45.8)	24.9 (14.0	, NE)	0.54	0.31,	0.96	0.0348*
NED/CR [IDS]	74	27 (36.5)	26.3 (22.1,	NE)	38	13 (34.2)	NE (NE	, NE)	1.05	0.55,	2.09	0.8933
NED/CR [Chemo]	40	18 (45.0)	22.1 (11.3,	NE)	20	8 (40.0)	NE (NE	, NE)	1.36	0.61,	3.32	0.4616
PR	49	19 (38.8)	NE (NE,	NE)	26	13 (50.0)	23.5 (12.7	, NE)	0.68	0.34,	1.42	0.2992
Interaction p-value												0.2288
Screening laboratory tBRG	CA statu	s (IVRS)										
tBRCAm	150	57 (38.0)	NE (NE,	NE)	65	27 (41.5)	NE (NE	, NE)	0.84	0.53,	1.34	0.4515
non-tBRCAm	105	36 (34.3)	52.5 (24.8,	NE)	67	29 (43.3)	25.4 (18.7	, NE)	0.73	0.45,	1.20	0.2125
Interaction p-value												0.6884
First line treatment out	come (eCl	RF)										
NED [PDS]	89	29 (32.6)	52.5 (NE,	NE)	47	21 (44.7)	27.8 (14.0	, NE)	0.59	0.34,	1.05	0.0729
NED/CR [IDS]	74	25 (33.8)	26.3 (23.2,	NE)	32	13 (40.6)	NE (NE	, NE)	0.76	0.40,	1.54	0.4348
NED/CR [Chemo]	39	19 (48.7)	21.8 (8.5,	NE)	18	8 (44.4)	19.9 (5.7	, NE)	1.16	0.52,	2.80	0.7296
PR	50	18 (36.0)	NE (NE,	NE)	34	14 (41.2)	25.4 (19.7	, NE)	0.87	0.43,	1.78	0.6921
Interaction p-value												0.5898
Screening laboratory tBR0	CA statu	s (eCRF)										
tBRCAm	147	56 (38.1)	NE (NE,	NE)	67	27 (40.3)	NE (NE	, NE)	0.86	0.55,	1.38	0.5249
non-tBRCAm	108	37 (34.3)	52.5 (24.8,	NE)	65	29 (44.6)	25.4 (18.7	, NE)	0.71	0.43,	1.16	0.1706
Interaction p-value												0.5691
Age group												
<65 years	185	67 (36.2)	NE (NE,	NE)	98	39 (39.8)	27.8 (23.5	, NE)	0.88	0.59,	1.31	0.5123
>=65 years	70	26 (37.1)	52.5 (22.1,	NE)	34	17 (50.0)	18.9 (11.1	, NE)	0.58	0.32,	1.10	0.0934
Interaction p-value												0.2787

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aab 24MAR2022:14:14 kvbv306

Table 2.2.3.2 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Physical (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	66 (36.3)	52.5 (24.8,	NE)	90	36 (40.0)	27.8 (18.7, NE)	0.80	0.54,	1.22	0.2957
IV	73	27 (37.0)	NE (NE,	NE)	42	20 (47.6)	25.4 (16.2, NE)	0.77	0.43,	1.39	0.3801
Interaction p-value											0.9090
Region											
Europe	245	92 (37.6)	52.5 (26.3,	NE)	126	54 (42.9)	25.4 (19.7,30.7)	0.79	0.57,	1.11	0.1765
Japan	10	1 (10.0)	NE (NE,	NE)	6	2 (33.3)	NE (NE, NE)	0.36	0.02,	3.75	0.3864
Interaction p-value											0.5112
ECOG performance status at	Basel	ine									
(0) Normal activity	190	73 (38.4)	52.5 (24.6,	NE)	100	44 (44.0)	27.8 (19.9,30.7)	0.85	0.59,	1.25	0.4149
(1) Restricted activity	61	18 (29.5)	NE (NE,	NE)	31	12 (38.7)	21.2 (12.3, NE)	0.58	0.28,	1.24	0.1565
Interaction p-value											0.3641
Baseline CA-125 value											
<=ULN	228	80 (35.1)	52.5 (NE,	NE)	118	49 (41.5)	27.8 (19.9, NE)	0.76	0.53,		0.1276
>ULN	27	13 (48.1)	19.9 (5.6,	NE)	14	7 (50.0)	21.2 (9.3, NE)	1.13	0.46,	3.02	0.7889
Interaction p-value											0.4150
Histological grade											
High grade	255	93 (36.5)	52.5 (26.3,	NE)	132	56 (42.4)	25.4 (19.9,30.7)	0.79	0.57,	1.10	0.1646
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	56 (33.7)	52.5 (26.3,	NE)	80	34 (42.5)	27.8 (16.6, NE)	0.68	0.44,	1.05	0.0806
Residue	79	33 (41.8)	NE (NE,	NE)	44	18 (40.9)	28.0 (19.9, NE)	1.07	0.61,	1.94	0.8244
Interaction p-value											0.2130

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aab 24MAR2022:14:14 kvbv306

Table 2.2.3.2 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Physical (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + h (N=2					oevacizumab 132)					
Subgroup		Number (%) of patients with events	Median tin (95% CI) (months) [Number (%) of patients with events)	Hazard ratio [b]	95% C	I [b]	2-sided p-value [b]
Timing of cytoreductive s	urgery											
Upfront	146	50 (34.2)	52.5 (NE,	NE)	79	34 (43.0)	27.8 (16.6,	NE)	0.67	0.43,	1.04	0.0755
Interval	99	39 (39.4)	26.3 (20.0,	NE)	45	18 (40.0)	25.4 (19.9,	NE)	1.04	0.60,	1.86	0.8971
Interaction p-value												0.2211
Myriad tumour BRCA mutati	on stat	us										
tBRCAm	158	58 (36.7)	NE (NE,	NE)	77	30 (39.0)	27.8 (19.7,	NE)	0.87	0.56,	1.36	0.5246
Non-tBRCAm	97	35 (36.1)	52.5 (23.3,	NE)	55	26 (47.3)	22.2 (16.4,	NE)	0.70	0.42,	1.17	0.1704
Interaction p-value												0.5284
Status somatic BRCA mutat	ions											
sBRCAm	22	4 (18.2)	NE (NE,	NE)	7	1 (14.3)	NE (NE,	NE)	1.22	0.18,	23.86	0.8559
gBRCAm	66	28 (42.4)	26.3 (13.9,	NE)	31	12 (38.7)	24.9 (16.2,	NE)	1.09	0.57,	2.22	0.8054
Non-BRCAm	41	13 (31.7)	52.5 (23.3,	NE)	22	8 (36.4)	NE (NE,	NE)	0.82	0.34,	2.08	0.6597
Interaction p-value												0.8662

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aab 24MAR2022:14:14 kvbv306

Table 2.2.3.3 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Role (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1				oevacizumab 132)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)								
NED [PDS]	92	58 (63.0)	11.1 (5.8,18.0)	48	29 (60.4)	11.3 (3.2,24.1)	0.95	0.61,	1.50	0.8191
NED/CR [IDS]	74	51 (68.9)	5.6 (3.0,11.1)	38	20 (52.6)	9.6 (8.3, NE)	1.70	1.03,	2.92	0.0374*
NED/CR [Chemo]	40	23 (57.5)	8.5 (3.3, NE)	20	14 (70.0)	4.6 (2.9,13.8)	0.76	0.39,	1.51	0.4181
PR	49	35 (71.4)	8.5 (4.9,13.8)	26	19 (73.1)	11.1 (2.9,19.9)	0.88	0.51,	1.57	0.6567
Interaction p-value										0.1710
Screening laboratory tBR0	CA statu	s (IVRS)								
tBRCAm	150	92 (61.3)	8.9 (5.8,16.8)	65	37 (56.9)	11.9 (5.7,27.7)	1.12	0.77,	1.66	0.5529
non-tBRCAm	105	75 (71.4)	8.3 (4.8,11.1)	67	45 (67.2)	8.3 (5.6,14.9)	1.06	0.74,	1.55	0.7386
Interaction p-value										0.8488
First line treatment out	come (eC	RF)								
NED [PDS]	89	54 (60.7)	11.1 (8.1,24.4)	47	28 (59.6)	8.5 (3.0,24.1)	0.88	0.56,	1.41	0.5802
NED/CR [IDS]	74	52 (70.3)	5.5 (2.9,11.1)	32	17 (53.1)	9.6 (8.3, NE)	1.79	1.06,	3.19	0.0297*
NED/CR [Chemo]	39	25 (64.1)	8.2 (3.0,15.2)	18	13 (72.2)	5.7 (2.9,13.8)	0.86	0.45,	1.74	0.6711
PR	50	35 (70.0)	8.9 (5.7,15.4)	34	23 (67.6)	11.1 (3.0,22.1)	0.97	0.58,	1.66	0.9064
Interaction p-value										0.1821
Screening laboratory tBR0	CA statu	s (eCRF)								
tBRCAm	147	89 (60.5)	8.9 (5.8,17.3)	67	38 (56.7)	11.3 (5.7,27.7)	1.08	0.75,	1.60	0.6790
non-tBRCAm	108	78 (72.2)	8.3 (4.8,11.1)	65	44 (67.7)	8.3 (5.6,14.9)	1.09	0.76,	1.60	0.6343
Interaction p-value										0.9713
Age group										
<65 years	185	121 (65.4)	8.5 (6.2,11.3)	98	61 (62.2)	8.5 (5.6,14.9)	1.03	0.76,	1.41	0.8594
>=65 years	70	46 (65.7)	5.8 (3.3,11.9)	34	21 (61.8)	13.8 (5.6,20.2)	1.17	0.71,	2.00	0.5526
Interaction p-value										0.6768

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aac 24MAR2022:14:14 kvbv306

Table 2.2.3.3 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Role (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			pevacizumab 132)				_
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	117 (64.3)	8.4 (5.7,11.2)	90	56 (62.2)	11.1 (5.7,19.9)	1.08	0.79,	1.50	0.6282
IV	73	50 (68.5)	10.4 (5.4,14.4)	42	26 (61.9)	8.3 (5.5,22.1)	1.02	0.64,	1.67	0.9201
Interaction p-value										0.8527
Region										
Europe	245	163 (66.5)	8.3 (5.8,11.1)	126	79 (62.7)	9.3 (5.7,16.2)	1.07	0.82,	1.40	0.6370
Japan	10	4 (40.0)	NE (NE, NE)	6	3 (50.0)	NE (NE, NE)	0.84	0.18,	4.25	0.8163
Interaction p-value										0.7558
ECOG performance status at	Basel	ine								
(0) Normal activity	190	129 (67.9)	8.4 (5.8,11.1)	100	64 (64.0)	8.5 (5.6,16.4)	1.09	0.81,	1.48	0.5740
(1) Restricted activity	61	36 (59.0)	7.2 (3.0,25.3)	31	18 (58.1)	9.6 (5.5, NE)	1.04	0.60,	1.88	0.8851
Interaction p-value										0.8931
Baseline CA-125 value										
<=ULN	228		8.4 (5.8,11.2)	118		8.5 (5.7,18.7)	1.07	0.81,		0.6454
>ULN	27	18 (66.7)	10.4 (3.5,19.9)	14	9 (64.3)	9.6 (2.8, NE)	1.03	0.47,	2.40	0.9481
Interaction p-value										0.9282
Histological grade										
High grade	255	167 (65.5)	8.4 (5.8,11.2)	132	82 (62.1)	9.3 (6.1,16.2)	1.06	0.82,	1.39	0.6480
Interaction p-value										NC
Cytoreductive surgery outco	me									
No residue	166	107 (64.5)	8.3 (5.6,11.3)	80	46 (57.5)	9.3 (8.1,21.9)	1.17	0.83,	1.66	0.3804
Residue	79	53 (67.1)	8.5 (5.8,13.8)	44	30 (68.2)	7.9 (3.5,19.9)	0.92	0.59,	1.46	0.7197
Interaction p-value										0.4156

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aac 24MAR2022:14:14 kvbv306

Table 2.2.3.3 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Role (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2				oevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	,	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery								
Upfront	146	92 (63.0)	11.1 (8.2,14.0)	79	50 (63.3)	8.2 (3.5,16.4)	0.86	0.61, 1.22	0.4003
Interval	99	68 (68.7)	5.7 (3.1,11.1)	45	26 (57.8)	9.6 (8.3, NE)	1.52	0.98, 2.42	0.0636
Interaction p-value									0.0490*
Myriad tumour BRCA mutati	on stat	us							
tBRCAm	158	100 (63.3)	8.5 (5.8,13.1)	77	45 (58.4)	11.1 (5.6,22.1)	1.06	0.75, 1.53	0.7257
Non-tBRCAm	97	67 (69.1)	8.3 (4.8,11.2)	55	37 (67.3)	8.3 (5.6,18.7)	1.08	0.73, 1.64	0.6910
Interaction p-value									0.9458
Status somatic BRCA mutat	ions								
sBRCAm	22	11 (50.0)	11.1 (3.0, NE)	7	2 (28.6)	NE (NE, NE)	1.90	0.51, 12.26	0.3718
gBRCAm	66	46 (69.7)	5.6 (3.0,12.5)	31	16 (51.6)	23.5 (8.3, NE)	1.70	0.98, 3.10	0.0583
Non-BRCAm	41	27 (65.9)	9.7 (3.0,19.9)	22	16 (72.7)	8.2 (3.5,21.9)	0.94	0.51, 1.78	0.8444
Interaction p-value									0.3462

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aac 24MAR2022:14:14 kvbv306

Table 2.2.3.4 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Cognitive (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

Subgroup	Olaparib + bevacizumab (N=255)			Placebo + bevacizumab (N=132)						
	n	Number (%) of patients with events	(95% CI)		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]	
First line treatment out	come (IV	RS)								
NED [PDS]	92	63 (68.5)	11.1 (6.3,14.0)	48	31 (64.6)	8.5 (5.6,16.4)	0.91	0.60,	1.42	0.6812
NED/CR [IDS]	74	47 (63.5)	14.8 (8.3,19.4)	38	21 (55.3)	16.3 (5.6, NE)	1.07	0.65,	1.83	0.7821
NED/CR [Chemo]	40	28 (70.0)	15.2 (8.3,19.4)	20	17 (85.0)	5.6 (2.9, 8.5)	0.52	0.29,	0.97	0.0406*
PR	49	36 (73.5)	8.4 (3.7,11.3)	26	16 (61.5)	12.7 (5.5,15.4)	1.12	0.63,	2.08	0.6976
Interaction p-value										0.2601
Screening laboratory tBR	CA statu	ıs (IVRS)								
tBRCAm	150	104 (69.3)	11.3 (8.7,16.8)	65	44 (67.7)	8.3 (5.7,13.6)	0.84	0.59,	1.20	0.3237
non-tBRCAm	105	70 (66.7)	8.4 (5.8,14.8)	67	41 (61.2)	11.1 (5.7,14.6)	1.01	0.69,	1.50	0.9397
Interaction p-value										0.4651
First line treatment out	come (eC	CRF)								
NED [PDS]	89	59 (66.3)	11.1 (6.2,14.0)	47	30 (63.8)	8.7 (5.8,16.4)	0.92	0.60,	1.45	0.7268
NED/CR [IDS]	74	48 (64.9)	11.5 (8.3,17.3)	32	19 (59.4)	14.1 (5.6,22.2)	1.03	0.62,	1.80	0.9067
NED/CR [Chemo]	39	27 (69.2)	15.2 (8.3,22.1)	18	13 (72.2)	8.2 (5.5,13.1)	0.67	0.35,	1.34	0.2472
PR	50	37 (74.0)	8.5 (4.9,13.8)	34	22 (64.7)	8.6 (3.0,15.4)	1.03	0.61,	1.77	0.9151
Interaction p-value										0.7522
Screening laboratory tBR	CA statu	ıs (eCRF)								
tBRCAm	147	102 (69.4)	11.3 (8.7,16.8)	67	45 (67.2)	8.3 (5.7,13.6)	0.83	0.59,	1.20	0.3161
non-tBRCAm	108	72 (66.7)	8.4 (5.8,14.8)	65	40 (61.5)	11.1 (5.6,16.4)	1.02	0.70,	1.51	0.9299
Interaction p-value										0.4546
Age group										
<65 years	185	127 (68.6)	11.2 (8.5,14.8)	98	65 (66.3)	8.5 (5.7,13.1)	0.89	0.67,	1.21	0.4654
>=65 years	70	47 (67.1)	10.3 (5.8,16.6)	34	20 (58.8)	12.7 (5.7,21.4)	0.97	0.58,	1.67	0.9001
Interaction p-value										0.7981

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aad 24MAR2022:14:14 kvbv306

Table 2.2.3.4 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Cognitive (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)		Placebo + k (N=1					
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	128 (70.3)	11.0 (8.3,13.8)	90	56 (62.2)	8.7 (5.7,14.4)	1.04	0.76,	1.43	0.8185
IV	73	46 (63.0)	14.8 (8.4,22.3)	42	29 (69.0)	8.3 (5.6,15.4)	0.67	0.42,	1.08	0.1014
Interaction p-value										0.1331
Region										
Europe	245	170 (69.4)	11.1 (8.4,13.8)	126	82 (65.1)	8.5 (5.8,12.7)	0.91	0.70,	1.18	0.4654
Japan	10	4 (40.0)	NE (NE, NE)	6	3 (50.0)	NE (NE, NE)	0.78	0.17,	3.94	0.7407
Interaction p-value										0.8416
ECOG performance status at	Basel	ine								
(0) Normal activity	190	125 (65.8)	11.4 (8.5,16.3)	100	67 (67.0)	8.3 (5.7,12.5)	0.81	0.60,	1.09	0.1579
(1) Restricted activity	61	46 (75.4)	8.6 (5.6,14.0)	31	18 (58.1)	14.1 (5.6,21.4)	1.31	0.77,	2.31	0.3266
Interaction p-value										0.1194
Baseline CA-125 value										
<=ULN			11.2 (8.5,14.3)	118		8.5 (5.9,13.1)	0.88	0.67,		0.3674
>ULN	27	19 (70.4)	6.1 (3.0,16.6)	14	8 (57.1)	21.2 (2.8, NE)	1.23	0.56,	2.99	0.6143
Interaction p-value										0.4413
Histological grade										
High grade	255	174 (68.2)	11.1 (8.5,14.0)	132	85 (64.4)	8.5 (5.9,13.6)	0.91	0.71,	1.19	0.4856
Interaction p-value										NC
Cytoreductive surgery outco	ome									
No residue	166	110 (66.3)	11.1 (8.3,15.9)	80	50 (62.5)	8.7 (5.8,16.4)	0.96	0.69,	1.34	0.7879
Residue	79	56 (70.9)	12.1 (8.5,16.6)	44	29 (65.9)	8.3 (5.6,13.1)	0.85	0.55,	1.35	0.4796
Interaction p-value										0.6809

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aad 24MAR2022:14:14 kvbv306

Table 2.2.3.4 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Cognitive (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			oevacizumab 132)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% C	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	100 (68.5)	11.1 (8.3,14.0)	79	52 (65.8)	8.5 (5.6,12.5)	0.86	0.62,	1.21	0.3738
Interval	99	66 (66.7)	13.0 (8.3,16.8)	45	27 (60.0)	11.1 (5.7,21.2)	1.03	0.67,	1.64	0.8939
Interaction p-value										0.5177
Myriad tumour BRCA mutati	ion stat	us								
tBRCAm	158	110 (69.6)	11.2 (8.7,16.3)	77	50 (64.9)	8.3 (5.8,14.1)	0.86	0.62,	1.21	0.3790
Non-tBRCAm	97	64 (66.0)	8.6 (5.8,15.2)	55	35 (63.6)	11.1 (5.6,14.6)	1.00	0.66,	1.52	0.9839
Interaction p-value										0.5854
Status somatic BRCA mutat	ions									
sBRCAm	22	16 (72.7)	8.5 (2.9,19.4)	7	4 (57.1)	9.5 (2.8, NE)	1.10	0.40,	3.88	0.8596
gBRCAm	66	43 (65.2)	16.8 (8.7,20.3)	31	20 (64.5)	8.7 (5.6,19.2)	0.80	0.48,	1.40	0.4260
Non-BRCAm	41	31 (75.6)	8.3 (5.6,16.6)	22	14 (63.6)	14.0 (5.6,22.2)	1.21	0.66,	2.35	0.5436
Interaction p-value										0.6004

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aad 24MAR2022:14:14 kvbv306

Table 2.2.3.5 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Emotional (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			- bevacizumab =255)			oevacizumab 132)				
Subgroup	n	Number (% of patient with event	s (95% CI)		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment outc	ome (IV	RS)								
NED [PDS]	92	52 (56.5) 14.0 (8.7,24.2)	48	30 (62.5)	11.3 (5.7,27.8)	0.84	0.54,	1.33	0.4482
NED/CR [IDS]	74	50 (67.6) 8.3 (5.6,15.6)	38	27 (71.1)	11.0 (5.9,14.1)	0.93	0.59,	1.51	0.7698
NED/CR [Chemo]	40	23 (57.5) 19.7 (11.1,25.0)	20	11 (55.0)	22.0 (8.3, NE)	1.12	0.56,	2.40	0.7485
PR	49	32 (65.3) 13.8 (5.8,22.1)	26	17 (65.4)	11.1 (2.9,14.0)	0.72	0.41,	1.34	0.2946
Interaction p-value										0.8097
Screening laboratory tBRC	!A statu	ıs (IVRS)								
tBRCAm	150	91 (60.7) 16.9 (11.1,22.1)	65	40 (61.5)	11.3 (8.3,22.2)	0.92	0.64,	1.34	0.6508
non-tBRCAm	105	66 (62.9) 11.1 (5.7,16.7)	67	45 (67.2)	11.1 (8.3,13.9)	0.89	0.61,	1.31	0.5622
Interaction p-value										0.9232
First line treatment outc	ome (eC	CRF)								
NED [PDS]	89	53 (59.6) 14.0 (8.7,22.8)	47	29 (61.7)	11.2 (5.7,27.8)	0.92	0.59,	1.46	0.7166
NED/CR [IDS]	74	48 (64.9) 10.6 (5.6,19.8)	32	23 (71.9)	11.0 (6.6,14.1)	0.86	0.53,	1.45	0.5682
NED/CR [Chemo]	39	22 (56.4) 19.6 (8.5,25.0)	18	10 (55.6)	22.0 (3.5, NE)	0.97	0.47,	2.15	0.9414
PR	50	33 (66.0) 12.1 (6.3,22.1)	34	22 (64.7)	11.3 (5.6,16.8)	0.87	0.51,	1.51	0.6059
Interaction p-value										0.9926
Screening laboratory tBRC	'A statu	ıs (eCRF)								
tBRCAm	147	90 (61.2) 15.9 (11.1,22.1)	67	41 (61.2)	11.3 (8.3,21.4)	0.93	0.65,	1.36	0.7168
non-tBRCAm	108	67 (62.0) 11.2 (5.8,19.7)	65	44 (67.7)	10.2 (8.3,13.9)	0.86	0.59,	1.26	0.4349
Interaction p-value										0.7567
Age group										
<65 years	185	121 (65.4) 11.1 (6.0,15.9)	98	65 (66.3)	11.1 (8.3,13.8)	0.95	0.71,	1.29	0.7481
>=65 years	70	36 (51.4) 22.4 (12.1, NE)	34	20 (58.8)	14.0 (9.0,26.5)	0.74	0.43,	1.31	0.2953
Interaction p-value										0.4413

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aae 24MAR2022:14:14 kvbv306

Table 2.2.3.5 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Emotional (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

				pevacizumab 55)			bevacizumab 132)				
Subgroup	n	Number (of patier with ever	ts	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% C	[b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	114 (62.	6)	13.8 (11.0,19.7)	90	56 (62.2)	11.3 (8.3,14.1	L) 0.94	0.69,	1.31	0.7154
IV	73	43 (58.	9)	12.1 (6.3,24.1)	42	29 (69.0)	11.1 (6.9,16.8	3) 0.77	0.49,	1.25	0.2932
Interaction p-value											0.5025
Region											
Europe	245	153 (62.	4)	13.8 (10.0,19.4)	126	82 (65.1)	11.1 (8.3,13.8	3) 0.86	0.66,	1.14	0.2927
Japan	10	4 (40.	0)	NE (NE, NE)	6	3 (50.0)	24.0 (13.8, NE	E) 1.09	0.24,	5.54	0.9083
Interaction p-value											0.7629
ECOG performance status at	Basel	ine									
(0) Normal activity	190	117 (61.	6)	13.8 (8.5,19.4)	100	63 (63.0)	11.2 (8.3,14.0	0.94	0.70,	1.29	0.7097
(1) Restricted activity	61	37 (60.	7)	13.8 (8.5,24.0)	31	22 (71.0)	9.0 (8.3,15.5	5) 0.71	0.42,	1.23	0.2179
Interaction p-value											0.3729
Baseline CA-125 value											
<=ULN	228			13.8 (11.0,19.6)	118		11.2 (8.7,13.9			1.18	0.4073
>ULN	27	16 (59.	3)	9.0 (5.7, NE)	14	9 (64.3)	6.9 (5.6,21.4	1) 0.87	0.39,	2.05	0.7355
Interaction p-value											0.9585
Histological grade											
High grade	255	157 (61.	6)	13.8 (11.0,19.6)	132	85 (64.4)	11.2 (8.3,13.9	0.89	0.68,	1.16	0.3716
Interaction p-value											NC
Cytoreductive surgery outco	me										
No residue	166	102 (61.	4)	11.5 (7.9,20.4)	80	53 (66.3)	11.1 (6.9,13.9	0.89	0.64,	1.24	0.4813
Residue	79	49 (62.	0)	16.7 (8.5,22.1)	44	26 (59.1)	11.3 (5.7,23.0	0.92	0.58,	1.50	0.7373
Interaction p-value											0.8968

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aae 24MAR2022:14:14 kvbv306

Table 2.2.3.5 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Emotional (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2				pevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	86 (58.9)	17.3 (11.4,22.3)	79	47 (59.5)	11.3 (8.3,23.0)	0.88	0.62,	1.27	0.4937
Interval	99	65 (65.7)	8.4 (5.7,15.6)	45	32 (71.1)	11.0 (5.9,14.0)	0.91	0.60,	1.41	0.6703
Interaction p-value										0.9085
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	97 (61.4)	15.6 (11.1,20.7)	77	47 (61.0)	11.4 (8.4,21.4)	0.93	0.66,	1.33	0.6772
Non-tBRCAm	97	60 (61.9)	11.1 (5.7,20.4)	55	38 (69.1)	9.0 (6.2,13.8)	0.84	0.56,	1.28	0.4165
Interaction p-value										0.7268
Status somatic BRCA mutat	ions									
sBRCAm	22	12 (54.5)	11.1 (5.6, NE)	7	5 (71.4)	5.6 (2.8, NE)	0.53	0.19,	1.67	0.2562
gBRCAm	66	42 (63.6)	14.0 (6.2,22.1)	31	18 (58.1)	13.9 (6.9,35.0)	1.09	0.63,	1.94	0.7657
Non-BRCAm	41	23 (56.1)	19.7 (5.7, NE)	22	15 (68.2)	11.1 (8.3,13.9)	0.76	0.40,	1.48	0.4083
Interaction p-value										0.4477

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aae 24MAR2022:14:14 kvbv306

Table 2.2.3.6 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Social (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	0]	laparib + (N=2	bevacizumab 255)			pevacizumab 132)				
Subgroup	of	umber (%) patients th events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment outcom	ne (IVRS)								
NED [PDS]	92	54 (58.7)	13.9 (8.3,23.1)	48	34 (70.8)	10.4 (6.2,14.2)	0.70	0.46,	1.08	0.1088
NED/CR [IDS]	74	47 (63.5)	11.1 (5.7,19.8)	38	21 (55.3)	13.9 (5.9,29.3)	1.27	0.77,	2.16	0.3596
NED/CR [Chemo]	40	19 (47.5)	24.0 (8.6, NE)	20	13 (65.0)	14.3 (3.5, NE)	0.72	0.36,	1.49	0.3631
PR	49	28 (57.1)	8.5 (5.3, NE)	26	13 (50.0)	13.9 (2.9, NE)	0.94	0.50,	1.88	0.8546
Interaction p-value										0.3281
Screening laboratory tBRCA	status	(IVRS)								
tBRCAm	150	86 (57.3)	15.9 (8.6,23.1)	65	37 (56.9)	14.0 (11.1,24.1)	1.01	0.69,	1.50	0.9627
non-tBRCAm	105	62 (59.0)	11.2 (5.8,19.8)	67	44 (65.7)	9.9 (5.7,14.6)	0.78	0.53,	1.16	0.2221
Interaction p-value										0.3641
First line treatment outcom	ne (eCRF)								
NED [PDS]	89	53 (59.6)	13.5 (8.3,23.1)	47	33 (70.2)	10.4 (6.2,16.4)	0.73	0.47,	1.14	0.1589
NED/CR [IDS]	74	46 (62.2)	11.1 (5.6,19.9)	32	19 (59.4)	11.5 (5.8, NE)	1.10	0.66,	1.92	0.7216
NED/CR [Chemo]	39	19 (48.7)	23.7 (8.6, NE)	18	10 (55.6)	16.7 (8.3, NE)	0.86	0.41,	1.93	0.7066
PR	50	28 (56.0)	8.5 (5.5, NE)	34	18 (52.9)	13.9 (3.0, NE)	0.99	0.55,	1.82	0.9670
Interaction p-value										0.6684
Screening laboratory tBRCA	status	(eCRF)								
tBRCAm	147	83 (56.5)	15.9 (8.5,23.2)	67	38 (56.7)	14.0 (9.3,24.1)	0.98	0.67,	1.45	0.9046
non-tBRCAm	108	65 (60.2)	11.3 (5.8,19.6)	65	43 (66.2)	9.9 (5.7,15.5)	0.81	0.55,	1.19	0.2803
Interaction p-value										0.4913
Age group										
<65 years	185 1	05 (56.8)	13.6 (8.4,19.6)	98	60 (61.2)	13.9 (8.5,19.4)	0.92	0.67,	1.27	0.5964
>=65 years	70	43 (61.4)	11.3 (6.3,23.2)	34	21 (61.8)	9.9 (5.8,15.5)	0.77	0.46,	1.33	0.3483
Interaction p-value										0.5912

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aaf 24MAR2022:14:14 kvbv306

Table 2.2.3.6 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Social (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olapar	ib + (N=2	bevacizumab 255)			bevacizumab 132)				
Subgroup	n	Number of pati with ev	ients	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	108 (5	59.3)	11.3 (8.4,18.0)	90	54 (60.0)	11.5 (8.5,20.0)	0.95	0.69,	1.32	0.7495
IV	73	40 (5	54.8)	16.0 (7.9, NE)	42	27 (64.3)	11.3 (5.8,19.4)	0.74	0.46,	1.22	0.2392
Interaction p-value											0.4186
Region											
Europe	245	145 (5	59.2)	11.4 (8.5,18.0)	126	79 (62.7)	11.2 (8.5,15.5)	0.87	0.66,	1.14	0.3071
Japan	10	3 (3	30.0)	NE (NE, NE)	6	2 (33.3)	NE (NE, NE)	1.02	0.17,	7.77	0.9800
Interaction p-value											0.8557
ECOG performance status at	Basel	ine									
(0) Normal activity	190	105 (5	55.3)	16.0 (11.2,23.1)	100	67 (67.0)	11.2 (8.3,14.6)	0.72	0.53,	0.99	0.0421*
(1) Restricted activity	61	41 (6	57.2)	5.7 (3.0,11.9)	31	14 (45.2)	19.4 (6.1, NE)	1.74	0.97,	3.31	0.0628
Interaction p-value											0.0088*
Baseline CA-125 value											
<=ULN	228			14.1 (11.1,20.7)	118		13.8 (8.7,16.7)	0.86	0.65,		0.3268
>ULN	27	19 (7	70.4)	5.7 (3.5,16.6)	14	9 (64.3)	5.6 (2.8, NE)	1.00	0.46,	2.32	0.9984
Interaction p-value											0.7354
Histological grade											
High grade	255	148 (5	58.0)	13.5 (8.6,19.6)	132	81 (61.4)	11.3 (8.5,16.4)	0.88	0.67,	1.16	0.3632
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	101 (6	50.8)	13.5 (8.3,17.3)	80	53 (66.3)	10.4 (8.3,14.2)	0.85	0.61,	1.19	0.3403
Residue	79	41 (5	51.9)	19.6 (8.4, NE)	44	24 (54.5)	16.6 (11.1, NE)	0.92	0.56,	1.54	0.7403
Interaction p-value											0.7998

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aaf 24MAR2022:14:14 kvbv306

Table 2.2.3.6 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Functional scale: Social (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + N=2				bevacizumab 132)			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery								
Upfront	146	82 (56.2)	16.0 (10.6,24.0)	79	51 (64.6)	11.3 (8.5,16.6)	0.76	0.54, 1.08	0.1257
Interval	99	60 (60.6)	11.1 (5.8,19.9)	45	26 (57.8)	13.9 (6.1,29.3)	1.12	0.71, 1.80	0.6296
Interaction p-value									0.1835
Myriad tumour BRCA mutati	ion stat	us							
tBRCAm	158	89 (56.3)	15.9 (10.6,23.2)	77	44 (57.1)	14.0 (9.3,24.1)	0.93	0.65, 1.35	0.6942
Non-tBRCAm	97	59 (60.8)	11.1 (5.6,16.6)	55	37 (67.3)	9.9 (5.7,14.6)	0.83	0.56, 1.27	0.3941
Interaction p-value									0.6998
Status somatic BRCA mutat	cions								
sBRCAm	22	8 (36.4)	NE (NE, NE)	7	1 (14.3)	NE (NE, NE)	2.94	0.54, 54.64	0.2436
gBRCAm	66	44 (66.7)	8.5 (4.2,14.1)	31	15 (48.4)	24.1 (11.3, NE)	1.79	1.02, 3.33	0.0425*
Non-BRCAm	41	26 (63.4)	11.9 (5.4, NE)	22	16 (72.7)	11.0 (5.6,15.5)	0.70	0.38, 1.35	0.2807
Interaction p-value									0.0693

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aaf 24MAR2022:14:14 kvbv306

Table 2.2.3.7 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Loss of appetite (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2				oevacizumab 132)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)								
NED [PDS]	92	44 (47.8)	25.5 (11.2, NE)	48	23 (47.9)	25.6 (16.4, NE)	1.01	0.62,	1.71	0.9650
NED/CR [IDS]	74	50 (67.6)	9.7 (6.2,19.3)	38	17 (44.7)	29.8 (13.9, NE)	2.08	1.23,	3.72	0.0059*
NED/CR [Chemo]	40	24 (60.0)	11.3 (8.1,24.2)	20	10 (50.0)	22.0 (8.7, NE)	1.51	0.74,	3.32	0.2585
PR	49	28 (57.1)	12.1 (5.8,25.7)	26	15 (57.7)	14.0 (6.4,21.7)	0.97	0.52,	1.87	0.9265
Interaction p-value										0.1822
Screening laboratory tBRG	CA statu	ıs (IVRS)								
tBRCAm	150	86 (57.3)	13.6 (10.9,23.1)	65	30 (46.2)	25.6 (16.7, NE)	1.50	1.003,	2.31	0.0484*
non-tBRCAm	105	60 (57.1)	13.8 (8.3,24.2)	67	35 (52.2)	17.4 (14.4,28.7)	1.18	0.78,	1.80	0.4476
Interaction p-value										0.4151
First line treatment out	come (eC	RF)								
NED [PDS]	89	43 (48.3)	24.2 (11.3, NE)	47	23 (48.9)	25.6 (16.4, NE)	0.99	0.60,	1.68	0.9836
NED/CR [IDS]	74	52 (70.3)	8.3 (5.6,13.6)	32	14 (43.8)	29.8 (11.3, NE)	2.39	1.36,	4.49	0.0019*
NED/CR [Chemo]	39	21 (53.8)	11.3 (8.2, NE)	18	7 (38.9)	NE (NE, NE)	1.67	0.74,	4.24	0.2214
PR	50	28 (56.0)	13.7 (5.8,25.7)	34	21 (61.8)	14.7 (8.3,21.7)	0.90	0.51,	1.60	0.7041
Interaction p-value										0.0571
Screening laboratory tBR0	CA statu	ıs (eCRF)								
tBRCAm	147	84 (57.1)	13.6 (8.7,23.1)	67	30 (44.8)	25.6 (16.7, NE)	1.53	1.02,	2.36	0.0395*
non-tBRCAm	108	62 (57.4)	13.8 (8.3,24.2)	65	35 (53.8)	17.4 (14.4,28.7)	1.15	0.76,	1.76	0.5118
Interaction p-value										0.3402
Age group										
<65 years	185	104 (56.2)	13.8 (11.0,22.1)	98	47 (48.0)	24.0 (16.7,28.7)	1.34	0.95,	1.90	0.0941
>=65 years	70	42 (60.0)	11.5 (7.9,24.2)	34	18 (52.9)	16.6 (12.7, NE)	1.28	0.75,	2.29	0.3743
Interaction p-value										0.9004

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aag 24MAR2022:14:14 kvbv306

Table 2.2.3.7 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Loss of appetite (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib (N	- bevac =255)	izumab		Placebo + N=1	oevaci 132)	izumab				
Subgroup	n	Number (% of patient with event	s	dian time (95% CI) onths) [a]		Number (%) of patients with events		edian time (95% CI) onths) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	100 (54.9) 19.2	(11.1,24.2)	90	43 (47.8)	25.6	(16.7, NE)	1.32	0.93,	1.90	0.1250
IV	73	46 (63.0) 11.2	(6.1,14.8)	42	22 (52.4)	19.7	(11.2, NE)	1.36	0.83,	2.31	0.2245
Interaction p-value												0.9137
Region												
Europe	245	140 (57.1) 13.4	(11.0,21.7)	126	61 (48.4)	22.3	(16.4,29.8)	1.33	0.99,	1.82	0.0560
Japan	10	6 (60.0) 22.1	(2.8, NE)	6	4 (66.7)	23.0	(2.8, NE)	1.15	0.33,	4.49	0.8293
Interaction p-value												0.8219
ECOG performance status at	Basel	ine										
(0) Normal activity	190	106 (55.8) 13.6	(11.0,22.1)	100	48 (48.0)	25.6	(16.6,29.8)	1.38	0.99,	1.96	0.0591
(1) Restricted activity	61	36 (59.0) 12.8	(5.6,25.7)	31	17 (54.8)	16.7	(8.4, NE)	1.10	0.63,	2.01	0.7395
Interaction p-value												0.5129
Baseline CA-125 value												
<=ULN	228	129 (56.6) 13.7	(11.1,22.4)	118	59 (50.0)	22.3	(16.4,28.7)	1.26	0.93,		0.1390
>ULN	27	17 (63.0) 11.2	(5.6,22.1)	14	6 (42.9)	21.7	(9.3, NE)	2.03	0.84,	5.62	0.1185
Interaction p-value												0.3276
Histological grade												
High grade	255	146 (57.3) 13.6	(11.1,22.1)	132	65 (49.2)	22.3	(16.6,28.7)	1.32	0.99,	1.78	0.0571
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	97 (58.4) 13.6	(9.7,22.1)	80	37 (46.3)	25.6	(17.4,29.8)	1.49	1.03,	2.21	0.0333*
Residue	79	41 (51.9) 22.1	(10.9, NE)	44	21 (47.7)	21.7	(14.6, NE)	1.18	0.70,	2.03	0.5373
Interaction p-value												0.4773

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aag 24MAR2022:14:14 kvbv306

Table 2.2.3.7 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Loss of appetite (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	,)	Hazard ratio [b]	95% C	I [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	70 (47.9)	24.2 (15.2, NE)	79	37 (46.8)	25.6 (17.4,	NE)	1.02	0.69,	1.53	0.9306
Interval	99	68 (68.7)	8.3 (5.7,11.5)	45	21 (46.7)	21.7 (14.6,	NE)	2.17	1.36,	3.63	0.0010*
Interaction p-value											0.0172*
Myriad tumour BRCA mutati	ion stat	us									
tBRCAm	158	92 (58.2)	11.5 (8.7,22.4)	77	34 (44.2)	27.8 (19.7,	NE)	1.60	1.09,	2.40	0.0160*
Non-tBRCAm	97	54 (55.7)	15.2 (8.4,25.7)	55	31 (56.4)	16.4 (12.7,	22.0)	1.02	0.66,	1.61	0.9283
Interaction p-value											0.1393
Status somatic BRCA mutat	ions										
sBRCAm	22	12 (54.5)	11.0 (2.9, NE)	7	1 (14.3)	NE (NE,	NE)	5.15	1.01,	93.99	0.0478*
gBRCAm	66	49 (74.2)	8.3 (4.6,11.3)	31	15 (48.4)	24.0 (11.3,	NE)	2.16	1.24,	3.99	0.0059*
Non-BRCAm	41	24 (58.5)	11.9 (4.8, NE)	22	14 (63.6)	14.7 (8.7,	NE)	1.03	0.54,	2.06	0.9239
Interaction p-value											0.1189

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aag 24MAR2022:14:14 kvbv306

Table 2.2.3.8 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Constipation (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2			Placebo + k				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVI	RS)							
NED [PDS]	92	42 (45.7)	NE (NE, NE)	48	27 (56.3)	19.4 (11.3,27.8)	0.72	0.44, 1.1	.8 0.1839
NED/CR [IDS]	74	46 (62.2)	19.8 (14.8,22.5)	38	19 (50.0)	19.7 (13.7, NE)	1.21	0.72, 2.1	1 0.4885
NED/CR [Chemo]	40	22 (55.0)	13.8 (8.3, NE)	20	9 (45.0)	NE (NE, NE)	1.61	0.76, 3.6	0.2177
PR	49	23 (46.9)	19.5 (14.0, NE)	26	14 (53.8)	19.9 (14.0,24.9)	0.84	0.44, 1.6	0.6017
Interaction p-value									0.2597
Screening laboratory tBR0	CA status	s (IVRS)							
tBRCAm	150	76 (50.7)	20.0 (16.3,25.3)	65	33 (50.8)	19.7 (13.7, NE)	0.99	0.67, 1.5	0.9781
non-tBRCAm	105	57 (54.3)	19.9 (13.9,24.2)	67	36 (53.7)	19.4 (13.9,24.9)	0.98	0.65, 1.5	0.9325
Interaction p-value									0.9670
First line treatment out	come (eCI	RF)							
NED [PDS]	89	41 (46.1)	NE (NE, NE)	47	26 (55.3)	19.4 (11.3,27.8)	0.78	0.48, 1.2	0.3285
NED/CR [IDS]	74	43 (58.1)	19.8 (15.9,24.0)	32	17 (53.1)	17.7 (10.8, NE)	0.97	0.57, 1.7	0.9223
NED/CR [Chemo]	39	20 (51.3)	19.2 (8.6, NE)	18	8 (44.4)	14.0 (6.0, NE)	1.22	0.56, 2.9	5 0.6276
PR	50	27 (54.0)	15.2 (13.8, NE)	34	18 (52.9)	22.2 (14.0, NE)	1.13	0.63, 2.0	0.6917
Interaction p-value									0.7305
Screening laboratory tBR0	CA status	s (eCRF)							
tBRCAm	147	74 (50.3)	20.0 (16.3,25.3)	67	33 (49.3)	19.7 (13.7, NE)	1.01	0.68, 1.5	0.9603
non-tBRCAm	108	59 (54.6)	19.8 (14.0,24.2)	65	36 (55.4)	19.4 (13.9,24.9)	0.96	0.64, 1.4	7 0.8536
Interaction p-value									0.8679
Age group									
<65 years	185	99 (53.5)	19.1 (14.8,22.5)	98	52 (53.1)	19.7 (13.8,24.9)	1.04	0.75, 1.4	7 0.8048
>=65 years	70	34 (48.6)	22.6 (16.6, NE)	34	17 (50.0)	18.7 (12.3, NE)	0.81	0.46, 1.4	9 0.4959
Interaction p-value									0.4748

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aah 24MAR2022:14:14 kvbv306

Table 2.2.3.8 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Constipation (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olapa	arib + (N=2		İzumab		Place	ebo + 1 (N=1	oevaci 132)	zumab				
Subgroup	n	of pa	er (%) atients events	(dian time 95% CI) onths) [a]		of pat	r (%) tients events		dian time (95% CI) onths) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)														
III	182	103	(56.6)	17.3	(14.0,22.1)	90	47	(52.2)	19.4	(13.7,24.1)	1.06	0.75,	1.50	0.7580
IV	73	30	(41.1)	24.9	(19.4, NE)	42	22	(52.4)	19.7	(13.8, NE)	0.77	0.45,	1.35	0.3561
Interaction p-value														0.3431
Region														
Europe	245	128	(52.2)	19.8	(16.5,23.4)	126	65	(51.6)	19.7	(15.4,24.1)	1.01	0.75,	1.36	0.9692
Japan	10	5	(50.0)	25.1	(5.6, NE)	6	4	(66.7)	8.3	(2.8, NE)	0.53	0.14,	2.15	0.3557
Interaction p-value														0.3624
ECOG performance status at	Basel	ine												
(0) Normal activity	190	100	(52.6)	19.8	(16.3,22.5)	100	51	(51.0)	22.1	(14.0,27.8)	1.09	0.78,	1.54	0.6077
(1) Restricted activity	61	30	(49.2)	23.4	(13.8, NE)	31	18	(58.1)	15.4	(11.1,21.2)	0.63	0.35,	1.15	0.1261
Interaction p-value														0.1123
Baseline CA-125 value														
<=ULN	228	118	(51.8)	20.0	(16.5,24.2)	118	63	(53.4)	19.4	(13.9,22.3)	0.92	0.68,	1.25	0.5822
>ULN	27	15	(55.6)	19.9	(11.1,24.2)	14	6	(42.9)	22.1	(8.3, NE)	1.70	0.69,	4.78	0.2543
Interaction p-value														0.2087
Histological grade														
High grade	255	133	(52.2)	19.9	(16.6,23.4)	132	69	(52.3)	19.7	(14.0,22.3)	0.98	0.73,	1.31	0.8786
Interaction p-value														NC
Cytoreductive surgery outco	ome													
No residue	166	86	(51.8)	20.0	(16.8,25.1)	80	43	(53.8)	19.4	(13.7,22.3)	0.88	0.61,	1.28	0.5011
Residue	79	43	(54.4)	15.2	(13.8,24.2)	44	23	(52.3)	22.2	(13.9, NE)	1.19	0.72,	2.00	0.5005
Interaction p-value														0.3464

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aah 24MAR2022:14:14 kvbv306

Table 2.2.3.8 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Constipation (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + N=2				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	72 (49.3)	20.0 (14.5, NE)	79	41 (51.9)	19.7 (13.9,27.8)	0.96	0.66,	1.42	0.8331
Interval	99	57 (57.6)	19.8 (16.3,22.6)	45	25 (55.6)	19.4 (11.1, NE)	0.99	0.63,	1.61	0.9674
Interaction p-value										0.9190
Myriad tumour BRCA mutat:	ion stat	us								
tBRCAm	158	80 (50.6)	19.5 (15.2,25.3)	77	37 (48.1)	21.2 (13.7, NE)	1.02	0.70,	1.53	0.9119
Non-tBRCAm	97	53 (54.6)	19.9 (14.5,24.2)	55	32 (58.2)	18.7 (13.8,24.1)	0.93	0.61,	1.46	0.7645
Interaction p-value										0.7658
Status somatic BRCA mutat	tions									
sBRCAm	22	10 (45.5)	15.2 (7.9, NE)	7	4 (57.1)	9.6 (2.8, NE)	0.73	0.24,	2.67	0.6046
gBRCAm	66	38 (57.6)	21.4 (14.5,25.1)	31	18 (58.1)	16.2 (11.3, NE)	0.91	0.53,	1.64	0.7568
Non-BRCAm	41	22 (53.7)	20.0 (13.8, NE)	22	14 (63.6)	13.8 (8.3, NE)	0.74	0.38,	1.48	0.3863
Interaction p-value										0.8729

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aah 24MAR2022:14:14 kvbv306

Table 2.2.3.9 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Diarrhoea (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2				oevacizumab 132)					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median tir (95% CI) (months) [Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
			, , , , , , , , , , , , , , , , , , , ,			, <u> </u>					
First line treatment out		,									
NED [PDS]	92	,	25.5 (16.8, NE)	48	, ,	22.3 (14.0,	NE)	0.94	0.57,		0.8216
NED/CR [IDS]	74	32 (43.2)	NE (NE, NE)	38		22.1 (13.8,	NE)	0.93	0.52,		0.8080
NED/CR [Chemo]	40	, ,	22.1 (8.3, NE)	20	, ,	25.2 (17.1,	,	1.72	0.79,		0.1744
PR	49	30 (61.2)	14.2 (11.3,21.6)	26	11 (42.3)	23.5 (8.3,	NE)	1.36	0.70,	2.84	0.3765
Interaction p-value											0.5236
Screening laboratory tBRG	CA statu	s (IVRS)									
tBRCAm	150	74 (49.3)	24.0 (16.8,27.4)	65	29 (44.6)	25.2 (20.0,	NE)	1.13	0.75,	1.77	0.5671
non-tBRCAm	105	50 (47.6)	21.6 (11.5, NE)	67	29 (43.3)	22.1 (14.0,	NE)	1.11	0.71,	1.78	0.6406
Interaction p-value											0.9611
First line treatment out	come (eCi	RF)									
NED [PDS]	89	39 (43.8)	25.9 (16.6, NE)	47	20 (42.6)	35.0 (19.4,	NE)	1.02	0.60,	1.78	0.9448
NED/CR [IDS]	74	34 (45.9)	24.0 (13.0, NE)	32	15 (46.9)	22.1 (13.8,	NE)	1.00	0.55,	1.89	0.9961
NED/CR [Chemo]	39	19 (48.7)	24.0 (8.7, NE)	18	7 (38.9)	25.2 (14.0,	NE)	1.44	0.63,	3.69	0.3984
PR	50	30 (60.0)	15.3 (11.3,25.5)	34	15 (44.1)	23.5 (11.3,	NE)	1.32	0.72,	2.53	0.3727
Interaction p-value											0.8377
Screening laboratory tBR0	CA statu	s (eCRF)									
tBRCAm	147	72 (49.0)	24.0 (16.9,27.4)	67	29 (43.3)	25.2 (20.0,	NE)	1.14	0.75,	1.78	0.5531
non-tBRCAm	108		21.6 (11.4, NE)	65		22.1 (14.0,	NE)	1.11	0.71,		0.6480
Interaction p-value											0.9406
Age group											
<65 years	185	86 (46.5)	25.3 (16.9,27.4)	98	40 (40.8)	35.0 (20.7,	NE)	1.15	0.80,	1.69	0.4605
>=65 years	70		16.6 (11.3,30.9)	34		19.9 (12.7,2		1.03	0.59,	1.84	0.9316
Interaction p-value		, -,			, , ,	, ,	,		,		0.7381

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aai 24MAR2022:14:14 kvbv306

Table 2.2.3.9 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Diarrhoea (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			bevacizumab 132)				
Subgroup	n	Number (%) of patients with events			Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	91 (50.0)	23.3 (15.9,25.9)	90	40 (44.4)	22.2 (19.8, NE	1.16	0.81,	1.70	0.4296
IV	73	33 (45.2)	25.0 (11.4, NE)	42	18 (42.9)	25.2 (13.9, NE	1.02	0.58,	1.85	0.9529
Interaction p-value										0.7081
Region										
Europe	245	119 (48.6)	24.0 (16.6,25.9)	126	54 (42.9)	25.2 (19.8, NE	1.13	0.82,	1.57	0.4584
Japan	10	5 (50.0)	16.6 (2.8, NE)	6	4 (66.7)	22.1 (5.5, NE	1.01	0.27,	4.08	0.9893
Interaction p-value										0.8716
ECOG performance status at	Basel	ine								
(0) Normal activity	190	88 (46.3)	24.0 (16.8, NE)	100	44 (44.0)	25.2 (20.7, NE	1.07	0.75,	1.55	0.7142
(1) Restricted activity	61	35 (57.4)	14.5 (8.7,25.7)	31	14 (45.2)	19.4 (12.3, NE	1.32	0.72,	2.53	0.3741
Interaction p-value										0.5665
Baseline CA-125 value										
<=ULN			25.5 (21.6,30.9)	118		25.2 (19.8, NE	,	0.75,		0.8282
>ULN	27	20 (74.1)	11.1 (5.7,16.6)	14	7 (50.0)	20.7 (5.7, NE	1.96	0.87,	5.01	0.1079
Interaction p-value										0.1626
Histological grade										
High grade	255	124 (48.6)	24.0 (16.6,25.9)	132	58 (43.9)	23.5 (19.9, NE	1.12	0.82,	1.54	0.4826
Interaction p-value										NC
Cytoreductive surgery outco	ome									
No residue	166	75 (45.2)	25.9 (16.8, NE)	80	36 (45.0)	22.2 (19.4, NE	1.00	0.68,		0.9954
Residue	79	43 (54.4)	21.9 (11.3,25.7)	44	17 (38.6)	25.2 (17.1, NE	1.46	0.85,	2.64	0.1759
Interaction p-value										0.2785

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aai 24MAR2022:14:14 kvbv306

Table 2.2.3.9 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Diarrhoea (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	71 (48.6)	24.0 (16.6,25.9)	79	32 (40.5)	25.2 (20.0,	NE)	1.23	0.82,	1.89	0.3246
Interval	99	47 (47.5)	24.0 (13.4, NE)	45	21 (46.7)	22.2 (13.9,	NE)	1.00	0.61,	1.71	NC
Interaction p-value											0.5412
Myriad tumour BRCA mutat:	ion stat	us									
tBRCAm	158	74 (46.8)	25.3 (18.0,30.9)	77	32 (41.6)	25.2 (20.0,	NE)	1.08	0.72,	1.66	0.7041
Non-tBRCAm	97	50 (51.5)	15.3 (11.1, NE)	55	26 (47.3)	22.1 (14.0,	NE)	1.20	0.76,	1.96	0.4420
Interaction p-value											0.7450
Status somatic BRCA mutat	tions										
sBRCAm	22	10 (45.5)	21.0 (5.6, NE)	7	2 (28.6)	NE (NE,	NE)	2.18	0.57, 1	14.22	0.2788
gBRCAm	66	31 (47.0)	25.3 (14.5, NE)	31	13 (41.9)	23.5 (19.4,	NE)	1.29	0.69,	2.56	0.4346
Non-BRCAm	41	24 (58.5)	14.0 (5.7, NE)	22	13 (59.1)	19.4 (12.7,	NE)	1.11	0.58,	2.25	0.7533
Interaction p-value											0.7106

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aai 24MAR2022:14:14 kvbv306

Table 2.2.3.10 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Dyspnoea (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% C	[b]	2-sided p-value [b]
First line treatment out	come (IVI	RS)									
NED [PDS]	92	39 (42.4)	52.5 (17.9, NE)	48	26 (54.2)	16.7 (8.5,	NE)	0.63	0.39,	1.06	0.0797
NED/CR [IDS]	74	41 (55.4)	17.3 (11.5,24.2)	38	19 (50.0)	18.9 (11.1,	NE)	1.15	0.67,	2.02	0.6226
NED/CR [Chemo]	40	20 (50.0)	22.1 (8.2, NE)	20	8 (40.0)	NE (NE,	NE)	1.59	0.73,	3.84	0.2522
PR	49	25 (51.0)	15.4 (11.3, NE)	26	14 (53.8)	11.1 (2.9,	NE)	0.71	0.37,	1.40	0.3087
Interaction p-value											0.1644
Screening laboratory tBR0	CA status	s (IVRS)									
tBRCAm	150	67 (44.7)	31.0 (17.3, NE)	65	34 (52.3)	17.0 (11.1,	NE)	0.76	0.51,	1.16	0.1967
non-tBRCAm	105	58 (55.2)	15.4 (11.3,23.4)	67	33 (49.3)	18.9 (11.1,	NE)	1.12	0.73,	1.73	0.6156
Interaction p-value											0.2043
First line treatment out	come (eCI	RF)									
NED [PDS]	89	37 (41.6)	52.5 (18.0, NE)	47	24 (51.1)	17.0 (8.6,	NE)	0.68	0.41,	1.15	0.1444
NED/CR [IDS]	74	41 (55.4)	14.8 (8.3,31.0)	32	17 (53.1)	18.9 (8.5,	NE)	1.13	0.65,	2.04	0.6772
NED/CR [Chemo]	39	21 (53.8)	15.2 (4.5, NE)	18	8 (44.4)	11.1 (5.6,	NE)	1.29	0.59,	3.09	0.5383
PR	50	24 (48.0)	22.3 (12.1, NE)	34	17 (50.0)	19.7 (9.0,	NE)	0.82	0.45,	1.56	0.5463
Interaction p-value											0.4539
Screening laboratory tBR0	CA status	s (eCRF)									
tBRCAm	147	64 (43.5)	31.0 (18.0, NE)	67	35 (52.2)	17.0 (11.1,	NE)	0.72	0.48,	1.10	0.1250
non-tBRCAm	108	61 (56.5)	15.3 (11.2,22.1)	65	32 (49.2)	18.9 (11.1,	NE)	1.18	0.77,	1.83	0.4562
Interaction p-value											0.1061
Age group											
<65 years	185	89 (48.1)	22.1 (16.6, NE)	98	49 (50.0)	19.7 (11.3,	NE)	0.90	0.64,	1.28	0.5527
>=65 years	70	36 (51.4)	19.7 (11.3, NE)	34	18 (52.9)	13.8 (8.4,	NE)	0.87	0.50,	1.57	0.6407
Interaction p-value											0.9293

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aaj 24MAR2022:14:14 kvbv306

Table 2.2.3.10 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Dyspnoea (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olap	arib + (N=	bevac 255)	izumab			Plac	ebo + 1 (N=	oevaci 132)	zumab					
Subgroup	n	of pa	er (%) atients events		dian ti (95% CI onths))		of pa	er (%) tients events		dian ti (95% CI onths))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)																
III	182	85	(46.7)	24.0	(17.9,	NE)	90	47	(52.2)	16.7	(11.1,	NE)	0.81	0.57,	1.17	0.2532
IV	73	40	(54.8)	14.8	(8.3,	NE)	42	20	(47.6)	21.2	(13.8,	NE)	1.13	0.67,	1.98	0.6425
Interaction p-value																0.3026
Region																
Europe	245	121	(49.4)	20.7	(16.0,	NE)	126	64	(50.8)	18.7	(12.3,	24.9)	0.90	0.66,	1.22	0.4878
Japan	10	4	(40.0)	NE	(NE,	NE)	6	3	(50.0)	NE	(NE,	NE)	0.81	0.18,	4.10	0.7814
Interaction p-value																0.8931
ECOG performance status at	Basel	ine														
(0) Normal activity	190	96	(50.5)	19.4	(15.2,	NE)	100	53	(53.0)	18.7	(11.1,	NE)	0.92	0.66,	1.30	0.6406
(1) Restricted activity	61	27	(44.3)	NE	(NE,	NE)	31	14	(45.2)	18.9	(8.5,	NE)	0.82	0.44,	1.61	0.5549
Interaction p-value																0.7549
Baseline CA-125 value																
<=ULN	228	110	(48.2)	22.3	(16.0,	NE)	118	58	(49.2)	19.7	(11.1,	NE)	0.90	0.66,	1.25	0.5363
>ULN	27	15	(55.6)	19.7	(5.6,	NE)	14	9	(64.3)	13.8	(8.4,	21.2)	0.85	0.38,	2.01	0.6936
Interaction p-value																0.8839
Histological grade																
High grade	255	125	(49.0)	20.7	(16.0,	NE)	132	67	(50.8)	18.7	(12.3,	24.9)	0.89	0.67,	1.21	0.4678
Interaction p-value																NC
Cytoreductive surgery outco	ome															
No residue	166	80	(48.2)	20.7	(16.6,	NE)	80	42	(52.5)	17.0	(12.1,	24.9)	0.85	0.59,	1.24	0.3948
Residue	79	38	(48.1)	23.4	(12.1,	NE)	44	21	(47.7)	19.7	(8.7,	NE)	0.97	0.57,	1.68	0.8997
Interaction p-value																0.6959

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aaj 24MAR2022:14:14 kvbv306

Table 2.2.3.10 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Dyspnoea (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	, ,		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	66 (45.2)	52.5 (16.7, NE)	79	40 (50.6)	18.7 (8.7,	NE)	0.79	0.53,	1.17	0.2355
Interval	99	52 (52.5)	19.9 (12.5,31.0)	45	23 (51.1)	18.9 (11.1,	NE)	1.06	0.66,	1.76	0.8160
Interaction p-value											0.3489
Myriad tumour BRCA mutat:	ion stat	us									
tBRCAm	158	73 (46.2)	24.2 (16.6, NE)	77	38 (49.4)	19.7 (12.1,	NE)	0.85	0.58,	1.27	0.4165
Non-tBRCAm	97	52 (53.6)	17.9 (12.5,24.2)	55	29 (52.7)	13.8 (8.5,	NE)	0.98	0.62,	1.56	0.9207
Interaction p-value											0.6458
Status somatic BRCA mutat	cions										
sBRCAm	22	7 (31.8)	NE (NE, NE)	7	4 (57.1)	19.5 (2.8,	NE)	0.54	0.16,	2.05	0.3379
gBRCAm	66	33 (50.0)	17.3 (11.3, NE)	31	17 (54.8)	16.7 (3.0,	NE)	0.80	0.45,	1.47	0.4661
Non-BRCAm	41	21 (51.2)	23.4 (14.8, NE)	22	10 (45.5)	NE (NE,	NE)	1.09	0.52,	2.43	0.8225
Interaction p-value											0.6141

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aaj 24MAR2022:14:14 kvbv306

Table 2.2.3.11 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Fatigue (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2				oevacizumab 132)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)								
NED [PDS]	92	52 (56.5)	16.6 (11.1, NE)	48	31 (64.6)	12.5 (8.3,24.0)	0.84	0.54,	1.33	0.4545
NED/CR [IDS]	74	43 (58.1)	16.6 (8.3,22.1)	38	21 (55.3)	16.8 (9.6, NE)	1.17	0.70,	2.01	0.5531
NED/CR [Chemo]	40	21 (52.5)	22.1 (11.2, NE)	20	12 (60.0)	11.2 (3.0, NE)	0.79	0.39,	1.66	0.5191
PR	49	28 (57.1)	16.8 (11.0,25.7)	26	20 (76.9)	8.4 (2.8,16.6)	0.46	0.26,	0.83	0.0110*
Interaction p-value										0.1392
Screening laboratory tBRG	CA statu	ıs (IVRS)								
tBRCAm	150	82 (54.7)	19.8 (12.5,24.0)	65	41 (63.1)	11.2 (5.8,22.1)	0.76	0.52,	1.11	0.1569
non-tBRCAm	105	62 (59.0)	14.8 (11.2,22.1)	67	43 (64.2)	16.4 (8.5,18.9)	0.93	0.63,	1.38	0.7009
Interaction p-value										0.4719
First line treatment out	come (eC	CRF)								
NED [PDS]	89	51 (57.3)	14.5 (11.1,22.1)	47	31 (66.0)	12.5 (8.2,22.1)	0.83	0.54,	1.31	0.4234
NED/CR [IDS]	74	43 (58.1)	16.6 (7.9,22.1)	32	17 (53.1)	13.9 (9.3, NE)	1.27	0.74,	2.30	0.3901
NED/CR [Chemo]	39	21 (53.8)	16.8 (5.8, NE)	18	10 (55.6)	16.6 (3.5, NE)	0.89	0.43,	1.98	0.7731
PR	50	28 (56.0)	22.0 (11.1,27.6)	34	26 (76.5)	13.8 (2.9,16.8)	0.50	0.29,	0.86	0.0121*
Interaction p-value										0.1214
Screening laboratory tBR0	CA statu	ıs (eCRF)								
tBRCAm	147	79 (53.7)	20.0 (13.8,24.2)	67	41 (61.2)	11.3 (5.8,23.5)	0.76	0.52,	1.11	0.1556
non-tBRCAm	108	65 (60.2)	13.8 (11.1,22.1)	65	43 (66.2)	16.4 (8.5,18.7)	0.93	0.63,	1.38	0.7119
Interaction p-value										0.4576
Age group										
<65 years	185	105 (56.8)	18.0 (13.8,22.1)	98	61 (62.2)	13.9 (8.5,22.1)	0.82	0.60,	1.13	0.2152
>=65 years	70	39 (55.7)	11.3 (8.4,27.6)	34	23 (67.6)	13.9 (8.5,18.9)	0.83	0.50,	1.42	0.4946
Interaction p-value										0.9473

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aak 24MAR2022:14:14 kvbv306

Table 2.2.3.11 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Fatigue (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olapa	arib + (N=)	bevaci 255)	izumab		Placebo		oevaci 132)	zumab				
Subgroup	n	of pa	er (%) tients events	(dian time 95% CI) onths) [a]		Number of pation with eve	ents		dian time (95% CI) onths) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)														
III	182	100	(54.9)	17.0	(11.3,22.1)	90	57 (63	3.3)	13.9	(11.1,19.9)	0.84	0.61,	1.17	0.2933
IV	73	44	(60.3)	19.3	(11.2,24.2)	42	27 (64	4.3)	13.9	(5.7,16.8)	0.78	0.49,	1.28	0.3272
Interaction p-value														0.8227
Region														
Europe	245	141	(57.6)	16.8	(12.1, 21.7)	126	80 (63	3.5)	13.8	(8.8,17.5)	0.83	0.63,	1.10	0.1852
Japan	10	3	(30.0)	NE	(NE, NE)	6	4 (66	5.7)	20.2	(8.5, NE)	0.49	0.10,	2.22	0.3450
Interaction p-value														0.4926
ECOG performance status at	Basel	ine												
(0) Normal activity	190	113	(59.5)	15.2	(11.3,19.7)	100	65 (69	5.0)	13.9	(8.5,19.4)	0.91	0.68,	1.25	0.5650
(1) Restricted activity	61	28	(45.9)	25.7	(14.5, NE)	31	19 (63	1.3)	12.3	(8.5,24.0)	0.56	0.31,	1.02	0.0576
Interaction p-value														0.1497
Baseline CA-125 value														
<=ULN	228				(12.5,22.1)	118				(8.8,18.7)	0.84	0.63,		0.2328
>ULN	27	15	(55.6)	16.6	(5.6, NE)	14	10 (7)	1.4)	11.1	(2.8,19.4)	0.69	0.31,	1.58	0.3637
Interaction p-value														0.6448
Histological grade														
High grade	255	144	(56.5)	17.0	(12.5,22.0)	132	84 (63	3.6)	13.9	(9.6,17.5)	0.82	0.63,	1.08	0.1557
Interaction p-value														NC
Cytoreductive surgery outco	ome													
No residue	166	95	(57.2)	16.6	(11.2,21.7)	80	48 (60	0.0)	13.9	(8.8,24.0)	0.99	0.70,	1.41	0.9432
Residue	79	43	(54.4)	21.9	(11.3,27.6)	44	29 (65	5.9)	16.6	(5.7,19.4)	0.67	0.42,	1.09	0.1057
Interaction p-value														0.2022

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aak 24MAR2022:14:14 kvbv306

Table 2.2.3.11 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Fatigue (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	,	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	82 (56.2)	17.0 (11.3,25.0)	79	50 (63.3)	11.3 (8.2,18.7)	0.76	0.54,	1.09	0.1362
Interval	99	56 (56.6)	16.8 (11.1,22.1)	45	27 (60.0)	16.8 (11.1,28.0)	1.06	0.68,	1.71	0.7954
Interaction p-value										0.2593
Myriad tumour BRCA mutati	ion stat	us								
tBRCAm	158	85 (53.8)	20.0 (13.8,25.0)	77	44 (57.1)	15.3 (8.5,25.8)	0.85	0.59,	1.23	0.3808
Non-tBRCAm	97	59 (60.8)	13.7 (8.5,19.7)	55	40 (72.7)	13.9 (8.5,16.6)	0.82	0.55,	1.23	0.3333
Interaction p-value										0.8976
Status somatic BRCA mutat	ions									
sBRCAm	22	8 (36.4)	NE (NE, NE)	7	3 (42.9)	NE (NE, NE)	0.64	0.18,	2.93	0.5256
gBRCAm	66	39 (59.1)	14.1 (8.5,22.4)	31	22 (71.0)	11.3 (5.7,23.5)	0.78	0.47,	1.33	0.3545
Non-BRCAm	41	22 (53.7)	19.7 (5.6, NE)	22	19 (86.4)	11.1 (5.5,14.7)	0.52	0.28,	0.98	0.0434*
Interaction p-value										0.6290

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aak 24MAR2022:14:14 kvbv306

Table 2.2.3.12 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Financial difficulties (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	1	Olaparib (1	+ beva =255)	cizumab			Plac		bevacizumab 132)					
Subgroup	c	Number (% of patient vith event	s	Median ti (95% CI Months))		of pa	er (%) atients events		()	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment outcom	e (IVF	RS)												
NED [PDS]	92	24 (26.1) N	E (NE,	NE)	48	17	(35.4)	NE (NE,	NE)	0.59	0.32,	1.11	0.1007
NED/CR [IDS]	74	24 (32.4) N	E (NE,	NE)	38	9	(23.7)	NE (NE,	NE)	1.36	0.65,	3.09	0.4205
NED/CR [Chemo]	40	14 (35.0) N	E (NE,	NE)	20	11	(55.0)	13.9 (5.7,	NE)	0.63	0.28,	1.41	0.2517
PR	49	15 (30.6) 38.	4 (22.3,	NE)	26	11	(42.3)	19.2 (4.7,	NE)	0.52	0.24,	1.15	0.1046
Interaction p-value														0.2519
Screening laboratory tBRCA	status	(IVRS)												
tBRCAm	150	50 (33.3) 38.	4 (38.4,	NE)	65	25	(38.5)	NE (NE,	NE)	0.79	0.49,	1.30	0.3424
non-tBRCAm	105	27 (25.7) N	E (NE,	NE)	67	23	(34.3)	NE (NE,	NE)	0.61	0.35,	1.08	0.0872
Interaction p-value														0.4950
First line treatment outcom	e (eCF	RF)												
NED [PDS]	89	25 (28.1) N	E (NE,	NE)	47	16	(34.0)	NE (NE,	NE)	0.68	0.37,	1.30	0.2407
NED/CR [IDS]	74	24 (32.4) N	E (NE,	NE)	32	7	(21.9)	NE (NE,	NE)	1.49	0.68,	3.75	0.3352
NED/CR [Chemo]	39	10 (25.6) N	E (NE,	NE)	18	10	(55.6)	13.3 (5.7,	NE)	0.38	0.15,	0.92	0.0326*
PR	50	17 (34.0) 38.	4 (22.3,	NE)	34	15	(44.1)	19.4 (8.8,	NE)	0.62	0.31,	1.27	0.1869
Interaction p-value														0.1401
Screening laboratory tBRCA	status	(eCRF)												
tBRCAm	147	50 (34.0) 38.	4 (38.4,	NE)	67	25	(37.3)	NE (NE,	NE)	0.83	0.52,	1.36	0.4409
non-tBRCAm	108	27 (25.0) N	E (NE,	NE)	65	23	(35.4)	NE (NE,	NE)	0.58	0.33,	1.02	0.0563
Interaction p-value														0.3381
Age group														
<65 years	185	58 (31.4) N	E (NE,	NE)	98	38	(38.8)	NE (NE,	NE)	0.72	0.48,	1.10	0.1269
>=65 years	70	19 (27.1) 38.	4 (38.4,	NE)	34	10	(29.4)	NE (NE,	NE)	0.72	0.34,	1.62	0.4114
Interaction p-value														0.9903

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aal 24MAR2022:14:14 kvbv306

Table 2.2.3.12 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Financial difficulties (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1					bevacizumab 132)					
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events	Median ti (95% CI) (months) [Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	58 (31.9)	NE (NE,	NE)	90	32 (35.6)	NE (NE,	NE)	0.77	0.50,	1.20	0.2466
IV	73	19 (26.0)	38.4 (38.4,	NE)	42	16 (38.1)	NE (NE,	NE)	0.59	0.30,	1.16	0.1236
Interaction p-value												0.5027
Region												
Europe	245	74 (30.2)	38.4 (38.4,	NE)	126	47 (37.3)	NE (NE,	NE)	0.68	0.48,	0.99	0.0450*
Japan	10	3 (30.0)	NE (NE,	NE)	6	1 (16.7)	NE (NE,	NE)	2.31	0.30,	46.77	0.4406
Interaction p-value												0.2612
ECOG performance status at	Basel	ine										
(0) Normal activity	190	58 (30.5)	38.4 (38.4,	NE)	100	38 (38.0)	NE (NE,	NE)	0.72	0.48,	1.09	0.1191
(1) Restricted activity	61	17 (27.9)	NE (NE,	NE)	31	10 (32.3)	NE (NE,	NE)	0.67	0.31,	1.51	0.3186
Interaction p-value												0.8661
Baseline CA-125 value												
<=ULN	228		38.4 (38.4,	NE)	118	42 (35.6)	NE (NE,	NE)	0.72	0.49,		0.1036
>ULN	27	8 (29.6)	NE (NE,	NE)	14	6 (42.9)	21.2 (11.3,	NE)	0.68	0.24,	2.07	0.4808
Interaction p-value												0.9153
Histological grade												
High grade	255	77 (30.2)	38.4 (38.4,	NE)	132	48 (36.4)	NE (NE,	NE)	0.72	0.50,	1.03	0.0746
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	50 (30.1)	NE (NE,	NE)	80	23 (28.8)	NE (NE,	NE)	0.94	0.58,	1.57	0.8038
Residue	79	24 (30.4)	38.4 (38.4,	NE)	44	21 (47.7)	19.4 (11.3,	NE)	0.53	0.30,	0.97	0.0399*
Interaction p-value												0.1497

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aal 24MAR2022:14:14 kvbv306

Table 2.2.3.12 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Financial difficulties (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		-	bevacizumab 255)		Placebo + bo (N=1		nab					
Subgroup	Number (%) of patients n with events		(95% CI	Median time (95% CI) (months) [a]		Number (%) of patients n with events		Median time (95% CI) (months) [a]		95% CI [b]		2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	40 (27.4)	NE (NE,	NE)	79	32 (40.5)	NE (NE, NE)	0.54	0.34,	0.86	0.0107*
Interval	99	34 (34.3)	NE (NE,	NE)	45	12 (26.7)	NE (NE, NE)	1.31	0.70,	2.64	0.4063
Interaction p-value												0.0264*
Myriad tumour BRCA mutati	on stat	us										
tBRCAm	158	54 (34.2)	38.4 (38.4,	NE)	77	31 (40.3)	NE (NE, NE)	0.72	0.47,	1.14	0.1617
Non-tBRCAm	97	23 (23.7)	NE (NE,	NE)	55	17 (30.9)	NE (NE, NE)	0.67	0.36,	1.28	0.2187
Interaction p-value												0.8434
Status somatic BRCA mutat	ions											
sBRCAm	22	7 (31.8)	38.4 (11.1,	NE)	7	3 (42.9)	NE (NE, NE)	0.63	0.17,	2.95	0.5192
gBRCAm	66	22 (33.3)	NE (NE,	NE)	31	9 (29.0)	NE (NE, NE)	1.11	0.53,	2.55	0.7828
Non-BRCAm	41	10 (24.4)	NE (NE,	NE)	22	7 (31.8)	NE (NE, NE)	0.72	0.28,	1.98	0.5072
Interaction p-value												0.6861

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aal 24MAR2022:14:14 kvbv306

Table 2.2.3.13 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Nausea and vomiting (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		bevacizumab =255)	Placeb	o + b (N=1	evacizumab 32)				
Subgroup	Number (%) of patients n with events	s (95% CI)	Number of pati n with ev	ents	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]	
First line treatment out	come (IVRS)								
NED [PDS]	92 66 (71.7)	5.7 (3.0,11.1)	48 26 (5	4.2)	16.9 (9.7, NE)	1.75	1.12,	2.80	0.0125*
NED/CR [IDS]	74 54 (73.0)	5.6 (2.9, 8.7)	38 20 (5	2.6)	19.2 (11.1, NE)	2.05	1.25,	3.51	0.0042*
NED/CR [Chemo]	40 27 (67.5)	5.7 (3.1,13.8)	20 11 (5	5.0)	19.3 (8.3, NE)	1.87	0.95,	3.94	0.0694
PR	49 31 (63.3)	11.1 (3.1,16.9)	26 13 (5	0.0)	19.9 (9.7, NE)	1.53	0.82,	3.02	0.1899
Interaction p-value									0.9149
Screening laboratory tBRG	CA status (IVRS)								
tBRCAm	150 104 (69.3)	5.8 (3.5,11.0)	65 34 (5	2.3)	19.7 (11.4,27.0)	1.90	1.31,	2.84	0.0006*
non-tBRCAm	105 74 (70.5)	5.7 (5.4,11.1)	67 36 (5	3.7)	17.4 (9.7,25.0)	1.72	1.16,	2.59	0.0062*
Interaction p-value									0.7212
First line treatment out	come (eCRF)								
NED [PDS]	89 65 (73.0)	5.6 (2.9,11.0)	47 25 (5	3.2)	17.4 (9.7, NE)	1.90	1.21,	3.06	0.0047*
NED/CR [IDS]	74 53 (71.6)	5.6 (2.9, 8.7)	32 19 (5	9.4)	16.2 (9.6,24.4)	1.74	1.05,	3.01	0.0322*
NED/CR [Chemo]	39 25 (64.1)	7.1 (4.5,16.8)	18 9 (5	0.0)	19.3 (11.1, NE)	1.84	0.89,	4.16	0.1034
PR	50 32 (64.0)	11.0 (3.0,19.4)	34 17 (5	0.0)	19.9 (9.7, NE)	1.57	0.88,	2.89	0.1256
Interaction p-value									0.9678
Screening laboratory tBR0	CA status (eCRF)								
tBRCAm	147 102 (69.4)	5.8 (3.4,11.0)	67 34 (5	0.7)	20.7 (14.3, NE)	1.95	1.34,	2.91	0.0004*
non-tBRCAm	108 76 (70.4)	5.7 (5.5,11.1)	65 36 (5	5.4)	16.6 (9.7,24.4)	1.67	1.13,	2.51	0.0093*
Interaction p-value									0.5883
Age group									
<65 years	185 133 (71.9)	5.7 (3.2, 8.4)	98 54 (5	5.1)	19.2 (11.2,24.4)	1.81	1.32,	2.50	0.0002*
>=65 years	70 45 (64.3)	11.1 (5.6,14.1)	34 16 (4	7.1)	19.9 (11.3, NE)	1.83	1.06,	3.34	0.0303*
Interaction p-value									0.9661

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aam 24MAR2022:14:14 kvbv306

Table 2.2.3.13 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Nausea and vomiting (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			bevacizumab 132)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	125 (68.7)	6.2 (5.6,11.1)	90	46 (51.1)	19.7 (12.7,25.0)	1.87	1.35,	2.65	0.0002*
IV	73	53 (72.6)	5.6 (3.0,11.1)	42	24 (57.1)	16.6 (8.3, NE)	1.69	1.05,	2.78	0.0294*
Interaction p-value										0.7267
Region										
Europe	245	171 (69.8)	5.8 (5.5, 8.7)	126	66 (52.4)	18.7 (11.4,23.5)	1.80	1.36,	2.41	<0.0001*
Japan	10	7 (70.0)	5.7 (2.8, NE)	6	4 (66.7)	23.6 (8.1, NE)	1.83	0.55,	7.00	0.3252
Interaction p-value										0.9780
ECOG performance status at	Basel	ine								
(0) Normal activity	190	140 (73.7)	5.7 (4.2, 8.3)	100	55 (55.0)	18.7 (11.3,25.0)	2.02	1.49,	2.79	<0.0001*
(1) Restricted activity	61	36 (59.0)	10.1 (4.5,25.7)	31	15 (48.4)	19.7 (11.1, NE)	1.33	0.74,	2.51	0.3406
Interaction p-value										0.2385
Baseline CA-125 value										
<=ULN	228		5.7 (5.5, 8.6)	118		18.7 (12.3,22.3)	1.74	1.31,		<0.0001*
>ULN	27	17 (63.0)	11.1 (3.0,20.0)	14	5 (35.7)	NE (NE, NE)	2.55	1.01,	7.76	0.0482*
Interaction p-value										0.4614
Histological grade										
High grade	255	178 (69.8)	5.8 (5.6, 8.7)	132	70 (53.0)	19.2 (12.7,23.5)	1.80	1.37,	2.39	<0.0001*
Interaction p-value										NC
Cytoreductive surgery outco	ome									
No residue	166	121 (72.9)	5.6 (3.1, 8.4)	80	44 (55.0)	17.4 (11.1,22.1)	1.85	1.32,	2.65	0.0003*
Residue	79	52 (65.8)	8.3 (5.6,12.1)	44	23 (52.3)	18.7 (11.1, NE)	1.62	1.005,	2.70	0.0478*
Interaction p-value										0.6613

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aam 24MAR2022:14:14 kvbv306

Table 2.2.3.13 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Nausea and vomiting (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			bevacizumab 132)			
Subgroup	n	Number (%) of patients with events			Number (%) of patients with events		Hazard ratio [b]	95% CI [2-sided b] p-value[b]
Timing of cytoreductive s	surgery								
Upfront	146	105 (71.9)	5.8 (4.5,11.1)	79	41 (51.9)	18.7 (11.2, NE)	1.82	1.28, 2	.64 0.0008*
Interval	99	68 (68.7)	5.7 (3.1,11.1)	45	26 (57.8)	16.2 (11.1,24.4)	1.72	1.11, 2	.75 0.0152*
Interaction p-value									0.8490
Myriad tumour BRCA mutati	on stat	tus							
tBRCAm	158	109 (69.0)	5.8 (3.5,11.0)	77	37 (48.1)	22.1 (16.6, NE)	2.05	1.43, 3	.02 <0.0001*
Non-tBRCAm	97	69 (71.1)	5.7 (5.4,11.1)	55	33 (60.0)	12.7 (8.5,19.9)	1.52	1.01, 2	.33 0.0424*
Interaction p-value									0.2939
Status somatic BRCA mutat	ions								
sBRCAm	22	11 (50.0)	13.8 (2.8, NE)	7	4 (57.1)	22.6 (2.8, NE)	1.02	0.35, 3	.69 0.9713
gBRCAm	66	48 (72.7)	3.8 (2.9, 6.2)	31	18 (58.1)	19.2 (11.3,27.0)	1.90	1.13, 3	.36 0.0156*
Non-BRCAm	41	33 (80.5)	5.6 (2.8, 8.3)	22	14 (63.6)	12.3 (8.1, NE)	1.91	1.04, 3	.70 0.0352*
Interaction p-value									0.6275

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aam 24MAR2022:14:14 kvbv306

Table 2.2.3.14 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Pain (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab -255)		Placebo + k					
Subgroup	n	Number (%) of patients with events	(95% CI)		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment outc	ome (IV	TRS)								
NED [PDS]	92	66 (71.7)	8.3 (5.6,11.2)	48	33 (68.8)	3.1 (2.9,11.5)	0.86	0.57,	1.33	0.4991
NED/CR [IDS]	74	58 (78.4)	5.6 (2.9, 5.9)	38	28 (73.7)	5.6 (2.9, 7.6)	1.12	0.72,	1.78	0.6241
NED/CR [Chemo]	40	29 (72.5)	5.6 (3.0,11.2)	20	13 (65.0)	7.3 (3.0, NE)	1.34	0.71,	2.67	0.3711
PR	49	30 (61.2)	8.3 (5.5,19.5)	26	21 (80.8)	5.6 (2.8,12.9)	0.55	0.31,	0.97	0.0382*
Interaction p-value										0.1467
Screening laboratory tBRC	A statı	ıs (IVRS)								
tBRCAm	150	105 (70.0	5.8 (4.2, 8.4)	65	45 (69.2)	6.0 (4.9,13.8)	1.07	0.76,	1.53	0.7137
non-tBRCAm	105	78 (74.3)	5.9 (5.6, 8.4)	67	50 (74.6)	3.0 (2.9, 6.0)	0.77	0.55,	1.11	0.1652
Interaction p-value										0.2083
First line treatment outc	ome (e0	CRF)								
NED [PDS]	89	66 (74.2)	8.1 (5.3,11.1)	47	33 (70.2)	3.0 (2.9,11.2)	0.90	0.60,	1.38	0.6175
NED/CR [IDS]	74	59 (79.7)	5.5 (2.8, 5.9)	32	24 (75.0)	5.7 (2.9, 8.1)	1.23	0.77,	2.01	0.3920
NED/CR [Chemo]	39	25 (64.1)	8.5 (3.0,15.2)	18	11 (61.1)	8.7 (3.5, NE)	1.05	0.53,	2.22	0.8981
PR	50	32 (64.0)	8.3 (5.5,18.7)	34	26 (76.5)	5.6 (2.9,12.9)	0.65	0.39,	1.10	0.1050
Interaction p-value										0.3445
Screening laboratory tBRC	A statı	ıs (eCRF)								
tBRCAm	147	102 (69.4)	5.8 (4.2, 8.4)	67	46 (68.7)	5.9 (4.9,13.8)	1.05	0.74,	1.50	0.7960
non-tBRCAm	108	81 (75.0)	5.9 (5.6, 8.4)	65	49 (75.4)	3.0 (2.9, 8.1)	0.79	0.55,	1.13	0.1890
Interaction p-value										0.2586
Age group										
<65 years	185	138 (74.6)	5.8 (4.9, 8.3)	98	70 (71.4)	5.6 (3.0, 8.7)	0.97	0.73,	1.31	0.8627
>=65 years	70	45 (64.3)	5.8 (5.6,11.1)	34	25 (73.5)	5.6 (2.9,11.1)	0.75	0.47,	1.25	0.2664
Interaction p-value										0.3795

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aan 24MAR2022:14:14 kvbv306

Table 2.2.3.14 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Pain (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k				bevacizumab 132)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI [b]	[b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	127 (69.8)	7.8 (5.6, 8.5)	90	67 (74.4)	5.6 (3.0, 7.6)	0.81	0.60,	1.09	0.1571
IV	73	56 (76.7)	5.6 (3.0, 8.3)	42	28 (66.7)	5.6 (3.0,13.9)	1.23	0.79,	1.97	0.3571
Interaction p-value										0.1194
Region										
Europe	245	179 (73.1)	5.8 (5.6, 8.3)	126	91 (72.2)	5.6 (3.0, 7.6)	0.93	0.72,	1.20	0.5599
Japan	10	4 (40.0)	NE (NE, NE)	6	4 (66.7)	13.5 (2.8, NE)	0.51	0.12,	2.16	0.3469
Interaction p-value										0.4102
ECOG performance status at	Basel	ine								
(0) Normal activity	190	140 (73.7)	5.7 (5.6, 8.3)	100	73 (73.0)	5.6 (3.0, 7.6)	0.94	0.71,	1.26	0.6922
(1) Restricted activity	61	40 (65.6)	8.3 (3.0,14.5)	31	22 (71.0)	5.6 (3.0,16.9)	0.85	0.51,	1.45	0.5429
Interaction p-value										0.7276
Baseline CA-125 value										
<=ULN	228	166 (72.8)	6.2 (5.6, 8.4)	118	86 (72.9)	5.6 (3.0, 8.1)	0.92	0.71,		0.5313
>ULN	27	17 (63.0)	5.6 (3.0, NE)	14	9 (64.3)	6.6 (2.8, NE)	0.85	0.39,	2.00	0.6954
Interaction p-value										0.8553
Histological grade										
High grade	255	183 (71.8)	5.8 (5.6, 8.3)	132	95 (72.0)	5.6 (3.0, 8.1)	0.91	0.71,	1.17	0.4717
Interaction p-value										NC
Cytoreductive surgery outco	me									
No residue	166	126 (75.9)	5.7 (4.2, 8.3)	80	58 (72.5)	5.5 (3.0, 7.6)	0.99	0.73,	1.36	0.9652
Residue	79	50 (63.3)	8.5 (5.7,15.2)	44	30 (68.2)	6.0 (3.0,11.2)	0.78	0.50,	1.23	0.2779
Interaction p-value										0.3806

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aan 24MAR2022:14:14 kvbv306

Table 2.2.3.14 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Symptom scale: Pain (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2			Placebo + b (N=1				
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	urgery								
Upfront	146	102 (69.9)	8.4 (5.7,11.2)	79	53 (67.1)	5.7 (3.0,11.1)	0.90	0.65, 1.27	0.5516
Interval	99	74 (74.7)	5.6 (3.2, 6.3)	45	35 (77.8)	5.6 (2.9, 8.1)	0.94	0.63, 1.42	0.7522
Interaction p-value									0.8914
Myriad tumour BRCA mutati	on stat	us							
tBRCAm	158	111 (70.3)	6.2 (4.9, 8.4)	77	50 (64.9)	6.0 (5.2,13.8)	1.11	0.80, 1.57	0.5234
Non-tBRCAm	97	72 (74.2)	5.8 (5.6, 8.3)	55	45 (81.8)	2.9 (2.9, 5.8)	0.68	0.47, 0.998	0.0488*
Interaction p-value									0.0557
Status somatic BRCA mutat	ions								
sBRCAm	22	14 (63.6)	7.9 (2.9,19.4)	7	3 (42.9)	NE (NE, NE)	1.63	0.53, 7.09	0.4207
gBRCAm	66	48 (72.7)	4.9 (3.0,11.1)	31	21 (67.7)	11.5 (5.8,22.2)	1.26	0.76, 2.15	0.3716
Non-BRCAm	41	28 (68.3)	5.9 (4.9,14.8)	22	20 (90.9)	2.8 (2.8, 2.9)	0.39	0.22, 0.70	0.0021*
Interaction p-value									0.0062*

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aan 24MAR2022:14:14 kvbv306

Table 2.2.3.15 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Insomnia (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2				pevacizumab 132)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)								
NED [PDS]	92	57 (62.0)	11.0 (5.8,16.6)	48	36 (75.0)	5.6 (3.0, 9.2)	0.60	0.40,	0.92	0.0199*
NED/CR [IDS]	74	41 (55.4)	14.0 (7.9, NE)	38	26 (68.4)	8.5 (3.1,21.2)	0.72	0.44,	1.18	0.1886
NED/CR [Chemo]	40	29 (72.5)	11.3 (5.7,21.9)	20	15 (75.0)	7.8 (2.9,22.0)	0.89	0.48,	1.70	0.7120
PR	49	32 (65.3)	11.3 (5.8,22.3)	26	14 (53.8)	11.1 (5.6, NE)	1.05	0.57,	2.04	0.8717
Interaction p-value										0.4678
Screening laboratory tBR0	CA statu	ıs (IVRS)								
tBRCAm	150	91 (60.7)	11.3 (8.4,16.6)	65	43 (66.2)	10.3 (5.6,13.9)	0.85	0.60,	1.23	0.3873
non-tBRCAm	105	68 (64.8)	11.2 (8.3,16.6)	67	48 (71.6)	5.7 (3.0,11.1)	0.67	0.46,	0.97	0.0351*
Interaction p-value										0.3588
First line treatment out	come (eC	RF)								
NED [PDS]	89	53 (59.6)	11.0 (5.8,23.3)	47	35 (74.5)	5.6 (3.0, 8.5)	0.57	0.37,	0.88	0.0113*
NED/CR [IDS]	74	41 (55.4)	13.9 (8.3, NE)	32	22 (68.8)	8.4 (3.1,21.2)	0.69	0.42,	1.19	0.1766
NED/CR [Chemo]	39	27 (69.2)	11.3 (5.6,21.9)	18	13 (72.2)	10.3 (3.0,11.2)	0.81	0.42,	1.62	0.5310
PR	50	35 (70.0)	11.0 (5.6,17.4)	34	20 (58.8)	11.3 (5.6,22.3)	1.19	0.70,	2.10	0.5280
Interaction p-value										0.2071
Screening laboratory tBR0	CA statu	ıs (eCRF)								
tBRCAm	147	88 (59.9)	13.4 (8.4,19.8)	67	44 (65.7)	10.3 (5.6,14.1)	0.83	0.58,	1.21	0.3282
non-tBRCAm	108	71 (65.7)	11.2 (8.3,14.0)	65	47 (72.3)	5.6 (2.9,11.1)	0.68	0.47,	0.99	0.0430*
Interaction p-value										0.4370
Age group										
<65 years	185	117 (63.2)	11.3 (8.4,15.2)	98	69 (70.4)	5.6 (3.0,10.3)	0.71	0.53,	0.96	0.0285*
>=65 years	70	42 (60.0)	11.3 (5.8,19.3)	34	22 (64.7)	11.1 (8.3,17.0)	0.87	0.53,	1.49	0.6127
Interaction p-value										0.4997

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aao 24MAR2022:14:14 kvbv306

Table 2.2.3.15 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Insomnia (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)			bevacizumab 132)			2-sided p-value [b]
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI [b]	
FIGO Stage (Disease state)									
III	182	117 (64.3)	11.3 (8.3,15.2)	90	61 (67.8)	8.5 (5.6,12.7)	0.85	0.62, 1.16	0.2989
IV	73	42 (57.5)	11.2 (8.3,22.3)	42	30 (71.4)	8.3 (2.8,11.2)	0.56	0.35, 0.91	0.0193*
Interaction p-value									0.1579
Region									
Europe	245	156 (63.7)	11.1 (8.4,13.9)	126	86 (68.3)	8.3 (5.6,11.1)	0.77	0.60, 1.01	0.0589
Japan	10	3 (30.0)	NE (NE, NE)	6	5 (83.3)	22.1 (2.8, NE)	0.30	0.06, 1.22	0.0928
Interaction p-value									0.1942
ECOG performance status at	Basel	ine							
(0) Normal activity	190	119 (62.6)	11.4 (8.4,16.6)	100	70 (70.0)	7.4 (5.2,11.1)	0.74	0.55, 1.001	0.0509
(1) Restricted activity	61	36 (59.0)	11.1 (5.8,25.7)	31	21 (67.7)	11.1 (3.2,16.9)	0.73	0.43, 1.28	0.2691
Interaction p-value									0.9756
Baseline CA-125 value									
<=ULN	228		11.4 (8.4,14.1)	118		8.3 (5.6,11.2)	0.76	0.58, 1.00002	0.0500
>ULN	27	16 (59.3)	11.0 (5.7,20.0)	14	10 (71.4)	8.3 (2.9,21.2)	0.70	0.32, 1.61	0.3906
Interaction p-value									0.8612
Histological grade									
High grade	255	159 (62.4)	11.3 (8.4,14.0)	132	91 (68.9)	8.3 (5.6,11.1)	0.75	0.58, 0.98	0.0330*
Interaction p-value									NC
Cytoreductive surgery outco	ome								
No residue	166	97 (58.4)	11.3 (8.4,17.3)	80	58 (72.5)	5.7 (3.0, 8.5)	0.62	0.45, 0.86	0.0045*
Residue	79	58 (73.4)	11.1 (5.7,14.0)	44	29 (65.9)	11.1 (3.5,17.0)	1.07	0.69, 1.69	0.7705
Interaction p-value									0.0490*

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aao 24MAR2022:14:14 kvbv306

Table 2.2.3.15 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-C30 Single item symptom scale: Insomnia (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + bevacizumab (N=255)				oevacizumab 132)				
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	96 (65.8)	11.0 (8.3,13.8)	79	56 (70.9)	5.7 (3.0,10.3)	0.74	0.53,	1.03	0.0748
Interval	99	59 (59.6)	13.8 (8.3,18.6)	45	31 (68.9)	11.1 (5.6,16.9)	0.77	0.50,	1.20	0.2473
Interaction p-value										0.8742
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	95 (60.1)	11.3 (8.4,15.9)	77	50 (64.9)	8.5 (5.2,11.3)	0.79	0.56,	1.12	0.1840
Non-tBRCAm	97	64 (66.0)	11.2 (5.8,17.4)	55	41 (74.5)	8.3 (3.2,11.1)	0.71	0.48,	1.06	0.0909
Interaction p-value										0.6831
Status somatic BRCA mutat	ions									
sBRCAm	22	15 (68.2)	8.5 (3.0,14.1)	7	5 (71.4)	2.8 (2.8, NE)	0.46	0.18,	1.42	0.1613
gBRCAm	66	43 (65.2)	8.4 (5.8,14.1)	31	20 (64.5)	13.9 (5.6,22.3)	1.09	0.65,	1.89	0.7512
Non-BRCAm	41	26 (63.4)	13.8 (8.5,25.7)	22	17 (77.3)	8.3 (2.8,12.7)	0.61	0.33,	1.15	0.1225
Interaction p-value										0.2111

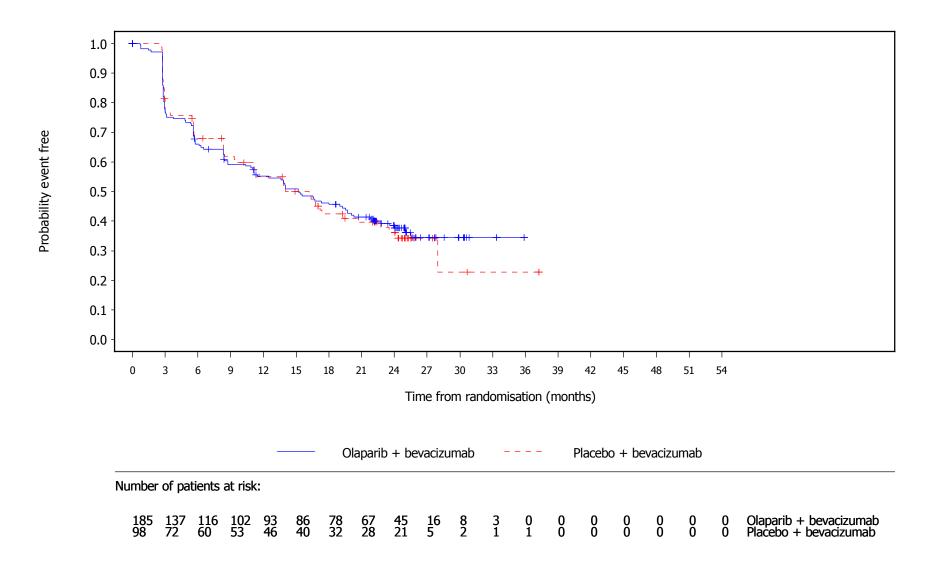
Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3aao 24MAR2022:14:14 kvbv306

Figure 2.2.4.1 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Global QoL/health status (MID = 15) time to clinically meaningful deterioration for Age group=<65 years

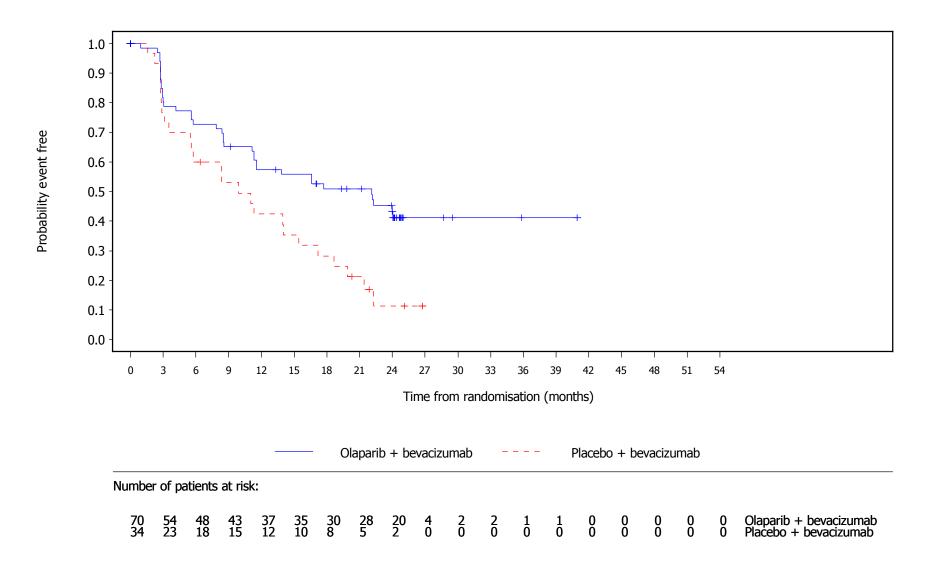
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3daa 24MAR2022:14:14 kvbv306

Figure 2.2.4.2 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Global QoL/health status (MID = 15) time to clinically meaningful deterioration for Age group=>=65 years

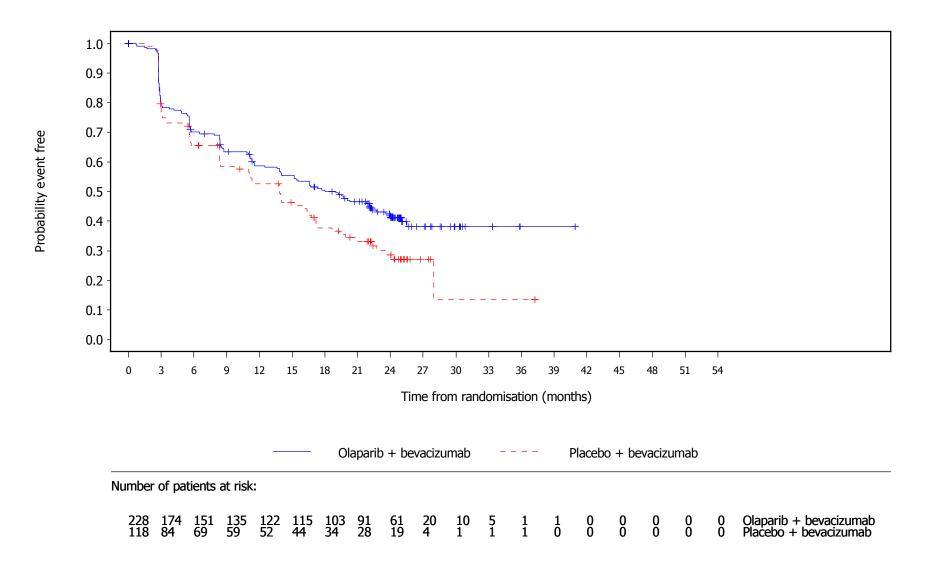
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dab 24MAR2022:14:14 kvbv306

Figure 2.2.4.3 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Global QoL/health status (MID = 15) time to clinically meaningful deterioration for Baseline CA-125 value=<=ULN

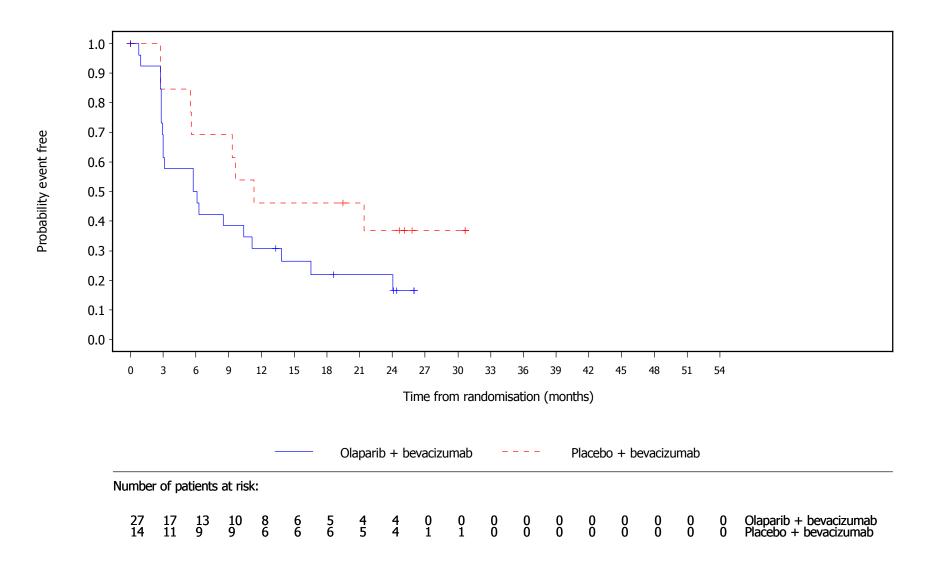
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dac 24MAR2022:14:14 kvbv306

Figure 2.2.4.4 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Global QoL/health status (MID = 15) time to clinically meaningful deterioration for Baseline CA-125 value=>ULN

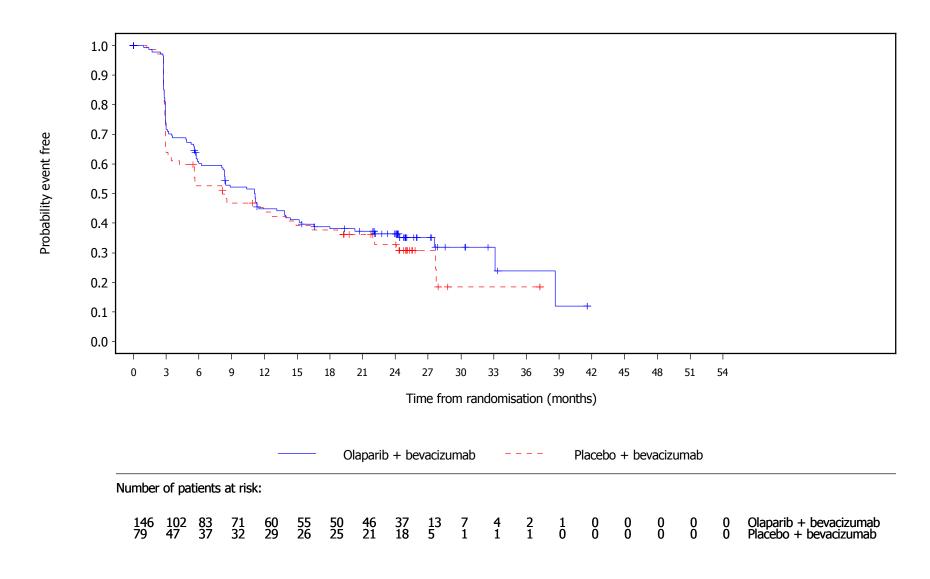
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dad 24MAR2022:14:14 kvbv306

Figure 2.2.4.5 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Role (MID = 15) time to clinically meaningful deterioration for Timing of cytoreductive surgery=Upfront

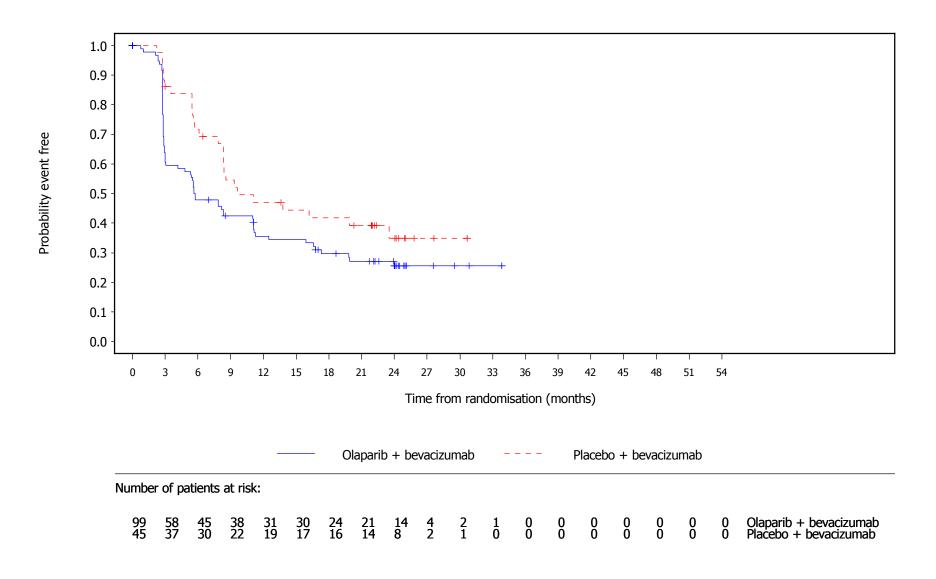
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dae 24MAR2022:14:14 kvbv306

Figure 2.2.4.6 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Role (MID = 15) time to clinically meaningful deterioration for Timing of cytoreductive surgery=Interval

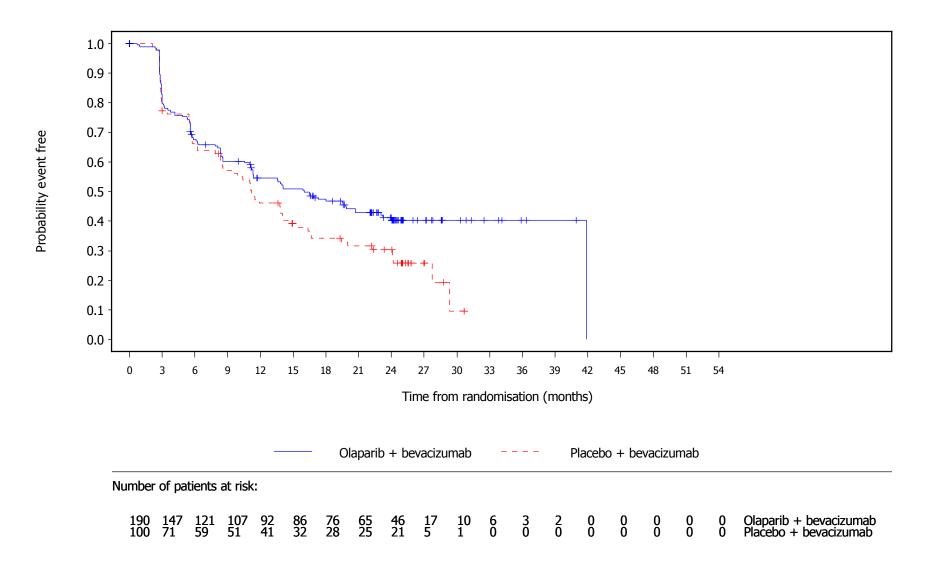
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3daf 24MAR2022:14:14 kvbv306

Figure 2.2.4.7 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Social (MID = 15) time to clinically meaningful deterioration for ECOG performance status at Baseline=(0) Normal activity

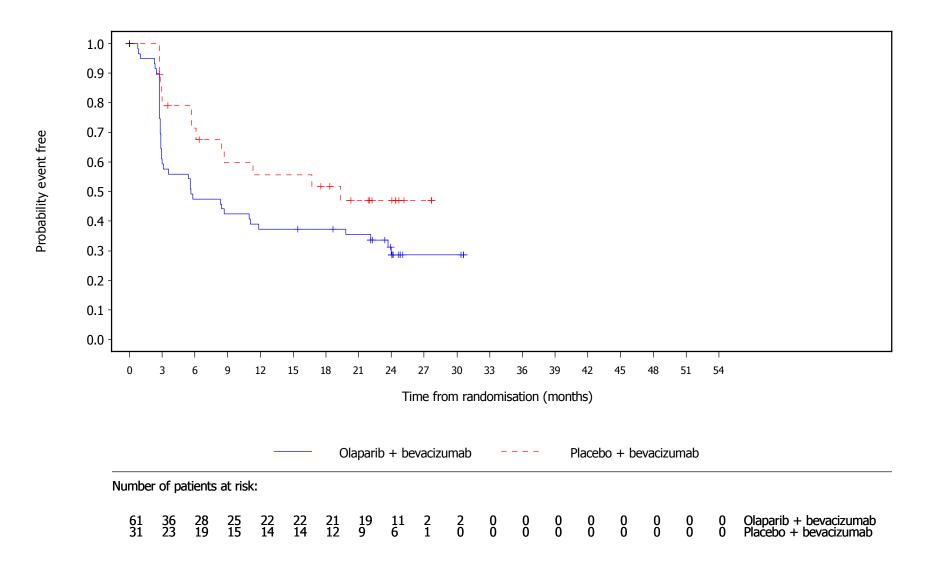
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dag 24MAR2022:14:14 kvbv306

Figure 2.2.4.8 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Social (MID = 15) time to clinically meaningful deterioration for ECOG performance status at Baseline=(1) Restricted activity

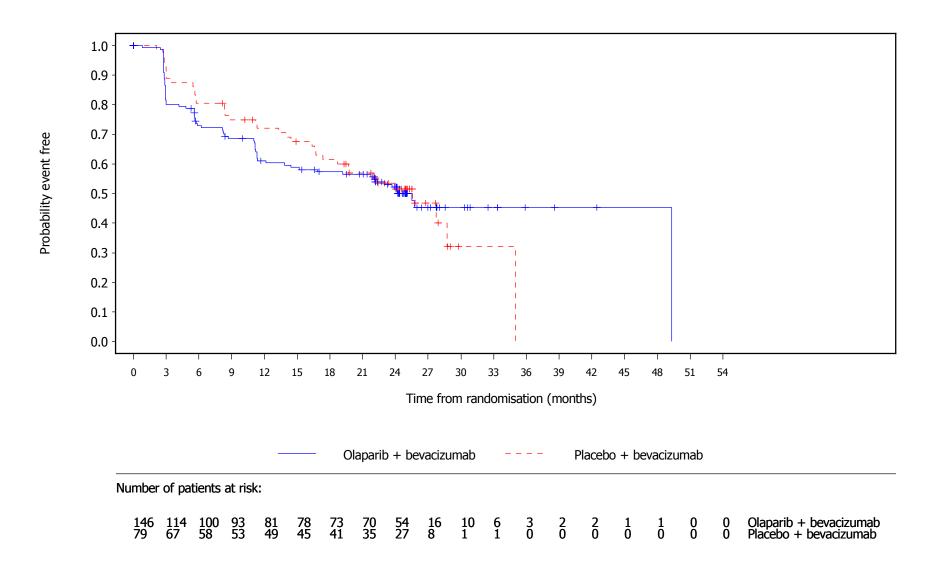
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dah 24MAR2022:14:14 kvbv306

Figure 2.2.4.9 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Loss of appetite (MID = 15) time to clinically meaningful deterioration for Timing of cytoreductive surgery=Upfront

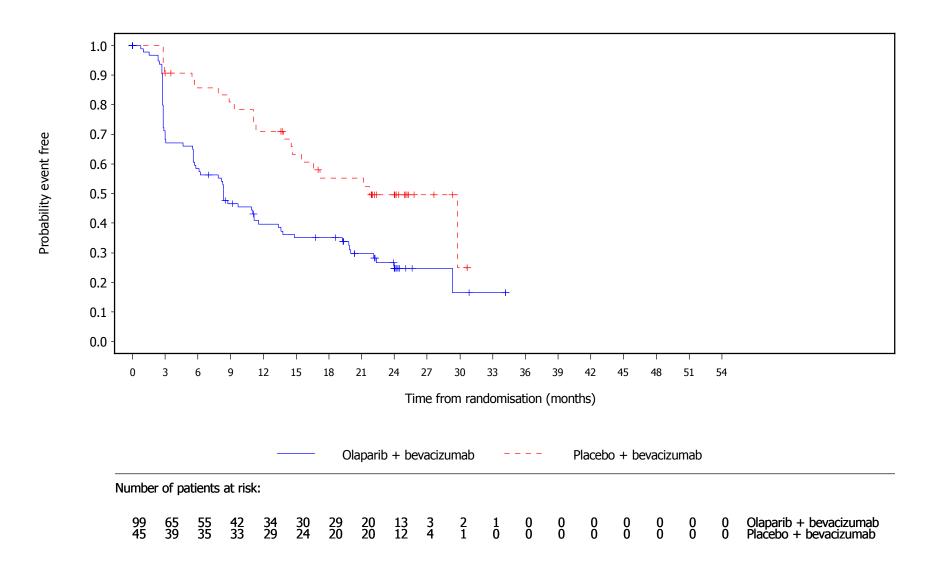
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dai 24MAR2022:14:14 kvbv306

Figure 2.2.4.10 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Loss of appetite (MID = 15) time to clinically meaningful deterioration for Timing of cytoreductive surgery=Interval

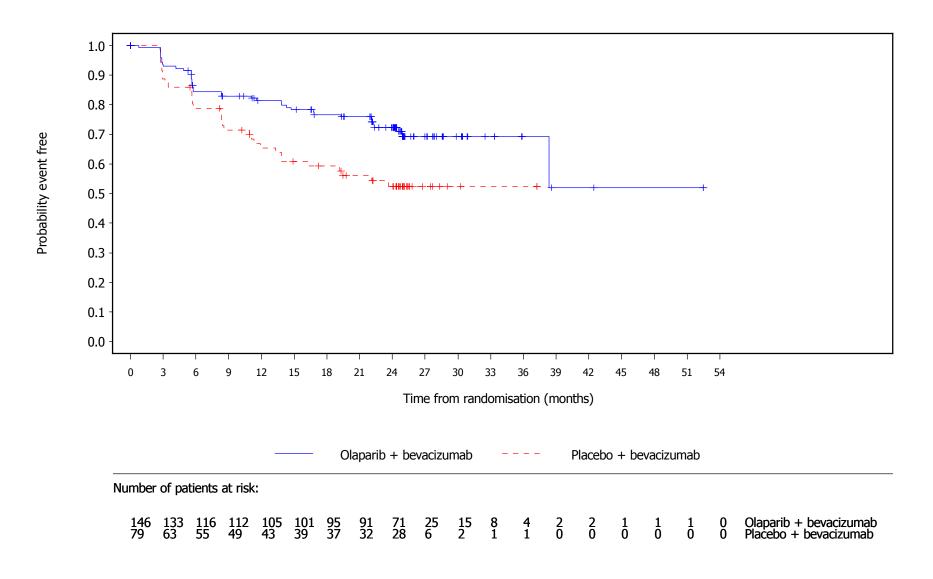
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3daj 24MAR2022:14:14 kvbv306

Figure 2.2.4.11 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Financial difficulties (MID = 15) time to clinically meaningful deterioration for Timing of cytoreductive surgery=Upfront

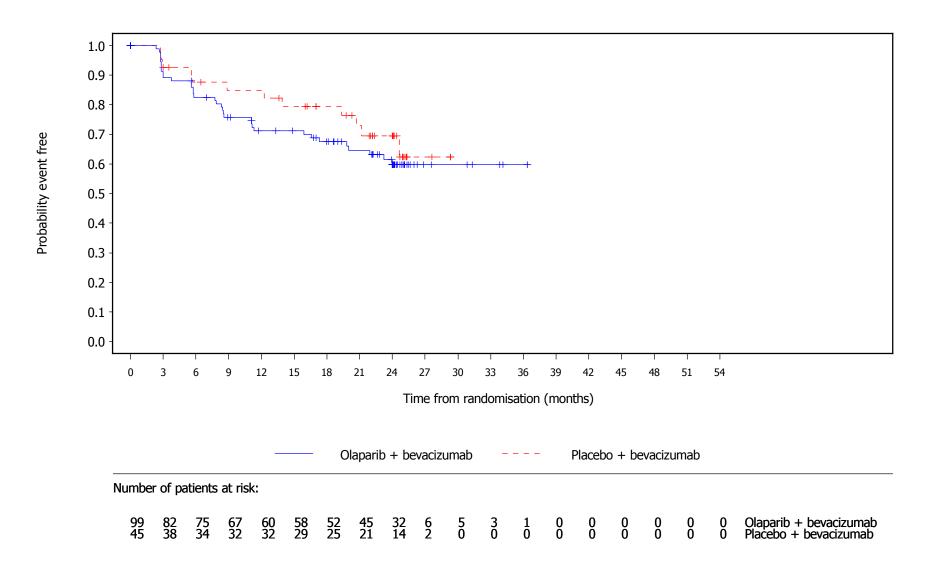
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dak 24MAR2022:14:14 kvbv306

Figure 2.2.4.12 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Financial difficulties (MID = 15) time to clinically meaningful deterioration for Timing of cytoreductive surgery=Interval

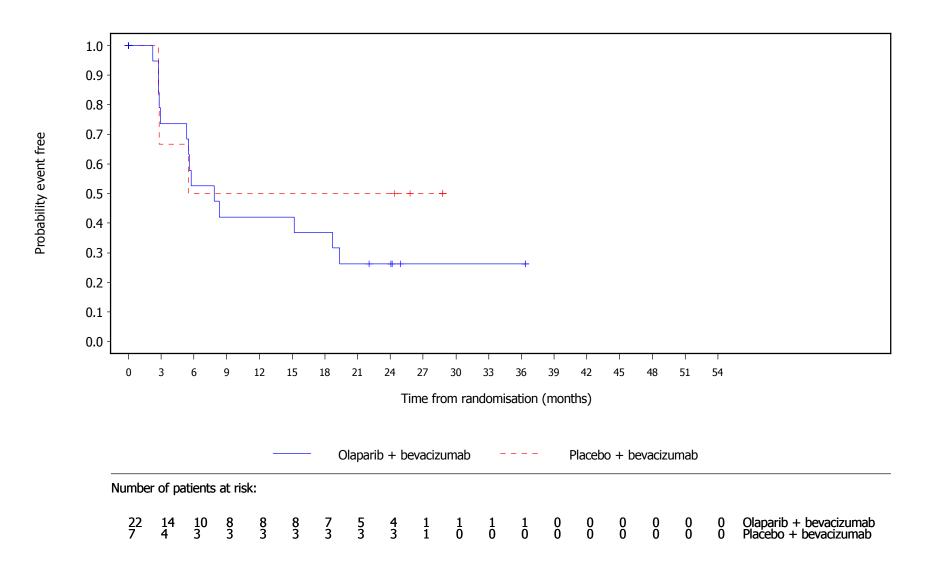
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dal 24MAR2022:14:14 kvbv306

Figure 2.2.4.13 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Pain (MID = 15) time to clinically meaningful deterioration for Status somatic BRCA mutations=sBRCAm

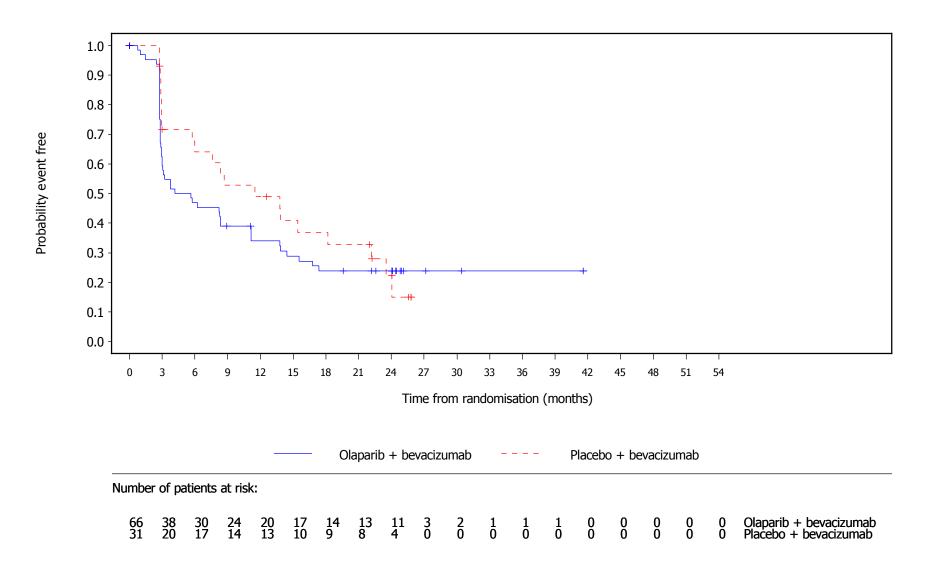
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dam 24MAR2022:14:14 kvbv306

Figure 2.2.4.14 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Pain (MID = 15) time to clinically meaningful deterioration for Status somatic BRCA mutations=gBRCAm

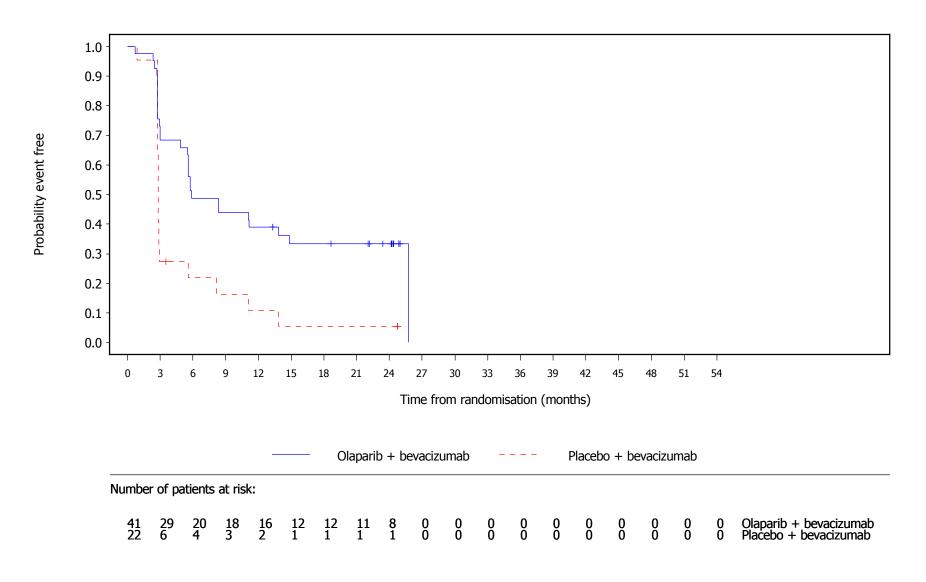
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dan 24MAR2022:14:14 kvbv306

Figure 2.2.4.15 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Pain (MID = 15) time to clinically meaningful deterioration for Status somatic BRCA mutations=Non-BRCAm

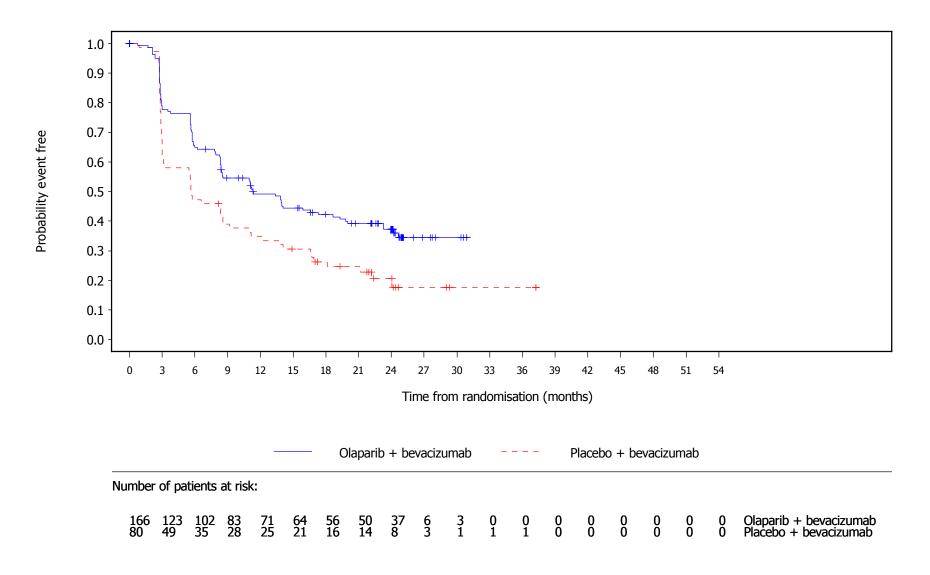
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dao 24MAR2022:14:14 kvbv306

Figure 2.2.4.16 PAOLAl Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Insomnia (MID = 15) time to clinically meaningful deterioration for Cytoreductive surgery outcome=No residue

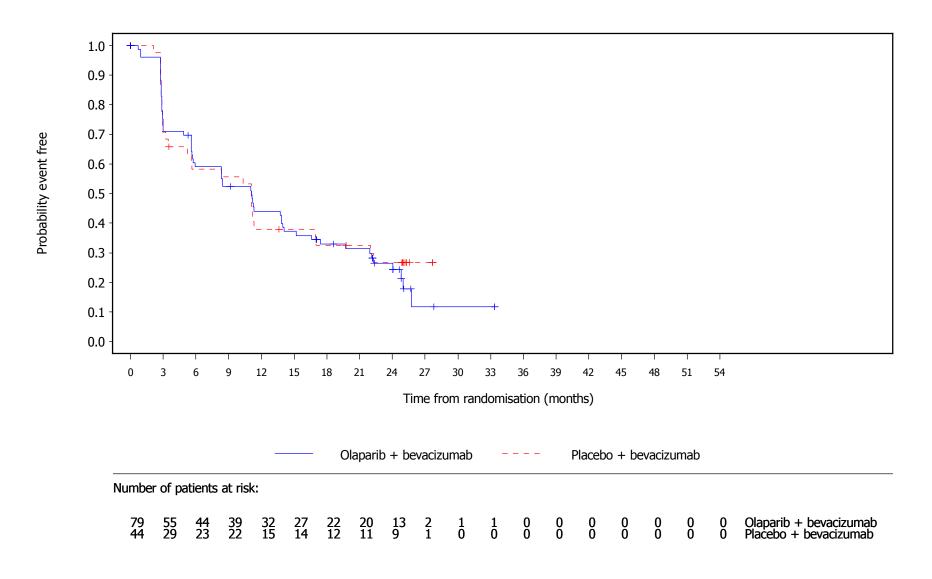
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3dap 24MAR2022:14:14 kvbv306

Figure 2.2.4.17 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Insomnia (MID = 15) time to clinically meaningful deterioration for Cytoreductive surgery outcome=Residue

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3daq 24MAR2022:14:14 kvbv306

Table 2.3.3.1 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	•	Olap	arib + (N=	bevac 255)	izumab			bevacizumab 132)				
Subgroup		of pa	er (%) atients events		edian time (95% CI) onths) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)										
NED [PDS]	92	51	(55.4)	16.8	(11.1, NE)	48	32 (66.7)	9.7 (5.6,19.4)	0.63	0.41,	0.99	0.0463*
NED/CR [IDS]	74	39	(52.7)	19.8	(9.7, NE)	38	20 (52.6)	13.9 (7.6, NE)	0.96	0.56,	1.67	0.8712
NED/CR [Chemo]	40	22	(55.0)	16.6	(11.1, NE)	20	12 (60.0)	16.7 (5.7, NE)	0.93	0.47,	1.95	0.8474
PR	49	27	(55.1)	16.7	(11.0,27.6)	26	17 (65.4)	14.0 (2.9,21.8)	0.65	0.36,	1.22	0.1751
Interaction p-value												0.5814
Screening laboratory tBR	CA statu	ıs (IV	/RS)									
tBRCAm	150	83	(55.3)	16.6	(11.1,25.5)	65	40 (61.5)	11.1 (7.6,19.3)	0.81	0.56,	1.19	0.2792
non-tBRCAm	105	56	(53.3)	19.9	(13.7,27.6)	67	41 (61.2)	13.8 (6.1,21.8)	0.71	0.47,	1.07	0.0969
Interaction p-value												0.6318
First line treatment out	come (eC	RF)										
NED [PDS]	89	47	(52.8)	18.0	(11.1, NE)	47	30 (63.8)	9.7 (5.7,24.8)	0.65	0.41,	1.04	0.0713
NED/CR [IDS]	74	40	(54.1)	19.6	(8.6, NE)	32	15 (46.9)	21.2 (7.6, NE)	1.14	0.65,	2.14	0.6547
NED/CR [Chemo]	39	22	(56.4)	16.6	(11.1, NE)	18	9 (50.0)	19.3 (5.7, NE)	1.20	0.57,	2.75	0.6399
PR	50	28	(56.0)	17.9	(11.0,27.6)	34	26 (76.5)	10.2 (5.5,15.4)	0.54	0.32,	0.93	0.0264*
Interaction p-value												0.1516
Screening laboratory tBR	CA statu	ıs (e0	CRF)									
tBRCAm	147	83	(56.5)	15.9	(11.1,20.7)	67	40 (59.7)	11.3 (8.2,21.2)	0.86	0.59,	1.26	0.4282
non-tBRCAm	108	56	(51.9)	22.0	(14.8, NE)	65	41 (63.1)	13.8 (5.7,21.8)	0.66	0.44,	0.99	0.0453*
Interaction p-value												0.3475
Age group												
<65 years	185	108	(58.4)	15.2	(11.1,19.8)	98	59 (60.2)	13.9 (8.3,21.8)	0.92	0.67,	1.27	0.5913
>=65 years	70	31	(44.3)	27.6	(19.6, NE)	34	22 (64.7)	11.1 (5.5,18.7)	0.44	0.26,	0.78	0.0053*
Interaction p-value												0.0278*

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3baa 24MAR2022:14:14 kvbv306

Table 2.3.3.1 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)									
III	182	97 (53.3)	17.3 (13.8,25.5)	90	54 (60.0)	11.3 (6.4,22.3)	0.76	0.55, 1.07	0.1174
IV Interaction p-value	73	42 (57.5)	17.9 (8.6,24.0)	42	27 (64.3)	13.9 (8.3,19.4)	0.76	0.47, 1.25	0.2797 NC
Region									
Europe	245	133 (54.3)	18.0 (13.8,23.4)	126	80 (63.5)	11.1 (8.3,16.8)	0.71	0.54, 0.94	0.0180*
Japan Interaction p-value	10	6 (60.0)	14.8 (3.1, NE)	6	1 (16.7)	NE (NE, NE)	4.84	0.83, 91.55	0.0848 0.0343*
ECOG performance status at	Basel	ine							
(0) Normal activity	190	108 (56.8)	16.7 (12.2,20.7)	100	63 (63.0)	11.1 (8.2,19.3)	0.80	0.59, 1.09	0.1578
(1) Restricted activity Interaction p-value	61	29 (47.5)	23.4 (11.1, NE)	31	18 (58.1)	14.0 (6.1, NE)	0.68	0.38, 1.25	0.2133 0.6525
Baseline CA-125 value									
<=ULN	228	126 (55.3)	16.8 (13.8,23.1)	118	72 (61.0)	13.8 (8.3,19.3)	0.78	0.58, 1.04	0.0930
>ULN	27	13 (48.1)	20.0 (5.8, NE)	14	9 (64.3)	10.3 (3.0, NE)	0.63	0.27, 1.53	0.2953
Interaction p-value									0.6484
Histological grade									
High grade	255	139 (54.5)	17.9 (13.8,23.1)	132	81 (61.4)	11.4 (8.3,18.7)	0.76	0.58, 1.005	0.0540
Interaction p-value									NC
Cytoreductive surgery outco	me								
No residue	166	89 (53.6)	18.0 (11.5,24.2)	80	46 (57.5)	13.8 (8.2,24.8)	0.80	0.56, 1.15	0.2279
Residue	79	46 (58.2)	16.7 (12.1,25.5)	44	29 (65.9)	11.3 (5.7,19.3)	0.76	0.48, 1.22	0.2519
Interaction p-value									0.8597

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3baa 24MAR2022:14:14 kvbv306

Table 2.3.3.1 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				pevacizumab 132)			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery								
Upfront	146	84 (57.5)	16.0 (11.3,23.4)	79	50 (63.3)	11.4 (5.8,19.3)	0.74	0.53, 1.06	0.1043
Interval	99	51 (51.5)	19.9 (11.5, NE)	45	25 (55.6)	13.9 (7.6, NE)	0.86	0.54, 1.42	0.5560
Interaction p-value									0.6189
Myriad tumour BRCA mutat:	ion stat	us							
tBRCAm	158	84 (53.2)	19.2 (13.7,27.6)	77	44 (57.1)	11.4 (8.3,21.2)	0.80	0.56, 1.16	0.2385
Non-tBRCAm	97	55 (56.7)	16.7 (11.3,23.4)	55	37 (67.3)	11.1 (5.7,19.9)	0.72	0.48, 1.10	0.1277
Interaction p-value									0.7061
Status somatic BRCA mutat	cions								
sBRCAm	22	11 (50.0)	16.6 (2.8, NE)	7	3 (42.9)	NE (NE, NE)	1.16	0.36, 5.14	0.8150
gBRCAm	66	38 (57.6)	13.8 (8.3, NE)	31	20 (64.5)	13.9 (8.2,21.2)	0.81	0.48, 1.41	0.4431
Non-BRCAm	41	27 (65.9)	16.7 (11.1,24.0)	22	18 (81.8)	7.1 (2.9,14.0)	0.57	0.31, 1.05	0.0723
Interaction p-value									0.5115

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3baa 24MAR2022:14:14 kvbv306

Table 2.3.3.2 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Body image (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	1	Olaparib + k (N=2			Placebo + h	pevacizumab 132)			
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IVF	RS)							
NED [PDS]	92	37 (40.2)	NE (NE, NE)	48	26 (54.2)	11.3 (3.0, NE)	0.55	0.33, 0.92	0.0224*
NED/CR [IDS]	74	41 (55.4)	14.8 (8.3, NE)	38	19 (50.0)	22.2 (8.8,26.5)	1.31	0.77, 2.31	0.3184
NED/CR [Chemo]	40	19 (47.5)	23.4 (11.6, NE)	20	12 (60.0)	12.4 (5.2, NE)	0.82	0.40, 1.74	0.6015
PR	49	29 (59.2)	11.2 (5.6,24.8)	26	14 (53.8)	17.5 (11.1,25.4)	1.24	0.67, 2.42	0.5046
Interaction p-value									0.0873
Screening laboratory tBRO	CA status	s (IVRS)							
tBRCAm	150	73 (48.7)	22.5 (11.6, NE)	65	33 (50.8)	21.2 (11.3, NE)	0.95	0.63, 1.45	0.8049
non-tBRCAm	105	53 (50.5)	19.5 (11.1, NE)	67	38 (56.7)	14.0 (8.5,25.4)	0.85	0.56, 1.30	0.4504
Interaction p-value									0.7143
First line treatment outo	come (eCF	RF)							
NED [PDS]	89	36 (40.4)	NE (NE, NE)	47	25 (53.2)	11.5 (5.6, NE)	0.58	0.35, 0.98	0.0405*
NED/CR [IDS]	74	38 (51.4)	19.9 (8.3, NE)	32	17 (53.1)	22.0 (8.3, NE)	1.08	0.62, 1.96	0.7991
NED/CR [Chemo]	39	20 (51.3)	21.9 (5.6, NE)	18	10 (55.6)	13.3 (3.5, NE)	0.92	0.44, 2.05	0.8303
PR	50	30 (60.0)	11.2 (5.7,24.8)	34	18 (52.9)	18.7 (12.7, NE)	1.36	0.76, 2.48	0.2999
Interaction p-value									0.1661
Screening laboratory tBRO	CA status	s (eCRF)							
tBRCAm	147	71 (48.3)	23.2 (12.5, NE)	67	34 (50.7)	21.2 (11.2, NE)	0.92	0.62, 1.40	0.6935
non-tBRCAm	108	55 (50.9)	18.7 (11.1, NE)	65	37 (56.9)	16.6 (8.5,25.4)	0.88	0.58, 1.34	0.5332
Interaction p-value									0.8653
Age group									
<65 years	185	90 (48.6)	22.5 (11.6, NE)	98	53 (54.1)	20.7 (11.1,25.4)	0.86	0.61, 1.21	0.3724
>=65 years	70	36 (51.4)	16.6 (8.4, NE)	34	18 (52.9)	18.7 (6.4, NE)	0.99	0.57, 1.78	0.9717
Interaction p-value									0.6649

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bab 24MAR2022:14:14 kvbv306

Table 2.3.3.2 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Body image (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2	bevacizu 255)	mab			bevacizumab 132)				
Subgroup		Number (%) of patients with events	(95%	n time % CI) ns) [a]		Number (%) of patients with events		Hazard ratio [b]			2-sided p-value [b]
FIGO Stage (Disease state)											_
III	182	87 (47.8)	23.4 (13	3.7, NE)	90	48 (53.3)	18.7 (8.8,27.8)	0.85	0.60,	1.22	0.3652
IV	73	39 (53.4)	14.8 (8	B.4, NE)	42	23 (54.8)	20.7 (9.7,25.4)	1.00	0.60,	1.70	0.9950
Interaction p-value											0.6090
Region											
Europe	245	122 (49.8)	19.9 (12	2.5, NE)	126	68 (54.0)	17.5 (11.5,25.1)	0.89	0.66,	1.20	0.4241
Japan	10	4 (40.0)	NE (NE, NE)	6	3 (50.0)	NE (NE, NE)	0.98	0.22,	4.96	0.9759
Interaction p-value											0.8990
ECOG performance status at	Basel	ine									
(0) Normal activity	190	95 (50.0)	18.0 (11	1.3, NE)	100	55 (55.0)	20.7 (11.2,25.4)	0.92	0.66,	1.29	0.6149
(1) Restricted activity	61	30 (49.2)	23.4 (8	B.4, NE)	31	16 (51.6)	17.5 (6.1, NE)	0.87	0.48,	1.64	0.6597
Interaction p-value											0.8836
Baseline CA-125 value											
<=ULN	228	112 (49.1)	22.5 (13	3.7, NE)	118	64 (54.2)	16.6 (11.1,25.1)	0.85	0.63,	1.16	0.3098
>ULN	27	14 (51.9)	11.3 (3	3.1, NE)	14	7 (50.0)	21.2 (2.9, NE)	1.30	0.54,	3.44	0.5632
Interaction p-value											0.3776
Histological grade											
High grade	255	126 (49.4)	21.9 (12	2.7, NE)	132	71 (53.8)	18.7 (11.5,25.1)	0.89	0.67,	1.20	0.4356
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	76 (45.8)	NE (NE, NE)	80	43 (53.8)	20.7 (8.3,26.5)	0.76	0.52,	1.11	0.1522
Residue	79	44 (55.7)	16.4 (8	3.4,24.8)	44	24 (54.5)	18.7 (11.2,25.4)	1.12	0.69,	1.86	0.6618
Interaction p-value											0.2202

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bab 24MAR2022:14:14 kvbv306

Table 2.3.3.2 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Body image (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery								
Upfront	146	67 (45.9)	NE (NE, NE)	79	42 (53.2)	14.0 (8.3, NE)	0.74	0.50, 1.09	0.1248
Interval	99	53 (53.5)	16.6 (9.7,24.0)	45	25 (55.6)	22.1 (8.8,25.4)	1.11	0.70, 1.81	0.6735
Interaction p-value									0.1894
Myriad tumour BRCA mutati	ion stat	us							
tBRCAm	158	78 (49.4)	21.9 (11.3, NE)	77	39 (50.6)	21.2 (11.2,27.8)	0.94	0.64, 1.39	0.7375
Non-tBRCAm	97	48 (49.5)	19.9 (9.7, NE)	55	32 (58.2)	16.6 (9.7,25.4)	0.84	0.54, 1.32	0.4354
Interaction p-value									0.7067
Status somatic BRCA mutat	cions								
sBRCAm	22	5 (22.7)	NE (NE, NE)	7	3 (42.9)	NE (NE, NE)	0.43	0.11, 2.11	0.2737
gBRCAm	66	37 (56.1)	12.5 (8.3, NE)	31	16 (51.6)	22.3 (11.3, NE)	1.23	0.70, 2.28	0.4784
Non-BRCAm	41	21 (51.2)	19.9 (5.8, NE)	22	13 (59.1)	16.6 (9.7, NE)	0.93	0.47, 1.91	0.8398
Interaction p-value									0.4216

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bab 24MAR2022:14:14 kvbv306

Table 2.3.3.3 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		-	bevacizumab 255)				bevacizumab 132)					
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events		[)	Hazard ratio [b]		[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)										
NED [PDS]	92	33 (35.9)	NE (NE,	NE)	48	19 (39.6)	25.5 (13.8	NE)	0.74	0.42,	1.32	0.3006
NED/CR [IDS]	74	31 (41.9)	NE (NE,	NE)	38	, ,	26.5 (11.2		1.00	0.56,		0.9973
NED/CR [Chemo]	40	14 (35.0)	NE (NE,	NE)	20	12 (60.0)	19.1 (3.0	, NE)	0.52	0.24,	1.15	0.1049
PR	49	17 (34.7)	NE (NE,	NE)	26	12 (46.2)	22.2 (11.4	, NE)	0.63	0.30,	1.36	0.2335
Interaction p-value												0.5868
Screening laboratory tBR	CA statu	s (IVRS)										
tBRCAm	150	52 (34.7)	NE (NE,	NE)	65	29 (44.6)	22.9 (13.6	, NE)	0.65	0.42,	1.04	0.0696
non-tBRCAm	105	43 (41.0)	25.7 (22.3,	NE)	67	30 (44.8)	25.4 (12.7	, NE)	0.86	0.54,	1.39	0.5391
Interaction p-value												0.3938
First line treatment out	come (eC	RF)										
NED [PDS]	89	29 (32.6)	NE (NE,	NE)	47	18 (38.3)	NE (NE	, NE)	0.71	0.40,	1.30	0.2571
NED/CR [IDS]	74	31 (41.9)	NE (NE,	NE)	32	14 (43.8)	26.5 (11.0	, NE)	0.94	0.51,	1.82	0.8511
NED/CR [Chemo]	39	13 (33.3)	NE (NE,	NE)	18	9 (50.0)	22.3 (5.7	, NE)	0.62	0.27,	1.51	0.2833
PR	50	21 (42.0)	25.7 (15.4,	NE)	34	17 (50.0)	22.2 (8.6	, NE)	0.72	0.38,	1.37	0.3095
Interaction p-value												0.8616
Screening laboratory tBR	CA statu	s (eCRF)										
tBRCAm	147	51 (34.7)	NE (NE,	NE)	67	30 (44.8)	22.9 (11.4	, NE)	0.64	0.41,	1.01	0.0573
non-tBRCAm	108	44 (40.7)	25.7 (22.3,	NE)	65	29 (44.6)	25.4 (12.7	, NE)	0.87	0.55,	1.41	0.5719
Interaction p-value												0.3482
Age group												
<65 years	185	69 (37.3)	NE (NE,	NE)	98	, ,	25.4 (13.8		0.78	0.53,		0.2059
>=65 years	70	26 (37.1)	NE (NE,	NE)	34	17 (50.0)	22.2 (12.3	, NE)	0.64	0.35,	1.20	0.1568
Interaction p-value												0.5881

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bac 24MAR2022:14:14 kvbv306

Table 2.3.3.3 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2					bevacizumab 132)					
Subgroup		Number (%) of patients with events	Median ti (95% CI) (months) [)		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]		2-sided p-value [b]	
FIGO Stage (Disease state)												
III	182	72 (39.6)	NE (NE,	NE)	90	41 (45.6)	24.1 (13.8,	NE)	0.80	0.55,	1.19	0.2636
IV	73	23 (31.5)	NE (NE,	NE)	42	18 (42.9)	23.5 (8.3,	NE)	0.59	0.32,	1.10	0.0969
Interaction p-value												0.4045
Region												
Europe	245	94 (38.4)	NE (NE,	NE)	126	56 (44.4)	24.1 (16.4,	NE)	0.76	0.55,	1.06	0.1042
Japan	10	1 (10.0)	NE (NE,	NE)	6	3 (50.0)	NE (NE,	NE)	0.20	0.01,	1.55	0.1246
Interaction p-value												0.2131
ECOG performance status at	Basel	ine										
(0) Normal activity	190	69 (36.3)	NE (NE,	NE)	100	45 (45.0)	25.4 (17.0,	NE)	0.75	0.52,	1.11	0.1472
(1) Restricted activity	61	24 (39.3)	25.7 (14.3,	NE)	31	14 (45.2)	22.2 (5.9,	NE)	0.67	0.35,	1.32	0.2388
Interaction p-value												0.7495
Baseline CA-125 value												
<=ULN	228	83 (36.4)	NE (NE,	NE)	118	49 (41.5)	25.5 (18.7,	NE)	0.77	0.54,		0.1452
>ULN	27	12 (44.4)	16.6 (5.7,	NE)	14	10 (71.4)	17.0 (5.6,	25.4)	0.61	0.26,	1.45	0.2564
Interaction p-value												0.6275
Histological grade												
High grade	255	95 (37.3)	NE (NE,	NE)	132	59 (44.7)	24.1 (17.0,	NE)	0.74	0.53,	1.02	0.0691
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	61 (36.7)	NE (NE,	NE)	80	33 (41.3)	25.5 (13.8,	NE)	0.79	0.52,	1.22	0.2847
Residue	79	31 (39.2)	25.7 (22.3,	NE)	44	23 (52.3)	22.3 (11.1,	NE)	0.68	0.40,	1.18	0.1682
Interaction p-value												0.6689

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bac 24MAR2022:14:14 kvbv306

Table 2.3.3.3 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2					oevacizumab 132)					
Subgroup		Number (%) of patients with events	Median ti (95% CI) (months) [95% CI) of pa		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b] 95% CI [b]		[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	50 (34.2)	NE (NE,	NE)	79	36 (45.6)	22.9 (13.8,	NE)	0.62	0.40,	0.95	0.0293*
Interval	99	42 (42.4)	25.4 (16.6,	NE)	45	20 (44.4)	25.4 (12.3,	NE)	0.98	0.58,	1.71	0.9488
Interaction p-value												0.1772
Myriad tumour BRCA mutat:	ion stat	us										
tBRCAm	158	57 (36.1)	NE (NE,	NE)	77	29 (37.7)	25.5 (19.1,	NE)	0.85	0.55,	1.34	0.4715
Non-tBRCAm	97	38 (39.2)	NE (NE,	NE)	55	30 (54.5)	18.7 (10.4,	26.5)	0.64	0.39,	1.03	0.0671
Interaction p-value												0.3891
Status somatic BRCA mutat	cions											
sBRCAm	22	10 (45.5)	16.6 (5.6,	NE)	7	4 (57.1)	15.3 (2.8,	NE)	0.83	0.28,	3.03	0.7577
gBRCAm	66	27 (40.9)	NE (NE,	NE)	31	11 (35.5)	NE (NE,	NE)	0.95	0.48,	1.99	0.8782
Non-BRCAm	41	16 (39.0)	NE (NE,	NE)	22	13 (59.1)	13.8 (5.6,	NE)	0.55	0.27,	1.17	0.1195
Interaction p-value												0.5712

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bac 24MAR2022:14:14 kvbv306

Table 2.3.3.4 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment
(MID = 15) time to clinically meaningful deterioration
Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2	bevacizumab 255)	·			bevacizumab 132)					
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events	, ,		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IVI	RS)										
NED [PDS]	92	28 (30.4)	52.5 (NE,	NE)	48	19 (39.6)	NE (NE,	NE)	0.65	0.36,	1.19	0.1603
NED/CR [IDS]	74	33 (44.6)	24.0 (11.3,	NE)	38	9 (23.7)	NE (NE,	NE)	2.23	1.11,	4.96	0.0225*
NED/CR [Chemo]	40	13 (32.5)	NE (NE,	NE)	20	9 (45.0)	NE (NE,	NE)	0.77	0.33,	1.87	0.5516
PR	49	21 (42.9)	24.9 (11.3,	NE)	26	8 (30.8)	25.4 (8.3,	NE)	1.27	0.59,	3.06	0.5551
Interaction p-value												0.0532
Screening laboratory tBR0	CA status	s (IVRS)										
tBRCAm	150	57 (38.0)	NE (NE,	NE)	65	19 (29.2)	NE (NE,	NE)	1.32	0.80,	2.27	0.2870
non-tBRCAm	105	38 (36.2)	52.5 (22.5,	NE)	67	26 (38.8)	25.4 (13.9,	NE)	0.91	0.55,	1.51	0.6998
Interaction p-value												0.3064
First line treatment out	come (eCI	RF)										
NED [PDS]	89	29 (32.6)	52.5 (NE,	NE)	47	19 (40.4)	NE (NE,	NE)	0.68	0.38,	1.24	0.2058
NED/CR [IDS]	74	31 (41.9)	NE (NE,	NE)	32	8 (25.0)	NE (NE,	NE)	1.89	0.91,	4.42	0.0882
NED/CR [Chemo]	39	12 (30.8)	NE (NE,	NE)	18	7 (38.9)	NE (NE,	NE)	0.79	0.32,	2.13	0.6266
PR	50	22 (44.0)	24.9 (11.3,	NE)	34	11 (32.4)	NE (NE,	NE)	1.40	0.69,	3.00	0.3514
Interaction p-value												0.1423
Screening laboratory tBR0	CA status	s (eCRF)										
tBRCAm	147	57 (38.8)	NE (NE,	NE)	67	19 (28.4)	NE (NE,	NE)	1.38	0.84,	2.38	0.2132
non-tBRCAm	108	38 (35.2)	52.5 (NE,	NE)	65	26 (40.0)	25.4 (13.9,	NE)	0.85	0.52,	1.42	0.5300
Interaction p-value												0.1870
Age group												
<65 years	185	69 (37.3)	NE (NE,	NE)	98	34 (34.7)	NE (NE,	NE)	1.09	0.73,	1.66	0.6736
>=65 years	70	26 (37.1)	52.5 (23.2,	NE)	34	11 (32.4)	NE (NE,	NE)	1.03	0.52,	2.19	0.9246
Interaction p-value												0.8981

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bad 24MAR2022:14:14 kvbv306

Table 2.3.3.4 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment
(MID = 15) time to clinically meaningful deterioration
Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N	bevacizur =255)	nab	•			bevacizumab 132)					
Subgroup		Number (%) of patient with event	s (95%	cI)			Number (%) of patients with events		.)	Hazard ratio [b]	95% C]	[b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	69 (37.9) 52.5 (24	1.9,	NE)	90	29 (32.2)	NE (NE,	NE)	1.18	0.77,	1.84	0.4585
IV	73	26 (35.6) NE (NE,	NE)	42	16 (38.1)	25.4 (13.9,	NE)	0.89	0.48,	1.70	0.7244
Interaction p-value													0.4790
Region													
Europe	245	93 (38.0) 52.5 (NE,	NE)	126	43 (34.1)	NE (NE,	NE)	1.08	0.75,	1.56	0.6942
Japan	10	2 (20.0) NE (NE,	NE)	6	2 (33.3)	NE (NE,	NE)	0.85	0.10,	7.11	0.8742
Interaction p-value													0.8207
ECOG performance status at	Basel	ine											
(0) Normal activity	190	75 (39.5) 52.5 (23	3.2,	NE)	100	35 (35.0)	NE (NE,	NE)	1.16	0.78,	1.75	0.4778
(1) Restricted activity	61	18 (29.5) NE (NE,	NE)	31	10 (32.3)	NE (NE,	NE)	0.82	0.39,	1.86	0.6281
Interaction p-value													0.4526
Baseline CA-125 value													
<=ULN	228	85 (37.3	,	NE,	NE)	118	40 (33.9)	NE (NE,	NE)	1.07	0.74,		0.7395
>ULN	27	10 (37.0) NE (NE,	NE)	14	5 (35.7)	25.4 (9.3,	NE)	1.18	0.42,	3.80	0.7556
Interaction p-value													0.8553
Histological grade													
High grade	255	95 (37.3) 52.5 (NE,	NE)	132	45 (34.1)	NE (NE,	NE)	1.08	0.76,	1.55	0.6816
Interaction p-value													NC
Cytoreductive surgery outco	me												
No residue	166	61 (36.7) 52.5 (NE,	NE)	80	27 (33.8)	NE (NE,	NE)	1.05	0.67,	1.68	0.8337
Residue	79	28 (35.4) NE (NE,	NE)	44	16 (36.4)	NE (NE,	NE)	1.01	0.55,	1.91	0.9723
Interaction p-value													0.9231

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bad 24MAR2022:14:14 kvbv306

Table 2.3.3.4 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment
(MID = 15) time to clinically meaningful deterioration
Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2					bevacizumab 132)					
Subgroup		Number (%) of patients with events	Median ti (95% CI) (months) [Number (%) of patients with events)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	48 (32.9)	52.5 (NE,	NE)	79	31 (39.2)	NE (NE,	NE)	0.75	0.48,	1.19	0.2155
Interval	99	41 (41.4)	NE (NE,	NE)	45	12 (26.7)	NE (NE,	NE)	1.80	0.98,	3.59	0.0591
Interaction p-value												0.0246*
Myriad tumour BRCA mutati	ion stat	us										
tBRCAm	158	60 (38.0)	NE (NE,	NE)	77	22 (28.6)	NE (NE,	NE)	1.30	0.81,	2.17	0.2825
Non-tBRCAm	97	35 (36.1)	52.5 (22.5,	NE)	55	23 (41.8)	25.4 (13.9,	NE)	0.86	0.51,	1.48	0.5855
Interaction p-value												0.2628
Status somatic BRCA mutat	cions											
sBRCAm	22	3 (13.6)	NE (NE,	NE)	7	0	NE (NE,	NE)	NC	NC		NC
gBRCAm	66	32 (48.5)	24.9 (11.0,	NE)	31	6 (19.4)	NE (NE,	NE)	2.71	1.22,	7.20	0.0129*
Non-BRCAm	41	13 (31.7)	52.5 (22.5,	NE)	22	11 (50.0)	22.2 (8.3,	NE)	0.62	0.27,	1.42	0.2524
Interaction p-value												0.0126*

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bad 24MAR2022:14:14 kvbv306

Table 2.3.3.5 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Hormonal (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olap	earib + (N=2	bevac 255)	izumab				pevacizumab .32)				
Subgroup	n	of pa	er (%) atients events		dian time (95% CI) onths) [a]		Number (of patier with ever	ıts	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)											
NED [PDS]	92	50	(54.3)	17.0	(13.8,33.4)	48	29 (60.	.4)	8.5 (5.5,19.3)	0.69	0.44,	1.10	0.1124
NED/CR [IDS]	74	39	(52.7)	19.3	(11.1, NE)	38	18 (47.	4)	19.4 (5.6, NE)	1.07	0.62,	1.92	0.8006
NED/CR [Chemo]	40	20	(50.0)	21.9	(14.0, NE)	20	14 (70.	.0)	11.0 (2.9,24.6)	0.61	0.31,	1.23	0.1587
PR	49	26	(53.1)	19.5	(11.2,30.4)	26	15 (57.	.7)	5.6 (2.9,19.9)	0.62	0.33,	1.20	0.1480
Interaction p-value													0.4717
Screening laboratory tBR0	CA statı	ıs (IV	VRS)										
tBRCAm	150	83	(55.3)	19.3	(13.9,24.2)	65	37 (56.	9)	11.8 (5.6,24.1)	0.79	0.54,	1.18	0.2472
non-tBRCAm	105	52	(49.5)	18.6	(12.1, NE)	67	39 (58.	.2)	11.1 (5.6,19.3)	0.69	0.46,	1.06	0.0873
Interaction p-value													0.6425
First line treatment out	come (e0	CRF)											
NED [PDS]	89	46	(51.7)	18.9	(13.8, NE)	47	29 (61.	.7)	6.2 (5.5,19.3)	0.61	0.38,	0.97	0.0388*
NED/CR [IDS]	74	40	(54.1)	16.8	(8.7,25.3)	32	15 (46.	9)	19.4 (5.6, NE)	1.09	0.62,	2.04	0.7742
NED/CR [Chemo]	39	19	(48.7)	21.9	(14.3, NE)	18	10 (55.	6)	13.9 (3.0, NE)	0.74	0.35,	1.66	0.4521
PR	50	28	(56.0)	19.3	(10.9,30.4)	34	22 (64.	.7)	11.1 (4.7,19.9)	0.69	0.39,	1.21	0.1937
Interaction p-value													0.4754
Screening laboratory tBR0	CA statı	ıs (e0	CRF)										
tBRCAm	147	81	(55.1)	19.2	(13.9,24.2)	67	38 (56.	.7)	11.3 (5.6,24.1)	0.78	0.54,	1.16	0.2201
non-tBRCAm	108	54	(50.0)	19.1	(12.1,33.4)	65	38 (58.	.5)	11.1 (5.6,19.4)	0.70	0.46,	1.07	0.1001
Interaction p-value													0.7082
Age group													
<65 years	185	101	(54.6)	17.5	(13.9,22.1)	98	60 (61.	.2)	8.4 (5.5,13.9)	0.68	0.50,	0.95	0.0227*
>=65 years	70	34	(48.6)	24.0	(11.5, NE)	34	16 (47.	.1)	19.9 (5.7, NE)	0.97	0.54,	1.80	0.9188
Interaction p-value													0.3066

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bae 24MAR2022:14:14 kvbv306

Table 2.3.3.5 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Hormonal (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + (N=2				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	99 (54.4)	16.8 (13.9,24.0)	90	52 (57.8)	11.3 (5.7,19.4)	0.81	0.58,	1.14	0.2241
IV	73	36 (49.3)	22.3 (13.9, NE)	42	24 (57.1)	13.9 (3.0,24.6)	0.61	0.37,	1.04	0.0706
Interaction p-value										0.3820
Region										
Europe	245	130 (53.1)	19.1 (14.0,24.2)	126	72 (57.1)	11.3 (5.7,16.6)	0.75	0.57,		0.0579
Japan	10	5 (50.0)	22.1 (2.8, NE)	6	4 (66.7)	12.5 (2.9, NE)	0.62	0.16,	2.52	0.4866
Interaction p-value										0.7836
ECOG performance status at	Basel	ine								
(0) Normal activity	190	94 (49.5)	20.0 (16.0,30.4)	100	60 (60.0)	11.1 (5.6,19.3)	0.68	0.49,	0.95	0.0226*
(1) Restricted activity	61	37 (60.7)	11.5 (8.4,24.0)	31	16 (51.6)	11.3 (5.6, NE)	0.94	0.53,	1.73	0.8269
Interaction p-value										0.3471
Baseline CA-125 value										
<=ULN	228	124 (54.4)	17.5 (13.9,24.0)	118	66 (55.9)	12.3 (5.7,19.4)	0.82	0.61,		0.1926
>ULN	27	11 (40.7)	25.3 (8.5, NE)	14	10 (71.4)	5.6 (2.8, NE)	0.33	0.14,	0.79	0.0142*
Interaction p-value										0.0543
Histological grade										
High grade	255	135 (52.9)	19.1 (14.3,24.2)	132	76 (57.6)	11.3 (5.6,19.1)	0.75	0.57,	0.99	0.0460*
Interaction p-value										NC
Cytoreductive surgery outco	me									
No residue	166	88 (53.0)	17.5 (13.8,25.3)	80	44 (55.0)	11.8 (5.6,22.3)	0.79	0.55,	1.14	0.1994
Residue	79	42 (53.2)	21.9 (14.3,25.5)	44	28 (63.6)	11.3 (5.6,19.1)	0.68	0.42,	1.11	0.1233
Interaction p-value										0.6431

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bae 24MAR2022:14:14 kvbv306

Table 2.3.3.5 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Hormonal (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + N=2				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events		Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
Timing of cytoreductive s	urgery									
Upfront	146	77 (52.7)	19.1 (14.0,30.4)	79	49 (62.0)	8.7 (5.6,16.4)	0.64	0.45,	0.92	0.0174*
Interval	99	53 (53.5)	19.3 (11.1,24.4)	45	23 (51.1)	14.7 (5.6, NE)	0.97	0.60,	1.62	0.9141
Interaction p-value										0.1740
Myriad tumour BRCA mutati	on stat	us								
tBRCAm	158	87 (55.1)	19.3 (13.9,24.2)	77	42 (54.5)	11.3 (5.6,24.1)	0.81	0.56,	1.18	0.2711
Non-tBRCAm	97	48 (49.5)	18.6 (12.1, NE)	55	34 (61.8)	11.1 (5.6,19.3)	0.66	0.43,	1.03	0.0695
Interaction p-value										0.4860
Status somatic BRCA mutat	ions									
sBRCAm	22	9 (40.9)	24.4 (8.3, NE)	7	2 (28.6)	NE (NE, NE)	1.05	0.27,	6.92	0.9455
gBRCAm	66	42 (63.6)	11.2 (5.8,19.5)	31	18 (58.1)	8.5 (4.2,24.1)	0.94	0.55,	1.68	0.8286
Non-BRCAm	41	22 (53.7)	16.6 (7.8, NE)	22	16 (72.7)	12.3 (4.7,19.3)	0.63	0.33,	1.23	0.1706
Interaction p-value										0.6185

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bae 24MAR2022:14:14 kvbv306

Table 2.3.3.6 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + : (N=2					bevacizumab 132)					
Subgroup		Number (%) of patients with events	Median ti (95% CI) (months) [Number (%) of patients with events)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment outo	come (IV	RS)										
NED [PDS]	92	33 (35.9)	NE (NE,	NE)	48	22 (45.8)	16.4 (8.4,	NE)	0.63	0.37,	1.10	0.1055
NED/CR [IDS]	74	37 (50.0)	18.6 (13.8,	NE)	38	13 (34.2)	NE (NE,	NE)	1.62	0.88,	3.16	0.1218
NED/CR [Chemo]	40	21 (52.5)	19.4 (3.0,	NE)	20	11 (55.0)	17.2 (3.0,	NE)	1.07	0.52,	2.30	0.8612
PR	49	23 (46.9)	25.3 (11.1,	NE)	26	12 (46.2)	12.7 (3.0,	NE)	0.80	0.41,	1.67	0.5401
Interaction p-value												0.1485
Screening laboratory tBRO	CA statu	s (IVRS)										
tBRCAm	150	73 (48.7)	22.1 (14.5,	NE)	65	33 (50.8)	13.9 (8.3,	NE)	0.87	0.58,	1.33	0.5024
non-tBRCAm	105	41 (39.0)	NE (NE,	NE)	67	25 (37.3)	NE (NE,	NE)	1.00	0.62,	1.67	0.9865
Interaction p-value												0.6566
First line treatment outo	come (eC	RF)										
NED [PDS]	89	30 (33.7)	NE (NE,	NE)	47	21 (44.7)	16.9 (8.3,	NE)	0.61	0.35,	1.09	0.0931
NED/CR [IDS]	74	37 (50.0)	16.6 (11.5,	NE)	32	11 (34.4)	NE (NE,	NE)	1.62	0.86,	3.34	0.1422
NED/CR [Chemo]	39	24 (61.5)	13.7 (3.0,2	24.2)	18	10 (55.6)	17.2 (3.0,	NE)	1.19	0.58,	2.61	0.6425
PR	50	21 (42.0)	25.7 (11.1,	NE)	34	16 (47.1)	17.0 (8.3,	NE)	0.77	0.41,	1.50	0.4416
Interaction p-value												0.1290
Screening laboratory tBRO	CA statu	s (eCRF)										
tBRCAm	147	71 (48.3)	22.2 (14.5,	NE)	67	33 (49.3)	16.9 (8.3,	NE)	0.88	0.59,	1.34	0.5382
non-tBRCAm	108	43 (39.8)	NE (NE,	NE)	65	25 (38.5)	NE (NE,	NE)	1.01	0.62,	1.67	0.9767
Interaction p-value												0.6736
Age group												
<65 years	185	90 (48.6)	22.2 (15.9,	NE)	98	46 (46.9)	17.2 (10.4,	NE)	0.95	0.67,	1.36	0.7637
>=65 years	70	24 (34.3)	NE (NE,	NE)	34	12 (35.3)	NE (NE,	NE)	0.96	0.49,	1.99	0.9099
Interaction p-value												0.9708

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3baf 24MAR2022:14:14 kvbv306

Table 2.3.3.6 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2					bevacizumab 132)				
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]			Number (%) of patients with events	(95% CI	Median time (95% CI) (months) [a]		95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	79 (43.4)	25.7 (19.4,	NE)	90	39 (43.3)	NE (NE,	NE)	0.95	0.65, 1.41	0.7936
IV	73	35 (47.9)	19.6 (9.0,	NE)	42	19 (45.2)	17.5 (8.7,	NE)	0.96	0.56, 1.71	0.8834
Interaction p-value											0.9781
Region											
Europe	245	110 (44.9)	25.3 (18.0,	NE)	126	56 (44.4)	21.9 (11.9,	NE)	0.92	0.67, 1.28	0.6165
Japan	10	4 (40.0)	NE (NE,	NE)	6	2 (33.3)	NE (NE,	NE)	1.78	0.35, 12.82	0.4957
Interaction p-value											0.4436
ECOG performance status at	Basel	ine									
(0) Normal activity	190	87 (45.8)	24.2 (15.9,	NE)	100	46 (46.0)	21.9 (11.0,	NE)	0.97	0.68, 1.40	0.8810
(1) Restricted activity	61	25 (41.0)	25.7 (16.6,	NE)	31	12 (38.7)	NE (NE,	NE)	0.89	0.46, 1.85	0.7529
Interaction p-value											0.8324
Baseline CA-125 value											
<=ULN	228	100 (43.9)	NE (NE,	NE)	118	51 (43.2)	, ,	,	0.93	0.67, 1.31	0.6756
>ULN	27	14 (51.9)	19.6 (5.6,	NE)	14	7 (50.0)	17.0 (6.6,	NE)	1.14	0.47, 3.00	0.7781
Interaction p-value											0.6804
Histological grade											
High grade	255	114 (44.7)	25.3 (18.6,	NE)	132	58 (43.9)	23.0 (12.7,	NE)	0.95	0.70, 1.31	0.7478
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	69 (41.6)	NE (NE,	NE)	80	32 (40.0)	NE (NE,	NE)	0.97	0.64, 1.49	0.8808
Residue	79	40 (50.6)	22.1 (8.9,	NE)	44	21 (47.7)	17.2 (6.0,	NE)	1.04	0.62, 1.79	0.8981
Interaction p-value											0.8465

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3baf 24MAR2022:14:14 kvbv306

Table 2.3.3.6 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + N=2					oevacizumab 132)					
Subgroup		Number (%) of patients with events	(95% CI)	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]		Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	58 (39.7)	NE (NE,	NE)	79	35 (44.3)	23.0 (8.6,	NE)	0.77	0.51,	1.18	0.2307
Interval	99	51 (51.5)	16.6 (11.5,	NE)	45	18 (40.0)	NE (NE,	NE)	1.41	0.84,	2.47	0.2020
Interaction p-value												0.0806
Myriad tumour BRCA mutat:	ion stat	us										
tBRCAm	158	77 (48.7)	22.1 (14.1,	NE)	77	34 (44.2)	17.5 (8.7,	NE)	1.02	0.69,	1.55	0.9132
Non-tBRCAm	97	37 (38.1)	NE (NE,	NE)	55	24 (43.6)	NE (NE,	NE)	0.82	0.49,	1.39	0.4545
Interaction p-value												0.5099
Status somatic BRCA mutat	tions											
sBRCAm	22	10 (45.5)	8.5 (2.8,	NE)	7	3 (42.9)	NE (NE,	NE)	1.40	0.43,	6.24	0.5995
gBRCAm	66	32 (48.5)	24.2 (13.9,	NE)	31	14 (45.2)	17.2 (8.7,	NE)	0.89	0.49,	1.73	0.7254
Non-BRCAm	41	19 (46.3)	25.7 (13.7,	NE)	22	9 (40.9)	NE (NE,	NE)	1.11	0.52,	2.59	0.7881
Interaction p-value												0.7942

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3baf 24MAR2022:14:14 kvbv306

Table 2.3.3.7 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Other single items (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

			bevacizumab 255)				bevacizumab 132)					
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV)	RS)										
NED [PDS]	92	31 (33.7)	NE (NE,	NE)	48	17 (35.4)	30.7 (19.4,	NE)	0.93	0.52,	1.72	0.8112
NED/CR [IDS]	74	35 (47.3)	24.7 (16.0,	NE)	38	9 (23.7)	NE (NE,	NE)	2.23	1.12,	4.94	0.0211*
NED/CR [Chemo]	40	12 (30.0)	NE (NE,	NE)	20	6 (30.0)	NE (NE,	NE)	1.11	0.43,	3.18	0.8382
PR	49	18 (36.7)	NE (NE,	NE)	26	13 (50.0)	23.5 (15.4,	NE)	0.63	0.31,	1.31	0.2070
Interaction p-value												0.0836
Screening laboratory tBR0	CA statu	s (IVRS)										
tBRCAm	150	62 (41.3)	26.3 (24.7,	NE)	65	22 (33.8)	NE (NE,	NE)	1.23	0.77,	2.04	0.4014
non-tBRCAm	105	34 (32.4)	NE (NE,	NE)	67	23 (34.3)	28.0 (22.1,	NE)	0.97	0.58,	1.67	0.9226
Interaction p-value												0.5278
First line treatment out	come (eCl	RF)										
NED [PDS]	89	28 (31.5)	NE (NE,	NE)	47	16 (34.0)	30.7 (19.5,	NE)	0.94	0.51,	1.78	0.8435
NED/CR [IDS]	74	33 (44.6)	26.3 (16.0,	NE)	32	9 (28.1)	NE (NE,	NE)	1.68	0.84,	3.72	0.1512
NED/CR [Chemo]	39	13 (33.3)	NE (NE,	NE)	18	5 (27.8)	NE (NE,	NE)	1.20	0.45,	3.75	0.7221
PR	50	20 (40.0)	25.3 (22.0,	NE)	34	15 (44.1)	25.4 (19.4,	NE)	0.85	0.44,	1.69	0.6375
Interaction p-value												0.5352
Screening laboratory tBR0	CA statu	s (eCRF)										
tBRCAm	147	62 (42.2)	26.0 (24.7,	NE)	67	22 (32.8)	NE (NE,	NE)	1.29	0.80,	2.14	0.3024
non-tBRCAm	108	34 (31.5)	NE (NE,	NE)	65	23 (35.4)	28.0 (22.1,	NE)	0.91	0.54,	1.57	0.7397
Interaction p-value												0.3516
Age group												
<65 years	185	73 (39.5)	29.0 (25.0,	NE)	98	32 (32.7)	30.7 (25.4,	NE)	1.29	0.86,	1.97	0.2285
>=65 years	70	23 (32.9)	NE (NE,	NE)	34	13 (38.2)	NE (NE,	NE)	0.73	0.37,	1.48	0.3725
Interaction p-value												0.1703

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bag 24MAR2022:14:14 kvbv306

Table 2.3.3.7 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Other single items (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olapar	rib + 1 (N=2		zumab	•		Plac	ebo +] (N=)	oevaci 132)	zumab					
Subgroup		Number (%) of patients n with events		Median time (95% CI) (months) [a]		Number (%) of patients n with events			Median time (95% CI) (months) [a]			Hazard ratio [b]	95% CI [b]		2-sided p-value [b]	
FIGO Stage (Disease state)																
III	182	67 (36.8)	NE	(NE,	NE)	90	27	(30.0)	NE	(NE,	NE)	1.29	0.83,	2.05	0.2570
IV	73	29 (39.7)	26.0	(22.0,	NE)	42	18	(42.9)	25.4	(16.6,	NE)	0.85	0.48,	1.56	0.5941
Interaction p-value																0.2726
Region																
Europe	245	92 (37.6)	30.4	(25.3,	NE)	126	43	(34.1)	28.0	(25.4,	NE)	1.10	0.77,	1.59	0.6104
Japan	10	4 (4	40.0)	NE	(NE,	NE)	6	2	(33.3)	NE	(NE,	NE)	1.57	0.31,	11.29	0.5974
Interaction p-value																0.6845
ECOG performance status at	Basel	ine														
(0) Normal activity	190	73 (38.4)	29.0	(24.9,	NE)	100	35	(35.0)	30.7	(25.4,	NE)	1.21	0.81,	1.83	0.3525
(1) Restricted activity	61	23 (37.7)	26.3	(24.8,	NE)	31	10	(32.3)	NE	(NE,	NE)	0.95	0.47,	2.10	0.9007
Interaction p-value																0.5867
Baseline CA-125 value																
<=ULN	228	87 (3	38.2)	30.4	(25.0,	NE)	118	40	(33.9)	30.7	(25.8,	NE)	1.13	0.78,	1.65	0.5348
>ULN	27	9 (3	33.3)	25.3	(20.0,	NE)	14	5	(35.7)	25.4	(11.3,	NE)	1.02	0.35,	3.33	0.9660
Interaction p-value																0.8738
Histological grade																
High grade	255	96 (3	37.6)	30.4	(25.3,	NE)	132	45	(34.1)	28.0	(25.4,	NE)	1.11	0.79,	1.60	0.5482
Interaction p-value																NC
Cytoreductive surgery outco	me															
No residue	166	63 (38.0)	29.0	(26.0,	NE)	80	25	(31.3)	30.7	(25.8,	NE)	1.25	0.80,	2.03	0.3333
Residue	79	26 (32.9)	NE	(NE,	NE)	44	16	(36.4)	28.0	(22.1,	NE)	0.90	0.49,	1.71	0.7344
Interaction p-value																0.4015

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bag 24MAR2022:14:14 kvbv306

Table 2.3.3.7 PAOLA1 Appendix: Summary of subgroup analysis of EORTC QLQ-OV28 Symptom scale/items: Other single items (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + N=2				bevacizumab 132)						
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients n with events				Hazard ratio [b] 95% CI [b]		[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	48 (32.9)	NE (NE,	NE)	79	27 (34.2)	30.7 (22.1,	NE)	0.94	0.59,	1.53	0.7964
Interval	99	41 (41.4)	26.3 (21.9,	NE)	45	14 (31.1)	28.0 (23.5,	NE)	1.45	0.81,	2.76	0.2174
Interaction p-value												0.2644
Myriad tumour BRCA mutat:	ion stat	us										
tBRCAm	158	61 (38.6)	26.3 (24.9,	NE)	77	23 (29.9)	30.7 (25.8,	NE)	1.29	0.81,	2.12	0.2934
Non-tBRCAm	97	35 (36.1)	NE (NE,	NE)	55	22 (40.0)	25.4 (19.4,	NE)	0.94	0.55,	1.62	0.8096
Interaction p-value												0.3845
Status somatic BRCA mutat	cions											
sBRCAm	22	8 (36.4)	NE (NE,	NE)	7	2 (28.6)	NE (NE,	NE)	1.42	0.36,	9.43	0.6453
gBRCAm	66	28 (42.4)	26.3 (24.7,	NE)	31	8 (25.8)	NE (NE,	NE)	1.69	0.81,	3.98	0.1703
Non-BRCAm	41	16 (39.0)	NE (NE,	NE)	22	9 (40.9)	NE (NE,	NE)	0.92	0.41,	2.18	0.8429
Interaction p-value												0.5697

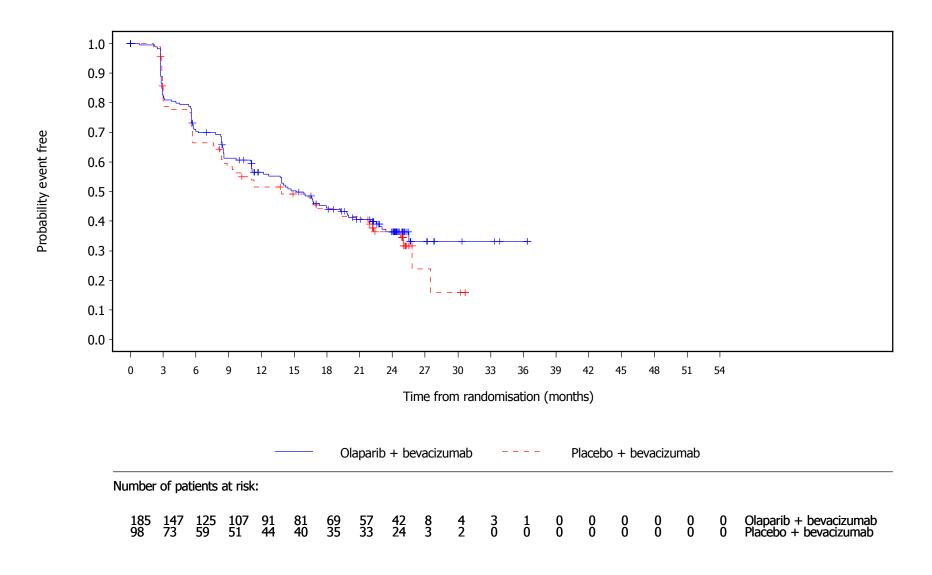
Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3bag 24MAR2022:14:14 kvbv306

Figure 2.3.4.1 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 15) time to clinically meaningful deterioration for Age group=<65 years

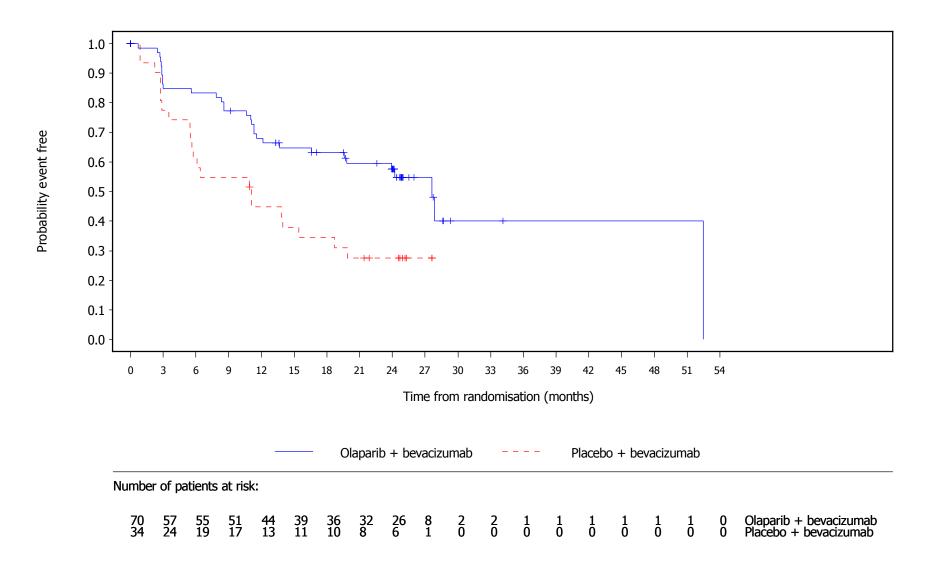
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3eaa 24MAR2022:14:14 kvbv306

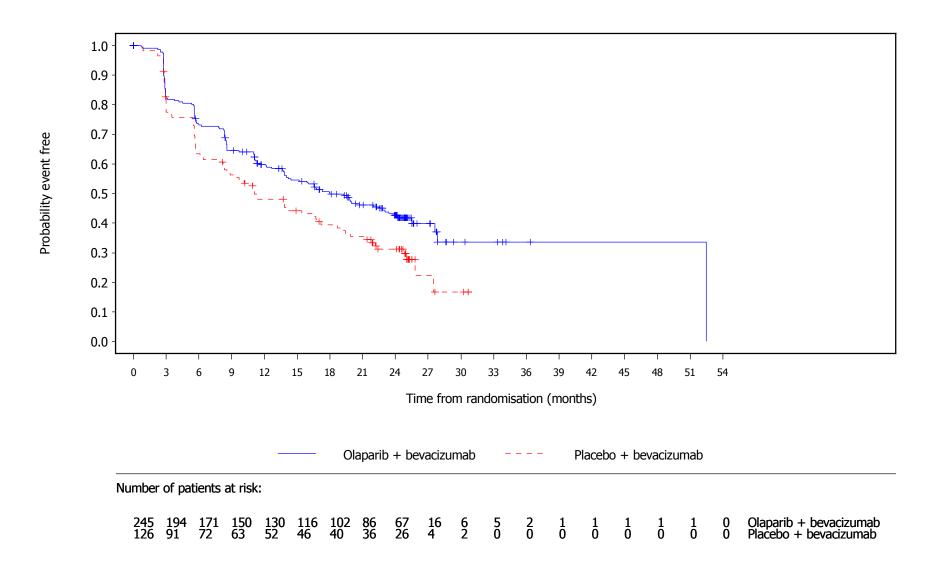
Figure 2.3.4.2 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 15) time to clinically meaningful deterioration for Age group=>=65 years

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



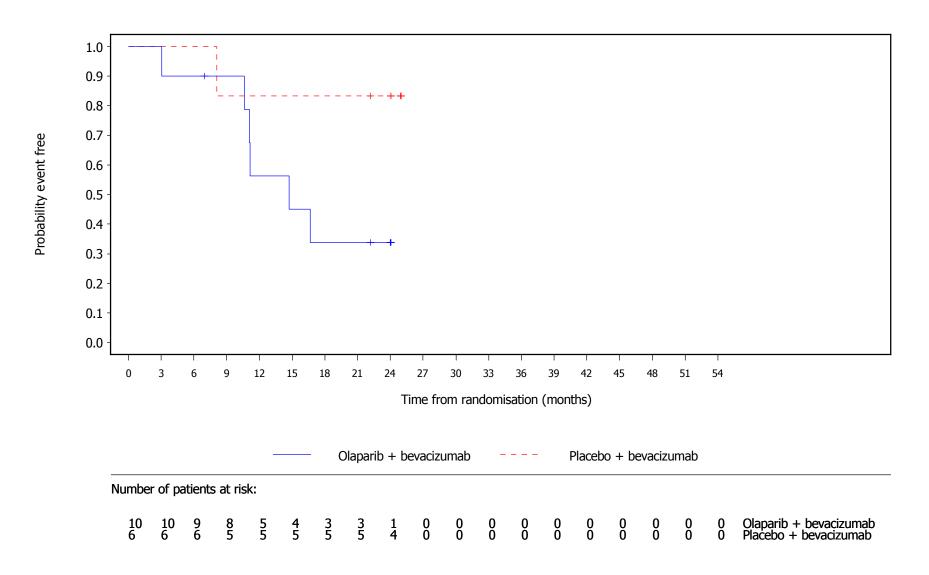
Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3eab 24MAR2022:14:14 kvbv306

Figure 2.3.4.3 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 15) time to clinically meaningful deterioration for Region=Europe
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3eac 24MAR2022:14:14 kvbv306

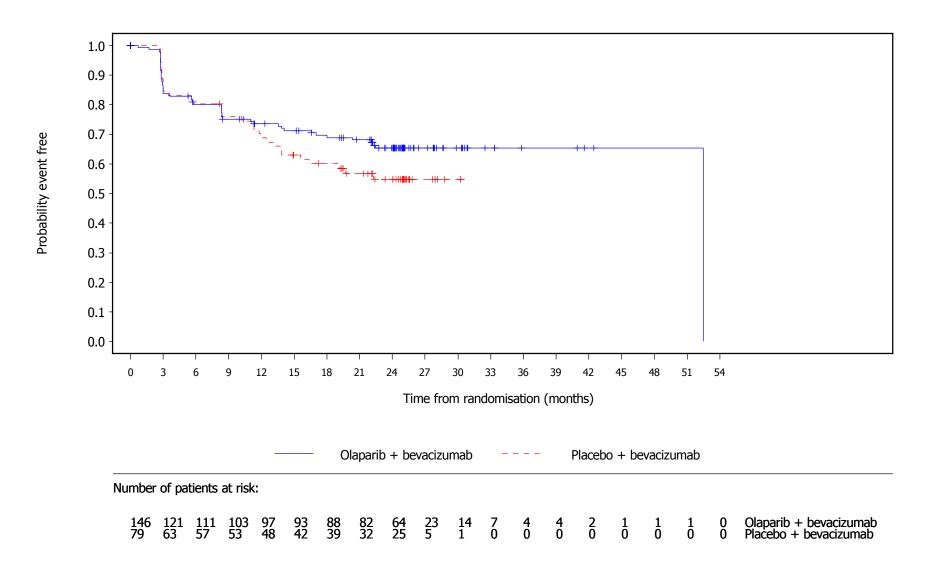
Figure 2.3.4.4 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 15) time to clinically meaningful deterioration for Region=Japan
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3ead 24MAR2022:14:14 kvbv306

Figure 2.3.4.5 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (MID = 15) time to clinically meaningful deterioration for Timing of cytoreductive surgery=Upfront

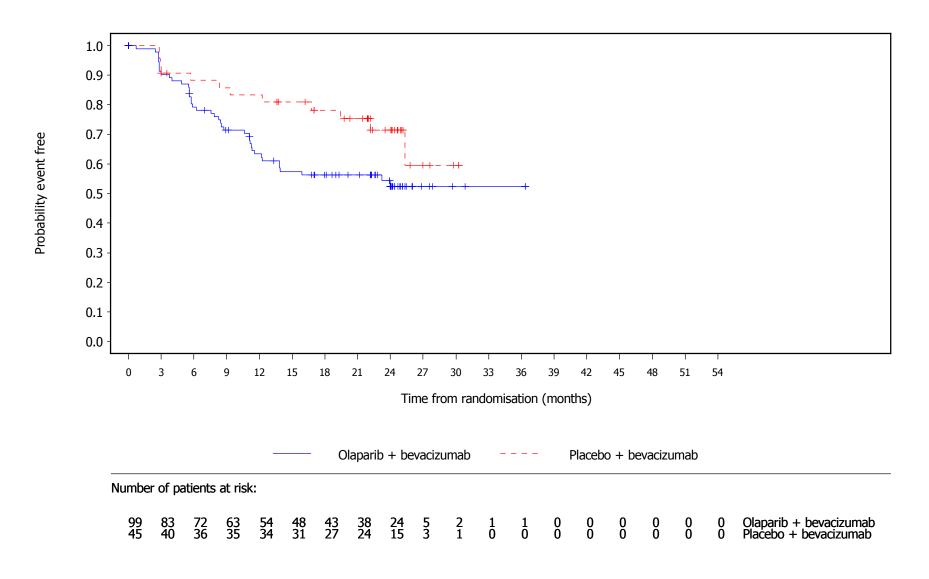
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3eae 24MAR2022:14:14 kvbv306

Figure 2.3.4.6 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (MID = 15) time to clinically meaningful deterioration for Timing of cytoreductive surgery=Interval

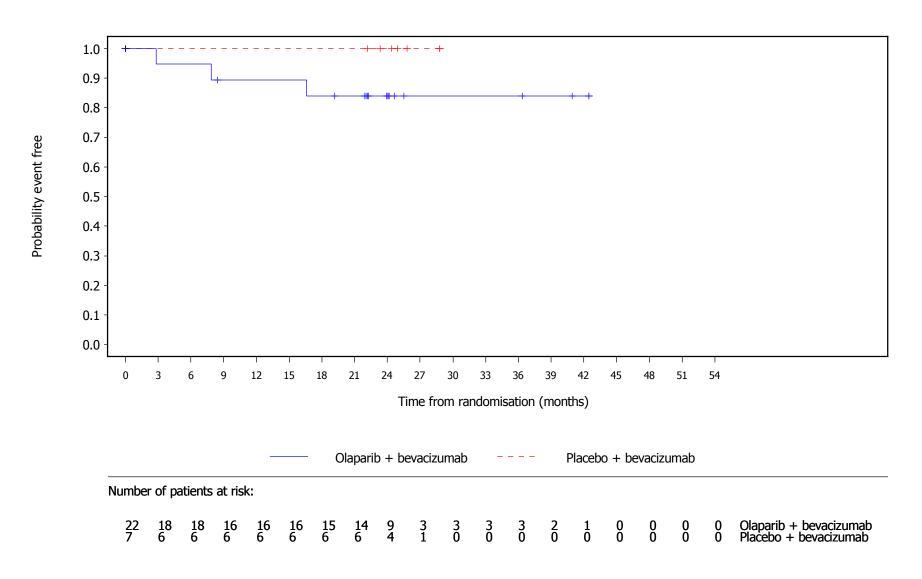
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3eaf 24MAR2022:14:14 kvbv306

Figure 2.3.4.7 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (MID = 15) time to clinically meaningful deterioration for Status somatic BRCA mutations=sBRCAm

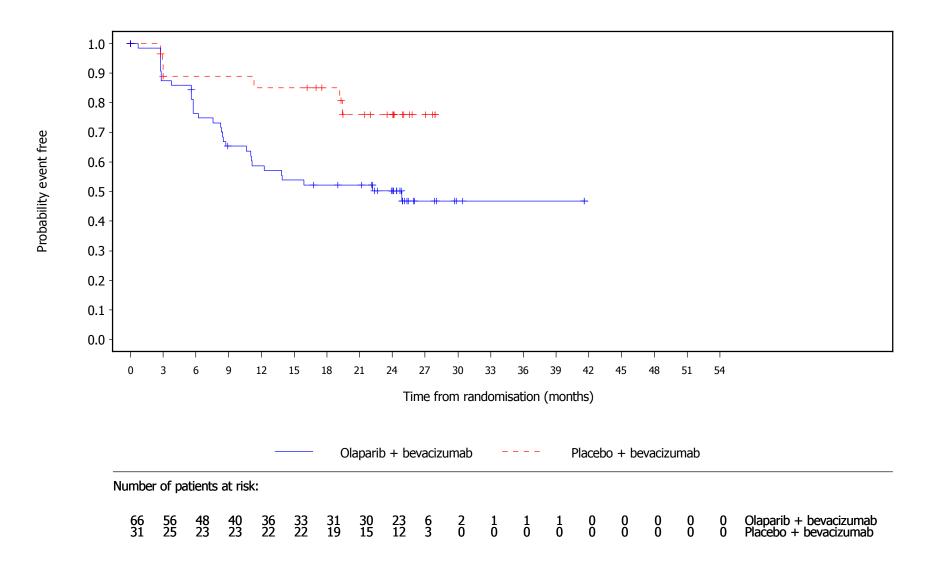
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3eag 24MAR2022:14:14 kvbv306

Figure 2.3.4.8 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (MID = 15) time to clinically meaningful deterioration for Status somatic BRCA mutations=gBRCAm

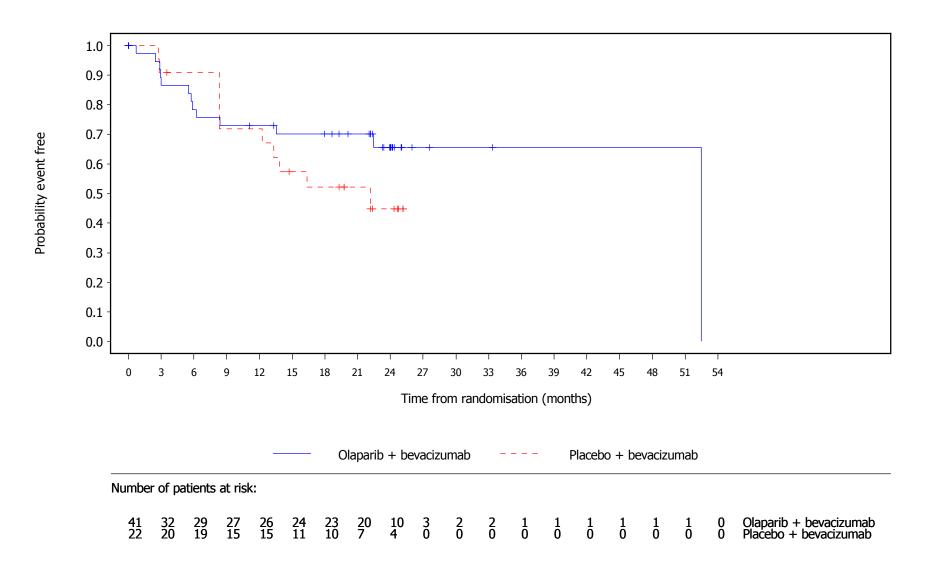
Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3eah 24MAR2022:14:14 kvbv306

Figure 2.3.4.9 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (MID = 15) time to clinically meaningful deterioration for Status somatic BRCA mutations=Non-BRCAm

Full Analysis Set, HRD[42] positive, DCO 22MAR2020



Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3eai 24MAR2022:14:14 kvbv306

Table 2.4.3.1 PAOLA1 Appendix: Summary of subgroup analysis of EQ-5D-5L Visual analogue scale (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	:	Olaparib + k (N=2				oevacizumab 132)			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVI	RS)							
NED [PDS]	92	32 (34.8)	NE (NE, NE)	48	22 (45.8)	24.1 (16.4, NE)	0.73	0.43, 1.28	0.2659
NED/CR [IDS]	74	41 (55.4)	18.6 (11.4,24.9)	38	16 (42.1)	22.1 (12.1, NE)	1.34	0.77, 2.46	0.3098
NED/CR [Chemo]	40	17 (42.5)	NE (NE, NE)	20	8 (40.0)	26.7 (5.7, NE)	1.17	0.52, 2.86	0.7146
PR	49	26 (53.1)	14.5 (5.7, NE)	26	12 (46.2)	19.9 (7.9, NE)	1.14	0.59, 2.35	0.7028
Interaction p-value									0.4791
Screening laboratory tBR0	CA status	s (IVRS)							
tBRCAm	150	66 (44.0)	NE (NE, NE)	65	25 (38.5)	NE (NE, NE)	1.19	0.76, 1.91	0.4636
non-tBRCAm	105	50 (47.6)	19.9 (13.6, NE)	67	33 (49.3)	21.9 (13.9,27.8)	0.96	0.62, 1.51	0.8638
Interaction p-value									0.5198
First line treatment out	come (eCI	RF)							
NED [PDS]	89	32 (36.0)	NE (NE, NE)	47	22 (46.8)	24.1 (13.8, NE)	0.74	0.43, 1.29	0.2783
NED/CR [IDS]	74	42 (56.8)	17.5 (11.3,24.2)	32	13 (40.6)	22.1 (12.1, NE)	1.42	0.79, 2.76	0.2541
NED/CR [Chemo]	39	16 (41.0)	NE (NE, NE)	18	8 (44.4)	26.7 (3.5, NE)	0.95	0.42, 2.33	0.8989
PR	50	25 (50.0)	17.5 (6.0, NE)	34	15 (44.1)	23.5 (11.3, NE)	1.20	0.64, 2.32	0.5825
Interaction p-value									0.4329
Screening laboratory tBR0	CA status	s (eCRF)							
tBRCAm	147	64 (43.5)	NE (NE, NE)	67	25 (37.3)	NE (NE, NE)	1.20	0.76, 1.94	0.4373
non-tBRCAm	108	52 (48.1)	19.9 (13.7, NE)	65	33 (50.8)	21.9 (13.9,27.8)	0.94	0.61, 1.47	0.7966
Interaction p-value									0.4611
Age group									
<65 years	185	84 (45.4)	25.3 (16.8, NE)	98	40 (40.8)	27.8 (22.1, NE)	1.14	0.78, 1.67	0.5068
>=65 years	70	32 (45.7)	24.2 (11.4, NE)	34	18 (52.9)	18.7 (5.8, NE)	0.82	0.47, 1.49	0.5058
Interaction p-value									0.3594

Time to worsening is defined as time from randomisation until earliest clinically meaningful worsening. Patients who have not shown a clinically meaningful worsening or who cannot experience a worsening will be censored at last PRO assessment. Patients who died <168 days of last PRO assessment will be censored at death. Those with no evaluable baseline will be censored at randomisation.

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3caa 24MAR2022:14:14 kvbv306

Table 2.4.3.1 PAOLA1 Appendix: Summary of subgroup analysis of EQ-5D-5L Visual analogue scale (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)							Placebo + bevacizumab (N=132)								
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]		Number (%) of patients n with events				Median time (95% CI) (months) [a]		Hazard ratio [b]	95% CI [b]		2-sided p-value [b]		
FIGO Stage (Disease state)																
III	182	85	(46.7)	24.0	(14.0,	NE)	90	39	(43.3)	27.8	(18.7,	NE)	1.12	0.77,	1.66	0.5478
IV	73	31	(42.5)	25.3	(16.8,	NE)	42	19	(45.2)	23.5	(8.5,	NE)	0.87	0.50,	1.57	0.6341
Interaction p-value																0.4678
Region																
Europe	245	112	(45.7)	24.9	(17.3,	NE)	126	56	(44.4)	26.7	(17.4,	NE)	1.02	0.74,	1.41	0.9269
Japan	10	4	(40.0)	NE	(NE,	NE)	6	2	(33.3)	NE	(NE,	NE)	1.63	0.32,	11.77	0.5629
Interaction p-value																0.5823
ECOG performance status at	Basel	ine														
(0) Normal activity	190	89	(46.8)	24.2	(16.4,	NE)	100	46	(46.0)	26.7	(17.0,	NE)	1.04	0.73,	1.50	0.8180
(1) Restricted activity	61	26	(42.6)	25.3	(13.7,	NE)	31	12	(38.7)	24.0	(11.3,	NE)	1.04	0.54,	2.13	0.9137
Interaction p-value																0.9920
Baseline CA-125 value																
<=ULN	228		(43.4)	NE		NE)	118				(21.9,		1.01		1.43	0.9602
>ULN	27	17	(63.0)	5.8	(3.0,	25.3)	14	8	(57.1)	17.0	(5.6,	NE)	1.38	0.61,	3.39	0.4440
Interaction p-value																0.4921
Histological grade																
High grade	255	116	(45.5)	25.3	(17.5,	NE)	132	58	(43.9)	26.7	(19.9,	NE)	1.04	0.76,	1.44	0.8041
Interaction p-value																NC
Cytoreductive surgery outco	me															
No residue	166	75	(45.2)	24.9	(16.6,	NE)	80	35	(43.8)	24.1	(17.4,	NE)	1.02		1.54	0.9323
Residue	79	35	(44.3)	NE	(NE,	NE)	44	18	(40.9)	26.7	(14.7,	NE)	1.16	0.67,	2.10	0.5962
Interaction p-value																0.7030

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[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3caa 24MAR2022:14:14 kvbv306

Table 2.4.3.1 PAOLA1 Appendix: Summary of subgroup analysis of EQ-5D-5L Visual analogue scale (MID = 15) time to clinically meaningful deterioration

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	•	-	bevacizumab 255)			bevacizumab 132)	:				
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]	n	Number (%) of patients with events			Hazard ratio [b] 95% CI [b]		[b]	2-sided p-value [b]
Timing of cytoreductive s	urgery										
Upfront	146	56 (38.4)	NE (NE, NE)	79	33 (41.8)	27.8 (18.7,	NE)	0.91	0.59,	1.41	0.6582
Interval	99	54 (54.5)	18.6 (11.5,24.9)	45	20 (44.4)	23.5 (13.9,	NE)	1.30	0.79,	2.22	0.3097
Interaction p-value											0.2912
Myriad tumour BRCA mutati	on stat	us									
tBRCAm	158	72 (45.6)	25.3 (16.8, NE)	77	29 (37.7)	27.8 (22.1,	NE)	1.24	0.81,	1.93	0.3244
Non-tBRCAm	97	44 (45.4)	24.9 (13.6, NE)	55	29 (52.7)	19.9 (8.5,	NE)	0.85	0.53,	1.37	0.4893
Interaction p-value											0.2419
Status somatic BRCA mutat	ions										
sBRCAm	22	6 (27.3)	NE (NE, NE)	7	2 (28.6)	NE (NE,	NE)	1.29	0.30,	8.80	0.7511
gBRCAm	66	33 (50.0)	22.6 (13.8, NE)	31	11 (35.5)	NE (NE,	NE)	1.54	0.80,	3.19	0.2013
Non-BRCAm	41	18 (43.9)	NE (NE, NE)	22	11 (50.0)	21.9 (3.5,	NE)	0.83	0.40,	1.81	0.6221
Interaction p-value											0.4843

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[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, model includes treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s2/tlf/prod/program/ttesubpr3.sas ettesubpr3caa 24MAR2022:14:14 kvbv306

Table 3.1.1 PAOLA1 Appendix: Summary of analysis of time to sustained worsening in EORTC QLQ-C30 global QoL/health status, functional, symptom and single item scales (MID=10)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + 1 (N=2				oevacizumab 132)				
		Number (%) of patients with events	Median time (95% CI) (months) [a]	(Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [c]
EORTC QLQ-C30 Global QoL/health status (MID = 10)	255	66 (25.9)	38.7 (30.9, NE)	132	40 (30.3)	28.0 (25.6, NE)	0.78	0.52,	1.17	0.2194
EORTC QLQ-C30 Functional scale: Physical (MID = 10)	255	38 (14.9)	52.5 (NE, NE)	132	37 (28.0)	28.0 (26.5, NE)	0.48	0.30,	0.77	0.0016*
EORTC QLQ-C30 Functional scale: Role (MID = 10)	255	59 (23.1)	44.9 (38.6, NE)	132	46 (34.8)	27.7 (24.9, NE)	0.58	0.39,	0.87	0.0073*
EORTC QLQ-C30 Functional scale: Cognitive (MID = 10)	255	78 (30.6)	30.4 (27.8, NE)	132	46 (34.8)	27.8 (24.9, NE)	0.80	0.55,	1.18	0.2553
EORTC QLQ-C30 Functional scale: Emotional (MID = 10)	255	72 (28.2)	41.9 (31.0, NE)	132	48 (36.4)	26.5 (24.9,30.7)	0.63	0.43,	0.92	0.0156*
EORTC QLQ-C30 Functional scale: Social (MID = 10)	255	62 (24.3)	41.9 (41.9, NE)	132	38 (28.8)	29.3 (25.4, NE)	0.74	0.49,	1.13	0.1504
EORTC QLQ-C30 Single item symptom scale: Loss of appetite (MID = 10)	255	41 (16.1)	49.3 (NE, NE)	132	32 (24.2)	28.7 (27.8, NE)	0.60	0.37,	0.97	0.0322*
EORTC QLQ-C30 Single item symptom scale: Constipation (MID = 10)	255	66 (25.9)	NE (NE, NE)	132	34 (25.8)	28.7 (25.5, NE)	0.99	0.65,	1.52	0.9394

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Table 3.1.1 PAOLA1 Appendix: Summary of analysis of time to sustained worsening in EORTC QLQ-C30 global QoL/health status, functional, symptom and single item scales (MID=10)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

		Olaparib + k (N=2				pevacizumab 132)				
	Number (%) of patients n with events		Median time (95% CI) (months) [a]	Number (%) of patients n with events			Hazard ratio [b]	95% CI [b]		2-sided p-value [c]
EORTC QLQ-C30 Single item symptom scale: Diarrhoea (MID = 10)	255	34 (13.3)	NE (NE, NE)	132	14 (10.6)	NE (NE, NE)	1.06	0.58,	2.06	0.8573
EORTC QLQ-C30 Single item symptom scale: Dyspnoea (MID = 10)	255	52 (20.4)	52.5 (NE, NE)	132	36 (27.3)	NE (NE, NE)	0.70	0.45,	1.09	0.1030
EORTC QLQ-C30 Symptom scale: Fatigue (MID = 10)	255	88 (34.5)	38.7 (25.7, NE)	132	57 (43.2)	25.8 (24.2,28.2)	0.71	0.50, 1.003		0.0479*
EORTC QLQ-C30 Single item symptom scale: Financial difficulties (MID = 10)	255	31 (12.2)	NE (NE, NE)	132	23 (17.4)	NE (NE, NE)	0.60	0.34,	1.06	0.0706
EORTC QLQ-C30 Symptom scale: Nausea and vomiting (MID = 10)	255	52 (20.4)	52.5 (NE, NE)	132	29 (22.0)	NE (NE, NE)	0.90	0.57,	1.46	0.6658
EORTC QLQ-C30 Symptom scale: Pain (MID = 10)	255	78 (30.6)	35.9 (29.9, NE)	132	56 (42.4)	24.9 (23.5,30.7)	0.57	0.40,	0.81	0.0014*
EORTC QLQ-C30 Single item symptom scale: Insomnia (MID = 10)	255	77 (30.2)	33.4 (26.0, NE)	132	41 (31.1)	28.2 (25.5, NE)	0.86	0.59,	1.28	0.4502

 [[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.
 [b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.
 [c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr3.sas ettemainpr3ae 04OCT2022:09:12 kpzx329

Figure 3.1.2.1 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Global QoL/health status (MID = 10) time to sustained worsening Full Analysis Set, HRD[42] positive, DCO 22Mar2020

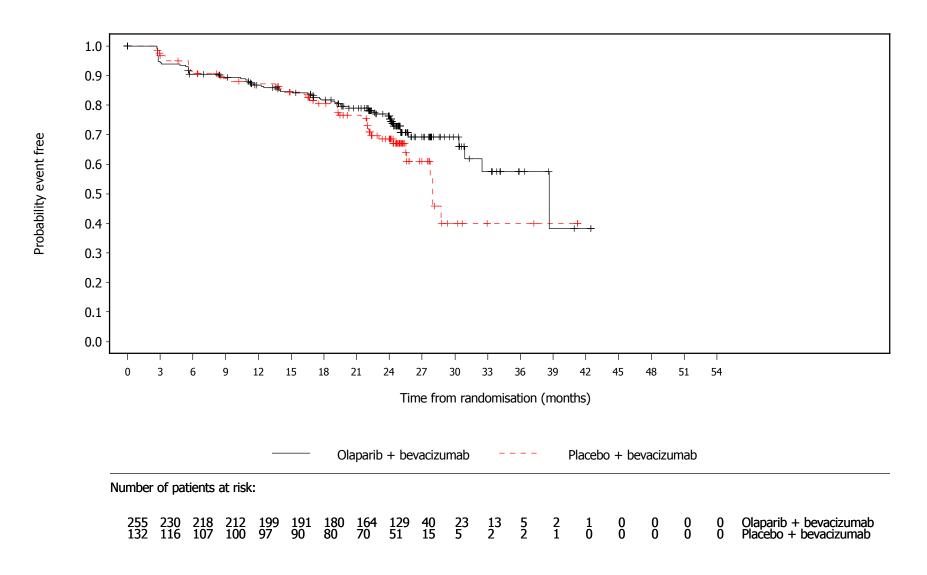


Figure 3.1.2.2 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Physical (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

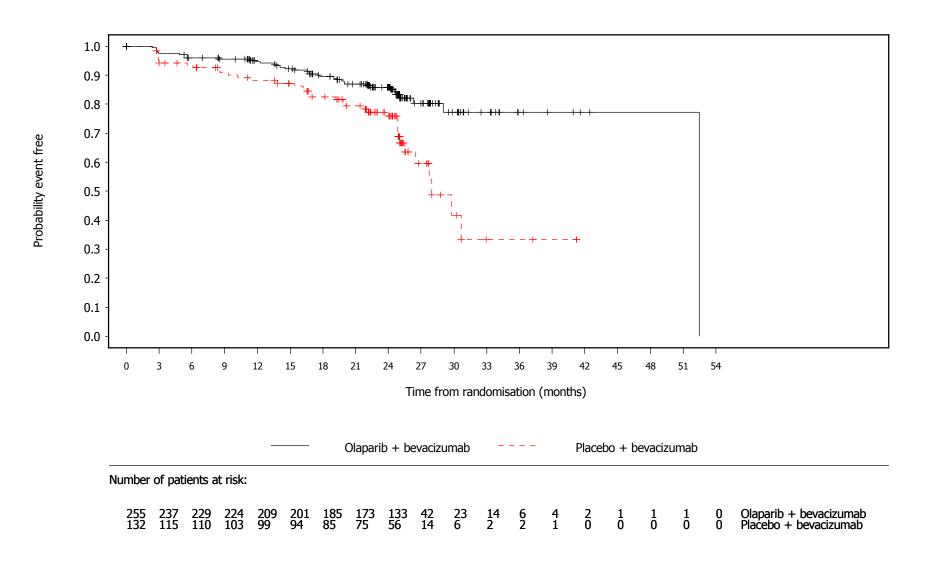


Figure 3.1.2.3 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Role (MID = 10) time to sustained worsening Full Analysis Set, HRD[42] positive, DCO 22Mar2020

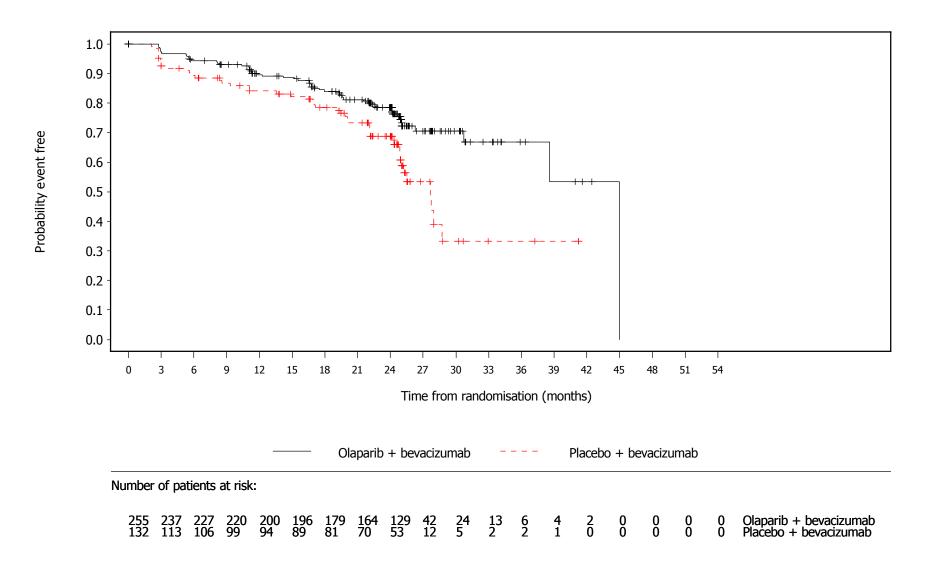


Figure 3.1.2.4 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Cognitive (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

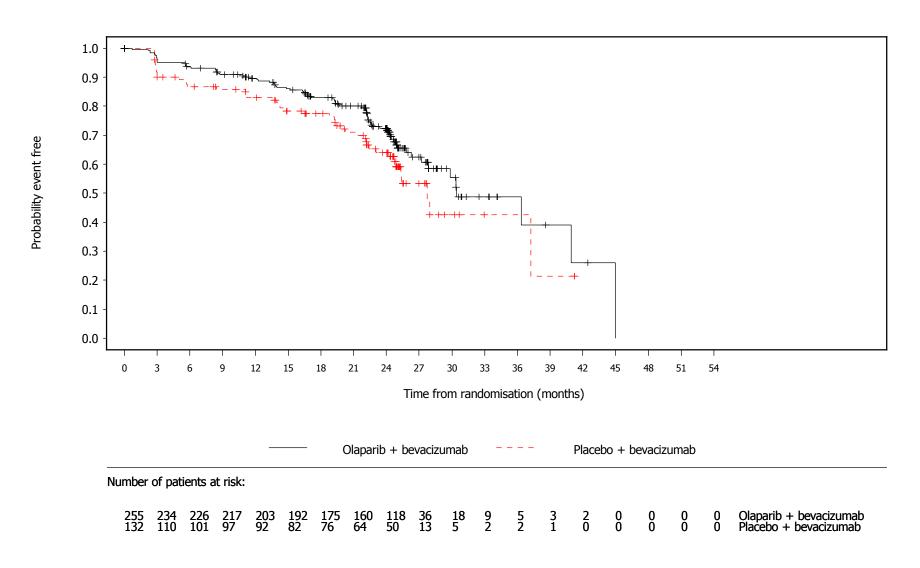


Figure 3.1.2.5 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Emotional (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

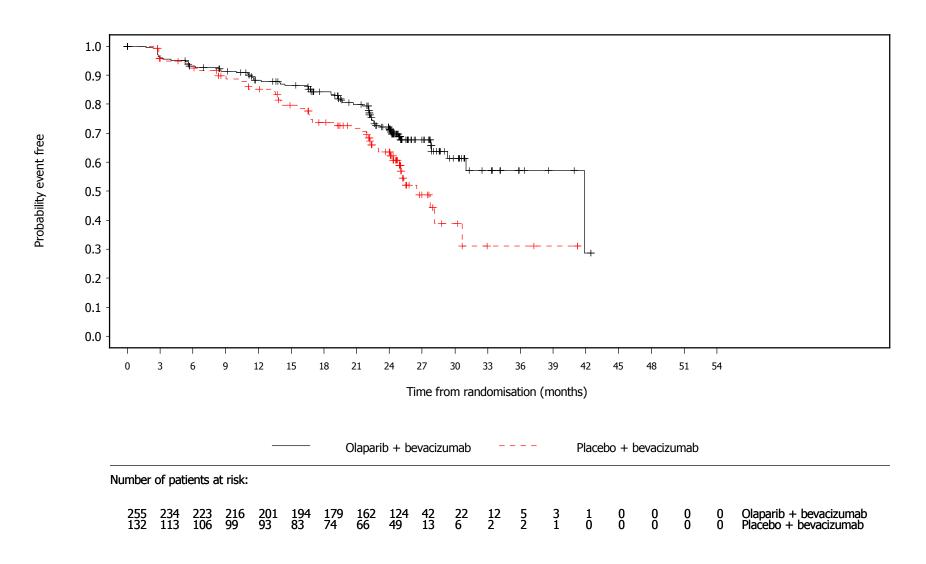


Figure 3.1.2.6 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Social (MID = 10) time to sustained worsening Full Analysis Set, HRD[42] positive, DCO 22Mar2020

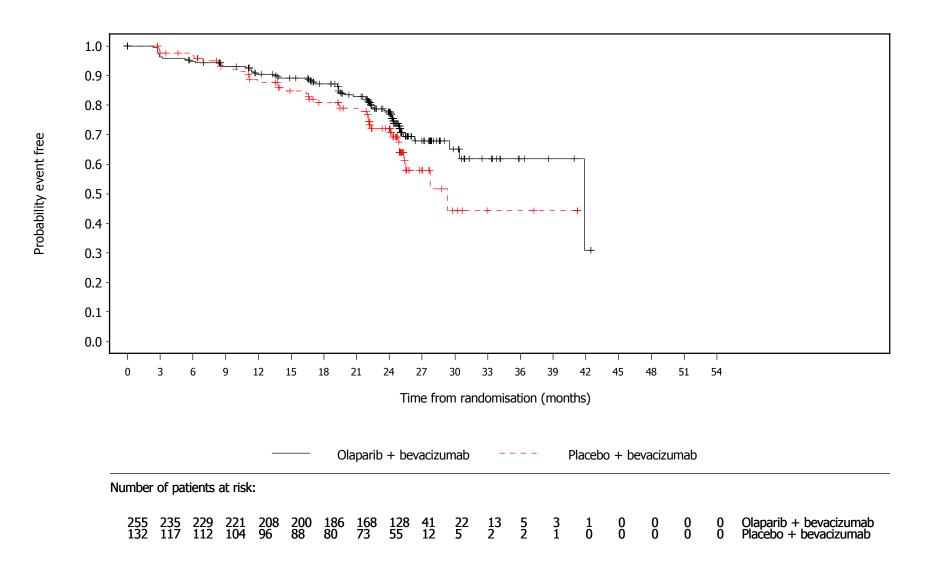


Figure 3.1.2.7 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Loss of appetite (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

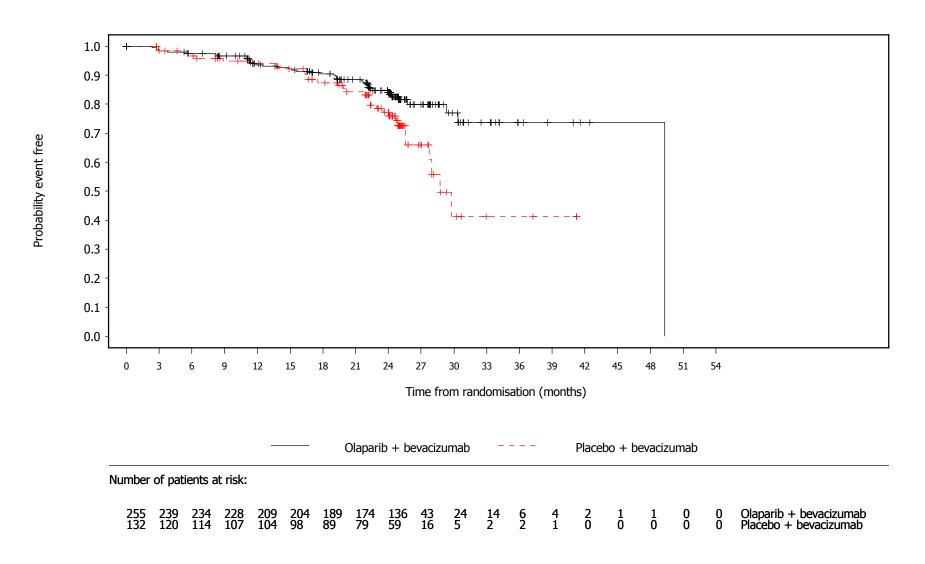


Figure 3.1.2.8 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Constipation (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

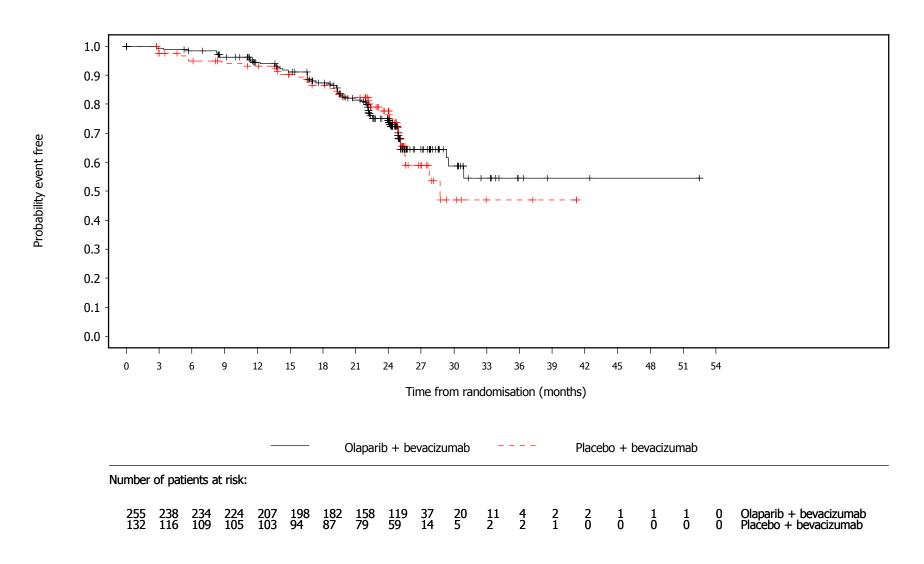


Figure 3.1.2.9 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Diarrhoea (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

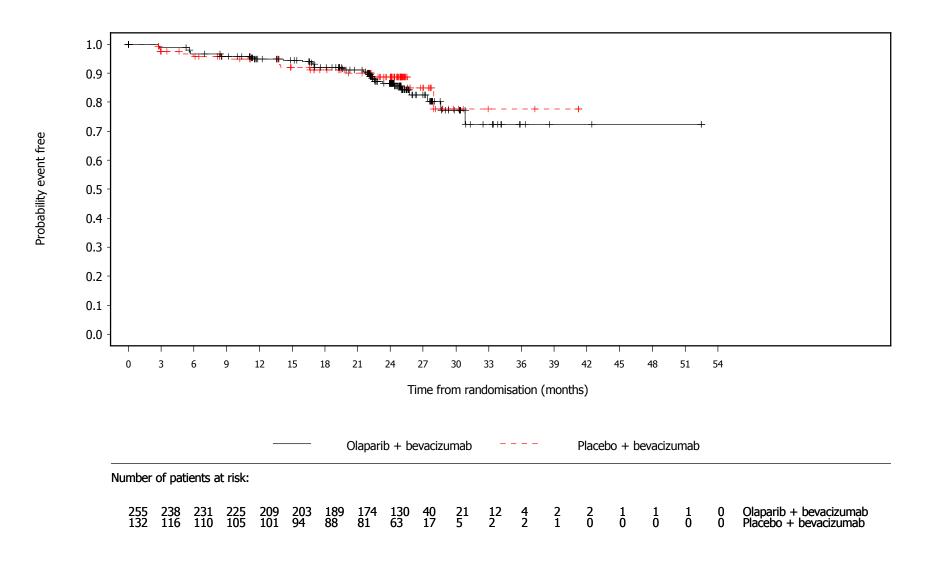


Figure 3.1.2.10 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Dyspnoea (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

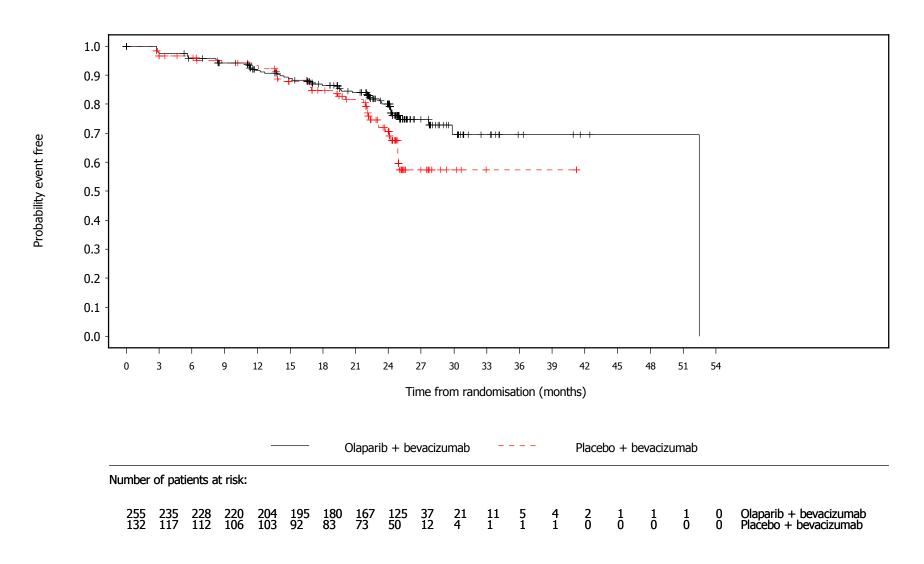


Figure 3.1.2.11 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Fatigue (MID = 10) time to sustained worsening Full Analysis Set, HRD[42] positive, DCO 22Mar2020

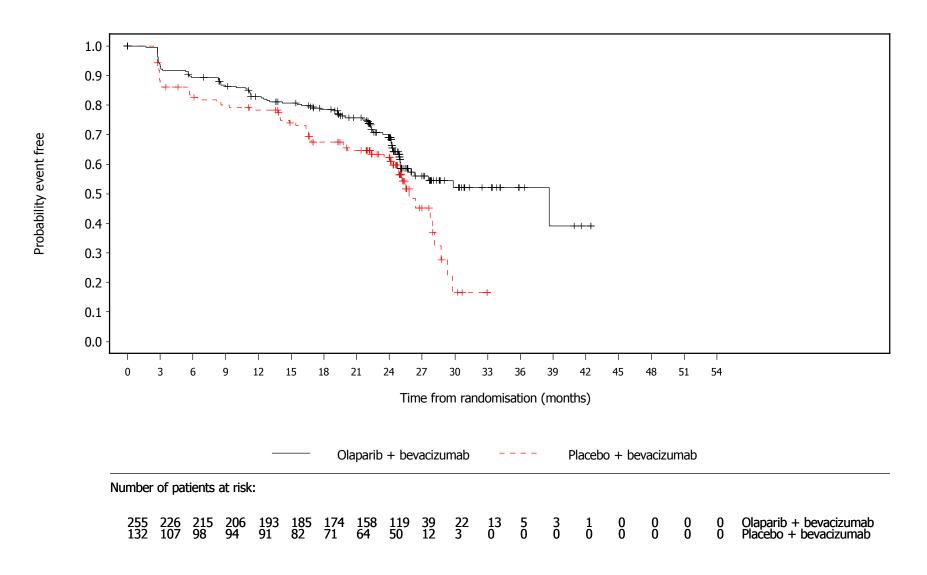


Figure 3.1.2.12 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Financial difficulties (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

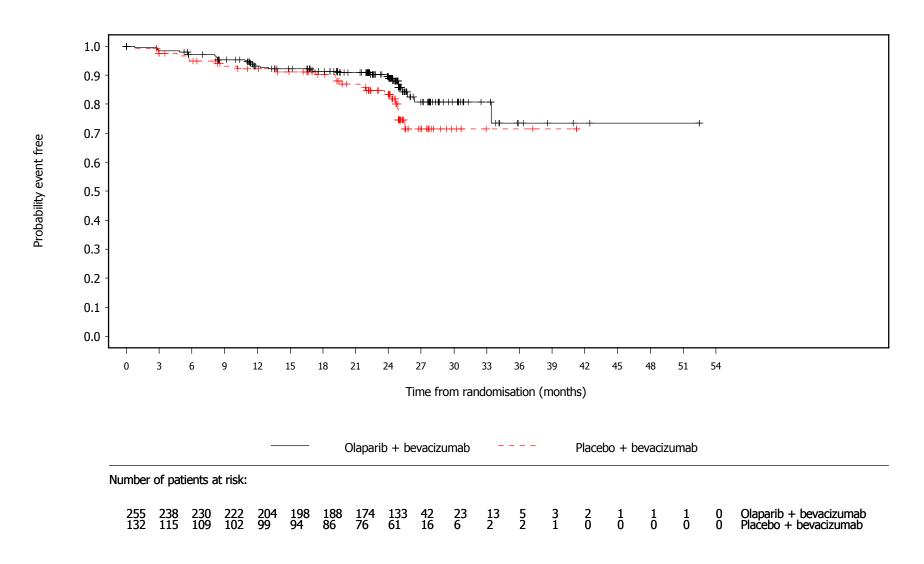


Figure 3.1.2.13 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Nausea and vomiting (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

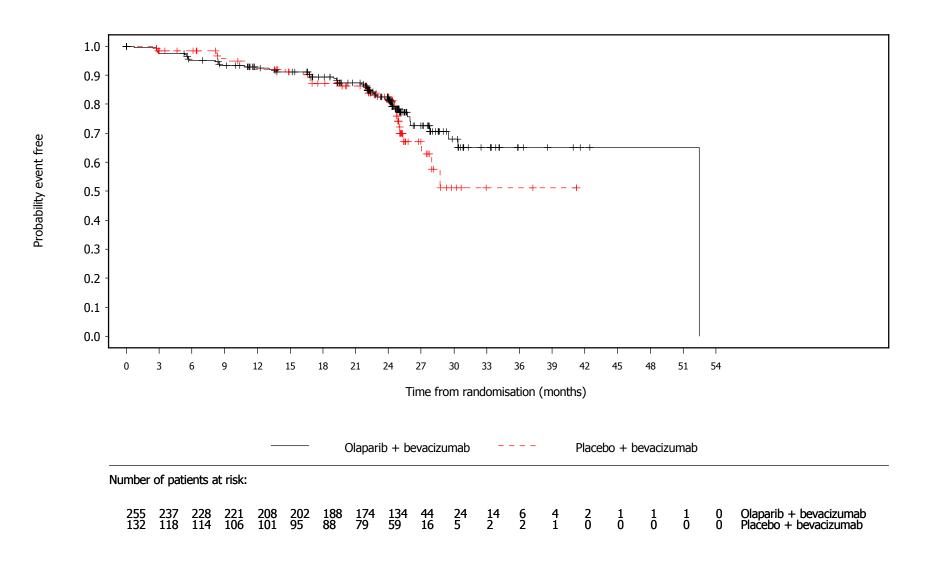


Figure 3.1.2.14 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Pain (MID = 10) time to sustained worsening
Full Analysis Set, HRD[42] positive, DCO 22Mar2020

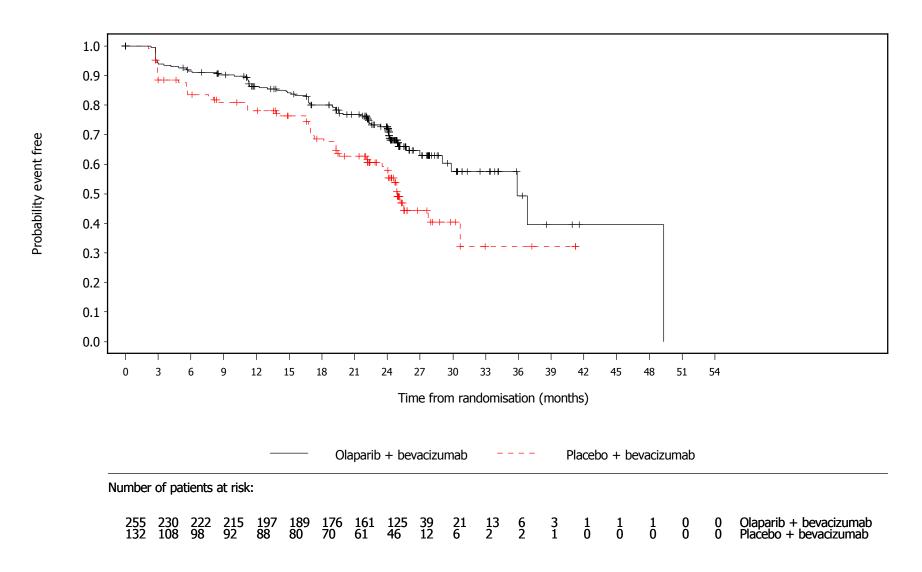


Figure 3.1.2.15 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Insomnia (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

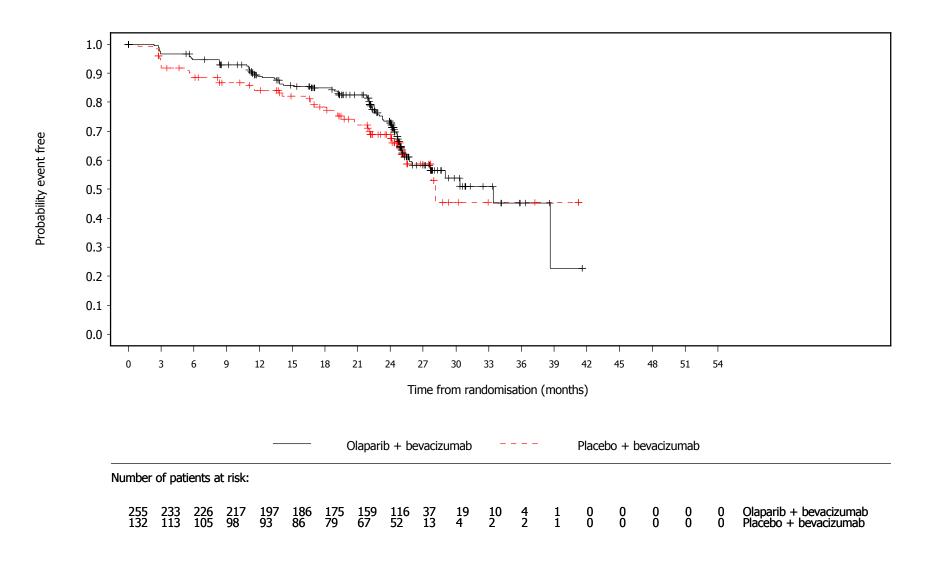


Table 3.1.3 PAOLA1 Appendix: Summary of analysis of time to sustained worsening in EORTC QLQ-C30 global QoL/health status, functional, symptom and single item scales (MID=10)

Sensitivity Analysis I (censoring patients with only one worsening post baseline and no subsequent observations)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olap	parib + bo (N=25	evacizumab 55)		Placebo + be						
	of pa	per (%) atients events	Median time (95% CI) (months) [a]	Number (%) of patients n with events		Median time (95% CI) (months) [a]		Hazard ratio [b]	95% CI [b]		2-sided p-value [c]
EORTC QLQ-C30 Global QoL/health status (MID = 10)	255 35	(13.7)	NE (NE, NE)	132	20 (15.2)	NE (NE,	NE)	0.83	0.48,	1.47	0.5046
EORTC QLQ-C30 Functional scale: Physical (MID = 10)	255 24	(9.4)	NE (NE, NE)	132	20 (15.2)	NE (NE,	NE)	0.57	0.31,	1.05	0.0649
EORTC QLQ-C30 Functional scale: Role (MID = 10)	255 41	(16.1)	44.9 (NE, NE)	132	22 (16.7)	NE (NE,	NE)	0.83	0.50,	1.43	0.4940
EORTC QLQ-C30 Functional scale: Cognitive (MID = 10)	255 48	(18.8)	44.9 (NE, NE)	132	32 (24.2)	NE (NE,	NE)	0.69	0.44,	1.10	0.1066
EORTC QLQ-C30 Functional scale: Emotional (MID = 10)	255 44	(17.3)	41.9 (41.9, NE)	132	31 (23.5)	NE (NE,	NE)	0.60	0.38,	0.97	0.0327*
EORTC QLQ-C30 Functional scale: Social (MID = 10)	255 36	(14.1)	41.9 (41.9, NE)	132	22 (16.7)	NE (NE,	NE)	0.69	0.41,	1.20	0.1809
EORTC QLQ-C30 Single item symptom scale: Loss of appetite (MID = 10)	255 23	(9.0)	49.3 (NE, NE)	132	13 (9.8)	NE (NE,	NE)	0.79	0.40,	1.61	0.4923

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Table 3.1.3 PAOLA1 Appendix: Summary of analysis of time to sustained worsening in EORTC QLQ-C30 global QoL/health status, functional, symptom and single item scales (MID=10)

Sensitivity Analysis I (censoring patients with only one worsening post baseline and no subsequent observations)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	0	laparib + b (N=25				Placebo + be					
	of	Tumber (%) f patients ith events	Median tin (95% CI) (months) [Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [c]
EORTC QLQ-C30 Single item symptom scale: Constipation (MID = 10)	255	34 (13.3)	NE (NE,	NE)	132	19 (14.4)	NE (NE, NE)	0.92	0.53,	1.64	0.7606
EORTC QLQ-C30 Single item symptom scale: Diarrhoea (MID = 10)	255	16 (6.3)	NE (NE,	NE)	132	8 (6.1)	NE (NE, NE)	0.88	0.39,	2.18	0.7734
EORTC QLQ-C30 Single item symptom scale: Dyspnoea (MID = 10)	255	32 (12.5)	NE (NE,	NE)	132	24 (18.2)	NE (NE, NE)	0.62	0.36,	1.07	0.0763
EORTC QLQ-C30 Symptom scale: Fatigue (MID = 10)	255	57 (22.4)	NE (NE,	NE)	132	33 (25.0)	NE (NE, NE)	0.77	0.50,	1.20	0.2332
EORTC QLQ-C30 Single item symptom scale: Financial difficulties (MID = 10)	255	17 (6.7)	NE (NE,	NE)	132	12 (9.1)	NE (NE, NE)	0.64	0.31,	1.38	0.2393
EORTC QLQ-C30 Symptom scale: Nausea and vomiting (MID = 10)	255	23 (9.0)	NE (NE,	NE)	132	13 (9.8)	NE (NE, NE)	0.92	0.47,	1.88	0.8210
EORTC QLQ-C30 Symptom scale: Pain (MID = 10)	255	48 (18.8)	19.3 (36.8,	NE)	132	41 (31.1)	NE (NE, NE)	0.47	0.31,	0.72	0.0004*
EORTC QLQ-C30 Single item symptom scale: Insomnia (MID = 10)	255	46 (18.0) 3	38.7 (33.4,	NE)	132	29 (22.0)	NE (NE, NE)	0.69	0.43,	1.12	0.1231

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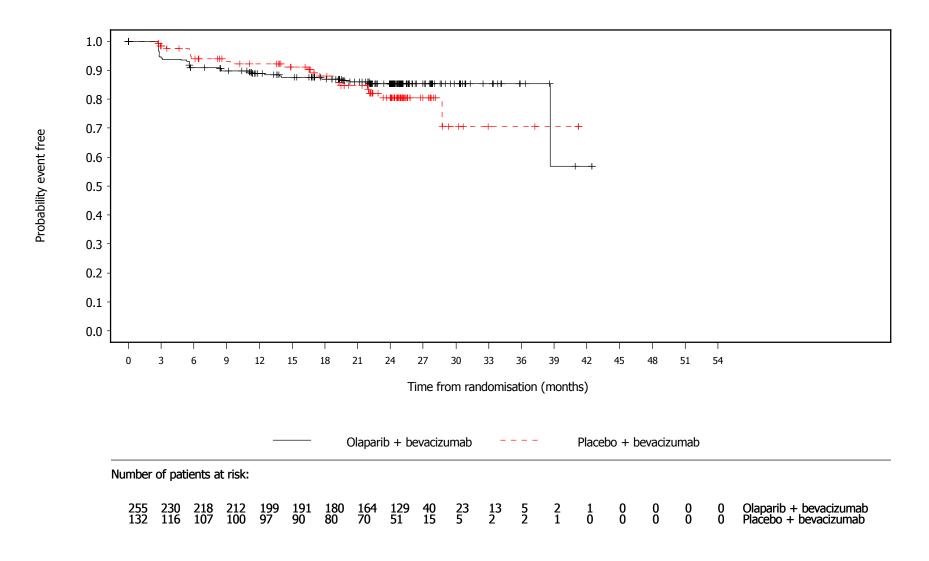
[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

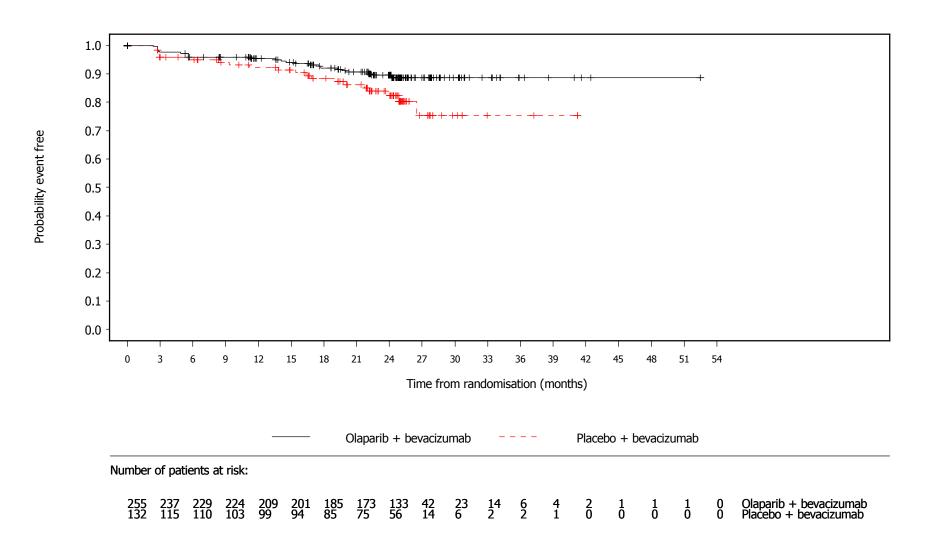
Figure 3.1.4.1 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Global QoL/health status (MID = 10) time to sustained worsening Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4faa 04OCT2022:09:13 kpzx329

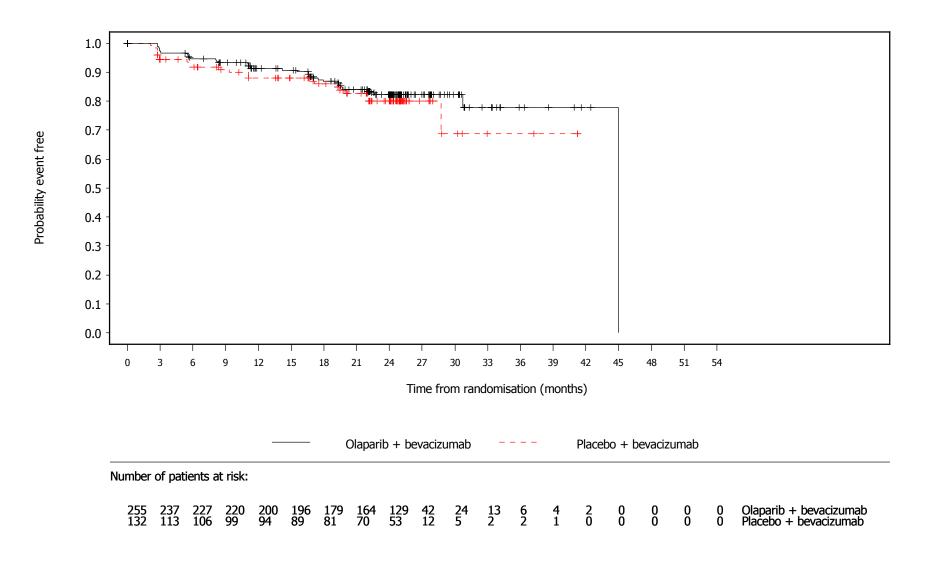
Figure 3.1.4.2 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Physical (MID = 10) time to sustained worsening

Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4fab 04OCT2022:09:13 kpzx329

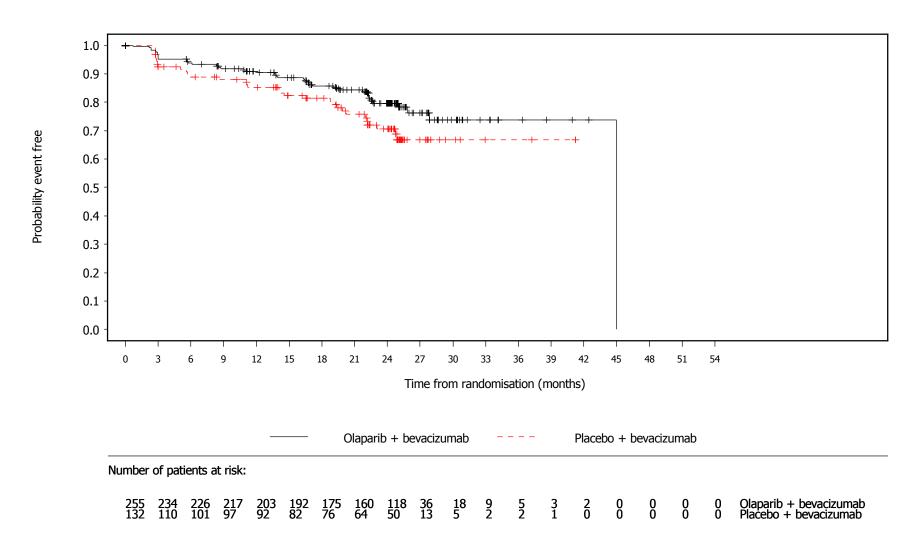
Figure 3.1.4.3 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Role (MID = 10) time to sustained worsening Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4fac 04OCT2022:09:13 kpzx329

Figure 3.1.4.4 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Cognitive (MID = 10) time to sustained worsening

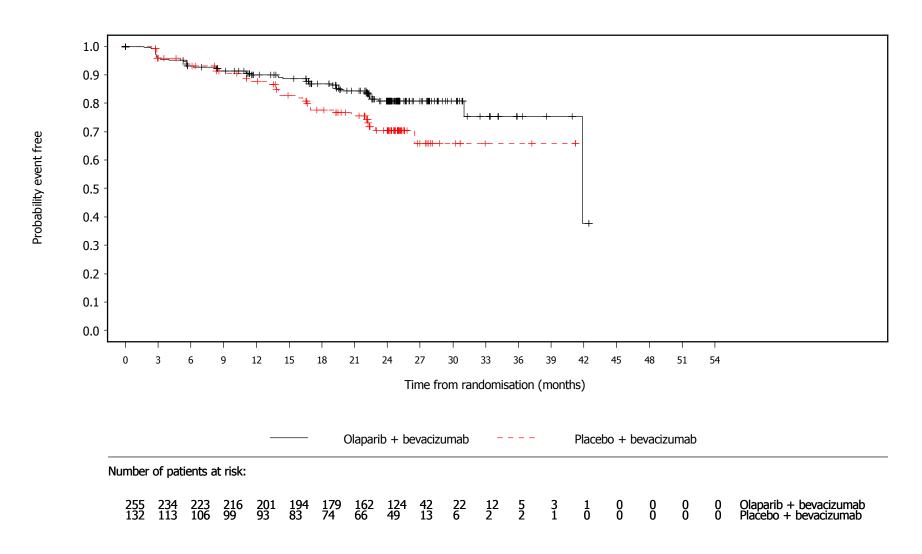
Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4fad 04OCT2022:09:13 kpzx329

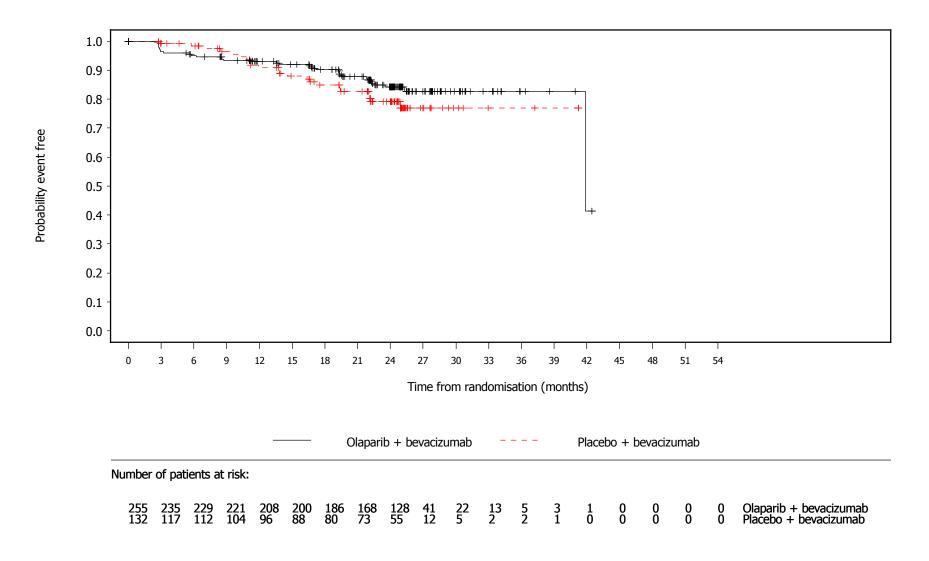
Figure 3.1.4.5 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Emotional (MID = 10) time to sustained worsening

Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4fae 04OCT2022:09:13 kpzx329

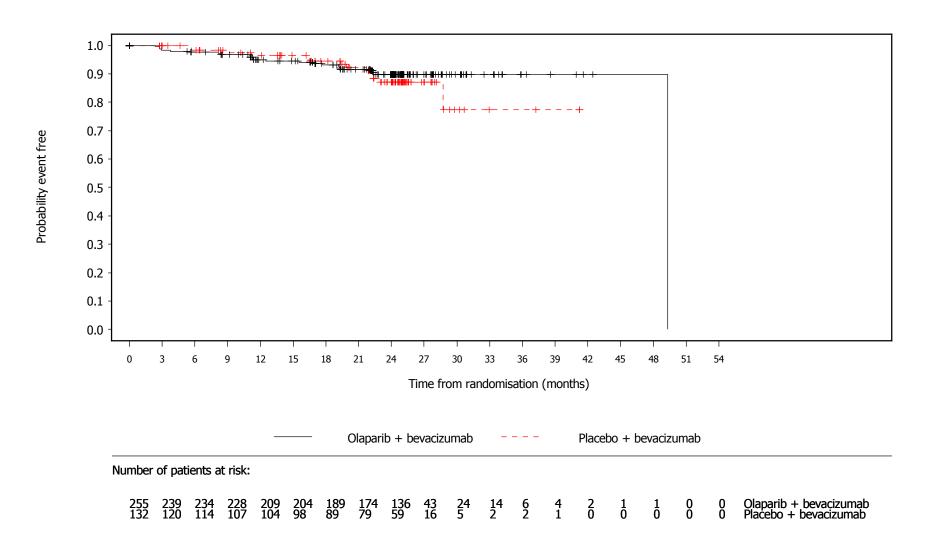
Figure 3.1.4.6 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Functional scale: Social (MID = 10) time to sustained worsening Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4faf 04OCT2022:09:13 kpzx329

Figure 3.1.4.7 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Loss of appetite (MID = 10) time to sustained worsening

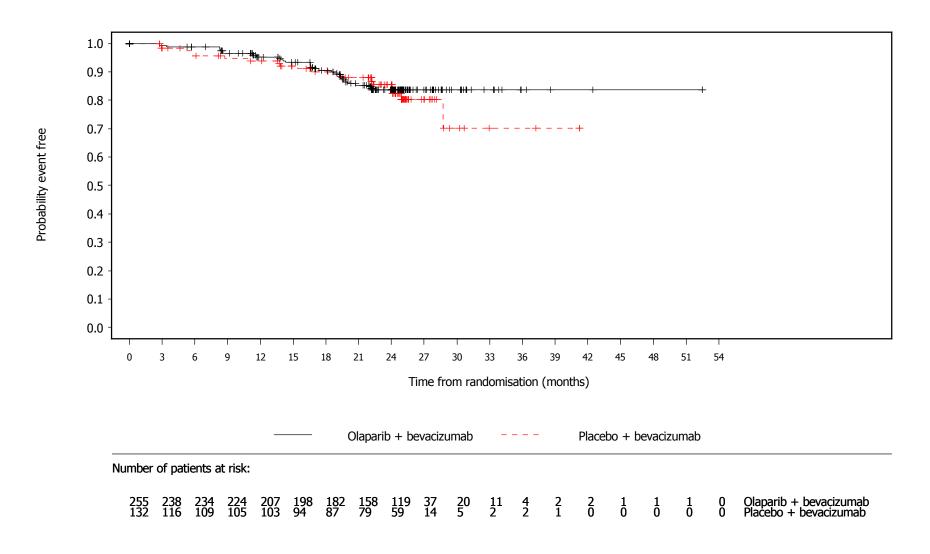
Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4fag 04OCT2022:09:13 kpzx329

Figure 3.1.4.8 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Constipation (MID = 10) time to sustained worsening

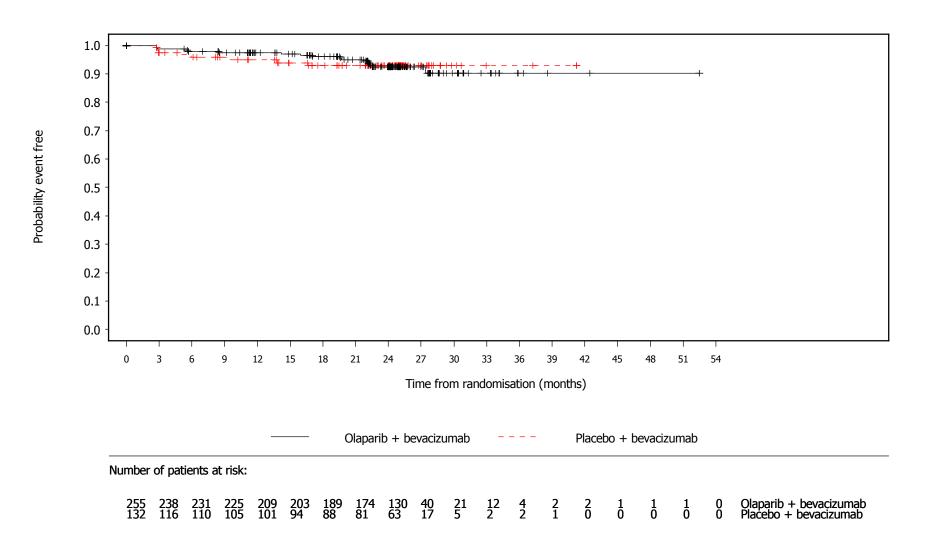
Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4fah 04OCT2022:09:13 kpzx329

Figure 3.1.4.9 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Diarrhoea (MID = 10) time to sustained worsening

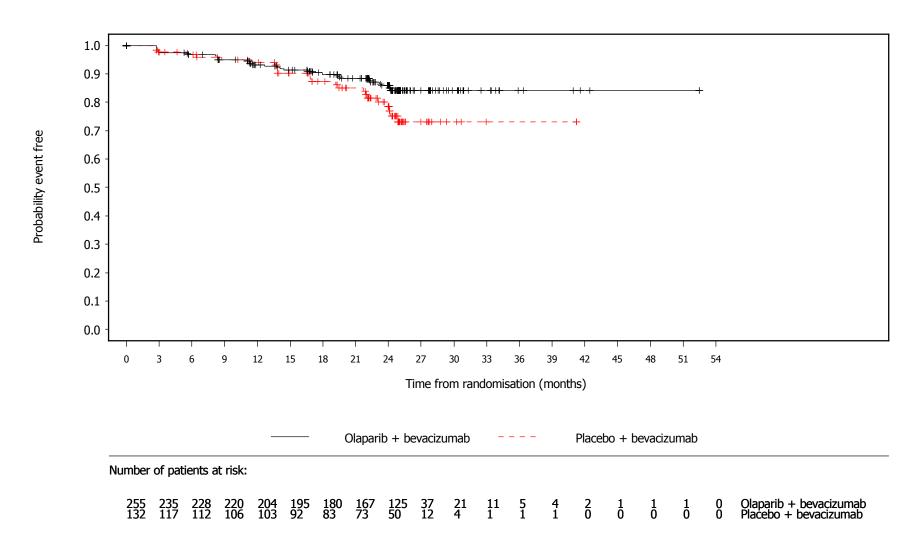
Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4fai 04OCT2022:09:13 kpzx329

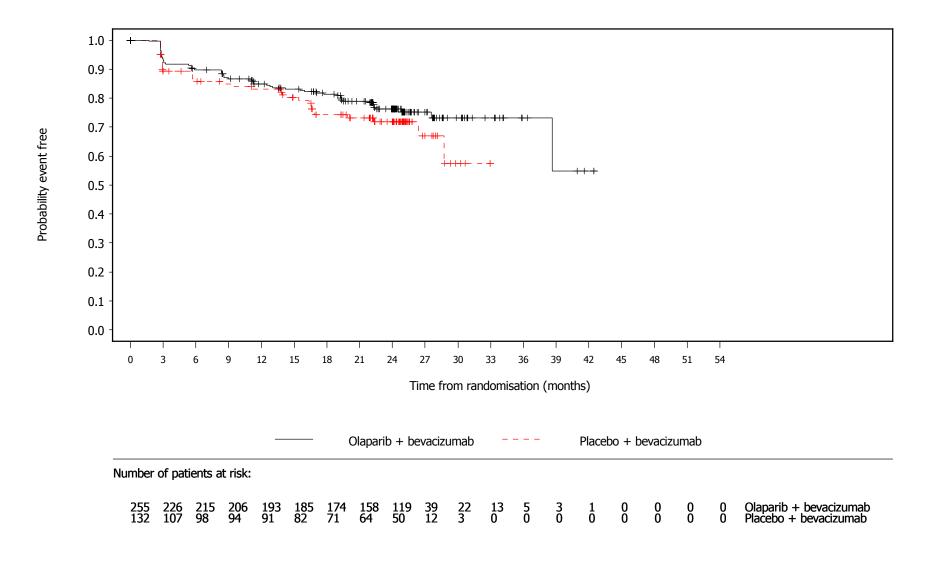
Figure 3.1.4.10 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Dyspnoea (MID = 10) time to sustained worsening

Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4faj 04OCT2022:09:13 kpzx329

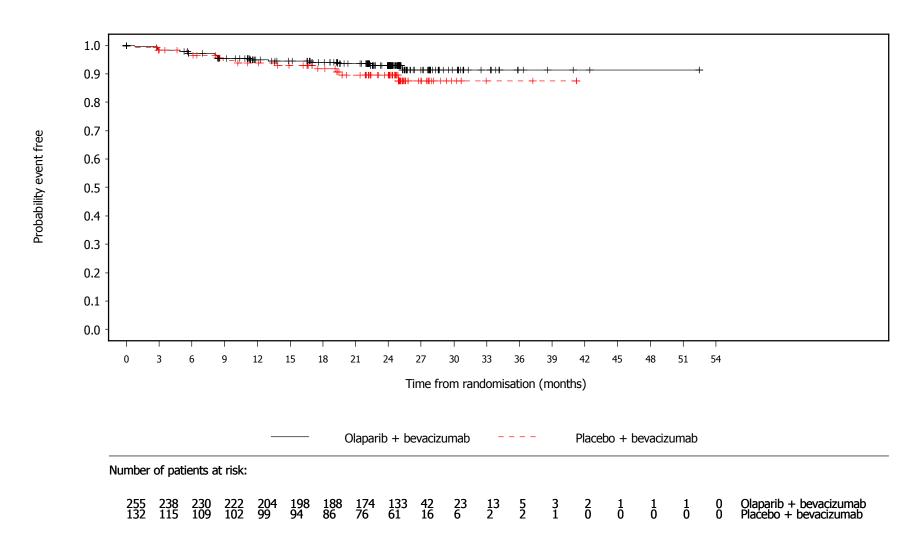
Figure 3.1.4.11 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Fatigue (MID = 10) time to sustained worsening Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4fak 04OCT2022:09:13 kpzx329

Figure 3.1.4.12 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Financial difficulties (MID = 10) time to sustained worsening

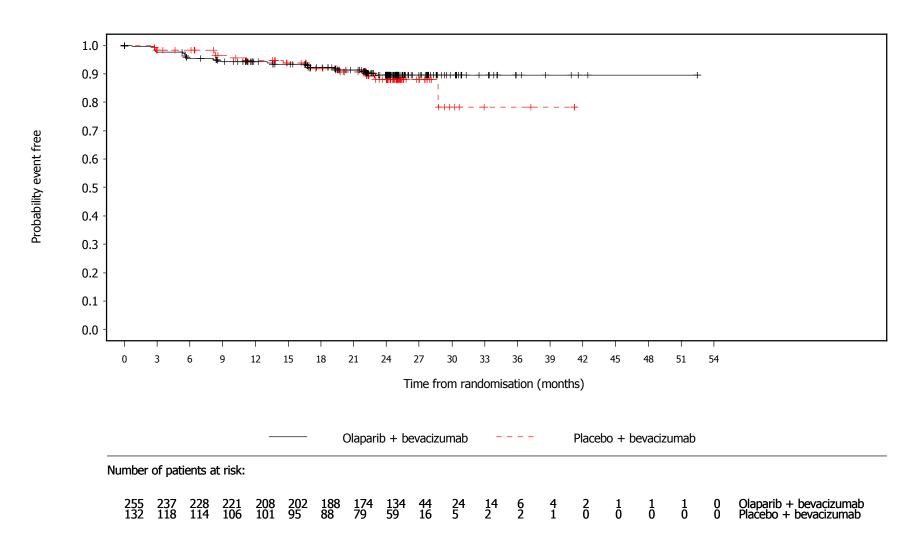
Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4fal 04OCT2022:09:13 kpzx329

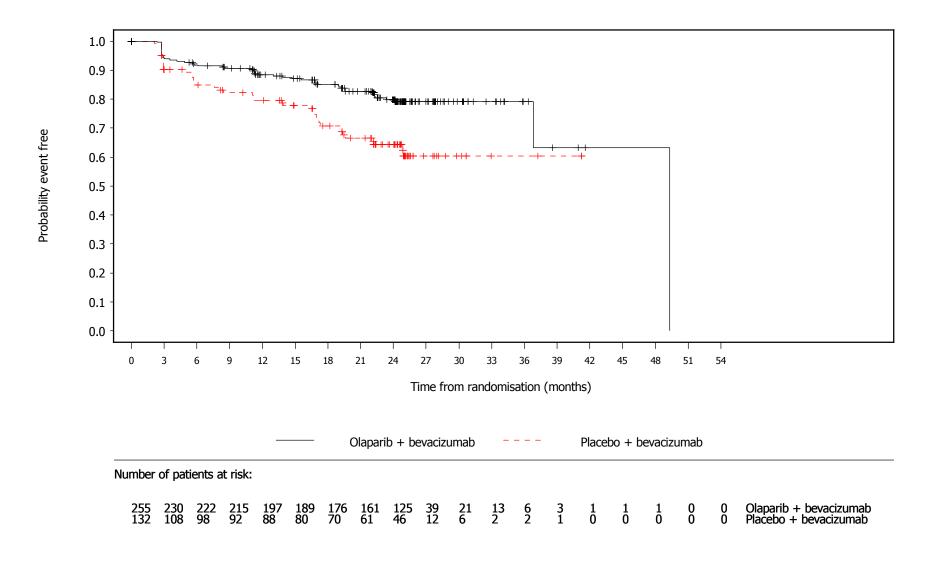
Figure 3.1.4.13 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Nausea and vomiting (MID = 10) time to sustained worsening

Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4fam 04OCT2022:09:13 kpzx329

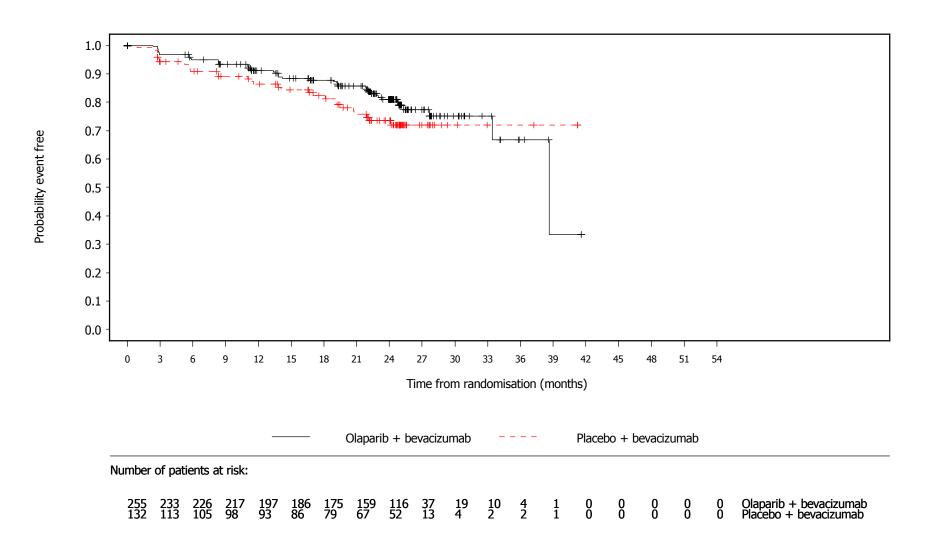
Figure 3.1.4.14 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Symptom scale: Pain (MID = 10) time to sustained worsening Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4fan 040CT2022:09:13 kpzx329

Figure 3.1.4.15 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-C30 Single item symptom scale: Insomnia (MID = 10) time to sustained worsening

Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4fao 04OCT2022:09:13 kpzx329

Table 3.2.1 PAOLA1 Appendix: Summary of analysis of time to sustained worsening in EORTC QLQ-OV28 symptom and single item scales (MID=10)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)		P	Placebo + bevacizumab (N=132)					
	Number of pati n with ev	ients (95% CI)	of	umber (%) patients th events	Median tim (95% CI) (months) [a	ratio		[b]	2-sided p-value [c]
EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 10)	255 77 (3	30.2) 49.3 (27.8, NE)	132	57 (43.2)	25.1 (23.8,29	9.3) 0.56	0.39,	0.80	0.0011*
EORTC QLQ-OV28 Symptom scale/items: Body image (MID = 10)	255 50 (1	.9.6) NE (NE, NE)	132	43 (32.6)	27.8 (25.1,	NE) 0.55	0.36,	0.84	0.0045*
EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (MID = 10)	255 39 (1	.5.3) 42.5 (42.5, NE)	132	32 (24.2)	NE (NE,	NE) 0.48	0.30,	0.79	0.0029*
EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (MID = 10)	255 57 (2	22.4) 52.5 (NE, NE)	132	38 (28.8)	NE (NE,	NE) 0.72	0.47,	1.10	0.1174
EORTC QLQ-OV28 Symptom scale/items: Hormonal (MID = 10)	255 66 (2	25.9) NE (NE, NE)	132	37 (28.0)	NE (NE,	NE) 0.80	0.54,	1.22	0.2901
EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (MID = 10)	255 49 (1	.9.2) NE (NE, NE)	132	18 (13.6)	NE (NE,	NE) 1.36	0.80,	2.42	0.2639
EORTC QLQ-OV28 Symptom scale/items: Other single items (MID = 10)	255 52 (2	20.4) NE (NE, NE)	132	28 (21.2)	NE (NE,	NE) 0.86	0.54,	1.40	0.5342

A sustained worsening is defined as a worsening of response (pts >= MID 10) without any recovery above response threshold compared to baseline value. Patients whose scores have not shown a sustained worsening will be censored at date of their last PRO assessment (also including those with too high/low baseline scores). Patients with no evaluable data at baseline or post baseline will be censored at date of randomisation.

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Figure 3.2.2.1 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

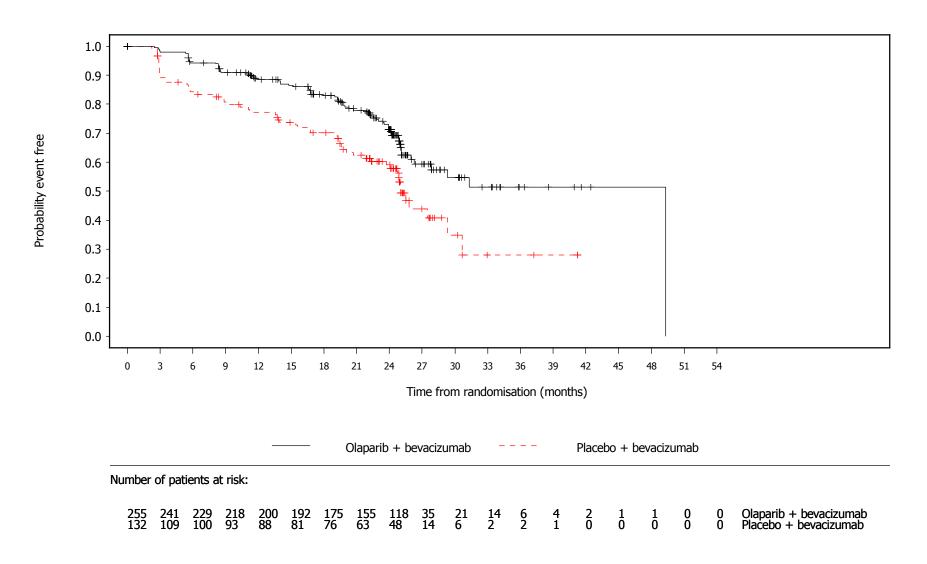


Figure 3.2.2.2 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Body image (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

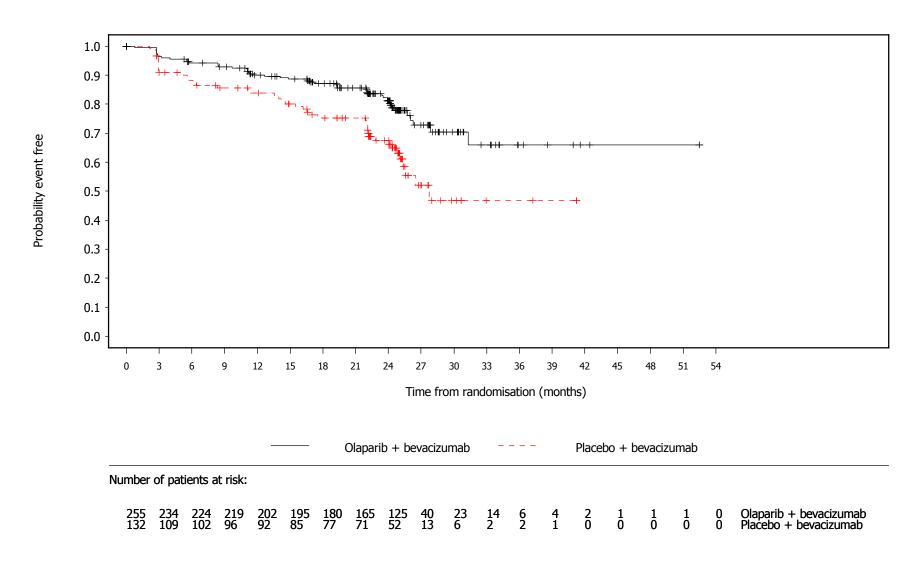


Figure 3.2.2.3 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

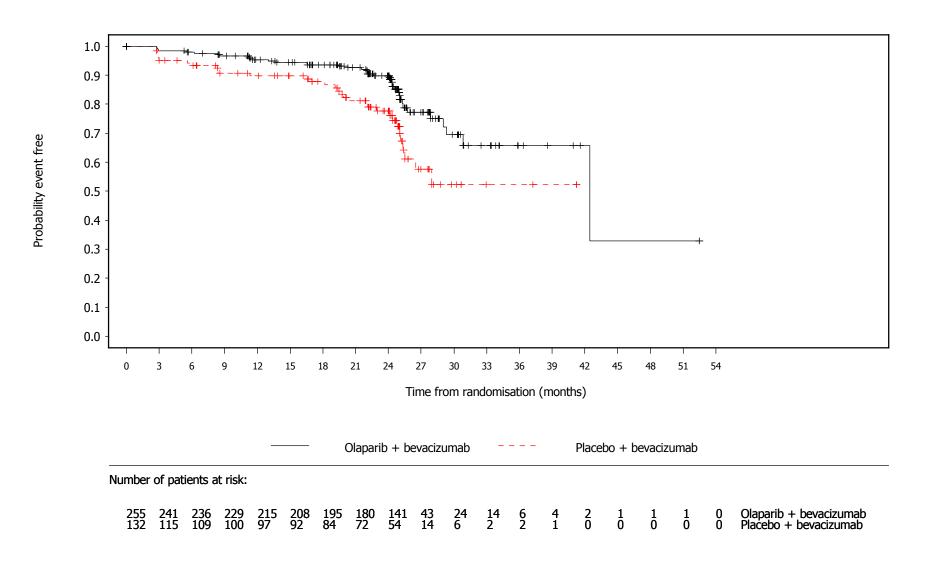


Figure 3.2.2.4 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

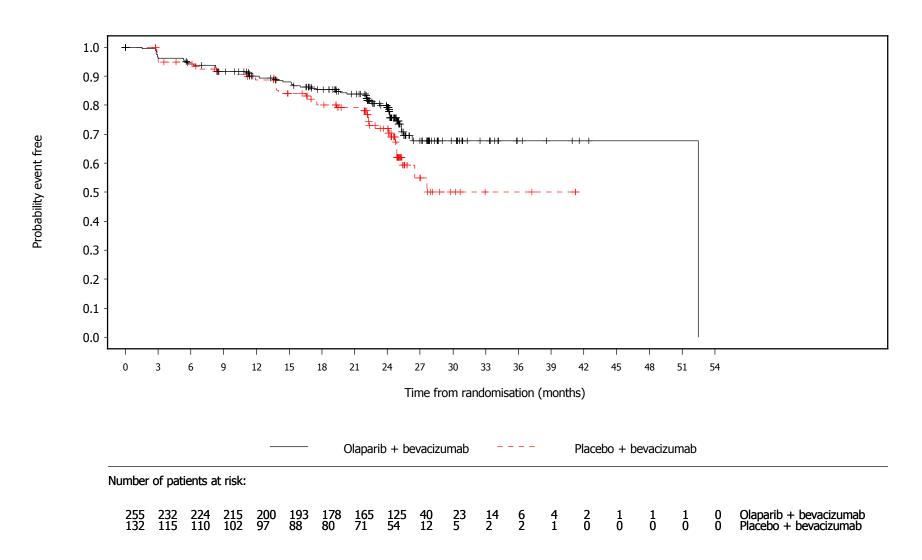


Figure 3.2.2.5 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Hormonal (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

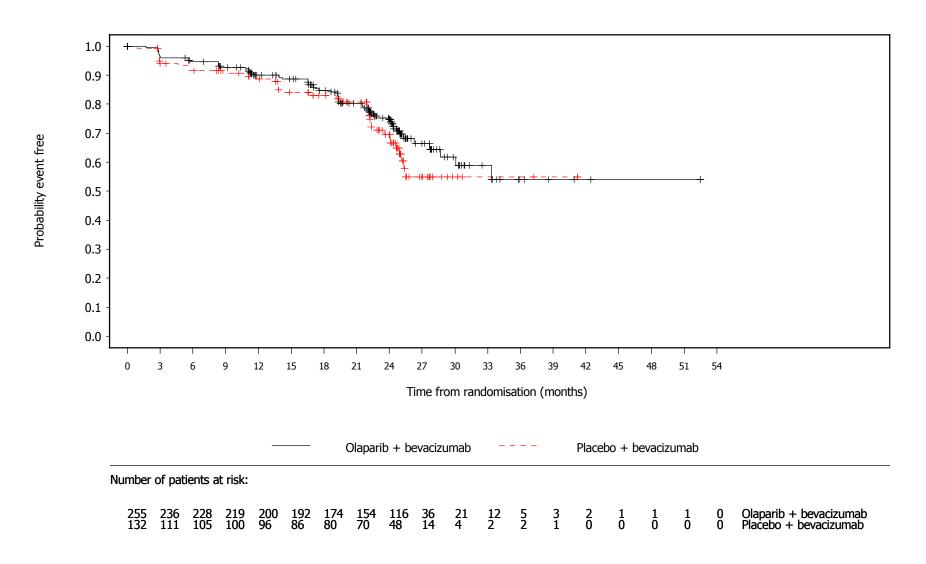


Figure 3.2.2.6 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

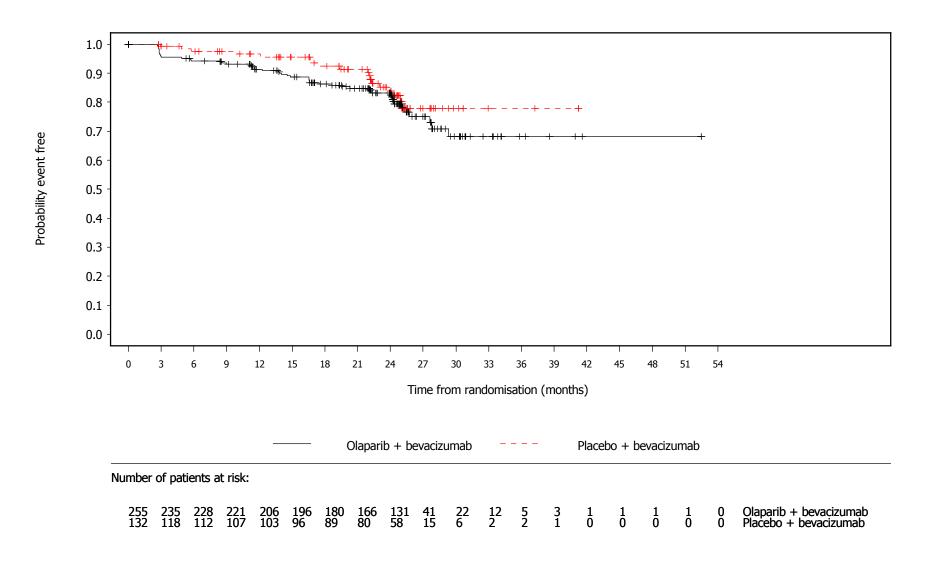


Figure 3.2.2.7 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Other single items (MID = 10) time to sustained worsening

Full Analysis Set, HRD[42] positive, DCO 22Mar2020

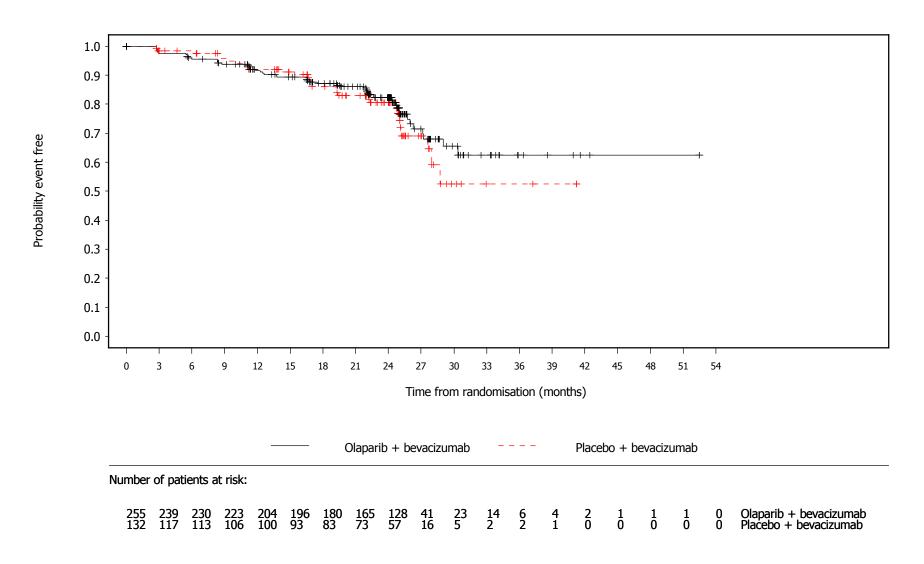


Table 3.2.3 PAOLA1 Appendix: Summary of analysis of time to sustained worsening in EORTC QLQ-OV28 symptom and single item scales (MID=10)

Sensitivity Analysis I (censoring patients with only one worsening post baseline and no subsequent observations)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)			Placebo + bevacizumab (N=132)								
	of]	nber (%) patients h events	Median tin (95% CI) (months) [(Number (%) of patients with events	Median ti (95% CI) (months) [)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [c]
EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 10)	255 50	0 (19.6)	49.3 (NE)	,	132	38 (28.8)	NE (NE,	NE)	0.54	0.35,	0.84	0.0046*
EORTC QLQ-OV28 Symptom scale/items: Body image (MID = 10)	255 29	9 (11.4)	NE (NE,	NE)	132	26 (19.7)	NE (NE,	NE)	0.57	0.33,	0.98	0.0379*
EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (MID = 10)	255 1	5 (5.9)	NE (NE,	NE)	132	19 (14.4)	NE (NE,	NE)	0.37	0.18,	0.72	0.0029*
EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (MID = 10)	255 3!	5 (13.7)	NE (NE,	NE)	132	21 (15.9)	NE (NE,	NE)	0.83	0.48,	1.46	0.4988
EORTC QLQ-OV28 Symptom scale/items: Hormonal (MID = 10)	255 42	2 (16.5)	NE (NE,	NE)	132	25 (18.9)	NE (NE,	NE)	0.74	0.45,	1.23	0.2357
EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (MID = 10)	255 30	0 (11.8)	NE (NE,	NE)	132	6 (4.5)	NE (NE,	NE)	2.56	1.14,	6.84	0.0303*
EORTC QLQ-OV28 Symptom scale/items: Other single items (MID = 10)	255 30	0 (11.8)	NE (NE,	NE)	132	15 (11.4)	NE (NE,	NE)	0.96	0.52,	1.84	0.8954

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

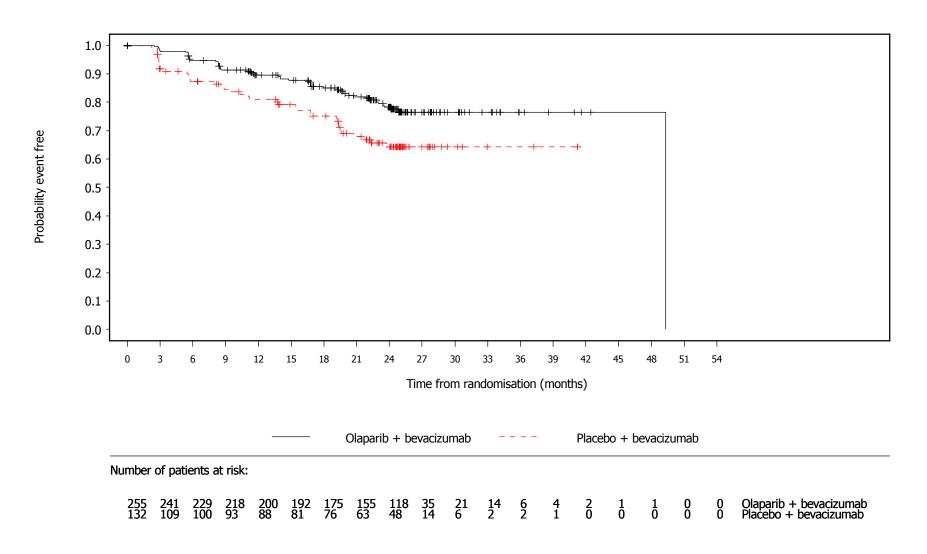
[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Figure 3.2.4.1 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Abdominal/GI (MID = 10) time to sustained worsening

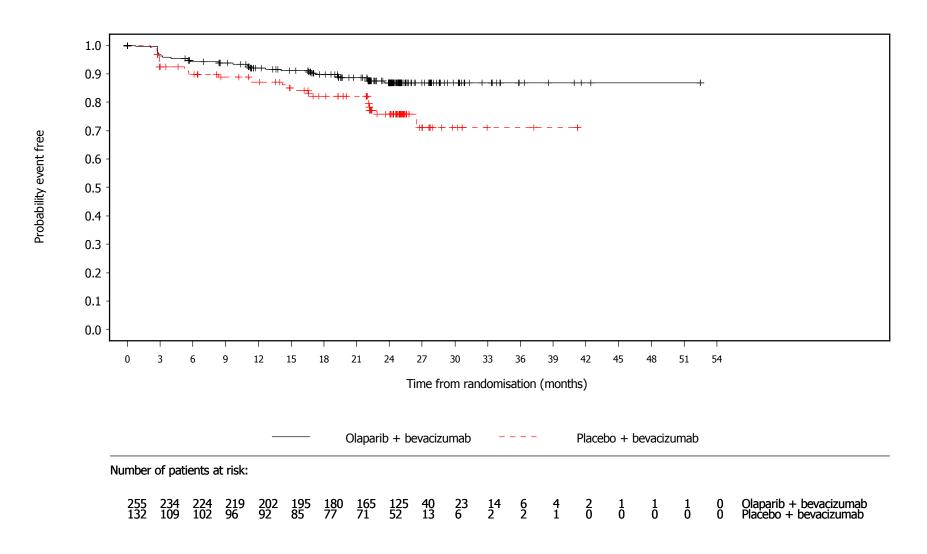
Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4gaa 04OCT2022:09:13 kpzx329

Figure 3.2.4.2 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Body image (MID = 10) time to sustained worsening

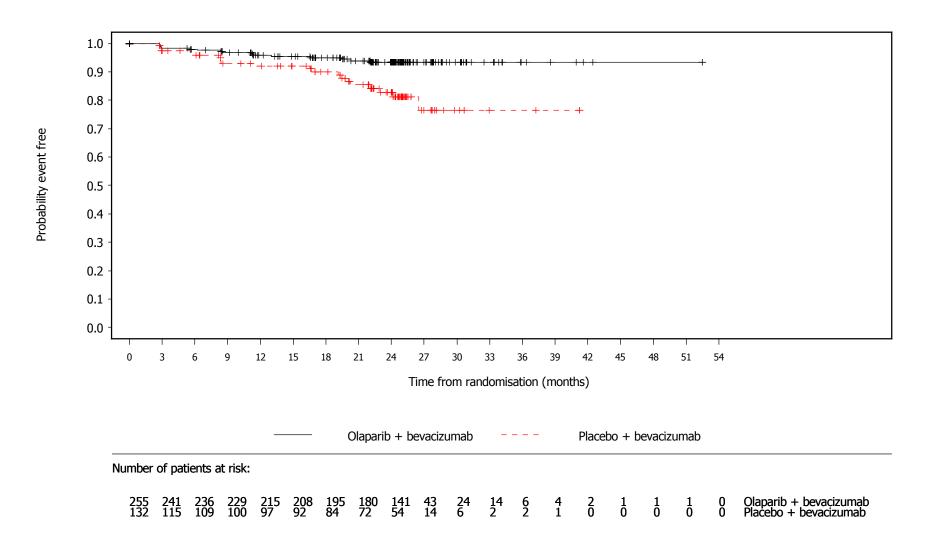
Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4gab 04OCT2022:09:13 kpzx329

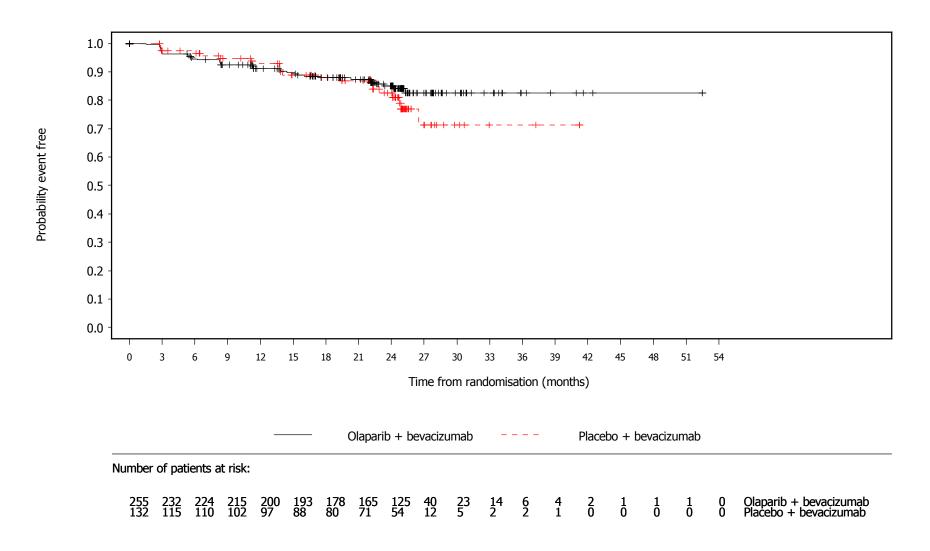
Figure 3.2.4.3 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Chemotherapy side effects (MID = 10) time to sustained worsening

Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4gac 04OCT2022:09:13 kpzx329

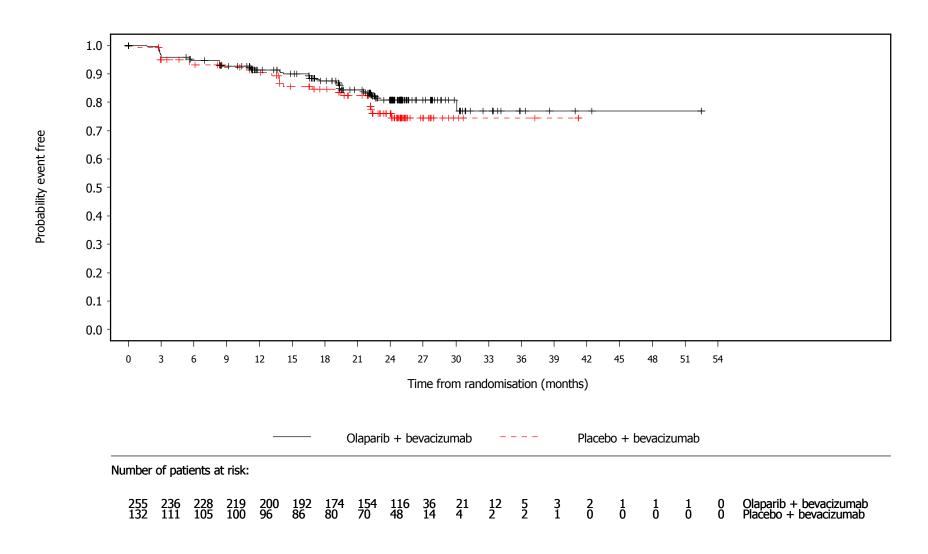
Figure 3.2.4.4 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Attitude to disease/treatment (MID = 10)
time to sustained worsening
Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4gad 04OCT2022:09:13 kpzx329

Figure 3.2.4.5 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Hormonal (MID = 10) time to sustained worsening

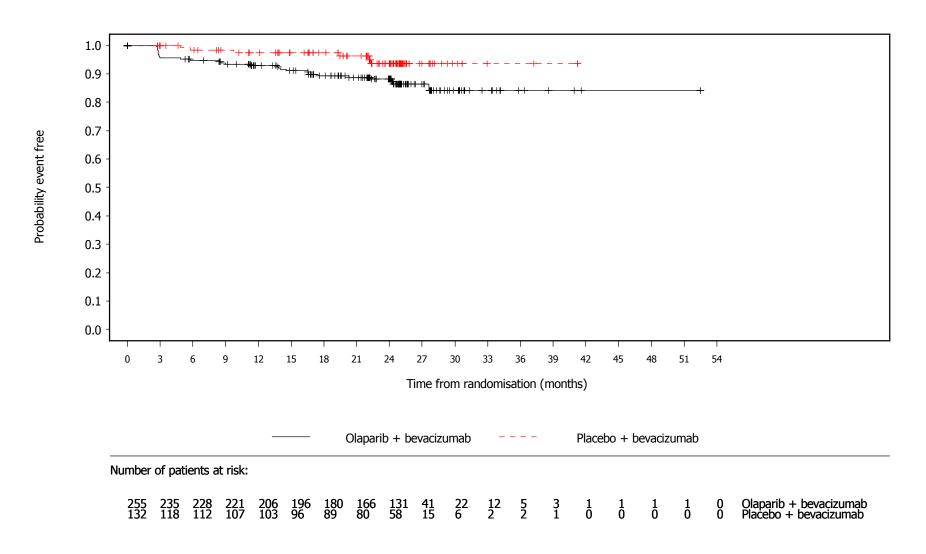
Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4gae 04OCT2022:09:13 kpzx329

Figure 3.2.4.6 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Peripheral neuropathy (MID = 10) time to sustained worsening

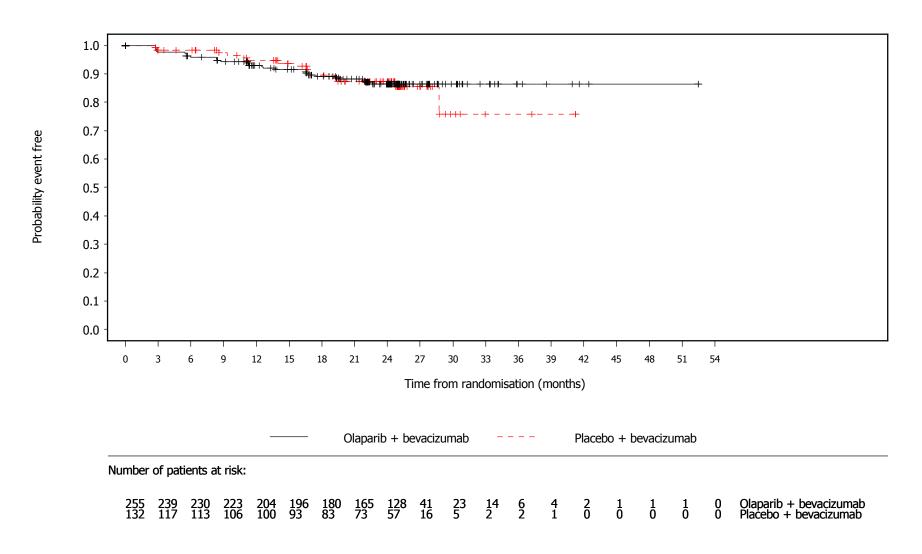
Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4gaf 04OCT2022:09:13 kpzx329

Figure 3.2.4.7 PAOLA1 Appendix: Kaplan-Meier plot of EORTC QLQ-OV28 Symptom scale/items: Other single items (MID = 10) time to sustained worsening

Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4gag 04OCT2022:09:13 kpzx329

Table 3.3.1 PAOLA1 Appendix: Summary of analysis of time to sustained worsening in EQ-5D VAS (MID = 15)

Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + b (N=2!		Placebo + be				
	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
EQ-5D-5L Visual analogue scale (MID = 15)	255 39 (15.3)	44.9 (NE, NE)	132 31 (23.5)	NE (NE, NE)	0.61	0.38, 0.995	0.0432*

A sustained worsening is defined as a worsening of response (pts >= MID 10) without any recovery above response threshold compared to baseline value. Patients whose scores have not shown a sustained worsening will be censored at date of their last PRO assessment (also including those with too high/low baseline scores). Patients with no evaluable data at baseline or post baseline will be censored at date of randomisation.

 [[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.
 [b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.
 [c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.
 Hazard ratio <1 favours olaparib. * p<0.05.</p>

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Figure 3.3.2.1 PAOLA1 Appendix: Kaplan-Meier plot of EQ-5D-5L Visual analogue scale (MID = 15) time to sustained worsening Full Analysis Set, HRD[42] positive, DCO 22Mar2020

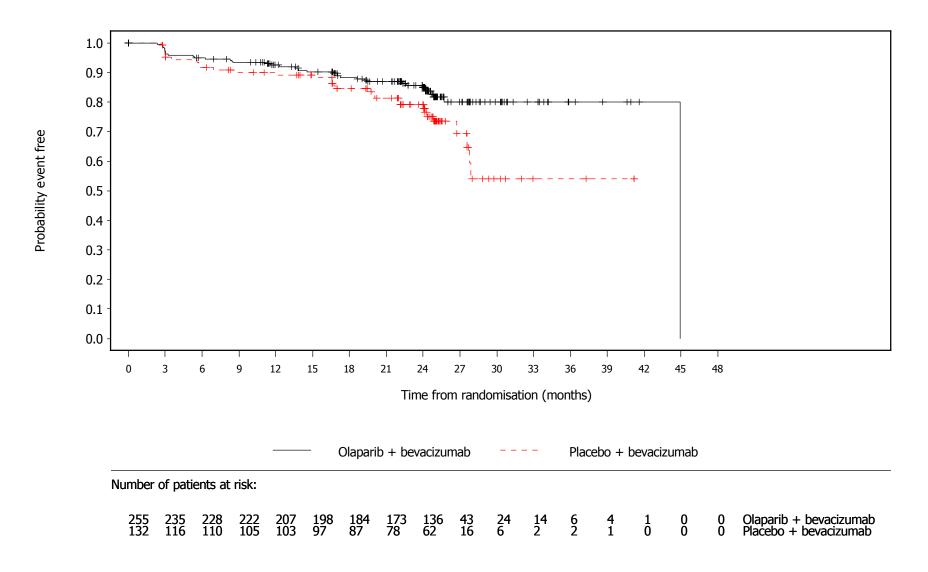


Table 3.3.3 PAOLA1 Appendix: Summary of analysis of time to sustained worsening in EQ-5D VAS (MID = 15)

Sensitivity Analysis I (censoring patients with only one worsening post baseline and no subsequent observations)

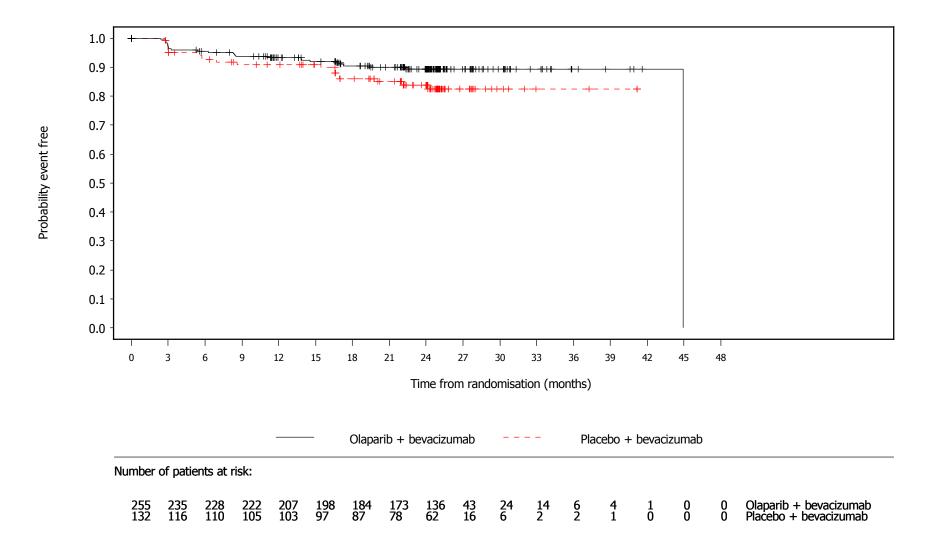
Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Olaparib + bevacizumab (N=255)		Placebo + be				
	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
EQ-5D-5L Visual analogue scale (MID = 15)	255 25 (9.8)	44.9 (NE, NE)	132 19 (14.4)	NE (NE, NE)	0.61	0.33, 1.13	0.1071

 [[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.
 [b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.
 [c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Figure 3.3.4.1 PAOLA1 Appendix: Kaplan-Meier plot of EQ-5D-5L Visual analogue scale (MID = 15) time to sustained worsening Sensitivity analysis I, Full Analysis Set, HRD[42] positive, DCO 22Mar2020



Sensitivity Analysis - censoring patients with only one worsening post baseline and no subsequent observations. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainpr4.sas ettemainpr4haa 04OCT2022:09:13 kpzx329

Table 3.4 PAOLA1 Appendix: Patients unblinded prior to last PRO observation Full Analysis Set, HRD[42] positive, DCO 22MAR2020

	Number (%) of patients						
	Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=132)	Total (N=387)				
Unblinded prior to last PRO observation	7 (2.7)	11 (8.3)	18 (4.7)				

Table 1.0.1 PAOLA1: Summary of observation period (months) for efficacy endpoints Full Analysis Set, HRD[42] positive, DCO 22MAR2022

		Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=132)
D		255	132
Progression-free survival		255	
	Median	38.87	16.80
	Min	0.0	0.0
	Max	76.5	71.1
Second Progression-free survival	n	255	132
	Median	53.59	35.30
	Min	0.0	0.0
	Max	76.5	72.0
Time to First Subsequent Cancer Therapy or Death	n	255	132
	Median	40.41	18.50
	Min	1.4	0.3
	Max	77.8	76.9
Time to Second Subsequent Cancer Therapy or Death	n	255	132
	Median	55.89	35.30
	Min	1.4	0.3
	Max	77.8	76.9
Overall Survival	n	255	132
	Median	58.74	55.38
	Min	1.4	0.3
	Max	77.8	76.9

Table 1.0.2 PAOLA1: Summary of observation period (months) for recurrence free survival Full Analysis Set, HRD[42] positive, DCO 22MAR2022

		Olaparib + bevacizumab (N=206)	Placebo + bevacizumab (N=106)
Time to recurrence or death	n	206	106
	Median	50.78	18.30
	Min	0.0	0.0
	Max	76.5	71.1

Table 1.1.1.1 PAOLA1: Summary of Progression-free Survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

	Olaparib + bevacizumab (N=255)		Placebo + k					
	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [c]
Progression-free survival	255 136 (53.3)	46.9 (36.4,65.7)	132 104 (78.8)	17.6 (15.8,20.3)	0.42	0.32,	0.55	<0.0001*

 [[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.
 [b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.
 [c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.
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Table 1.1.1.2 PAOLA1: Summary of Second Progression-free Survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

	Olaparib + bevacizumab (N=255)		Placebo + (N=				
	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	, ,	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
Second Progression-free survival	255 112 (43.9) 7	75.2 (57.1, NE)	132 90 (68.2)	37.7 (31.2,46.7)	0.56	0.42, 0.75	<0.0001*

 [[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.
 [b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.
 [c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Table 1.1.1.3 PAOLA1: Summary of Time to First Subsequent Cancer Therapy or Death Full Analysis Set, HRD[42] positive, DCO 22Mar2022

	Olaparib + b (N=25			pevacizumab 132)			
	Number (%) of patients n with events	of patients (95% CI) of patients (95% CI) ratio		95% CI [b]	2-sided p-value [c]		
Time to First Subsequent Cancer Therapy or Death	255 143 (56.1)	42.2 (35.7,60.3)	132 108 (81.8)	18.8 (16.1,20.5)	0.43	0.33, 0.56	<0.0001*

 [[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.
 [b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.
 [c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Table 1.1.1.4 PAOLA1: Summary of Time to Second Subsequent Cancer Therapy or Death Full Analysis Set, HRD[42] positive, DCO 22Mar2022

	Olaparib + b		Placebo + k (N=1				
	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
Time to Second Subsequent Cancer Therapy or Death	255 115 (45.1)	75.2 (56.2, NE)	132 92 (69.7)	35.4 (29.2,45.2)	0.54	0.40, 0.71	<0.0001*

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.
[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.
[c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.</pre>

Table 1.1.1.5 PAOLA1: Summary of Overall Survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

	Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=132)		
	Number (%) Median time of patients (95% CI) n with events (months) [a]	Number (%) Median time of patients (95% CI) n with events (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [c]
Overall Survival	255 93 (36.5) 75.2 (73.3, NE)	132 69 (52.3) 57.3 (51.6, NE)	0.68 0.50, 0.94	0.0169*

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.
[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.
[c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.</pre>

Figure 1.1.2.1 PAOLA1: Kaplan-Meier plot of Progression-free Survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

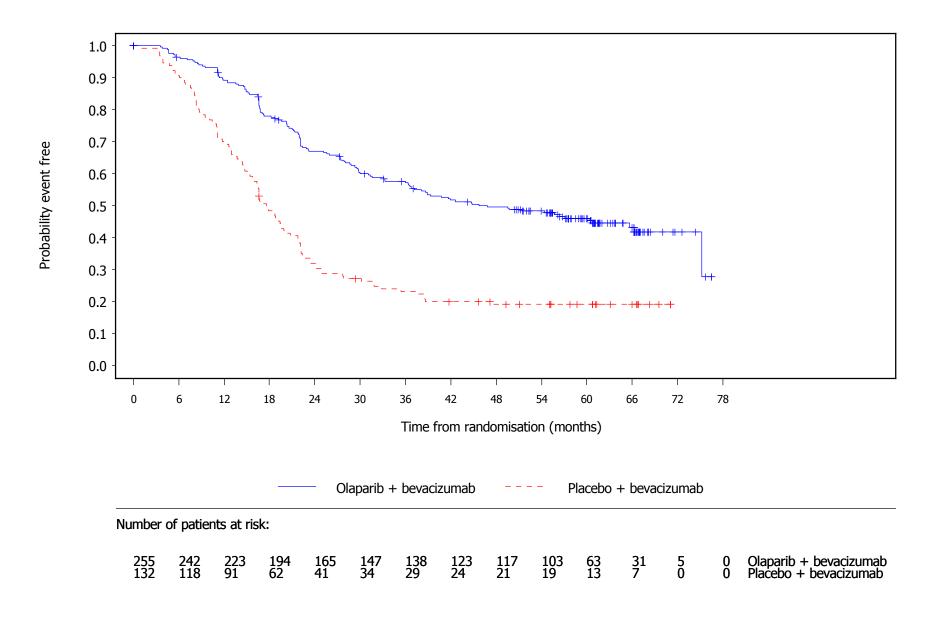


Figure 1.1.2.2 PAOLA1: Kaplan-Meier plot of Second Progression-free Survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

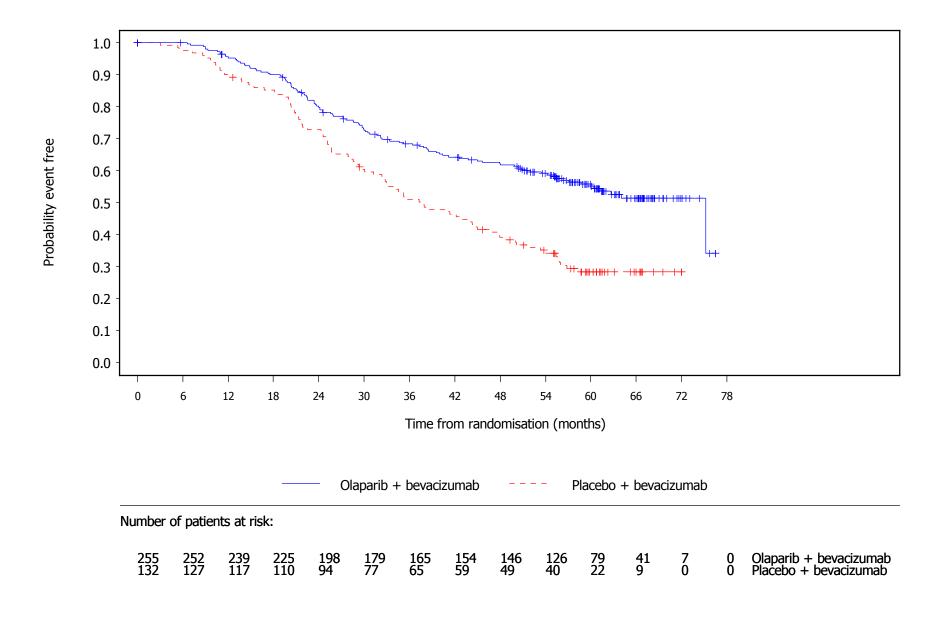


Figure 1.1.2.3 PAOLA1: Kaplan-Meier plot of Time to First Subsequent Cancer Therapy or Death Full Analysis Set, HRD[42] positive, DCO 22Mar2022

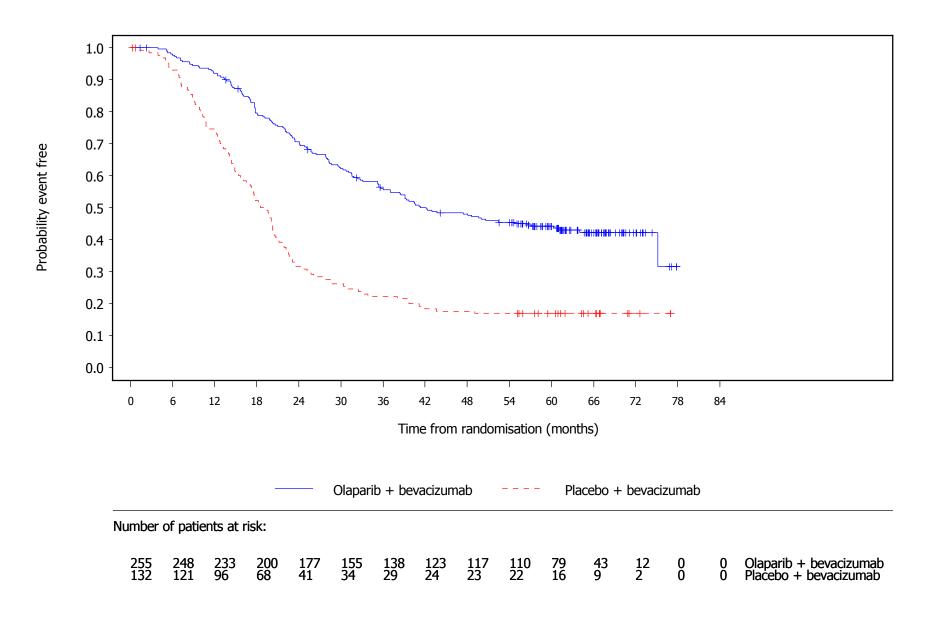


Figure 1.1.2.4 PAOLA1: Kaplan-Meier plot of Time to Second Subsequent Cancer Therapy or Death Full Analysis Set, HRD[42] positive, DCO 22Mar2022

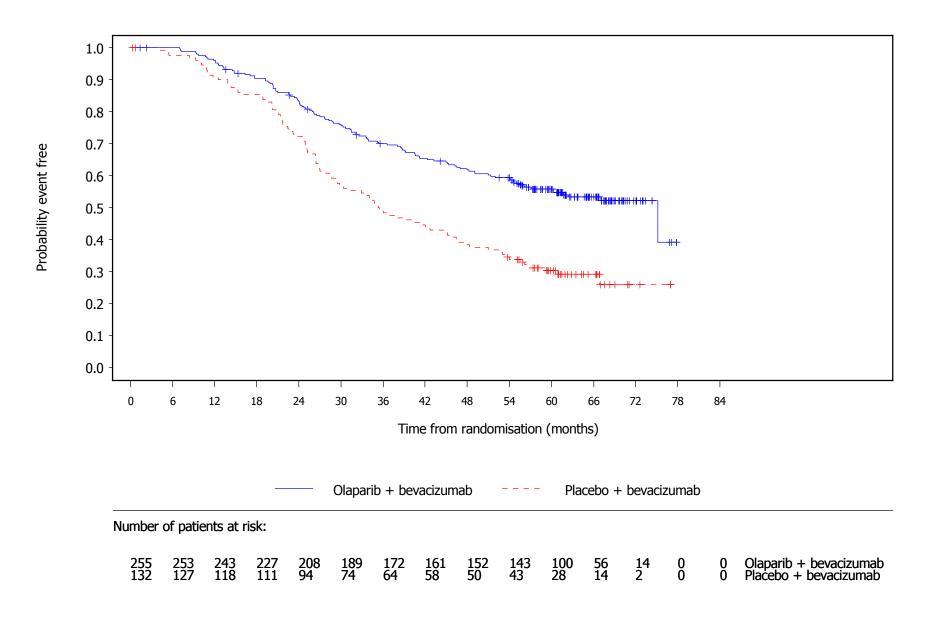


Figure 1.1.2.5 PAOLA1: Kaplan-Meier plot of Overall Survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

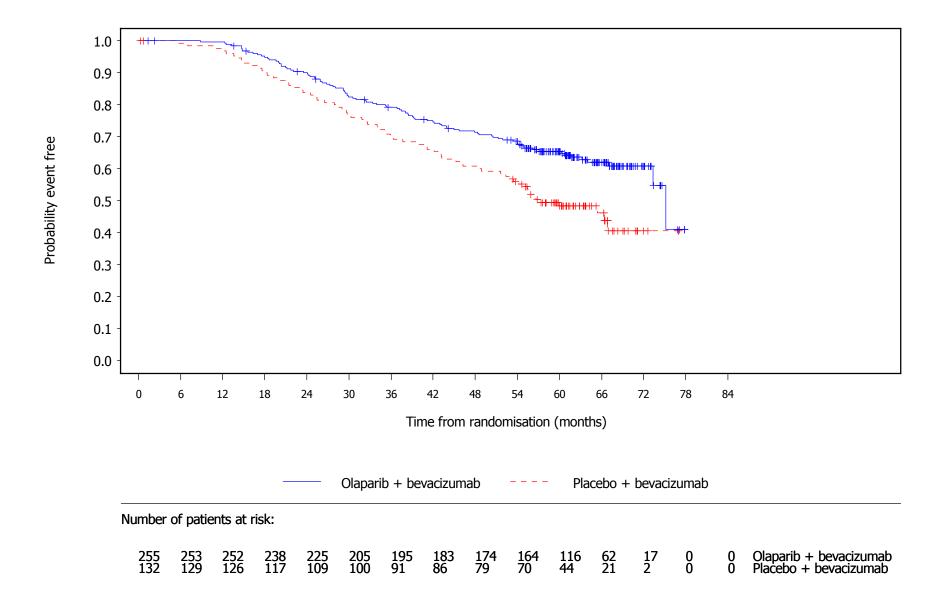


Table 1.1.3.1 PAOLA1: Summary of subgroup analysis of Progression-free survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

	:	Olaparib + N=2				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)								
NED [PDS]	92	32 (34.8)	75.2 (65.7, NE)	48	34 (70.8)	21.7 (16.6,24.9)	0.27	0.17,	0.45	<0.0001*
NED/CR [IDS]	74	46 (62.2)	31.3 (22.1,44.8)	38	31 (81.6)	17.2 (13.7,22.1)	0.52	0.33,	0.82	0.0059*
NED/CR [Chemo]	40	19 (47.5)	57.1 (25.9, NE)	20	19 (95.0)	14.5 (10.9,22.3)	0.25	0.13,	0.48	<0.0001*
PR	49	39 (79.6)	22.0 (16.6,27.5)	26	20 (76.9)	16.0 (8.1,19.4)	0.68	0.40,	1.18	0.1675
Interaction p-value										0.0243*
Screening laboratory tBR	CA statu	ıs (IVRS)								
tBRCAm	150	75 (50.0)	56.4 (41.9, NE)	65	46 (70.8)	20.3 (15.4,24.0)	0.45	0.31,	0.65	<0.0001*
non-tBRCAm	105	61 (58.1)	30.0 (22.0,60.3)	67	58 (86.6)	16.5 (12.9,18.6)	0.40	0.28,	0.57	<0.0001*
Interaction p-value										0.6313
First line treatment out	come (eC	RF)								
NED [PDS]	89	29 (32.6)	75.2 (66.3, NE)	47	33 (70.2)	21.9 (16.6,27.7)	0.26	0.16,	0.43	<0.0001*
NED/CR [IDS]	74	47 (63.5)	33.3 (22.1,44.8)	32	27 (84.4)	16.6 (12.9,22.1)	0.48	0.30,	0.78	0.0037*
NED/CR [Chemo]	39	21 (53.8)	38.9 (22.9, NE)	18	17 (94.4)	14.5 (10.9,21.9)	0.25	0.13,	0.49	<0.0001*
PR	50	37 (74.0)	22.1 (16.8,28.6)	34	26 (76.5)	16.7 (8.3,22.0)	0.67	0.41,	1.12	0.1282
Interaction p-value										0.0239*
Screening laboratory tBR	CA statu	s (eCRF)								
tBRCAm	147	74 (50.3)	56.4 (41.9, NE)	67	48 (71.6)	19.9 (15.4,23.5)	0.44	0.31,	0.63	<0.0001*
non-tBRCAm	108	62 (57.4)	33.3 (22.0,60.3)	65	56 (86.2)	16.5 (12.9,19.2)	0.40	0.28,	0.58	<0.0001*
Interaction p-value										0.7190
Age group										
<65 years	185	101 (54.6)	39.3 (31.1, NE)	98	77 (78.6)	18.7 (15.4,22.1)	0.45	0.33,	0.60	<0.0001*
>=65 years	70	35 (50.0)	60.3 (38.9, NE)	34	27 (79.4)	16.7 (10.9,19.5)	0.32	0.20,	0.54	<0.0001*
Interaction p-value										0.2873
FIGO Stage (Disease stat	e)									
III	182	89 (48.9)	60.7 (39.3, NE)	90	67 (74.4)	21.0 (17.6,23.4)	0.43	0.31,	0.59	<0.0001*

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

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Table 1.1.3.1 PAOLA1: Summary of subgroup analysis of Progression-free survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olapa	rib + (N=2	bevacizumab 255)		Placebo + (N=	bevac: 132)	izumab				
Subgroup	n	Number of pat with e	ients	Median time (95% CI) (months) [a]	n	Number (%) of patients with events	;	edian time (95% CI) onths) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
IV	73	47 (64.4)	28.1 (21.9,45.7)	42	37 (88.1)	12.8	(9.5,14.7)	0.35	0.22,	0.54	<0.0001*
Interaction p-value												0.4325
Region												
Europe	245	129 (52.7)	49.7 (37.2,66.3)	126	98 (77.8)	17.0	(15.1,20.3)	0.41	0.31,	0.53	<0.0001*
Japan	10	7 (70.0)	23.3 (5.6, NE)	6	6 (100)	20.7	(11.1, NE)	0.59	0.19,	1.83	0.3445
Interaction p-value												0.5175
ECOG performance status at	Basel	ine										
(0) Normal activity	190	102 (53.7)	42.6 (31.3,66.3)	100	78 (78.0)	18.7	(15.4,21.9)	0.45	0.34,	0.61	<0.0001*
(1) Restricted activity	61	31 (50.8)	60.3 (36.0, NE)	31	26 (83.9)	16.8	(10.1,22.2)	0.28	0.17,	0.48	<0.0001*
Interaction p-value												0.1319
Baseline CA-125 value												
<=ULN	228	111 (48.7)	60.7 (41.9, NE)	118	93 (78.8)	17.7	(16.0,20.3)	0.36	0.27,	0.47	<0.0001*
>ULN	27	25 (92.6)	16.6 (11.9,23.1)	14	11 (78.6)	11.4	(5.1,37.7)	1.12	0.56,	2.37	0.7512
Interaction p-value												0.0023*
Histological grade												
High grade	255	136 (53.3)	46.9 (36.4,65.7)	132	104 (78.8)	17.6	(15.8,20.3)	0.41	0.32,	0.53	<0.0001*
Interaction p-value												NC
Cytoreductive surgery outc	ome											
No residue	166	78 (47.0)	65.7 (45.7, NE)	80	61 (76.3)	19.5	(16.6,22.3)	0.38	0.27,	0.54	<0.0001*
Residue	79	48 (60.8)	30.0 (22.1,41.9)	44	36 (81.8)	14.5	(10.9,20.3)	0.45	0.29,	0.70	0.0005*
Interaction p-value												0.5466
Timing of cytoreductive su	rgery											
Upfront	146	62 (42.5)	75.2 (60.3, NE)	79	59 (74.7)	18.8	(16.4,22.2)	0.33	0.23,	0.47	<0.0001*
Interval	99	64 (64.6)	31.3 (22.1,40.8)	45	38 (84.4)	16.6	(12.9,20.3)	0.50	0.34,	0.75	0.0011*
Interaction p-value												0.1263

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

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Table 1.1.3.1 PAOLA1: Summary of subgroup analysis of Progression-free survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

		-	bevacizumab 255)				oevacizumab 132)				
Subgroup		Number (%) of patients with events				Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Myriad tumour BRCA mutati	on stati	ıs									
tBRCAm	158		56.4 (38.9,	NE)	77	58 (75.3)	18.8 (14.7,22.1)	0.39	0.28,	0.55	<0.0001*
Non-tBRCAm	97	58 (59.8)	30.0 (21.9,60).3)	55	46 (83.6)	16.6 (12.9,19.5)	0.45	0.31,	0.67	0.0001*
Interaction p-value											0.5701
Status somatic BRCA mutat	ions										
sBRCAm	25	10 (40.0)	NE (NE,	NE)	9	8 (88.9)	17.9 (5.3,37.7)	0.23	0.09,	0.60	0.0037*
gBRCAm	69	36 (52.2)	46.9 (36.0,	NE)	36	26 (72.2)	22.2 (16.6,31.9)	0.47	0.28,	0.79	0.0045*
Non-BRCAm	43	26 (60.5)	39.3 (21.2,	NE)	23	21 (91.3)	15.8 (9.5,19.5)	0.30	0.17,	0.54	<0.0001*
Interaction p-value											0.3060

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

Table 1.1.3.2 PAOLA1: Summary of subgroup analysis of Second Progression-free survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

	•	Olaparib + N=2				bevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV)	RS)								
NED [PDS]	92	21 (22.8)	NE (NE, NE)) 48	26 (54.2)	44.3 (33.2, NE)	0.29	0.16,	0.52	<0.0001*
NED/CR [IDS]	74	38 (51.4)	60.0 (32.3, NE)) 38	28 (73.7)	43.3 (24.5,55.6)	0.65	0.40,	1.07	0.0891
NED/CR [Chemo]	40	17 (42.5)	61.3 (37.8, NE)) 20	18 (90.0)	31.2 (24.5,41.0)	0.36	0.18,		0.0030*
PR	49	36 (73.5)	30.6 (23.7,45.0)) 26	18 (69.2)	29.3 (21.0,54.0)	0.97	0.56,	1.75	0.9280
Interaction p-value										0.0138*
Screening laboratory tBR	CA statu	s (IVRS)								
tBRCAm	150	56 (37.3)	75.2 (64.0, NE)) 65	40 (61.5)	46.7 (33.0,55.4)	0.50	0.34,	0.76	0.0013*
non-tBRCAm	105	56 (53.3)	52.0 (35.3, NE)	67	50 (74.6)	32.3 (25.7,41.3)	0.59	0.40,	0.87	0.0075*
Interaction p-value										0.5802
First line treatment out	come (eCl	RF)								
NED [PDS]	89	19 (21.3)	NE (NE, NE)) 47	24 (51.1)	51.9 (35.3, NE)	0.30	0.16,	0.55	0.0001*
NED/CR [IDS]	74	38 (51.4)	60.0 (37.3, NE)) 32	25 (78.1)	43.3 (24.5,55.4)	0.61	0.37,	1.02	0.0568
NED/CR [Chemo]	39	19 (48.7)	57.1 (25.9, NE)) 18	17 (94.4)	30.1 (16.8,38.3)	0.35	0.18,	0.68	0.0023*
PR	50	34 (68.0)	31.9 (28.6,50.8)	34	23 (67.6)	27.5 (21.0,54.0)	0.92	0.54,	1.58	0.7508
Interaction p-value										0.0240*
Screening laboratory tBR	CA statu	s (eCRF)								
tBRCAm	147	55 (37.4)	75.2 (64.0, NE)) 67	42 (62.7)	46.7 (33.0,55.6)	0.49	0.33,	0.74	0.0008*
non-tBRCAm	108	57 (52.8)	53.5 (35.3, NE)) 65	48 (73.8)	32.3 (25.7,38.3)	0.59	0.40,	0.87	0.0078*
Interaction p-value										0.5279
Age group										
<65 years	185	86 (46.5)	62.7 (52.0, NE)) 98	63 (64.3)	37.7 (31.2,47.6)	0.63	0.45,	0.87	0.0054*
>=65 years	70	26 (37.1)	NE (NE, NE)	34	27 (79.4)	34.6 (21.0,53.4)	0.32	0.19,	0.55	<0.0001*
Interaction p-value										0.0381*
FIGO Stage (Disease stat	e)									
III	182	69 (37.9)	75.2 (64.0, NE)	90	55 (61.1)	45.0 (35.4,55.6)	0.52	0.36,	0.74	0.0004*

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

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Table 1.1.3.2 PAOLA1: Summary of subgroup analysis of Second Progression-free survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olapa	rib + (N=2		izumab			Plac	cebo + 1 (N=1	oevaci 132)	izumab				
Subgroup	n	of pa	er (%) tients events		edian ti (95% CI onths))		of pa	er (%) atients events		edian time (95% CI) onths) [a]	Hazard ratio [b]	95% CI	[b]	2-sided o] p-value[b]
IV	73	43	(58.9)	37.8	(29.7,	58.7)	42	35	(83.3)	25.6	(21.8,33.0)	0.54	0.34,	0.84	0.0073*
Interaction p-value															0.9172
Region															
Europe	245	106	(43.3)	75.2	(58.7,	NE)	126	86	(68.3)	37.7	(31.2,45.0)	0.51	0.38,	0.68	<0.0001*
Japan	10	6	(60.0)	33.8	(9.3,	NE)	6	4	(66.7)	41.1	(18.5, NE)	1.02	0.29,	3.97	0.9813
Interaction p-value															0.2916
ECOG performance status at	Basel	ine													
(0) Normal activity	190	87	(45.8)	62.7	(50.8,	NE)	100	67	(67.0)	37.9	(29.8,48.6)	0.59	0.43,	0.82	0.0015*
(1) Restricted activity	61	22	(36.1)	NE	(NE,	NE)	31	23	(74.2)	34.6	(25.1,54.0)	0.34	0.19,	0.62	0.0004*
Interaction p-value															0.1061
Baseline CA-125 value															
<=ULN	228	87	(38.2)	75.2	(75.2,	NE)	118	79	(66.9)	41.0	(32.6,48.6)	0.45	0.33,	0.62	<0.0001*
>ULN	27	25	(92.6)	24.1	(16.3,	33.4)	14	11	(78.6)	22.7	(9.7,45.0)	1.21	0.61,	2.57	0.5927
Interaction p-value															0.0099*
Histological grade															
High grade	255	112	(43.9)	75.2	(57.1,	NE)	132	90	(68.2)	37.7	(31.2,46.7)	0.52	0.40,	0.70	<0.0001*
Interaction p-value															NC
Cytoreductive surgery outc	ome														
No residue	166	59	(35.5)	75.2	(75.2,	NE)	80	50	(62.5)	44.5	(34.6,55.8)	0.46	0.32,	0.68	<0.0001*
Residue	79	45	(57.0)	45.7	(29.7,	58.7)	44	33	(75.0)	29.1	(21.8,37.3)	0.60	0.38,	0.94	0.0265*
Interaction p-value															0.4047
Timing of cytoreductive su	rgery														
Upfront	146	50	(34.2)	75.2	(75.2,	NE)	79	48	(60.8)	37.7	(30.1,50.2)	0.42	0.28,	0.63	<0.0001*
Interval	99	54	(54.5)	53.5	(32.3,	64.0)	45	35	(77.8)	34.6	(24.5,50.0)	0.62	0.41,	0.96	0.0325*
Interaction p-value															0.1929

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

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Table 1.1.3.2 PAOLA1: Summary of subgroup analysis of Second Progression-free survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

		_	bevacizumab 255)	•			oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median ti (95% CI) (months) [Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Myriad tumour BRCA mutat:	ion stat	us									
tBRCAm	158		75.2 (64.0,	NE)	77	51 (66.2)	43.0 (33.0,51.9)	0.47	0.32,	0.68	0.0001*
Non-tBRCAm	97	52 (53.6)	52.0 (32.2,	NE)	55	39 (70.9)	30.1 (25.1,43.8)	0.62	0.41,	0.95	0.0278*
Interaction p-value											0.3191
Status somatic BRCA mutat	ions										
sBRCAm	25	7 (28.0)	NE (NE,	NE)	9	8 (88.9)	37.7 (9.7,50.0)	0.19	0.07,	0.53	0.0020*
gBRCAm	69	28 (40.6)	75.2 (60.0,	NE)	36	21 (58.3)	53.7 (44.5, NE)	0.64	0.37,	1.15	0.1336
Non-BRCAm	43	22 (51.2)	60.3 (29.7,	NE)	23	18 (78.3)	30.1 (20.3,44.3)	0.41	0.22,	0.78	0.0073*
Interaction p-value											0.1114

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

Table 1.1.3.3 PAOLA1: Summary of subgroup analysis of Time to First Subsequent Cancer Therapy or Death Full Analysis Set, HRD[42] positive, DCO 22Mar2022

	•	-		pevacizumab 55)			pevacizumab 132)			
Subgroup		Number (of paties with eve	nts	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outco	ome (IV	RS)								
NED [PDS]	92	•	8)	75.2 (75.2, NE)	48	35 (72.9)	21.2 (17.1,25.2)	0.25	0.16, 0.41	<0.0001*
NED/CR [IDS]	74			29.7 (22.8,39.7)	38		18.4 (15.0,23.1)	0.55	0.36, 0.86	0.0094*
NED/CR [Chemo]	40			54.8 (26.4, NE)	20		14.9 (10.5,22.6)	0.24	0.13, 0.46	<0.0001*
PR	49	,	,	23.2 (17.7,27.9)	26	, ,	15.7 (8.8,22.1)	0.66	0.39, 1.15	0.1384
Interaction p-value										0.0095*
Screening laboratory tBRCA	statu	s (IVRS)								
tBRCAm	150	82 (54	7)	49.6 (38.4, NE)	65	49 (75.4)	20.4 (14.9,23.1)	0.44	0.31, 0.63	<0.0001*
non-tBRCAm	105	61 (58	1)	35.4 (24.8,60.3)	67	59 (88.1)	18.0 (14.8,20.1)	0.39	0.27, 0.55	<0.0001*
Interaction p-value										0.6121
First line treatment outco	ome (eC	RF)								
NED [PDS]	89	30 (33	7)	75.2 (75.2, NE)	47	34 (72.3)	21.5 (17.1,27.8)	0.25	0.15, 0.41	<0.0001*
NED/CR [IDS]	74	53 (71	6)	35.3 (23.1,40.4)	32	27 (84.4)	17.7 (14.4,23.1)	0.55	0.35, 0.88	0.0142*
NED/CR [Chemo]	39	21 (53	8)	39.3 (26.4, NE)	18	17 (94.4)	14.9 (10.5,20.3)	0.22	0.11, 0.42	<0.0001*
PR	50	37 (74	0)	23.8 (17.8,30.8)	34	29 (85.3)	19.0 (9.5,22.1)	0.61	0.38, 1.004	0.0519
Interaction p-value										0.0099*
Screening laboratory tBRCA	statu	s (eCRF)								
tBRCAm	147	81 (55	1)	49.6 (38.4, NE)	67	51 (76.1)	20.0 (14.9,22.5)	0.43	0.30, 0.61	<0.0001*
non-tBRCAm	108	62 (57	4)	38.5 (24.8, NE)	65	57 (87.7)	18.4 (14.8,20.3)	0.39	0.27, 0.56	<0.0001*
Interaction p-value										0.6938
Age group										
<65 years	185	108 (58	4)	39.2 (31.9,50.7)	98	81 (82.7)	18.5 (14.9,21.7)	0.43	0.32, 0.58	<0.0001*
>=65 years	70	35 (50	0)	60.3 (39.1, NE)	34	27 (79.4)	19.1 (10.9,20.6)	0.33	0.20, 0.55	<0.0001*
Interaction p-value										0.3599
FIGO Stage (Disease state)	1									
III	182	95 (52	2)	52.5 (39.2, NE)	90	69 (76.7)	20.4 (18.5,23.1)	0.43	0.32, 0.59	<0.0001*

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

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Table 1.1.3.3 PAOLA1: Summary of subgroup analysis of Time to First Subsequent Cancer Therapy or Death Full Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olap	arib + (N=2	bevacizumab 255)		Placebo + (N=	bevac: 132)	izumab				
Subgroup	n	of pa	er (%) atients events	Median time (95% CI) (months) [a]	n	Number (%) of patients with events	;	edian time (95% CI) onths) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
IV	73	48	(65.8)	29.2 (22.8,41.3)	42	39 (92.9)	13.9	(10.9,15.4)	0.34	0.22,	0.52	<0.0001*
Interaction p-value												0.3547
Region												
Europe	245	136	(55.5)	43.0 (37.0,61.2)	126	102 (81.0)	18.5	(15.4,20.5)	0.40	0.31,	0.52	<0.0001*
Japan	10	7	(70.0)	23.9 (6.3, NE)	6	6 (100)	21.5	(12.0, NE)	0.58	0.19,	1.79	0.3286
Interaction p-value												0.5178
ECOG performance status at	Basel	ine										
(0) Normal activity	190	106	(55.8)	41.3 (32.4,64.0)	100	82 (82.0)	19.7	(15.7,21.2)	0.42	0.32,	0.57	<0.0001*
(1) Restricted activity	61	34	(55.7)	48.1 (28.6, NE)	31	26 (83.9)	17.5	(10.5,22.5)	0.31	0.19,	0.52	<0.0001*
Interaction p-value												0.3032
Baseline CA-125 value												
<=ULN	228	117	(51.3)	54.8 (40.5, NE)	118	96 (81.4)	19.7	(17.1,21.0)	0.36	0.27,	0.47	<0.0001*
>ULN	27	26	(96.3)	19.1 (12.4,24.1)	14	12 (85.7)	10.5	(7.3,25.2)	0.96	0.49,	1.97	0.9012
Interaction p-value												0.0069*
Histological grade												
High grade	255	143	(56.1)	42.2 (35.7,60.3)	132	108 (81.8)	18.8	(16.1,20.5)	0.40	0.31,	0.52	<0.0001*
Interaction p-value												NC
Cytoreductive surgery outc	ome											
No residue	166	85	(51.2)	56.6 (41.3, NE)	80	62 (77.5)	20.1	(17.3,23.1)	0.40	0.29,	0.55	<0.0001*
Residue	79	48	(60.8)	31.6 (25.3,43.7)	44	38 (86.4)	14.9	(10.8,20.3)	0.41	0.27,	0.64	<0.0001*
Interaction p-value												0.8754
Timing of cytoreductive su	rgery											
Upfront	146	64	(43.8)	75.2 (56.6, NE)	79	60 (75.9)	20.1	(14.9,22.2)	0.32	0.22,	0.46	<0.0001*
Interval	99	69	(69.7)	31.6 (23.5,39.2)	45	40 (88.9)	17.6	(14.4,20.3)	0.51	0.35,	0.76	0.0011*
Interaction p-value												0.0823

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

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Table 1.1.3.3 PAOLA1: Summary of subgroup analysis of Time to First Subsequent Cancer Therapy or Death Full Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + 1 (N=2				oevacizumab 132)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Myriad tumour BRCA mutat:	ion stati	ıs								
tBRCAm	158		49.6 (39.1, NE)	77	62 (80.5)	19.2 (14.9,22.1)	0.37	0.27,	0.52	<0.0001*
Non-tBRCAm	97	58 (59.8)	31.6 (23.5,60.3)	55	46 (83.6)	18.8 (14.8,20.3)	0.46	0.31,	0.68	0.0001*
Interaction p-value										0.4234
Status somatic BRCA mutat	ions									
sBRCAm	25	13 (52.0)	61.2 (35.2, NE)	9	8 (88.9)	16.1 (6.9,33.3)	0.24	0.10,	0.61	0.0039*
gBRCAm	69	37 (53.6)	48.7 (35.7, NE)	36	29 (80.6)	22.4 (15.4,32.5)	0.43	0.26,	0.70	0.0009*
Non-BRCAm	43	25 (58.1)	41.3 (21.9, NE)	23	21 (91.3)	17.6 (10.8,20.3)	0.30	0.16,	0.54	0.0001*
Interaction p-value										0.4535

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

Table 1.1.3.4 PAOLA1: Summary of subgroup analysis of Time to Second Subsequent Cancer Therapy or Death Full Analysis Set, HRD[42] positive, DCO 22Mar2022

		-	bevacizumab 255)	:	Placebo + k (N=1				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IV)	RS)							
NED [PDS]	92	22 (23.9)	NE (NE, NE)	48	28 (58.3)	46.5 (29.5, NE)	0.29	0.16, 0.50	<0.0001*
NED/CR [IDS]	74	, ,	47.0 (33.8, NE)	38		39.0 (24.9,55.6)	0.65	0.40, 1.0	0.0895
NED/CR [Chemo]	40	17 (42.5)	NE (NE, NE)	20		29.8 (24.6,40.1)	0.31	0.16, 0.61	0.0007*
PR	49	36 (73.5)	31.8 (25.6,45.2)	26	18 (69.2)	29.9 (21.4,54.0)	0.98	0.56, 1.76	0.9441
Interaction p-value									0.0058*
Screening laboratory tBR0	CA statu	s (IVRS)							
tBRCAm	150	61 (40.7)	75.2 (62.5, NE)	65	43 (66.2)	45.0 (28.6,54.0)	0.49	0.33, 0.73	0.0006*
non-tBRCAm	105	54 (51.4)	54.5 (38.6, NE)	67	49 (73.1)	32.4 (25.2,40.1)	0.56	0.38, 0.82	0.0034*
Interaction p-value									0.6663
First line treatment out	come (eCl	RF)							
NED [PDS]	89	20 (22.5)	NE (NE, NE)	47	26 (55.3)	49.7 (29.5, NE)	0.29	0.16, 0.52	<0.0001*
NED/CR [IDS]	74	40 (54.1)	54.5 (38.0, NE)	32	24 (75.0)	39.0 (24.9,55.6)	0.63	0.38, 1.07	0.0842
NED/CR [Chemo]	39	19 (48.7)	57.1 (35.3, NE)	18	17 (94.4)	29.2 (18.9,36.8)	0.30	0.16, 0.59	0.0006*
PR	50	34 (68.0)	33.9 (28.8,52.0)	34	24 (70.6)	29.1 (21.4,54.0)	0.88	0.52, 1.49	0.6190
Interaction p-value									0.0128*
Screening laboratory tBR0	CA statu	s (eCRF)							
tBRCAm	147	60 (40.8)	75.2 (62.5, NE)	67	44 (65.7)	45.0 (28.6,54.0)	0.49	0.34, 0.73	0.0006*
non-tBRCAm	108	55 (50.9)	55.5 (38.8, NE)	65	48 (73.8)	32.4 (25.2,40.1)	0.55	0.37, 0.83	0.0026*
Interaction p-value									0.7263
Age group									
<65 years	185	88 (47.6)	67.1 (54.2, NE)	98	65 (66.3)	35.3 (28.6,45.8)	0.58	0.42, 0.83	0.0012*
>=65 years	70	27 (38.6)	NE (NE, NE)	34	27 (79.4)	36.8 (25.1,51.0)	0.35	0.20, 0.59	
Interaction p-value									0.0999
FIGO Stage (Disease state	e)								
III	182	72 (39.6)	75.2 (67.1, NE)	90	57 (63.3)	45.5 (35.3,55.6)	0.51	0.36, 0.72	0.0002*

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

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Table 1.1.3.4 PAOLA1: Summary of subgroup analysis of Time to Second Subsequent Cancer Therapy or Death Full Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib (+ bev 1=255				Place		oevaci 132)	zumab				
Subgroup	n	Number (9 of patien with even	s	Median ti (95% CI (months))		Numbe: of pat with e	ients		edian time (95% CI) onths) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
IV	73	43 (58.	9) 40	.0 (30.3,	NE)	42	35 (83.3)	26.8	(22.0,33.6)	0.51	0.32,	0.80	0.0037*
Interaction p-value														0.9998
Region														
Europe	245	109 (44.	5) 75	.2 (57.1,	NE)	126	87 (69.0)	36.4	(29.5,46.5)	0.51	0.38,	0.67	<0.0001*
Japan	10	6 (60.)) 35	.6 (7.0,	NE)	6	5 (83.3)	26.9	(20.2, NE)	0.58	0.17,	2.02	0.3772
Interaction p-value														0.8240
ECOG performance status at	Basel	ine												
(0) Normal activity	190	88 (46.	3)	NE (NE,	NE)	100	68 (68.0)	36.8	(29.5,47.5)	0.57	0.41,	0.78	0.0005*
(1) Restricted activity	61	24 (39.	3)	NE (NE,	NE)	31	24 (77.4)	31.6	(23.2,46.5)	0.33	0.19,	0.59	0.0002*
Interaction p-value														0.1105
Baseline CA-125 value														
<=ULN	228	90 (39.	5) 75	.2 (75.2,	NE)	118	80 (67.8)	37.4	(29.9,47.5)	0.44	0.33,	0.60	<0.0001*
>ULN	27	25 (92.	5) 27	.7 (17.1,	33.8)	14	12 (85.7)	24.1	(11.8,45.8)	1.10	0.56,	2.27	0.7839
Interaction p-value														0.0154*
Histological grade														
High grade	255	115 (45.	L) 75	.2 (56.2,	NE)	132	92 (69.7)	35.4	(29.2,45.2)	0.51	0.39,	0.67	<0.0001*
Interaction p-value														NC
Cytoreductive surgery outc	ome													
No residue	166	62 (37.	3) 75	.2 (75.2,	NE)	80	51 (63.8)	45.2	(32.9,56.2)	0.47	0.32,	0.68	<0.0001*
Residue	79	45 (57.)) 51	.3 (33.8,	NE)	44	34 (77.3)	26.5	(21.4,36.8)	0.54	0.35,	0.85	0.0078*
Interaction p-value														0.6202
Timing of cytoreductive su	rgery													
Upfront	146	51 (34.	9) 75	.2 (75.2,	NE)	79	50 (63.3)	36.0	(29.2,46.5)	0.40	0.27,	0.60	<0.0001*
Interval	99	56 (56.	5) 47	.0 (34.0,	67.1)	45	35 (77.8)	34.2	(24.9,51.0)	0.61	0.40,	0.94	0.0257*
Interaction p-value														0.1522

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

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Table 1.1.3.4 PAOLA1: Summary of subgroup analysis of Time to Second Subsequent Cancer Therapy or Death Full Analysis Set, HRD[42] positive, DCO 22Mar2022

	•	Olap	arib + (N=	bevac: 255)	izumab	•		Placebo + (N	bevac =132)	izumab							
Subgroup		of pa	er (%) atients events	(dian ti (95% CI onths))		Number (%) of patient with event	3	edian time (95% CI) onths) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]			
Myriad tumour BRCA mutati	on stat	นร															
tBRCAm	158	64	(40.5)	75.2	(67.1,	NE)	77	53 (68.8	40.0	(28.6,52.5)	0.45	0.32,	0.65	<0.0001*			
Non-tBRCAm	97	51	(52.6)	54.2	(35.3,	NE)	55	39 (70.9	32.2	(25.2,41.4)	0.61	0.40,	0.93	0.0226*			
Interaction p-value														0.2925			
Status somatic BRCA mutat	ions																
sBRCAm	25	9	(36.0)	NE	(NE,	NE)	9	8 (88.9	32.9	(10.2,52.5)	0.23	0.09,	0.61	0.0042*			
gBRCAm	69	27	(39.1)	75.2	(61.7,	NE)	36	23 (63.9	49.7	(26.8,66.9)	0.52	0.30,	0.92	0.0254*			
Non-BRCAm	43	22	(51.2)	60.3	(32.0,	NE)	23	18 (78.3	30.4	(20.8,46.5)	0.44	0.24,	0.84	0.0129*			
Interaction p-value														0.3444			

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

Table 1.1.3.5 PAOLA1: Summary of subgroup analysis of Overall Survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

		-	bevacizumab 255)				bevacizumab 132)			
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	ome (IV	RS)								
NED [PDS]	92	15 (16.3)	NE (NE,	NE)	48	21 (43.8)	NE (NE, NE)	0.29	0.15, 0.5	7 0.0003*
NED/CR [IDS]	74	, ,	73.3 (45.0,	NE)	38	• •	57.3 (45.2, NE)	0.88	0.51, 1.5	
NED/CR [Chemo]	40	15 (37.5)	NE (NE,	NE)	20		56.9 (31.8,66.4)	0.56	0.26, 1.2	
PR	49	29 (59.2)	50.4 (32.3,	NE)	26	16 (61.5)	43.0 (25.2, NE)	0.88	0.48, 1.6	0.6789
Interaction p-value										0.0501
Screening laboratory tBRC	!A statu	s (IVRS)								
tBRCAm	150	46 (30.7)	75.2 (73.3,	NE)	65	30 (46.2)	66.9 (54.9, NE)	0.61	0.38, 0.9	7 0.0370*
non-tBRCAm	105	47 (44.8)	NE (NE,	NE)	67	39 (58.2)	52.0 (41.0,66.4)	0.68	0.44, 1.0	0.0774
Interaction p-value										0.7229
First line treatment outo	ome (eC	RF)								
NED [PDS]	89	14 (15.7)	NE (NE,	NE)	47	19 (40.4)	NE (NE, NE)	0.31	0.15, 0.6	2 0.0009*
NED/CR [IDS]	74	32 (43.2)	73.3 (45.7,	NE)	32	18 (56.3)	55.6 (43.1, NE)	0.75	0.42, 1.3	0.3364
NED/CR [Chemo]	39	17 (43.6)	NE (NE,	NE)	18	11 (61.1)	48.4 (25.5, NE)	0.63	0.30, 1.3	0.2366
PR	50	28 (56.0)	54.0 (37.2,	NE)	34	20 (58.8)	48.5 (30.3, NE)	0.89	0.51, 1.6	1 0.7003
Interaction p-value										0.1187
Screening laboratory tBRC	'A statu	s (eCRF)								
tBRCAm	147	45 (30.6)	75.2 (73.3,	NE)	67	31 (46.3)	66.9 (54.9, NE)	0.60	0.38, 0.9	5 0.0300*
non-tBRCAm	108	48 (44.4)	NE (NE,	NE)	65	38 (58.5)	52.0 (41.0,66.4)	0.68	0.44, 1.0	0.0744
Interaction p-value										0.6961
Age group										
<65 years	185	69 (37.3)	75.2 (73.3,	NE)	98	49 (50.0)	65.4 (48.9, NE)	0.69	0.48, 1.00	0.0500
>=65 years	70	24 (34.3)	NE (NE,	NE)	34	20 (58.8)	54.0 (34.0, NE)	0.46	0.25, 0.8	
Interaction p-value										0.2496
FIGO Stage (Disease state	·)									
III	182	56 (30.8)	75.2 (75.2,	NE)	90	41 (45.6)	66.4 (55.6, NE)	0.60	0.40, 0.9	0.0147*

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

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Table 1.1.3.5 PAOLA1: Summary of subgroup analysis of Overall Survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olap	arib + (N=2	bevac 255)	izumab	•		Plac	ebo + 1 (N=)	bevaci 132)	izumab					
Subgroup		of pa	er (%) atients events		edian ti (95% CI onths))		of pa	er (%) atients events		edian ti (95% CI onths))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
IV	73	37	(50.7)	63.2	(38.5,	NE)	42	28	(66.7)	45.7	(34.3,	55.7)	0.69	0.42,	1.13	0.1368
Interaction p-value																0.6798
Region																
Europe	245	87	(35.5)	75.2	(73.3,	NE)	126	66	(52.4)	57.3	(51.6,	NE)	0.60	0.43,	0.83	0.0020*
Japan	10	6	(60.0)	57.1	(12.3,	NE)	6	3	(50.0)	NE	(NE,	NE)	1.21	0.32,	5.73	0.7862
Interaction p-value																0.3194
ECOG performance status at	Basel	ine														
(0) Normal activity	190	73	(38.4)	NE	(NE,	NE)	100	52	(52.0)	59.8	(48.9,	NE)	0.69	0.48,	0.99	0.0428*
(1) Restricted activity	61	17	(27.9)	NE	(NE,	NE)	31	17	(54.8)	54.0	(35.2,	NE)	0.39	0.20,	0.78	0.0079*
Interaction p-value																0.1525
Baseline CA-125 value																
<=ULN	228	72	(31.6)	75.2	(73.3,	NE)	118	59	(50.0)	59.8	(52.4,	NE)	0.55	0.39,	0.77	0.0007*
>ULN	27	21	(77.8)	38.0	(21.7,	56.2)	14	10	(71.4)	42.0	(16.0,	NE)	1.28	0.62,	2.83	0.5207
Interaction p-value																0.0395*
Histological grade																
High grade	255	93	(36.5)	75.2	(73.3,	NE)	132	69	(52.3)	57.3	(51.6,	NE)	0.62	0.45,	0.85	0.0031*
Interaction p-value																NC
Cytoreductive surgery outc	ome															
No residue	166	48	(28.9)	75.2	(73.3,	NE)	80	38	(47.5)	NE	(NE,	NE)	0.54	0.35,	0.83	0.0054*
Residue	79	37	(46.8)	60.8	(50.5,	NE)	44	26	(59.1)	42.2	(30.3,	66.4)	0.69	0.42,	1.15	0.1516
Interaction p-value																0.4650
Timing of cytoreductive su	rgery															
Upfront	146	41	(28.1)	NE	(NE,	NE)	79	38	(48.1)	65.4	(46.3,	NE)	0.48	0.31,	0.75	0.0014*
Interval	99	44	(44.4)	73.3	(47.9,	NE)	45	26	(57.8)	55.6	(41.0,	NE)	0.75	0.46,	1.23	0.2450
Interaction p-value																0.1868

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

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Table 1.1.3.5 PAOLA1: Summary of subgroup analysis of Overall Survival Full Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olap		bevacizu 255)	ımab			Placebo + (N=	bevaciz 132)	zumab	•				
Subgroup		of pa	er (%) atients events	Media (95) (mont)	% CI))		Number (%) of patients with events	(!	lian ti 95% CI nths))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Myriad tumour BRCA mutat:	on stat	นร													
tBRCAm	158	49	(31.0)	75.2 (7	3.3,	NE)	77	37 (48.1)	66.9	(54.0,	NE)	0.57	0.37,	0.88	0.0117*
Non-tBRCAm	97	44	(45.4)	NE (NE,	NE)	55	32 (58.2)				0.70	0.45,	1.12	0.1369
Interaction p-value															0.5114
Status somatic BRCA mutat	ions														
sBRCAm	25	5	(20.0)	NE (NE,	NE)	9	4 (44.4)	NE	(NE,	NE)	0.36	0.09,	1.45	0.1412
gBRCAm	69	19	(27.5)	75.2 (7	3.3,	NE)	36	13 (36.1)	NE	(NE,	NE)	0.74	0.37,	1.54	0.4113
Non-BRCAm	43	17	(39.5)	NE (NE,	NE)	23	15 (65.2)	52.4	(35.2,	NE)	0.45	0.22,	0.91	0.0273*
Interaction p-value															0.4888

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and, if appropriate, interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only.

* Interaction p-value <0.05. HR <1 favours olaparib. NC = not calculable.

Figure 1.1.4.1 PAOLA1: Kaplan-Meier plot of Progression-free survival for First line treatment outcome (IVRS) = NED [PDS]
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

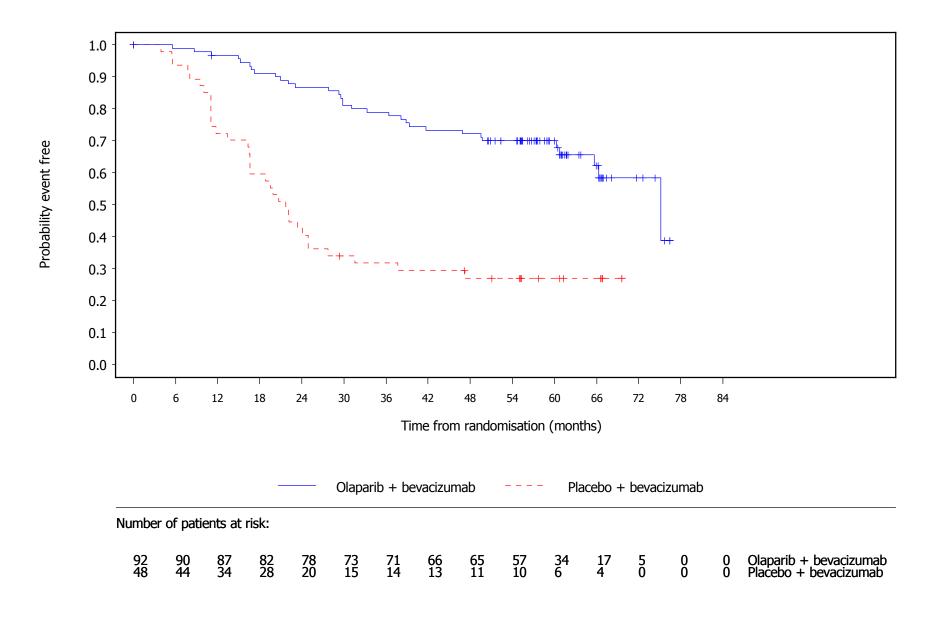


Figure 1.1.4.2 PAOLA1: Kaplan-Meier plot of Progression-free survival for First line treatment outcome (IVRS) = NED/CR [IDS]
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

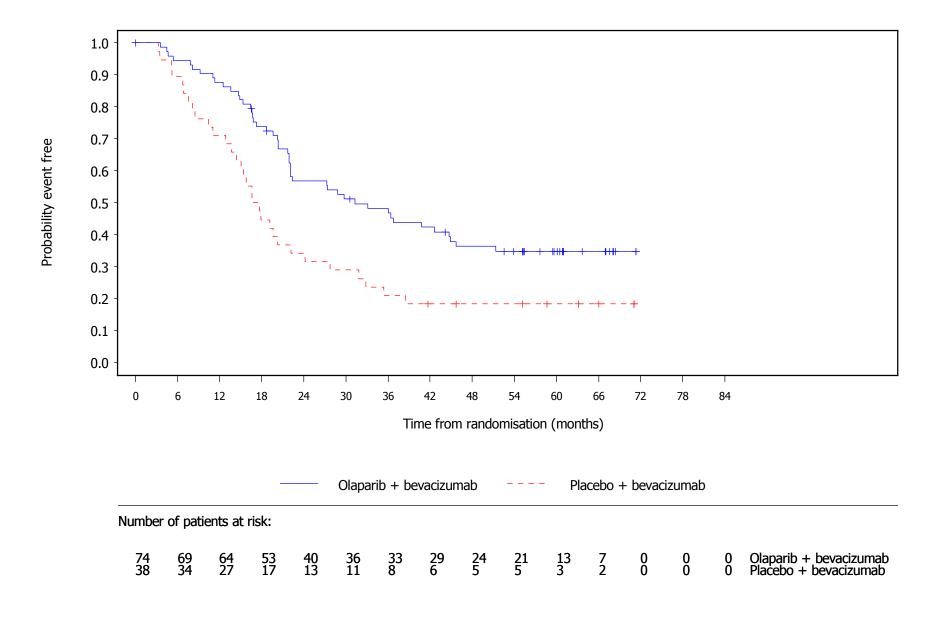


Figure 1.1.4.3 PAOLA1: Kaplan-Meier plot of Progression-free survival for First line treatment outcome (IVRS) = NED/CR [Chemo]
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

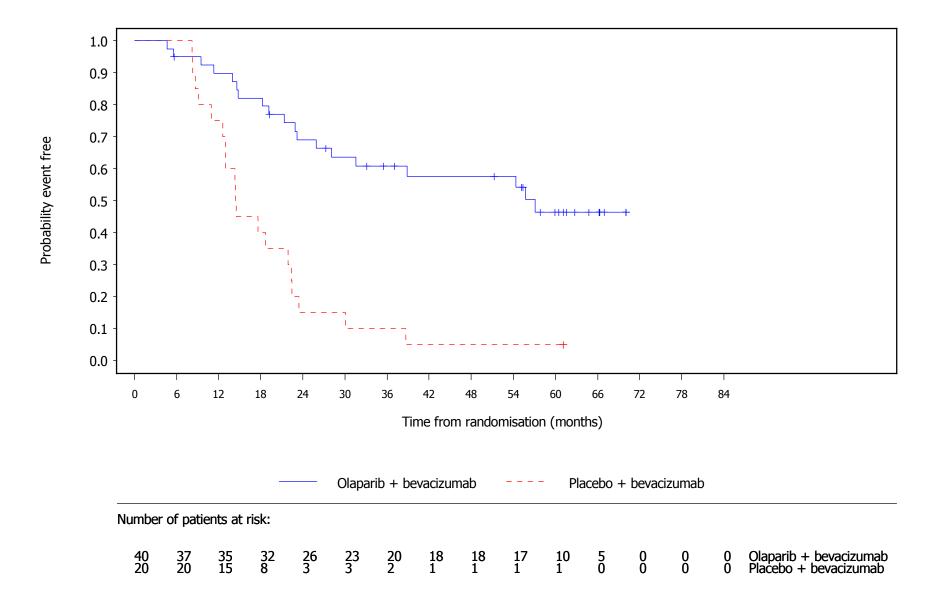


Figure 1.1.4.4 PAOLA1: Kaplan-Meier plot of Progression-free survival for First line treatment outcome (IVRS) = PR
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

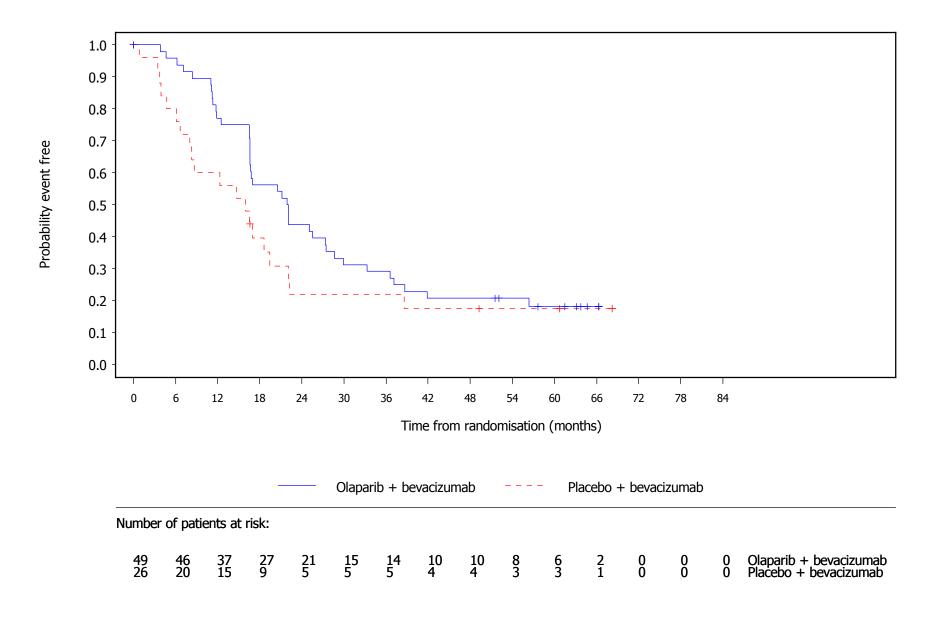


Figure 1.1.4.5 PAOLA1: Kaplan-Meier plot of Progression-free survival for First line treatment outcome (eCRF) = NED [PDS]
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

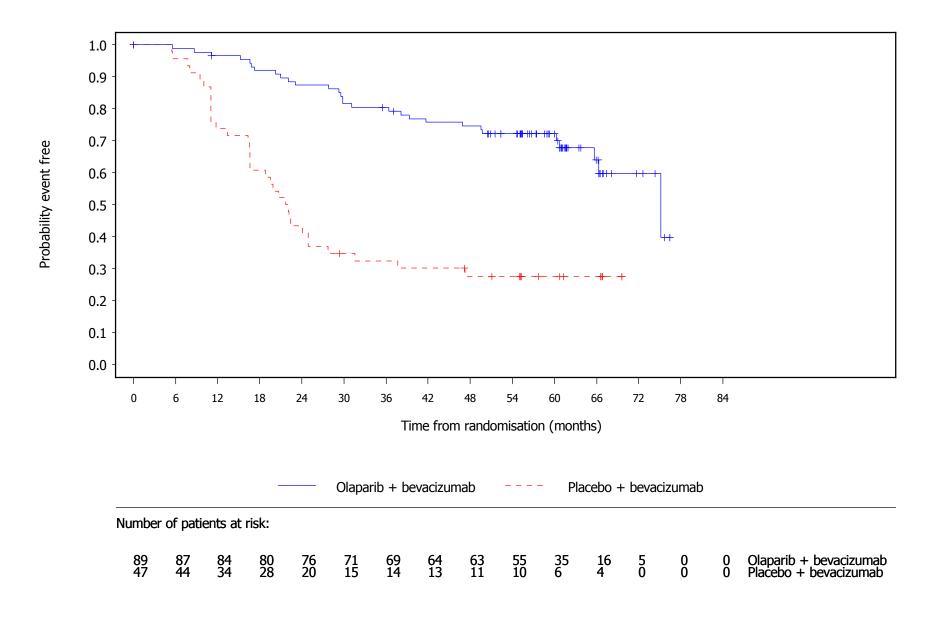


Figure 1.1.4.6 PAOLA1: Kaplan-Meier plot of Progression-free survival for First line treatment outcome (eCRF) = NED/CR [IDS]
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

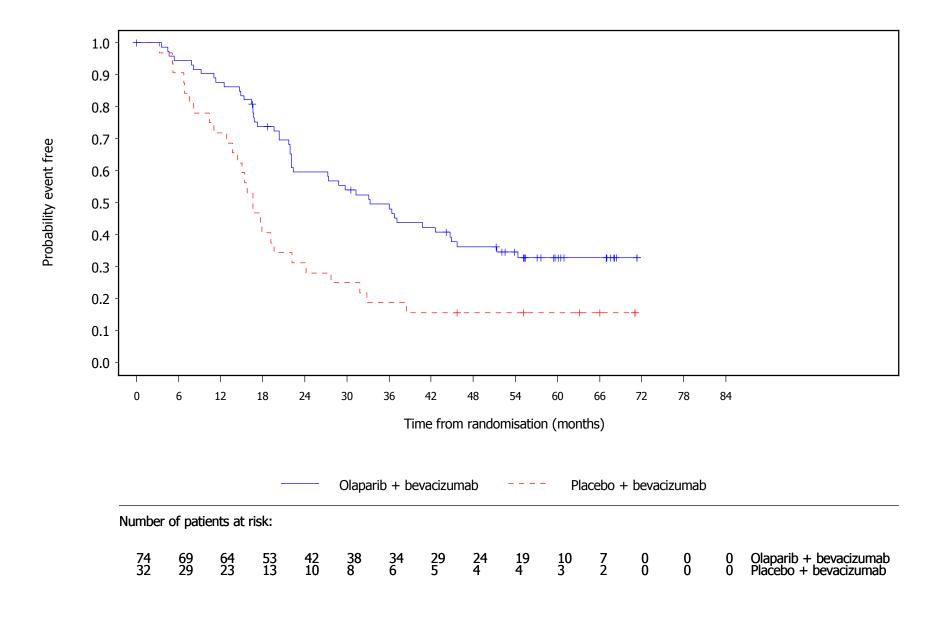


Figure 1.1.4.7 PAOLA1: Kaplan-Meier plot of Progression-free survival for First line treatment outcome (eCRF) = NED/CR [Chemo]
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

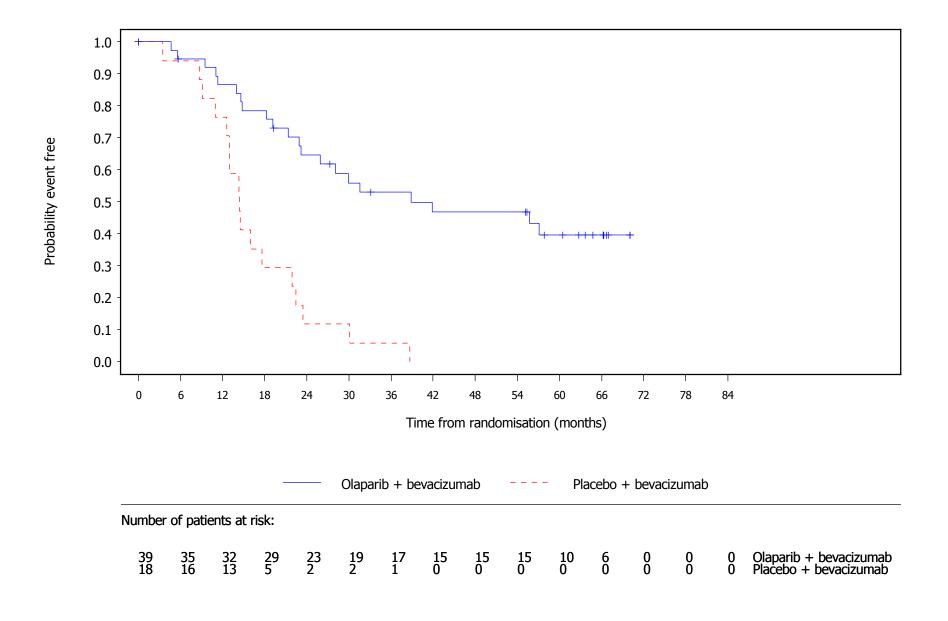


Figure 1.1.4.8 PAOLA1: Kaplan-Meier plot of Progression-free survival for First line treatment outcome (eCRF) = PR
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

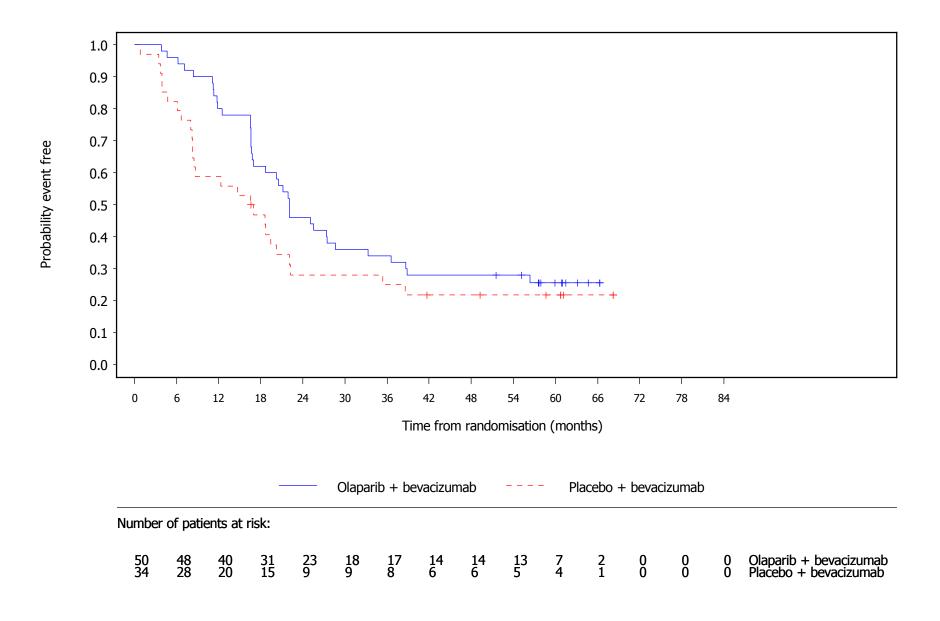


Figure 1.1.4.9 PAOLA1: Kaplan-Meier plot of Progression-free survival for Baseline CA-125 value = <=ULN Full Analysis Set, HRD[42] positive, DCO 22Mar2022

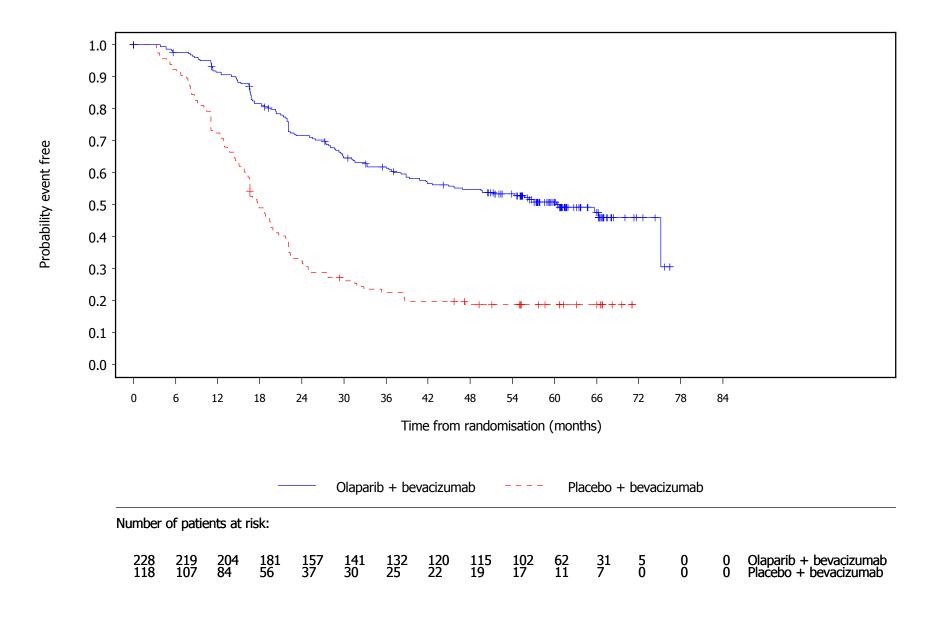


Figure 1.1.4.10 PAOLA1: Kaplan-Meier plot of Progression-free survival for Baseline CA-125 value = >ULN Full Analysis Set, HRD[42] positive, DCO 22Mar2022

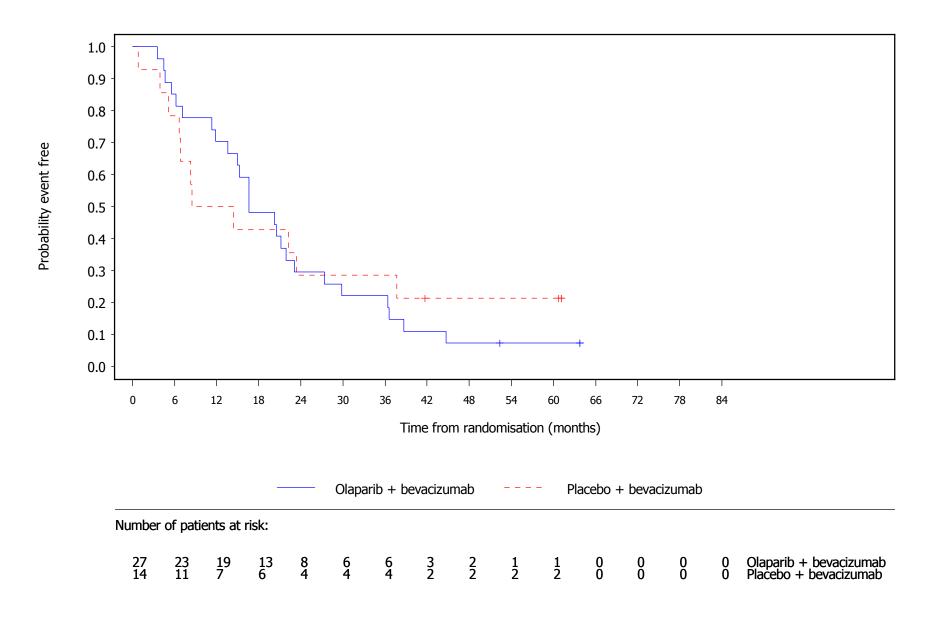


Figure 1.1.4.11 PAOLA1: Kaplan-Meier plot of Second Progression-free survival for First line treatment outcome (IVRS) = NED [PDS]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

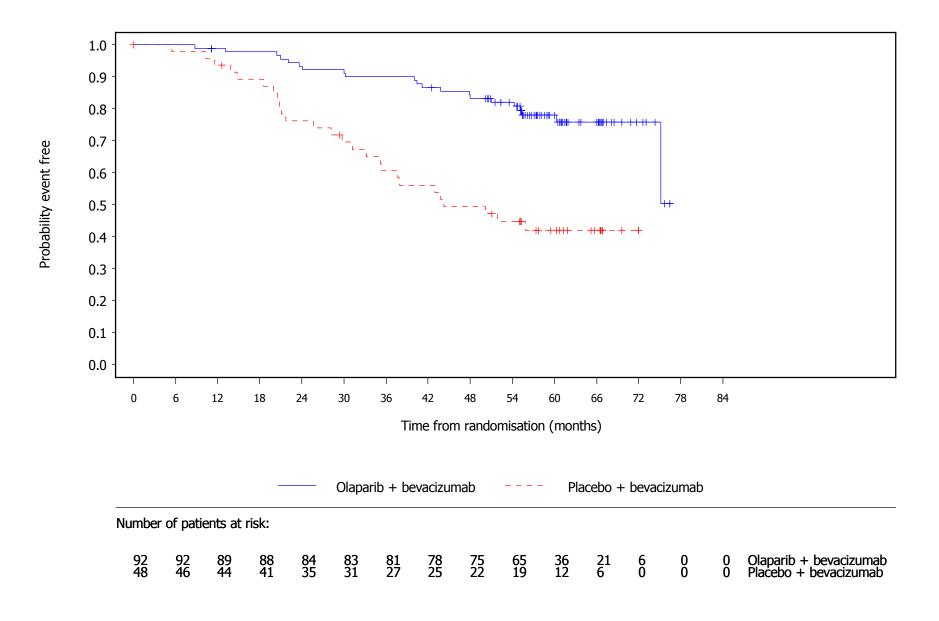


Figure 1.1.4.12 PAOLA1: Kaplan-Meier plot of Second Progression-free survival for First line treatment outcome (IVRS) = NED/CR
[IDS]
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

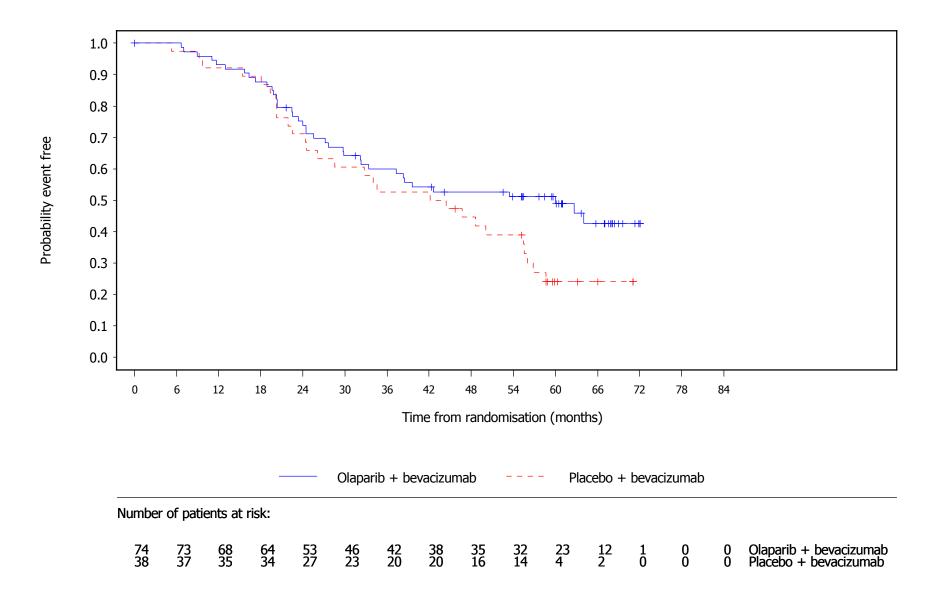


Figure 1.1.4.13 PAOLA1: Kaplan-Meier plot of Second Progression-free survival for First line treatment outcome (IVRS) = NED/CR [Chemo]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

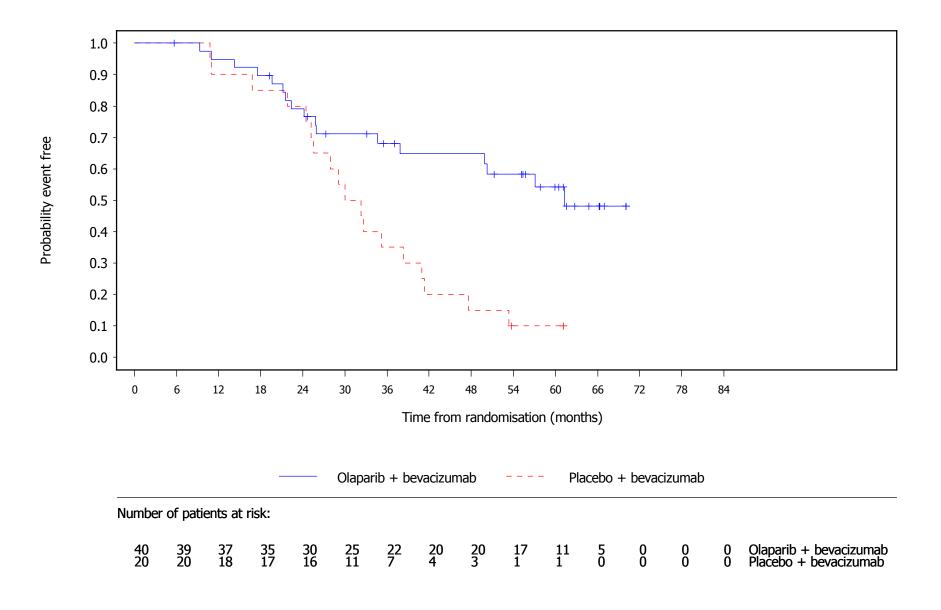


Figure 1.1.4.14 PAOLA1: Kaplan-Meier plot of Second Progression-free survival for First line treatment outcome (IVRS) = PR
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

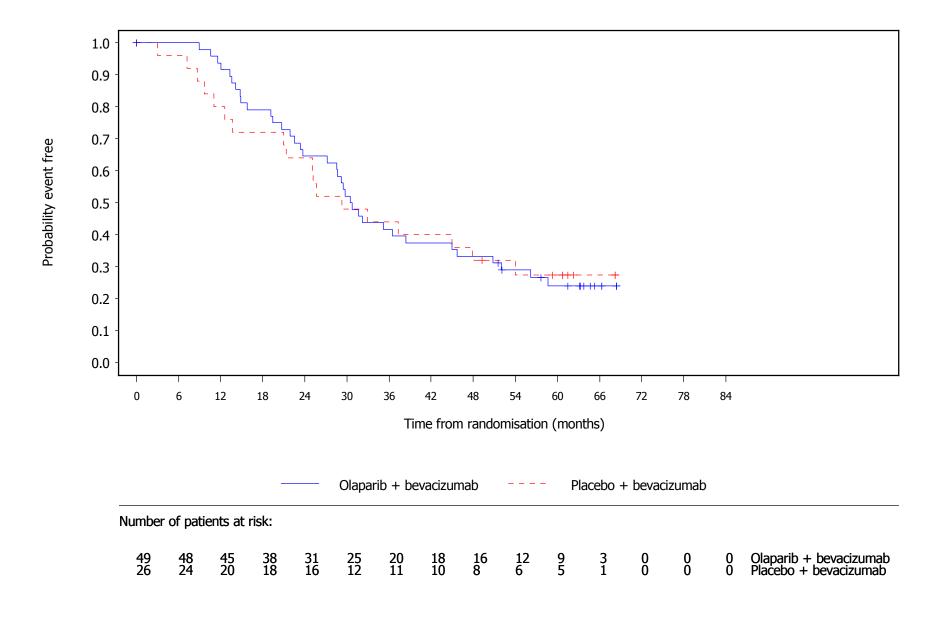


Figure 1.1.4.15 PAOLA1: Kaplan-Meier plot of Second Progression-free survival for First line treatment outcome (eCRF) = NED [PDS]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

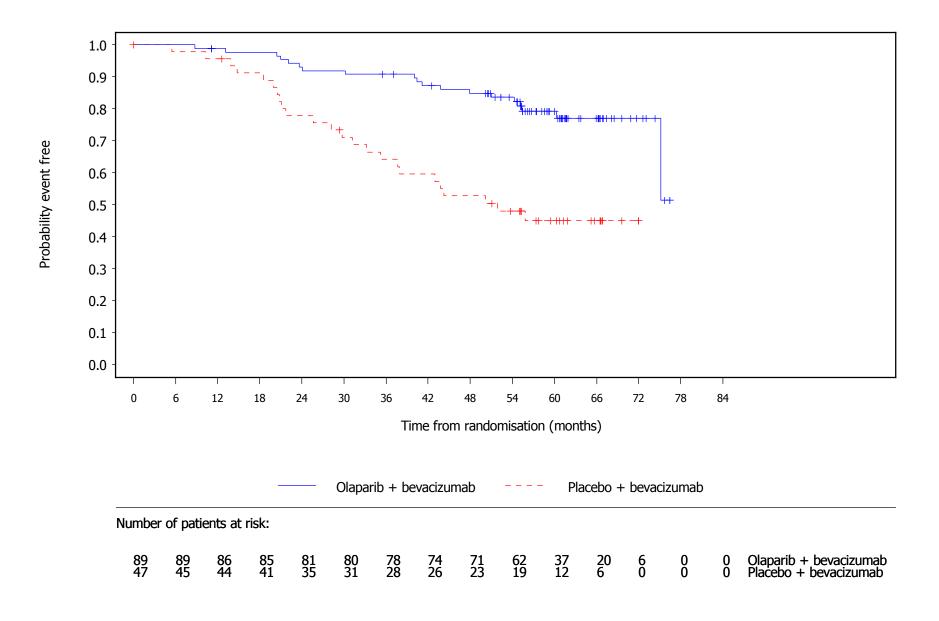


Figure 1.1.4.16 PAOLA1: Kaplan-Meier plot of Second Progression-free survival for First line treatment outcome (eCRF) = NED/CR [IDS]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

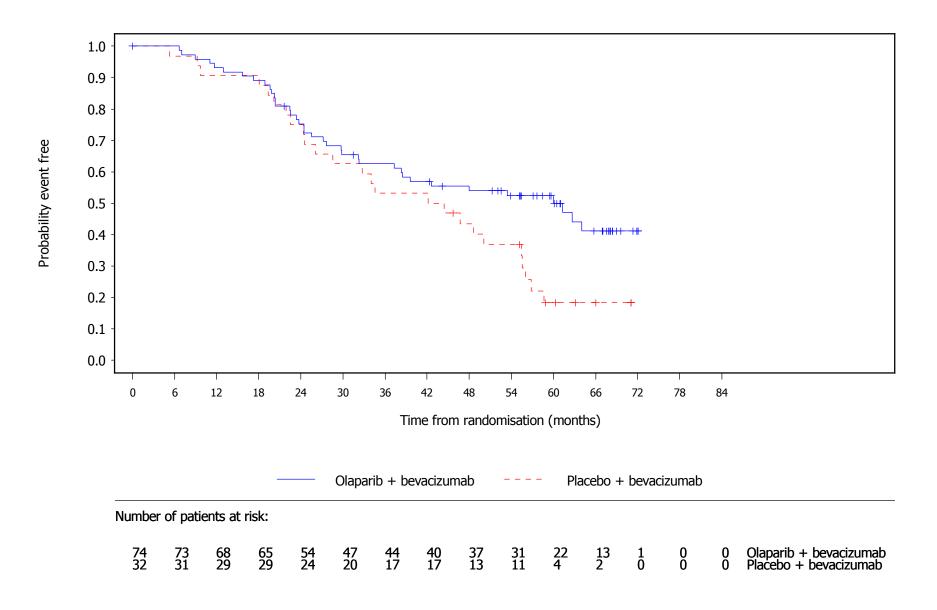


Figure 1.1.4.17 PAOLA1: Kaplan-Meier plot of Second Progression-free survival for First line treatment outcome (eCRF) = NED/CR [Chemo]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

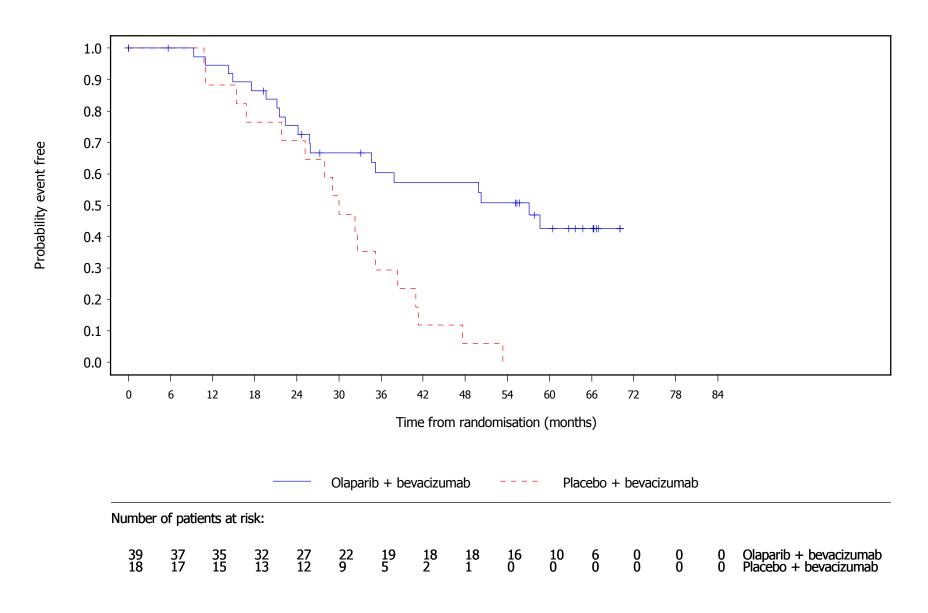


Figure 1.1.4.18 PAOLA1: Kaplan-Meier plot of Second Progression-free survival for First line treatment outcome (eCRF) = PR
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

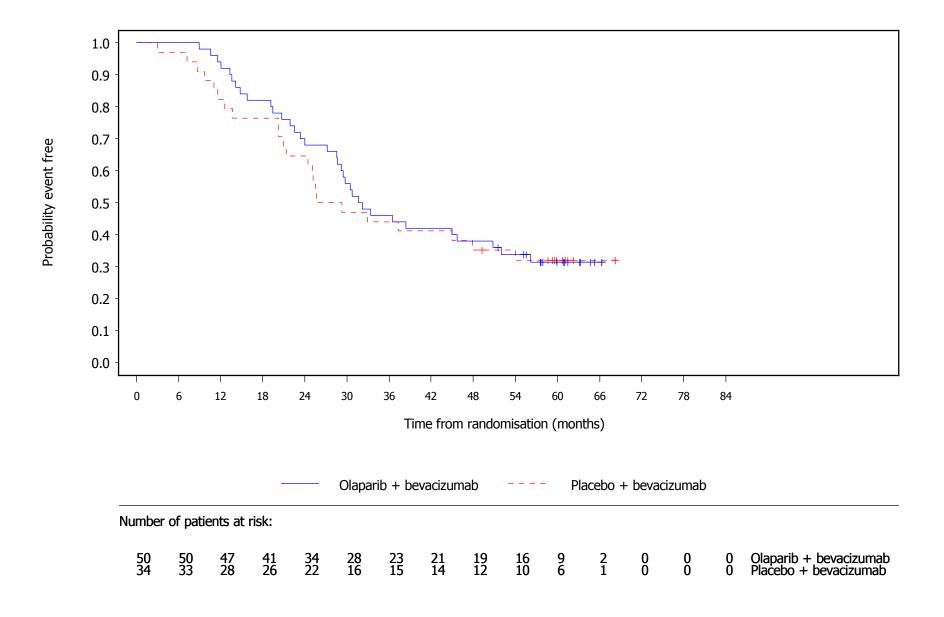


Figure 1.1.4.19 PAOLA1: Kaplan-Meier plot of Second Progression-free survival for Age group = <65 years
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

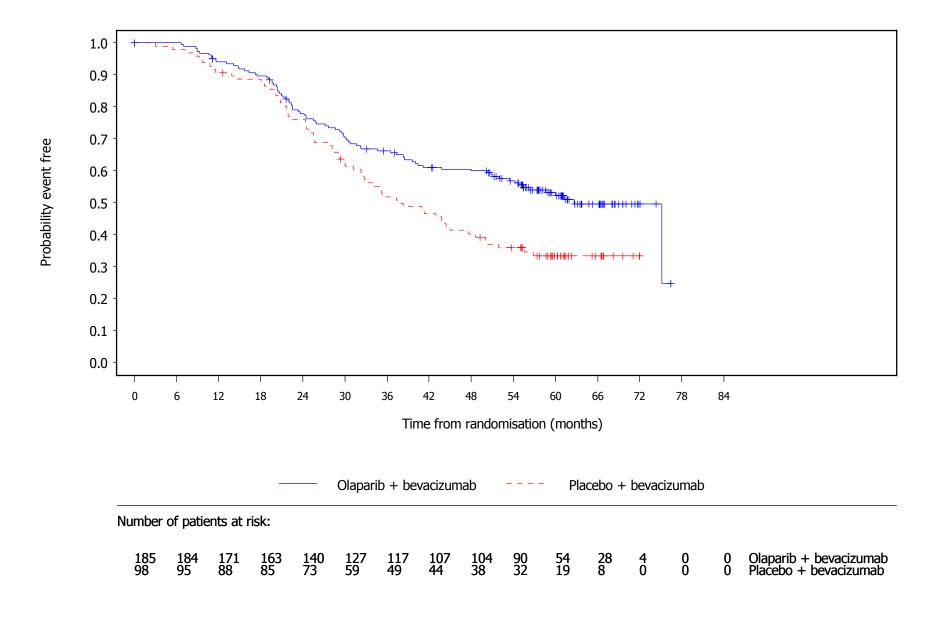


Figure 1.1.4.20 PAOLA1: Kaplan-Meier plot of Second Progression-free survival for Age group = >=65 years
Full Analysis Set, HRD[42] positive, DCO 22Mar2022

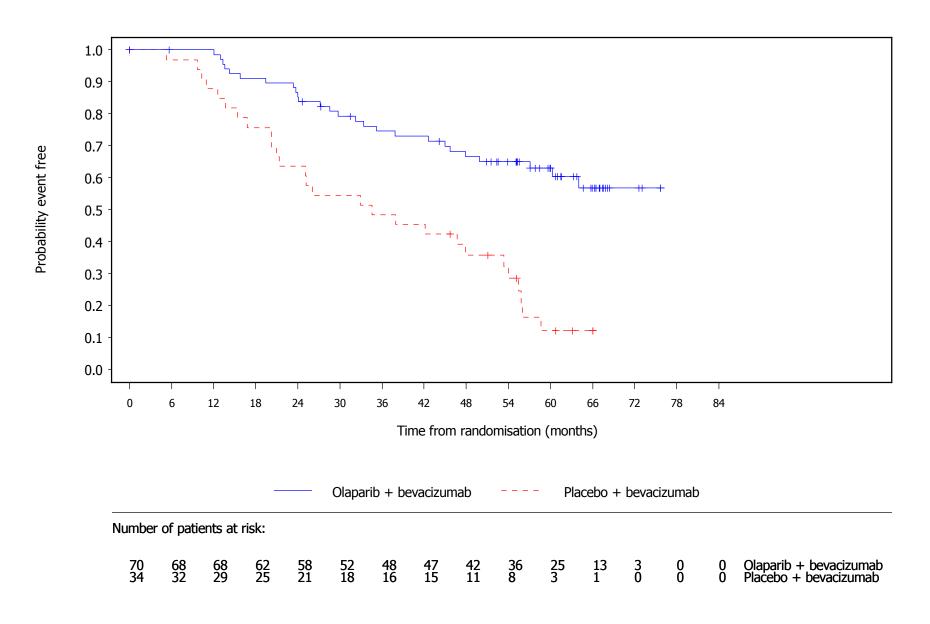


Figure 1.1.4.21 PAOLA1: Kaplan-Meier plot of Second Progression-free survival for Baseline CA-125 value = <=ULN Full Analysis Set, HRD[42] positive, DCO 22Mar2022

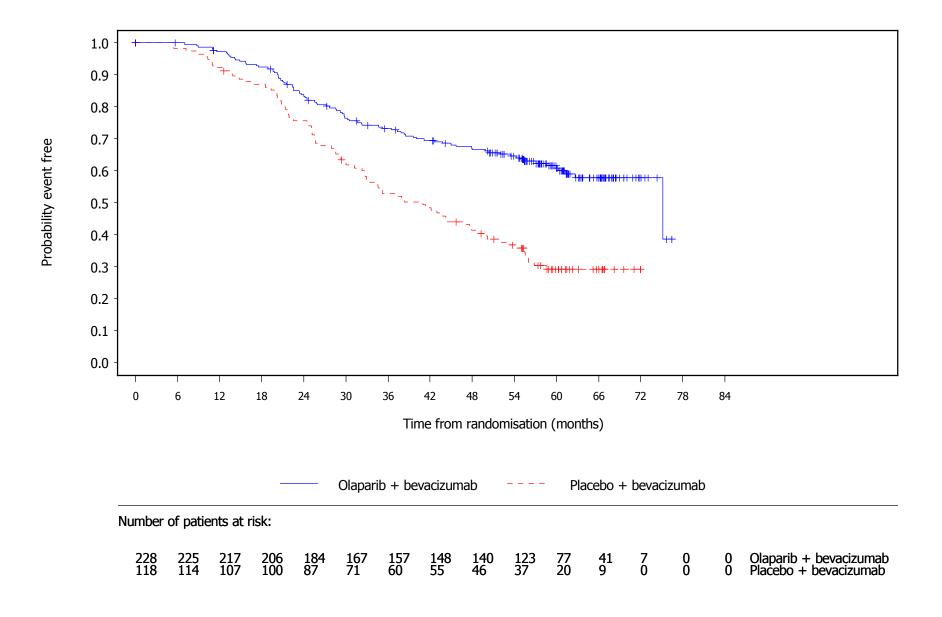


Figure 1.1.4.22 PAOLA1: Kaplan-Meier plot of Second Progression-free survival for Baseline CA-125 value = >ULN Full Analysis Set, HRD[42] positive, DCO 22Mar2022

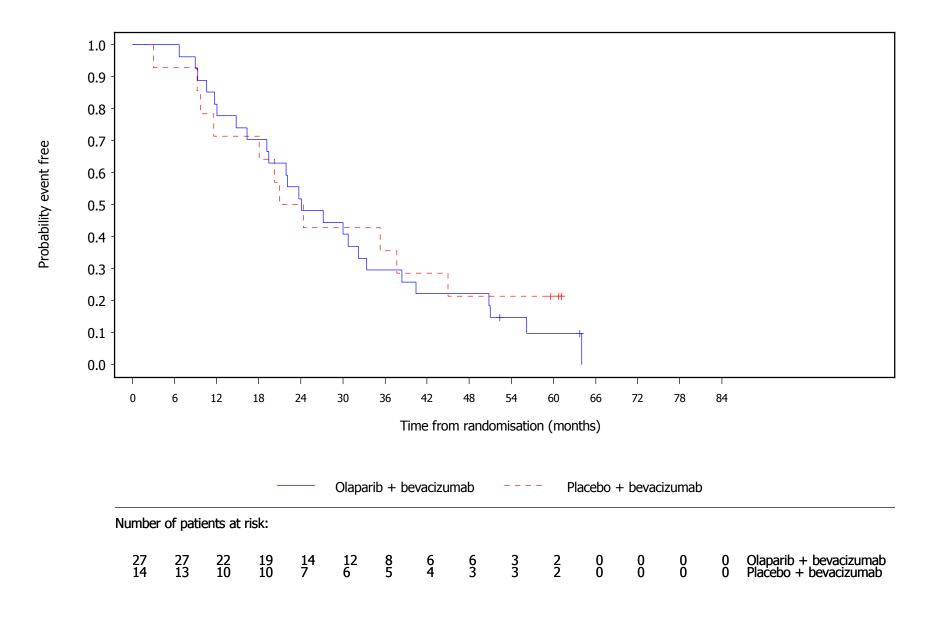


Figure 1.1.4.23 PAOLA1: Kaplan-Meier plot of Time to First Subsequent Cancer Therapy or Death for First line treatment outcome (IVRS) = NED [PDS]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

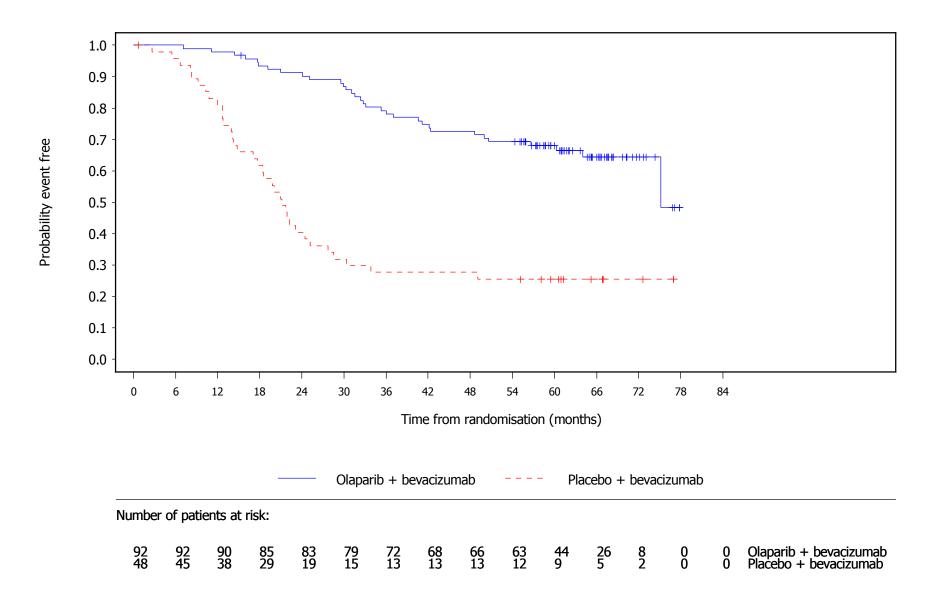


Figure 1.1.4.24 PAOLA1: Kaplan-Meier plot of Time to First Subsequent Cancer Therapy or Death for First line treatment outcome (IVRS) = NED/CR [IDS]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

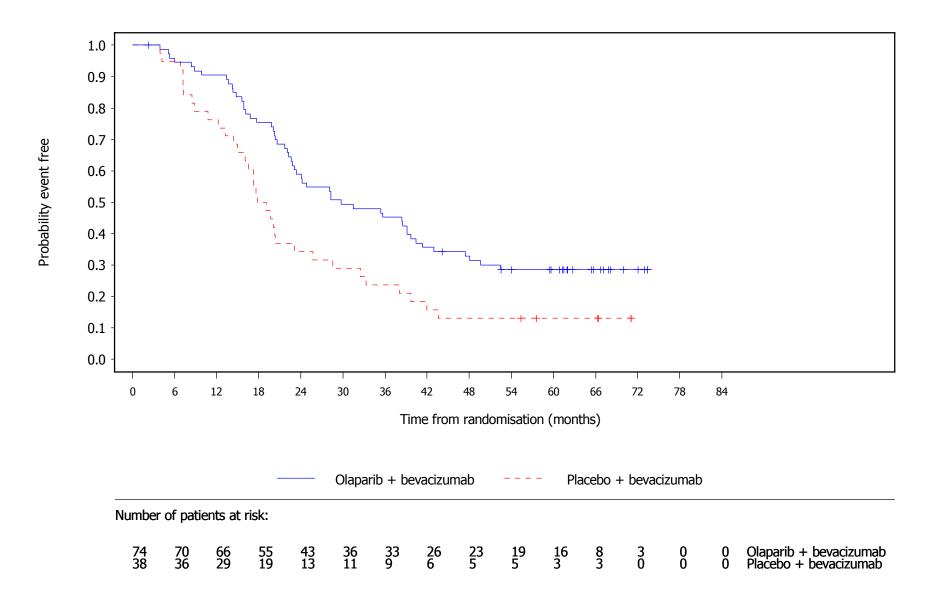


Figure 1.1.4.25 PAOLA1: Kaplan-Meier plot of Time to First Subsequent Cancer Therapy or Death for First line treatment outcome (IVRS) = NED/CR [Chemo]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

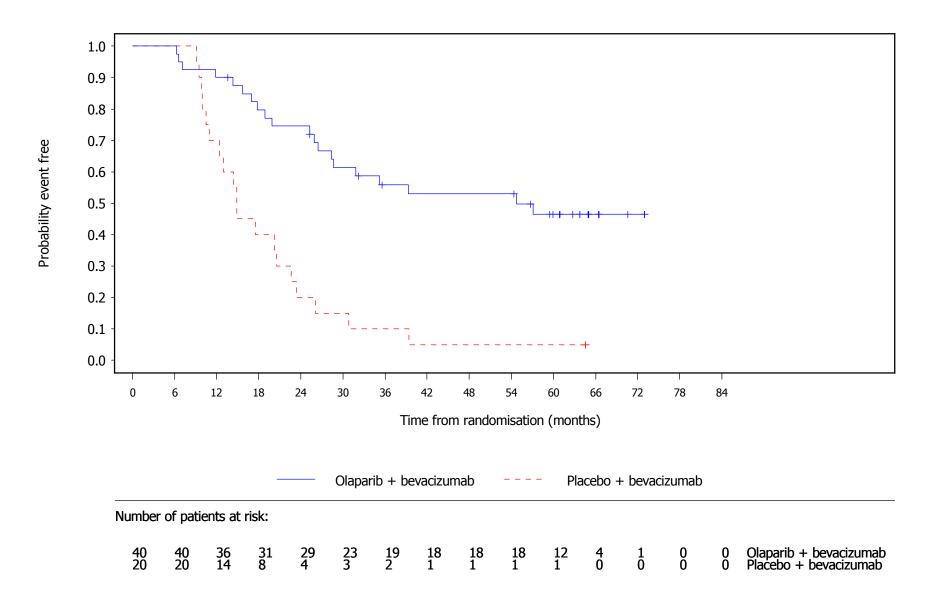


Figure 1.1.4.26 PAOLA1: Kaplan-Meier plot of Time to First Subsequent Cancer Therapy or Death for First line treatment outcome (IVRS) = PR

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

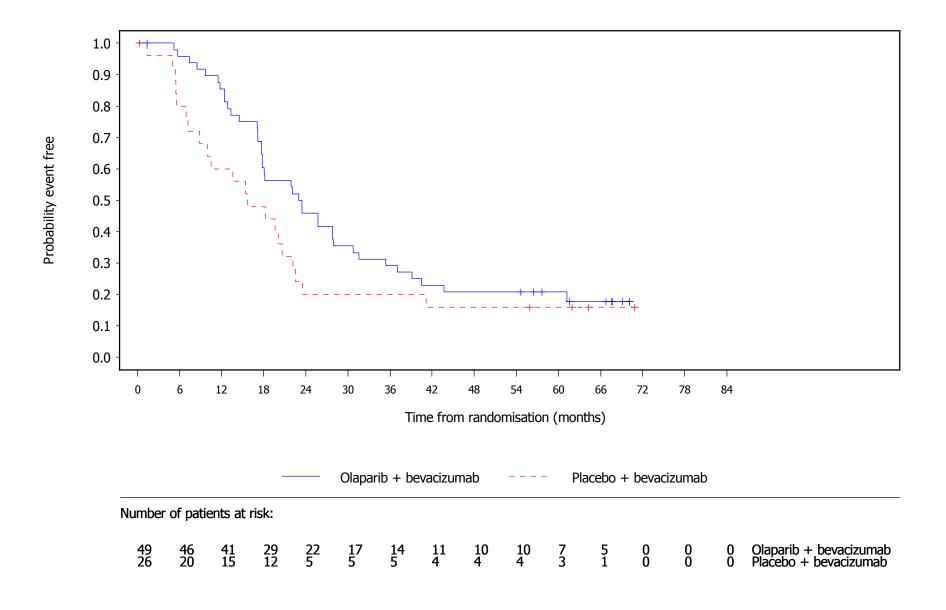


Figure 1.1.4.27 PAOLA1: Kaplan-Meier plot of Time to First Subsequent Cancer Therapy or Death for First line treatment outcome (eCRF) = NED [PDS]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

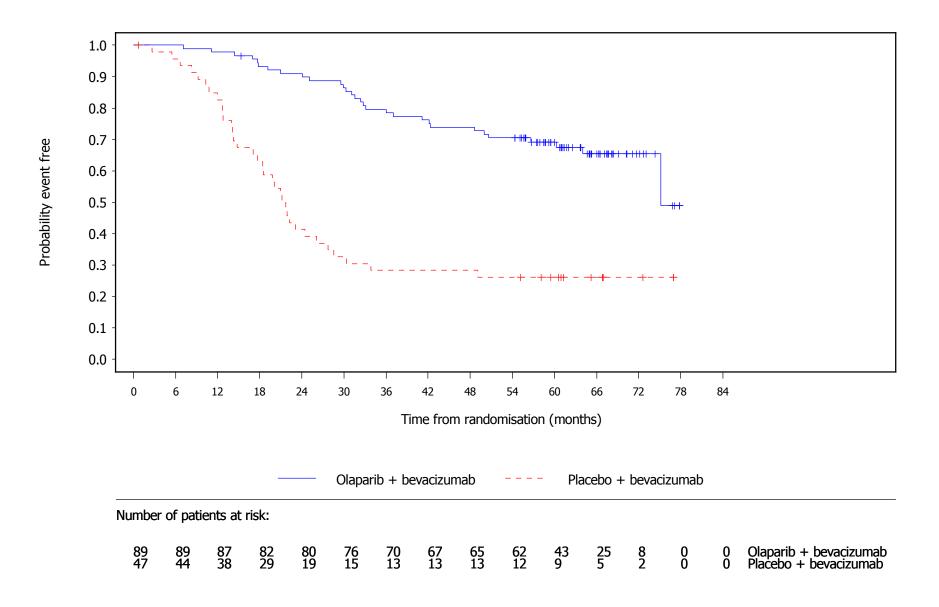


Figure 1.1.4.28 PAOLA1: Kaplan-Meier plot of Time to First Subsequent Cancer Therapy or Death for First line treatment outcome (eCRF) = NED/CR [IDS]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

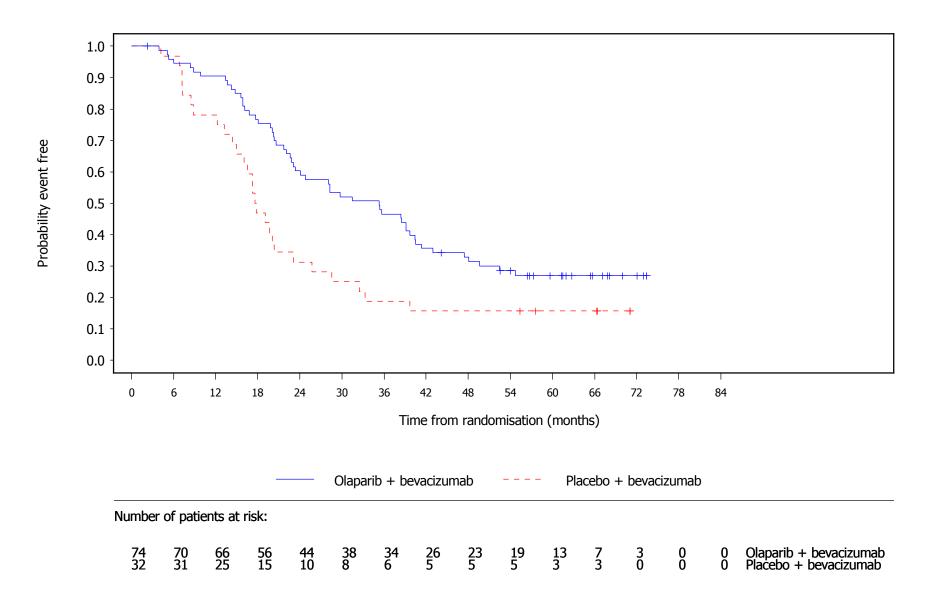


Figure 1.1.4.29 PAOLA1: Kaplan-Meier plot of Time to First Subsequent Cancer Therapy or Death for First line treatment outcome (eCRF) = NED/CR [Chemo]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

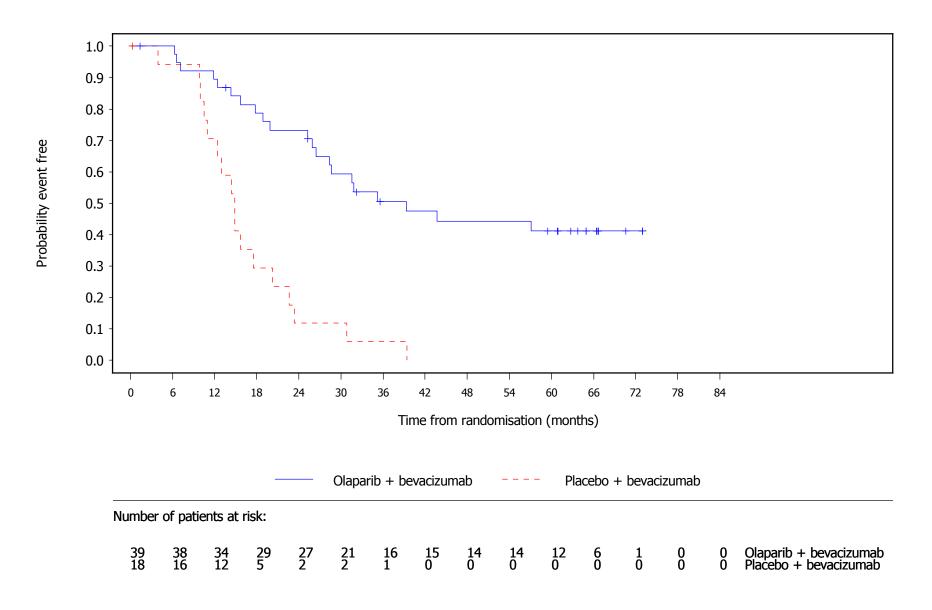


Figure 1.1.4.30 PAOLA1: Kaplan-Meier plot of Time to First Subsequent Cancer Therapy or Death for First line treatment outcome (eCRF) = PR

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

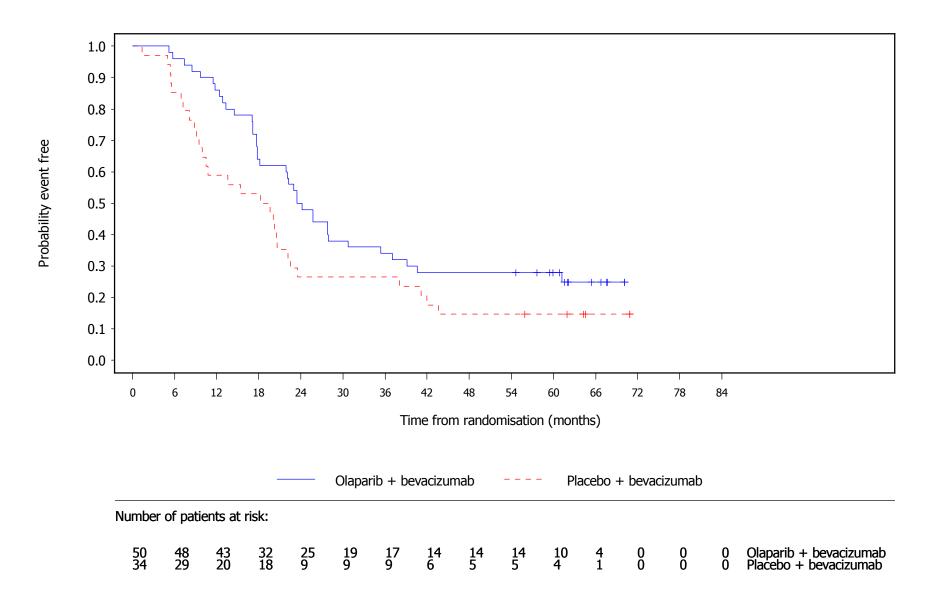


Figure 1.1.4.31 PAOLA1: Kaplan-Meier plot of Time to First Subsequent Cancer Therapy or Death for Baseline CA-125 value = <=ULN Full Analysis Set, HRD[42] positive, DCO 22Mar2022

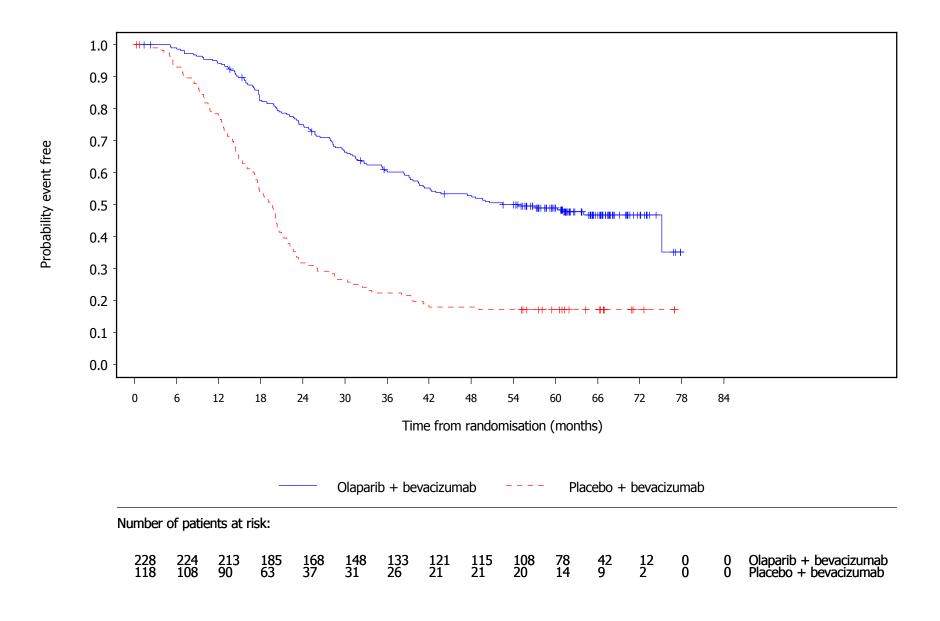


Figure 1.1.4.32 PAOLA1: Kaplan-Meier plot of Time to First Subsequent Cancer Therapy or Death for Baseline CA-125 value = >ULN Full Analysis Set, HRD[42] positive, DCO 22Mar2022

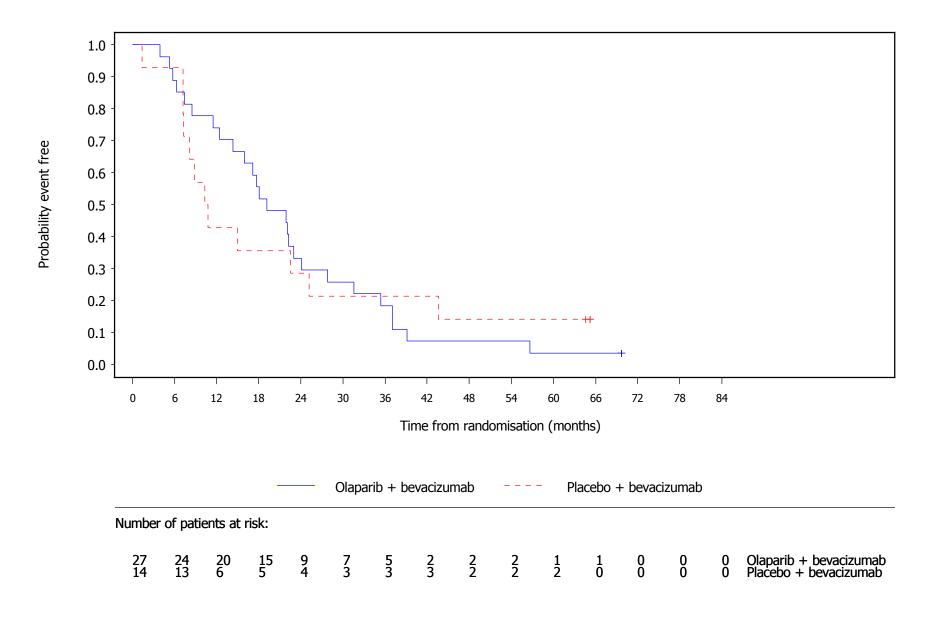


Figure 1.1.4.33 PAOLA1: Kaplan-Meier plot of Time to Second Subsequent Cancer Therapy or Death for First line treatment outcome (IVRS) = NED [PDS]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

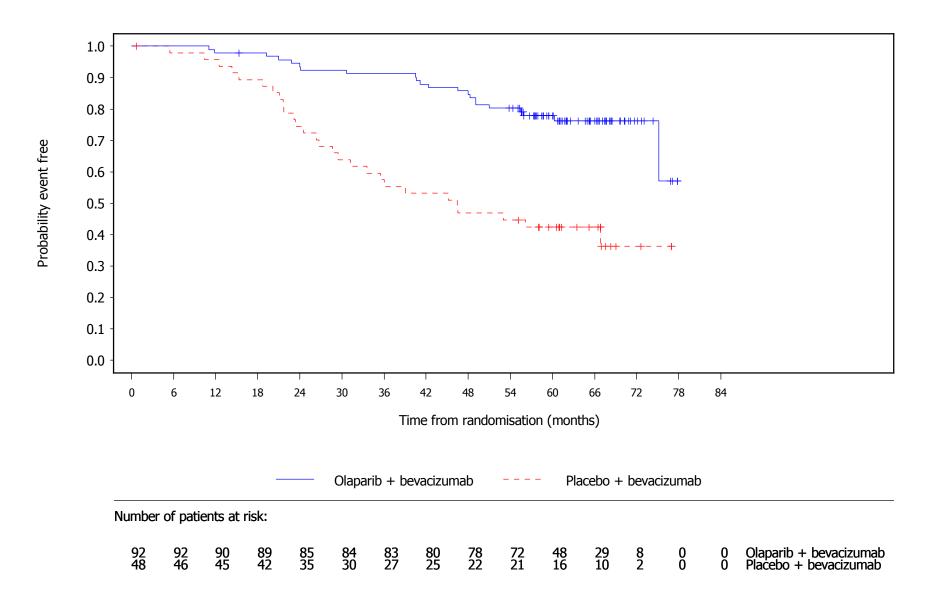


Figure 1.1.4.34 PAOLA1: Kaplan-Meier plot of Time to Second Subsequent Cancer Therapy or Death for First line treatment outcome (IVRS) = NED/CR [IDS]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

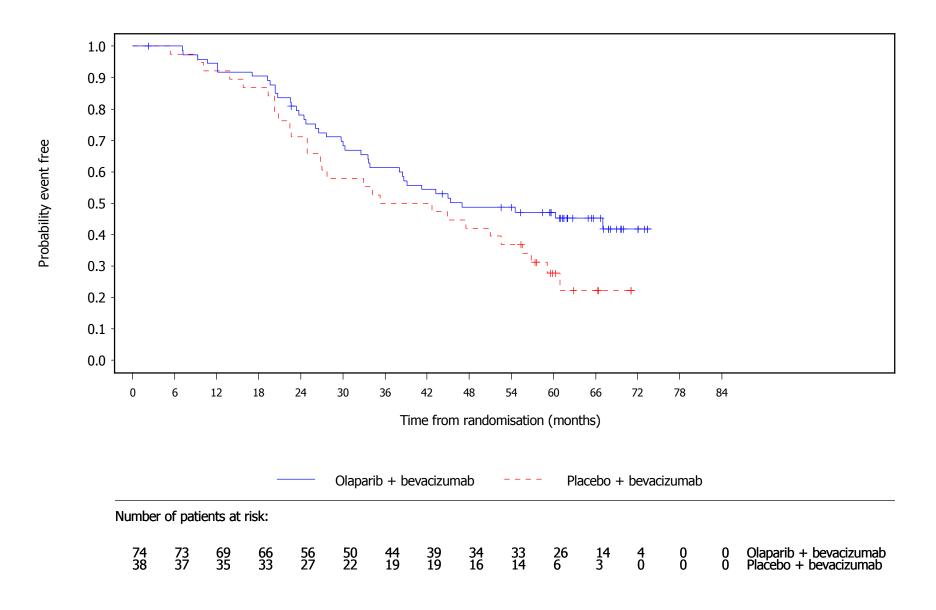


Figure 1.1.4.35 PAOLA1: Kaplan-Meier plot of Time to Second Subsequent Cancer Therapy or Death for First line treatment outcome (IVRS) = NED/CR [Chemo]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

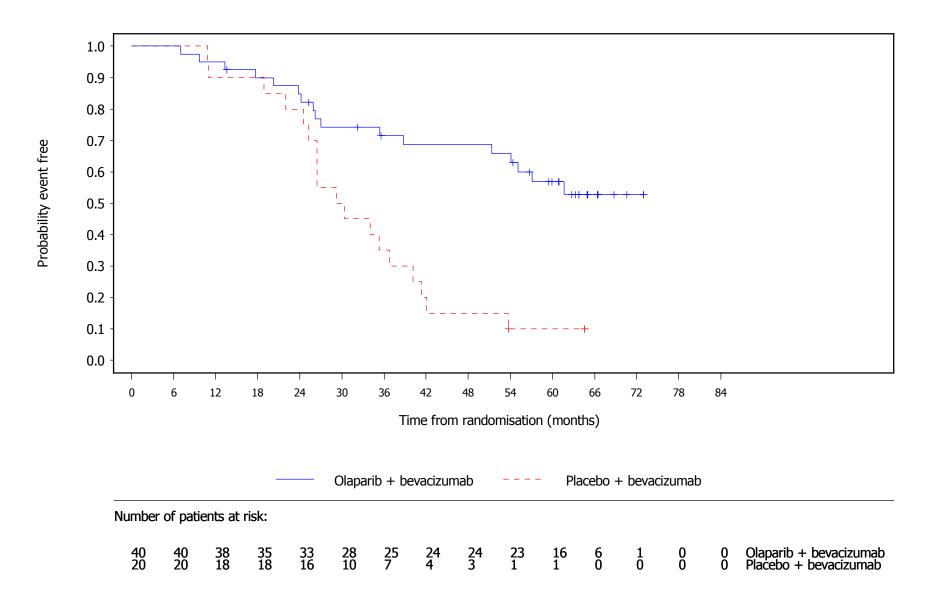


Figure 1.1.4.36 PAOLA1: Kaplan-Meier plot of Time to Second Subsequent Cancer Therapy or Death for First line treatment outcome (IVRS) = PR

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

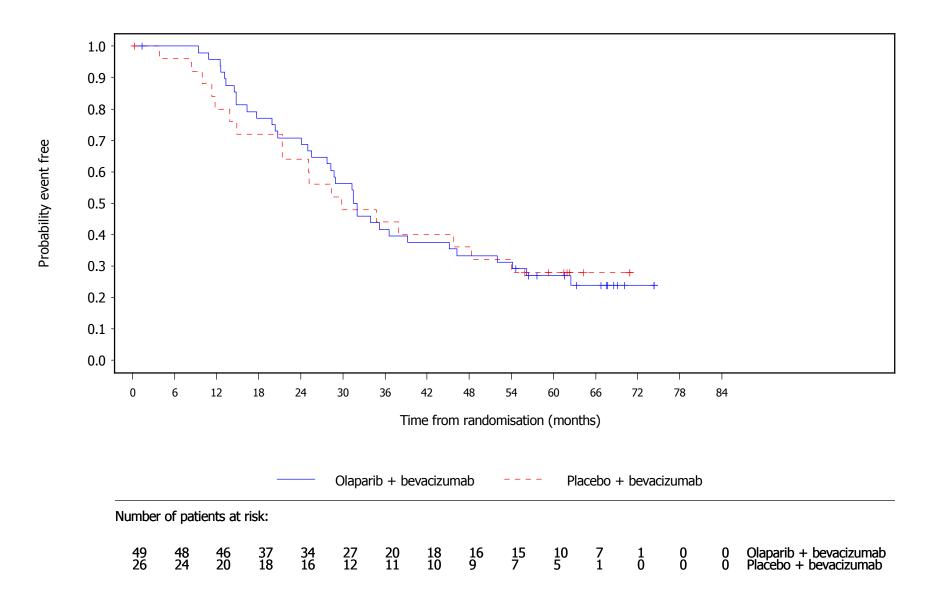


Figure 1.1.4.37 PAOLA1: Kaplan-Meier plot of Time to Second Subsequent Cancer Therapy or Death for First line treatment outcome (eCRF) = NED [PDS]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

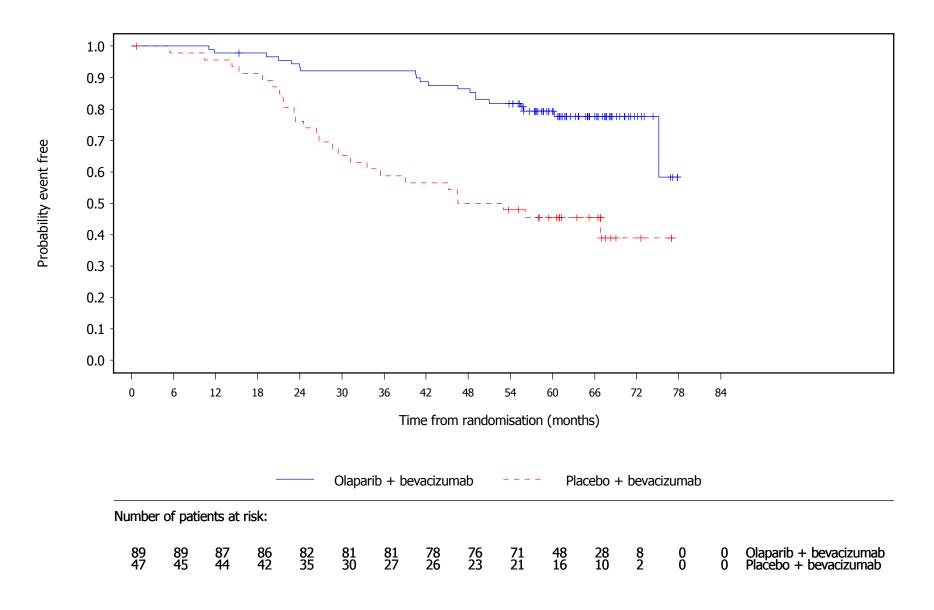


Figure 1.1.4.38 PAOLA1: Kaplan-Meier plot of Time to Second Subsequent Cancer Therapy or Death for First line treatment outcome (eCRF) = NED/CR [IDS]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

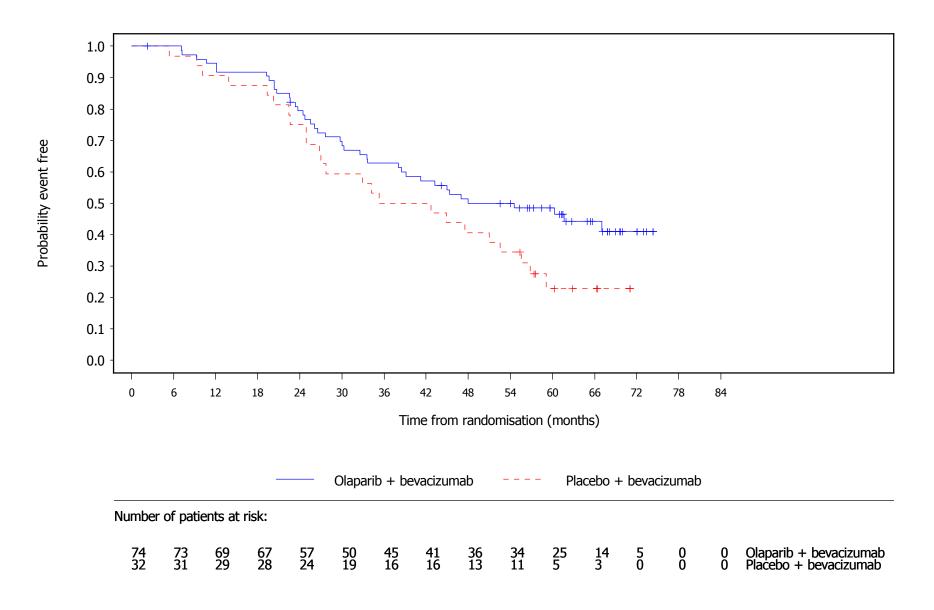


Figure 1.1.4.39 PAOLA1: Kaplan-Meier plot of Time to Second Subsequent Cancer Therapy or Death for First line treatment outcome (eCRF) = NED/CR [Chemo]

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

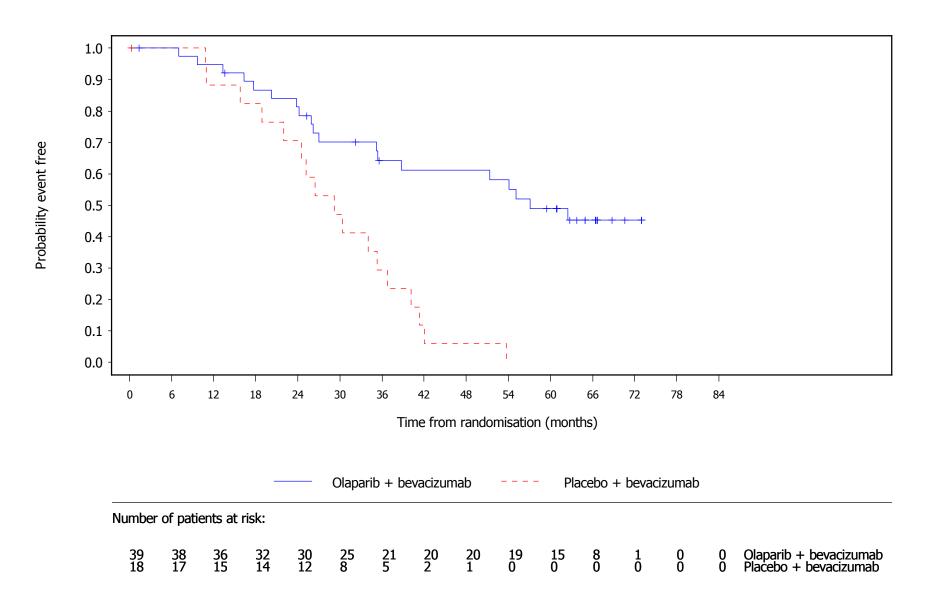


Figure 1.1.4.40 PAOLA1: Kaplan-Meier plot of Time to Second Subsequent Cancer Therapy or Death for First line treatment outcome (eCRF) = PR

Full Analysis Set, HRD[42] positive, DCO 22Mar2022

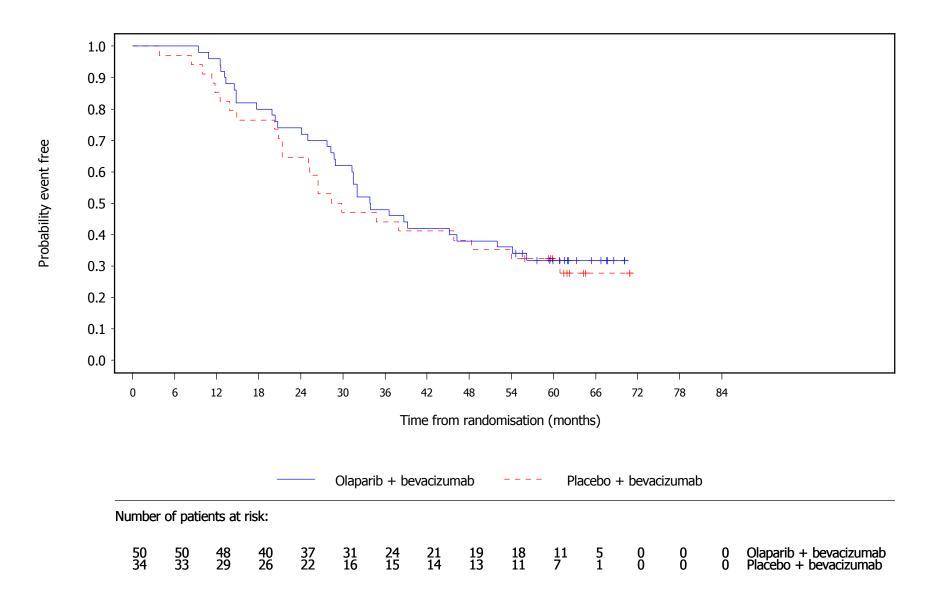


Figure 1.1.4.41 PAOLA1: Kaplan-Meier plot of Time to Second Subsequent Cancer Therapy or Death for Baseline CA-125 value = <=ULN Full Analysis Set, HRD[42] positive, DCO 22Mar2022

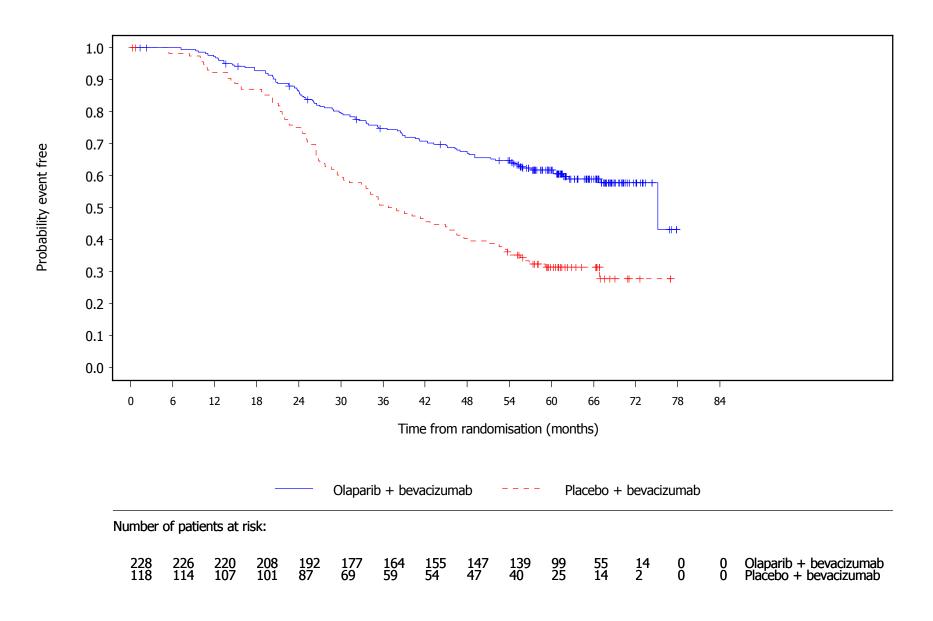


Figure 1.1.4.42 PAOLA1: Kaplan-Meier plot of Time to Second Subsequent Cancer Therapy or Death for Baseline CA-125 value = >ULN Full Analysis Set, HRD[42] positive, DCO 22Mar2022

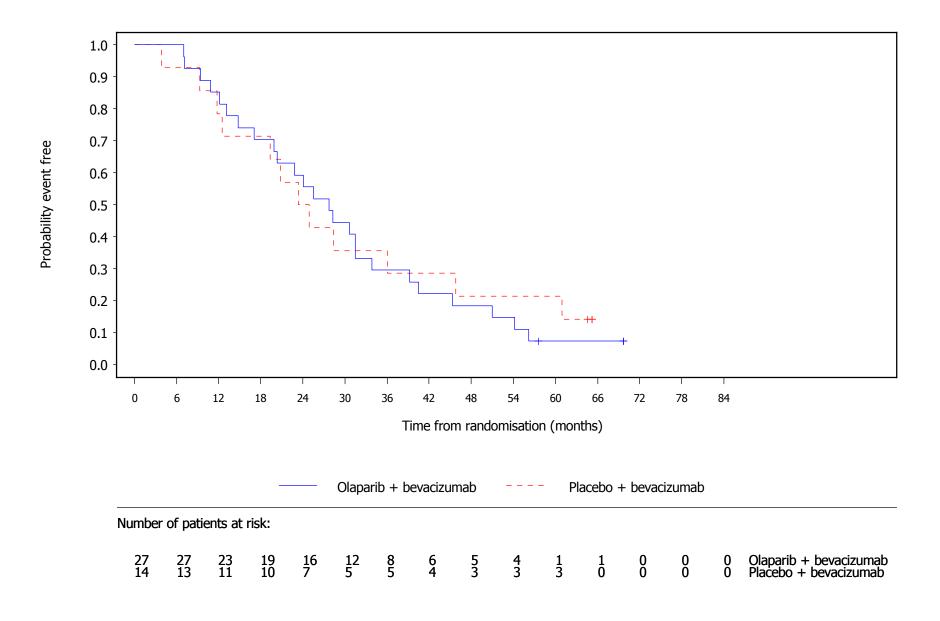


Figure 1.1.4.43 PAOLA1: Kaplan-Meier plot of Overall Survival for Baseline CA-125 value = <=ULN Full Analysis Set, HRD[42] positive, DCO 22Mar2022

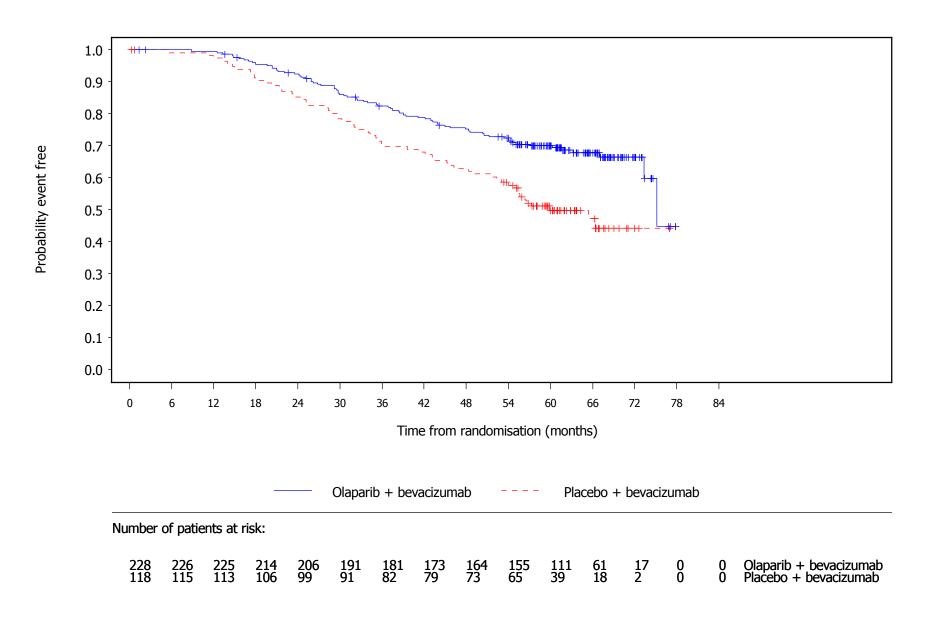


Figure 1.1.4.44 PAOLA1: Kaplan-Meier plot of Overall Survival for Baseline CA-125 value = >ULN Full Analysis Set, HRD[42] positive, DCO 22Mar2022

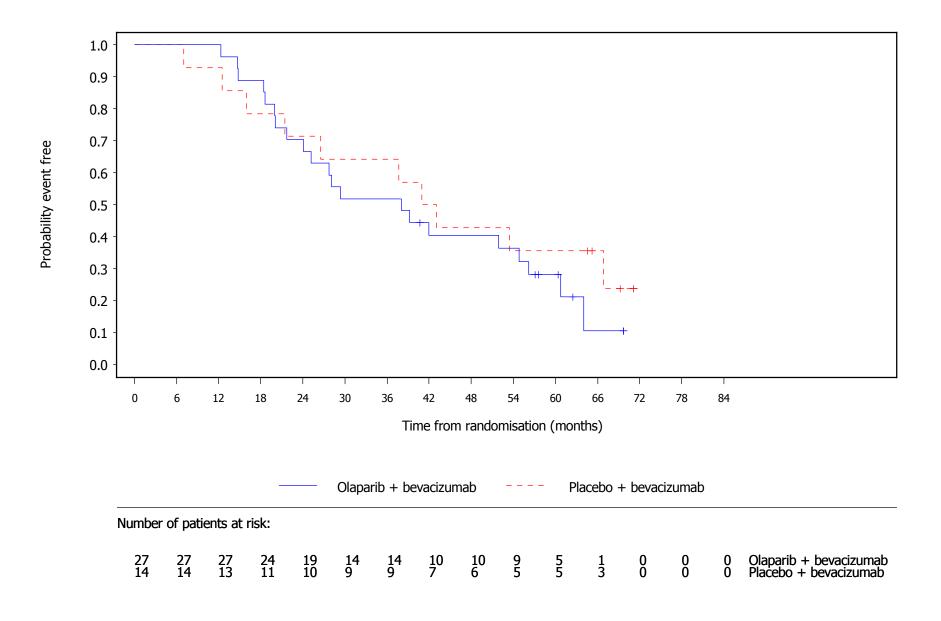


Table 1.2.1.1 PAOLA1: Summary of time to recurrence or death free survival NED and CR only, HRD[42] positive, DCO 22Mar2022

	Olaparib + be (N=20		Placebo + 1				
	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
Time to recurrence or death	206 97 (47.1) 6	55.7 (45.7, NE)	106 84 (79.2)	18.7 (15.8,22.1)	0.36	0.27, 0.49	<0.0001*

Only patients with NED and complete response (Stratification groups 1, 2 and 3).

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05.

Figure 1.2.2.1 PAOLA1: Kaplan-Meier plot of recurrence or death free survival NED and CR only, HRD[42] positive, DCO 22Mar2022

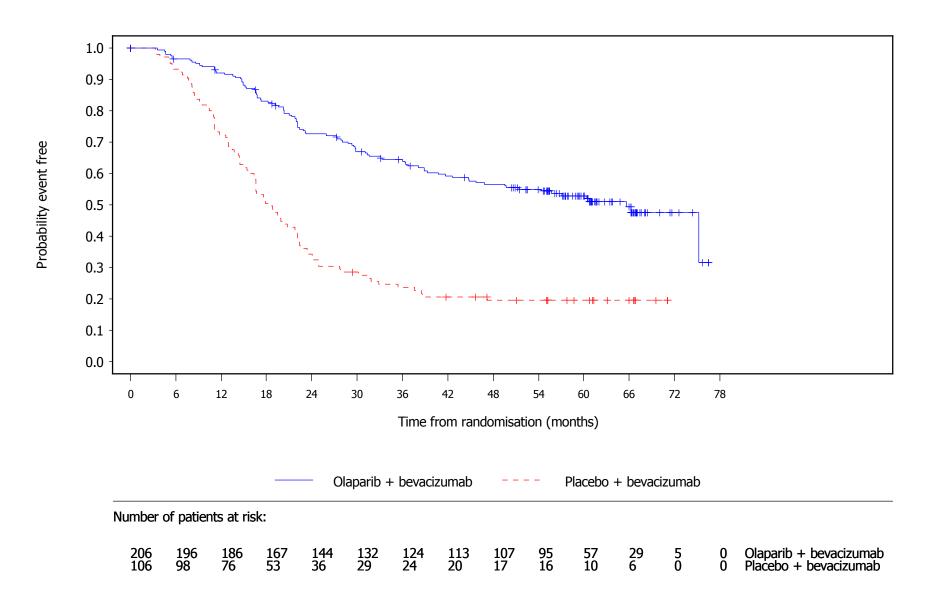


Table 1.2.3.1 PAOLA1: Summary of recurrence or death free rate (odds ratio, relative risk and risk difference) - NED and CR only Full Analysis Set, HRD[42] positive, DCO 22Mar2022

					Treatment effect										
	Olaparib + bevacizumab (N=206)	Placebo + bevacizumab (N=106)	Odds Ratio			Relative Risk		R:	Risk Difference						
	n	Number (%) of patients with events	n	Number (%) of patients with events		stimat 95% CI		2- sided p- value		stimat 95% CI		2- sided p- value		stimate 95% CI)	2- sided p- value
Rate of recurrence or death [a][d][g]	206	97(47.1)	106	84(79.2)	0.23(0.13,	0.40)	<0.0001	0.59(0.50,	0.71)	<0.0001	-0.32(-0).42, -0.21)	<0.0001

Only patients with NED and complete response (Stratification groups 1, 2 and 3 based on IVRS).

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [b] as [a] but with Firth method. [c] OR NC via [a] or [b].

[d] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [e] RR, 95% CI, p-value via modified poisson regression. [f] RR NC via [d] or [e]. [g] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [h] RD NC via [g].

Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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Table 3.1 PAOLA1: Summary of observation period (months) for adverse events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + bevacizumab (N=255)	Placebo + bevacizumab (N=131)		
All AESI endpoints	n	255	131		
	Median	58.74	55.23		
	Min	1.2	0.7		
	Max	77.8	76.9		

Table 3.2.1 PAOLA1: Summary of analysis of time to first occurrence of adverse events of special interest Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		+ bevacizumab N=255)		bevacizumab 131)			
	Number (of patie n with eve	nts (95% CI)	Number (%) of patients n with events		Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AESI: Anaemia	255 102 (40	0) NE (NE, NE)	131 12 (9.2)	NE (NE, NE)	5.62	3.21, 10.80	<0.0001*
AESI: Neutropenia	255 53 (20	8) NE (NE, NE)	131 22 (16.8)	NE (NE, NE)	1.28	0.79, 2.16	0.3420
AESI: Thrombocytopenia	255 18 (7	1) NE (NE, NE)	131 7 (5.3)	NE (NE, NE)	1.29	0.56, 3.36	0.5650
AESI: Nausea	255 144 (56	5) 2.9 (0.8,16.0)	131 34 (26.0)	NE (NE, NE)	2.99	2.07, 4.44	<0.0001*
AESI: Vomiting	255 55 (21	6) NE (NE, NE)	131 18 (13.7)	NE (NE, NE)	1.60	0.96, 2.81	0.0820
AESI: Fatigue and Asthenia	255 142 (55	7) 11.0 (4.1,30.1)	131 47 (35.9)	NE (NE, NE)	1.89	1.36, 2.66	0.0002*
AESI: Hypertension	255 127 (49	8) 30.5 (9.7, NE)	131 78 (59.5)	5.5 (3.4,11.3)	0.74	0.56, 0.99	0.0424*
AESI: Proteinuria	255 20 (7	8) NE (NE, NE)	131 19 (14.5)	NE (NE, NE)	0.48	0.25, 0.91	0.0202*
AESI: GI perforations, abscesses and fistulae	255 3 (1	2) NE (NE, NE)	131 0	NE (NE, NE)	NC	NC	0.2141
AESI: Wound healing complications	255 2 (0	8) NE (NE, NE)	131 3 (2.3)	NE (NE, NE)	0.31	0.04, 1.89	0.1790
AESI: Haemorrhage	255 30 (11	8) NE (NE, NE)	131 12 (9.2)	NE (NE, NE)	1.21	0.63, 2.47	0.5786
AESI: Arterial thromboembolic events	255 3 (1	2) NE (NE, NE)	131 4 (3.1)	NE (NE, NE)	0.35	0.07, 1.61	0.1564

The time to event endpoint is the time to first AE of special interest or the time to censoring if the AE of special interest has not occurred prior to the date of DCO. Includes AEs with an onset date on or after the date of first dose.

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.1 PAOLA1: Summary of analysis of time to first occurrence of adverse events of special interest Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	Olaparib + bevacizumab (N=255)					Placebo + be (N=1)				
	C	Number (%) of patients with events	(95%	n time (CI) (S) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	— Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AESI: Venous thromboembolic events	255	10 (3.9)	NE (NE, NE	131	1 (0.8)	NE (NE, 1	E) 5.16	0.98, 94.87	0.0815
AESI: Posterior Reversible Encephalopathy Syndrome (PRES)	255	0	NE (NE, NE	131	1 (0.8)	NE (NE, 1	E) NC	NC	0.1830
AESI: Congestive heart failure	255	0	NE (NE, NE	131	0	NE (NE, 1	E) NC	NC	NC
AESI: Non-GI fistula or abscess	255	0	NE (NE, NE	131	2 (1.5)	NE (NE, 1	E) NC	NC	0.0468*
AESI: MDS/AML	255	4 (1.6)	NE (NE, NE	131	3 (2.3)	NE (NE, 1	E) 0.67	0.14, 3.52	0.6109
AESI: Myelodysplastic syndrome and Acute myeloid leukaemia	255	4 (1.6)	NE (NE, NE	131	4 (3.1)	NE (NE, 1	E) 0.51	0.12, 2.19	0.3360
AESI: Secondary cancer	255	15 (5.9)	NE (NE, NE	131	4 (3.1)	NE (NE, 1	E) 1.43	0.51, 5.06	0.5272
AESI: Pneumonitis	255	3 (1.2)	NE (NE, NE	131	0	NE (NE, 1	E) NC	NC	0.1935

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.2 PAOLA1: Summary of analysis of time to first occurrence of serious adverse events of special interest Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	Olaparib + bevacizumab (N=255)					Placebo + be					
	(Number (%) of patients with events	Median t: (95% CI (months))		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
Serious AESI: Anaemia	255	13 (5.1)	NE (NE,	NE)	131	1 (0.8)	NE (NE,	NE)	6.96	1.38,126.60	0.0298*
Serious AESI: Neutropenia	255	2 (0.8)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	0.2766
Serious AESI: Thrombocytopenia	255	4 (1.6)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	0.1218
Serious AESI: Vomiting	255	0	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Serious AESI: Hypertension	255	22 (8.6)	NE (NE,	NE)	131	16 (12.2)	NE (NE,	NE)	0.64	0.33, 1.24	0.1667
Serious AESI: Proteinuria	255	1 (0.4)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	0.5101
Serious AESI: GI perforations, abscesses and fistulae	255	2 (0.8)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	0.2971
Serious AESI: Wound healing complications	255	0	NE (NE,	NE)	131	1 (0.8)	NE (NE,	NE)	NC	NC	0.1373
Serious AESI: Haemorrhage	255	2 (0.8)	NE (NE,	NE)	131	1 (0.8)	NE (NE,	NE)	1.09	0.10, 23.54	0.9416
Serious AESI: Arterial thromboembolic events	255	1 (0.4)	NE (NE,	NE)	131	3 (2.3)	NE (NE,	NE)	0.18	0.01, 1.39	0.0916
Serious AESI: Venous thromboembolic events	255	2 (0.8)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	0.3092

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.2 PAOLA1: Summary of analysis of time to first occurrence of serious adverse events of special interest Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	0.	laparib + bo (N=25			Ī	Placebo + be (N=13						
	of	umber (%) patients th events	Median time (95% CI) (months) [0	Number (%) f patients ith events	Median tir (95% CI) (months) [Hazard ratio [b]	95% CI	[b]	2-sided p-value [c]
Serious AESI: Non-GI fistula or abscess	255	0	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC		NC
Serious AESI: MDS/AML	255	3 (1.2)	NE (NE,	NE)	131	3 (2.3)	NE (NE,	NE)	0.46	0.08,	2.53	0.3326
Serious AESI: Myelodysplastic syndrome and Acute myeloid leukaemia	255	3 (1.2)	NE (NE,	NE)	131	4 (3.1)	NE (NE,	NE)	0.35	0.07,	1.62	0.1561
Serious AESI: Secondary cancer	255	15 (5.9)	NE (NE,	NE)	131	4 (3.1)	NE (NE,	NE)	1.43	0.51,	5.06	0.5272
Serious AESI: Pneumonitis	255	2 (0.8)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC		0.2504

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.3 PAOLA1: Summary of analysis of time to first occurrence of severe adverse events of special interest with max. CTCAE grade >=3 including grade 5
Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	Olaparib + bevacizumab (N=255)					Placebo + be					
	(Number (%) of patients with events	(95% C	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]		Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AESI G>=3: Anaemia	255	47 (18.4)	NE (NE	, NE)	131	1 (0.8)	NE (NE,	NE)	27.87	6.09,493.95	<0.0001*
AESI G>=3: Neutropenia	255	21 (8.2)	NE (NE	, NE)	131	4 (3.1)	NE (NE,	NE)	2.86	1.08, 9.85	0.0457*
AESI G>=3: Thrombocytopenia	255	5 (2.0)	NE (NE	, NE)	131	4 (3.1)	NE (NE,	NE)	0.59	0.15, 2.43	0.4338
AESI G>=3: Nausea	255	9 (3.5)	NE (NE	, NE)	131	4 (3.1)	NE (NE,	NE)	1.07	0.35, 4.00	0.9065
AESI G>=3: Vomiting	255	4 (1.6)	NE (NE	, NE)	131	6 (4.6)	NE (NE,	NE)	0.31	0.08, 1.09	0.0553
AESI G>=3: Fatigue and Asthenia	255	17 (6.7)	NE (NE	, NE)	131	3 (2.3)	NE (NE,	NE)	3.06	1.02, 13.17	0.0611
AESI G>=3: Hypertension	255	50 (19.6)	NE (NE	, NE)	131	42 (32.1)	NE (NE,	NE)	0.52	0.34, 0.79	0.0016*
AESI G>=3: Proteinuria	255	3 (1.2)	NE (NE	, NE)	131	0	NE (NE,	NE)	NC	NC	0.2630
AESI G>=3: GI perforations, abscesses and fistulae	255	3 (1.2)	NE (NE	, NE)	131	0	NE (NE,	NE)	NC	NC	0.2141
AESI G>=3: Wound healing complications	255	0	NE (NE	, NE)	131	0	NE (NE,	NE)	NC	NC	NC
AESI G>=3: Haemorrhage	255	2 (0.8)	NE (NE	, NE)	131	1 (0.8)	NE (NE,	NE)	1.09	0.10, 23.54	0.9416

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.3 PAOLA1: Summary of analysis of time to first occurrence of severe adverse events of special interest with max. CTCAE grade >=3 including grade 5
Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	Olaparib + bevacizumab (N=255)					Placebo + be						
		Number (%) of patients with events	(95% C	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]		Hazard ratio [b]	95% CI	[b]	2-sided p-value [c]
AESI G>=3: Arterial thromboembolic events	255	1 (0.4)	NE (NE	, NE)	131	4 (3.1)	NE (NE,	NE)	0.12	0.01,	0.84	0.0271*
AESI G>=3: Venous thromboembolic events	255	3 (1.2)	NE (NE	, NE)	131	0	NE (NE,	NE)	NC	NC		0.1894
AESI G>=3: Congestive heart failure	255	0	NE (NE	, NE)	131	0	NE (NE,	NE)	NC	NC		NC
AESI G>=3: Non-GI fistula or abscess	255	0	NE (NE	, NE)	131	1 (0.8)	NE (NE,	NE)	NC	NC		0.2367
AESI G>=3: MDS/AML	255	4 (1.6)	NE (NE	, NE)	131	3 (2.3)	NE (NE,	NE)	0.67	0.14,	3.52	0.6109
AESI G>=3: Myelodysplastic syndrome and Acute myeloid leukaemia	255	4 (1.6)	NE (NE	, NE)	131	4 (3.1)	NE (NE,	NE)	0.51	0.12,	2.19	0.3360
AESI G>=3: Secondary cancer	255	11 (4.3)	NE (NE	, NE)	131	3 (2.3)	NE (NE,	NE)	1.47	0.45,	6.54	0.5572

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.4 PAOLA1: Summary of analysis of time to first occurrence of non-severe adverse events of special interest with max. CTCAE grade 1 or 2

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	Olaparib + bevacizumab (N=255)					Placebo + bo				
	n	Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [c]
AESI G1-2: Anaemia	255	86 (33.7)	NE (NE,	NE)	131	11 (8.4)	NE (NE, NE)	4.90	2.73, 9.76	<0.0001*
AESI G1-2: Neutropenia	255	39 (15.3)	NE (NE,	NE)	131	19 (14.5)	NE (NE, NE)	1.05	0.61, 1.87	0.8637
AESI G1-2: Thrombocytopenia	255	14 (5.5)	NE (NE,	NE)	131	4 (3.1)	NE (NE, NE)	1.80	0.64, 6.40	0.2958
AESI G1-2: Nausea	255	142 (55.7)	3.4 (1.1,	NE)	131	30 (22.9)	NE (NE, NE)	3.29	2.24, 4.99	<0.0001*
AESI G1-2: Vomiting	255	52 (20.4)	NE (NE,	NE)	131	14 (10.7)	NE (NE, NE)	1.98	1.13, 3.73	0.0213*
AESI G1-2: Fatigue and Asthenia	255	133 (52.2)	13.4 (6.0,	NE)	131	44 (33.6)	NE (NE, NE)	1.82	1.30, 2.59	0.0006*
AESI G1-2: Hypertension	255	98 (38.4)	NE (NE,	NE)	131	59 (45.0)	NE (NE, NE)	0.87	0.63, 1.20	0.3807
AESI G1-2: Proteinuria	255	18 (7.1)	NE (NE,	NE)	131	19 (14.5)	NE (NE, NE)	0.43	0.22, 0.83	0.0092*
AESI G1-2: Wound healing complications	255	2 (0.8)	NE (NE,	NE)	131	3 (2.3)	NE (NE, NE)	0.31	0.04, 1.89	0.1790
AESI G1-2: Haemorrhage	255	28 (11.0)	NE (NE,	NE)	131	11 (8.4)	NE (NE, NE)	1.27	0.65, 2.67	0.5073
AESI G1-2: Arterial thromboembolic events	255	2 (0.8)	NE (NE,	NE)	131	0	NE (NE, NE)	NC	NC	0.3737
AESI G1-2: Venous thromboembolic events	255	8 (3.1)	NE (NE,	NE)	131	1 (0.8)	NE (NE, NE)	4.02	0.73, 74.64	0.1568

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[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Table 3.2.4 PAOLA1: Summary of analysis of time to first occurrence of non-severe adverse events of special interest with max. CTCAE grade 1 or 2

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	Olaparib + bevacizumab (N=255)				Placebo + bo				
	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	n	Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]		2-sided p-value [c]
AESI G1-2: Posterior Reversible Encephalopathy Syndrome (PRES)	255	0	NE (NE, NE) 131	1 (0.8)	NE (NE, NE)	NC	NC	0.1830
AESI G1-2: Non-GI fistula or abscess	255	0	NE (NE, NE) 131	1 (0.8)	NE (NE, NE)	NC	NC	0.0973
AESI G1-2: Secondary cancer	255	3 (1.2)	NE (NE, NE) 131	1 (0.8)	NE (NE, NE)	1.09	0.14, 22.17	0.9432
AESI G1-2: Pneumonitis	255	3 (1.2)	NE (NE, NE) 131	0	NE (NE, NE)	NC	NC	0.1935

root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttemainae.sas ettemainaead 11AUG2022:11:28 kpzx329

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] Estimated from Cox proportional hazards model including treatment only. First line treatment outcome and tBRCA status included as stratification factors. Efron method for handling ties. 95% CI from profile likelihood estimation.

[[]c] Determined using log-rank test stratified by First line treatment outcome and tBRCA status. Breslow method for handling ties.

Hazard ratio <1 favours olaparib. * p<0.05. NC = not calculable.

Figure 3.3.1 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

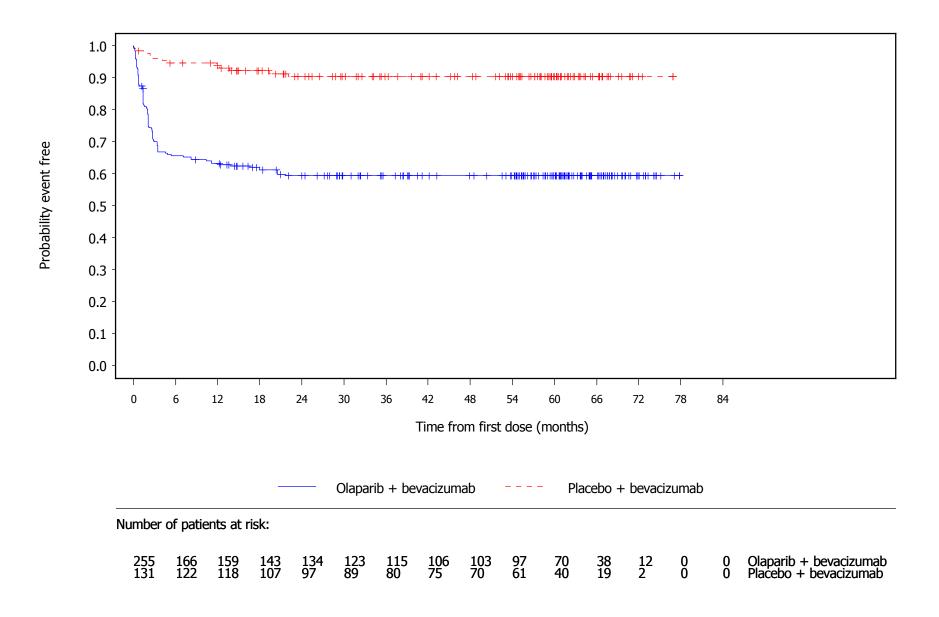


Figure 3.3.2 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

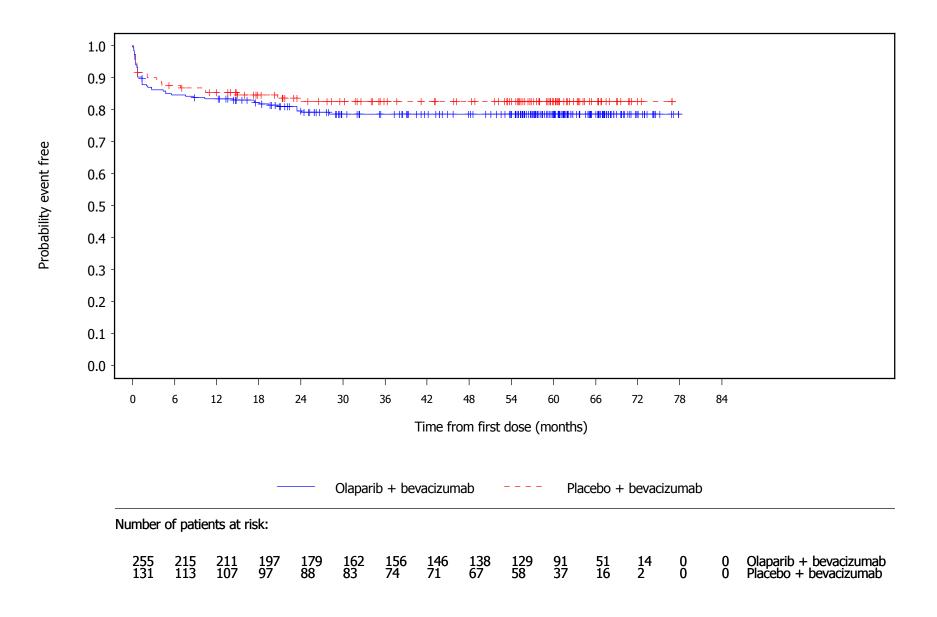


Figure 3.3.3 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

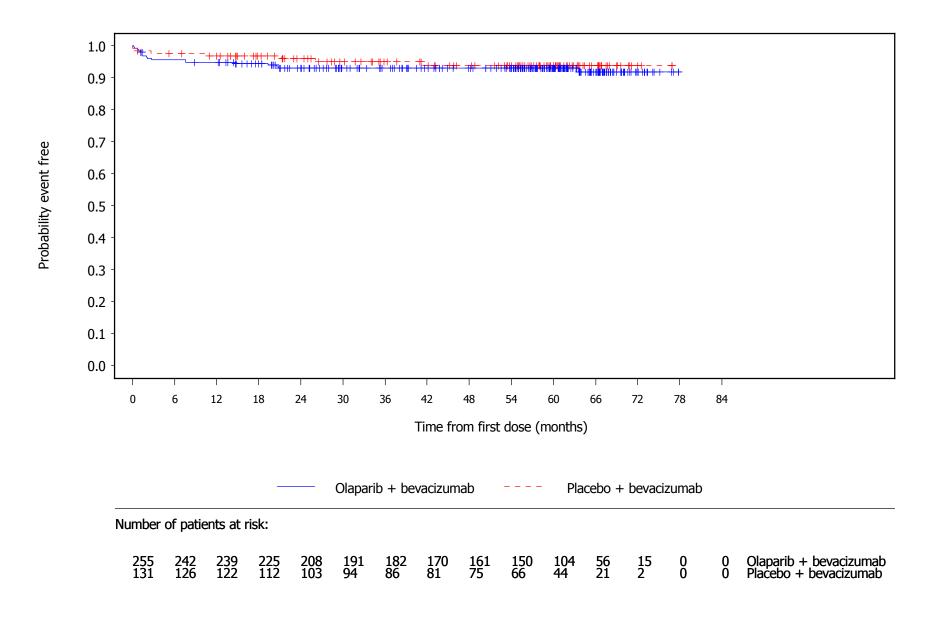


Figure 3.3.4 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Nausea Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

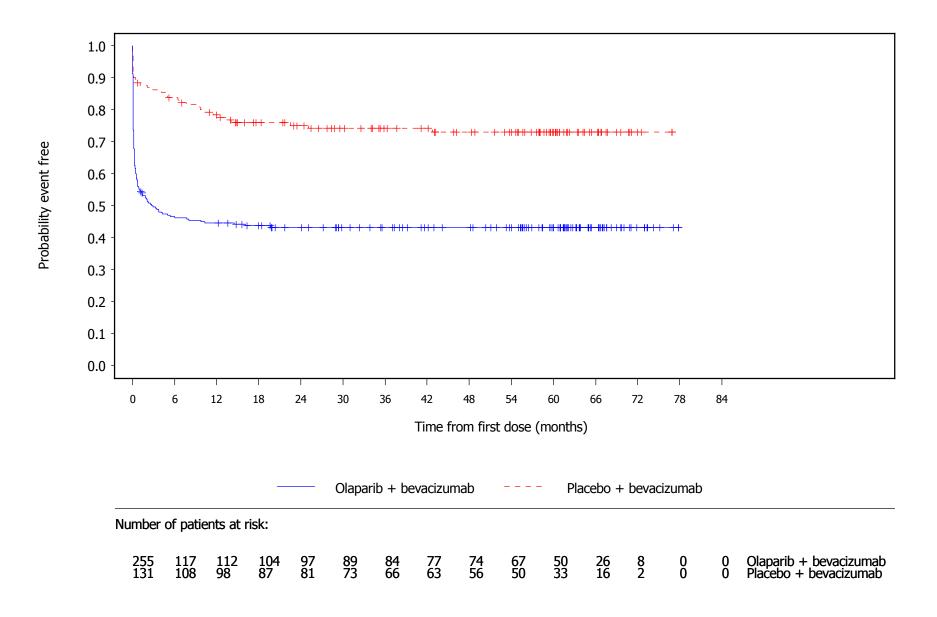


Figure 3.3.5 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

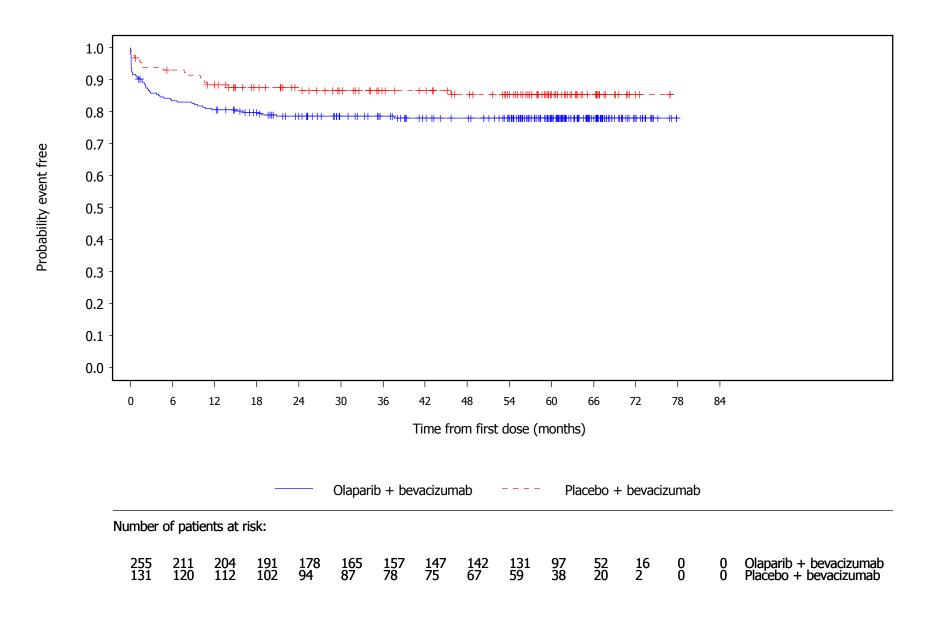


Figure 3.3.6 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Fatigue and Asthenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

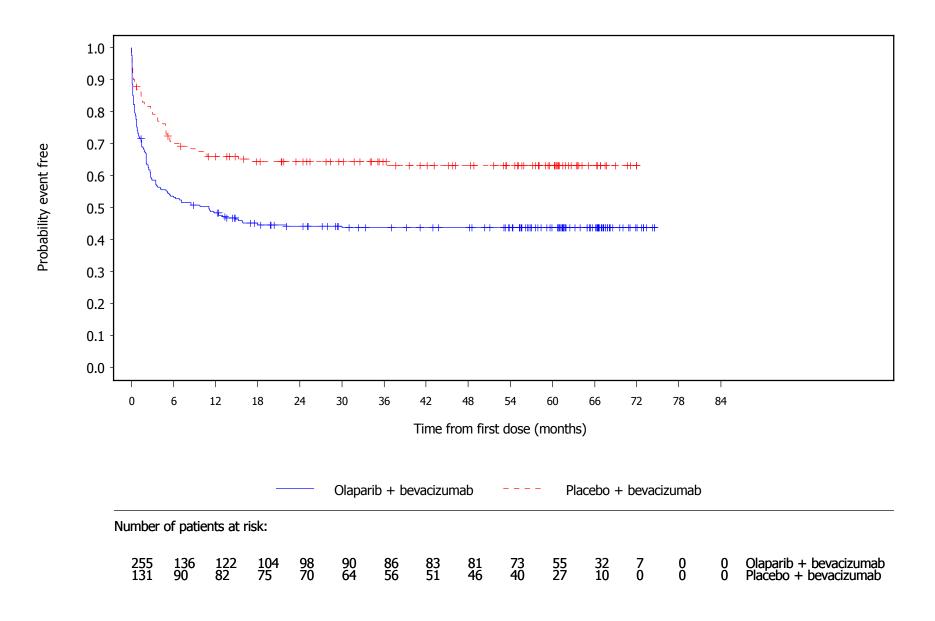


Figure 3.3.7 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

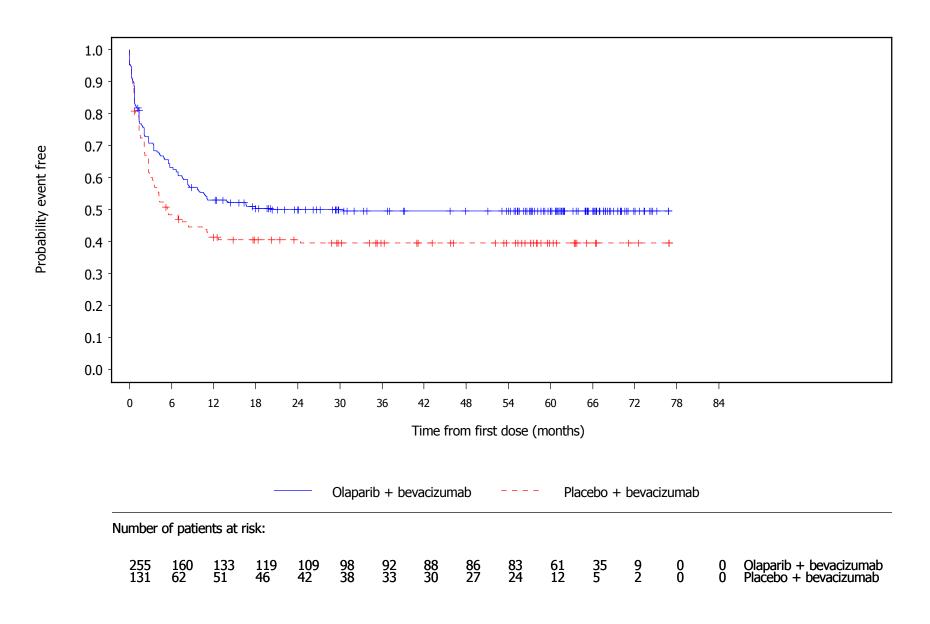


Figure 3.3.8 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

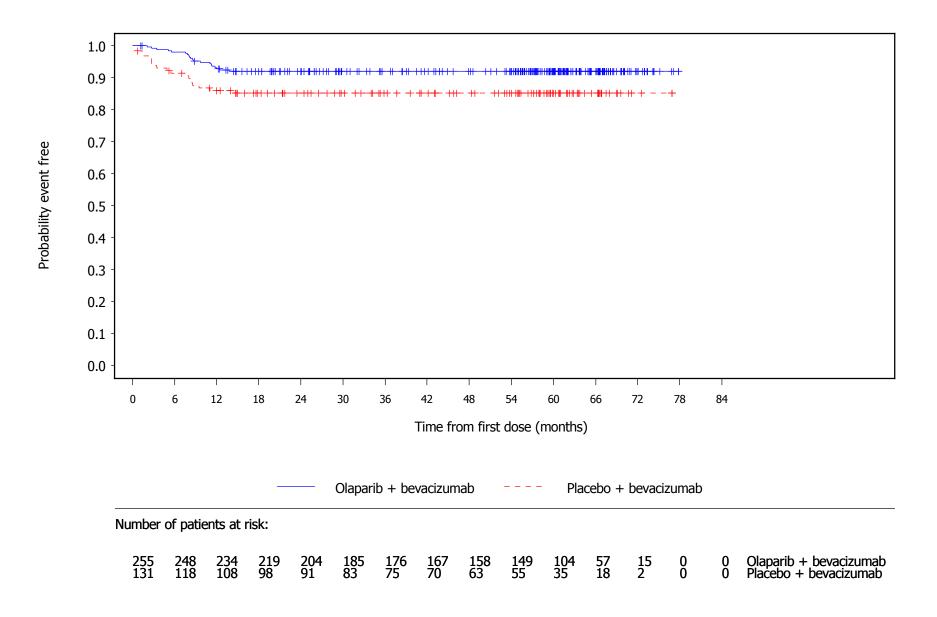


Figure 3.3.9 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: GI perforations, abscesses and fistulae Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

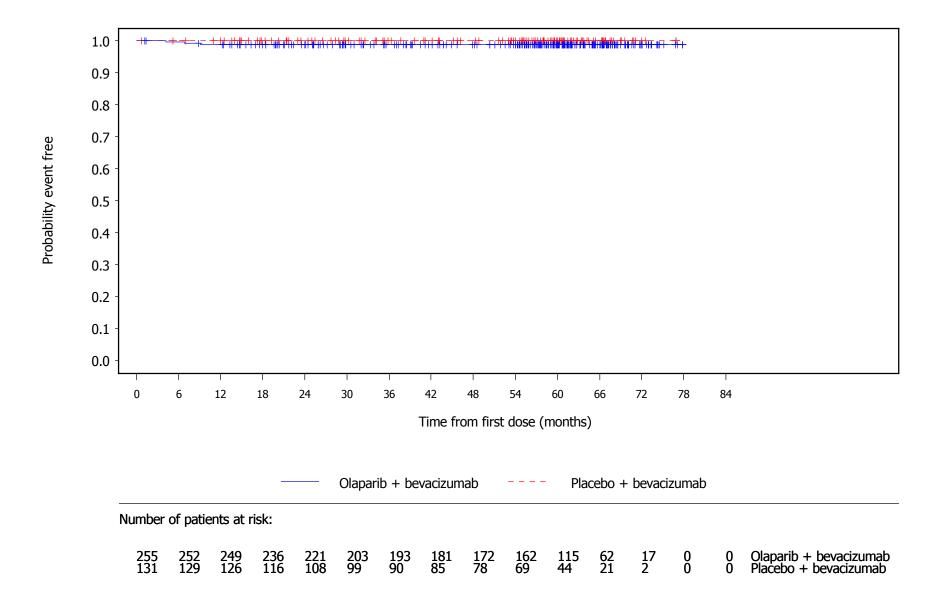


Figure 3.3.10 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

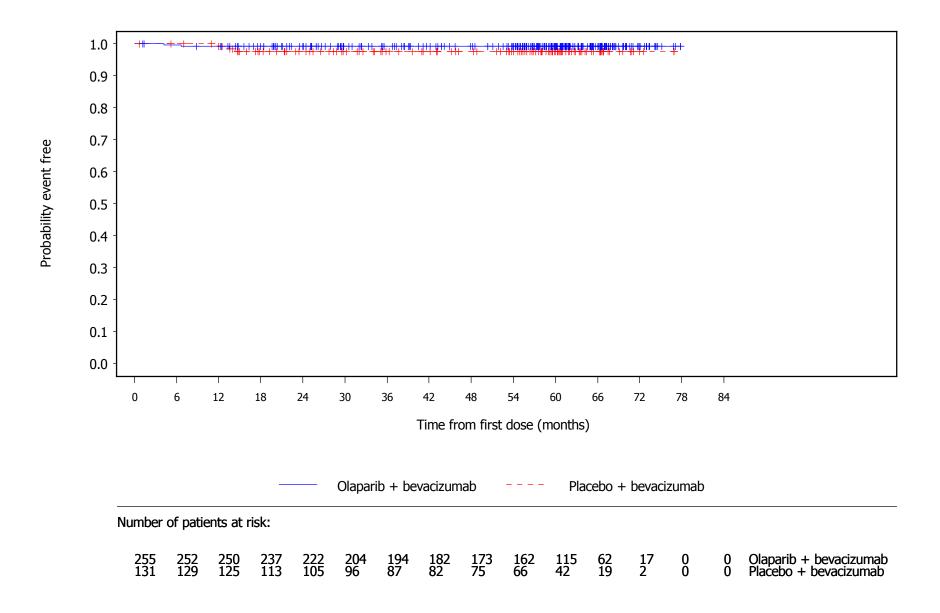


Figure 3.3.11 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

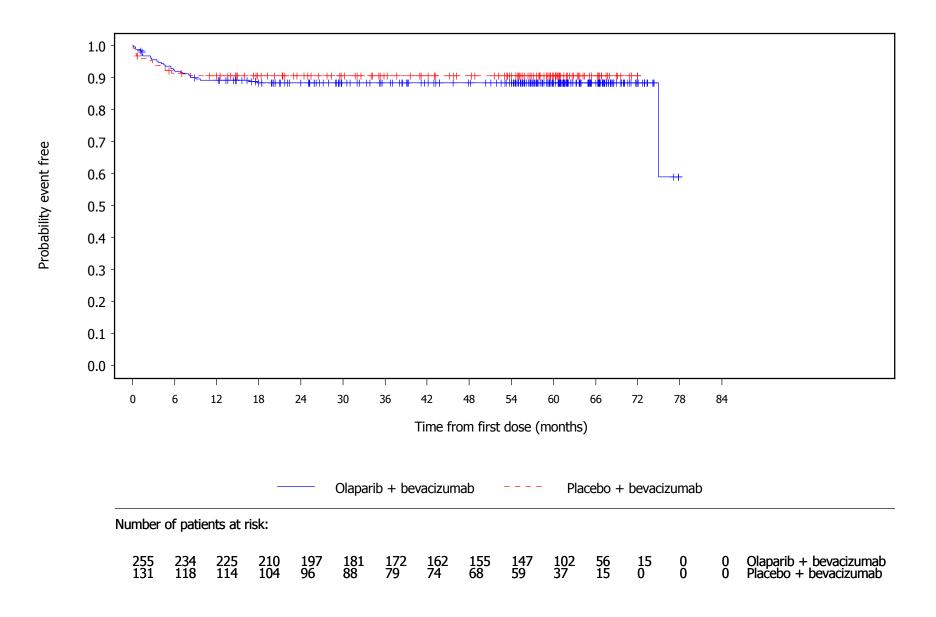


Figure 3.3.12 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

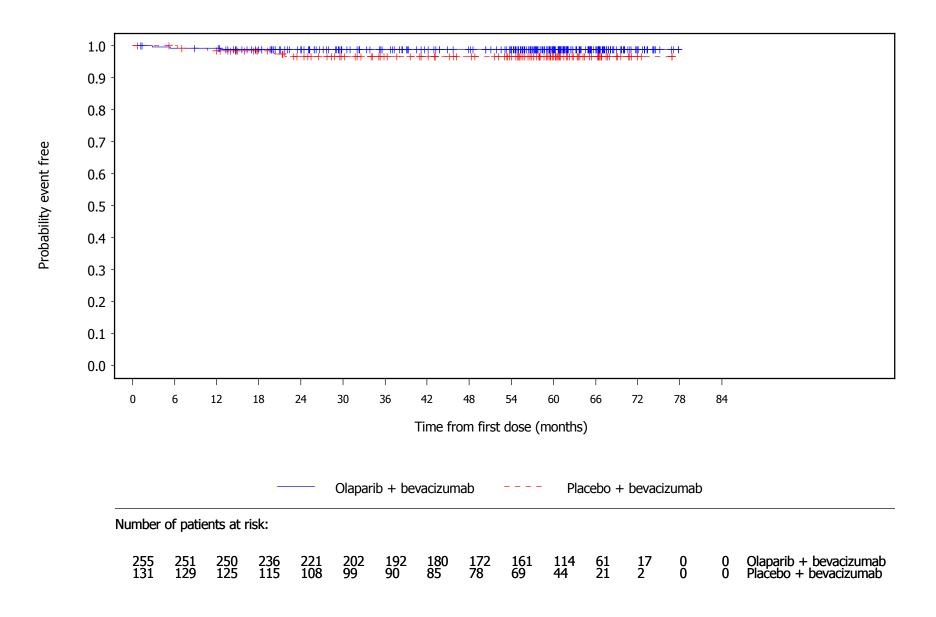


Figure 3.3.13 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

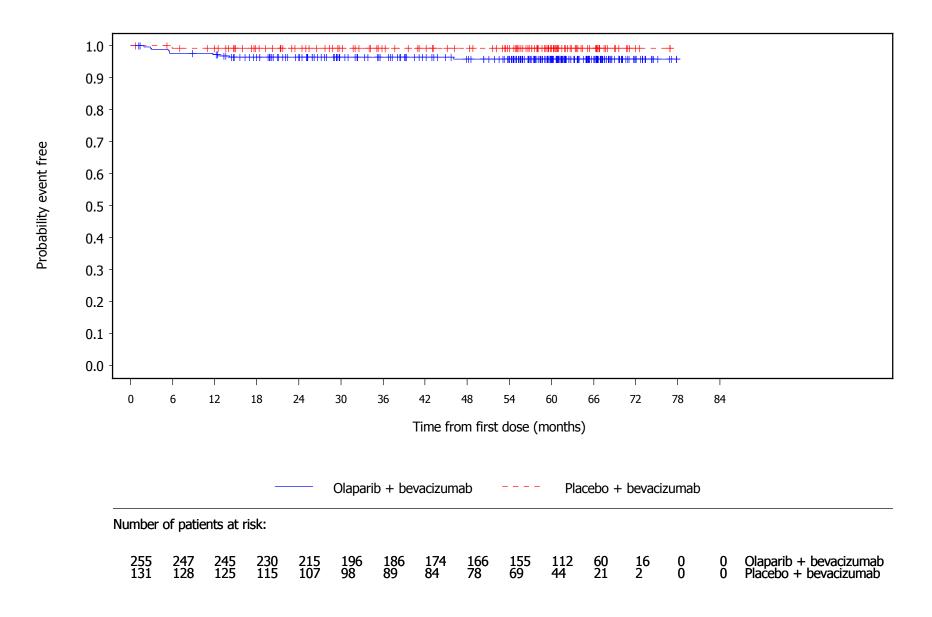


Figure 3.3.14 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Posterior Reversible Encephalopathy Syndrome (PRES)

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

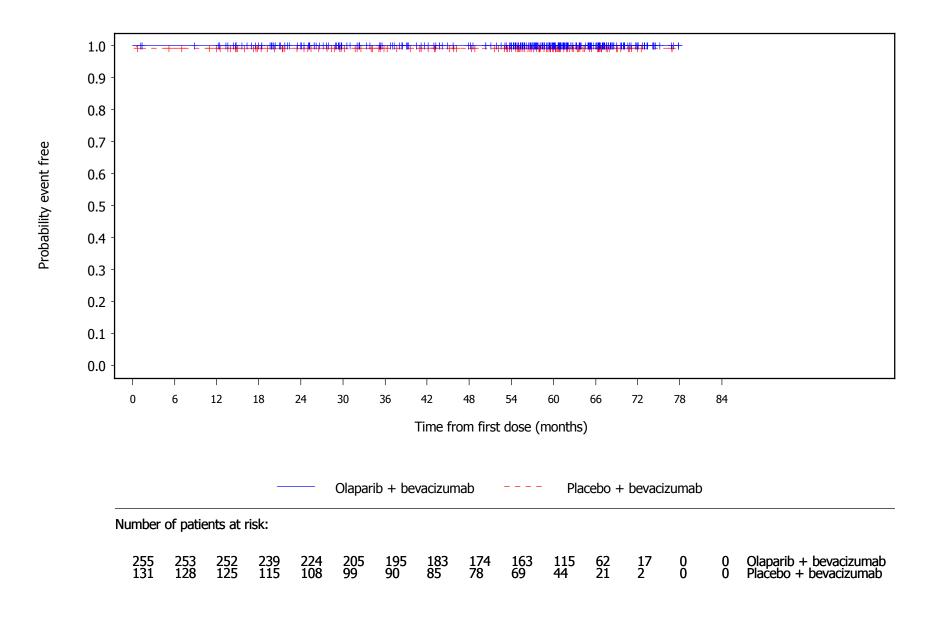


Figure 3.3.15 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Congestive heart failure Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

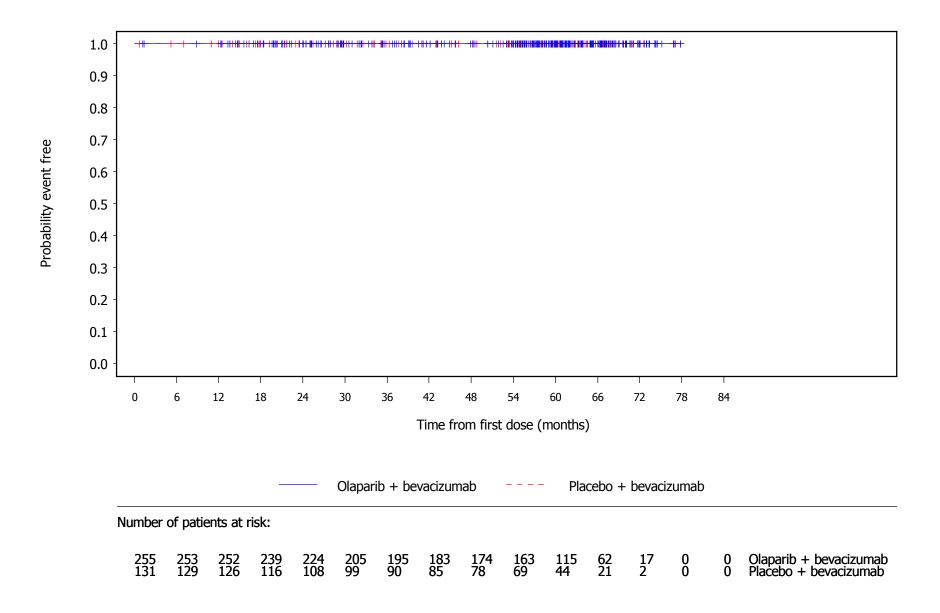


Figure 3.3.16 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

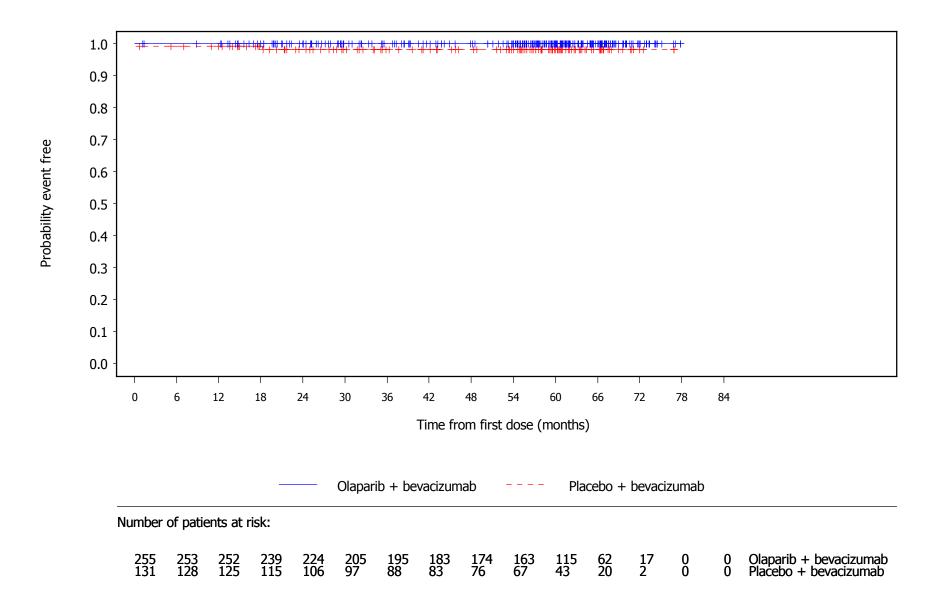


Figure 3.3.17 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: MDS/AML Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

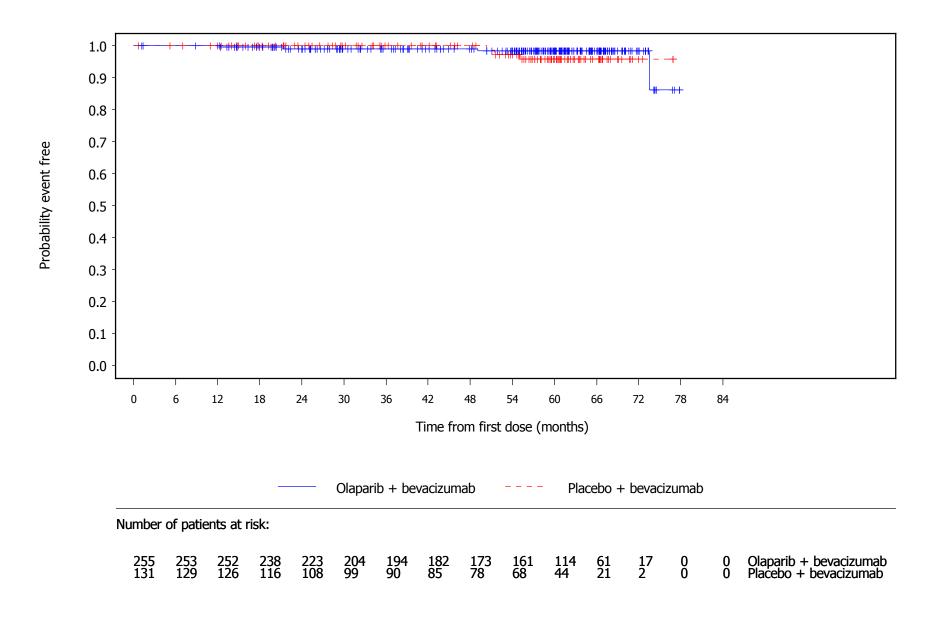


Figure 3.3.18 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Myelodysplastic syndrome and Acute myeloid leukaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

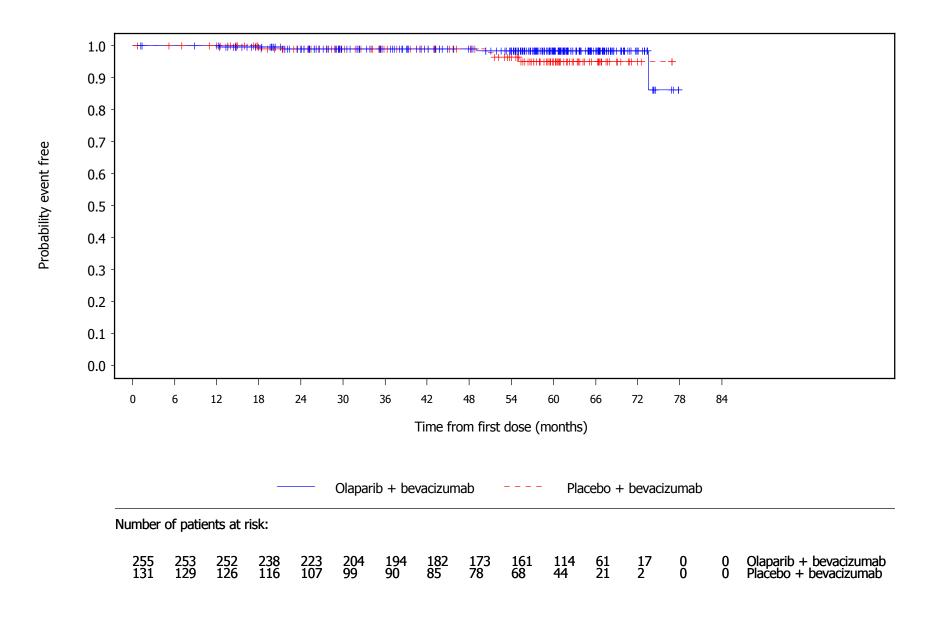


Figure 3.3.19 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

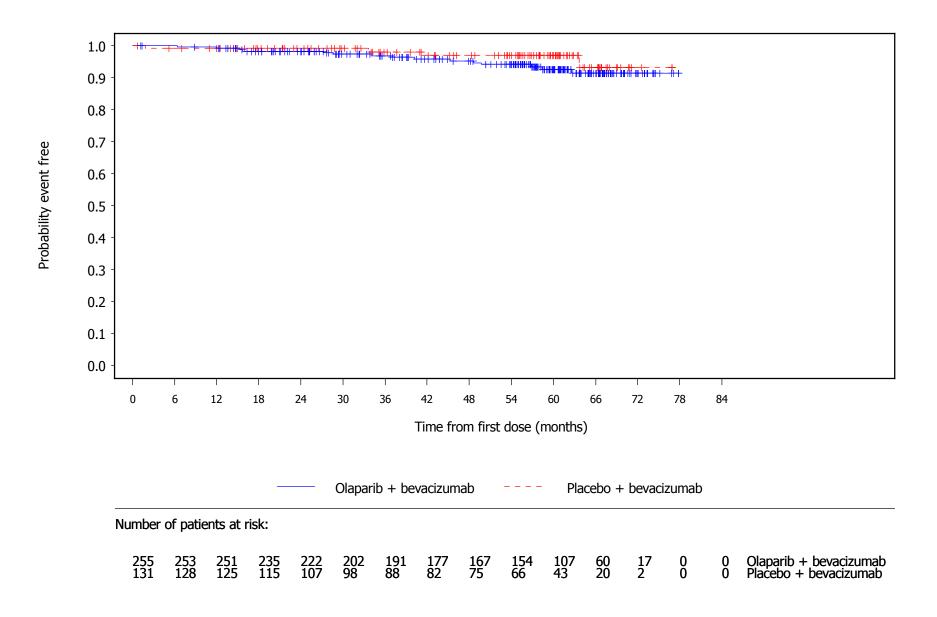


Figure 3.3.20 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI: Pneumonitis Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

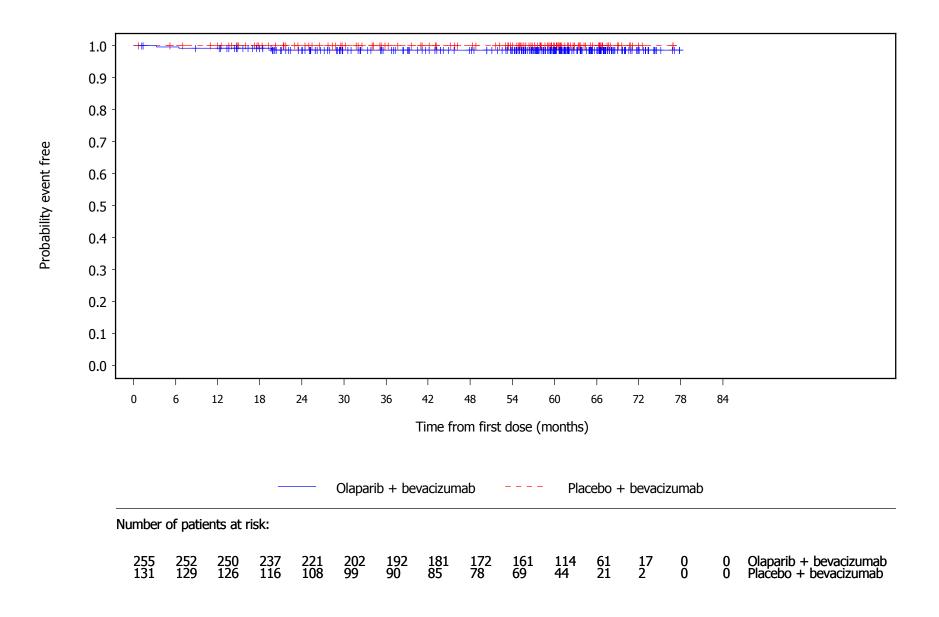


Figure 3.3.21 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

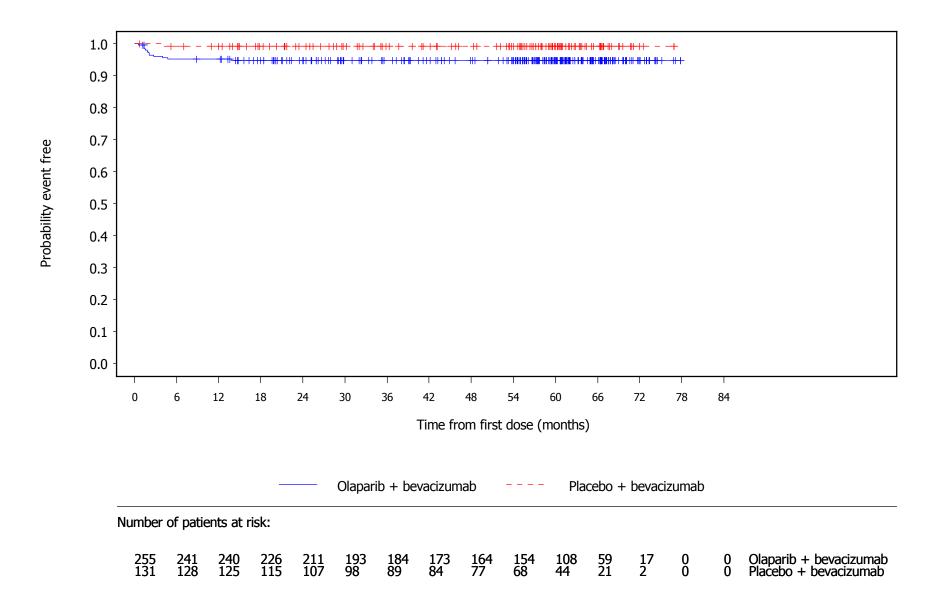


Figure 3.3.22 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

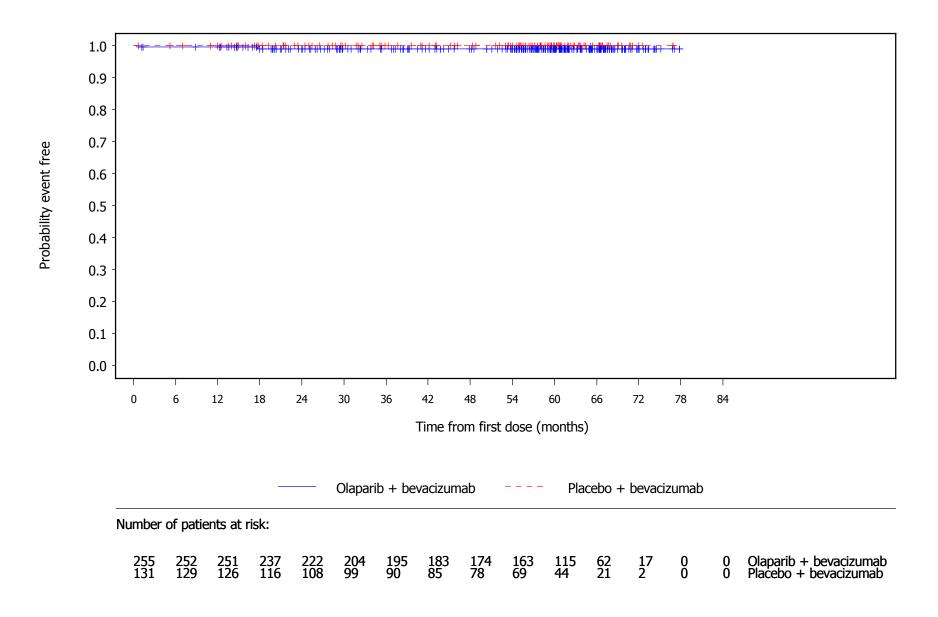


Figure 3.3.23 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

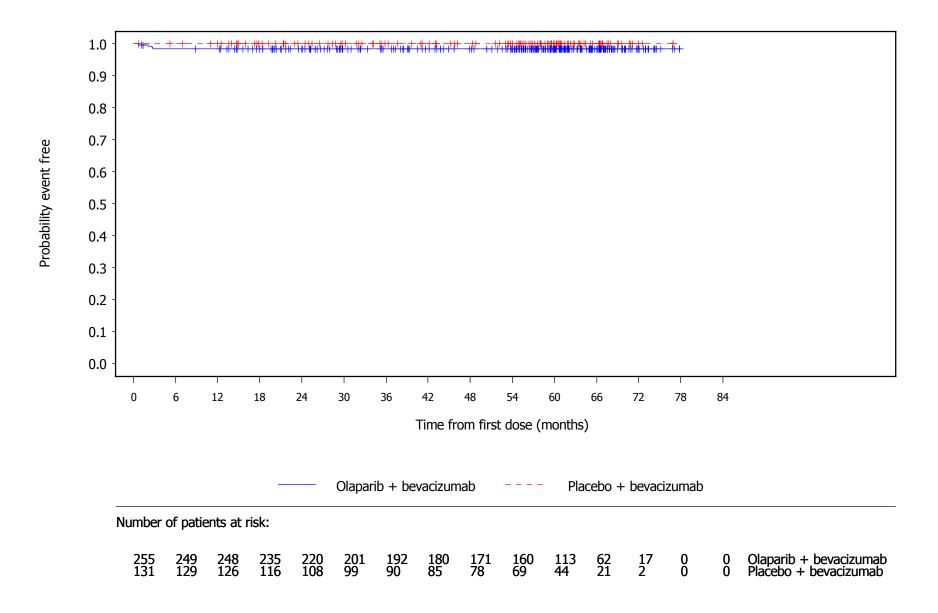


Figure 3.3.24 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

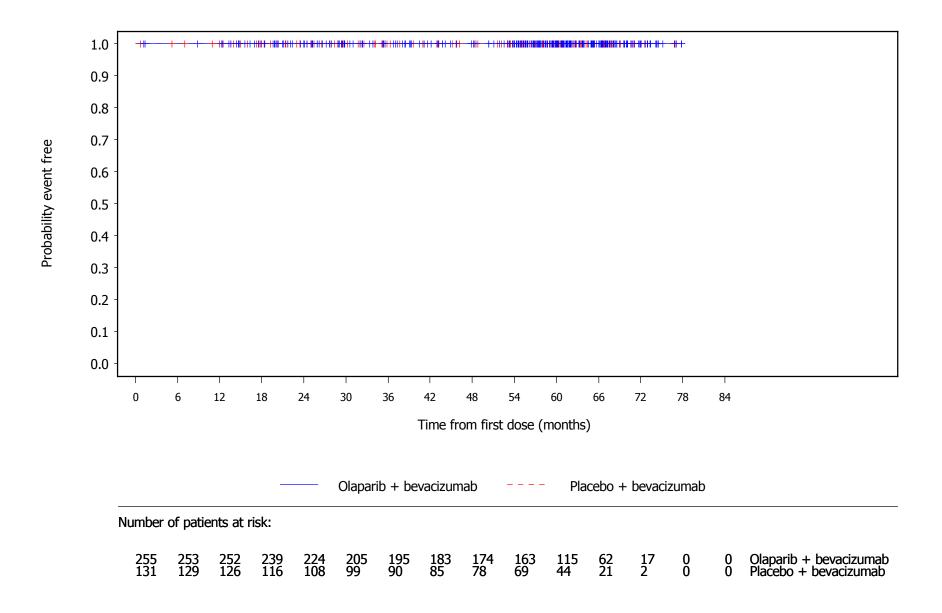


Figure 3.3.25 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

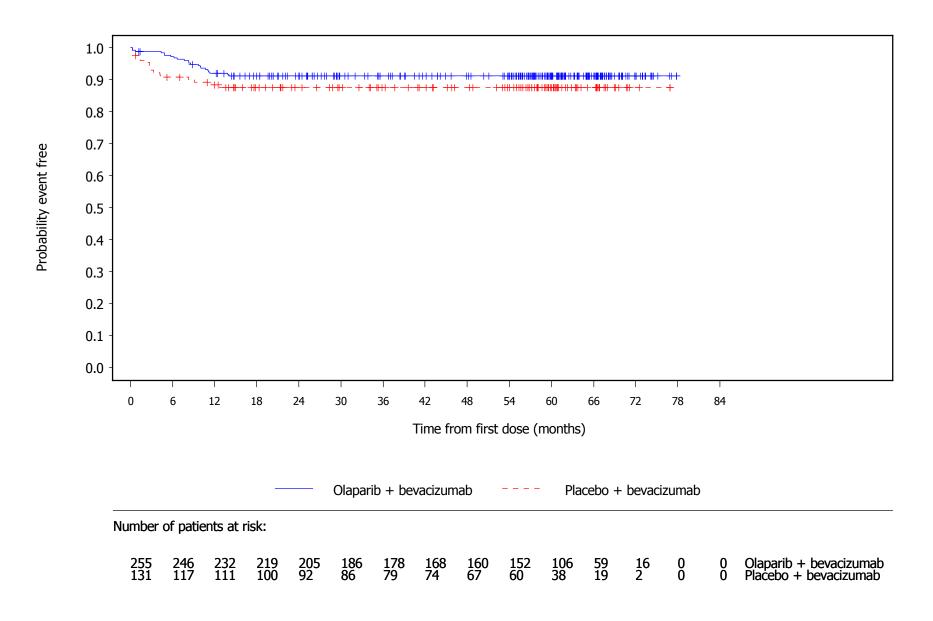


Figure 3.3.26 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

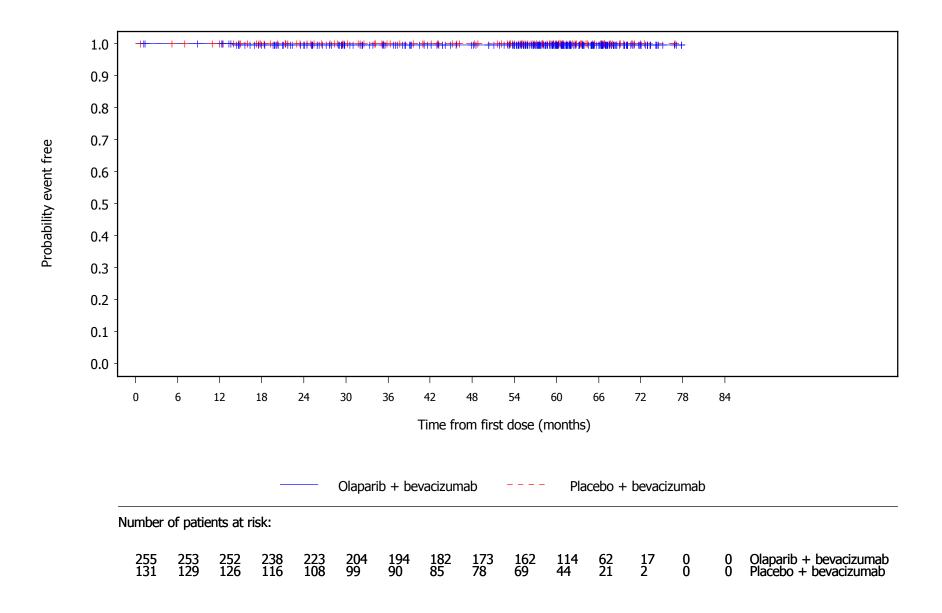


Figure 3.3.27 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: GI perforations, abscesses and fistulae Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

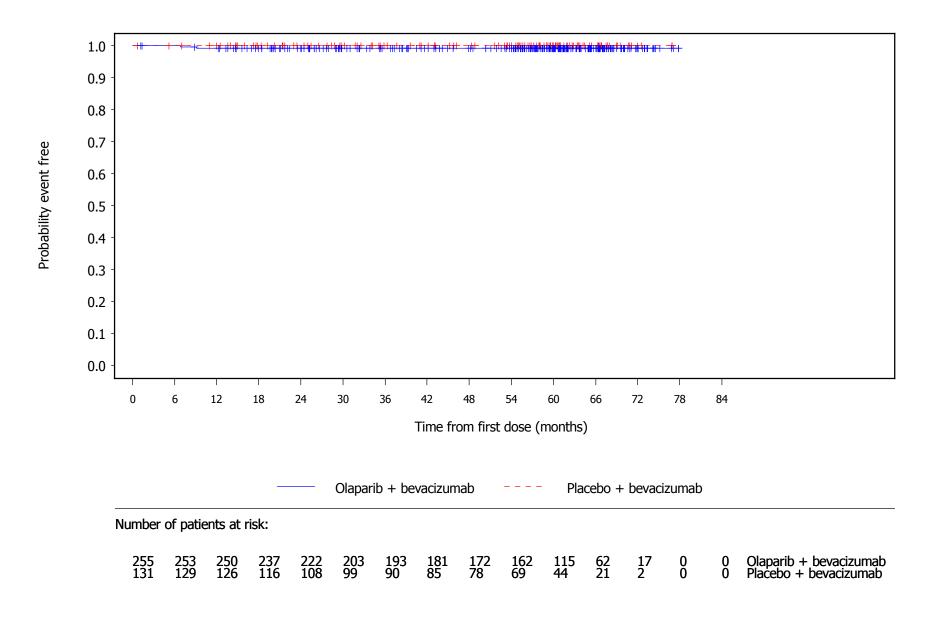


Figure 3.3.28 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

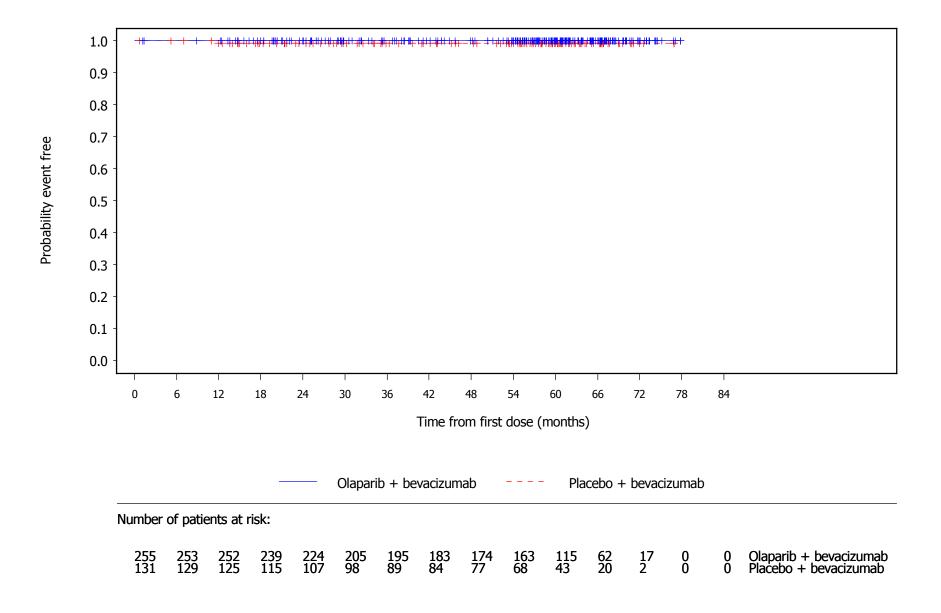


Figure 3.3.29 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

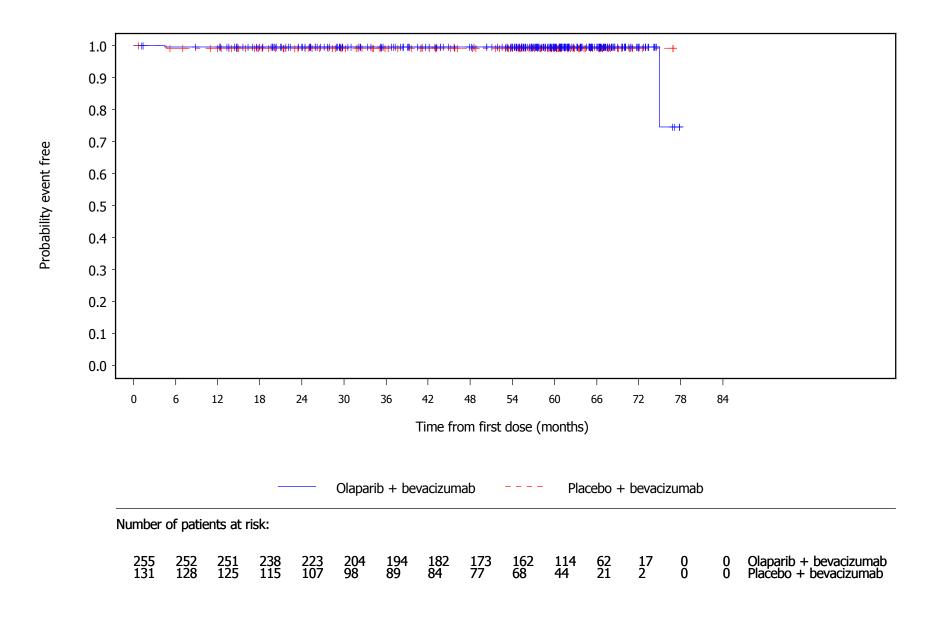


Figure 3.3.30 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

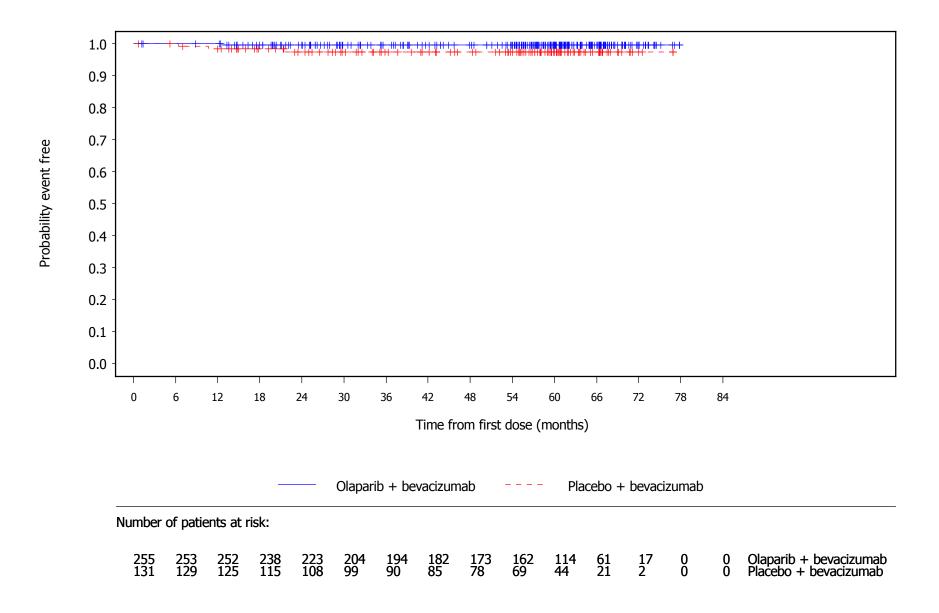


Figure 3.3.31 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

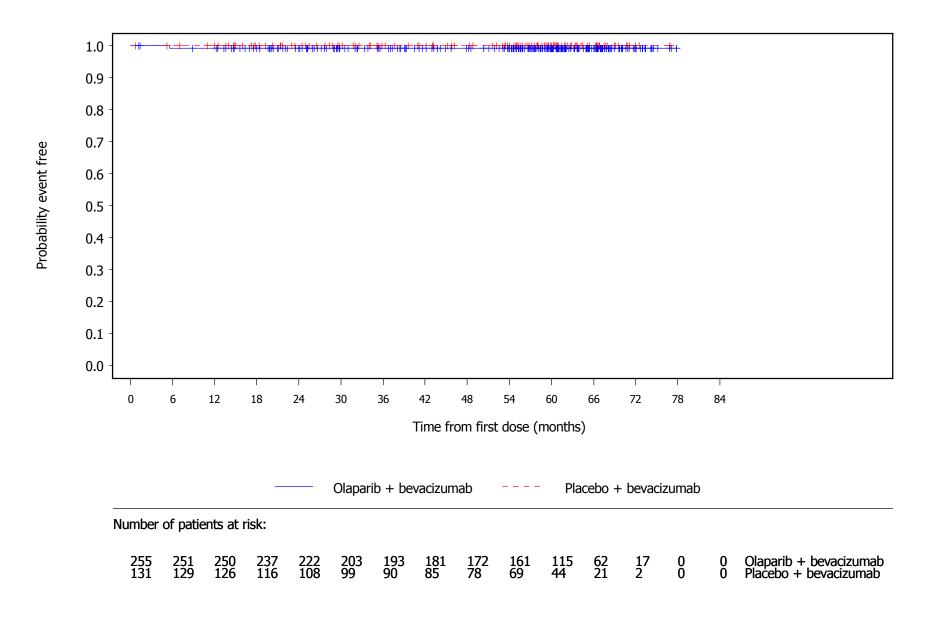


Figure 3.3.32 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

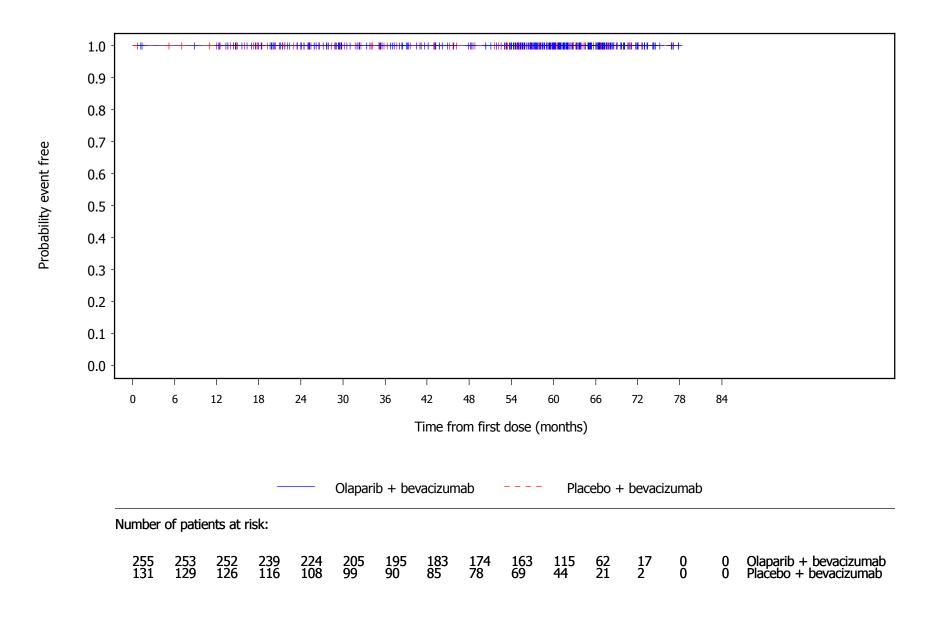


Figure 3.3.33 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: MDS/AML Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

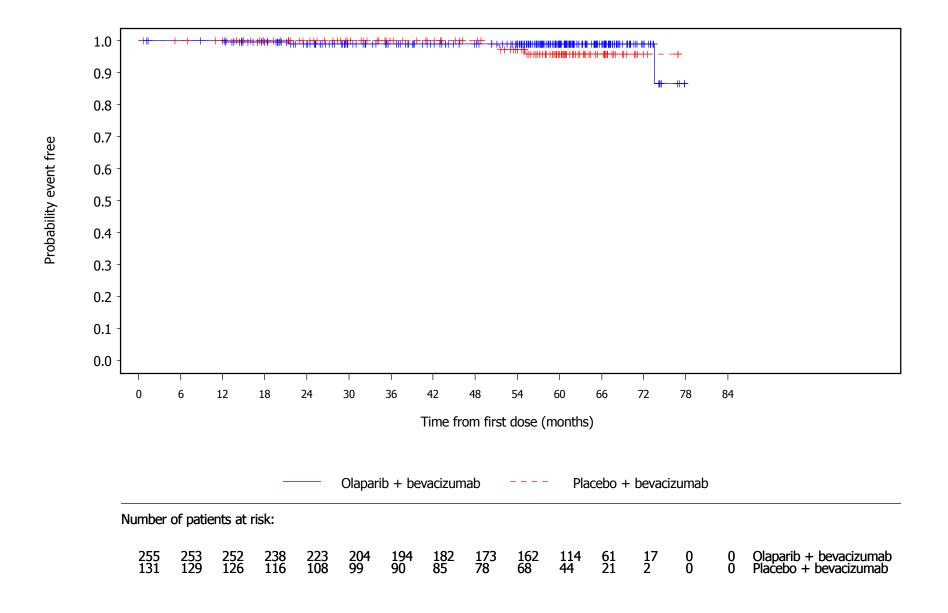


Figure 3.3.34 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Myelodysplastic syndrome and Acute myeloid leukaemia

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

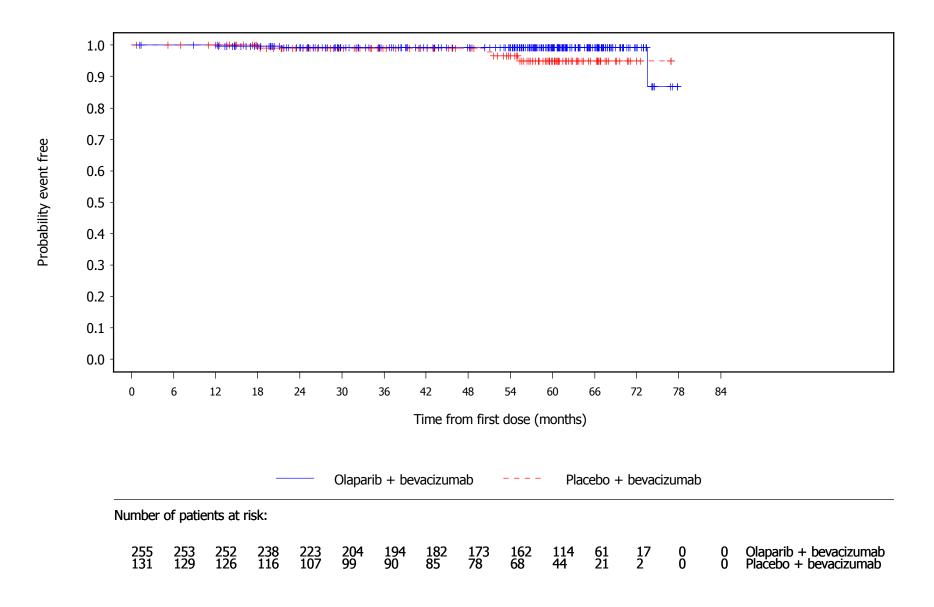


Figure 3.3.35 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

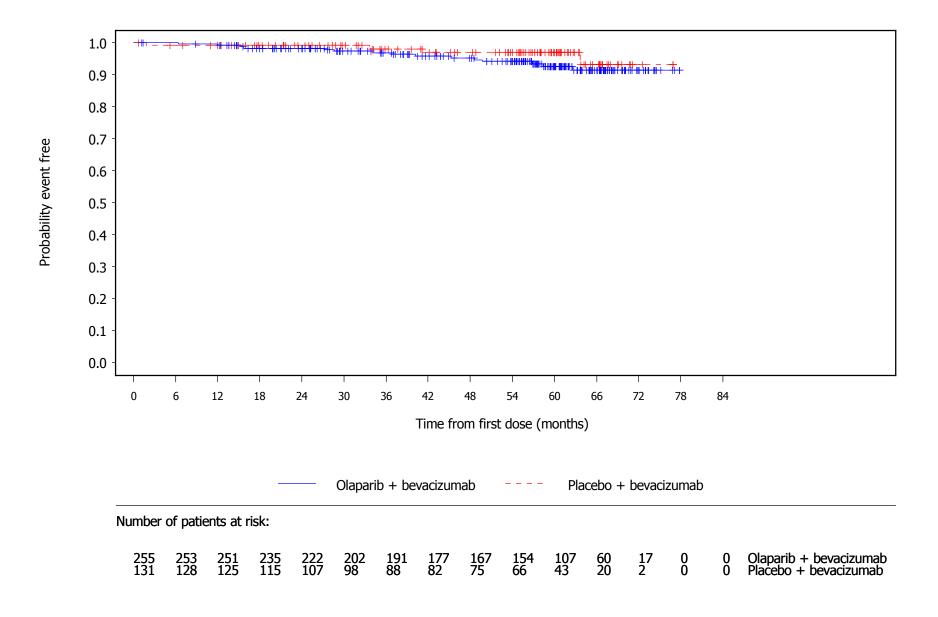


Figure 3.3.36 PAOLA1: Kaplan-Meier plot of time to first occurrence of Serious AESI: Pneumonitis Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

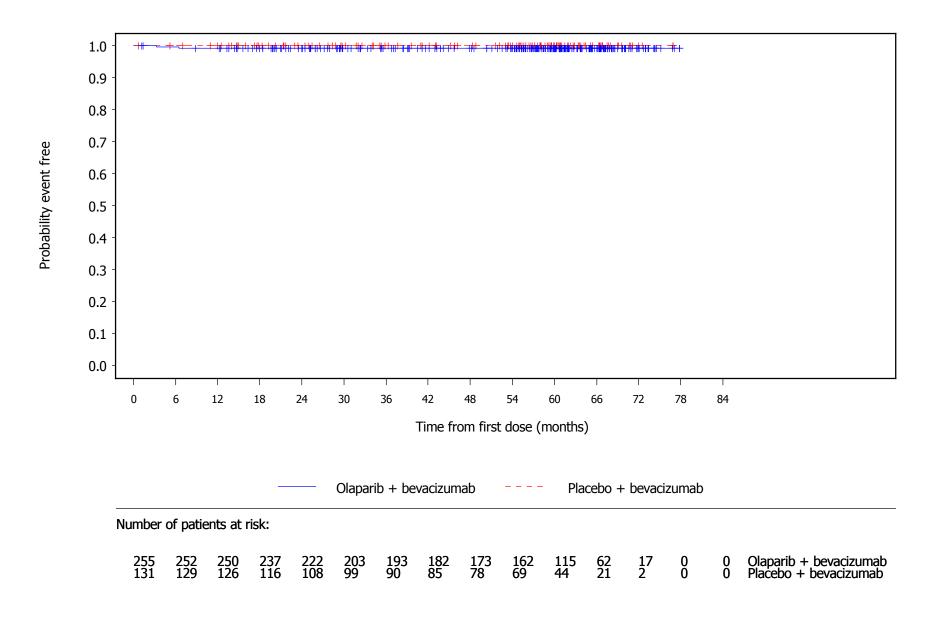


Figure 3.3.37 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

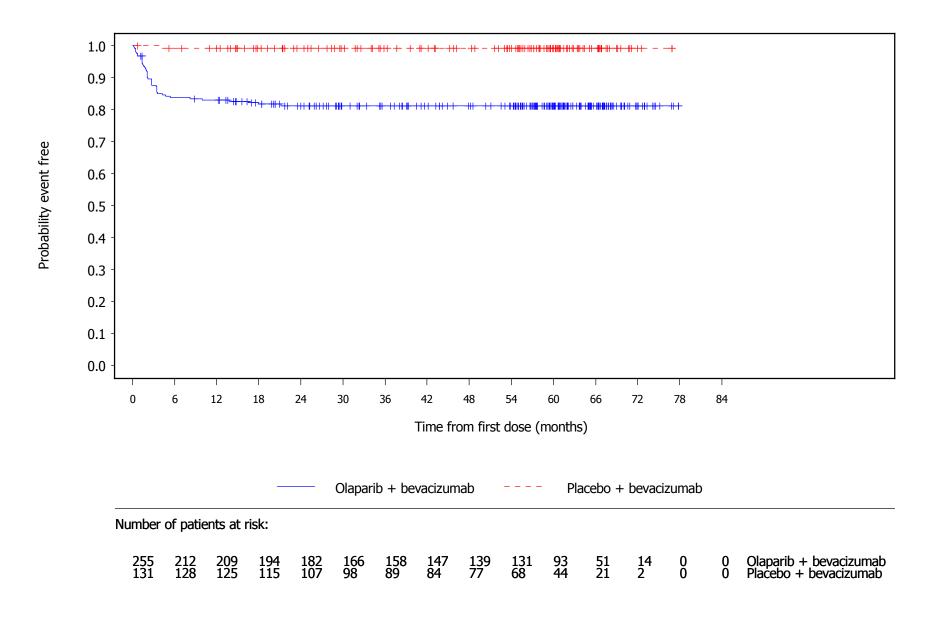


Figure 3.3.38 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

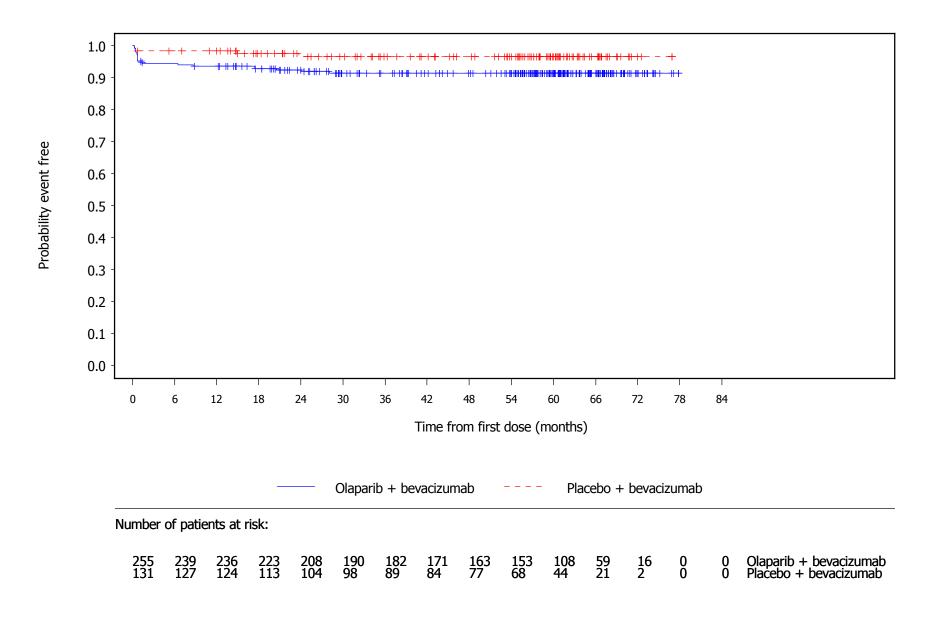


Figure 3.3.39 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

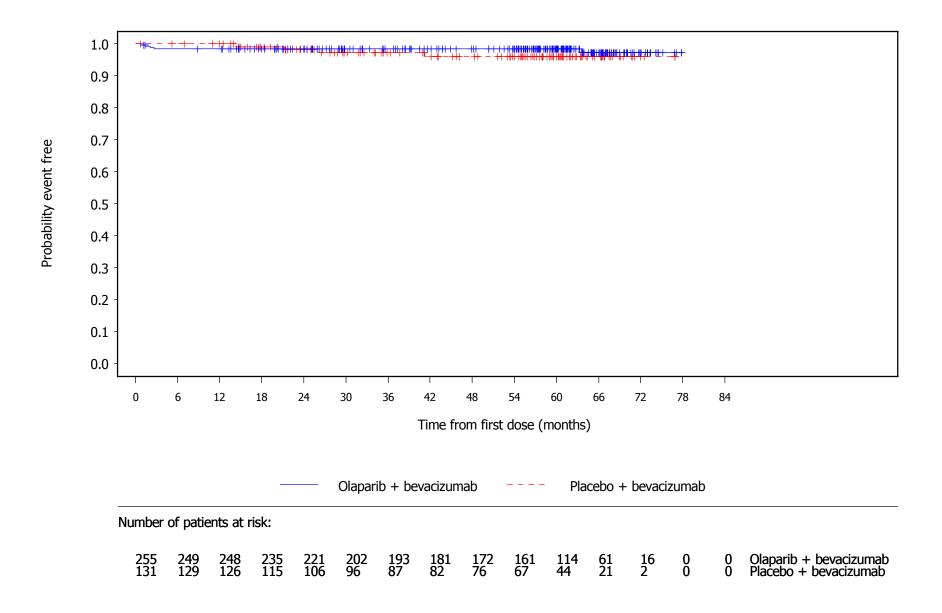


Figure 3.3.40 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Nausea Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

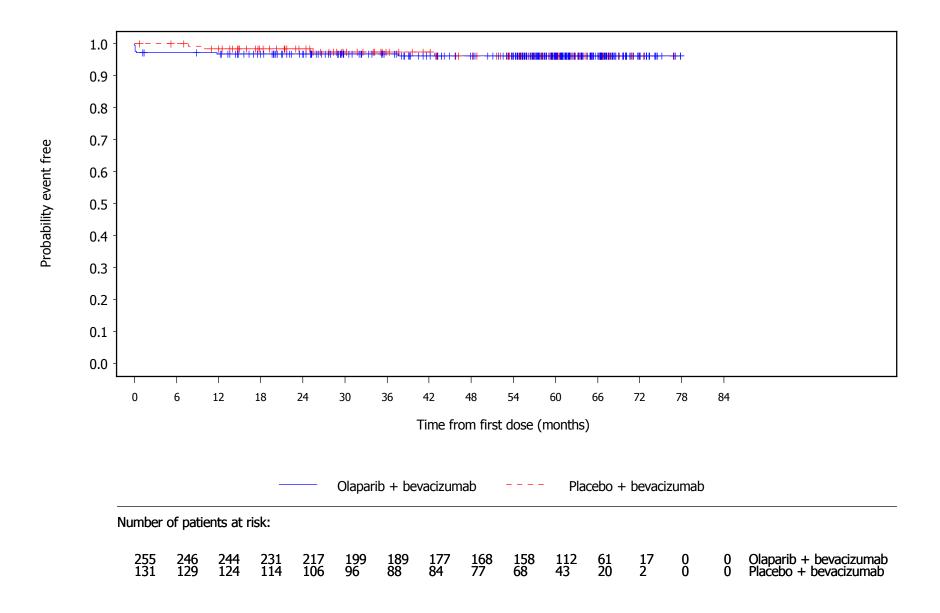


Figure 3.3.41 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

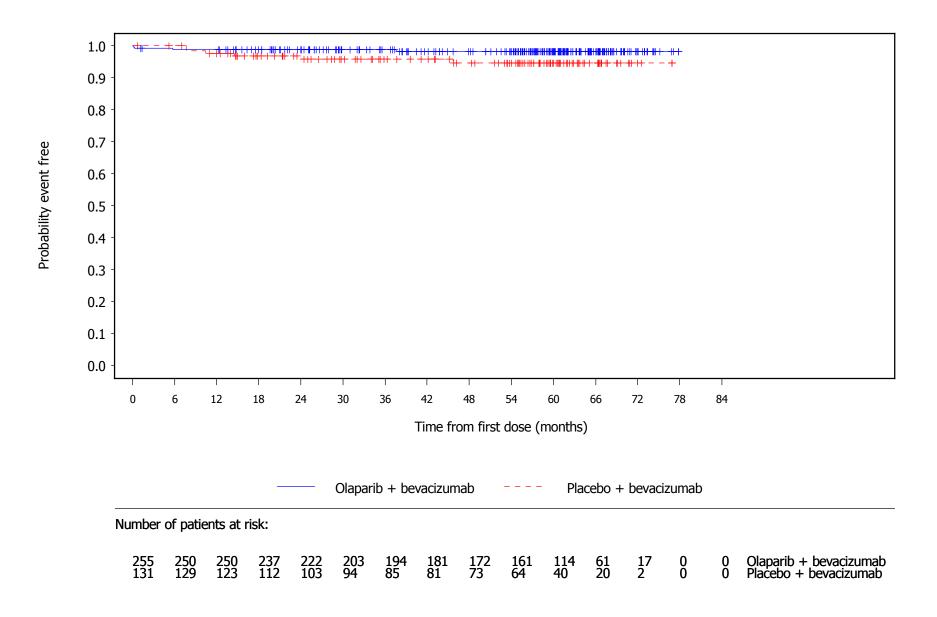


Figure 3.3.42 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Fatigue and Asthenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

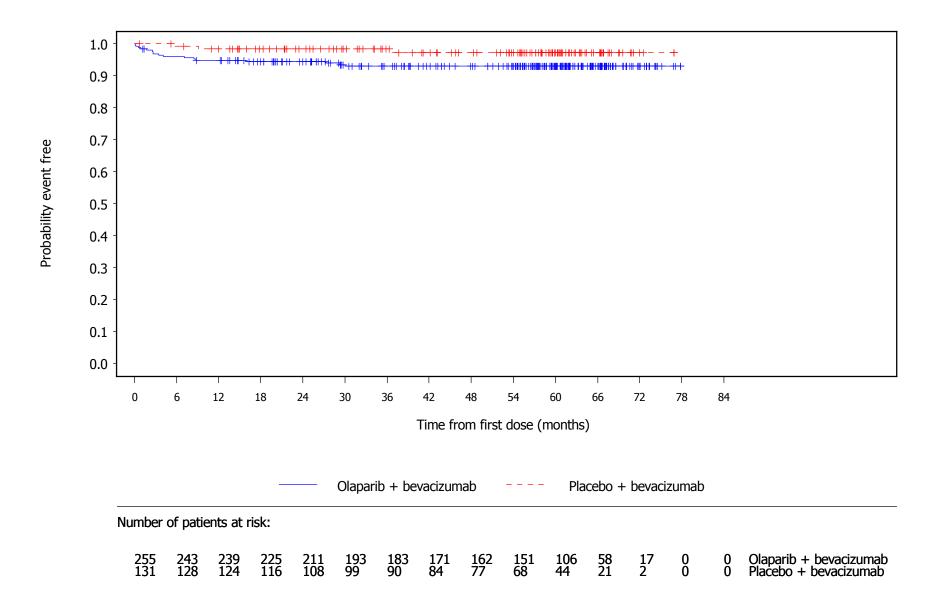


Figure 3.3.43 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

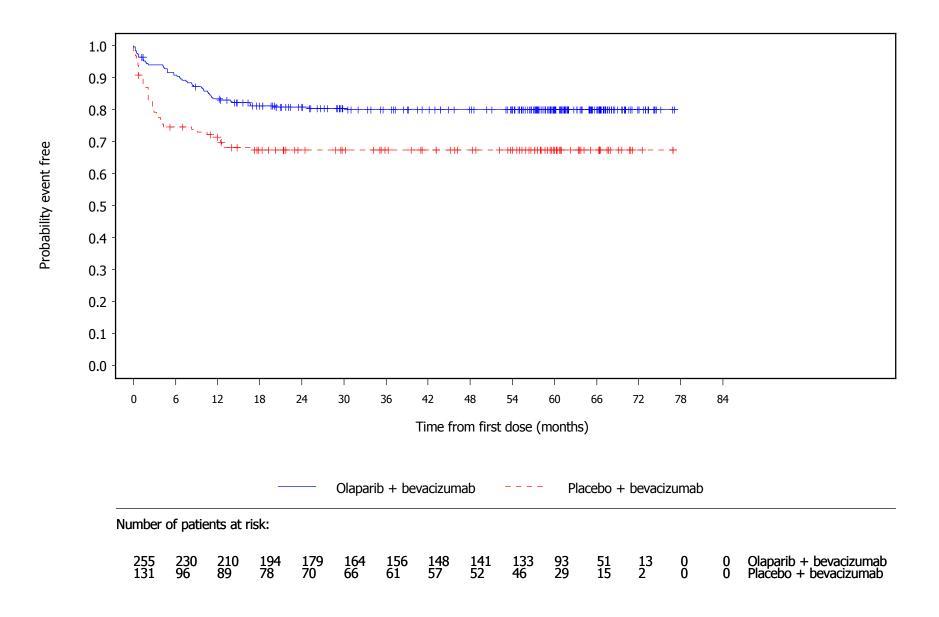


Figure 3.3.44 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

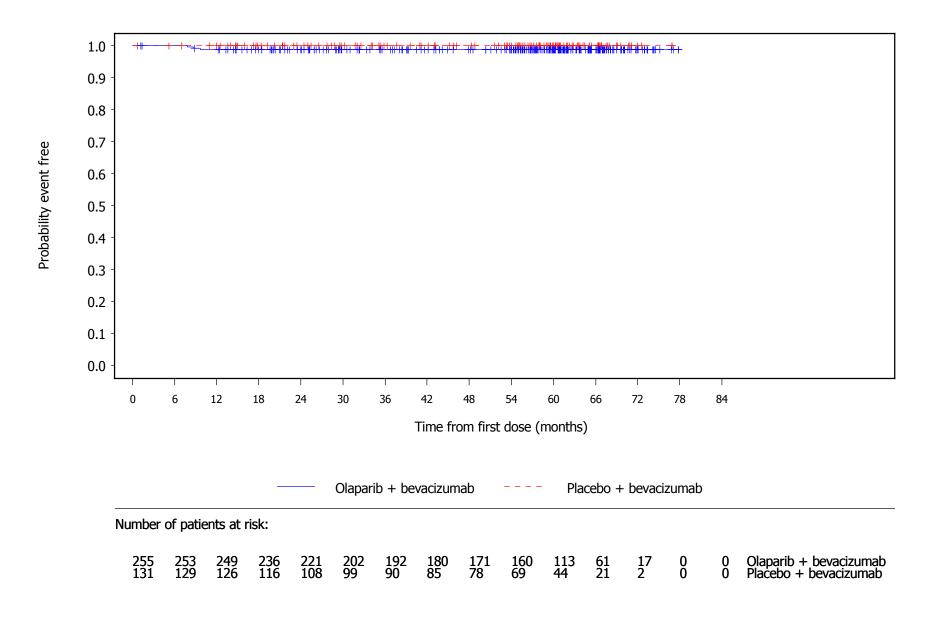


Figure 3.3.45 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: GI perforations, abscesses and fistulae Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

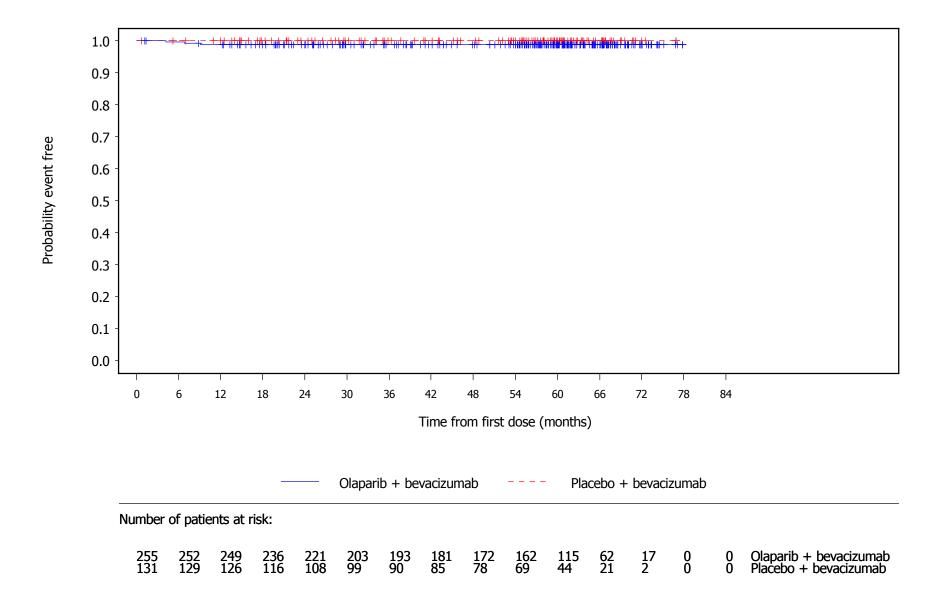


Figure 3.3.46 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

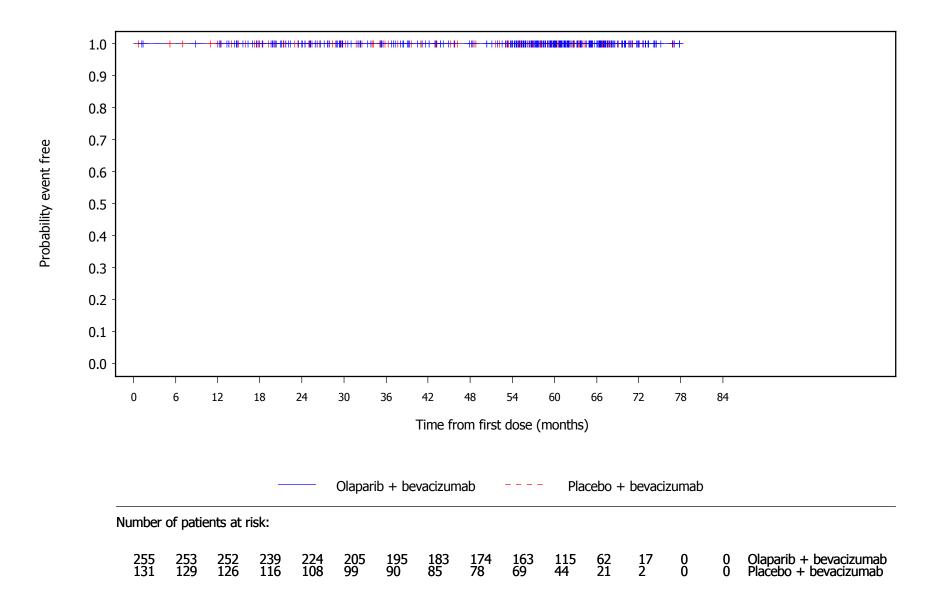


Figure 3.3.47 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

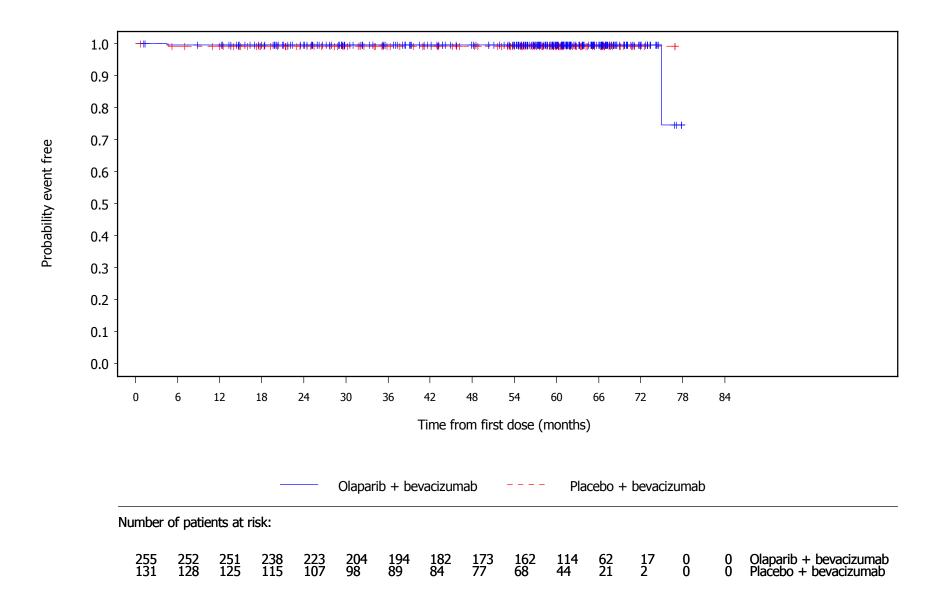


Figure 3.3.48 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

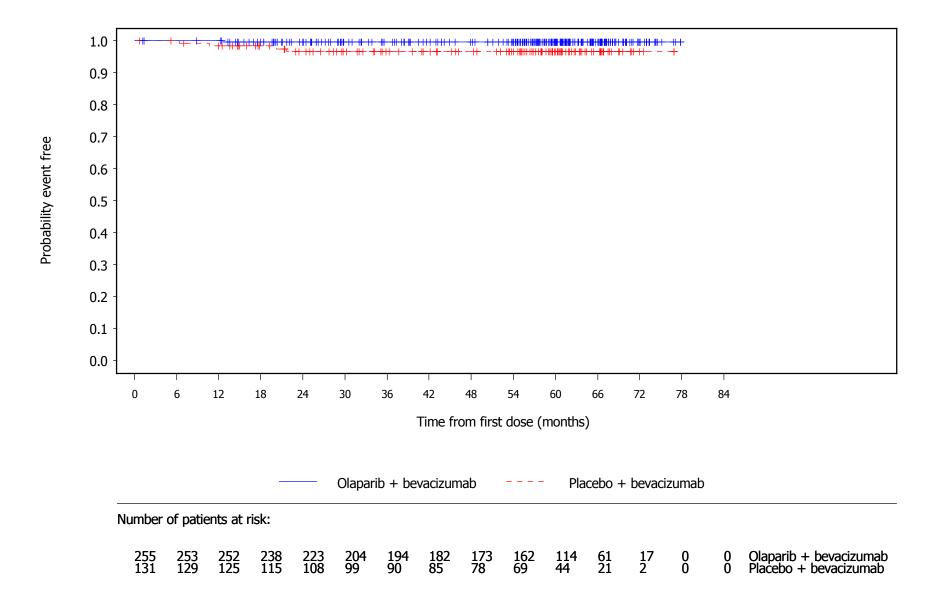


Figure 3.3.49 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

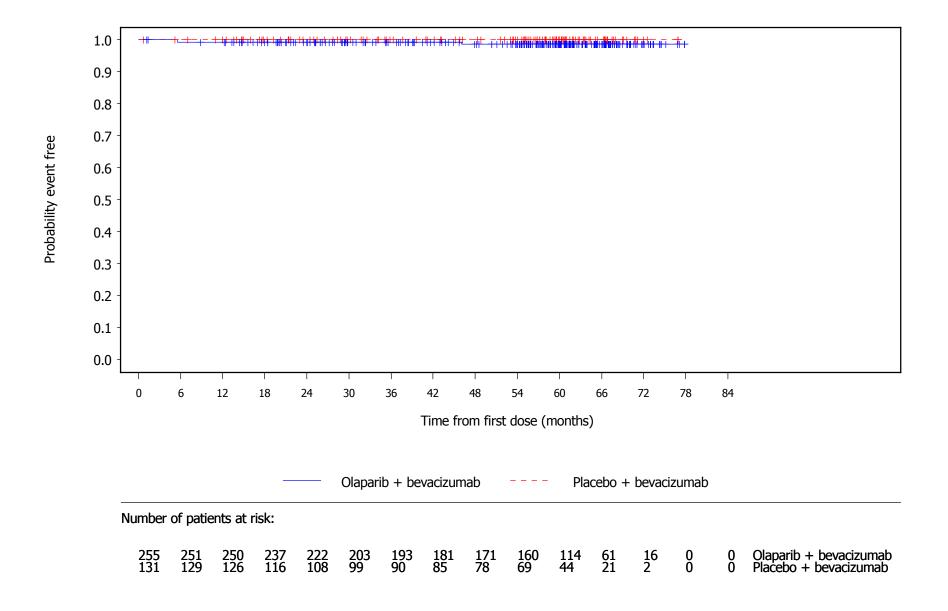


Figure 3.3.50 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Congestive heart failure Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

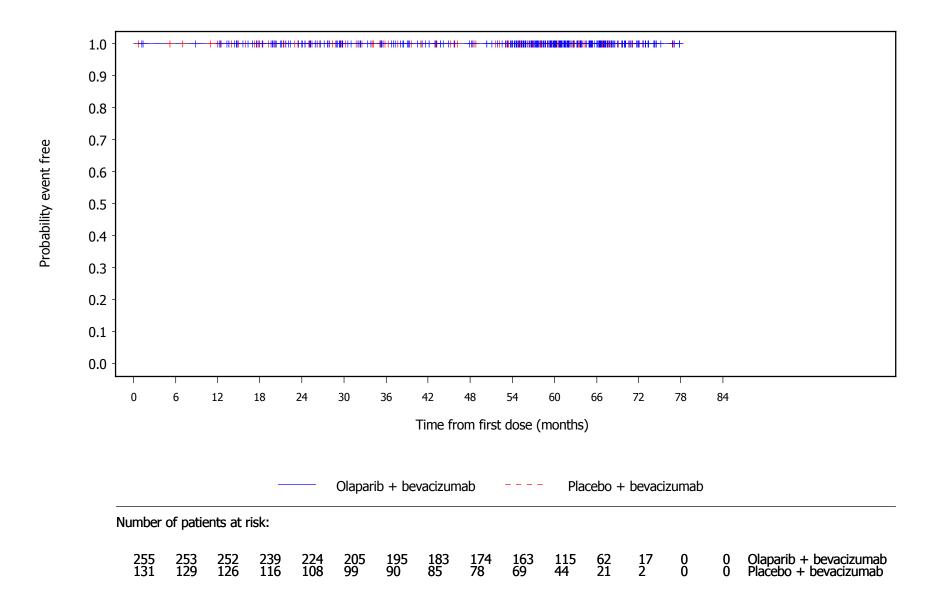


Figure 3.3.51 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

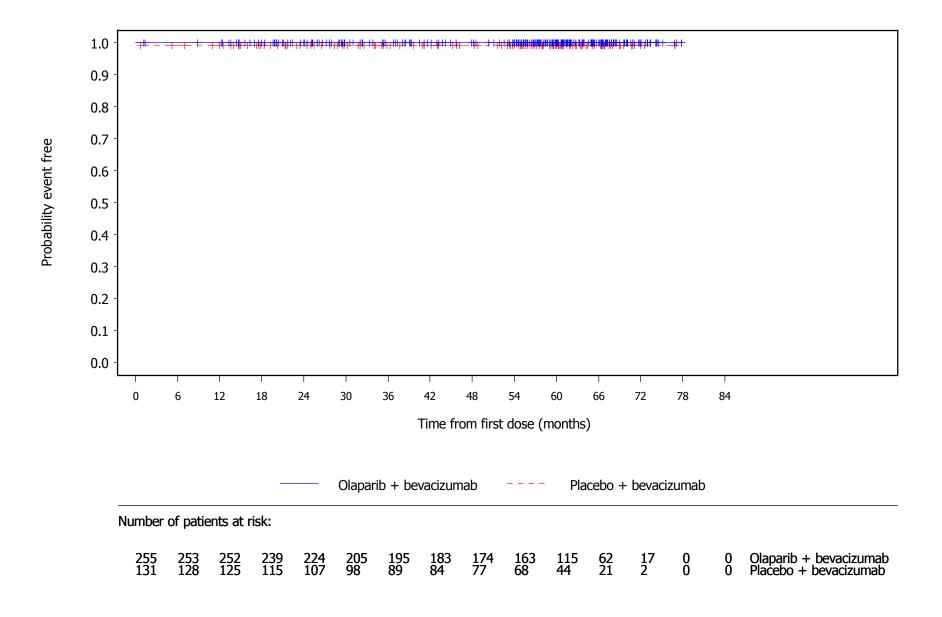


Figure 3.3.52 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: MDS/AML Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

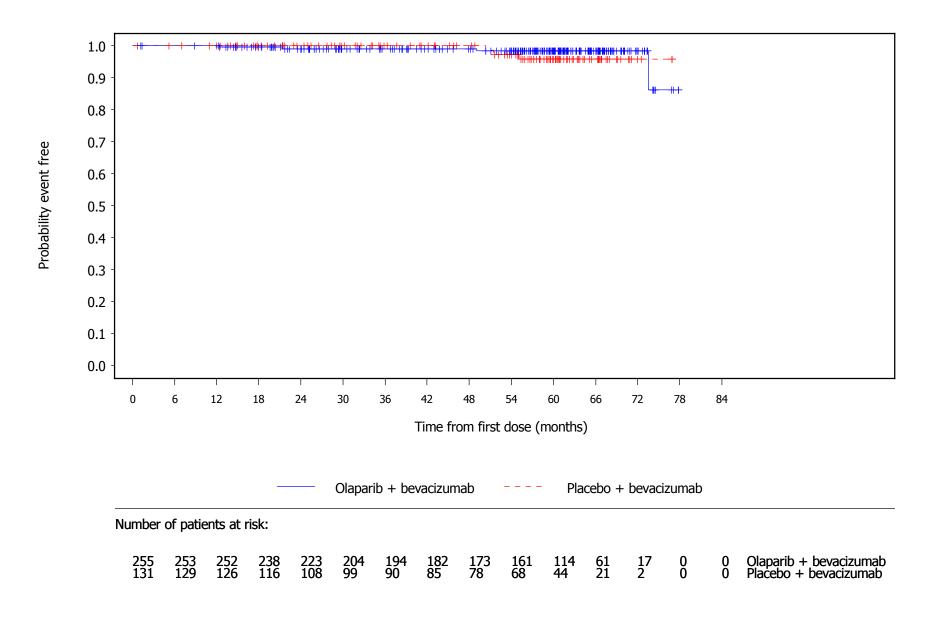


Figure 3.3.53 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Myelodysplastic syndrome and Acute myeloid leukaemia

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

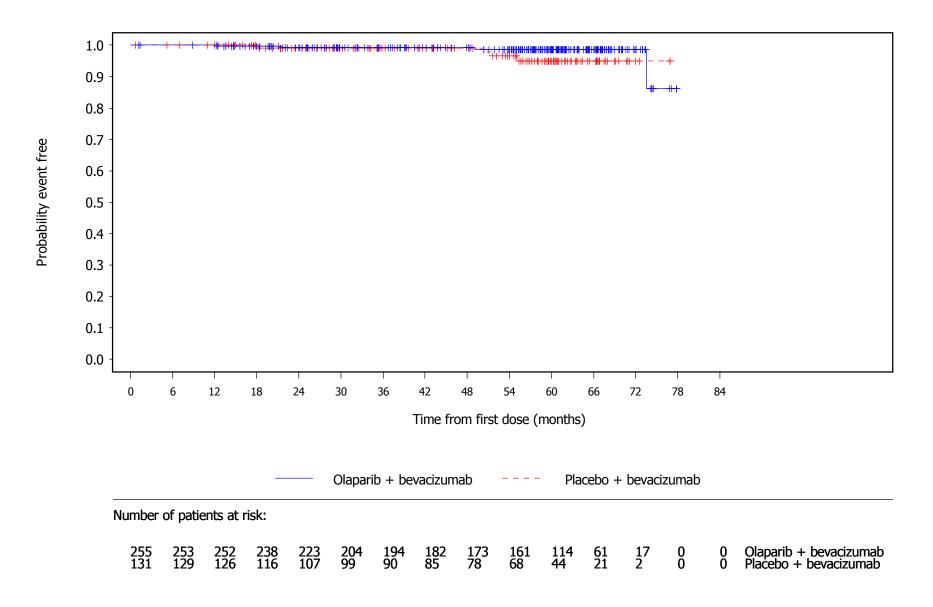


Figure 3.3.54 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G>=3: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

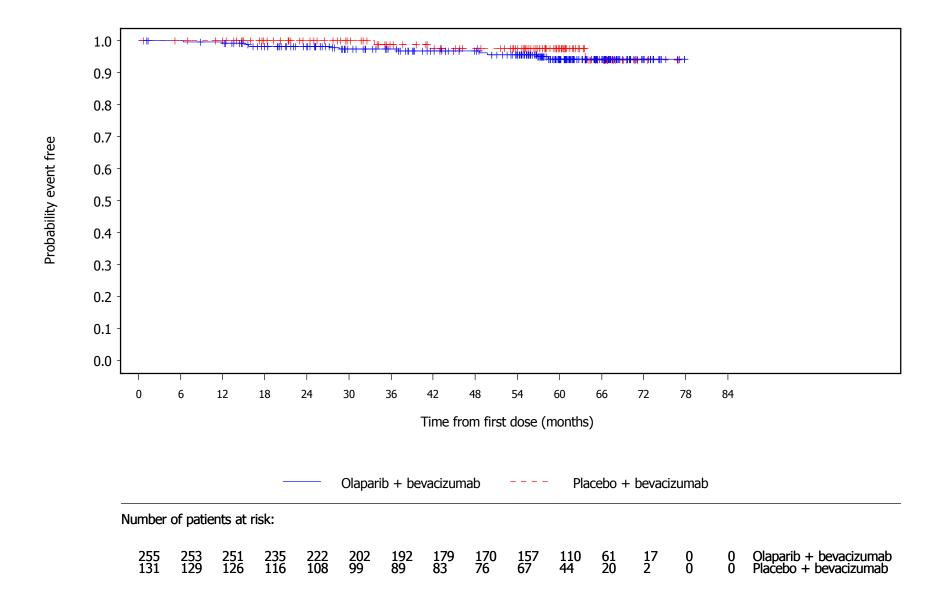


Figure 3.3.55 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

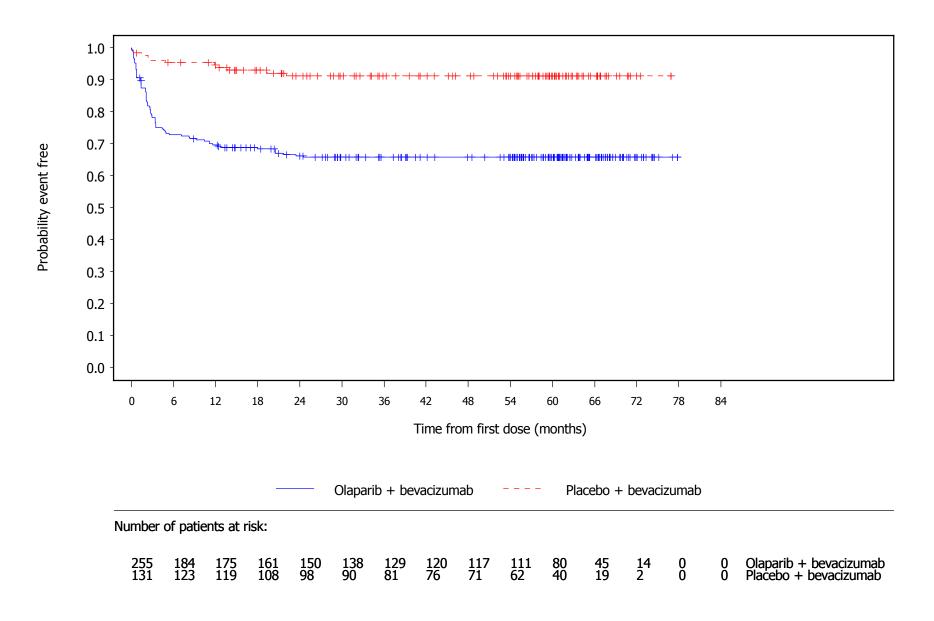


Figure 3.3.56 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

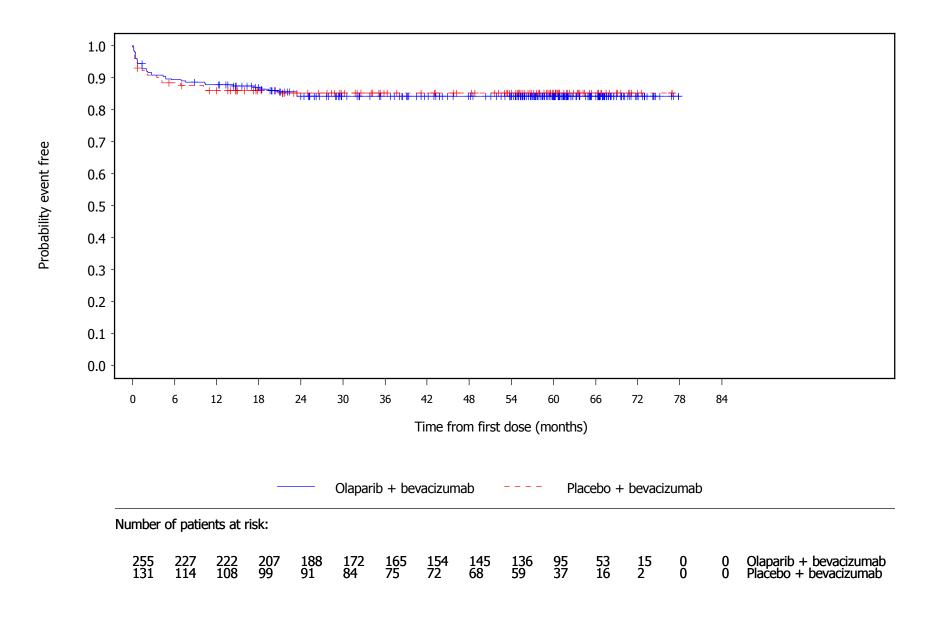


Figure 3.3.57 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

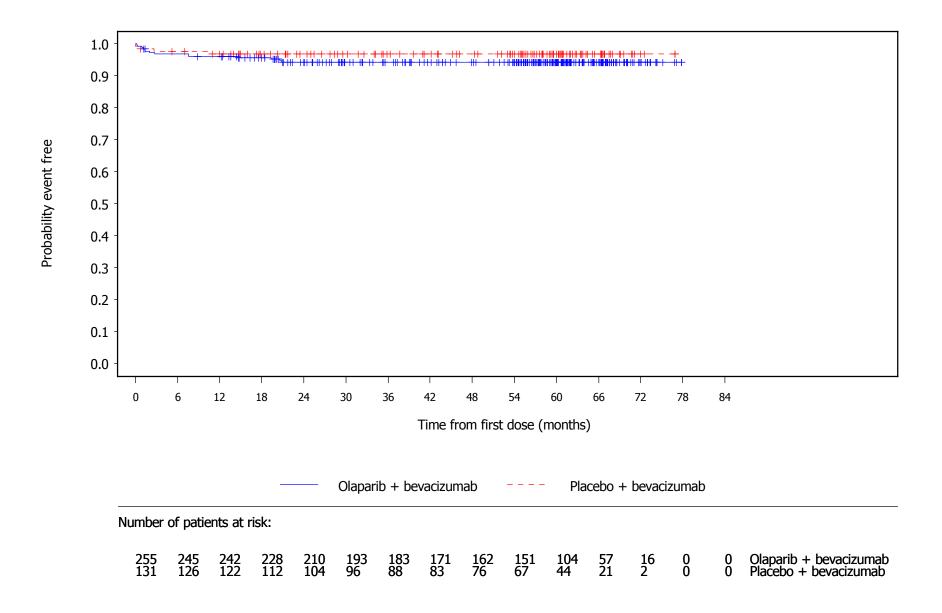


Figure 3.3.58 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Nausea Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

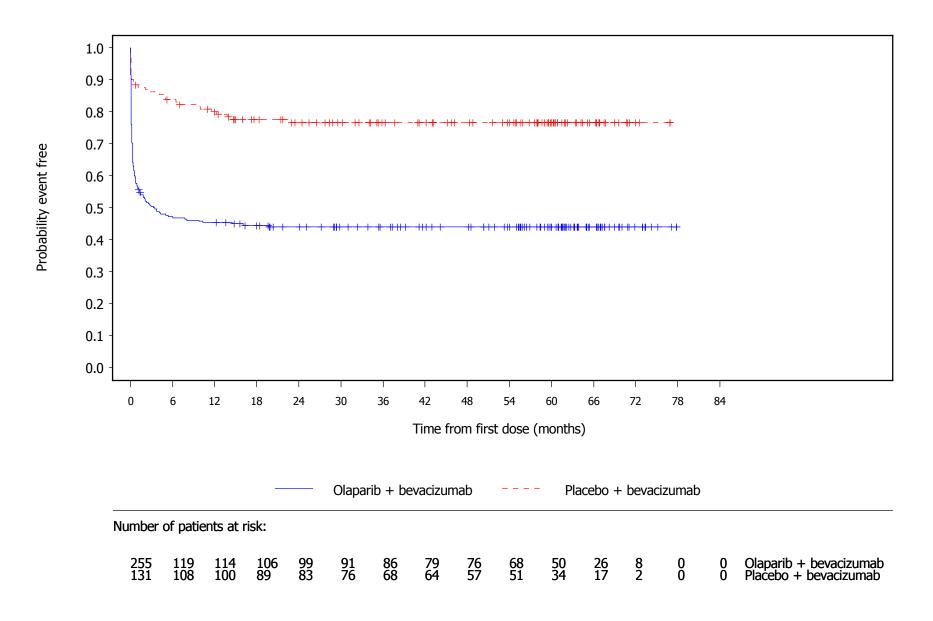


Figure 3.3.59 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

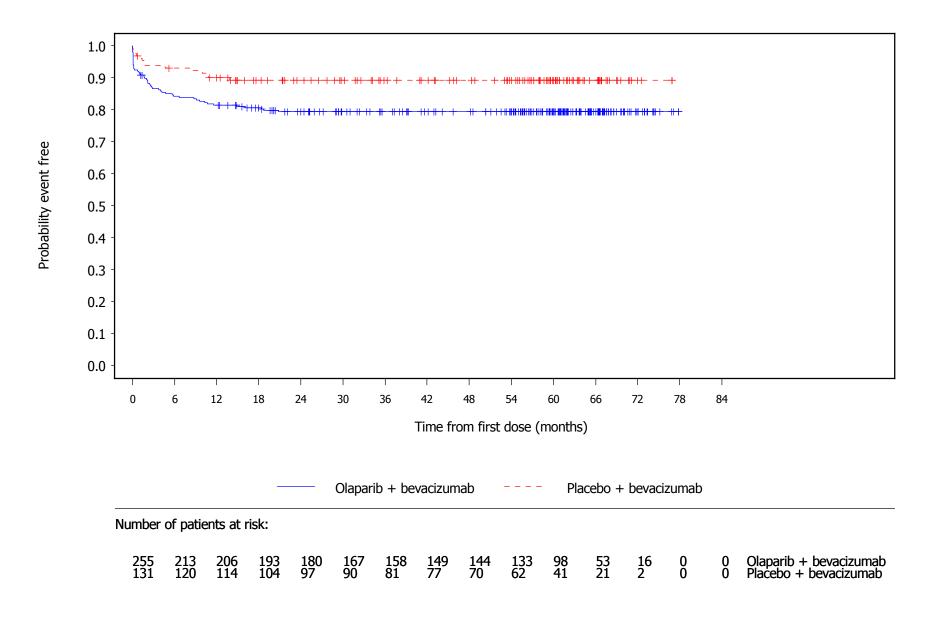


Figure 3.3.60 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Fatigue and Asthenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

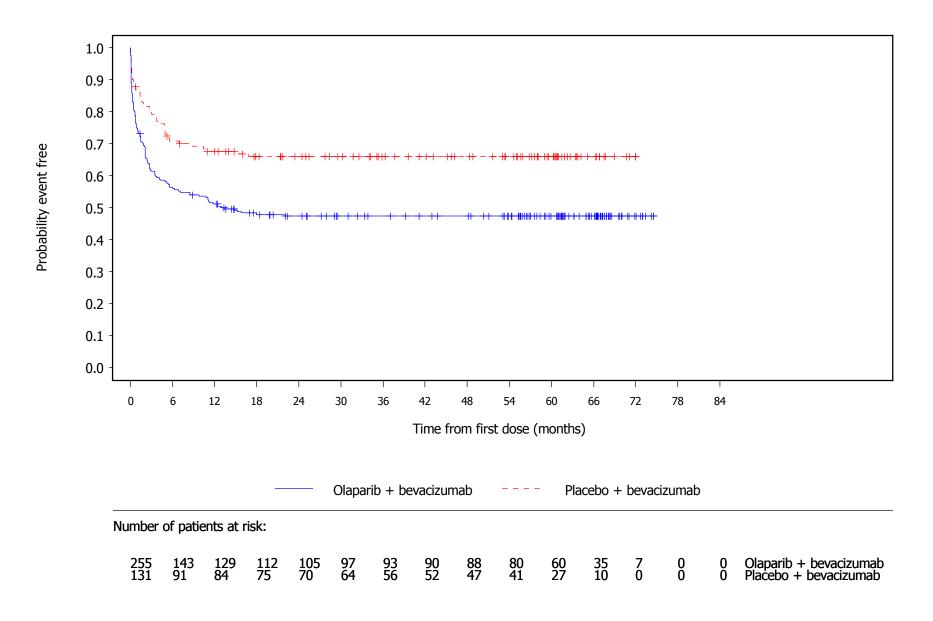


Figure 3.3.61 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

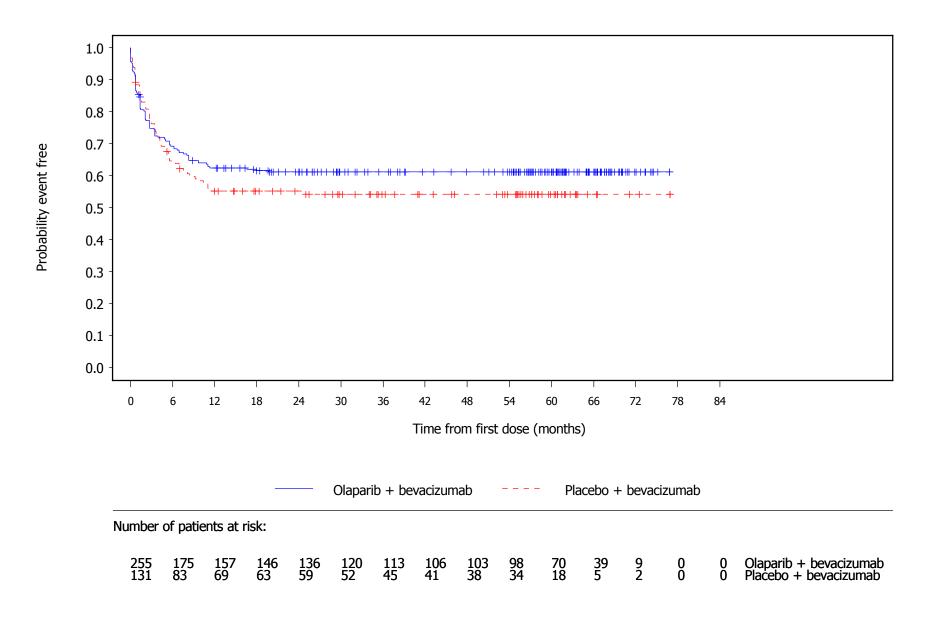


Figure 3.3.62 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

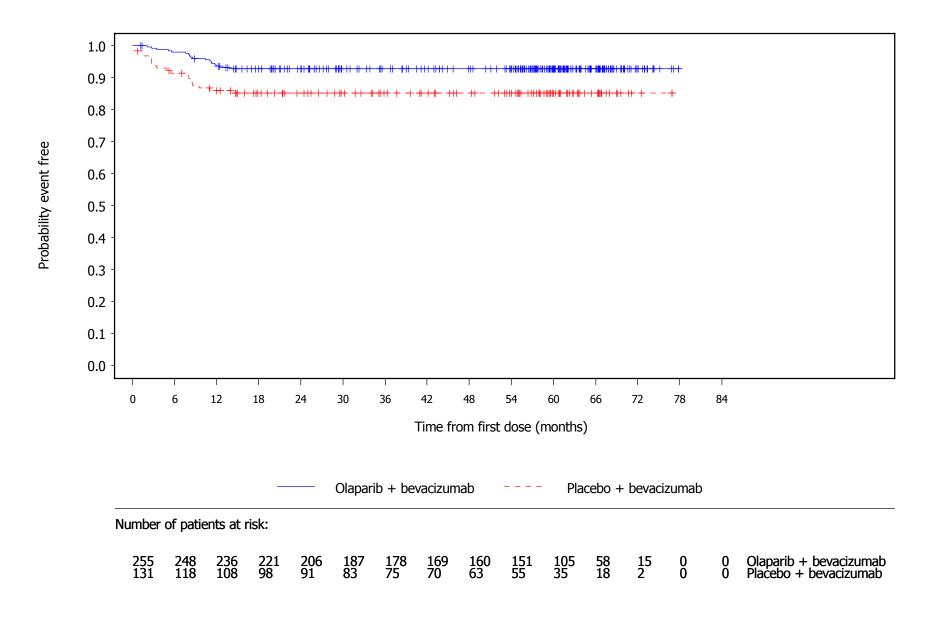


Figure 3.3.63 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

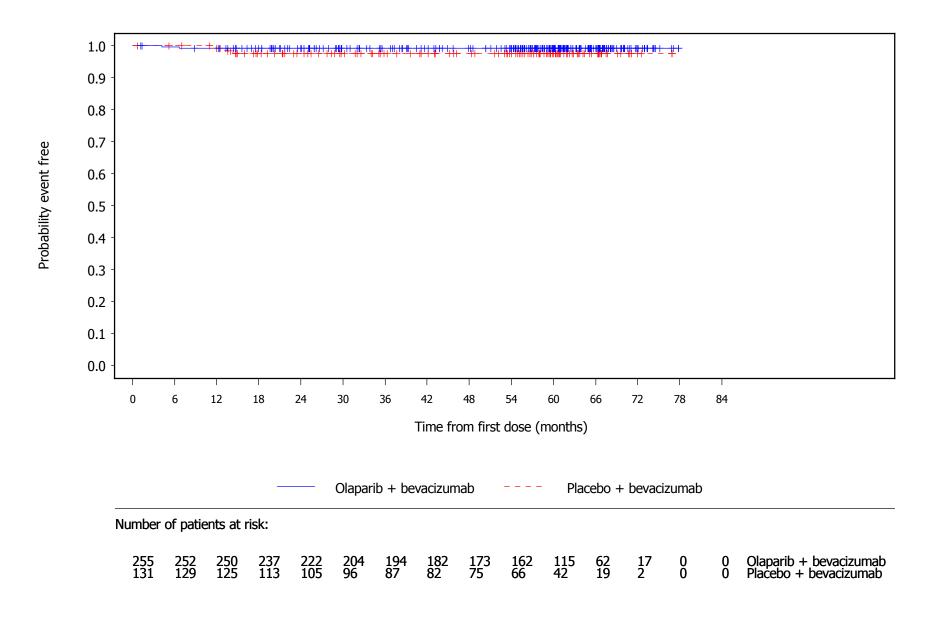


Figure 3.3.64 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

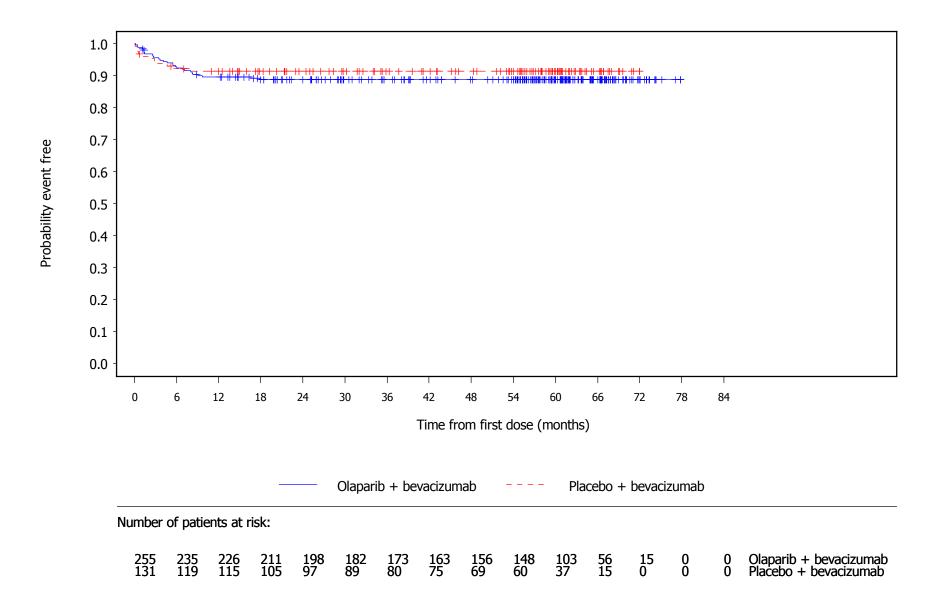


Figure 3.3.65 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

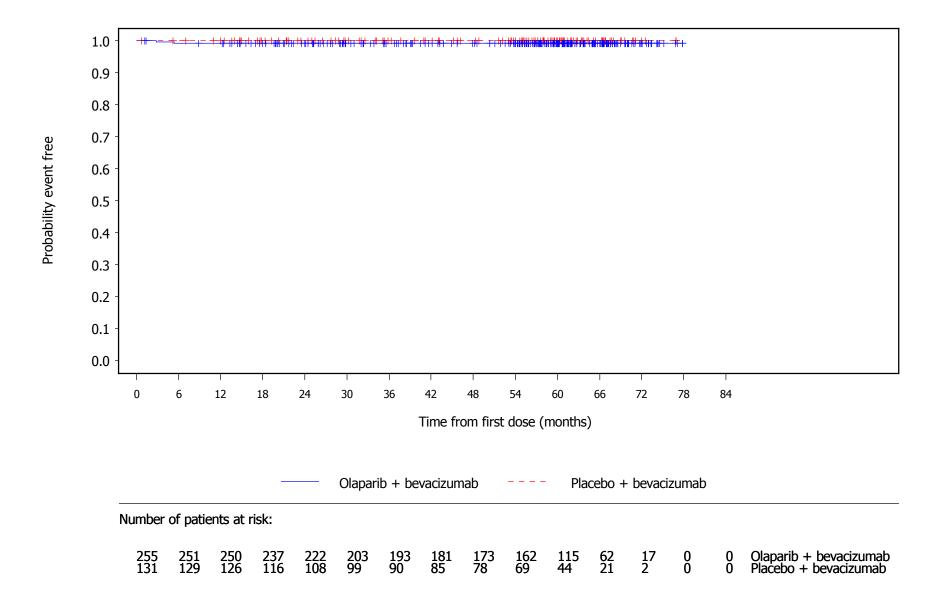


Figure 3.3.66 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

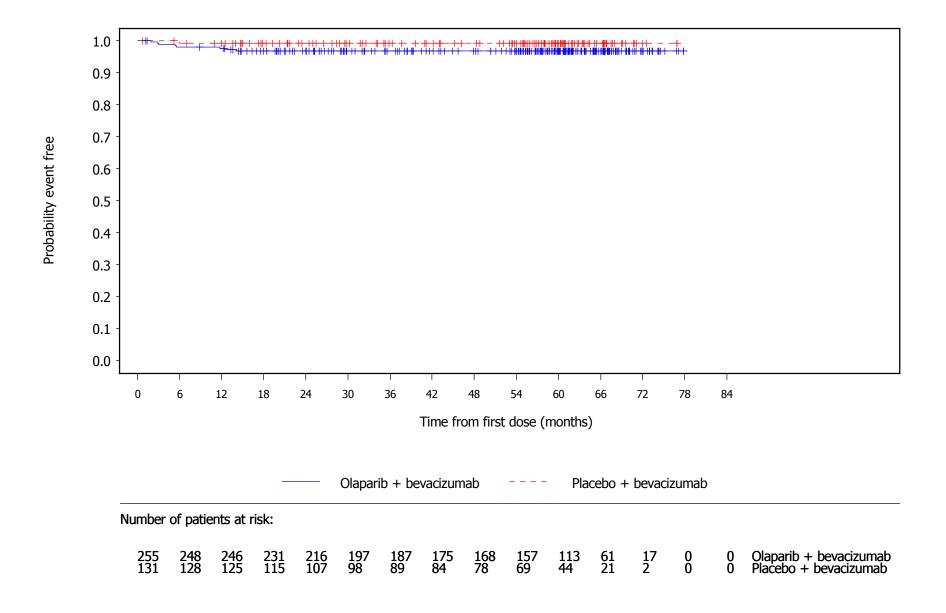


Figure 3.3.67 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Posterior Reversible Encephalopathy Syndrome (PRES)

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

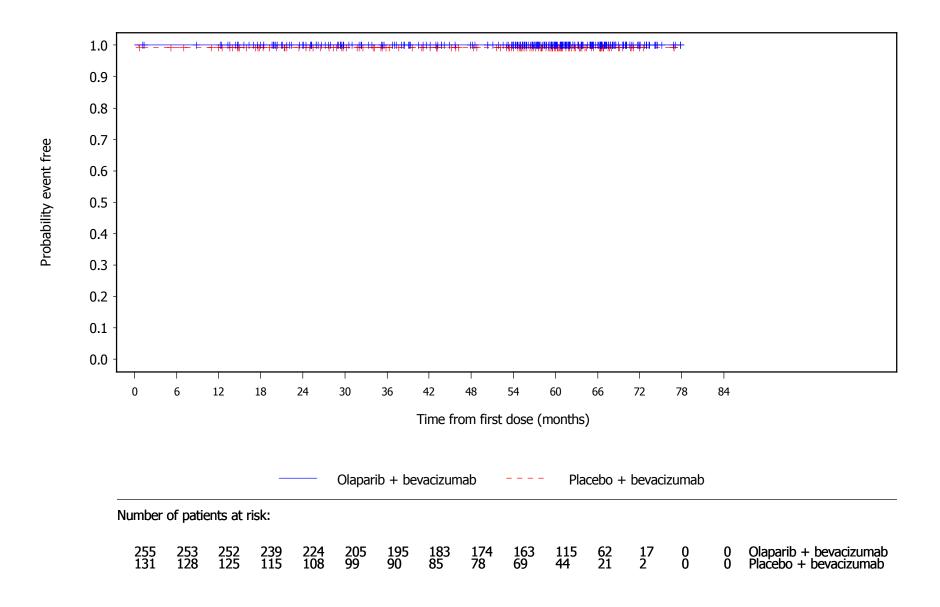


Figure 3.3.68 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

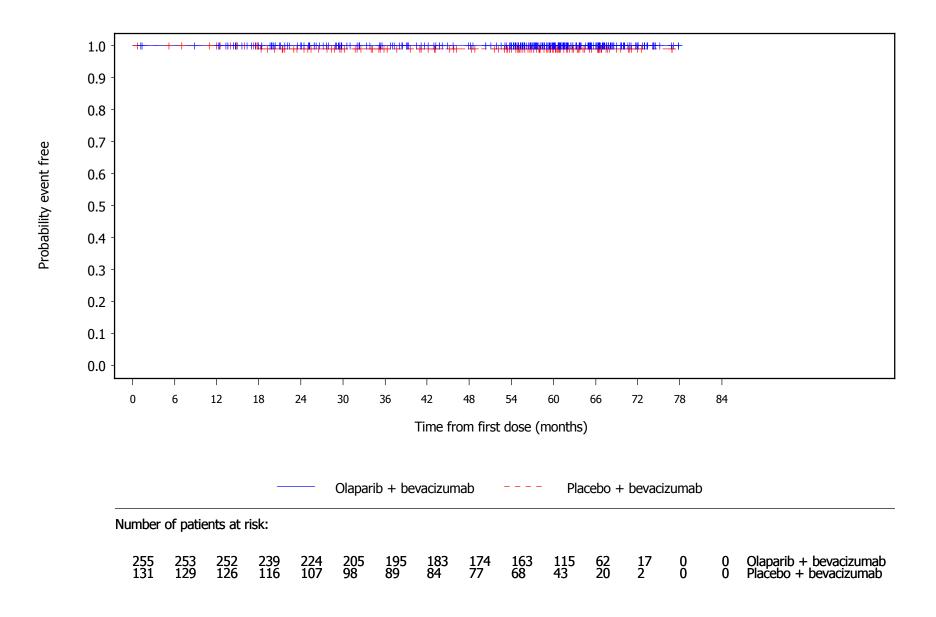


Figure 3.3.69 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

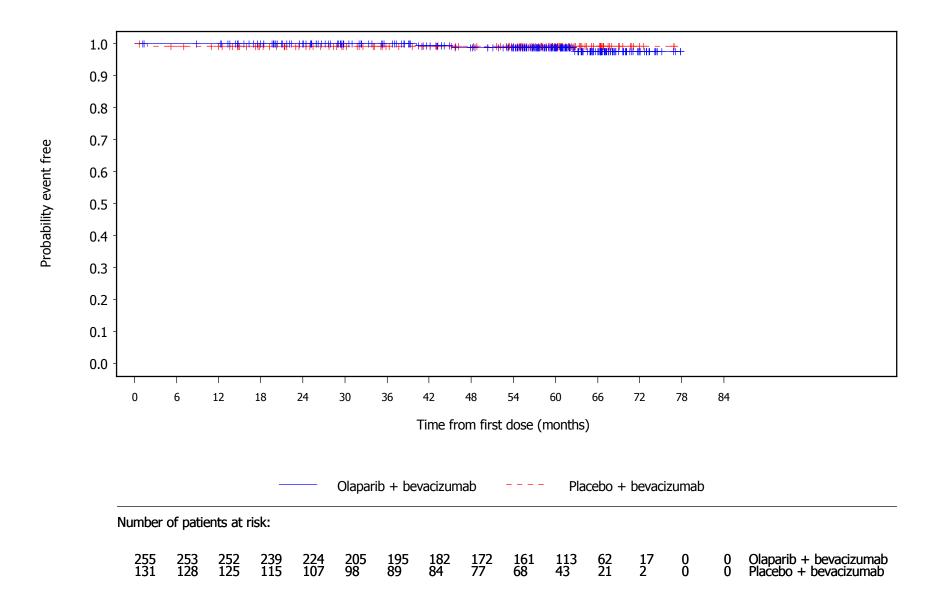


Figure 3.3.70 PAOLA1: Kaplan-Meier plot of time to first occurrence of AESI G1-2: Pneumonitis Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

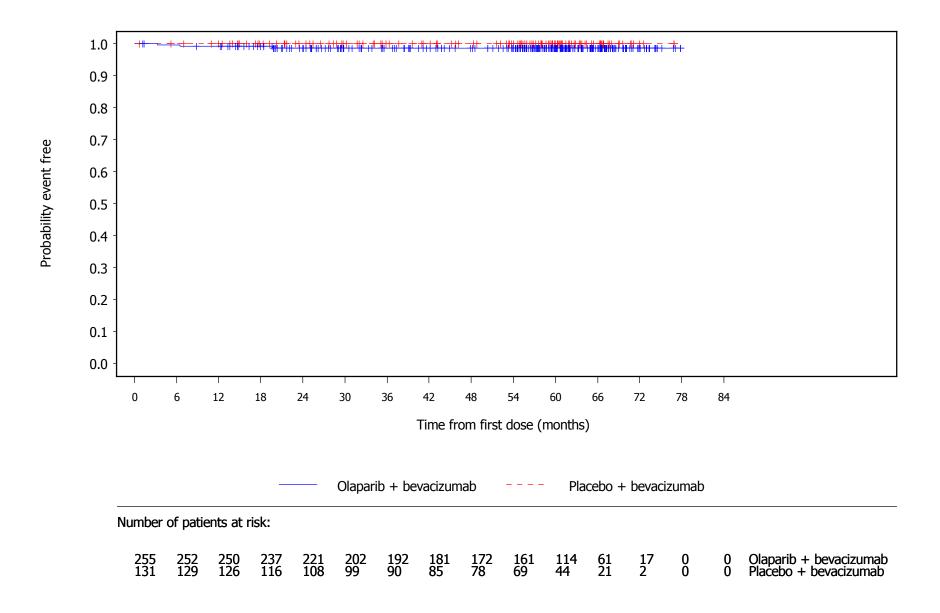


Table 3.4.1 PAOLA1: Summary of subgroup analysis of AESI: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	:	Olaparib + b (N=2		ab		Placebo + be (N=1)		ıb	:		2-sided p-value [b]
Subgroup		Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	Median (95% (month)	CI)	Hazard ratio [b]	95% CI [b]	
First line treatment out	come (IV	RS)									
NED [PDS]	92	37 (40.2)	NE (N	IE, NE)	48	8 (16.7)	NE (NE, NE)	2.92	1.43, 6.75	0.0023*
NED/CR [IDS]	74	34 (45.9)	NE (N	IE, NE)	38	0	NE (NE, NE)	NC	NC	NC
NED/CR [Chemo]	40	13 (32.5)	NE (N	IE, NE)	20	1 (5.0)	NE (NE, NE)	7.23	1.44,131.47	0.0117*
PR	49	18 (36.7)	NE (N	IE, NE)	25	3 (12.0)	NE (NE, NE)	3.81	1.29, 16.25	0.0132*
Interaction p-value											0.6593
Screening laboratory tBR0	CA statu	s (IVRS)									
tBRCAm	150	55 (36.7)	NE (N	IE, NE)	65	7 (10.8)	NE (NE, NE)	4.01	1.95, 9.66	<0.0001*
non-tBRCAm	105	47 (44.8)	NE (N	IE, NE)	66	5 (7.6)	NE (NE, NE)	7.79	3.41, 22.45	<0.0001*
Interaction p-value											0.2788
First line treatment out	come (eC	RF)									
NED [PDS]	89	35 (39.3)	NE (N	IE, NE)	47	8 (17.0)	NE (NE, NE)	2.78	1.36, 6.46	0.0041*
NED/CR [IDS]	74	32 (43.2)	NE (N	IE, NE)	32	0	NE (NE, NE)	NC	NC	NC
NED/CR [Chemo]	39	11 (28.2)	NE (N	IE, NE)	17	1 (5.9)	NE (NE, NE)	5.35	1.04, 97.72	0.0438*
PR	50	22 (44.0)	NE (N	IE, NE)	34	3 (8.8)	NE (NE, NE)	6.28	2.18, 26.54	0.0002*
Interaction p-value											0.4810
Screening laboratory tBR0	CA statu	s (eCRF)									
tBRCAm	147	55 (37.4)	NE (N	IE, NE)	67	7 (10.4)	NE (NE, NE)	4.24	2.07, 10.22	<0.0001*
non-tBRCAm	108	47 (43.5)	NE (1	IE, NE)	64	5 (7.8)	NE (NE, NE)	7.28	3.19, 20.98	<0.0001*
Interaction p-value											0.3785
Age group											
<65 years	185	70 (37.8)	NE (N	IE, NE)	98	9 (9.2)	NE (NE, NE)	4.94	2.60, 10.61	<0.0001*
>=65 years	70	32 (45.7)	NE (1	IE, NE)	33	3 (9.1)	NE (NE, NE)	6.84	2.45, 28.45	<0.0001*
Interaction p-value											0.6348

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaa 11AUG2022:11:30 kpzx329

Table 3.4.1 PAOLA1: Summary of subgroup analysis of AESI: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + (N=	bevacizuma 255)	ıb		Placebo + bo (N=1					
Subgroup		Number (%) of patients with events	Median (95% (months	CI)		Number (%) of patients with events	Median t (95% CI (months)	:)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	73 (40.1)	NE (N	E, NE)	89	9 (10.1)	NE (NE,	NE)	4.90	2.59, 10.51	<0.0001*
IV	73	29 (39.7)	NE (N	E, NE)	42	3 (7.1)	NE (NE,	NE)	6.93	2.46, 28.94	<0.0001*
Interaction p-value											0.6129
Region											
Europe	245	96 (39.2)	NE (N	E, NE)	125	12 (9.6)	NE (NE,	NE)	5.00	2.86, 9.61	<0.0001*
Japan	10	6 (60.0)	2.8 (0.	3, NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	74 (38.9)	NE (N	E, NE)	100	8 (8.0)	NE (NE,	NE)	6.10	3.13, 13.75	<0.0001*
(1) Restricted activity	61	28 (45.9)	NE (N	E, NE)	30	4 (13.3)	NE (NE,	NE)	4.17	1.63, 14.08	0.0016*
Interaction p-value											0.5651
Baseline CA-125 value											
<=ULN	228	89 (39.0)	NE (N	E, NE)	117	11 (9.4)	NE (NE,	NE)	5.09	2.84, 10.08	<0.0001*
>ULN	27	13 (48.1)	NE (N	E, NE)	14	1 (7.1)	NE (NE,	NE)	9.22	1.84,167.51	0.0036*
Interaction p-value											0.5569
Histological grade											
High grade	255	102 (40.0)	NE (N	E, NE)	131	12 (9.2)	NE (NE,	NE)	5.41	3.10, 10.37	<0.0001*
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	69 (41.6)	NE (N	E, NE)	80	8 (10.0)	NE (NE,	NE)	5.29	2.70, 11.93	<0.0001*
Residue	79	28 (35.4)	NE (N	E, NE)	43	2 (4.7)	NE (NE,	NE)	8.99	2.71, 55.75	<0.0001*
Interaction p-value											0.4989

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaa 11AUG2022:11:30 kpzx329

Table 3.4.1 PAOLA1: Summary of subgroup analysis of AESI: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		Placebo + b (N=1			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	52 (35.6)	NE (NE, NE)	78 10 (12.8)	NE (NE, NE)	3.25 1.73, 6.80	0.0001*
Interval	99	45 (45.5)	NE (NE, NE)	45 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Myriad tumour BRCA mutat:	ion stat	us					
tBRCAm	158	63 (39.9)	NE (NE, NE)	77 6 (7.8)	NE (NE, NE)	6.25 2.94, 16.18	<0.0001*
Non-tBRCAm	97	39 (40.2)	NE (NE, NE)	54 6 (11.1)	NE (NE, NE)	4.57 2.09, 12.03	<0.0001*
Interaction p-value							0.6098
Status somatic BRCA muta	tions						
sBRCAm	25	10 (40.0)	NE (NE, NE)	9 0	NE (NE, NE)	NC NC	NC
gBRCAm	69	27 (39.1)	NE (NE, NE)	36 4 (11.1)	NE (NE, NE)	4.37 1.71, 14.79	0.0012*
Non-BRCAm	43	20 (46.5)	NE (NE, NE)	23 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaa 11AUG2022:11:30 kpzx329

Table 3.4.2 PAOLA1: Summary of subgroup analysis of AESI: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ımab			Placebo + b (N=1		mab					2-sided p-value [b]
Subgroup		Number (%) of patients with events		an ti % CI; hs) [)		Number (%) of patients with events	(95	an tir 5% CI) :hs) [Hazard ratio [b]	95% CI	[b]	
First line treatment out	come (IV)	RS)												
NED [PDS]	92	16 (17.4)	NE (NE,	NE)	48	8 (16.7)	NE (NE,	NE)	1.01	0.44,	2.49	0.9840
NED/CR [IDS]	74	21 (28.4)	NE (NE,	NE)	38	7 (18.4)	NE (NE,	NE)	1.57	0.70,	3.98	0.2872
NED/CR [Chemo]	40	9 (22.5)	NE (NE,	NE)	20	5 (25.0)	NE (NE,	NE)	0.89	0.31,	2.91	0.8421
PR	49	7 (14.3)	NE (NE,	NE)	25	2 (8.0)	NE (NE,	NE)	1.89	0.46,	12.66	0.4036
Interaction p-value														0.7661
Screening laboratory tBR	CA statu	s (IVRS)												
tBRCAm	150	29 (19.3)	NE (NE,	NE)	65	11 (16.9)	NE (NE,	NE)	1.13	0.58,	2.36	0.7326
non-tBRCAm	105	24 (22.9)	NE (NE,	NE)	66	11 (16.7)	NE (NE,	NE)	1.41	0.71,	2.99	0.3388
Interaction p-value														0.6626
First line treatment out	come (eCl	RF)												
NED [PDS]	89	19 (21.3)	NE (NE,	NE)	47	7 (14.9)	NE (NE,	NE)	1.44	0.63,	3.70	0.3940
NED/CR [IDS]	74	19 (25.7)	NE (NE,	NE)	32	6 (18.8)	NE (NE,	NE)	1.37	0.58,	3.77	0.4870
NED/CR [Chemo]	39	5 (12.8)	NE (NE,	NE)	17	4 (23.5)	NE (NE,	NE)	0.51	0.13,	2.04	0.3195
PR	50	10 (20.0)	NE (NE,	NE)	34	5 (14.7)	NE (NE,	NE)	1.41	0.50,	4.53	0.5225
Interaction p-value														0.5923
Screening laboratory tBR	CA statu	s (eCRF)												
tBRCAm	147	29 (19.7)	NE (NE,	NE)	67	12 (17.9)	NE (NE,	NE)	1.09	0.57,	2.21	0.8091
non-tBRCAm	108	24 (22.2)	NE (NE,	NE)	64	10 (15.6)	NE (NE,	NE)	1.46	0.72,	3.19	0.3055
Interaction p-value														0.5630
Age group														
<65 years	185	38 (20.5)	NE (NE,	NE)	98	17 (17.3)	NE (NE,	NE)	1.20	0.69,	2.18	0.5295
>=65 years	70	15 (21.4)	NE (NE,	NE)	33	5 (15.2)	NE (NE,	NE)	1.37	0.53,	4.23	0.5275
Interaction p-value														0.8167

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaab 11AUG2022:11:30 kpzx329

Table 3.4.2 PAOLA1: Summary of subgroup analysis of AESI: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + k (N=2				Placebo + be (N=1)						
Subgroup		Number (%) of patients with events	Median t (95% CI (months)	:)		Number (%) of patients with events	Median t (95% CI (months)	()	Hazard ratio [b]	-	[b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	38 (20.9)	NE (NE,	NE)	89	14 (15.7)	NE (NE,	NE)	1.36	0.75,	2.59	0.3201
IV	73	15 (20.5)	NE (NE,	NE)	42	8 (19.0)	NE (NE,	NE)	1.04	0.45,	2.57	0.9349
Interaction p-value												0.6194
Region												
Europe	245	46 (18.8)	NE (NE,	NE)	125	21 (16.8)	NE (NE,	NE)	1.10	0.67,	1.88	0.7108
Japan	10	7 (70.0)	2.1 (0.5,	NE)	6	1 (16.7)	NE (NE,	NE)	6.62	1.17,123	3.99	0.0299*
Interaction p-value												0.0536
ECOG performance status at	Basel	ine										
(0) Normal activity	190	40 (21.1)	NE (NE,	NE)	100	15 (15.0)	NE (NE,	NE)	1.41	0.80,	2.64	0.2444
(1) Restricted activity	61	12 (19.7)	NE (NE,	NE)	30	7 (23.3)	NE (NE,	NE)	0.83	0.33,	2.22	0.6901
Interaction p-value												0.3477
Baseline CA-125 value												
<=ULN	228	48 (21.1)	NE (NE,	,	117	19 (16.2)	NE (NE,	NE)	1.31	•	2.28	0.3139
>ULN	27	5 (18.5)	NE (NE,	NE)	14	3 (21.4)	NE (NE,	NE)	0.82	0.20,	4.00	0.7887
Interaction p-value												0.5570
Histological grade												
High grade	255	53 (20.8)	NE (NE,	NE)	131	22 (16.8)	NE (NE,	NE)	1.24	0.77,	2.08	0.3894
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	38 (22.9)	NE (NE,	NE)	80	13 (16.3)	NE (NE,	NE)	1.42	0.78,	2.77	0.2628
Residue	79	12 (15.2)	NE (NE,	NE)	43	8 (18.6)	NE (NE,	NE)	0.79	0.33,	2.03	0.6175
Interaction p-value												0.3016

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaab 11AUG2022:11:30 kpzx329

Table 3.4.2 PAOLA1: Summary of subgroup analysis of AESI: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		Placebo + k (N=1			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	27 (18.5)	NE (NE, NE)	78 12 (15.4)	NE (NE, NE)	1.21 0.63, 2.47	0.5853
Interval	99	23 (23.2)	NE (NE, NE)	45 9 (20.0)	NE (NE, NE)	1.14 0.54, 2.59	0.7425
Interaction p-value							0.9103
Myriad tumour BRCA mutat:	ion stat	us					
tBRCAm	158	31 (19.6)	NE (NE, NE)	77 12 (15.6)	NE (NE, NE)	1.24 0.65, 2.52	0.5178
Non-tBRCAm	97	22 (22.7)	NE (NE, NE)	54 10 (18.5)	NE (NE, NE)	1.26 0.61, 2.78	0.5383
Interaction p-value							0.9766
Status somatic BRCA muta	cions						
sBRCAm	25	8 (32.0)	NE (NE, NE)	9 1 (11.1)	NE (NE, NE)	2.80 0.51, 51.90	0.2692
gBRCAm	69	12 (17.4)	NE (NE, NE)	36 8 (22.2)	NE (NE, NE)	0.77 0.32, 1.97	0.5767
Non-BRCAm	43	14 (32.6)	NE (NE, NE)	23 4 (17.4)	NE (NE, NE)	2.01 0.72, 7.10	0.1913
Interaction p-value							0.2846

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaab 11AUG2022:11:30 kpzx329

Table 3.4.3 PAOLA1: Summary of subgroup analysis of AESI: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		:		Placebo + be (N=1)					2-sided p-value [b]
Subgroup	(Number (%) of patients with events	Median to (95% CI (months))	C	Number (%) of patients with events	Median (95% ((months)	I)	Hazard ratio [b]	95% CI [b]	
First line treatment out	come (IVF	RS)									
NED [PDS]	92	7 (7.6)	NE (NE,	NE)	48	1 (2.1)	NE (NE	, NE)	3.47	0.62, 64.78	0.1783
NED/CR [IDS]	74	7 (9.5)	NE (NE,	NE)	38	6 (15.8)	NE (NE	, NE)	0.59	0.20, 1.83	0.3487
NED/CR [Chemo]	40	3 (7.5)	NE (NE,	NE)	20	0	NE (NE	, NE)	NC	NC	NC
PR	49	1 (2.0)	NE (NE,	NE)	25	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											0.1009
Screening laboratory tBR0	CA status	s (IVRS)									
tBRCAm	150	9 (6.0)	NE (NE,	NE)	65	4 (6.2)	NE (NE	, NE)	0.96	0.31, 3.53	0.9400
non-tBRCAm	105	9 (8.6)	NE (NE,	NE)	66	3 (4.5)	NE (NE	, NE)	1.88	0.56, 8.46	0.3222
Interaction p-value											0.4497
First line treatment out	come (eCF	RF)									
NED [PDS]	89	5 (5.6)	NE (NE,	NE)	47	1 (2.1)	NE (NE	, NE)	2.49	0.40, 47.65	0.3608
NED/CR [IDS]	74	9 (12.2)	NE (NE,	NE)	32	5 (15.6)	NE (NE	, NE)	0.77	0.27, 2.52	0.6511
NED/CR [Chemo]	39	2 (5.1)	NE (NE,	NE)	17	0	NE (NE	, NE)	NC	NC	NC
PR	50	2 (4.0)	NE (NE,	NE)	34	1 (2.9)	NE (NE	, NE)	1.35	0.13, 29.09	0.8021
Interaction p-value											0.5863
Screening laboratory tBR0	CA status	s (eCRF)									
tBRCAm	147	9 (6.1)	NE (NE,	NE)	67	5 (7.5)	NE (NE	, NE)	0.80	0.28, 2.59	0.6876
non-tBRCAm	108	9 (8.3)	NE (NE,	NE)	64	2 (3.1)	NE (NE	, NE)	2.68	0.69, 17.55	0.1666
Interaction p-value											0.1906
Age group											
<65 years	185	12 (6.5)	NE (NE,	NE)	98	6 (6.1)	NE (NE	, NE)	1.04	0.40, 2.98	0.9397
>=65 years	70	6 (8.6)	NE (NE,	NE)	33	1 (3.0)	NE (NE	, NE)	2.82	0.48, 53.22	0.2818
Interaction p-value											0.3687

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaac 11AUG2022:11:30 kpzx329

Table 3.4.3 PAOLA1: Summary of subgroup analysis of AESI: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=25		ımab			Placebo + be (N=13		mab	•				
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI)		C	Number (%) of patients with events	(95	an tim % CI) hs) [a		Hazard ratio [b]	ratio	2-sided p-value [b]	
FIGO Stage (Disease state)														
III	182	15 (8.2)	NE (NE,	NE)	89	5 (5.6)	NE (NE,	NE)	1.45	0.56,	4.45	0.4610
IV	73	3 (4.1)	NE (NE,	NE)	42	2 (4.8)	NE (NE,	NE)	0.85	0.14,	6.44	0.8579
Interaction p-value														0.6141
Region														
Europe	245	17 (6.9)	NE (NE,	NE)	125	7 (5.6)	NE (NE,	NE)	1.21	0.52,	3.14	0.6609
Japan	10	1 (10.0)	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
ECOG performance status at	Basel	ine												
(0) Normal activity	190	13 (6.8)	NE (NE,	NE)	100	5 (5.0)	NE (NE,	NE)	1.35	0.51,	4.21	0.5583
(1) Restricted activity	61	5 (8.2)	NE (NE,	NE)	30	2 (6.7)	NE (NE,	NE)	1.20	0.26,	8.36	0.8281
Interaction p-value														0.9020
Baseline CA-125 value														
<=ULN	228	17 (7.5)	NE (NE,	NE)	117	7 (6.0)	NE (NE,	NE)	1.21	0.52,	3.14	0.6619
>ULN	27	1 (3.7)	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
Histological grade														
High grade	255	18 (7.1)	NE (NE,	NE)	131	7 (5.3)	NE (NE,	NE)	1.30	0.57,	3.35	0.5484
Interaction p-value														NC
Cytoreductive surgery outco	me													
No residue	166	14 (8.4)	NE (NE,	NE)	80	6 (7.5)	NE (NE,	NE)	1.09	0.44,	3.09	0.8549
Residue	79	3 (3.8)	NE (NE,	NE)	43	1 (2.3)	NE (NE,	NE)	1.62	0.21,	32.76	0.6641
Interaction p-value														0.7478

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaac 11AUG2022:11:30 kpzx329

Table 3.4.3 PAOLA1: Summary of subgroup analysis of AESI: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2			Placebo + be (N=1)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	(Number (%) of patients with events	Median time (95% CI) (months) [a]	— Hazard ratio [b]	ratio	
Timing of cytoreductive s	surgery								
Upfront	146	7 (4.8)	NE (NE, NE)	78	1 (1.3)	NE (NE, N	E) 3.57	0.63, 66.67	0.1672
Interval	99	10 (10.1)	NE (NE, NE)	45	6 (13.3)	NE (NE, N	E) 0.77	0.28, 2.25	0.6087
Interaction p-value									0.1507
Myriad tumour BRCA mutati	on stat	us							
tBRCAm	158	10 (6.3)	NE (NE, NE)	77	6 (7.8)	NE (NE, N	E) 0.78	0.29, 2.29	0.6312
Non-tBRCAm	97	8 (8.2)	NE (NE, NE)	54	1 (1.9)	NE (NE, N	E) 4.58	0.84, 84.87	0.0844
Interaction p-value									0.0906
Status somatic BRCA mutat	ions								
sBRCAm	25	2 (8.0)	NE (NE, NE)	9	1 (11.1)	NE (NE, N	E) 0.69	0.07, 14.86	0.7676
gBRCAm	69	5 (7.2)	NE (NE, NE)	36	5 (13.9)	NE (NE, N	E) 0.50	0.14, 1.79	0.2751
Non-BRCAm	43	5 (11.6)	NE (NE, NE)	23	1 (4.3)	NE (NE, N	E) 2.60	0.42, 49.84	0.3370
Interaction p-value									0.3626

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaac 11AUG2022:11:30 kpzx329

Table 3.4.4 PAOLA1: Summary of subgroup analysis of AESI: Nausea Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	:	Olaparib + N=2			Placebo + be (N=1)		lb	:		
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IV	RS)								
NED [PDS]	92	52 (56.5)	3.7 (0.6, NE)	48	11 (22.9)	NE (1	NE, NE)	3.30	1.79, 6.68	<0.0001*
NED/CR [IDS]	74	48 (64.9)	1.3 (0.3, 4.3)	38	15 (39.5)	NE (1	NE, NE)	2.15	1.23, 3.98	0.0061*
NED/CR [Chemo]	40	21 (52.5)	0.6 (0.2, NE)	20	2 (10.0)	NE (1	NE, NE)	8.02	2.35, 50.17	0.0002*
PR	49	23 (46.9)	NE (NE, NE)	25	6 (24.0)	NE (1	NE, NE)	2.49	1.08, 6.74	0.0315*
Interaction p-value										0.2920
Screening laboratory tBRO	CA statu	ıs (IVRS)								
tBRCAm	150	87 (58.0)	2.6 (0.7,19.7)	65	14 (21.5)	NE (1	NE, NE)	3.69	2.17, 6.78	<0.0001*
non-tBRCAm	105	57 (54.3)	3.3 (0.3, NE)	66	20 (30.3)	NE (1	NE, NE)	2.35	1.44, 4.01	0.0005*
Interaction p-value										0.2420
First line treatment outo	come (eC	RF)								
NED [PDS]	89	51 (57.3)	2.7 (0.5, NE)	47	10 (21.3)	NE (1	NE, NE)	3.65	1.94, 7.64	<0.0001*
NED/CR [IDS]	74	49 (66.2)	1.0 (0.3, 3.4)	32	12 (37.5)	NE (1	NE, NE)	2.45	1.35, 4.82	0.0026*
NED/CR [Chemo]	39	19 (48.7)	NE (NE, NE)	17	2 (11.8)	NE (1	NE, NE)	6.22	1.80, 39.02	0.0019*
PR	50	24 (48.0)	NE (NE, NE)	34	9 (26.5)	NE (1	NE, NE)	2.15	1.03, 4.89	0.0399*
Interaction p-value										0.4619
Screening laboratory tBRO	CA statu	s (eCRF)								
tBRCAm	147	85 (57.8)	2.9 (0.8,19.7)	67	15 (22.4)	NE (1	NE, NE)	3.51	2.09, 6.32	<0.0001*
non-tBRCAm	108	59 (54.6)	2.7 (0.3, NE)	64	19 (29.7)	NE (1	NE, NE)	2.44	1.48, 4.21	0.0003*
Interaction p-value										0.3452
Age group										
<65 years	185	109 (58.9)	2.1 (0.6, 9.8)	98	24 (24.5)	NE (1	NE, NE)	3.32	2.17, 5.29	<0.0001*
>=65 years	70	35 (50.0)	19.7 (0.8, NE)	33	10 (30.3)	NE (1	NE, NE)	2.07	1.07, 4.42	0.0307*
Interaction p-value										0.2767

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaad 11AUG2022:11:30 kpzx329

Table 3.4.4 PAOLA1: Summary of subgroup analysis of AESI: Nausea Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + k			Placebo + be (N=1)			:			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median (95% C (months)	I)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	100 (54.9)	3.7 (0.8, NE)	89	28 (31.5)	NE (NE	, NE)	2.27	1.51, 3	.51	<0.0001*
IV	73	44 (60.3)	2.1 (0.3, NE)	42	6 (14.3)	NE (NE	, NE)	6.09	2.80, 15	.94	<0.0001*
Interaction p-value											0.0292*
Region											
Europe	245	141 (57.6)	2.6 (0.8,10.3)	125	33 (26.4)	NE (NE	, NE)	2.94	2.04, 4	.38	<0.0001*
Japan	10	3 (30.0)	NE (NE, NE)	6	1 (16.7)	NE (NE	, NE)	2.15	0.28, 43	.49	0.4830
Interaction p-value											0.7942
ECOG performance status at	Basel	ine									
(0) Normal activity	190	113 (59.5)	2.0 (0.5, 5.4)	100	22 (22.0)	NE (NE	, NE)	3.74	2.42, 6	.07	<0.0001*
(1) Restricted activity	61	30 (49.2)	NE (NE, NE)	30	12 (40.0)	NE (NE	, NE)	1.53	0.80, 3	3.10	0.2020
Interaction p-value											0.0351*
Baseline CA-125 value											
<=ULN	228		2.7 (0.8,19.7)	117	30 (25.6)	NE (NE	, NE)	2.95	2.01, 4	.48	<0.0001*
>ULN	27	15 (55.6)	4.1 (0.1, NE)	14	4 (28.6)	NE (NE	, NE)	2.70	0.98, 9	.48	0.0550
Interaction p-value											0.8824
Histological grade											
High grade	255	144 (56.5)	2.9 (0.8,16.0)	131	34 (26.0)	NE (NE	, NE)	2.92	2.04, 4	.32	<0.0001*
Interaction p-value											NC
Cytoreductive surgery outco	me										
No residue	166	101 (60.8)	1.4 (0.4, 5.4)	80	23 (28.8)	NE (NE	, NE)	2.89	1.87, 4	.67	<0.0001*
Residue	79	39 (49.4)	NE (NE, NE)	43	10 (23.3)	NE (NE	, NE)	2.74	1.42, 5	.80	0.0020*
Interaction p-value											0.8946

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaad 11AUG2022:11:30 kpzx329

Table 3.4.4 PAOLA1: Summary of subgroup analysis of AESI: Nausea Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + N=2			Placebo + be (N=1)		mab					
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]	Number (%) of patients n with events		Median time (95% CI) (months) [a]		Hazard ratio [b]	95% CI [b]		2-sided p-value [b]	
Timing of cytoreductive s	surgery											
Upfront	146	77 (52.7)	6.6 (0.6, NE)	78	14 (17.9)	NE (NE,	NE)	4.01	2.35,	7.40	<0.0001*
Interval	99	63 (63.6)	1.9 (0.4, 5.0)	45	19 (42.2)	NE (NE,	NE)	1.95	1.19,	3.34	0.0074*
Interaction p-value												0.0626
Myriad tumour BRCA mutat:	ion stat	us										
tBRCAm	158	92 (58.2)	2.3 (0.7,16.0)	77	17 (22.1)	NE (NE,	NE)	3.58	2.19,	6.22	<0.0001*
Non-tBRCAm	97	52 (53.6)	3.4 (0.4, NE)	54	17 (31.5)	NE (NE,	NE)	2.25	1.33,	4.01	0.0021*
Interaction p-value												0.2269
Status somatic BRCA mutat	cions											
sBRCAm	25	16 (64.0)	0.8 (0.1, NE)	9	2 (22.2)	NE (NE,	NE)	3.94	1.12,	24.91	0.0306*
gBRCAm	69	44 (63.8)	2.1 (0.5, 5.4)	36	10 (27.8)	NE (NE,	NE)	3.27	1.71,	6.88	0.0002*
Non-BRCAm	43	23 (53.5)	2.7 (0.2, NE)	23	8 (34.8)	NE (NE,	NE)	2.07	0.97,	4.95	0.0620
Interaction p-value												0.6294

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaad 11AUG2022:11:30 kpzx329

Table 3.4.5 PAOLA1: Summary of subgroup analysis of AESI: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		mab			Placebo + be (N=1		mab					
Subgroup		Number (%) of patients with events	Media (95: (mont)	% CI)			Number (%) of patients with events		an ti % CI hs))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)												
NED [PDS]	92	23 (25.0)	NE (NE,	NE)	48	7 (14.6)	NE (NE,	NE)	1.82	0.82,	4.58	0.1471
NED/CR [IDS]	74	18 (24.3)	NE (NE,	NE)	38	7 (18.4)	NE (NE,	NE)	1.39	0.61,	3.58	0.4470
NED/CR [Chemo]	40	4 (10.0)	NE (NE,	NE)	20	1 (5.0)	NE (NE,	NE)	2.05	0.30,	40.11	0.4922
PR	49	10 (20.4)	NE (NE,	NE)	25	3 (12.0)	NE (NE,	NE)	1.85	0.57,	8.24	0.3266
Interaction p-value														0.9675
Screening laboratory tBRG	CA statu	s (IVRS)												
tBRCAm	150	34 (22.7)	NE (NE,	NE)	65	8 (12.3)	NE (NE,	NE)	1.96	0.95,	4.55	0.0681
non-tBRCAm	105	21 (20.0)	NE (NE,	NE)	66	10 (15.2)	NE (NE,	NE)	1.39	0.67,	3.08	0.3865
Interaction p-value														0.5287
First line treatment out	come (eC	RF)												
NED [PDS]	89	21 (23.6)	NE (NE,	NE)	47	7 (14.9)	NE (NE,	NE)	1.68	0.75,	4.26	0.2183
NED/CR [IDS]	74	17 (23.0)	NE (NE,	NE)	32	5 (15.6)	NE (NE,	NE)	1.59	0.63,	4.83	0.3457
NED/CR [Chemo]	39	5 (12.8)	NE (NE,	NE)	17	2 (11.8)	NE (NE,	NE)	1.14	0.25,	7.96	0.8738
PR	50	10 (20.0)	NE (NE,	NE)	34	4 (11.8)	NE (NE,	NE)	1.75	0.58,	6.37	0.3290
Interaction p-value														0.9793
Screening laboratory tBR0	CA statu	s (eCRF)												
tBRCAm	147	33 (22.4)	NE (NE,	NE)	67	8 (11.9)	NE (NE,	NE)	2.00	0.97,	4.65	0.0608
non-tBRCAm	108	22 (20.4)	NE (NE,	NE)	64	10 (15.6)	NE (NE,	NE)	1.37	0.67,	3.03	0.4001
Interaction p-value														0.4904
Age group														
<65 years	185	44 (23.8)	NE (NE,	NE)	98	12 (12.2)	NE (NE,	NE)	2.11	1.15,	4.18	0.0146*
>=65 years	70	11 (15.7)	NE (NE,	NE)	33	6 (18.2)	NE (NE,	NE)	0.85	0.32,	2.47	0.7543
Interaction p-value														0.1420

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaae 11AUG2022:11:30 kpzx329

Table 3.4.5 PAOLA1: Summary of subgroup analysis of AESI: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ab			bevacizumab 131)		:		
Subgroup		Number (%) of patients with events	Median (95% (months	CI)		Number (%) of patients with events)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	42 (23.1)	NE (1	NE, NE)	89	13 (14.6)	NE (NE,	NE)	1.68	0.93, 3.26	0.0867
IV	73	13 (17.8)	NE (1	NE, NE)	42	5 (11.9)	NE (NE,	NE)	1.55	0.59, 4.84	0.3899
Interaction p-value											0.8942
Region											
Europe	245	55 (22.4)	NE (1	NE, NE)	125	15 (12.0)	NE (NE,	NE)	1.99	1.16, 3.66	0.0119*
Japan	10	0	NE (1	NE, NE)	6	3 (50.0)	45.7 (1.4,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	42 (22.1)	NE (NE, NE)	100	13 (13.0)	NE (NE,	NE)	1.79	0.99, 3.46	0.0556
(1) Restricted activity	61	13 (21.3)	NE (1	NE, NE)	30	5 (16.7)	NE (NE,	NE)	1.39	0.52, 4.33	0.5218
Interaction p-value											0.6864
Baseline CA-125 value											
<=ULN	228	44 (19.3)	NE (1	NE, NE)	117	16 (13.7)	NE (NE,	NE)	1.47	0.85, 2.69	0.1732
>ULN	27	11 (40.7)	NE (1	NE, NE)	14	2 (14.3)	NE (NE,	NE)	3.41	0.91, 22.03	0.0698
Interaction p-value											0.2754
Histological grade											
High grade	255	55 (21.6)	NE (NE, NE)	131	18 (13.7)	NE (NE,	NE)	1.66	0.99, 2.90	0.0524
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	40 (24.1)	NE (1	NE, NE)	80	12 (15.0)	NE (NE,	NE)	1.72	0.93, 3.43	0.0841
Residue	79	13 (16.5)	NE (1	NE, NE)	43	6 (14.0)	NE (NE,	NE)	1.19	0.47, 3.40	0.7161
Interaction p-value											0.5409

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaae 11AUG2022:11:30 kpzx329

Table 3.4.5 PAOLA1: Summary of subgroup analysis of AESI: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2			Placebo + be (N=1)						
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median to (95% CI (months))	Hazard ratio [b]	95% C	I [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	30 (20.5)	NE (NE, NE)	78	9 (11.5)	NE (NE,	NE)	1.88	0.93,	4.20	0.0808
Interval	99	23 (23.2)	NE (NE, NE)	45	9 (20.0)	NE (NE,	NE)	1.21	0.58,	2.77	0.6173
Interaction p-value											0.4251
Myriad tumour BRCA mutat:	ion stat	us									
tBRCAm	158	36 (22.8)	NE (NE, NE)	77	9 (11.7)	NE (NE,	NE)	2.08	1.05,	4.60	0.0357*
Non-tBRCAm	97	19 (19.6)	NE (NE, NE)	54	9 (16.7)	NE (NE,	NE)	1.23	0.57,	2.84	0.6112
Interaction p-value											0.3368
Status somatic BRCA muta	tions										
sBRCAm	25	7 (28.0)	NE (NE, NE)	9	1 (11.1)	NE (NE,	NE)	2.77	0.49,	51.69	0.2823
gBRCAm	69	16 (23.2)	NE (NE, NE)	36	5 (13.9)	NE (NE,	NE)	1.84	0.72,	5.63	0.2110
Non-BRCAm	43	6 (14.0)	NE (NE, NE)	23	5 (21.7)	NE (NE,	NE)	0.61	0.18,	2.11	0.4172
Interaction p-value											0.2762

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaae 11AUG2022:11:30 kpzx329

Table 3.4.6 PAOLA1: Summary of subgroup analysis of AESI: Fatigue and Asthenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + 1			Placebo + be (N=1)			:			
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median t (95% Ci (months)	[)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)									
NED [PDS]	92	49 (53.3)	12.7 (2.9, NE)	48	18 (37.5)	NE (NE	, NE)	1.69	1.01,	2.99	0.0471*
NED/CR [IDS]	74	50 (67.6)	3.5 (1.8,11.8)	38	15 (39.5)	NE (NE	, NE)	2.06	1.18,	3.79	0.0096*
NED/CR [Chemo]	40	19 (47.5)	NE (NE, NE)	20	5 (25.0)	NE (NE	, NE)	2.34	0.94,	7.05	0.0690
PR	49	24 (49.0)	14.9 (3.5, NE)	25	9 (36.0)	NE (NE	, NE)	1.57	0.76,	3.57	0.2348
Interaction p-value											0.8897
Screening laboratory tBRG	CA statu	s (IVRS)									
tBRCAm	150	82 (54.7)	9.8 (3.5, NE)	65	25 (38.5)	NE (NE	, NE)	1.62	1.05,	2.59	0.0276*
non-tBRCAm	105	60 (57.1)	11.0 (2.7, NE)	66	22 (33.3)	NE (NE	, NE)	2.14	1.33,	3.56	0.0013*
Interaction p-value											0.4121
First line treatment out	come (eC	RF)									
NED [PDS]	89	48 (53.9)	12.5 (2.8, NE)	47	18 (38.3)	NE (NE	, NE)	1.67	0.99,	2.94	0.0552
NED/CR [IDS]	74	49 (66.2)	3.5 (1.4,11.0)	32	9 (28.1)	NE (NE	, NE)	3.30	1.70,	7.18	0.0002*
NED/CR [Chemo]	39	18 (46.2)	NE (NE, NE)	17	4 (23.5)	NE (NE	, NE)	2.40	0.90,	8.32	0.0839
PR	50	26 (52.0)	15.3 (3.5, NE)	34	16 (47.1)	NE (NE	, NE)	1.09	0.59,	2.07	0.7935
Interaction p-value											0.1155
Screening laboratory tBR0	CA statu	s (eCRF)									
tBRCAm	147	80 (54.4)	11.8 (3.5, NE)	67	25 (37.3)	NE (NE	, NE)	1.68	1.09,	2.68	0.0189*
non-tBRCAm	108	62 (57.4)	11.0 (2.7, NE)	64	22 (34.4)	NE (NE	, NE)	2.07	1.30,	3.45	0.0020*
Interaction p-value											0.5283
Age group											
<65 years	185	105 (56.8)	9.8 (2.9,30.1)	98	32 (32.7)	NE (NE	, NE)	2.13	1.45,	3.22	<0.0001*
>=65 years	70	37 (52.9)	11.8 (3.5, NE)	33	15 (45.5)	NE (NE	, NE)	1.27	0.71,	2.39	0.4252
Interaction p-value											0.1650

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaf 11AUG2022:11:30 kpzx329

Table 3.4.6 PAOLA1: Summary of subgroup analysis of AESI: Fatigue and Asthenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olapa	arib + (N=2	bevaci 255)	zumab			Placebo (+ be N=13		mab	•				
Subgroup	n	of pa	er (%) atients events	(9	lian ti 95% CI nths))		Number (of patier with ever	nts	(95	an ti % CI hs)		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)																
III	182	100	(54.9)	11.1	(3.7,	NE)	89	32 (36.	0)	NE (NE,	NE)	1.77	1.20,	2.68	0.0032*
IV	73	42	(57.5)	11.0	(2.1,	NE)	42	15 (35.	7)	NE (NE,	NE)	2.05	1.16,	3.82	0.0121*
Interaction p-value																0.6848
Region																
Europe	245	142	(58.0)	7.2	(3.5,	15.3)	125	47 (37.	6)	NE (NE,	NE)	1.85	1.34,	2.61	0.0001*
Japan	10	0		NE	(NE,	NE)	6	0		NE (NE,	NE)	NC	NC		NC
Interaction p-value																NC
ECOG performance status at	Basel	ine														
(0) Normal activity	190	106	(55.8)	8.5	(3.5,	NE)	100	33 (33.	0)	NE (NE,	NE)	2.03	1.39,	3.05	0.0002*
(1) Restricted activity	61	32	(52.5)	18.0	(2.9,	NE)	30	13 (43.	3)	NE (NE,	NE)	1.37	0.74,	2.71	0.3282
Interaction p-value																0.3119
Baseline CA-125 value																
<=ULN	228		(55.3)				117	41 (35.		NE (NE,	NE)	1.86	1.32,		0.0003*
>ULN	27	16	(59.3)	2.1	(0.5,	NE)	14	6 (42.	9)	NE (NE,	NE)	1.82	0.75,	5.07	0.1933
Interaction p-value																0.9656
Histological grade																
High grade	255	142	(55.7)	11.0	(4.1,	30.1)	131	47 (35.	9)	NE (NE,	NE)	1.85	1.34,	2.60	0.0001*
Interaction p-value																NC
Cytoreductive surgery outco	ome															
No residue	166	98	(59.0)	6.2	(2.5,	15.9)	80	27 (33.	8)	NE (NE,	NE)	2.22	1.47,	3.46	<0.0001*
Residue	79	39	(49.4)	18.0	(3.7,	NE)	43	15 (34.	9)	NE (NE,	NE)	1.52	0.85,	2.84	0.1580
Interaction p-value																0.3144

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaf 11AUG2022:11:30 kpzx329

Table 3.4.6 PAOLA1: Summary of subgroup analysis of AESI: Fatigue and Asthenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + k (N=2			Placebo + k (N=1	oevacizumab 131)					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median t: (95% CI (months))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	75 (51.4)	14.9 (5.1, NE)	78	25 (32.1)	NE (NE,	NE)	1.90	1.23,	3.05	0.0035*
Interval	99	62 (62.6)	4.1 (2.1,15.3)	45	17 (37.8)	NE (NE,	NE)	2.00	1.20,	3.52	0.0074*
Interaction p-value											0.8947
Myriad tumour BRCA mutati	ion stat	us									
tBRCAm	158	87 (55.1)	9.8 (3.5, NE)	77	28 (36.4)	NE (NE,	NE)	1.74	1.15,	2.70	0.0081*
Non-tBRCAm	97	55 (56.7)	11.0 (2.7, NE)	54	19 (35.2)	NE (NE,	NE)	2.03	1.23,	3.51	0.0052*
Interaction p-value											0.6483
Status somatic BRCA mutat	cions										
sBRCAm	25	13 (52.0)	15.8 (2.1, NE)	9	0	NE (NE,	NE)	NC	NC		NC
gBRCAm	69	48 (69.6)	2.1 (1.0, 6.9)	36	18 (50.0)	36.5 (4.4,	NE)	1.87	1.11,	3.30	0.0189*
Non-BRCAm	43	25 (58.1)	6.0 (1.4, NE)	23	12 (52.2)	10.4 (2.9,	NE)	1.28	0.65,	2.63	0.4830
Interaction p-value											0.3969

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaf 11AUG2022:11:30 kpzx329

Table 3.4.7 PAOLA1: Summary of subgroup analysis of AESI: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	;	Olaparib + : (N=2	bevacizumab 255)	:-			bevacizumab 131)				
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IVI	RS)									
NED [PDS]	92	46 (50.0)	17.9 (7.6,	NE)	48	26 (54.2)	11.0 (4.2, NE)	0.93	0.58,	1.52	0.7607
NED/CR [IDS]	74	36 (48.6)	30.5 (8.3,	NE)	38	22 (57.9)	5.2 (2.1, NE)	0.70	0.42,	1.21	0.2016
NED/CR [Chemo]	40	21 (52.5)	13.9 (3.4,	NE)	20	13 (65.0)	4.1 (0.8, NE)	0.72	0.36,	1.47	0.3518
PR	49	24 (49.0)	NE (NE,	NE)	25	17 (68.0)	2.8 (0.7,11.3)	0.54	0.29,	1.03	0.0613
Interaction p-value											0.6093
Screening laboratory tBR0	CA status	s (IVRS)									
tBRCAm	150	70 (46.7)	NE (NE,	NE)	65	40 (61.5)	6.9 (2.9,24.3)	0.66	0.45,	0.98	0.0415*
non-tBRCAm	105	57 (54.3)	11.1 (6.6,	NE)	66	38 (57.6)	4.6 (2.2, NE)	0.86	0.58,	1.31	0.4903
Interaction p-value											0.3532
First line treatment out	come (eCI	RF)									
NED [PDS]	89	44 (49.4)	NE (NE,	NE)	47	25 (53.2)	11.0 (4.2, NE)	0.94	0.58,	1.55	0.7981
NED/CR [IDS]	74	33 (44.6)	NE (NE,	NE)	32	17 (53.1)	9.8 (2.8, NE)	0.73	0.41,	1.34	0.3008
NED/CR [Chemo]	39	23 (59.0)	5.6 (0.9,	NE)	17	12 (70.6)	4.1 (0.8, NE)	0.84	0.43,	1.75	0.6372
PR	50	26 (52.0)	16.6 (5.7,	NE)	34	23 (67.6)	2.5 (0.7,11.3)	0.53	0.30,	0.94	0.0311*
Interaction p-value											0.5104
Screening laboratory tBR0	CA status	s (eCRF)									
tBRCAm	147	68 (46.3)	NE (NE,	NE)	67	40 (59.7)	6.9 (3.3, NE)	0.68	0.47,	1.02	0.0620
non-tBRCAm	108	59 (54.6)	10.6 (6.7,	NE)	64	38 (59.4)	4.3 (2.1, NE)	0.83	0.56,	1.26	0.3797
Interaction p-value											0.4984
Age group											
<65 years	185	83 (44.9)	NE (NE,	NE)	98	54 (55.1)	8.4 (4.1, NE)	0.72	0.52,	1.03	0.0693
>=65 years	70	44 (62.9)	5.8 (3.4,	14.1)	33	24 (72.7)	2.8 (1.4, 5.4)	0.75	0.46,	1.25	0.2602
Interaction p-value											0.9181

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaag 11AUG2022:11:30 kpzx329

Table 3.4.7 PAOLA1: Summary of subgroup analysis of AESI: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		-	bevacizumab 255)			Placebo + k (N=1				
Subgroup	n	Number (%) of patients with events)		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	90 (49.5)	NE (NE,	NE)	89	53 (59.6)	7.0 (4.1,24.3)	0.75	0.54, 1.06	0.1058
IV	73	37 (50.7)	16.7 (5.6,	NE)	42	25 (59.5)	3.5 (2.1, NE)	0.73	0.44, 1.22	0.2266
Interaction p-value										0.9158
Region										
Europe	245	124 (50.6)	17.9 (9.7,	NE)	125	72 (57.6)	5.6 (3.5, NE)	0.80	0.60, 1.07	0.1283
Japan	10	3 (30.0)	NE (NE,	NE)	6	6 (100)	2.1 (0.3, NE)	0.17	0.04, 0.65	0.0095*
Interaction p-value										0.0268*
ECOG performance status at	Basel	ine								
(0) Normal activity	190	89 (46.8)	NE (NE,	NE)	100	59 (59.0)	4.2 (2.8, 8.1)	0.68	0.49, 0.95	0.0234*
(1) Restricted activity	61	35 (57.4)	9.8 (5.8,	NE)	30	19 (63.3)	11.0 (2.8, NE)	0.91	0.53, 1.63	0.7475
Interaction p-value										0.3695
Baseline CA-125 value										
<=ULN	228	112 (49.1)	, ,	NE)	117	70 (59.8)	, , ,	0.73	0.55, 0.99	0.0459*
>ULN	27	15 (55.6)	6.9 (2.8,	NE)	14	8 (57.1)	4.9 (0.7, NE)	0.82	0.36, 2.05	0.6623
Interaction p-value										0.8023
Histological grade										
High grade	255	127 (49.8)	30.5 (9.7,	NE)	131	78 (59.5)	5.5 (3.4,11.3)	0.74	0.56, 0.99	0.0429*
Interaction p-value										NC
Cytoreductive surgery outco	ome									
No residue	166	78 (47.0)	NE (NE,	NE)	80	43 (53.8)	11.0 (4.2, NE)	0.83	0.57, 1.21	0.3220
Residue	79	41 (51.9)	16.6 (5.6,	NE)	43	27 (62.8)	4.2 (2.1, NE)	0.73	0.45, 1.21	0.2167
Interaction p-value										0.7006

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaag 11AUG2022:11:30 kpzx329

Table 3.4.7 PAOLA1: Summary of subgroup analysis of AESI: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		_	bevacizumab 255)			Placebo + b (N=1						
Subgroup		Number (%) of patients with events				Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
Timing of cytoreductive :	surgery											
Upfront	146	71 (48.6)	NE (NE,	NE)	78	45 (57.7)	8.1 (4.2,	NE)	0.81	0.56,	1.18	0.2650
Interval	99	48 (48.5)	NE (NE,	NE)	45	25 (55.6)	5.6 (2.8,	NE)	0.77	0.48,	1.26	0.2903
Interaction p-value												0.8716
Myriad tumour BRCA mutat:	ion stat	us										
tBRCAm	158	77 (48.7)	NE (NE,	NE)	77	45 (58.4)	7.0 (3.5,	NE)	0.75	0.52,	1.10	0.1369
Non-tBRCAm	97	50 (51.5)	17.9 (7.5,	NE)	54	33 (61.1)	4.3 (2.1,	NE)	0.73	0.47,	1.15	0.1735
Interaction p-value												0.9289
Status somatic BRCA muta	cions											
sBRCAm	25	11 (44.0)	NE (NE,	NE)	9	4 (44.4)	NE (NE,	NE)	0.86	0.30,	3.12	0.8036
gBRCAm	69	33 (47.8)	NE (NE,	NE)	36	22 (61.1)	6.2 (2.8,	NE)	0.66	0.39,	1.15	0.1368
Non-BRCAm	43	22 (51.2)	20.3 (7.5,	NE)	23	15 (65.2)	3.4 (1.4,	NE)	0.59	0.31,	1.16	0.1209
Interaction p-value												0.8447

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaag 11AUG2022:11:30 kpzx329

Table 3.4.8 PAOLA1: Summary of subgroup analysis of AESI: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		mab			Placebo + be (N=1)		mab					
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI)			Number (%) of patients with events	Medi (95 (mont	% CI)	Hazard ratio [b]	95% C	I [b]	2-sided p-value [b]
First line treatment out	come (IVI	RS)												
NED [PDS]	92	6 (6.5)	NE (NE,	NE)	48	9 (18.8)	NE (NE,	NE)	0.31	0.10,	0.85	0.0229*
NED/CR [IDS]	74	7 (9.5)	NE (NE,	NE)	38	3 (7.9)	NE (NE,	NE)	1.20	0.33,	5.57	0.7888
NED/CR [Chemo]	40	3 (7.5)	NE (NE,	NE)	20	3 (15.0)	NE (NE,	NE)	0.44	0.08,	2.36	0.3173
PR	49	4 (8.2)	NE (NE,	NE)	25	4 (16.0)	NE (NE,	NE)	0.49	0.12,	2.07	0.3173
Interaction p-value														0.4427
Screening laboratory tBR0	CA statu	s (IVRS)												
tBRCAm	150	15 (10.0)	NE (NE,	NE)	65	9 (13.8)	NE (NE,	NE)	0.67	0.30,	1.60	0.3532
non-tBRCAm	105	5 (4.8)	NE (NE,	NE)	66	10 (15.2)	NE (NE,	NE)	0.29	0.09,	0.82	0.0190*
Interaction p-value														0.2172
First line treatment out	come (eCI	RF)												
NED [PDS]	89	5 (5.6)	NE (NE,	NE)	47	8 (17.0)	NE (NE,	NE)	0.29	0.09,	0.88	0.0285*
NED/CR [IDS]	74	9 (12.2)	NE (NE,	NE)	32	2 (6.3)	NE (NE,	NE)	1.98	0.51,	13.00	0.3489
NED/CR [Chemo]	39	1 (2.6)	NE (NE,	NE)	17	2 (11.8)	NE (NE,	NE)	0.20	0.01,	2.07	0.1704
PR	50	4 (8.0)	NE (NE,	NE)	34	6 (17.6)	NE (NE,	NE)	0.41	0.11,	1.45	0.1648
Interaction p-value														0.1346
Screening laboratory tBR0	CA status	s (eCRF)												
tBRCAm	147	14 (9.5)	NE (NE,	NE)	67	9 (13.4)	NE (NE,	NE)	0.66	0.29,	1.58	0.3362
non-tBRCAm	108	6 (5.6)	NE (NE,	NE)	64	10 (15.6)	NE (NE,	NE)	0.33	0.11,	0.89	0.0281*
Interaction p-value														0.2966
Age group														
<65 years	185	14 (7.6)	NE (NE,	NE)	98	17 (17.3)	NE (NE,	NE)	0.39	0.19,	0.80	0.0101*
>=65 years	70	6 (8.6)	NE (NE,	NE)	33	2 (6.1)	NE (NE,	NE)	1.41	0.33,	9.65	0.6628
Interaction p-value														0.1294

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaah 11AUG2022:11:30 kpzx329

Table 3.4.8 PAOLA1: Summary of subgroup analysis of AESI: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b		ıb		Placebo + be (N=1)						
Subgroup		Number (%) of patients with events	Median (95% (months	CI)		Number (%) of patients with events	Median t (95% C (months)	I)	Hazard ratio [b]	95% C]	[b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	15 (8.2)	NE (N	E, NE)	89	12 (13.5)	NE (NE	, NE)	0.56	0.26,	1.23	0.1473
IV	73	5 (6.8)	NE (N	E, NE)	42	7 (16.7)	NE (NE	, NE)	0.39	0.11,	1.21	0.1006
Interaction p-value												0.5834
Region												
Europe	245	20 (8.2)	NE (N	E, NE)	125	16 (12.8)	NE (NE	, NE)	0.60	0.31,	1.17	0.1293
Japan	10	0	NE (N	E, NE)	6	3 (50.0)	NE (NE	, NE)	NC	NC		NC
Interaction p-value												NC
ECOG performance status at	Basel	ine										
(0) Normal activity	190	15 (7.9)	NE (N	E, NE)	100	13 (13.0)	NE (NE	, NE)	0.57	0.27,	1.22	0.1459
(1) Restricted activity	61	5 (8.2)	NE (N	E, NE)	30	6 (20.0)	NE (NE	, NE)	0.36	0.10,	1.20	0.0948
Interaction p-value												0.5191
Baseline CA-125 value												
<=ULN	228	18 (7.9)	NE (N	E, NE)	117	17 (14.5)	NE (NE	, NE)	0.51	0.26,		0.0465*
>ULN	27	2 (7.4)	NE (N	E, NE)	14	2 (14.3)	NE (NE	, NE)	0.47	0.06,	3.91	0.4548
Interaction p-value												0.9439
Histological grade												
High grade	255	20 (7.8)	NE (N	E, NE)	131	19 (14.5)	NE (NE	, NE)	0.50	0.27,	0.95	0.0336*
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	15 (9.0)	NE (N	E, NE)	80	11 (13.8)	NE (NE	, NE)	0.62	0.29,	1.38	0.2329
Residue	79	4 (5.1)	NE (N	E, NE)	43	7 (16.3)	NE (NE	, NE)	0.28	0.07,	0.92	0.0354*
Interaction p-value												0.2710

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaah 11AUG2022:11:30 kpzx329

Table 3.4.8 PAOLA1: Summary of subgroup analysis of AESI: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		•		Placebo + be (N=1)		mab				
Subgroup		Number (%) of patients with events	Median t (95% CI (months))		Number (%) of patients with events	(95	an time % CI) hs) [a]	- Hazard ratio [b]	95% C	I [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	8 (5.5)	NE (NE,	NE)	78	14 (17.9)	NE (NE, N	E) 0.27	0.11,	0.62	0.0022*
Interval	99	11 (11.1)	NE (NE,	NE)	45	4 (8.9)	NE (NE, N	E) 1.26	0.43,	4.54	0.6900
Interaction p-value												0.0270*
Myriad tumour BRCA mutat:	ion stat	us										
tBRCAm	158	16 (10.1)	NE (NE,	NE)	77	10 (13.0)	NE (NE, N	0.72	0.33,	1.65	0.4272
Non-tBRCAm	97	4 (4.1)	NE (NE,	NE)	54	9 (16.7)	NE (NE, N	0.23	0.06,	0.71	0.0099*
Interaction p-value												0.1026
Status somatic BRCA mutat	cions											
sBRCAm	25	2 (8.0)	NE (NE,	NE)	9	2 (22.2)	NE (NE, N	E) 0.29	0.04,	2.46	0.2356
gBRCAm	69	7 (10.1)	NE (NE,	NE)	36	7 (19.4)	NE (NE, N	E) 0.49	0.17,	1.43	0.1882
Non-BRCAm	43	2 (4.7)	NE (NE,	NE)	23	2 (8.7)	NE (NE, N	E) 0.53	0.06,	4.41	0.5280
Interaction p-value												0.8913

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaah 11AUG2022:11:30 kpzx329

Table 3.4.9 PAOLA1: Summary of subgroup analysis of AESI: GI perforations, abscesses and fistulae Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2)		umab			Placebo + be (N=13		mab	•			
Subgroup	С	Number (%) of patients with events	(95	an ti 5% CI; :hs) [)	C	Number (%) of patients with events	(95	an ti 5% CI :hs))	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]	
First line treatment outo	come (IVR	S)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	2 (5.0)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	1 (2.0)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA status	(IVRS)											
tBRCAm	150	2 (1.3)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	1 (1.0)	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment outo	come (eCR	F)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	2 (5.1)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	1 (2.0)	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA status	(eCRF)											
tBRCAm	147	2 (1.4)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	1 (0.9)	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	2 (1.1)	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	1 (1.4)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaai 11AUG2022:11:30 kpzx329

Table 3.4.9 PAOLA1: Summary of subgroup analysis of AESI: GI perforations, abscesses and fistulae Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2!		nab		Placebo + bo					
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]			Number (%) of patients with events	Median t (95% C (months)	I)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	1 (0.5)	NE (NE, NE) 89	0	NE (NE	, NE)	NC	NC	NC
IV	73	2 (2.7)	NE (NE, NE) 42	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	3 (1.2)	NE (NE, NE) 125	0	NE (NE	, NE)	NC	NC	NC
Japan	10	0	NE (NE, NE) 6	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Baseli	.ne									
(0) Normal activity	190	2 (1.1)	NE (NE, NE) 100	0	NE (NE	, NE)	NC	NC	NC
(1) Restricted activity	61	1 (1.6)	NE (NE, NE) 30	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	2 (0.9)	NE (NE, NE) 117	0	NE (NE	, NE)	NC	NC	NC
>ULN	27	1 (3.7)	NE (NE, NE) 14	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	3 (1.2)	NE (NE, NE) 131	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	0	NE (NE, NE) 80	0	NE (NE	, NE)	NC	NC	NC
Residue	79	3 (3.8)	NE (NE, NE) 43	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaai 11AUG2022:11:30 kpzx329

Table 3.4.9 PAOLA1: Summary of subgroup analysis of AESI: GI perforations, abscesses and fistulae Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2			Placebo + be (N=1		mab					
Subgroup	C	Number (%) of patients with events	(959	n time % CI) ns) [a]		Number (%) of patients with events	(95	an tim % CI) hs) [Hazard ratio [b] 95% CI [b]		2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	2 (1.4)	NE (NE, NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	1 (1.0)	NE (NE, NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Myriad tumour BRCA mutat:	ion statu	ıs										
tBRCAm	158	2 (1.3)	NE (NE, NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	1 (1.0)	NE (NE, NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Status somatic BRCA muta	tions											
sBRCAm	25	0	NE (NE, NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE, NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE, NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaai 11AUG2022:11:30 kpzx329

Table 3.4.10 PAOLA1: Summary of subgroup analysis of AESI: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2!		ımab			Placebo + bo (N=1		ımab				
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI)	C	Number (%) of patients with events	(95	an ti 5% CI: ths))	Hazard ratio [b] 95% CI [b]		2-sided p-value [b]
First line treatment out	come (IVR	.S)											
NED [PDS]	92	1 (1.1)	NE (NE,	NE)	48	3 (6.3)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	1 (2.0)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(IVRS)											
tBRCAm	150	2 (1.3)	NE (NE,	NE)	65	2 (3.1)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	1 (1.1)	NE (NE,	NE)	47	3 (6.4)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	1 (2.0)	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(eCRF)											
tBRCAm	147	2 (1.4)	NE (NE,	NE)	67	2 (3.0)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	2 (1.1)	NE (NE,	NE)	98	2 (2.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	1 (3.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaj 11AUG2022:11:30 kpzx329

Table 3.4.10 PAOLA1: Summary of subgroup analysis of AESI: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ımab	-		Placebo + be (N=1)		b			
Subgroup	C	Number (%) of patients with events	(95	an ti % CI) hs) [)	C	Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	ratio	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	1 (0.5)	NE (NE,	NE)	89	3 (3.4)	NE (N	E, NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE,	NE)	42	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	2 (0.8)	NE (NE,	NE)	125	3 (2.4)	NE (E, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (E, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	1 (0.5)	NE (NE,	NE)	100	2 (2.0)	NE (E, NE)	NC	NC	NC
(1) Restricted activity	61	1 (1.6)	NE (NE,	NE)	30	1 (3.3)	NE (E, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	1 (0.4)	NE (NE,	NE)	117	3 (2.6)	NE (E, NE)	NC	NC	NC
>ULN	27	1 (3.7)	NE (NE,	NE)	14	0	NE (E, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	2 (0.8)	NE (NE,	NE)	131	3 (2.3)	NE (E, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outc	ome											
No residue	166	1 (0.6)	NE (NE,	NE)	80	3 (3.8)	NE (E, NE)	NC	NC	NC
Residue	79	1 (1.3)	NE (NE,	NE)	43	0	NE (E, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaj 11AUG2022:11:30 kpzx329

Table 3.4.10 PAOLA1: Summary of subgroup analysis of AESI: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		ab		Placebo + be (N=1)		nab			
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]		Number (%) of patients n with events		Median time (95% CI) (months) [a]		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	1 (0.7)	NE (NE, NE)	78	3 (3.8)	NE (NE, NE) NC	NC	NC
Interval	99	1 (1.0)	NE (NE, NE)	45	0	NE (NE, NE) NC	NC	NC
Interaction p-value											NC
Myriad tumour BRCA mutat:	ion statu	ıs									
tBRCAm	158	1 (0.6)	NE (NE, NE)	77	2 (2.6)	NE (NE, NE) NC	NC	NC
Non-tBRCAm	97	1 (1.0)	NE (NE, NE)	54	1 (1.9)	NE (NE, NE) NC	NC	NC
Interaction p-value											NC
Status somatic BRCA mutat	tions										
sBRCAm	25	0	NE (NE, NE)	9	0	NE (NE, NE) NC	NC	NC
gBRCAm	69	0	NE (NE, NE)	36	0	NE (NE, NE) NC	NC	NC
Non-BRCAm	43	0	NE (NE, NE)	23	0	NE (NE, NE) NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaj 11AUG2022:11:30 kpzx329

Table 3.4.11 PAOLA1: Summary of subgroup analysis of AESI: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		-	bevacizumab 255)			Placebo + be (N=1)					2-sided p-value [b]
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))	C	Number (%) of patients with events	Median t (95% CI (months))	Hazard ratio [b]	95% CI [b]	
First line treatment out	come (IV)	RS)									
NED [PDS]	92	12 (13.0)	NE (NE,	NE)	48	7 (14.6)	NE (NE,	NE)	0.77	0.30, 2.09	0.5923
NED/CR [IDS]	74	9 (12.2)	NE (NE,	NE)	38	3 (7.9)	NE (NE,	NE)	1.55	0.46, 7.00	0.4938
NED/CR [Chemo]	40	4 (10.0)	NE (NE,	NE)	20	1 (5.0)	NE (NE,	NE)	1.95	0.29, 38.03	0.5266
PR	49	5 (10.2)	NE (NE,	NE)	25	1 (4.0)	NE (NE,	NE)	2.70	0.44, 51.77	0.3150
Interaction p-value											0.6104
Screening laboratory tBR0	CA statu	s (IVRS)									
tBRCAm	150	18 (12.0)	75.0 (NE,	NE)	65	9 (13.8)	NE (NE,	NE)	0.81	0.37, 1.90	0.6210
non-tBRCAm	105	12 (11.4)	NE (NE,	NE)	66	3 (4.5)	NE (NE,	NE)	2.40	0.76, 10.56	0.1447
Interaction p-value											0.1420
First line treatment out	come (eCl	RF)									
NED [PDS]	89	11 (12.4)	NE (NE,	NE)	47	7 (14.9)	NE (NE,	NE)	0.71	0.27, 1.95	0.4882
NED/CR [IDS]	74	9 (12.2)	NE (NE,	NE)	32	2 (6.3)	NE (NE,	NE)	1.99	0.51, 13.07	0.3446
NED/CR [Chemo]	39	4 (10.3)	NE (NE,	NE)	17	1 (5.9)	NE (NE,	NE)	1.75	0.26, 34.22	0.5975
PR	50	5 (10.0)	NE (NE,	NE)	34	2 (5.9)	NE (NE,	NE)	1.71	0.37, 11.95	0.5065
Interaction p-value											0.5984
Screening laboratory tBR0	CA statu	s (eCRF)									
tBRCAm	147	17 (11.6)	75.0 (NE,	NE)	67	9 (13.4)	NE (NE,	NE)	0.81	0.37, 1.90	0.6106
non-tBRCAm	108	13 (12.0)	NE (NE,	NE)	64	3 (4.7)	NE (NE,	NE)	2.47	0.79, 10.80	0.1258
Interaction p-value											0.1265
Age group											
<65 years	185	21 (11.4)	75.0 (75.0,	NE)	98	9 (9.2)	NE (NE,	NE)	1.17	0.55, 2.69	0.6927
>=65 years	70	9 (12.9)	NE (NE,	NE)	33	3 (9.1)	NE (NE,	NE)	1.38	0.41, 6.24	0.6200
Interaction p-value											0.8291

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaak 11AUG2022:11:30 kpzx329

Table 3.4.11 PAOLA1: Summary of subgroup analysis of AESI: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		umab			Placebo + be (N=1)		ab	,		
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI)		Number (%) of patients with events	Mediar (95% (month	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	24 (13.2)	NE (NE,	NE)	89	11 (12.4)	NE (NE, NE)	0.99	0.49, 2.11	0.9732
IV	73	6 (8.2)	NE (NE,	NE)	42	1 (2.4)	NE (NE, NE)	3.55	0.61, 67.09	0.1776
Interaction p-value												0.2116
Region												
Europe	245	30 (12.2)	NE (NE,	NE)	125	12 (9.6)	NE (NE, NE)	1.21	0.64, 2.47	0.5659
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Basel:	ine										
(0) Normal activity	190	24 (12.6)	NE (NE,	NE)	100	10 (10.0)	NE (NE, NE)	1.24	0.61, 2.72	0.5588
(1) Restricted activity	61	5 (8.2)	NE (NE,	NE)	30	1 (3.3)	NE (NE, NE)	2.47	0.40, 47.23	0.3657
Interaction p-value												0.5311
Baseline CA-125 value												
<=ULN	228	25 (11.0)	NE (NE,	NE)	117	12 (10.3)	NE (NE, NE)	1.00	0.51, 2.07	1.0000
>ULN	27	5 (18.5)	NE (NE,	NE)	14	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	30 (11.8)	NE (NE,	NE)	131	12 (9.2)	NE (NE, NE)	1.22	0.64, 2.49	0.5492
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	21 (12.7)	NE (NE,	NE)	80	9 (11.3)	NE (NE, NE)	1.04	0.49, 2.41	0.9149
Residue	79	8 (10.1)	NE (NE,	NE)	43	3 (7.0)	NE (NE, NE)	1.45	0.42, 6.62	0.5738
Interaction p-value												0.6732

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaak 11AUG2022:11:30 kpzx329

Table 3.4.11 PAOLA1: Summary of subgroup analysis of AESI: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + (N=	bevacizuma 255)	b		Placebo + b (N=1		ab			
Subgroup		Number (%) of patients with events		CI)		Number (%) of patients with events	(95%	n time CI) s) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	17 (11.6)	NE (N	E, NE)	78	8 (10.3)	NE (NE, NE)	1.04	0.46, 2.56	0.9296
Interval	99	12 (12.1)	NE (N	E, NE)	45	4 (8.9)	NE (NE, NE)	1.36	0.48, 4.88	0.5817
Interaction p-value											0.7042
Myriad tumour BRCA mutat:	ion stat	us									
tBRCAm	158	20 (12.7)	75.0 (N	E, NE)	77	10 (13.0)	NE (NE, NE)	0.92	0.44, 2.05	0.8298
Non-tBRCAm	97	10 (10.3)	NE (N	E, NE)	54	2 (3.7)	NE (NE, NE)	2.66	0.70, 17.37	0.1644
Interaction p-value											0.1920
Status somatic BRCA muta	cions										
sBRCAm	25	2 (8.0)	NE (N	E, NE)	9	0	NE (NE, NE)	NC	NC	NC
gBRCAm	69	11 (15.9)	75.0 (N	E, NE)	36	6 (16.7)	NE (NE, NE)	0.90	0.34, 2.63	0.8411
Non-BRCAm	43	5 (11.6)	NE (N	E, NE)	23	1 (4.3)	NE (NE, NE)	2.36	0.37, 45.47	0.3955
Interaction p-value											0.3979

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaak 11AUG2022:11:30 kpzx329

Table 3.4.12 PAOLA1: Summary of subgroup analysis of AESI: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + (N=2		ab		Placebo + be (N=1					
Subgroup	C	Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVF	RS)									
NED [PDS]	92	1 (1.1)	NE (I	NE, NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (I	NE, NE)	38	1 (2.6)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (I	NE, NE)	20	1 (5.0)	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (I	NE, NE)	25	1 (4.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	s (IVRS)									
tBRCAm	150	3 (2.0)	NE (I	NE, NE)	65	2 (3.1)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (I	NE, NE)	66	2 (3.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
First line treatment out	come (eCF	RF)									
NED [PDS]	89	1 (1.1)	NE (I	NE, NE)	47	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (I	NE, NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (I	NE, NE)	17	1 (5.9)	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (I	NE, NE)	34	2 (5.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	s (eCRF)									
tBRCAm	147	3 (2.0)	NE (I	NE, NE)	67	2 (3.0)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (I	NE, NE)	64	2 (3.1)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Age group											
<65 years	185	1 (0.5)	NE (I	NE, NE)	98	1 (1.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	2 (2.9)	NE (I	NE, NE)	33	3 (9.1)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaal 11AUG2022:11:30 kpzx329

Table 3.4.12 PAOLA1: Summary of subgroup analysis of AESI: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		ımab	•		Placebo + be (N=13		ab			
Subgroup	c	Number (%) of patients with events		an ti % CI) hs) [c	Number (%) of patients with events	Mediar (95% (month	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	3 (1.6)	NE (NE,	NE)	89	2 (2.2)	NE (NE, NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	2 (4.8)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	3 (1.2)	NE (NE,	NE)	125	4 (3.2)	NE (NE, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	2 (1.1)	NE (NE,	NE)	100	2 (2.0)	NE (NE, NE)	NC	NC	NC
(1) Restricted activity	61	1 (1.6)	NE (NE,	NE)	30	2 (6.7)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	3 (1.3)	NE (NE,	NE)	117	4 (3.4)	NE (NE, NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	3 (1.2)	NE (NE,	NE)	131	4 (3.1)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	me											
No residue	166	3 (1.8)	NE (NE,	NE)	80	1 (1.3)	NE (NE, NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	3 (7.0)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaal 11AUG2022:11:30 kpzx329

Table 3.4.12 PAOLA1: Summary of subgroup analysis of AESI: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	. (Olaparib + b (N=2			Placebo + be (N=1)					
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive	surgery									
Upfront	146	1 (0.7)	NE (NE, NE)	78	3 (3.8)	NE (NE,	NE)	NC	NC	NC
Interval	99	2 (2.0)	NE (NE, NE)	45	1 (2.2)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Myriad tumour BRCA mutat	ion statu	ıs								
tBRCAm	158	3 (1.9)	NE (NE, NE)	77	3 (3.9)	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE, NE)	54	1 (1.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Status somatic BRCA muta	tions									
sBRCAm	25	0	NE (NE, NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE, NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE, NE)	23	1 (4.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaal 11AUG2022:11:30 kpzx329

Table 3.4.13 PAOLA1: Summary of subgroup analysis of AESI: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	. (Olaparib (+ be N=255		ımab			Placebo + bo (N=1		mab				
Subgroup	c	Number (of patier vith ever	its	Media (95 (mont	% CI))	C	Number (%) of patients with events	(95	an ti % CI; hs) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	LS)												
NED [PDS]	92	4 (4.	3)	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	4 (5.	4)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0		NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	2 (4.	1)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value														NC
Screening laboratory tBR	CA status	(IVRS)												
tBRCAm	150	7 (4.	7)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	3 (2.	9)	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value														NC
First line treatment out	come (eCR	2F)												
NED [PDS]	89	3 (3.	4)	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	7 (9.	5)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0		NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0		NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value														NC
Screening laboratory tBR	CA status	(eCRF)												
tBRCAm	147	7 (4.	8)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	3 (2.	8)	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	NC	NC	NC
Interaction p-value														NC
Age group														
<65 years	185	6 (3.	2)	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	4 (5.	7)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value														NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaam 11AUG2022:11:30 kpzx329

Table 3.4.13 PAOLA1: Summary of subgroup analysis of AESI: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ab		Placebo + be (N=1)					
Subgroup		Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	Median t (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	7 (3.8)	NE (N	E, NE)	89	1 (1.1)	NE (NE	NE)	NC	NC	NC
IV	73	3 (4.1)	NE (N	E, NE)	42	0	NE (NE	NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	10 (4.1)	NE (N	E, NE)	125	1 (0.8)	NE (NE	NE)	5.07	0.97, 93.09	0.0550
Japan	10	0	NE (N	E, NE)	6	0	NE (NE	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	7 (3.7)	NE (N	E, NE)	100	1 (1.0)	NE (NE	NE)	NC	NC	NC
(1) Restricted activity	61	3 (4.9)	NE (N	E, NE)	30	0	NE (NE	NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	10 (4.4)	NE (N	E, NE)	117	1 (0.9)	NE (NE	NE)	5.12	0.98, 93.91	0.0534
>ULN	27	0	NE (N	E, NE)	14	0	NE (NE	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	10 (3.9)	NE (N	E, NE)	131	1 (0.8)	NE (NE	NE)	5.12	0.98, 93.88	0.0535
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	10 (6.0)	NE (N	E, NE)	80	1 (1.3)	NE (NE	NE)	4.81	0.92, 88.29	0.0650
Residue	79	0	NE (N	E, NE)	43	0	NE (NE	NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaam 11AUG2022:11:30 kpzx329

Table 3.4.13 PAOLA1: Summary of subgroup analysis of AESI: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	1	Olaparib + b (N=2!		Placebo + b (N=1				
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery							
Upfront	146	3 (2.1)	NE (NE, NE)	78 1 (1.3)	NE (NE, NE)	NC	NC	NC
Interval	99	7 (7.1)	NE (NE, NE)	45 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC
Myriad tumour BRCA mutati	ion statu	ıs						
tBRCAm	158	8 (5.1)	NE (NE, NE)	77 0	NE (NE, NE)	NC	NC	NC
Non-tBRCAm	97	2 (2.1)	NE (NE, NE)	54 1 (1.9)	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC
Status somatic BRCA mutat	cions							
sBRCAm	25	1 (4.0)	NE (NE, NE)	9 0	NE (NE, NE)	NC	NC	NC
gBRCAm	69	5 (7.2)	NE (NE, NE)	36 0	NE (NE, NE)	NC	NC	NC
Non-BRCAm	43	2 (4.7)	NE (NE, NE)	23 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaam 11AUG2022:11:30 kpzx329

Table 3.4.14 PAOLA1: Summary of subgroup analysis of AESI: Posterior Reversible Encephalopathy Syndrome (PRES) Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	. (Olaparib + b (N=2)		umab			Placebo + be (N=1)		ımab				
Subgroup	c	Number (%) of patients with events	(95	an ti 5% CI :hs))	C	Number (%) of patients with events	(95	an ti 5% CI ths))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	LS)											
NED [PDS]	92	0	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(IVRS)											
tBRCAm	150	0	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	EF)											
NED [PDS]	89	0	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(eCRF)											
tBRCAm	147	0	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	0	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaan 11AUG2022:11:30 kpzx329

Table 3.4.14 PAOLA1: Summary of subgroup analysis of AESI: Posterior Reversible Encephalopathy Syndrome (PRES)
Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + be (N=1)		nab			
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI ths))	C	Number (%) of patients with events	(95	n time % CI) ns) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	0	NE (NE,	NE)	89	1 (1.1)	NE (NE, NE) NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	0	NE (NE,	NE)	125	1 (0.8)	NE (NE, NE) NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	0	NE (NE,	NE)	100	0	NE (NE, NE) NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	1 (3.3)	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	0	NE (NE,	NE)	117	1 (0.9)	NE (NE, NE) NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	0	NE (NE,	NE)	131	1 (0.8)	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	0	NE (NE,	NE)	80	1 (1.3)	NE (NE, NE) NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaan 11AUG2022:11:30 kpzx329

Table 3.4.14 PAOLA1: Summary of subgroup analysis of AESI: Posterior Reversible Encephalopathy Syndrome (PRES)
Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		ımab			Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events		an ti % CI) hs) [C	Number (%) of patients with events	(95	an ti % CI) hs) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery												
Upfront	146	0	NE (NE,	NE)	78	1 (1.3)	NE (NE,	NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Myriad tumour BRCA mutat:	ion statu	ıs											
tBRCAm	158	0	NE (NE,	NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	1 (1.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Status somatic BRCA mutat	tions												
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE,	NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaan 11AUG2022:11:30 kpzx329

Table 3.4.15 PAOLA1: Summary of subgroup analysis of AESI: Congestive heart failure Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		umab			Placebo + be (N=13		mab				
Subgroup	c	Number (%) of patients with events	(95	an ti 5% CI :hs))	C	Number (%) of patients with events	Media (95 (mont	% CI))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	LS)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(IVRS)											
tBRCAm	150	0	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	EF)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(eCRF)											
tBRCAm	147	0	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	0	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaao 11AUG2022:11:30 kpzx329

Table 3.4.15 PAOLA1: Summary of subgroup analysis of AESI: Congestive heart failure Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		umab	•		Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI ths))	(Number (%) of patients with events	Media (95 (mont)	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	0	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	0	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	ne											
(0) Normal activity	190	0	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	0	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	0	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	0	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaao 11AUG2022:11:30 kpzx329

Table 3.4.15 PAOLA1: Summary of subgroup analysis of AESI: Congestive heart failure Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + b (N=1		mab				
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI) :hs) [)	(Number (%) of patients with events	(95	an ti % CI; hs) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery												
Upfront	146	0	NE (NE,	NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Myriad tumour BRCA mutat:	ion statu	ıs											
tBRCAm	158	0	NE (NE,	NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Status somatic BRCA muta	tions												
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE,	NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaao 11AUG2022:11:30 kpzx329

Table 3.4.16 PAOLA1: Summary of subgroup analysis of AESI: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + be (N=1)		mab	•			
Subgroup	C	Number (%) of patients with events	(95	an ti % CI hs) [)	C	Number (%) of patients with events	Medi (95 (mont	% CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	S)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	1 (2.6)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	1 (5.0)	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(IVRS)											
tBRCAm	150	0	NE (NE,	NE)	65	1 (1.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	1 (5.9)	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	1 (2.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(eCRF)											
tBRCAm	147	0	NE (NE,	NE)	67	1 (1.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	0	NE (NE,	NE)	98	2 (2.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaap 11AUG2022:11:30 kpzx329

Table 3.4.16 PAOLA1: Summary of subgroup analysis of AESI: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2		umab			Placebo + be (N=1)		0	:		
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI ths))	C	Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	0	NE (NE,	NE)	89	2 (2.2)	NE (N	E, NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	0	NE (NE,	NE)	125	2 (1.6)	NE (N	E, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	0	NE (NE,	NE)	100	2 (2.0)	NE (N	E, NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	0	NE (NE,	NE)	117	1 (0.9)	NE (N	E, NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	1 (7.1)	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	0	NE (NE,	NE)	131	2 (1.5)	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	0	NE (NE,	NE)	80	0	NE (N	E, NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	2 (4.7)	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaap 11AUG2022:11:30 kpzx329

Table 3.4.16 PAOLA1: Summary of subgroup analysis of AESI: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		umab	:		Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events		an ti % CI) .hs) [)	C	Number (%) of patients with events	(95	an time % CI) .hs) [a]	ra	zard tio b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery												
Upfront	146	0	NE (NE,	NE)	78	1 (1.3)	NE (NE, N	E) 1	JC	NC	NC
Interval	99	0	NE (NE,	NE)	45	1 (2.2)	NE (NE, N	E) 1	1C	NC	NC
Interaction p-value													NC
Myriad tumour BRCA mutati	ion statu	ıs											
tBRCAm	158	0	NE (NE,	NE)	77	1 (1.3)	NE (NE, N	E) 1	1C	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	1 (1.9)	NE (NE, N	E) 1	1C	NC	NC
Interaction p-value													NC
Status somatic BRCA mutat	cions												
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE, N	E) 1	1C	NC	NC
gBRCAm	69	0	NE (NE,	NE)	36	1 (2.8)	NE (NE, N	E) 1	1C	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE, N	E) 1	1C	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaap 11AUG2022:11:30 kpzx329

Table 3.4.17 PAOLA1: Summary of subgroup analysis of AESI: MDS/AML Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2)		ımab	,		Placebo + be (N=1)		ımab				
Subgroup	С	Number (%) of patients with events	Media (95 (mont)	% CI))	C	Number (%) of patients with events	(95	an ti 5% CI :hs))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	S)											
NED [PDS]	92	2 (2.2)	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (NE,	NE)	38	2 (5.3)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	1 (2.5)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA status	(IVRS)											
tBRCAm	150	2 (1.3)	NE (NE,	NE)	65	1 (1.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	2 (1.9)	NE (NE,	NE)	66	2 (3.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	2 (2.2)	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (NE,	NE)	32	2 (6.3)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	1 (2.6)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA status	(eCRF)											
tBRCAm	147	2 (1.4)	NE (NE,	NE)	67	1 (1.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	2 (1.9)	NE (NE,	NE)	64	2 (3.1)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	3 (1.6)	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	1 (1.4)	NE (NE,	NE)	33	2 (6.1)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaq 11AUG2022:11:30 kpzx329

Table 3.4.17 PAOLA1: Summary of subgroup analysis of AESI: MDS/AML Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2!		mab			Placebo + be (N=13		ab			
Subgroup	C	Number (%) of patients with events	Media (95% (mont)	% CI)		C	Number (%) of patients with events	Mediar (95% (month	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	3 (1.6)	NE (NE,	NE)	89	3 (3.4)	NE (NE, NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE,	NE)	42	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	4 (1.6)	NE (NE,	NE)	125	3 (2.4)	NE (NE, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	1 (0.5)	NE (NE,	NE)	100	2 (2.0)	NE (NE, NE)	NC	NC	NC
(1) Restricted activity	61	2 (3.3)	NE (NE,	NE)	30	1 (3.3)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	4 (1.8)	NE (NE,	NE)	117	3 (2.6)	NE (NE, NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	4 (1.6)	NE (NE,	NE)	131	3 (2.3)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	3 (1.8)	NE (NE,	NE)	80	3 (3.8)	NE (NE, NE)	NC	NC	NC
Residue	79	1 (1.3)	NE (NE,	NE)	43	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaq 11AUG2022:11:30 kpzx329

Table 3.4.17 PAOLA1: Summary of subgroup analysis of AESI: MDS/AML Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2!			Placebo + be (N=1					
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive	surgery									
Upfront	146	3 (2.1)	NE (NE, NE) 78	1 (1.3)	NE (NE,	NE)	NC	NC	NC
Interval	99	1 (1.0)	NE (NE, NE) 45	2 (4.4)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Myriad tumour BRCA mutat	ion statu	ıs								
tBRCAm	158	2 (1.3)	NE (NE, NE) 77	2 (2.6)	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	2 (2.1)	NE (NE, NE) 54	1 (1.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Status somatic BRCA muta	tions									
sBRCAm	25	1 (4.0)	NE (NE, NE) 9	1 (11.1)	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE, NE) 36	1 (2.8)	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE, NE) 23	1 (4.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaq 11AUG2022:11:30 kpzx329

Table 3.4.18 PAOLA1: Summary of subgroup analysis of AESI: Myelodysplastic syndrome and Acute myeloid leukaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2!		nab	:	Placebo + b (N=1		ab	:		
Subgroup	C	Number (%) of patients with events	(95%	n time CI) s) [a]	(Number (%) of patients with events		n time CI) s) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVF	RS)									
NED [PDS]	92	2 (2.2)	NE (NE, NE)	48	2 (4.2)	NE (NE, NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (NE, NE)	38	2 (5.3)	NE (NE, NE)	NC	NC	NC
NED/CR [Chemo]	40	1 (2.5)	NE (NE, NE)	20	0	NE (NE, NE)	NC	NC	NC
PR	49	0	NE (NE, NE)	25	0	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	s (IVRS)									
tBRCAm	150	2 (1.3)	NE (NE, NE)	65	1 (1.5)	NE (NE, NE)	NC	NC	NC
non-tBRCAm	105	2 (1.9)	NE (NE, NE)	66	3 (4.5)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
First line treatment out	come (eCF	RF)									
NED [PDS]	89	2 (2.2)	NE (NE, NE)	47	2 (4.3)	NE (NE, NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (NE, NE)	32	2 (6.3)	NE (NE, NE)	NC	NC	NC
NED/CR [Chemo]	39	1 (2.6)	NE (NE, NE)	17	0	NE (NE, NE)	NC	NC	NC
PR	50	0	NE (NE, NE)	34	0	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	s (eCRF)									
tBRCAm	147	2 (1.4)	NE (NE, NE)	67	1 (1.5)	NE (NE, NE)	NC	NC	NC
non-tBRCAm	108	2 (1.9)	NE (NE, NE)	64	3 (4.7)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Age group											
<65 years	185	3 (1.6)	NE (NE, NE)	98	2 (2.0)	NE (NE, NE)	NC	NC	NC
>=65 years	70	1 (1.4)	NE (NE, NE)	33	2 (6.1)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaar 11AUG2022:11:30 kpzx329

Table 3.4.18 PAOLA1: Summary of subgroup analysis of AESI: Myelodysplastic syndrome and Acute myeloid leukaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2				Placebo + be (N=1)					
Subgroup	C	Number (%) of patients with events	Median t (95% C (months)	[)	(Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	3 (1.6)	NE (NE	, NE)	89	4 (4.5)	NE (NE,	NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE	, NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	4 (1.6)	NE (NE	, NE)	125	4 (3.2)	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE	, NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Baseli	.ne									
(0) Normal activity	190	1 (0.5)	NE (NE	, NE)	100	3 (3.0)	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	2 (3.3)	NE (NE	, NE)	30	1 (3.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	4 (1.8)	NE (NE	, NE)	117	4 (3.4)	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE	, NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	4 (1.6)	NE (NE	, NE)	131	4 (3.1)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	3 (1.8)	NE (NE	, NE)	80	4 (5.0)	NE (NE,	NE)	NC	NC	NC
Residue	79	1 (1.3)	NE (NE	, NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaar 11AUG2022:11:30 kpzx329

Table 3.4.18 PAOLA1: Summary of subgroup analysis of AESI: Myelodysplastic syndrome and Acute myeloid leukaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		Placebo + b (N=1			
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95%	2-sided CI [b] p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	3 (2.1)	NE (NE, NE)	78 2 (2.6)	NE (NE, NE)	NC NC	NC
Interval	99	1 (1.0)	NE (NE, NE)	45 2 (4.4)	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Myriad tumour BRCA mutat	ion statu	ıs					
tBRCAm	158	2 (1.3)	NE (NE, NE)	77 3 (3.9)	NE (NE, NE)	NC NC	NC
Non-tBRCAm	97	2 (2.1)	NE (NE, NE)	54 1 (1.9)	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Status somatic BRCA mutat	cions						
sBRCAm	25	1 (4.0)	NE (NE, NE)	9 1 (11.1)	NE (NE, NE)	NC NC	NC
gBRCAm	69	1 (1.4)	NE (NE, NE)	36 2 (5.6)	NE (NE, NE)	NC NC	NC
Non-BRCAm	43	0	NE (NE, NE)	23 1 (4.3)	NE (NE, NE)	NC NC	NC
Interaction p-value							NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaar 11AUG2022:11:30 kpzx329

Table 3.4.19 PAOLA1: Summary of subgroup analysis of AESI: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	;	Olaparib + b (N=2		ab		Placebo + be (N=1			:		
Subgroup		Number (%) of patients with events	Median (95% (month)	CI)	(Number (%) of patients with events	Median t (95% C: (months)	[)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVI	RS)									
NED [PDS]	92	5 (5.4)	NE (NE, NE)	48	2 (4.2)	NE (NE	NE)	NC	NC	NC
NED/CR [IDS]	74	6 (8.1)	NE (NE, NE)	38	1 (2.6)	NE (NE	NE)	NC	NC	NC
NED/CR [Chemo]	40	4 (10.0)	NE (NE, NE)	20	1 (5.0)	NE (NE	NE)	NC	NC	NC
PR	49	0	NE (NE, NE)	25	0	NE (NE	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	s (IVRS)									
tBRCAm	150	13 (8.7)	NE (NE, NE)	65	3 (4.6)	NE (NE	NE)	1.68	0.54, 7.31	0.3956
non-tBRCAm	105	2 (1.9)	NE (NE, NE)	66	1 (1.5)	NE (NE	NE)	1.14	0.11, 24.54	0.9137
Interaction p-value											0.7833
First line treatment out	come (eCI	RF)									
NED [PDS]	89	6 (6.7)	NE (NE, NE)	47	2 (4.3)	NE (NE	NE)	NC	NC	NC
NED/CR [IDS]	74	6 (8.1)	NE (NE, NE)	32	0	NE (NE	NE)	NC	NC	NC
NED/CR [Chemo]	39	3 (7.7)	NE (NE, NE)	17	1 (5.9)	NE (NE	NE)	NC	NC	NC
PR	50	0	NE (NE, NE)	34	1 (2.9)	NE (NE	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	s (eCRF)									
tBRCAm	147	13 (8.8)	NE (NE, NE)	67	3 (4.5)	NE (NE	NE)	1.74	0.56, 7.60	0.3605
non-tBRCAm	108	2 (1.9)	NE (NE, NE)	64	1 (1.6)	NE (NE	NE)	1.09	0.10, 23.35	0.9463
Interaction p-value											0.7367
Age group											
<65 years	185	13 (7.0)	NE (NE, NE)	98	3 (3.1)	NE (NE	NE)	2.13	0.69, 9.29	0.2042
>=65 years	70	2 (2.9)	NE (NE, NE)	33	1 (3.0)	NE (NE	NE)	0.74	0.07, 15.91	0.8085
Interaction p-value											0.4631

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaas 11AUG2022:11:30 kpzx329

Table 3.4.19 PAOLA1: Summary of subgroup analysis of AESI: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2				Placebo + be (N=13					
Subgroup		Number (%) of patients with events	Median to (95% CI (months))	C	Number (%) of patients with events	Median tin (95% CI) (months) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	10 (5.5)	NE (NE,	NE)	89	3 (3.4)	NE (NE,	NE)	1.46	0.45, 6.51	0.5539
IV	73	5 (6.8)	NE (NE,	NE)	42	1 (2.4)	NE (NE,	NE)	2.57	0.41, 49.34	0.3422
Interaction p-value											0.6481
Region											
Europe	245	14 (5.7)	NE (NE,	NE)	125	4 (3.2)	NE (NE,	NE)	1.59	0.57, 5.62	0.3937
Japan	10	1 (10.0)	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	8 (4.2)	NE (NE,	NE)	100	3 (3.0)	NE (NE,	NE)	1.27	0.37, 5.80	0.7190
(1) Restricted activity	61	7 (11.5)	NE (NE,	NE)	30	1 (3.3)	NE (NE,	NE)	2.91	0.52, 54.37	0.2574
Interaction p-value											0.4960
Baseline CA-125 value											
<=ULN	228	15 (6.6)	NE (NE,	,	117	4 (3.4)	NE (NE,	NE)	1.66	0.60, 5.82	0.3477
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	15 (5.9)	NE (NE,	NE)	131	4 (3.1)	NE (NE,	NE)	1.71	0.62, 6.01	0.3157
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	12 (7.2)	NE (NE,	NE)	80	2 (2.5)	NE (NE,	NE)	2.55	0.69, 16.40	0.1732
Residue	79	3 (3.8)	NE (NE,	NE)	43	2 (4.7)	NE (NE,	NE)	0.73	0.12, 5.52	0.7299
Interaction p-value											0.2929

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaas 11AUG2022:11:30 kpzx329

Table 3.4.19 PAOLA1: Summary of subgroup analysis of AESI: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + N=2		Placebo + b (N=1				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery							
Upfront	146	9 (6.2)	NE (NE, NE)	78 3 (3.8)	NE (NE, NE)	1.33	0.40, 6.01	0.6596
Interval	99	6 (6.1)	NE (NE, NE)	45 1 (2.2)	NE (NE, NE)	2.62	0.45, 49.47	0.3204
Interaction p-value								0.5821
Myriad tumour BRCA mutati	ion stat	us						
tBRCAm	158	13 (8.2)	NE (NE, NE)	77 4 (5.2)	NE (NE, NE)	1.37	0.48, 4.86	0.5734
Non-tBRCAm	97	2 (2.1)	NE (NE, NE)	54 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC
Status somatic BRCA mutat	ions							
sBRCAm	25	1 (4.0)	NE (NE, NE)	9 0	NE (NE, NE)	NC	NC	NC
gBRCAm	69	6 (8.7)	NE (NE, NE)	36 2 (5.6)	NE (NE, NE)	NC	NC	NC
Non-BRCAm	43	1 (2.3)	NE (NE, NE)	23 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaas 11AUG2022:11:30 kpzx329

Table 3.4.20 PAOLA1: Summary of subgroup analysis of AESI: Pneumonitis Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	. (Olaparib + N (N=2		mab			Placebo + be (N=13		nab					
Subgroup	c	Number (%) of patients with events	Media (95: (mont)	% CI))	C	Number (%) of patients vith events	Media (95 (mont	% CI)		Hazard ratio [b]	95% C	!I [b]	2-sided p-value [b]
First line treatment out	come (IVR	RS)												
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC		NC
NED/CR [IDS]	74	2 (2.7)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC		NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC		NC
PR	49	1 (2.0)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
Screening laboratory tBR0	CA status	s (IVRS)												
tBRCAm	150	1 (0.7)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC		NC
non-tBRCAm	105	2 (1.9)	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
First line treatment out	come (eCR	RF)												
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC		NC
NED/CR [IDS]	74	3 (4.1)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC		NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC		NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
Screening laboratory tBR0	CA status	(eCRF)												
tBRCAm	147	1 (0.7)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC		NC
non-tBRCAm	108	2 (1.9)	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
Age group														
<65 years	185	1 (0.5)	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC		NC
>=65 years	70	2 (2.9)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaat 11AUG2022:11:30 kpzx329

Table 3.4.20 PAOLA1: Summary of subgroup analysis of AESI: Pneumonitis Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2!		ımab			Placebo + be (N=1)		mab	•			
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI))	(Number (%) of patients with events		an tir % CI) hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	2 (1.1)	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	3 (1.2)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	.ne											
(0) Normal activity	190	3 (1.6)	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	2 (0.9)	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	1 (3.7)	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	3 (1.2)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	3 (1.8)	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaat 11AUG2022:11:30 kpzx329

Table 3.4.20 PAOLA1: Summary of subgroup analysis of AESI: Pneumonitis Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2				oevacizumab 131)			
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients n with events		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery								
Upfront	146	0	NE (NE, N	IE)	78 0	NE (NE, NE)	NC	NC	NC
Interval	99	3 (3.0)	NE (NE, N	IE)	45 0	NE (NE, NE)	NC	NC	NC
Interaction p-value									NC
Myriad tumour BRCA mutat:	ion statu	ıs							
tBRCAm	158	2 (1.3)	NE (NE, N	IE)	77 0	NE (NE, NE)	NC	NC	NC
Non-tBRCAm	97	1 (1.0)	NE (NE, N	IE)	54 0	NE (NE, NE)	NC	NC	NC
Interaction p-value									NC
Status somatic BRCA mutat	tions								
sBRCAm	25	1 (4.0)	NE (NE, N	IE)	9 0	NE (NE, NE)	NC	NC	NC
gBRCAm	69	0	NE (NE, N	IE)	36 0	NE (NE, NE)	NC	NC	NC
Non-BRCAm	43	1 (2.3)	NE (NE, N	JE)	23 0	NE (NE, NE)	NC	NC	NC
Interaction p-value									NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaat 11AUG2022:11:30 kpzx329

Table 3.4.21 PAOLA1: Summary of subgroup analysis of Serious AESI: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	. (Olaparib (1	+ bev N=255		ımab			Placebo + bo (N=1		mab				
Subgroup	c	Number (% of patien vith even	ts	Media (95 (mont	% CI))	С	Number (%) of patients with events	(95	an ti 5% CI; :hs) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	RS)												
NED [PDS]	92	5 (5.	4)	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.	7)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	2 (5.	0)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	4 (8.	2)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value														NC
Screening laboratory tBR	CA status	(IVRS)												
tBRCAm	150	6 (4.	0)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	7 (6.	7)	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value														NC
First line treatment out	come (eCR	RF)												
NED [PDS]	89	5 (5.	6)	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	3 (4.	1)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	2 (5.	1)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	3 (6.	0)	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value														NC
Screening laboratory tBR	CA status	(eCRF)												
tBRCAm	147	6 (4.	1)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	7 (6.	5)	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	NC	NC	NC
Interaction p-value														NC
Age group														
<65 years	185	9 (4.	9)	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	4.84	0.91, 89.20	0.0674
>=65 years	70	4 (5.	7)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value														NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaau 11AUG2022:11:30 kpzx329

Table 3.4.21 PAOLA1: Summary of subgroup analysis of Serious AESI: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		nab	•		Placebo + be (N=13		ıab	-			
Subgroup		Number (%) of patients with events	Media (95% (month	CI)	-	C	Number (%) of patients with events		n time k CI) ns) [a	_	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	8 (4.4)	NE (NE,	NE)	89	1 (1.1)	NE (NE,	NE)	NC	NC	NC
IV	73	5 (6.8)	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	11 (4.5)	NE (NE,	NE)	125	1 (0.8)	NE (NE,	NE)	5.72	1.11,104.45	0.0345*
Japan	10	2 (20.0)	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Basel	ine											
(0) Normal activity	190	9 (4.7)	NE (NE,	NE)	100	1 (1.0)	NE (NE,	NE)	4.84	0.91, 89.22	0.0674
(1) Restricted activity	61	4 (6.6)	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	12 (5.3)	NE (NE,	NE)	117	1 (0.9)	NE (NE,	NE)	6.31	1.24,114.86	0.0224*
>ULN	27	1 (3.7)	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	13 (5.1)	NE (NE,	NE)	131	1 (0.8)	NE (NE,	NE)	6.83	1.36,124.06	0.0151*
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	8 (4.8)	NE (NE,	NE)	80	1 (1.3)	NE (NE,	NE)	NC	NC	NC
Residue	79	3 (3.8)	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaau 11AUG2022:11:30 kpzx329

Table 3.4.21 PAOLA1: Summary of subgroup analysis of Serious AESI: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		Placebo + b (N=1				
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery							
Upfront	146	8 (5.5)	NE (NE, NE)	78 1 (1.3)	NE (NE, NE)	NC I	NC	NC
Interval	99	3 (3.0)	NE (NE, NE)	45 0	NE (NE, NE)	NC I	NC	NC
Interaction p-value								NC
Myriad tumour BRCA mutat:	ion statu	ıs						
tBRCAm	158	8 (5.1)	NE (NE, NE)	77 0	NE (NE, NE)	NC I	NC	NC
Non-tBRCAm	97	5 (5.2)	NE (NE, NE)	54 1 (1.9)	NE (NE, NE)	NC I	NC	NC
Interaction p-value								NC
Status somatic BRCA mutat	tions							
sBRCAm	25	2 (8.0)	NE (NE, NE)	9 0	NE (NE, NE)	NC I	NC	NC
gBRCAm	69	3 (4.3)	NE (NE, NE)	36 0	NE (NE, NE)	NC I	NC	NC
Non-BRCAm	43	2 (4.7)	NE (NE, NE)	23 0	NE (NE, NE)	NC I	NC	NC
Interaction p-value								NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaau 11AUG2022:11:30 kpzx329

Table 3.4.22 PAOLA1: Summary of subgroup analysis of Serious AESI: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		mab			Placebo + be (N=13		nab	•			
Subgroup	c	Number (%) of patients with events	Media (95% (month	k CI))	C	Number (%) of patients with events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	RS)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	1 (2.5)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	1 (2.0)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(IVRS)											
tBRCAm	150	0	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	2 (1.9)	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	RF)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	1 (2.6)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	1 (2.0)	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(eCRF)											
tBRCAm	147	0	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	2 (1.9)	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	1 (0.5)	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	1 (1.4)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaav 11AUG2022:11:30 kpzx329

Table 3.4.22 PAOLA1: Summary of subgroup analysis of Serious AESI: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

-		Olaparib + b (N=2!		umab			Placebo + be (N=1)		mab	•			
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI; ths) [)	(Number (%) of patients with events		an ti: % CI) hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	1 (0.5)	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	2 (0.8)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	.ne											
(0) Normal activity	190	2 (1.1)	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	1 (0.4)	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	1 (3.7)	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	2 (0.8)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	0	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	1 (1.3)	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaav 11AUG2022:11:30 kpzx329

Table 3.4.22 PAOLA1: Summary of subgroup analysis of Serious AESI: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	1	Olaparib + b (N=2				Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	Median ti (95% CI (months))	C	Number (%) of patients with events	(95	an time % CI) hs) [a]	ra	zard atio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	1 (0.7)	NE (NE,	NE)	78	0	NE (NE, N	E)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (NE, N	E)	NC	NC	NC
Interaction p-value												NC
Myriad tumour BRCA mutat:	ion statu	ıs										
tBRCAm	158	1 (0.6)	NE (NE,	NE)	77	0	NE (NE, N	E)	NC	NC	NC
Non-tBRCAm	97	1 (1.0)	NE (NE,	NE)	54	0	NE (NE, N	E)	NC	NC	NC
Interaction p-value												NC
Status somatic BRCA mutat	tions											
sBRCAm	25	1 (4.0)	NE (NE,	NE)	9	0	NE (NE, N	E)	NC	NC	NC
gBRCAm	69	0	NE (NE,	NE)	36	0	NE (NE, N	E)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE, N	E)	NC	NC	NC
Interaction p-value												NC

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaav 11AUG2022:11:30 kpzx329

Table 3.4.23 PAOLA1: Summary of subgroup analysis of Serious AESI: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		mab	:		Placebo + be (N=13		ımab				
Subgroup	C	Number (%) of patients with events	Media (95% (month	È CI)		C	Number (%) of patients with events	(95	an ti 5% CI :hs))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)											
NED [PDS]	92	1 (1.1)	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	1 (2.5)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(IVRS)											
tBRCAm	150	1 (0.7)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	3 (2.9)	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	1 (1.1)	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	1 (2.6)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(eCRF)											
tBRCAm	147	1 (0.7)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	3 (2.8)	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	2 (1.1)	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	2 (2.9)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaw 11AUG2022:11:30 kpzx329

Table 3.4.23 PAOLA1: Summary of subgroup analysis of Serious AESI: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		umab			Placebo + be (N=1)		mab	•			
Subgroup	c	Number (%) of patients with events		an ti 5% CI; hs) [)	C	Number (%) of patients with events		an ti: % CI) hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	3 (1.6)	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	3 (1.2)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	1 (10.0)	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	.ne											
(0) Normal activity	190	3 (1.6)	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	1 (1.6)	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	4 (1.8)	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	4 (1.6)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	3 (1.8)	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaw 11AUG2022:11:30 kpzx329

Table 3.4.23 PAOLA1: Summary of subgroup analysis of Serious AESI: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2!				Placebo + be (N=13		mab				
Subgroup	C	Number (%) of patients with events	Median tim (95% CI) (months) [a		C	Number (%) of patients with events	(95	an time % CI) hs) [a		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	1 (0.7)	NE (NE,	NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	2 (2.0)	NE (NE,	NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Myriad tumour BRCA mutat:	ion statu	ıs										
tBRCAm	158	1 (0.6)	NE (NE,	NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	3 (3.1)	NE (NE,	NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Status somatic BRCA mutat	cions											
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE,	NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	1 (2.3)	NE (NE,	NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaw 11AUG2022:11:30 kpzx329

Table 3.4.24 PAOLA1: Summary of subgroup analysis of Serious AESI: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=25		umab	:-		Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI .hs))	C	Number (%) of patients with events	(95	an ti 5% CI; :hs) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	S)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(IVRS)											
tBRCAm	150	0	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(eCRF)											
tBRCAm	147	0	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	0	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaax 11AUG2022:11:30 kpzx329

Table 3.4.24 PAOLA1: Summary of subgroup analysis of Serious AESI: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2!		umab	•		Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	(9!	an ti 5% CI ths))	(Number (%) of patients with events		an tir % CI) hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	0	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	0	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	ine											
(0) Normal activity	190	0	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	0	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	0	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outc	ome												
No residue	166	0	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaax 11AUG2022:11:30 kpzx329

Table 3.4.24 PAOLA1: Summary of subgroup analysis of Serious AESI: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + b (N=1		mab				
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI) :hs) [)	(Number (%) of patients with events	(95	an ti % CI; hs) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery												
Upfront	146	0	NE (NE,	NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Myriad tumour BRCA mutat:	ion statu	ıs											
tBRCAm	158	0	NE (NE,	NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Status somatic BRCA muta	tions												
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE,	NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaax 11AUG2022:11:30 kpzx329

Table 3.4.25 PAOLA1: Summary of subgroup analysis of Serious AESI: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ımab			Placebo + bo (N=1		ımab					
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI)		Number (%) of patients with events	(95	an ti 5% CI ths))	Hazard ratio [b]	95% C]	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)												
NED [PDS]	92	7 (7.6)	NE (NE,	NE)	48	2 (4.2)	NE (NE,	NE)	1.75	0.42,	11.72	0.4655
NED/CR [IDS]	74	3 (4.1)	NE (NE,	NE)	38	6 (15.8)	NE (NE,	NE)	0.24	0.05,	0.90	0.0346*
NED/CR [Chemo]	40	3 (7.5)	NE (NE,	NE)	20	5 (25.0)	NE (NE,	NE)	0.25	0.05,	1.03	0.0553
PR	49	9 (18.4)	NE (NE,	NE)	25	3 (12.0)	NE (NE,	NE)	1.61	0.48,	7.27	0.4573
Interaction p-value														0.0506
Screening laboratory tBR0	CA statu	s (IVRS)												
tBRCAm	150	14 (9.3)	NE (NE,	NE)	65	10 (15.4)	NE (NE,	NE)	0.55	0.25,	1.29	0.1638
non-tBRCAm	105	8 (7.6)	NE (NE,	NE)	66	6 (9.1)	NE (NE,	NE)	0.82	0.28,	2.49	0.7128
Interaction p-value														0.5661
First line treatment out	come (eCl	RF)												
NED [PDS]	89	6 (6.7)	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	3.08	0.53,	58.13	0.2383
NED/CR [IDS]	74	4 (5.4)	NE (NE,	NE)	32	5 (15.6)	NE (NE,	NE)	0.33	0.08,	1.24	0.0975
NED/CR [Chemo]	39	4 (10.3)	NE (NE,	NE)	17	4 (23.5)	NE (NE,	NE)	0.40	0.09,	1.69	0.2016
PR	50	8 (16.0)	NE (NE,	NE)	34	5 (14.7)	NE (NE,	NE)	1.04	0.35,	3.45	0.9424
Interaction p-value														0.1743
Screening laboratory tBR0	CA statu	s (eCRF)												
tBRCAm	147	12 (8.2)	NE (NE,	NE)	67	10 (14.9)	NE (NE,	NE)	0.50	0.21,	1.18	0.1097
non-tBRCAm	108	10 (9.3)	NE (NE,	NE)	64	6 (9.4)	NE (NE,	NE)	0.98	0.36,	2.87	0.9626
Interaction p-value														0.3119
Age group														
<65 years	185	12 (6.5)	NE (NE,	NE)	98	12 (12.2)	NE (NE,	NE)	0.49	0.22,	1.10	0.0844
>=65 years	70	10 (14.3)	NE (NE,	NE)	33	4 (12.1)	NE (NE,	NE)	1.19	0.40,	4.33	0.7674
Interaction p-value														0.2076

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaay 11AUG2022:11:30 kpzx329

Table 3.4.25 PAOLA1: Summary of subgroup analysis of Serious AESI: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		umab			Placebo + be (N=1)		mab	,				
Subgroup		Number (%) of patients with events	(95	an ti 5% CI :hs))		Number (%) of patients with events	(95	n tim % CI) hs) [a		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)														
III	182	16 (8.8)	NE (NE,	NE)	89	9 (10.1)	NE (NE,	NE)	0.84	0.38,	1.98	0.6770
IV	73	6 (8.2)	NE (NE,	NE)	42	7 (16.7)	NE (NE,	NE)	0.45	0.14,	1.35	0.1512
Interaction p-value														0.3656
Region														
Europe	245	22 (9.0)	NE (NE,	NE)	125	16 (12.8)	NE (NE,	NE)	0.66	0.35,	1.28	0.2163
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
ECOG performance status at	Basel	ine												
(0) Normal activity	190	12 (6.3)	NE (NE,	NE)	100	12 (12.0)	NE (NE,	NE)	0.49	0.22,	1.11	0.0856
(1) Restricted activity	61	9 (14.8)	NE (NE,	NE)	30	4 (13.3)	NE (NE,	NE)	1.10	0.36,	4.05	0.8779
Interaction p-value														0.2611
Baseline CA-125 value														
<=ULN	228	20 (8.8)	NE (NE,	NE)	117	14 (12.0)	NE (NE,	NE)	0.70	0.36,		0.3108
>ULN	27	2 (7.4)	NE (NE,	NE)	14	2 (14.3)	NE (NE,	NE)	0.46	0.06,	3.86	0.4471
Interaction p-value														0.6981
Histological grade														
High grade	255	22 (8.6)	NE (NE,	NE)	131	16 (12.2)	NE (NE,	NE)	0.67	0.35,	1.30	0.2279
Interaction p-value														NC
Cytoreductive surgery outco	ome													
No residue	166	10 (6.0)	NE (NE,	NE)	80	7 (8.8)	NE (NE,	NE)	0.66	0.25,	1.81	0.4034
Residue	79	8 (10.1)	NE (NE,	NE)	43	9 (20.9)	NE (NE,	NE)	0.44	0.16,	1.15	0.0925
Interaction p-value														0.5579

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaay 11AUG2022:11:30 kpzx329

Table 3.4.25 PAOLA1: Summary of subgroup analysis of Serious AESI: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olap	arib + k (N=2		umab			Placebo + b (N=1		mab					
Subgroup		of pa	er (%) atients events	(9!	an ti 5% CI ths))		Number (%) of patients with events	(95	an ti 5% CI :hs))	Hazard ratio [b]	95% C	I [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery														
Upfront	146	12	(8.2)	NE (NE,	NE)	78	9 (11.5)	NE (NE,	NE)	0.66	0.28,	1.63	0.3600
Interval	99	6	(6.1)	NE (NE,	NE)	45	7 (15.6)	NE (NE,	NE)	0.37	0.12,	1.11	0.0756
Interaction p-value															0.4070
Myriad tumour BRCA mutat:	ion stat	us													
tBRCAm	158	14	(8.9)	NE (NE,	NE)	77	11 (14.3)	NE (NE,	NE)	0.57	0.26,	1.28	0.1678
Non-tBRCAm	97	8	(8.2)	NE (NE,	NE)	54	5 (9.3)	NE (NE,	NE)	0.88	0.29,	2.92	0.8286
Interaction p-value															0.5249
Status somatic BRCA mutat	cions														
sBRCAm	25	1	(4.0)	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC		NC
gBRCAm	69	4	(5.8)	NE (NE,	NE)	36	6 (16.7)	NE (NE,	NE)	0.32	0.08,	1.12	0.0733
Non-BRCAm	43	4	(9.3)	NE (NE,	NE)	23	1 (4.3)	NE (NE,	NE)	2.22	0.33,	43.39	0.4433
Interaction p-value															0.1018

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaay 11AUG2022:11:30 kpzx329

Table 3.4.26 PAOLA1: Summary of subgroup analysis of Serious AESI: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + be (N=1)		ımab	•			
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI ths))	C	Number (%) of patients with events	(9!	an ti 5% CI ths))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)											
NED [PDS]	92	1 (1.1)	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(IVRS)											
tBRCAm	150	1 (0.7)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	1 (2.0)	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(eCRF)											
tBRCAm	147	1 (0.7)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	1 (0.5)	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaz 11AUG2022:11:30 kpzx329

Table 3.4.26 PAOLA1: Summary of subgroup analysis of Serious AESI: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		mab	•		Placebo + be (N=1)		nab	•			
Subgroup	C	Number (%) of patients with events	Media (95: (montl	% CI)		(Number (%) of patients with events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	1 (0.5)	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	1 (0.4)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	.ne											
(0) Normal activity	190	1 (0.5)	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	1 (0.4)	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	1 (0.4)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	0	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	1 (1.3)	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaz 11AUG2022:11:30 kpzx329

Table 3.4.26 PAOLA1: Summary of subgroup analysis of Serious AESI: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	-	Olaparib + b (N=2!		ab		Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	(95	an tim % CI) hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	0	NE (1	NE, NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	1 (1.0)	NE (I	NE, NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Myriad tumour BRCA mutati	ion statu	ıs										
tBRCAm	158	1 (0.6)	NE (NE, NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE, NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Status somatic BRCA mutat	cions											
sBRCAm	25	0	NE (NE, NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (I	NE, NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (I	NE, NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaaz 11AUG2022:11:30 kpzx329

Table 3.4.27 PAOLA1: Summary of subgroup analysis of Serious AESI: GI perforations, abscesses and fistulae Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + (N=	bevacizu 255)	mab			Placebo + be (N=1)		mab	•			
Subgroup	C	Number (%) of patients with events	Media (95 (mont)	% CI))	C	Number (%) of patients with events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IVF	RS)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	2 (5.0)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA status	s (IVRS)											
tBRCAm	150	1 (0.7)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	1 (1.0)	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment outo	come (eCF	RF)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	2 (5.1)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA status	s (eCRF)											
tBRCAm	147	1 (0.7)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	1 (0.9)	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	1 (0.5)	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	1 (1.4)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaba 11AUG2022:11:30 kpzx329

Table 3.4.27 PAOLA1: Summary of subgroup analysis of Serious AESI: GI perforations, abscesses and fistulae Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2!)		Placebo + be (N=1		0	:		
Subgroup	C	Number (%) of patients with events	Median (95% ((months)	!I)	(Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	1 (0.5)	NE (NE	, NE)	89	0	NE (N	E, NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE	, NE)	42	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	2 (0.8)	NE (NE	, NE)	125	0	NE (N	E, NE)	NC	NC	NC
Japan	10	0	NE (NE	, NE)	6	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Baseli	.ne									
(0) Normal activity	190	2 (1.1)	NE (NE	, NE)	100	0	NE (N	E, NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE	, NE)	30	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	2 (0.9)	NE (NE	, NE)	117	0	NE (N	E, NE)	NC	NC	NC
>ULN	27	0	NE (NE	, NE)	14	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	2 (0.8)	NE (NE	, NE)	131	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	0	NE (NE	, NE)	80	0	NE (N	E, NE)	NC	NC	NC
Residue	79	2 (2.5)	NE (NE	, NE)	43	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaba 11AUG2022:11:30 kpzx329

Table 3.4.27 PAOLA1: Summary of subgroup analysis of Serious AESI: GI perforations, abscesses and fistulae Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		mab		Placebo + be (N=1		mab				
Subgroup	C	Number (%) of patients with events	(959	n time % CI) ns) [a]		Number (%) of patients with events	(95	an tir 5% CI) :hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive	surgery											
Upfront	146	2 (1.4)	NE (NE, NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	0	NE (NE, NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Myriad tumour BRCA mutat	ion statu	s										
tBRCAm	158	1 (0.6)	NE (NE, NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	1 (1.0)	NE (NE, NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Status somatic BRCA muta	tions											
sBRCAm	25	0	NE (NE, NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE, NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE, NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaba 11AUG2022:11:30 kpzx329

Table 3.4.28 PAOLA1: Summary of subgroup analysis of Serious AESI: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		umab			Placebo + be (N=1)		b			
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI :hs))	C	Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	LS)										
NED [PDS]	92	0	NE (NE,	NE)	48	1 (2.1)	NE (N	E, NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (N	E, NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (N	E, NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR	CA status	(IVRS)										
tBRCAm	150	0	NE (NE,	NE)	65	1 (1.5)	NE (N	E, NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
First line treatment out	come (eCR	EF)										
NED [PDS]	89	0	NE (NE,	NE)	47	1 (2.1)	NE (N	E, NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (N	E, NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (N	E, NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR	CA status	(eCRF)										
tBRCAm	147	0	NE (NE,	NE)	67	1 (1.5)	NE (N	E, NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Age group												
<65 years	185	0	NE (NE,	NE)	98	1 (1.0)	NE (N	E, NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabb 11AUG2022:11:30 kpzx329

Table 3.4.28 PAOLA1: Summary of subgroup analysis of Serious AESI: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2		umab			Placebo + be (N=1)		b	:		
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI ths))	c	Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	0	NE (NE,	NE)	89	1 (1.1)	NE (N	E, NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	0	NE (NE,	NE)	125	1 (0.8)	NE (E, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (E, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	0	NE (NE,	NE)	100	0	NE (N	E, NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	1 (3.3)	NE (E, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	0	NE (NE,	NE)	117	1 (0.9)	NE (N	E, NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	0	NE (NE,	NE)	131	1 (0.8)	NE (E, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	0	NE (NE,	NE)	80	1 (1.3)	NE (E, NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (E, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabb 11AUG2022:11:30 kpzx329

Table 3.4.28 PAOLA1: Summary of subgroup analysis of Serious AESI: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		ımab			Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events		an ti % CI) .hs) [C	Number (%) of patients with events	(95	an ti % CI) .hs) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery												
Upfront	146	0	NE (NE,	NE)	78	1 (1.3)	NE (NE,	NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Myriad tumour BRCA mutat:	ion statu	ıs											
tBRCAm	158	0	NE (NE,	NE)	77	1 (1.3)	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Status somatic BRCA mutat	tions												
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE,	NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabb 11AUG2022:11:30 kpzx329

Table 3.4.29 PAOLA1: Summary of subgroup analysis of Serious AESI: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + k (N=2		b		Placebo + be (N=1)		•			
Subgroup	C	Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	Median tin (95% CI) (months) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVF	RS)									
NED [PDS]	92	2 (2.2)	NE (N	E, NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (N	E, NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (N	E, NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (N	E, NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	(IVRS)									
tBRCAm	150	1 (0.7)	NE (N	E, NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	1 (1.0)	NE (N	E, NE)	66	1 (1.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
First line treatment out	come (eCF	RF)									
NED [PDS]	89	2 (2.2)	NE (N	E, NE)	47	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (N	E, NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (N	E, NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (N	E, NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	(eCRF)									
tBRCAm	147	1 (0.7)	NE (N	E, NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	1 (0.9)	NE (N	E, NE)	64	1 (1.6)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Age group											
<65 years	185	1 (0.5)	NE (N	E, NE)	98	1 (1.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	1 (1.4)	NE (N	E, NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabc 11AUG2022:11:30 kpzx329

Table 3.4.29 PAOLA1: Summary of subgroup analysis of Serious AESI: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2!		ımab	•		Placebo + be (N=1)		ab			
Subgroup	C	Number (%) of patients with events	Media (95 (mont)	% CI)		c	Number (%) of patients with events	Mediar (95% (month	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	2 (1.1)	NE (NE,	NE)	89	1 (1.1)	NE (NE, NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	2 (0.8)	NE (NE,	NE)	125	1 (0.8)	NE (NE, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	0	NE (NE,	NE)	100	1 (1.0)	NE (NE, NE)	NC	NC	NC
(1) Restricted activity	61	1 (1.6)	NE (NE,	NE)	30	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	2 (0.9)	NE (NE,	NE)	117	1 (0.9)	NE (NE, NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	2 (0.8)	NE (NE,	NE)	131	1 (0.8)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	2 (1.2)	NE (NE,	NE)	80	1 (1.3)	NE (NE, NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabc 11AUG2022:11:30 kpzx329

Table 3.4.29 PAOLA1: Summary of subgroup analysis of Serious AESI: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + 1 (N=2	bevacizumab 255)			Placebo + be (N=1)		ab			
Subgroup	C	Number (%) of patients with events	Median ti (95% CI) (months) [C	Number (%) of patients with events	Median (95% (month)	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	2 (1.4)	NE (NE,	NE)	78	1 (1.3)	NE (NE, NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Myriad tumour BRCA mutat:	ion statu	ıs									
tBRCAm	158	2 (1.3)	75.0 (75.0,	NE)	77	0	NE (NE, NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	1 (1.9)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Status somatic BRCA mutat	cions										
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE, NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE,	NE)	36	0	NE (NE, NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabc 11AUG2022:11:30 kpzx329

Table 3.4.30 PAOLA1: Summary of subgroup analysis of Serious AESI: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		ımab			Placebo + be (N=13		mab				
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI)	C	Number (%) of patients with events	(95	an ti 5% CI :hs))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)											
NED [PDS]	92	1 (1.1)	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	1 (5.0)	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	1 (4.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(IVRS)											
tBRCAm	150	1 (0.7)	NE (NE,	NE)	65	1 (1.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	2 (3.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	1 (1.1)	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	1 (5.9)	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	1 (2.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(eCRF)											
tBRCAm	147	1 (0.7)	NE (NE,	NE)	67	1 (1.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	2 (3.1)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	1 (0.5)	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	2 (6.1)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabd 11AUG2022:11:30 kpzx329

Table 3.4.30 PAOLA1: Summary of subgroup analysis of Serious AESI: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=25		ab	÷	Placebo + be (N=1)					
Subgroup	c	Number (%) of patients with events	Median (95% (month	CI)	(Number (%) of patients with events	Median ti (95% CI) (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	1 (0.5)	NE (NE, NE)	89	1 (1.1)	NE (NE,	NE)	NC	NC	NC
IV	73	0	NE (NE, NE)	42	2 (4.8)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	1 (0.4)	NE (NE, NE)	125	3 (2.4)	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE, NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Baseli	ne									
(0) Normal activity	190	1 (0.5)	NE (NE, NE)	100	2 (2.0)	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE, NE)	30	1 (3.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	1 (0.4)	NE (NE, NE)	117	3 (2.6)	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE, NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	1 (0.4)	NE (NE, NE)	131	3 (2.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	1 (0.6)	NE (NE, NE)	80	1 (1.3)	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE, NE)	43	2 (4.7)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabd 11AUG2022:11:30 kpzx329

Table 3.4.30 PAOLA1: Summary of subgroup analysis of Serious AESI: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	1	Olaparib + b (N=2		mab		Placebo + be (N=1)		ıb			
Subgroup	C	Number (%) of patients with events	(959	n time % CI) ns) [a]	(Number (%) of patients with events	Median (95% (month)	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	1 (0.7)	NE (NE, NE)	78	3 (3.8)	NE (NE, NE)	NC	NC	NC
Interval	99	0	NE (NE, NE)	45	0	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Myriad tumour BRCA mutat:	ion statu	ıs									
tBRCAm	158	1 (0.6)	NE (NE, NE)	77	2 (2.6)	NE (NE, NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE, NE)	54	1 (1.9)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Status somatic BRCA muta	cions										
sBRCAm	25	0	NE (NE, NE)	9	0	NE (NE, NE)	NC	NC	NC
gBRCAm	69	0	NE (NE, NE)	36	0	NE (NE, NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE, NE)	23	1 (4.3)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabd 11AUG2022:11:30 kpzx329

Table 3.4.31 PAOLA1: Summary of subgroup analysis of Serious AESI: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		ımab			Placebo + be (N=1)		ımab	•			
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI)	C	Number (%) of patients with events	(95	an ti 5% CI ths))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)											
NED [PDS]	92	1 (1.1)	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	1 (2.0)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(IVRS)											
tBRCAm	150	1 (0.7)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	1 (1.0)	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	1 (1.1)	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(eCRF)											
tBRCAm	147	1 (0.7)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	1 (0.9)	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	2 (1.1)	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabe 11AUG2022:11:30 kpzx329

Table 3.4.31 PAOLA1: Summary of subgroup analysis of Serious AESI: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		ımab			Placebo + be (N=1)		mab	•			
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI; hs) [)	C	Number (%) of patients with events	Media (95 (mont	% CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	1 (0.5)	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	2 (0.8)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	.ne											
(0) Normal activity	190	1 (0.5)	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	1 (1.6)	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	2 (0.9)	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	2 (0.8)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	2 (1.2)	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabe 11AUG2022:11:30 kpzx329

Table 3.4.31 PAOLA1: Summary of subgroup analysis of Serious AESI: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	1	Olaparib + b (N=2			Placebo + bo (N=1					
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	n	Number (%) of patients with events	Median t (95% CI (months)	[)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	1 (0.7)	NE (NE, NE) 78	0	NE (NE	NE)	NC	NC	NC
Interval	99	1 (1.0)	NE (NE, NE) 45	0	NE (NE	NE)	NC	NC	NC
Interaction p-value										NC
Myriad tumour BRCA mutat:	ion statu	ıs								
tBRCAm	158	2 (1.3)	NE (NE, NE) 77	0	NE (NE	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE, NE) 54	0	NE (NE	NE)	NC	NC	NC
Interaction p-value										NC
Status somatic BRCA mutat	cions									
sBRCAm	25	1 (4.0)	NE (NE, NE) 9	0	NE (NE	NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE, NE) 36	0	NE (NE	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE, NE) 23	0	NE (NE	NE)	NC	NC	NC
Interaction p-value										NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabe 11AUG2022:11:30 kpzx329

Table 3.4.32 PAOLA1: Summary of subgroup analysis of Serious AESI: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		ımab			Placebo + be (N=1)		mab				
Subgroup	0	Number (%) f patients ith events	(95	an ti % CI hs))	C	Number (%) of patients with events		an ti % CI; hs) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	S)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(IVRS)											
tBRCAm	150	0	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(eCRF)											
tBRCAm	147	0	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	0	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabf 11AUG2022:11:30 kpzx329

Table 3.4.32 PAOLA1: Summary of subgroup analysis of Serious AESI: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		umab			Placebo + be (N=1)		mab	•			
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI ths))	(Number (%) of patients with events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	0	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	0	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	ne											
(0) Normal activity	190	0	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	0	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	0	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	0	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabf 11AUG2022:11:30 kpzx329

Table 3.4.32 PAOLA1: Summary of subgroup analysis of Serious AESI: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + b (N=1		mab				
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI) :hs) [)	(Number (%) of patients with events	(95	an ti % CI; hs) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery												
Upfront	146	0	NE (NE,	NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Myriad tumour BRCA mutat:	ion statu	ıs											
tBRCAm	158	0	NE (NE,	NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Status somatic BRCA muta	tions												
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE,	NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabf 11AUG2022:11:30 kpzx329

Table 3.4.33 PAOLA1: Summary of subgroup analysis of Serious AESI: MDS/AML Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + (N=	bevacizu 255)	mab			Placebo + be (N=13		ab			
Subgroup	c	Number (%) of patients with events	Media (959 (mont)	% CI)		c	Number (%) of patients with events	Mediar (95% (month	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	RS)										
NED [PDS]	92	2 (2.2)	NE (NE,	NE)	48	1 (2.1)	NE (NE, NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (NE,	NE)	38	2 (5.3)	NE (NE, NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE, NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR	CA status	s (IVRS)										
tBRCAm	150	2 (1.3)	NE (NE,	NE)	65	1 (1.5)	NE (NE, NE)	NC	NC	NC
non-tBRCAm	105	1 (1.0)	NE (NE,	NE)	66	2 (3.0)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
First line treatment out	come (eCR	RF)										
NED [PDS]	89	2 (2.2)	NE (NE,	NE)	47	1 (2.1)	NE (NE, NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (NE,	NE)	32	2 (6.3)	NE (NE, NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE, NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR	CA status	s (eCRF)										
tBRCAm	147	2 (1.4)	NE (NE,	NE)	67	1 (1.5)	NE (NE, NE)	NC	NC	NC
non-tBRCAm	108	1 (0.9)	NE (NE,	NE)	64	2 (3.1)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Age group												
<65 years	185	3 (1.6)	NE (NE,	NE)	98	1 (1.0)	NE (NE, NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	2 (6.1)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabg 11AUG2022:11:30 kpzx329

Table 3.4.33 PAOLA1: Summary of subgroup analysis of Serious AESI: MDS/AML Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

-	(Olaparib + b (N=2!		umab	,		Placebo + be (N=13		ab			
Subgroup	c	Number (%) of patients with events		an ti % CI; hs) [)	c	Number (%) of patients with events	Mediar (95% (month	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	3 (1.6)	NE (NE,	NE)	89	3 (3.4)	NE (NE, NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	3 (1.2)	NE (NE,	NE)	125	3 (2.4)	NE (NE, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	0	NE (NE,	NE)	100	2 (2.0)	NE (NE, NE)	NC	NC	NC
(1) Restricted activity	61	2 (3.3)	NE (NE,	NE)	30	1 (3.3)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	3 (1.3)	NE (NE,	NE)	117	3 (2.6)	NE (NE, NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	3 (1.2)	NE (NE,	NE)	131	3 (2.3)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	me											
No residue	166	3 (1.8)	NE (NE,	NE)	80	3 (3.8)	NE (NE, NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabg 11AUG2022:11:30 kpzx329

Table 3.4.33 PAOLA1: Summary of subgroup analysis of Serious AESI: MDS/AML Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	1	Olaparib + b (N=2			Placebo + b (N=1				
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive :	surgery								
Upfront	146	2 (1.4)	NE (NE, 1	IE) 7	3 1 (1.3)	NE (NE, NI	E) NC	NC	NC
Interval	99	1 (1.0)	NE (NE, 1	IE) 4	5 2 (4.4)	NE (NE, NE	E) NC	NC	NC
Interaction p-value									NC
Myriad tumour BRCA mutat:	ion statu	ıs							
tBRCAm	158	2 (1.3)	NE (NE, 1	IE) 7	7 2 (2.6)	NE (NE, NE	E) NC	NC	NC
Non-tBRCAm	97	1 (1.0)	NE (NE, 1	IE) 5	1 (1.9)	NE (NE, NE	E) NC	NC	NC
Interaction p-value									NC
Status somatic BRCA muta	tions								
sBRCAm	25	1 (4.0)	NE (NE, 1	IE)	9 1 (11.1)	NE (NE, N	E) NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE, 1	IE) 3	5 1 (2.8)	NE (NE, NE	E) NC	NC	NC
Non-BRCAm	43	0	NE (NE, 1	IE) 2	3 1 (4.3)	NE (NE, NE	E) NC	NC	NC
Interaction p-value									NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabg 11AUG2022:11:30 kpzx329

Table 3.4.34 PAOLA1: Summary of subgroup analysis of Serious AESI: Myelodysplastic syndrome and Acute myeloid leukaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2!		ab		Placebo + bo (N=1		ab			
Subgroup	C	Number (%) of patients vith events	Median (95% (months	CI)	C	Number (%) of patients with events	Median (95% (month)	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	LS)									
NED [PDS]	92	2 (2.2)	NE (1	IE, NE)	48	2 (4.2)	NE (NE, NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (1	IE, NE)	38	2 (5.3)	NE (NE, NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (1	IE, NE)	20	0	NE (NE, NE)	NC	NC	NC
PR	49	0	NE (1	IE, NE)	25	0	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	(IVRS)									
tBRCAm	150	2 (1.3)	NE (1	IE, NE)	65	1 (1.5)	NE (NE, NE)	NC	NC	NC
non-tBRCAm	105	1 (1.0)	NE (1	IE, NE)	66	3 (4.5)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
First line treatment out	come (eCR	EF)									
NED [PDS]	89	2 (2.2)	NE (1	IE, NE)	47	2 (4.3)	NE (NE, NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (1	IE, NE)	32	2 (6.3)	NE (NE, NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (1	IE, NE)	17	0	NE (NE, NE)	NC	NC	NC
PR	50	0	NE (IE, NE)	34	0	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	(eCRF)									
tBRCAm	147	2 (1.4)	NE (1	IE, NE)	67	1 (1.5)	NE (NE, NE)	NC	NC	NC
non-tBRCAm	108	1 (0.9)	NE (1	IE, NE)	64	3 (4.7)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Age group											
<65 years	185	3 (1.6)	NE (1	IE, NE)	98	2 (2.0)	NE (NE, NE)	NC	NC	NC
>=65 years	70	0	NE (1	IE, NE)	33	2 (6.1)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabh 11AUG2022:11:30 kpzx329

Table 3.4.34 PAOLA1: Summary of subgroup analysis of Serious AESI: Myelodysplastic syndrome and Acute myeloid leukaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		umab	•		Placebo + be (N=13		ab			
Subgroup	c	Number (%) of patients with events		an ti % CI; hs) [)	C	Number (%) of patients with events	Mediar (95% (month	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	3 (1.6)	NE (NE,	NE)	89	4 (4.5)	NE (NE, NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	3 (1.2)	NE (NE,	NE)	125	4 (3.2)	NE (NE, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	0	NE (NE,	NE)	100	3 (3.0)	NE (NE, NE)	NC	NC	NC
(1) Restricted activity	61	2 (3.3)	NE (NE,	NE)	30	1 (3.3)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	3 (1.3)	NE (NE,	NE)	117	4 (3.4)	NE (NE, NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	3 (1.2)	NE (NE,	NE)	131	4 (3.1)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	3 (1.8)	NE (NE,	NE)	80	4 (5.0)	NE (NE, NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabh 11AUG2022:11:30 kpzx329

Table 3.4.34 PAOLA1: Summary of subgroup analysis of Serious AESI: Myelodysplastic syndrome and Acute myeloid leukaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2				Placebo + be (N=1)		b			
Subgroup	C	Number (%) of patients with events	Median tim (95% CI) (months) [a		C	Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	2 (1.4)	NE (NE,	NE)	78	2 (2.6)	NE (N	E, NE)	NC	NC	NC
Interval	99	1 (1.0)	NE (NE,	NE)	45	2 (4.4)	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Myriad tumour BRCA mutat:	ion statu	ıs									
tBRCAm	158	2 (1.3)	NE (NE,	NE)	77	3 (3.9)	NE (N	E, NE)	NC	NC	NC
Non-tBRCAm	97	1 (1.0)	NE (NE,	NE)	54	1 (1.9)	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Status somatic BRCA mutat	cions										
sBRCAm	25	1 (4.0)	NE (NE,	NE)	9	1 (11.1)	NE (N	E, NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE,	NE)	36	2 (5.6)	NE (N	E, NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	1 (4.3)	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabh 11AUG2022:11:30 kpzx329

Table 3.4.35 PAOLA1: Summary of subgroup analysis of Serious AESI: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	;	Olaparib + b (N=2		ab		Placebo + be (N=1)	:		
Subgroup	(Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	Median (95% ((months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVF	RS)									
NED [PDS]	92	5 (5.4)	NE (N	IE, NE)	48	2 (4.2)	NE (N	E, NE)	NC	NC	NC
NED/CR [IDS]	74	6 (8.1)	NE (N	IE, NE)	38	1 (2.6)	NE (N	E, NE)	NC	NC	NC
NED/CR [Chemo]	40	4 (10.0)	NE (N	IE, NE)	20	1 (5.0)	NE (N	E, NE)	NC	NC	NC
PR	49	0	NE (N	IE, NE)	25	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	s (IVRS)									
tBRCAm	150	13 (8.7)	NE (N	IE, NE)	65	3 (4.6)	NE (N	E, NE)	1.68	0.54, 7.31	0.3956
non-tBRCAm	105	2 (1.9)	NE (N	IE, NE)	66	1 (1.5)	NE (N	E, NE)	1.14	0.11, 24.54	0.9137
Interaction p-value											0.7833
First line treatment out	come (eCF	RF)									
NED [PDS]	89	6 (6.7)	NE (N	IE, NE)	47	2 (4.3)	NE (N	E, NE)	NC	NC	NC
NED/CR [IDS]	74	6 (8.1)	NE (N	IE, NE)	32	0	NE (N	E, NE)	NC	NC	NC
NED/CR [Chemo]	39	3 (7.7)	NE (N	IE, NE)	17	1 (5.9)	NE (N	E, NE)	NC	NC	NC
PR	50	0	NE (N	IE, NE)	34	1 (2.9)	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	s (eCRF)									
tBRCAm	147	13 (8.8)	NE (N	IE, NE)	67	3 (4.5)	NE (N	E, NE)	1.74	0.56, 7.60	0.3605
non-tBRCAm	108	2 (1.9)	NE (N	IE, NE)	64	1 (1.6)	NE (N	E, NE)	1.09	0.10, 23.35	0.9463
Interaction p-value											0.7367
Age group											
<65 years	185	13 (7.0)	NE (N	IE, NE)	98	3 (3.1)	NE (N	E, NE)	2.13	0.69, 9.29	0.2042
>=65 years	70	2 (2.9)	NE (N	IE, NE)	33	1 (3.0)	NE (N	E, NE)	0.74	0.07, 15.91	0.8085
Interaction p-value											0.4631

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabi 11AUG2022:11:30 kpzx329

Table 3.4.35 PAOLA1: Summary of subgroup analysis of Serious AESI: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2				Placebo + be (N=13					
Subgroup		Number (%) of patients with events	Median to (95% CI (months))	C	Number (%) of patients with events	Median tin (95% CI) (months) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	10 (5.5)	NE (NE,	NE)	89	3 (3.4)	NE (NE,	NE)	1.46	0.45, 6.51	0.5539
IV	73	5 (6.8)	NE (NE,	NE)	42	1 (2.4)	NE (NE,	NE)	2.57	0.41, 49.34	0.3422
Interaction p-value											0.6481
Region											
Europe	245	14 (5.7)	NE (NE,	NE)	125	4 (3.2)	NE (NE,	NE)	1.59	0.57, 5.62	0.3937
Japan	10	1 (10.0)	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel	ine									
(0) Normal activity	190	8 (4.2)	NE (NE,	NE)	100	3 (3.0)	NE (NE,	NE)	1.27	0.37, 5.80	0.7190
(1) Restricted activity	61	7 (11.5)	NE (NE,	NE)	30	1 (3.3)	NE (NE,	NE)	2.91	0.52, 54.37	0.2574
Interaction p-value											0.4960
Baseline CA-125 value											
<=ULN	228	15 (6.6)	NE (NE,	,	117	4 (3.4)	NE (NE,	NE)	1.66	0.60, 5.82	0.3477
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	15 (5.9)	NE (NE,	NE)	131	4 (3.1)	NE (NE,	NE)	1.71	0.62, 6.01	0.3157
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	12 (7.2)	NE (NE,	NE)	80	2 (2.5)	NE (NE,	NE)	2.55	0.69, 16.40	0.1732
Residue	79	3 (3.8)	NE (NE,	NE)	43	2 (4.7)	NE (NE,	NE)	0.73	0.12, 5.52	0.7299
Interaction p-value											0.2929

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabi 11AUG2022:11:30 kpzx329

Table 3.4.35 PAOLA1: Summary of subgroup analysis of Serious AESI: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		Placebo + b (N=1				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery							
Upfront	146	9 (6.2)	NE (NE, NE)	78 3 (3.8)	NE (NE, NE)	1.33	0.40, 6.01	0.6596
Interval	99	6 (6.1)	NE (NE, NE)	45 1 (2.2)	NE (NE, NE)	2.62	0.45, 49.47	0.3204
Interaction p-value								0.5821
Myriad tumour BRCA mutat:	ion stat	us						
tBRCAm	158	13 (8.2)	NE (NE, NE)	77 4 (5.2)	NE (NE, NE)	1.37	0.48, 4.86	0.5734
Non-tBRCAm	97	2 (2.1)	NE (NE, NE)	54 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC
Status somatic BRCA muta	cions							
sBRCAm	25	1 (4.0)	NE (NE, NE)	9 0	NE (NE, NE)	NC	NC	NC
gBRCAm	69	6 (8.7)	NE (NE, NE)	36 2 (5.6)	NE (NE, NE)	NC	NC	NC
Non-BRCAm	43	1 (2.3)	NE (NE, NE)	23 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabi 11AUG2022:11:30 kpzx329

Table 3.4.36 PAOLA1: Summary of subgroup analysis of Serious AESI: Pneumonitis Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		ımab			Placebo + be (N=1)		ımab	•			
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI))	C	Number (%) of patients with events	(95	an ti 5% CI ths))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	1 (2.0)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(IVRS)											
tBRCAm	150	0	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	2 (1.9)	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(eCRF)											
tBRCAm	147	0	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	2 (1.9)	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	0	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	2 (2.9)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabj 11AUG2022:11:30 kpzx329

Table 3.4.36 PAOLA1: Summary of subgroup analysis of Serious AESI: Pneumonitis Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2!		ımab	,		Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI))	(Number (%) of patients with events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	2 (1.1)	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	2 (0.8)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	.ne											
(0) Normal activity	190	2 (1.1)	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	1 (0.4)	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	1 (3.7)	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	2 (0.8)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	2 (1.2)	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabj 11AUG2022:11:30 kpzx329

Table 3.4.36 PAOLA1: Summary of subgroup analysis of Serious AESI: Pneumonitis Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	1	Olaparib + b (N=2		ab		Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	(95	an tim % CI) hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	0	NE (1	IE, NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	2 (2.0)	NE (1	IE, NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Myriad tumour BRCA mutat:	ion statu	ıs										
tBRCAm	158	1 (0.6)	NE (1	IE, NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	1 (1.0)	NE (1	IE, NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Status somatic BRCA mutat	tions											
sBRCAm	25	0	NE (1	IE, NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (1	IE, NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	1 (2.3)	NE (1	IE, NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabj 11AUG2022:11:30 kpzx329

Table 3.4.37 PAOLA1: Summary of subgroup analysis of AESI G>=3: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	-	Olaparib + b (N=2		ımab			Placebo + be (N=1)		ımab				
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI)	C	Number (%) of patients with events	(95	an ti 5% CI :hs))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IV)	RS)											
NED [PDS]	92	18 (19.6)	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	10.12	2.09,182.11	0.0014*
NED/CR [IDS]	74	14 (18.9)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	7 (17.5)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	8 (16.3)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA statu	s (IVRS)											
tBRCAm	150	23 (15.3)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	24 (22.9)	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	17.16	3.63,306.70	<0.0001*
Interaction p-value													NC
First line treatment out	come (eCl	RF)											
NED [PDS]	89	19 (21.3)	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	10.94	2.27,196.60	0.0008*
NED/CR [IDS]	74	16 (21.6)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	5 (12.8)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	7 (14.0)	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA statu	s (eCRF)											
tBRCAm	147	23 (15.6)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	24 (22.2)	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	16.13	3.41,288.17	<0.0001*
Interaction p-value													NC
Age group													
<65 years	185	32 (17.3)	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	18.40	3.96,327.26	<0.0001*
>=65 years	70	15 (21.4)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabk 11AUG2022:11:30 kpzx329

Table 3.4.37 PAOLA1: Summary of subgroup analysis of AESI G>=3: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		•		Placebo + be (N=13		•			
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))	C	Number (%) of patients with events	Median tim (95% CI) (months) [a		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	35 (19.2)	NE (NE,	NE)	89	1 (1.1)	NE (NE,	NE)	18.77	4.06,333.36	<0.0001*
IV	73	12 (16.4)	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	43 (17.6)	NE (NE,	NE)	125	1 (0.8)	NE (NE,	NE)	23.95	5.23,424.41	<0.0001*
Japan	10	4 (40.0)	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Basel:	ine									
(0) Normal activity	190	33 (17.4)	NE (NE,	NE)	100	1 (1.0)	NE (NE,	NE)	18.95	4.09,336.90	<0.0001*
(1) Restricted activity	61	14 (23.0)	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	44 (19.3)	NE (NE,	NE)	117	1 (0.9)	NE (NE,	NE)	24.98	5.46,442.61	<0.0001*
>ULN	27	3 (11.1)	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	47 (18.4)	NE (NE,	NE)	131	1 (0.8)	NE (NE,	NE)	26.53	5.81,469.83	<0.0001*
Interaction p-value											NC
Cytoreductive surgery outco	me										
No residue	166	35 (21.1)	NE (NE,	NE)	80	1 (1.3)	NE (NE,	NE)	18.74	4.05,332.85	<0.0001*
Residue	79	8 (10.1)	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabk 11AUG2022:11:30 kpzx329

Table 3.4.37 PAOLA1: Summary of subgroup analysis of AESI G>=3: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		Placebo + b (N=1			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	26 (17.8)	NE (NE, NE)	78 1 (1.3)	NE (NE, NE)	15.00 3.19,267.6	0 <0.0001*
Interval	99	17 (17.2)	NE (NE, NE)	45 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Myriad tumour BRCA mutat:	ion stat	us					
tBRCAm	158	27 (17.1)	NE (NE, NE)	77 0	NE (NE, NE)	NC NC	NC
Non-tBRCAm	97	20 (20.6)	NE (NE, NE)	54 1 (1.9)	NE (NE, NE)	12.55 2.62,225.0	5 0.0003*
Interaction p-value							NC
Status somatic BRCA mutat	cions						
sBRCAm	25	3 (12.0)	NE (NE, NE)	9 0	NE (NE, NE)	NC NC	NC
gBRCAm	69	10 (14.5)	NE (NE, NE)	36 0	NE (NE, NE)	NC NC	NC
Non-BRCAm	43	10 (23.3)	NE (NE, NE)	23 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabk 11AUG2022:11:30 kpzx329

Table 3.4.38 PAOLA1: Summary of subgroup analysis of AESI G>=3: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ımab			Placebo + be (N=13		ımab				
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI)	C	Number (%) of patients with events	(95	an ti 5% CI ths))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVI	RS)											
NED [PDS]	92	5 (5.4)	NE (NE,	NE)	48	3 (6.3)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	8 (10.8)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	4 (10.0)	NE (NE,	NE)	20	1 (5.0)	NE (NE,	NE)	NC	NC	NC
PR	49	4 (8.2)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	s (IVRS)											
tBRCAm	150	11 (7.3)	NE (NE,	NE)	65	2 (3.1)	NE (NE,	NE)	2.39	0.64, 15.43	0.2133
non-tBRCAm	105	10 (9.5)	NE (NE,	NE)	66	2 (3.0)	NE (NE,	NE)	3.21	0.85, 20.86	0.0911
Interaction p-value													0.7873
First line treatment out	come (eCI	RF)											
NED [PDS]	89	7 (7.9)	NE (NE,	NE)	47	3 (6.4)	NE (NE,	NE)	1.20	0.33, 5.55	0.7921
NED/CR [IDS]	74	8 (10.8)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	1 (2.6)	NE (NE,	NE)	17	1 (5.9)	NE (NE,	NE)	0.45	0.02, 11.31	0.5748
PR	50	5 (10.0)	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													0.5354
Screening laboratory tBR0	CA status	s (eCRF)											
tBRCAm	147	11 (7.5)	NE (NE,	NE)	67	2 (3.0)	NE (NE,	NE)	2.51	0.67, 16.20	0.1865
non-tBRCAm	108	10 (9.3)	NE (NE,	NE)	64	2 (3.1)	NE (NE,	NE)	3.02	0.80, 19.68	0.1102
Interaction p-value													0.8634
Age group													
<65 years	185	14 (7.6)	NE (NE,	NE)	98	4 (4.1)	NE (NE,	NE)	1.87	0.67, 6.59	0.2450
>=65 years	70	7 (10.0)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabl 11AUG2022:11:30 kpzx329

Table 3.4.38 PAOLA1: Summary of subgroup analysis of AESI G>=3: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		mab			Placebo + be (N=13		ab			
Subgroup		Number (%) of patients with events	(95%	n time k CI) ns) [a]		С	Number (%) of patients with events	Median (95% (month		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	17 (9.3)	NE (NE, N	JE)	89	3 (3.4)	NE (NE, NE)	2.81	0.94, 12.03	0.0649
IV	73	4 (5.5)	NE (NE, N	JE)	42	1 (2.4)	NE (NE, NE)	2.30	0.34, 44.90	0.4226
Interaction p-value												0.8764
Region												
Europe	245	20 (8.2)	NE (NE,	1E)	125	4 (3.2)	NE (NE, NE)	2.56	0.97, 8.81	0.0584
Japan	10	1 (10.0)	NE (NE,	1E)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Basel	ine										
(0) Normal activity	190	11 (5.8)	NE (NE, N	1E)	100	4 (4.0)	NE (NE, NE)	1.43	0.49, 5.17	0.5263
(1) Restricted activity	61	9 (14.8)	NE (NE,	1E)	30	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	18 (7.9)	NE (NE, N	1E)	117	4 (3.4)	NE (NE, NE)	2.32	0.87, 8.04	0.0978
>ULN	27	3 (11.1)	NE (NE, N	1E)	14	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	21 (8.2)	NE (NE,	1E)	131	4 (3.1)	NE (NE, NE)	2.72	1.04, 9.33	0.0415*
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	15 (9.0)	NE (NE,	JE)	80	3 (3.8)	NE (NE, NE)	2.42	0.80, 10.44	0.1261
Residue	79	4 (5.1)	NE (NE,	1E)	43	1 (2.3)	NE (NE, NE)	2.19	0.32, 42.92	0.4499
Interaction p-value												0.9403

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabl 11AUG2022:11:30 kpzx329

Table 3.4.38 PAOLA1: Summary of subgroup analysis of AESI G>=3: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2				Placebo + be (N=1)		ab			
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))	C	Number (%) of patients with events	(95%	n time (CI) (s) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	urgery										
Upfront	146	10 (6.8)	NE (NE,	NE)	78	4 (5.1)	NE (NE, NE)	1.32	0.44, 4.81	0.6325
Interval	99	9 (9.1)	NE (NE,	NE)	45	0	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Myriad tumour BRCA mutati	on stat	us									
tBRCAm	158	12 (7.6)	NE (NE,	NE)	77	3 (3.9)	NE (NE, NE)	1.92	0.61, 8.45	0.2811
Non-tBRCAm	97	9 (9.3)	NE (NE,	NE)	54	1 (1.9)	NE (NE, NE)	5.23	0.98, 96.34	0.0531
Interaction p-value											0.3949
Status somatic BRCA mutat	ions										
sBRCAm	25	5 (20.0)	NE (NE,	NE)	9	0	NE (NE, NE)	NC	NC	NC
gBRCAm	69	3 (4.3)	NE (NE,	NE)	36	1 (2.8)	NE (NE, NE)	NC	NC	NC
Non-BRCAm	43	5 (11.6)	NE (NE,	NE)	23	0	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabl 11AUG2022:11:30 kpzx329

Table 3.4.39 PAOLA1: Summary of subgroup analysis of AESI G>=3: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	,	Olaparib + 1 (N=2				Placebo + bo (N=1					
Subgroup	C	Number (%) of patients with events	Median t (95% C (months)	I)	C	Number (%) of patients with events	Median ((95% C (months)	I)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVF	RS)									
NED [PDS]	92	2 (2.2)	NE (NE	, NE)	48	0	NE (NE	, NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (NE	, NE)	38	4 (10.5)	NE (NE	, NE)	NC	NC	NC
NED/CR [Chemo]	40	1 (2.5)	NE (NE	, NE)	20	0	NE (NE	, NE)	NC	NC	NC
PR	49	0	NE (NE	, NE)	25	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	(IVRS)									
tBRCAm	150	1 (0.7)	NE (NE	, NE)	65	3 (4.6)	NE (NE	, NE)	NC	NC	NC
non-tBRCAm	105	4 (3.8)	NE (NE	, NE)	66	1 (1.5)	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
First line treatment out	come (eCF	RF)									
NED [PDS]	89	2 (2.2)	NE (NE	, NE)	47	0	NE (NE	, NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (NE	, NE)	32	3 (9.4)	NE (NE	, NE)	NC	NC	NC
NED/CR [Chemo]	39	1 (2.6)	NE (NE	, NE)	17	0	NE (NE	, NE)	NC	NC	NC
PR	50	0	NE (NE	, NE)	34	1 (2.9)	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	s (eCRF)									
tBRCAm	147	1 (0.7)	NE (NE	, NE)	67	3 (4.5)	NE (NE	, NE)	NC	NC	NC
non-tBRCAm	108	4 (3.7)	NE (NE	, NE)	64	1 (1.6)	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Age group											
<65 years	185	4 (2.2)	NE (NE	, NE)	98	3 (3.1)	NE (NE	, NE)	NC	NC	NC
>=65 years	70	1 (1.4)	NE (NE	, NE)	33	1 (3.0)	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabm 11AUG2022:11:30 kpzx329

Table 3.4.39 PAOLA1: Summary of subgroup analysis of AESI G>=3: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=25		mab		Placebo + be (N=1)					
Subgroup	c	Number (%) of patients with events	(95%	n time k CI) ns) [a]		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	4 (2.2)	NE (NE, NE	3) 89	2 (2.2)	NE (NE,	NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE, NE	1) 42	2 (4.8)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	5 (2.0)	NE (NE, NE	125	4 (3.2)	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE, NE	:) 6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Baseli	.ne									
(0) Normal activity	190	3 (1.6)	NE (NE, NE	100	3 (3.0)	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	2 (3.3)	NE (NE, NE	30	1 (3.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	5 (2.2)	NE (NE, NE	117	4 (3.4)	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE, NE	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	5 (2.0)	NE (NE, NE	131	4 (3.1)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Cytoreductive surgery outco	me										
No residue	166	4 (2.4)	NE (NE, NE	80	3 (3.8)	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE, NE	3) 43	1 (2.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabm 11AUG2022:11:30 kpzx329

Table 3.4.39 PAOLA1: Summary of subgroup analysis of AESI G>=3: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	. (Olaparib + b (N=2!			Placebo + be (N=1					
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	(Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive	surgery									
Upfront	146	2 (1.4)	NE (NE, NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	2 (2.0)	NE (NE, NE)	45	4 (8.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Myriad tumour BRCA mutat	ion statu	ıs								
tBRCAm	158	1 (0.6)	NE (NE, NE)	77	3 (3.9)	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	4 (4.1)	NE (NE, NE)	54	1 (1.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Status somatic BRCA muta	tions									
sBRCAm	25	0	NE (NE, NE)	9	1 (11.1)	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE, NE)	36	2 (5.6)	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	2 (4.7)	NE (NE, NE)	23	1 (4.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabm 11AUG2022:11:30 kpzx329

Table 3.4.40 PAOLA1: Summary of subgroup analysis of AESI G>=3: Nausea Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		b		Placebo + be (N=1)					
Subgroup	c	Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	Median t (95% CI (months)	()	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	RS)									
NED [PDS]	92	1 (1.1)	NE (N	E, NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	5 (6.8)	NE (N	E, NE)	38	2 (5.3)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	1 (2.5)	NE (N	E, NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	2 (4.1)	NE (N	E, NE)	25	2 (8.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	(IVRS)									
tBRCAm	150	7 (4.7)	NE (N	E, NE)	65	2 (3.1)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	2 (1.9)	NE (N	E, NE)	66	2 (3.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
First line treatment out	come (eCR	RF)									
NED [PDS]	89	1 (1.1)	NE (N	E, NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	4 (5.4)	NE (N	E, NE)	32	2 (6.3)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	1 (2.6)	NE (N	E, NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	3 (6.0)	NE (N	E, NE)	34	2 (5.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	(eCRF)									
tBRCAm	147	7 (4.8)	NE (N	E, NE)	67	2 (3.0)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	2 (1.9)	NE (N	E, NE)	64	2 (3.1)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Age group											
<65 years	185	8 (4.3)	NE (N	E, NE)	98	3 (3.1)	NE (NE,	NE)	1.41	0.41, 6.42	0.6053
>=65 years	70	1 (1.4)	NE (N	E, NE)	33	1 (3.0)	NE (NE,	NE)	0.46	0.02, 11.57	0.5855
Interaction p-value											0.4792

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabn 11AUG2022:11:30 kpzx329

Table 3.4.40 PAOLA1: Summary of subgroup analysis of AESI G>=3: Nausea Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2!		nab	:	Placebo + bo (N=1					
Subgroup	C	Number (%) of patients with events	(95%	n time (CI) (s) [a]		Number (%) of patients with events	Median to (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	7 (3.8)	NE (NE, NI	E) 89	3 (3.4)	NE (NE,	NE)	1.13	0.31, 5.26	0.8552
IV	73	2 (2.7)	NE (NE, NI	E) 42	1 (2.4)	NE (NE,	NE)	1.13	0.11, 24.41	0.9170
Interaction p-value											0.9990
Region											
Europe	245	9 (3.7)	NE (NE, NI	E) 125	3 (2.4)	NE (NE,	NE)	1.52	0.45, 6.86	0.5156
Japan	10	0	NE (NE, NI	E) 6	1 (16.7)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Baseli	.ne									
(0) Normal activity	190	7 (3.7)	NE (NE, NI	E) 100	3 (3.0)	NE (NE,	NE)	1.23	0.34, 5.69	0.7653
(1) Restricted activity	61	2 (3.3)	NE (NE, NI	E) 30	1 (3.3)	NE (NE,	NE)	0.95	0.09, 20.42	0.9663
Interaction p-value											0.8570
Baseline CA-125 value											
<=ULN	228	8 (3.5)	NE (NE, N	E) 117	2 (1.7)	NE (NE,	NE)	2.03	0.51, 13.46	0.3386
>ULN	27	1 (3.7)	NE (NE, NI	E) 14	2 (14.3)	NE (NE,	NE)	0.25	0.01, 2.66	0.2452
Interaction p-value											0.1366
Histological grade											
High grade	255	9 (3.5)	NE (NE, NI	E) 131	4 (3.1)	NE (NE,	NE)	1.14	0.37, 4.22	0.8218
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	5 (3.0)	NE (NE, NI	E) 80	2 (2.5)	NE (NE,	NE)	NC	NC	NC
Residue	79	4 (5.1)	NE (NE, NI	E) 43	1 (2.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabn 11AUG2022:11:30 kpzx329

Table 3.4.40 PAOLA1: Summary of subgroup analysis of AESI G>=3: Nausea Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		Placebo + b (N=1			
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI	2-sided [b] p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	4 (2.7)	NE (NE, NE)	78 0	NE (NE, NE)	NC NC	NC
Interval	99	5 (5.1)	NE (NE, NE)	45 3 (6.7)	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Myriad tumour BRCA mutat:	ion statu	ıs					
tBRCAm	158	7 (4.4)	NE (NE, NE)	77 2 (2.6)	NE (NE, NE)	NC NC	NC
Non-tBRCAm	97	2 (2.1)	NE (NE, NE)	54 2 (3.7)	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Status somatic BRCA mutat	tions						
sBRCAm	25	0	NE (NE, NE)	9 0	NE (NE, NE)	NC NC	NC
gBRCAm	69	4 (5.8)	NE (NE, NE)	36 2 (5.6)	NE (NE, NE)	NC NC	NC
Non-BRCAm	43	2 (4.7)	NE (NE, NE)	23 1 (4.3)	NE (NE, NE)	NC NC	NC
Interaction p-value							NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabn 11AUG2022:11:30 kpzx329

Table 3.4.41 PAOLA1: Summary of subgroup analysis of AESI G>=3: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(-	bevacizumab 255)			Placebo + be (N=1)					
Subgroup	C	Number (%) of patients vith events	Median ti (95% CI (months))	C	Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	RS)									
NED [PDS]	92	0	NE (NE,	NE)	48	2 (4.2)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	3 (4.1)	NE (NE,	NE)	38	2 (5.3)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	1 (5.0)	NE (NE,	NE)	NC	NC	NC
PR	49	1 (2.0)	NE (NE,	NE)	25	1 (4.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	s (IVRS)									
tBRCAm	150	3 (2.0)	NE (NE,	NE)	65	3 (4.6)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	1 (1.0)	NE (NE,	NE)	66	3 (4.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
First line treatment out	come (eCR	RF)									
NED [PDS]	89	0	NE (NE,	NE)	47	2 (4.3)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	3 (4.1)	NE (NE,	NE)	32	2 (6.3)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	1 (5.9)	NE (NE,	NE)	NC	NC	NC
PR	50	1 (2.0)	NE (NE,	NE)	34	1 (2.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	(eCRF)									
tBRCAm	147	3 (2.0)	NE (NE,	NE)	67	3 (4.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	1 (0.9)	NE (NE,	NE)	64	3 (4.7)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Age group											
<65 years	185	4 (2.2)	NE (NE,	NE)	98	5 (5.1)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	1 (3.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabo 11AUG2022:11:30 kpzx329

Table 3.4.41 PAOLA1: Summary of subgroup analysis of AESI G>=3: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=25		ımab	•		Placebo + be (N=13		ab				
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI)		C	Number (%) of patients with events	(95%	n time (CI) (S) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	4 (2.2)	NE (NE,	NE)	89	5 (5.6)	NE (NE, NE) NC	NC		NC
IV	73	0	NE (NE,	NE)	42	1 (2.4)	NE (NE, NE) NC	NC		NC
Interaction p-value													NC
Region													
Europe	245	4 (1.6)	NE (NE,	NE)	125	4 (3.2)	NE (NE, NE) NC	NC		NC
Japan	10	0	NE (NE,	NE)	6	2 (33.3)	NE (NE, NE) NC	NC		NC
Interaction p-value													NC
ECOG performance status at	Baseli	ne											
(0) Normal activity	190	4 (2.1)	NE (NE,	NE)	100	4 (4.0)	NE (NE, NE) NC	NC		NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	2 (6.7)	NE (NE, NE) NC	NC		NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	4 (1.8)	NE (NE,	NE)	117	5 (4.3)	NE (NE, NE) NC	NC		NC
>ULN	27	0	NE (NE,	NE)	14	1 (7.1)	NE (NE, NE) NC	NC		NC
Interaction p-value													NC
Histological grade													
High grade	255	4 (1.6)	NE (NE,	NE)	131	6 (4.6)	NE (NE, NE	0.33	0.08,	1.14	0.0792
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	3 (1.8)	NE (NE,	NE)	80	4 (5.0)	NE (NE, NE) NC	NC		NC
Residue	79	1 (1.3)	NE (NE,	NE)	43	2 (4.7)	NE (NE, NE) NC	NC		NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabo 11AUG2022:11:30 kpzx329

Table 3.4.41 PAOLA1: Summary of subgroup analysis of AESI G>=3: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2				Placebo + be (N=13		ab			
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a		C	Number (%) of patients with events	(95%	n time (CI) (S) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive :	surgery										
Upfront	146	1 (0.7)	NE (NE,	NE)	78	3 (3.8)	NE (NE, NE)	NC	NC	NC
Interval	99	3 (3.0)	NE (NE,	NE)	45	3 (6.7)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Myriad tumour BRCA mutat:	ion statu	ıs									
tBRCAm	158	3 (1.9)	NE (NE,	NE)	77	3 (3.9)	NE (NE, NE)	NC	NC	NC
Non-tBRCAm	97	1 (1.0)	NE (NE,	NE)	54	3 (5.6)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Status somatic BRCA muta	tions										
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE, NE)	NC	NC	NC
gBRCAm	69	2 (2.9)	NE (NE,	NE)	36	2 (5.6)	NE (NE, NE)	NC	NC	NC
Non-BRCAm	43	1 (2.3)	NE (NE,	NE)	23	2 (8.7)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabo 11AUG2022:11:30 kpzx329

Table 3.4.42 PAOLA1: Summary of subgroup analysis of AESI G>=3: Fatigue and Asthenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	:	Olaparib + b (N=2		mab			Placebo + be (N=13		mab				
Subgroup	(Number (%) of patients with events	Media (95% (month	% CI)		C	Number (%) of patients with events	(95	an ti 5% CI :hs))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVF	RS)											
NED [PDS]	92	4 (4.3)	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	6 (8.1)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	3 (7.5)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	4 (8.2)	NE (NE,	NE)	25	2 (8.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	s (IVRS)											
tBRCAm	150	9 (6.0)	NE (NE,	NE)	65	2 (3.1)	NE (NE,	NE)	1.96	0.51, 12.87	0.3566
non-tBRCAm	105	8 (7.6)	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	5.15	0.94, 95.50	0.0597
Interaction p-value													0.4513
First line treatment out	come (eCF	RF)											
NED [PDS]	89	4 (4.5)	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	5 (6.8)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	3 (7.7)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	5 (10.0)	NE (NE,	NE)	34	3 (8.8)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	s (eCRF)											
tBRCAm	147	9 (6.1)	NE (NE,	NE)	67	2 (3.0)	NE (NE,	NE)	2.06	0.53, 13.52	0.3203
non-tBRCAm	108	8 (7.4)	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	4.85	0.89, 90.03	0.0712
Interaction p-value													0.5046
Age group													
<65 years	185	10 (5.4)	NE (NE,	NE)	98	2 (2.0)	NE (NE,	NE)	2.67	0.70, 17.34	0.1617
>=65 years	70	7 (10.0)	NE (NE,	NE)	33	1 (3.0)	NE (NE,	NE)	3.39	0.60, 63.46	0.1865
Interaction p-value													0.8534

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabp 11AUG2022:11:30 kpzx329

Table 3.4.42 PAOLA1: Summary of subgroup analysis of AESI G>=3: Fatigue and Asthenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ımab	•		Placebo + be (N=13		nab	,		
Subgroup		Number (%) of patients with events		an ti % CI) hs) [C	Number (%) of patients with events	(95	n time % CI) ns) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	14 (7.7)	NE (NE,	NE)	89	1 (1.1)	NE (NE, NE) 6.93	1.39,125.60	0.0133*
IV	73	3 (4.1)	NE (NE,	NE)	42	2 (4.8)	NE (NE, NE) 0.88	0.15, 6.65	0.8857
Interaction p-value												0.1190
Region												
Europe	245	17 (6.9)	NE (NE,	NE)	125	3 (2.4)	NE (NE, NE) 2.93	0.98, 12.54	0.0540
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Basel	ine										
(0) Normal activity	190	10 (5.3)	NE (NE,	NE)	100	1 (1.0)	NE (NE, NE) 5.28	1.01, 96.89	0.0482*
(1) Restricted activity	61	7 (11.5)	NE (NE,	NE)	30	2 (6.7)	NE (NE, NE) 1.80	0.44, 12.09	0.4403
Interaction p-value												0.4008
Baseline CA-125 value												
<=ULN	228	16 (7.0)	NE (NE,	NE)	117	2 (1.7)	NE (NE, NE	,	1.18, 26.29	0.0236*
>ULN	27	1 (3.7)	NE (NE,	NE)	14	1 (7.1)	NE (NE, NE) 0.53	0.02, 13.27	0.6516
Interaction p-value												0.2112
Histological grade												
High grade	255	17 (6.7)	NE (NE,	NE)	131	3 (2.3)	NE (NE, NE) 2.95	0.99, 12.63	0.0523
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	9 (5.4)	NE (NE,	NE)	80	0	NE (NE, NE) NC	NC	NC
Residue	79	6 (7.6)	NE (NE,	NE)	43	2 (4.7)	NE (NE, NE) NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabp 11AUG2022:11:30 kpzx329

Table 3.4.42 PAOLA1: Summary of subgroup analysis of AESI G>=3: Fatigue and Asthenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2				Placebo + be (N=13		ab			
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a		O	Number (%) of patients with events	(95%	n time (CI) (s) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	8 (5.5)	NE (NE,	NE)	78	2 (2.6)	NE (NE, NE)	2.10	0.53, 13.94	0.3136
Interval	99	7 (7.1)	NE (NE,	NE)	45	0	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Myriad tumour BRCA mutat:	ion statu	ıs									
tBRCAm	158	9 (5.7)	NE (NE,	NE)	77	2 (2.6)	NE (NE, NE)	2.19	0.56, 14.35	0.2793
Non-tBRCAm	97	8 (8.2)	NE (NE,	NE)	54	1 (1.9)	NE (NE, NE)	4.62	0.85, 85.68	0.0821
Interaction p-value											0.5614
Status somatic BRCA mutat	cions										
sBRCAm	25	1 (4.0)	NE (NE,	NE)	9	0	NE (NE, NE)	NC	NC	NC
gBRCAm	69	7 (10.1)	NE (NE,	NE)	36	2 (5.6)	NE (NE, NE)	NC	NC	NC
Non-BRCAm	43	3 (7.0)	NE (NE,	NE)	23	0	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabp 11AUG2022:11:30 kpzx329

Table 3.4.43 PAOLA1: Summary of subgroup analysis of AESI G>=3: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	:	Olaparib + b (N=2		ımab			Placebo + b		mab					
Subgroup		Number (%) of patients with events		an ti % CI; hs) [)		Number (%) of patients with events	Medi (95 (mont	% CI)	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)												
NED [PDS]	92	15 (16.3)	NE (NE,	NE)	48	11 (22.9)	NE (NE,	NE)	0.63	0.29,	1.41	0.2563
NED/CR [IDS]	74	14 (18.9)	NE (NE,	NE)	38	13 (34.2)	NE (NE,	NE)	0.47	0.22,	1.01	0.0543
NED/CR [Chemo]	40	9 (22.5)	NE (NE,	NE)	20	7 (35.0)	NE (NE,	NE)	0.57	0.21,	1.61	0.2791
PR	49	12 (24.5)	NE (NE,	NE)	25	11 (44.0)	NE (NE,	NE)	0.47	0.21,	1.08	0.0742
Interaction p-value														0.9388
Screening laboratory tBR	CA statu	s (IVRS)												
tBRCAm	150	29 (19.3)	NE (NE,	NE)	65	23 (35.4)	NE (NE,	NE)	0.44	0.26,	0.77	0.0046*
non-tBRCAm	105	21 (20.0)	NE (NE,	NE)	66	19 (28.8)	NE (NE,	NE)	0.65	0.35,	1.21	0.1735
Interaction p-value														0.3677
First line treatment out	come (eC	RF)												
NED [PDS]	89	14 (15.7)	NE (NE,	NE)	47	10 (21.3)	NE (NE,	NE)	0.67	0.30,	1.55	0.3344
NED/CR [IDS]	74	13 (17.6)	NE (NE,	NE)	32	9 (28.1)	NE (NE,	NE)	0.57	0.24,	1.37	0.1997
NED/CR [Chemo]	39	9 (23.1)	NE (NE,	NE)	17	7 (41.2)	NE (NE,	NE)	0.51	0.19,	1.43	0.1910
PR	50	14 (28.0)	NE (NE,	NE)	34	15 (44.1)	NE (NE,	NE)	0.50	0.24,	1.05	0.0667
Interaction p-value														0.9609
Screening laboratory tBR	CA statu	s (eCRF)												
tBRCAm	147	27 (18.4)	NE (NE,	NE)	67	23 (34.3)	NE (NE,	NE)	0.43	0.25,	0.76	0.0041*
non-tBRCAm	108	23 (21.3)	NE (NE,	NE)	64	19 (29.7)	NE (NE,	NE)	0.67	0.37,	1.25	0.2060
Interaction p-value														0.2958
Age group														
<65 years	185	28 (15.1)	NE (NE,	NE)	98	29 (29.6)	NE (NE,	NE)	0.44	0.26,	0.74	0.0022*
>=65 years	70	22 (31.4)	NE (NE,	NE)	33	13 (39.4)	NE (NE,	NE)	0.71	0.36,	1.46	0.3434
Interaction p-value														0.2669

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabg 11AUG2022:11:30 kpzx329

Table 3.4.43 PAOLA1: Summary of subgroup analysis of AESI G>=3: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ımab			Placebo + be (N=1)		ab				
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI)		Number (%) of patients with events	(95%	n time (CI) (s) [a]	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	38 (20.9)	NE (NE,	NE)	89	25 (28.1)	NE (NE, NE	0.68	0.41,	1.14	0.1368
IV	73	12 (16.4)	NE (NE,	NE)	42	17 (40.5)	NE (NE, NE	0.32	0.15,	0.66	0.0023*
Interaction p-value													0.0958
Region													
Europe	245	50 (20.4)	NE (NE,	NE)	125	41 (32.8)	NE (NE, NE	0.54	0.36,	0.82	0.0041*
Japan	10	0	NE (NE,	NE)	6	1 (16.7)	NE (NE, NE	NC	NC		NC
Interaction p-value													NC
ECOG performance status at	Basel	ine											
(0) Normal activity	190	30 (15.8)	NE (NE,	NE)	100	32 (32.0)	NE (NE, NE	0.42	0.26,	0.70	0.0008*
(1) Restricted activity	61	18 (29.5)	NE (NE,	NE)	30	10 (33.3)	NE (NE, NE	0.81	0.38,	1.82	0.5892
Interaction p-value													0.1658
Baseline CA-125 value													
<=ULN	228	44 (19.3)	NE (NE,	NE)	117	35 (29.9)	NE (NE, NE	0.57	0.37,	0.90	0.0162*
>ULN	27	6 (22.2)	NE (NE,	NE)	14	7 (50.0)	NE (NE, NE	0.31	0.10,	0.92	0.0363*
Interaction p-value													0.2959
Histological grade													
High grade	255	50 (19.6)	NE (NE,	NE)	131	42 (32.1)	NE (NE, NE	0.53	0.35,	0.81	0.0031*
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	27 (16.3)	NE (NE,	NE)	80	20 (25.0)	NE (NE, NE	0.58	0.33,	1.05	0.0739
Residue	79	16 (20.3)	NE (NE,	NE)	43	16 (37.2)	NE (NE, NE	0.47	0.24,	0.96	0.0371*
Interaction p-value													0.6502

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabg 11AUG2022:11:30 kpzx329

Table 3.4.43 PAOLA1: Summary of subgroup analysis of AESI G>=3: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2!			Placebo + be (N=1)				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive	surgery								
Upfront	146	25 (17.1)	NE (NE, NE)	78	21 (26.9)	NE (NE, NE)	0.57	0.32, 1.03	0.0634
Interval	99	18 (18.2)	NE (NE, NE)	45	15 (33.3)	NE (NE, NE)	0.47	0.24, 0.95	0.0357*
Interaction p-value									0.6728
Myriad tumour BRCA mutat	ion stat	us							
tBRCAm	158	31 (19.6)	NE (NE, NE)	77	25 (32.5)	NE (NE, NE)	0.50	0.30, 0.85	0.0115*
Non-tBRCAm	97	19 (19.6)	NE (NE, NE)	54	17 (31.5)	NE (NE, NE)	0.58	0.30, 1.13	0.1095
Interaction p-value									0.7189
Status somatic BRCA muta	tions								
sBRCAm	25	5 (20.0)	NE (NE, NE)	9	3 (33.3)	NE (NE, NE)	0.47	0.12, 2.31	0.3261
gBRCAm	69	13 (18.8)	NE (NE, NE)	36	12 (33.3)	NE (NE, NE)	0.48	0.22, 1.06	0.0677
Non-BRCAm	43	12 (27.9)	NE (NE, NE)	23	8 (34.8)	NE (NE, NE)	0.77	0.32, 1.96	0.5667
Interaction p-value									0.7047

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabg 11AUG2022:11:30 kpzx329

Table 3.4.44 PAOLA1: Summary of subgroup analysis of AESI G>=3: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		ımab			Placebo + be (N=1)		ımab				
Subgroup	C	Number (%) of patients with events		an ti % CI; .hs) [)	C	Number (%) of patients with events	(9!	an ti 5% CI ths))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)											
NED [PDS]	92	1 (1.1)	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(IVRS)											
tBRCAm	150	3 (2.0)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	1 (1.1)	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(eCRF)											
tBRCAm	147	3 (2.0)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	2 (1.1)	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	1 (1.4)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabr 11AUG2022:11:30 kpzx329

Table 3.4.44 PAOLA1: Summary of subgroup analysis of AESI G>=3: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		ımab	•		Placebo + be (N=1)		mab	•			
Subgroup	c	Number (%) of patients with events		an ti % CI) hs) [)	C	Number (%) of patients with events		an ti: % CI) hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	2 (1.1)	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	3 (1.2)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	ne											
(0) Normal activity	190	2 (1.1)	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	1 (1.6)	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	3 (1.3)	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	3 (1.2)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	3 (1.8)	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabr 11AUG2022:11:30 kpzx329

Table 3.4.44 PAOLA1: Summary of subgroup analysis of AESI G>=3: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2			Placebo + b (N=1					
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI) (months)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	1 (0.7)	NE (NE, NE) 78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	2 (2.0)	NE (NE, NE) 45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Myriad tumour BRCA mutati	ion statu	ıs								
tBRCAm	158	3 (1.9)	NE (NE, NE) 77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE, NE) 54	. 0	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Status somatic BRCA mutat	cions									
sBRCAm	25	0	NE (NE, NE) 9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	2 (2.9)	NE (NE, NE) 36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE, NE) 23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabr 11AUG2022:11:30 kpzx329

Table 3.4.45 PAOLA1: Summary of subgroup analysis of AESI G>=3: GI perforations, abscesses and fistulae Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2)		ab		Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	(95	an ti % CI hs))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)										
NED [PDS]	92	0	NE (N	IE, NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (N	IE, NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	2 (5.0)	NE (N	IE, NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	1 (2.0)	NE (N	IE, NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR0	CA status	(IVRS)										
tBRCAm	150	2 (1.3)	NE (N	IE, NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	1 (1.0)	NE (N	IE, NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
First line treatment out	come (eCR	F)										
NED [PDS]	89	0	NE (N	IE, NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (N	IE, NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	2 (5.1)	NE (N	IE, NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	1 (2.0)	NE (N	IE, NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR0	CA status	(eCRF)										
tBRCAm	147	2 (1.4)	NE (N	IE, NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	1 (0.9)	NE (N	IE, NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Age group												
<65 years	185	2 (1.1)	NE (N	IE, NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	1 (1.4)	NE (N	IE, NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabs 11AUG2022:11:30 kpzx329

Table 3.4.45 PAOLA1: Summary of subgroup analysis of AESI G>=3: GI perforations, abscesses and fistulae Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		ımab			Placebo + be (N=13		nab				
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI)		C	Number (%) of patients with events	Media (95 (mont)	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	1 (0.5)	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	2 (2.7)	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	3 (1.2)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	ne											
(0) Normal activity	190	2 (1.1)	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	1 (1.6)	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	2 (0.9)	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	1 (3.7)	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	3 (1.2)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	0	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	3 (3.8)	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabs 11AUG2022:11:30 kpzx329

Table 3.4.45 PAOLA1: Summary of subgroup analysis of AESI G>=3: GI perforations, abscesses and fistulae Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		mab		Placebo + be (N=1		mab				
Subgroup	C	Number (%) of patients with events	(959	n time % CI) ns) [a]		Number (%) of patients with events	(95	an tim % CI) hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	2 (1.4)	NE (NE, NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	1 (1.0)	NE (NE, NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Myriad tumour BRCA mutat:	ion statu	ıs										
tBRCAm	158	2 (1.3)	NE (NE, NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	1 (1.0)	NE (NE, NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Status somatic BRCA muta	tions											
sBRCAm	25	0	NE (NE, NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE, NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE, NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabs 11AUG2022:11:30 kpzx329

Table 3.4.46 PAOLA1: Summary of subgroup analysis of AESI G>=3: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + be (N=1)		ımab				
Subgroup	C	Number (%) of patients with events	(95	an ti % CI; :hs) [)	C	Number (%) of patients with events	(95	an ti 5% CI ths))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(IVRS)											
tBRCAm	150	0	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(eCRF)											
tBRCAm	147	0	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	0	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabt 11AUG2022:11:30 kpzx329

Table 3.4.46 PAOLA1: Summary of subgroup analysis of AESI G>=3: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		umab			Placebo + be (N=1)		mab	•			
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI ths))	(Number (%) of patients with events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	0	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	0	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	ne											
(0) Normal activity	190	0	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	0	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	0	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	0	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabt 11AUG2022:11:30 kpzx329

Table 3.4.46 PAOLA1: Summary of subgroup analysis of AESI G>=3: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + b (N=1		mab				
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI) :hs) [)	(Number (%) of patients with events	(95	an ti % CI; hs) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery												
Upfront	146	0	NE (NE,	NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Myriad tumour BRCA mutat:	ion statu	ıs											
tBRCAm	158	0	NE (NE,	NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Status somatic BRCA muta	tions												
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE,	NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabt 11AUG2022:11:30 kpzx329

Table 3.4.47 PAOLA1: Summary of subgroup analysis of AESI G>=3: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + k (N=2		ab		Placebo + b (N=1)			
Subgroup	C	Number (%) of patients with events	Mediar (95% (month	CI)		Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVF	RS)									
NED [PDS]	92	2 (2.2)	NE (NE, NE)	48	1 (2.1)	NE (N	E, NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE, NE)	38	0	NE (N	E, NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE, NE)	20	0	NE (N	E, NE)	NC	NC	NC
PR	49	0	NE (NE, NE)	25	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	(IVRS)									
tBRCAm	150	1 (0.7)	NE (NE, NE)	65	0	NE (N	E, NE)	NC	NC	NC
non-tBRCAm	105	1 (1.0)	NE (NE, NE)	66	1 (1.5)	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
First line treatment out	come (eCF	RF)									
NED [PDS]	89	2 (2.2)	NE (NE, NE)	47	1 (2.1)	NE (N	E, NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE, NE)	32	0	NE (N	E, NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE, NE)	17	0	NE (N	E, NE)	NC	NC	NC
PR	50	0	NE (NE, NE)	34	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	(eCRF)									
tBRCAm	147	1 (0.7)	NE (NE, NE)	67	0	NE (N	E, NE)	NC	NC	NC
non-tBRCAm	108	1 (0.9)	NE (NE, NE)	64	1 (1.6)	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Age group											
<65 years	185	1 (0.5)	NE (NE, NE)	98	1 (1.0)	NE (N	E, NE)	NC	NC	NC
>=65 years	70	1 (1.4)	NE (NE, NE)	33	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabu 11AUG2022:11:30 kpzx329

Table 3.4.47 PAOLA1: Summary of subgroup analysis of AESI G>=3: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		ımab	•		Placebo + be (N=13		ıb			
Subgroup	c	Number (%) of patients with events	Media (95 (mont	% CI))	c	Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	2 (1.1)	NE (NE,	NE)	89	1 (1.1)	NE (NE, NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	2 (0.8)	NE (NE,	NE)	125	1 (0.8)	NE (NE, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	0	NE (NE,	NE)	100	1 (1.0)	NE (I	NE, NE)	NC	NC	NC
(1) Restricted activity	61	1 (1.6)	NE (NE,	NE)	30	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	2 (0.9)	NE (NE,	NE)	117	1 (0.9)	NE (I	NE, NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (I	NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	2 (0.8)	NE (NE,	NE)	131	1 (0.8)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	2 (1.2)	NE (NE,	NE)	80	1 (1.3)	NE (I	NE, NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (I	NE, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabu 11AUG2022:11:30 kpzx329

Table 3.4.47 PAOLA1: Summary of subgroup analysis of AESI G>=3: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + 1 (N=2	bevacizumab 255)			Placebo + be (N=1)		b			
Subgroup	C	Number (%) of patients with events	Median ti (95% CI) (months) [)	C	Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	2 (1.4)	NE (NE,	NE)	78	1 (1.3)	NE (N	IE, NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (N	IE, NE)	NC	NC	NC
Interaction p-value											NC
Myriad tumour BRCA mutati	ion statu	ıs									
tBRCAm	158	2 (1.3)	75.0 (75.0,	NE)	77	0	NE (N	IE, NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	1 (1.9)	NE (N	IE, NE)	NC	NC	NC
Interaction p-value											NC
Status somatic BRCA mutat	cions										
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (N	IE, NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE,	NE)	36	0	NE (N	IE, NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (N	IE, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabu 11AUG2022:11:30 kpzx329

Table 3.4.48 PAOLA1: Summary of subgroup analysis of AESI G>=3: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ımab			Placebo + be (N=13		ıb			
Subgroup	c	Number (%) of patients with events	Media (95 (mont	% CI)	C	Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	RS)										
NED [PDS]	92	1 (1.1)	NE (NE,	NE)	48	1 (2.1)	NE (1	NE, NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	1 (2.6)	NE (1	NE, NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	1 (5.0)	NE (1	NE, NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	1 (4.0)	NE (1	NE, NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR	CA status	(IVRS)										
tBRCAm	150	1 (0.7)	NE (NE,	NE)	65	2 (3.1)	NE (1	NE, NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	2 (3.0)	NE (1	NE, NE)	NC	NC	NC
Interaction p-value												NC
First line treatment out	come (eCR	RF)										
NED [PDS]	89	1 (1.1)	NE (NE,	NE)	47	1 (2.1)	NE (1	NE, NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (1	NE, NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	1 (5.9)	NE (1	NE, NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	2 (5.9)	NE (1	NE, NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR	CA status	s (eCRF)										
tBRCAm	147	1 (0.7)	NE (NE,	NE)	67	2 (3.0)	NE (1	NE, NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	2 (3.1)	NE (1	NE, NE)	NC	NC	NC
Interaction p-value												NC
Age group												
<65 years	185	1 (0.5)	NE (NE,	NE)	98	1 (1.0)	NE (1	NE, NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	3 (9.1)	NE (1	NE, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabv 11AUG2022:11:30 kpzx329

Table 3.4.48 PAOLA1: Summary of subgroup analysis of AESI G>=3: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	I	Olaparib + b (N=2!		ımab	,		Placebo + be (N=13		b			
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI)		С	Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	1 (0.5)	NE (NE,	NE)	89	2 (2.2)	NE (1	E, NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	2 (4.8)	NE (1	E, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	1 (0.4)	NE (NE,	NE)	125	4 (3.2)	NE (1	E, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (1	E, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	ne										
(0) Normal activity	190	1 (0.5)	NE (NE,	NE)	100	2 (2.0)	NE (1	E, NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	2 (6.7)	NE (1	E, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	1 (0.4)	NE (NE,	NE)	117	4 (3.4)	NE (1	E, NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (1	E, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	1 (0.4)	NE (NE,	NE)	131	4 (3.1)	NE (1	E, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	1 (0.6)	NE (NE,	NE)	80	1 (1.3)	NE (1	E, NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	3 (7.0)	NE (1	E, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabv 11AUG2022:11:30 kpzx329

Table 3.4.48 PAOLA1: Summary of subgroup analysis of AESI G>=3: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		mab		Placebo + be (N=1		nab			
Subgroup	C	Number (%) of patients with events	(959	n time % CI) ns) [a]	(Number (%) of patients with events	(95	n time % CI) ns) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery										
Upfront	146	1 (0.7)	NE (NE, NE)	78	3 (3.8)	NE (NE, NE)	NC	NC	NC
Interval	99	0	NE (NE, NE)	45	1 (2.2)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Myriad tumour BRCA mutat:	ion statu	ıs									
tBRCAm	158	1 (0.6)	NE (NE, NE)	77	3 (3.9)	NE (NE, NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE, NE)	54	1 (1.9)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC
Status somatic BRCA muta	tions										
sBRCAm	25	0	NE (NE, NE)	9	0	NE (NE, NE)	NC	NC	NC
gBRCAm	69	0	NE (NE, NE)	36	0	NE (NE, NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE, NE)	23	1 (4.3)	NE (NE, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabv 11AUG2022:11:30 kpzx329

Table 3.4.49 PAOLA1: Summary of subgroup analysis of AESI G>=3: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + (N=	bevacizum 255)	mab			Placebo + be (N=13		nab	•			
Subgroup	C	Number (%) of patients vith events	Media (95% (month	k CI)		О	Number (%) of patients vith events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	LS)											
NED [PDS]	92	1 (1.1)	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	2 (4.1)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA status	(IVRS)											
tBRCAm	150	2 (1.3)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	1 (1.0)	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment outo	come (eCR	PF)											
NED [PDS]	89	1 (1.1)	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBRO	CA status	(eCRF)											
tBRCAm	147	2 (1.4)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	1 (0.9)	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	3 (1.6)	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabw 11AUG2022:11:30 kpzx329

Table 3.4.49 PAOLA1: Summary of subgroup analysis of AESI G>=3: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		ımab	•		Placebo + be (N=13		mab	•			
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI)		C	Number (%) of patients with events	(95	an ti % CI) hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	1 (0.5)	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	2 (2.7)	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	3 (1.2)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	.ne											
(0) Normal activity	190	1 (0.5)	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	2 (3.3)	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	3 (1.3)	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	3 (1.2)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	3 (1.8)	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabw 11AUG2022:11:30 kpzx329

Table 3.4.49 PAOLA1: Summary of subgroup analysis of AESI G>=3: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		·	Placebo + b (N=1					
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI) (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	1 (0.7)	NE (NE, NE	:) 78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	2 (2.0)	NE (NE, NE	1) 45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Myriad tumour BRCA mutat:	ion statu	ıs								
tBRCAm	158	3 (1.9)	NE (NE, NE	:) 77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE, NE	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Status somatic BRCA mutat	cions									
sBRCAm	25	1 (4.0)	NE (NE, NE	:) 9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	2 (2.9)	NE (NE, NE	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE, NE	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabw 11AUG2022:11:30 kpzx329

Table 3.4.50 PAOLA1: Summary of subgroup analysis of AESI G>=3: Congestive heart failure Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + be (N=1)		ımab				
Subgroup	C	Number (%) of patients with events	(95	an ti % CI; :hs) [)	C	Number (%) of patients with events	(95	an ti 5% CI ths))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(IVRS)											
tBRCAm	150	0	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(eCRF)											
tBRCAm	147	0	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	0	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabx 11AUG2022:11:30 kpzx329

Table 3.4.50 PAOLA1: Summary of subgroup analysis of AESI G>=3: Congestive heart failure Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		umab			Placebo + be (N=1)		mab	•			
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI ths))	(Number (%) of patients with events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	0	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	0	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	ne											
(0) Normal activity	190	0	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	0	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	0	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	0	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabx 11AUG2022:11:30 kpzx329

Table 3.4.50 PAOLA1: Summary of subgroup analysis of AESI G>=3: Congestive heart failure Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		umab			Placebo + be (N=1		mab				
Subgroup	C	Number (%) of patients with events		an ti 5% CI; ths) [)	C	Number (%) of patients with events	(95	an ti % CI) hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery												
Upfront	146	0	NE (NE,	NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Myriad tumour BRCA mutat:	ion statu	ıs											
tBRCAm	158	0	NE (NE,	NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Status somatic BRCA mutat	cions												
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE,	NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabx 11AUG2022:11:30 kpzx329

Table 3.4.51 PAOLA1: Summary of subgroup analysis of AESI G>=3: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		ımab	:-		Placebo + be (N=13		mab				
Subgroup	C	Number (%) of patients with events	(95	an ti % CI hs))	C	Number (%) of patients with events	(95	an ti % CI; hs) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	1 (5.0)	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(IVRS)											
tBRCAm	150	0	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	1 (5.9)	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(eCRF)											
tBRCAm	147	0	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	0	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaby 11AUG2022:11:30 kpzx329

Table 3.4.51 PAOLA1: Summary of subgroup analysis of AESI G>=3: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + be (N=1)		nab			
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI ths))	C	Number (%) of patients with events	(95	n time % CI) ns) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	0	NE (NE,	NE)	89	1 (1.1)	NE (NE, NE) NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	0	NE (NE,	NE)	125	1 (0.8)	NE (NE, NE) NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	0	NE (NE,	NE)	100	1 (1.0)	NE (NE, NE) NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	0	NE (NE,	NE)	117	1 (0.9)	NE (NE, NE) NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	0	NE (NE,	NE)	131	1 (0.8)	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	0	NE (NE,	NE)	80	0	NE (NE, NE) NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	1 (2.3)	NE (NE, NE) NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaby 11AUG2022:11:30 kpzx329

Table 3.4.51 PAOLA1: Summary of subgroup analysis of AESI G>=3: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		ımab			Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events		an ti % CI) hs) [C	Number (%) of patients with events	(95	an ti % CI) hs) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery												
Upfront	146	0	NE (NE,	NE)	78	1 (1.3)	NE (NE,	NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Myriad tumour BRCA mutat:	ion statu	ıs											
tBRCAm	158	0	NE (NE,	NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	1 (1.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Status somatic BRCA mutat	cions												
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE,	NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaby 11AUG2022:11:30 kpzx329

Table 3.4.52 PAOLA1: Summary of subgroup analysis of AESI G>=3: MDS/AML Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + k (N=2		b		Placebo + be (N=1)					
Subgroup	(Number (%) of patients with events	Median (95% ((months	CI)	(Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVF	RS)									
NED [PDS]	92	2 (2.2)	NE (N	E, NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (N	E, NE)	38	2 (5.3)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	1 (2.5)	NE (N	E, NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (N	E, NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	(IVRS)									
tBRCAm	150	2 (1.3)	NE (N	E, NE)	65	1 (1.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	2 (1.9)	NE (N	E, NE)	66	2 (3.0)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
First line treatment out	come (eCF	RF)									
NED [PDS]	89	2 (2.2)	NE (N	E, NE)	47	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (N	E, NE)	32	2 (6.3)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	1 (2.6)	NE (N	E, NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (N	E, NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR	CA status	s (eCRF)									
tBRCAm	147	2 (1.4)	NE (N	E, NE)	67	1 (1.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	2 (1.9)	NE (N	E, NE)	64	2 (3.1)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Age group											
<65 years	185	3 (1.6)	NE (N	E, NE)	98	1 (1.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	1 (1.4)	NE (N	E, NE)	33	2 (6.1)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabz 11AUG2022:11:30 kpzx329

Table 3.4.52 PAOLA1: Summary of subgroup analysis of AESI G>=3: MDS/AML Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2!		mab			Placebo + be (N=13		ab			
Subgroup	C	Number (%) of patients with events	Media (95% (mont)	% CI)		C	Number (%) of patients with events	Mediar (95% (month	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	3 (1.6)	NE (NE,	NE)	89	3 (3.4)	NE (NE, NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE,	NE)	42	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	4 (1.6)	NE (NE,	NE)	125	3 (2.4)	NE (NE, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	1 (0.5)	NE (NE,	NE)	100	2 (2.0)	NE (NE, NE)	NC	NC	NC
(1) Restricted activity	61	2 (3.3)	NE (NE,	NE)	30	1 (3.3)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	4 (1.8)	NE (NE,	NE)	117	3 (2.6)	NE (NE, NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	4 (1.6)	NE (NE,	NE)	131	3 (2.3)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	3 (1.8)	NE (NE,	NE)	80	3 (3.8)	NE (NE, NE)	NC	NC	NC
Residue	79	1 (1.3)	NE (NE,	NE)	43	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabz 11AUG2022:11:30 kpzx329

Table 3.4.52 PAOLA1: Summary of subgroup analysis of AESI G>=3: MDS/AML Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2			Placebo + be (N=1)					
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	(Number (%) of patients with events	Median ti (95% CI) (months) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery									
Upfront	146	3 (2.1)	NE (NE, NE)	78	1 (1.3)	NE (NE,	NE)	NC	NC	NC
Interval	99	1 (1.0)	NE (NE, NE)	45	2 (4.4)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Myriad tumour BRCA mutat:	ion statu	ıs								
tBRCAm	158	2 (1.3)	NE (NE, NE)	77	2 (2.6)	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	2 (2.1)	NE (NE, NE)	54	1 (1.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Status somatic BRCA mutat	tions									
sBRCAm	25	1 (4.0)	NE (NE, NE)	9	1 (11.1)	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE, NE)	36	1 (2.8)	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE, NE)	23	1 (4.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeabz 11AUG2022:11:30 kpzx329

Table 3.4.53 PAOLA1: Summary of subgroup analysis of AESI G>=3: Myelodysplastic syndrome and Acute myeloid leukaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ab		Placebo + be (N=1)					
Subgroup	C	Number (%) of patients with events	Mediar (95% (month	CI)	(Number (%) of patients with events	Median t (95% C (months)	Γ)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment outo	come (IVR	S)									
NED [PDS]	92	2 (2.2)	NE (NE, NE)	48	2 (4.2)	NE (NE	, NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (NE, NE)	38	2 (5.3)	NE (NE	, NE)	NC	NC	NC
NED/CR [Chemo]	40	1 (2.5)	NE (NE, NE)	20	0	NE (NE	, NE)	NC	NC	NC
PR	49	0	NE (NE, NE)	25	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBRO	CA status	(IVRS)									
tBRCAm	150	2 (1.3)	NE (NE, NE)	65	1 (1.5)	NE (NE	, NE)	NC	NC	NC
non-tBRCAm	105	2 (1.9)	NE (NE, NE)	66	3 (4.5)	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
First line treatment outo	come (eCR	F)									
NED [PDS]	89	2 (2.2)	NE (NE, NE)	47	2 (4.3)	NE (NE	, NE)	NC	NC	NC
NED/CR [IDS]	74	1 (1.4)	NE (NE, NE)	32	2 (6.3)	NE (NE	, NE)	NC	NC	NC
NED/CR [Chemo]	39	1 (2.6)	NE (NE, NE)	17	0	NE (NE	, NE)	NC	NC	NC
PR	50	0	NE (NE, NE)	34	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBRO	CA status	(eCRF)									
tBRCAm	147	2 (1.4)	NE (NE, NE)	67	1 (1.5)	NE (NE	, NE)	NC	NC	NC
non-tBRCAm	108	2 (1.9)	NE (NE, NE)	64	3 (4.7)	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC
Age group											
<65 years	185	3 (1.6)	NE (NE, NE)	98	2 (2.0)	NE (NE	, NE)	NC	NC	NC
>=65 years	70	1 (1.4)	NE (NE, NE)	33	2 (6.1)	NE (NE	, NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaca 11AUG2022:11:30 kpzx329

Table 3.4.53 PAOLA1: Summary of subgroup analysis of AESI G>=3: Myelodysplastic syndrome and Acute myeloid leukaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2!		mab	•		Placebo + be (N=13		ab	:		
Subgroup	C	Number (%) of patients with events	Media (95: (montl	% CI)		С	Number (%) of patients with events	Mediar (95% (month	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	3 (1.6)	NE (NE,	NE)	89	4 (4.5)	NE (NE, NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE,	NE)	42	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	4 (1.6)	NE (NE,	NE)	125	4 (3.2)	NE (NE, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	1 (0.5)	NE (NE,	NE)	100	3 (3.0)	NE (NE, NE)	NC	NC	NC
(1) Restricted activity	61	2 (3.3)	NE (NE,	NE)	30	1 (3.3)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	4 (1.8)	NE (NE,	NE)	117	4 (3.4)	NE (NE, NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	4 (1.6)	NE (NE,	NE)	131	4 (3.1)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	3 (1.8)	NE (NE,	NE)	80	4 (5.0)	NE (NE, NE)	NC	NC	NC
Residue	79	1 (1.3)	NE (NE,	NE)	43	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaca 11AUG2022:11:30 kpzx329

Table 3.4.53 PAOLA1: Summary of subgroup analysis of AESI G>=3: Myelodysplastic syndrome and Acute myeloid leukaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!				Placebo + be (N=1)		nab				
Subgroup	C	Number (%) of patients with events	Median t (95% C (months)	I)	C	Number (%) of patients with events	Media (95 (mont)	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	3 (2.1)	NE (NE	, NE)	78	2 (2.6)	NE (NE,	NE)	NC	NC	NC
Interval	99	1 (1.0)	NE (NE	, NE)	45	2 (4.4)	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Myriad tumour BRCA mutat:	ion statu	ıs										
tBRCAm	158	2 (1.3)	NE (NE	, NE)	77	3 (3.9)	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	2 (2.1)	NE (NE	, NE)	54	1 (1.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Status somatic BRCA muta	cions											
sBRCAm	25	1 (4.0)	NE (NE	, NE)	9	1 (11.1)	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE	, NE)	36	2 (5.6)	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE	, NE)	23	1 (4.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeaca 11AUG2022:11:30 kpzx329

Table 3.4.54 PAOLA1: Summary of subgroup analysis of AESI G>=3: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2				Placebo + be (N=1))			
Subgroup	C	Number (%) of patients with events	Median t (95% C (months)	I)	C	Number (%) of patients with events	Median (95% (months	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	RS)									
NED [PDS]	92	3 (3.3)	NE (NE	, NE)	48	1 (2.1)	NE (N	E, NE)	NC	NC	NC
NED/CR [IDS]	74	6 (8.1)	NE (NE	, NE)	38	1 (2.6)	NE (N	E, NE)	NC	NC	NC
NED/CR [Chemo]	40	2 (5.0)	NE (NE	, NE)	20	1 (5.0)	NE (N	E, NE)	NC	NC	NC
PR	49	0	NE (NE	, NE)	25	0	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	(IVRS)									
tBRCAm	150	9 (6.0)	NE (NE	, NE)	65	2 (3.1)	NE (N	E, NE)	1.76	0.45, 11.55	0.4436
non-tBRCAm	105	2 (1.9)	NE (NE	, NE)	66	1 (1.5)	NE (N	E, NE)	1.14	0.11, 24.55	0.9137
Interaction p-value											0.7676
First line treatment out	come (eCR	RF)									
NED [PDS]	89	4 (4.5)	NE (NE	, NE)	47	1 (2.1)	NE (N	E, NE)	NC	NC	NC
NED/CR [IDS]	74	6 (8.1)	NE (NE	, NE)	32	0	NE (N	E, NE)	NC	NC	NC
NED/CR [Chemo]	39	1 (2.6)	NE (NE	, NE)	17	1 (5.9)	NE (N	E, NE)	NC	NC	NC
PR	50	0	NE (NE	, NE)	34	1 (2.9)	NE (N	E, NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	(eCRF)									
tBRCAm	147	9 (6.1)	NE (NE	, NE)	67	2 (3.0)	NE (N	E, NE)	1.83	0.47, 12.00	0.4121
non-tBRCAm	108	2 (1.9)	NE (NE	, NE)	64	1 (1.6)	NE (N	E, NE)	1.08	0.10, 23.32	0.9474
Interaction p-value											0.7222
Age group											
<65 years	185	9 (4.9)	NE (NE	, NE)	98	2 (2.0)	NE (N	E, NE)	2.21	0.57, 14.49	0.2736
>=65 years	70	2 (2.9)	NE (NE	, NE)	33	1 (3.0)	NE (N	E, NE)	0.75	0.07, 16.21	0.8199
Interaction p-value											0.4733

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacb 11AUG2022:11:30 kpzx329

Table 3.4.54 PAOLA1: Summary of subgroup analysis of AESI G>=3: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		mab	,		Placebo + be (N=13		nab					
Subgroup		Number (%) of patients with events	Media (959 (mont)	% CI)		C	Number (%) of patients with events	Media (95: (montl	% CI)		Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
FIGO Stage (Disease state)														
III	182	7 (3.8)	NE (NE,	NE)	89	2 (2.2)	NE (NE,	NE)	NC	NC		NC
IV	73	4 (5.5)	NE (NE,	NE)	42	1 (2.4)	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
Region														
Europe	245	10 (4.1)	NE (NE,	NE)	125	3 (2.4)	NE (NE,	NE)	1.52	0.47,	6.81	0.5068
Japan	10	1 (10.0)	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
ECOG performance status at	Basel:	ine												
(0) Normal activity	190	6 (3.2)	NE (NE,	NE)	100	2 (2.0)	NE (NE,	NE)	NC	NC		NC
(1) Restricted activity	61	5 (8.2)	NE (NE,	NE)	30	1 (3.3)	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
Baseline CA-125 value														
<=ULN	228	11 (4.8)	,	NE,	NE)	117	3 (2.6)	NE (NE,	NE)	1.64	0.51,	7.24	0.4309
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value														NC
Histological grade														
High grade	255	11 (4.3)	NE (NE,	NE)	131	3 (2.3)	NE (NE,	NE)	1.68	0.52,	7.44	0.4035
Interaction p-value														NC
Cytoreductive surgery outco	ome													
No residue	166	10 (6.0)	NE (NE,	NE)	80	1 (1.3)	NE (NE,	NE)	4.30	0.82,	79.05	0.0910
Residue	79	1 (1.3)	NE (NE,	NE)	43	2 (4.7)	NE (NE,	NE)	0.24	0.01,	2.48	0.2223
Interaction p-value														0.0524

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacb 11AUG2022:11:30 kpzx329

Table 3.4.54 PAOLA1: Summary of subgroup analysis of AESI G>=3: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!			Placebo + be (N=1					
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive	surgery									
Upfront	146	5 (3.4)	NE (NE, NE) 78	2 (2.6)	NE (NE,	NE)	NC	NC	NC
Interval	99	6 (6.1)	NE (NE, NE) 45	1 (2.2)	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Myriad tumour BRCA mutat	ion statu	ıs								
tBRCAm	158	9 (5.7)	NE (NE, NE) 77	3 (3.9)	NE (NE,	NE)	1.27	0.38, 5.7	3 0.7149
Non-tBRCAm	97	2 (2.1)	NE (NE, NE) 54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC
Status somatic BRCA muta	tions									
sBRCAm	25	0	NE (NE, NE) 9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	5 (7.2)	NE (NE, NE) 36	2 (5.6)	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	1 (2.3)	NE (NE, NE) 23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value										NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacb 11AUG2022:11:30 kpzx329

Table 3.4.55 PAOLA1: Summary of subgroup analysis of AESI G1-2: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ımab			Placebo + be (N=13		ımab				
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI))	C	Number (%) of patients with events	(95	an ti 5% CI :hs))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IV)	RS)											
NED [PDS]	92	28 (30.4)	NE (NE,	NE)	48	7 (14.6)	NE (NE,	NE)	2.31	1.07, 5.74	0.0328*
NED/CR [IDS]	74	31 (41.9)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	11 (27.5)	NE (NE,	NE)	20	1 (5.0)	NE (NE,	NE)	5.95	1.16,108.78	0.0298*
PR	49	16 (32.7)	NE (NE,	NE)	25	3 (12.0)	NE (NE,	NE)	3.29	1.10, 14.15	0.0322*
Interaction p-value													0.6375
Screening laboratory tBR0	CA statu	s (IVRS)											
tBRCAm	150	46 (30.7)	NE (NE,	NE)	65	7 (10.8)	NE (NE,	NE)	3.18	1.53, 7.72	0.0011*
non-tBRCAm	105	40 (38.1)	NE (NE,	NE)	66	4 (6.1)	NE (NE,	NE)	7.81	3.15, 26.03	<0.0001*
Interaction p-value													0.1665
First line treatment out	come (eCl	RF)											
NED [PDS]	89	26 (29.2)	NE (NE,	NE)	47	7 (14.9)	NE (NE,	NE)	2.16	0.99, 5.39	0.0536
NED/CR [IDS]	74	28 (37.8)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	10 (25.6)	NE (NE,	NE)	17	1 (5.9)	NE (NE,	NE)	4.77	0.91, 87.51	0.0669
PR	50	20 (40.0)	NE (NE,	NE)	34	3 (8.8)	NE (NE,	NE)	5.59	1.92, 23.72	0.0008*
Interaction p-value													0.3918
Screening laboratory tBR0	CA statu	s (eCRF)											
tBRCAm	147	46 (31.3)	NE (NE,	NE)	67	7 (10.4)	NE (NE,	NE)	3.36	1.62, 8.16	0.0006*
non-tBRCAm	108	40 (37.0)	NE (NE,	NE)	64	4 (6.3)	NE (NE,	NE)	7.32	2.95, 24.38	<0.0001*
Interaction p-value													0.2316
Age group													
<65 years	185	57 (30.8)	NE (NE,	NE)	98	8 (8.2)	NE (NE,	NE)	4.34	2.20, 9.85	<0.0001*
>=65 years	70	29 (41.4)	NE (NE,	NE)	33	3 (9.1)	NE (NE,	NE)	5.66	2.01, 23.64	0.0003*
Interaction p-value													0.7057

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacc 11AUG2022:11:30 kpzx329

Table 3.4.55 PAOLA1: Summary of subgroup analysis of AESI G1-2: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + k				Placebo + be (N=1)		,		
Subgroup		Number (%) of patients with events	Median ti (95% CI (months))		Number (%) of patients with events	Median time (95% CI) (months) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	61 (33.5)	NE (NE,	NE)	89	8 (9.0)	NE (NE, N	E) 4.33	2.20, 9.81	<0.0001*
IV	73	25 (34.2)	NE (NE,	NE)	42	3 (7.1)	NE (NE, N	E) 5.77	2.03, 24.25	0.0004*
Interaction p-value										0.6836
Region										
Europe	245	80 (32.7)	NE (NE,	NE)	125	11 (8.8)	NE (NE, N	E) 4.30	2.39, 8.55	<0.0001*
Japan	10	6 (60.0)	2.8 (0.3,	NE)	6	0	NE (NE, N	E) NC	NC	NC
Interaction p-value										NC
ECOG performance status at	Basel:	ine								
(0) Normal activity	190	63 (33.2)	NE (NE,	NE)	100	7 (7.0)	NE (NE, N	E) 5.68	2.79, 13.63	<0.0001*
(1) Restricted activity	61	23 (37.7)	NE (NE,	NE)	30	4 (13.3)	NE (NE, N	E) 3.14	1.21, 10.71	0.0168*
Interaction p-value										0.3900
Baseline CA-125 value										
<=ULN	228	74 (32.5)	NE (NE,	NE)	117	10 (8.5)	NE (NE, N	E) 4.40	2.38, 9.07	<0.0001*
>ULN	27	12 (44.4)	NE (NE,	NE)	14	1 (7.1)	NE (NE, N	E) 8.32	1.64,151.60	0.0066*
Interaction p-value										0.5311
Histological grade										
High grade	255	86 (33.7)	NE (NE,	NE)	131	11 (8.4)	NE (NE, N	E) 4.71	2.63, 9.35	<0.0001*
Interaction p-value										NC
Cytoreductive surgery outco	ome									
No residue	166	56 (33.7)	NE (NE,	NE)	80	7 (8.8)	NE (NE, N	E) 4.55	2.22, 10.96	<0.0001*
Residue	79	26 (32.9)	NE (NE,	NE)	43	2 (4.7)	NE (NE, N	E) 8.36	2.50, 51.89	<0.0001*
Interaction p-value										0.4474

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacc 11AUG2022:11:30 kpzx329

Table 3.4.55 PAOLA1: Summary of subgroup analysis of AESI G1-2: Anaemia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		Placebo + b (N=1			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI	2-sided [b] p-value[b]
Timing of cytoreductive s	surgery						
Upfront	146	41 (28.1)	NE (NE, NE)	78 9 (11.5)	NE (NE, NE)	2.68 1.37,	5.90 0.0032*
Interval	99	41 (41.4)	NE (NE, NE)	45 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Myriad tumour BRCA mutat:	ion stat	us					
tBRCAm	158	52 (32.9)	NE (NE, NE)	77 6 (7.8)	NE (NE, NE)	4.86 2.26,	12.64 <0.0001*
Non-tBRCAm	97	34 (35.1)	NE (NE, NE)	54 5 (9.3)	NE (NE, NE)	4.59 1.96,	13.38 0.0002*
Interaction p-value							0.9296
Status somatic BRCA muta	tions						
sBRCAm	25	9 (36.0)	NE (NE, NE)	9 0	NE (NE, NE)	NC NC	NC
gBRCAm	69	25 (36.2)	NE (NE, NE)	36 4 (11.1)	NE (NE, NE)	3.93 1.52,	13.35 0.0032*
Non-BRCAm	43	17 (39.5)	NE (NE, NE)	23 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacc 11AUG2022:11:30 kpzx329

Table 3.4.56 PAOLA1: Summary of subgroup analysis of AESI G1-2: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	:	Olaparib + b (N=2		ımab			Placebo + be (N=1		mab					2-sided p-value [b]
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI))		Number (%) of patients with events	(95	an ti % CI hs))	Hazard ratio [b]	95% C	95% CI [b]	
First line treatment out	come (IV	RS)												
NED [PDS]	92	12 (13.0)	NE (NE,	NE)	48	6 (12.5)	NE (NE,	NE)	1.01	0.39,	2.91	0.9806
NED/CR [IDS]	74	16 (21.6)	NE (NE,	NE)	38	7 (18.4)	NE (NE,	NE)	1.14	0.49,	2.98	0.7645
NED/CR [Chemo]	40	6 (15.0)	NE (NE,	NE)	20	4 (20.0)	NE (NE,	NE)	0.70	0.20,	2.74	0.5863
PR	49	5 (10.2)	NE (NE,	NE)	25	2 (8.0)	NE (NE,	NE)	1.33	0.29,	9.28	0.7289
Interaction p-value														0.9193
Screening laboratory tBR	CA statu	s (IVRS)												
tBRCAm	150	21 (14.0)	NE (NE,	NE)	65	10 (15.4)	NE (NE,	NE)	0.87	0.42,	1.94	0.7272
non-tBRCAm	105	18 (17.1)	NE (NE,	NE)	66	9 (13.6)	NE (NE,	NE)	1.27	0.58,	2.96	0.5558
Interaction p-value														0.5062
First line treatment out	come (eC	RF)												
NED [PDS]	89	13 (14.6)	NE (NE,	NE)	47	5 (10.6)	NE (NE,	NE)	1.36	0.51,	4.25	0.5462
NED/CR [IDS]	74	14 (18.9)	NE (NE,	NE)	32	6 (18.8)	NE (NE,	NE)	0.97	0.39,	2.74	0.9516
NED/CR [Chemo]	39	5 (12.8)	NE (NE,	NE)	17	3 (17.6)	NE (NE,	NE)	0.67	0.16,	3.27	0.5909
PR	50	7 (14.0)	NE (NE,	NE)	34	5 (14.7)	NE (NE,	NE)	0.96	0.31,	3.24	0.9435
Interaction p-value														0.8833
Screening laboratory tBR	CA statu	s (eCRF)												
tBRCAm	147	21 (14.3)	NE (NE,	NE)	67	11 (16.4)	NE (NE,	NE)	0.83	0.41,	1.79	0.6300
non-tBRCAm	108	18 (16.7)	NE (NE,	NE)	64	8 (12.5)	NE (NE,	NE)	1.34	0.60,	3.28	0.4790
Interaction p-value														0.3971
Age group														
<65 years	185	28 (15.1)	NE (NE,	NE)	98	14 (14.3)	NE (NE,	NE)	1.05	0.56,	2.05	0.8866
>=65 years	70	11 (15.7)	NE (NE,	NE)	33	5 (15.2)	NE (NE,	NE)	0.99	0.36,	3.13	0.9805
Interaction p-value														0.9247

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacd 11AUG2022:11:30 kpzx329

Table 3.4.56 PAOLA1: Summary of subgroup analysis of AESI G1-2: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + k (N=2				Placebo + be (N=1)						
Subgroup		Number (%) of patients with events	Median to (95% CI (months))		Number (%) of patients with events)	Hazard ratio [b]	95% CI [b]		2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	27 (14.8)	NE (NE,	NE)	89	11 (12.4)	NE (NE,	NE)	1.19	0.61,	2.51	0.6181
IV	73	12 (16.4)	NE (NE,	NE)	42	8 (19.0)	NE (NE,	NE)	0.82	0.34,	2.09	0.6653
Interaction p-value												0.5185
Region												
Europe	245	33 (13.5)	NE (NE,	NE)	125	18 (14.4)	NE (NE,	NE)	0.90	0.51,	1.64	0.7303
Japan	10	6 (60.0)	5.2 (0.5,	NE)	6	1 (16.7)	NE (NE,	NE)	5.17	0.88,	97.70	0.0715
Interaction p-value												0.0703
ECOG performance status at	Basel	ine										
(0) Normal activity	190	31 (16.3)	NE (NE,	NE)	100	12 (12.0)	NE (NE,	NE)	1.35	0.71,	2.73	0.3679
(1) Restricted activity	61	7 (11.5)	NE (NE,	NE)	30	7 (23.3)	NE (NE,	NE)	0.46	0.16,	1.34	0.1483
Interaction p-value												0.0887
Baseline CA-125 value												
<=ULN	228	35 (15.4)	NE (NE,	NE)	117	16 (13.7)	NE (NE,	NE)	1.11	0.62,		0.7331
>ULN	27	4 (14.8)	NE (NE,	NE)	14	3 (21.4)	NE (NE,	NE)	0.64	0.14,	3.23	0.5590
Interaction p-value												0.5047
Histological grade												
High grade	255	39 (15.3)	NE (NE,	NE)	131	19 (14.5)	NE (NE,	NE)	1.03	0.61,	1.83	0.9077
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	27 (16.3)	NE (NE,	NE)	80	11 (13.8)	NE (NE,	NE)	1.16	0.59,	2.45	0.6702
Residue	79	10 (12.7)	NE (NE,	NE)	43	7 (16.3)	NE (NE,	NE)	0.74	0.29,	2.05	0.5512
Interaction p-value												0.4639

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacd 11AUG2022:11:30 kpzx329

Table 3.4.56 PAOLA1: Summary of subgroup analysis of AESI G1-2: Neutropenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2			Placebo + bo (N=1					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median time (95% CI) (months) [a]	- Hazard ratio [b]	95% CI [b]	2-sided p-value [b]	
Timing of cytoreductive s	surgery									
Upfront	146	20 (13.7)	NE (NE, NE)	78	9 (11.5)	NE (NE, N	E) 1.17	0.55, 2.71	0.6901	
Interval	99	17 (17.2)	NE (NE, NE)	45	9 (20.0)	NE (NE, N	E) 0.81	0.37, 1.91	0.6169	
Interaction p-value									0.5236	
Myriad tumour BRCA mutat:	ion stat	us								
tBRCAm	158	22 (13.9)	NE (NE, NE)	77	10 (13.0)	NE (NE, N	E) 1.04	0.51, 2.29	0.9193	
Non-tBRCAm	97	17 (17.5)	NE (NE, NE)	54	9 (16.7)	NE (NE, N	E) 1.05	0.48, 2.46	0.9056	
Interaction p-value									0.9855	
Status somatic BRCA mutat	cions									
sBRCAm	25	4 (16.0)	NE (NE, NE)	9	1 (11.1)	NE (NE, N	E) 1.35	0.20, 26.47	0.7800	
gBRCAm	69	11 (15.9)	NE (NE, NE)	36	7 (19.4)	NE (NE, N	E) 0.81	0.32, 2.19	0.6595	
Non-BRCAm	43	10 (23.3)	NE (NE, NE)	23	4 (17.4)	NE (NE, N	E) 1.38	0.46, 5.02	0.5818	
Interaction p-value									0.7553	

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacd 11AUG2022:11:30 kpzx329

Table 3.4.57 PAOLA1: Summary of subgroup analysis of AESI G1-2: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		:		Placebo + be (N=1)					2-sided p-value [b]
Subgroup	c	Number (%) of patients with events	Median ti (95% CI (months))	C	Number (%) of patients with events	Median to (95% CI (months))	Hazard ratio [b]	95% CI [b]	
First line treatment out	come (IVR	RS)									
NED [PDS]	92	5 (5.4)	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	5 (6.8)	NE (NE,	NE)	38	3 (7.9)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	3 (7.5)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	1 (2.0)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Screening laboratory tBR0	CA status	s (IVRS)									
tBRCAm	150	8 (5.3)	NE (NE,	NE)	65	2 (3.1)	NE (NE,	NE)	1.71	0.43, 11.34	0.4745
non-tBRCAm	105	6 (5.7)	NE (NE,	NE)	66	2 (3.0)	NE (NE,	NE)	1.88	0.43, 12.86	0.4159
Interaction p-value											0.9324
First line treatment out	come (eCR	RF)									
NED [PDS]	89	3 (3.4)	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	1.52	0.20, 30.77	0.7066
NED/CR [IDS]	74	7 (9.5)	NE (NE,	NE)	32	3 (9.4)	NE (NE,	NE)	1.00	0.28, 4.66	0.9959
NED/CR [Chemo]	39	2 (5.1)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	2 (4.0)	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											0.7529
Screening laboratory tBR0	CA status	s (eCRF)									
tBRCAm	147	8 (5.4)	NE (NE,	NE)	67	3 (4.5)	NE (NE,	NE)	1.19	0.35, 5.45	0.7908
non-tBRCAm	108	6 (5.6)	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	3.57	0.61, 67.41	0.1758
Interaction p-value											0.3672
Age group											
<65 years	185	8 (4.3)	NE (NE,	NE)	98	4 (4.1)	NE (NE,	NE)	1.03	0.32, 3.86	0.9598
>=65 years	70	6 (8.6)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeace 11AUG2022:11:30 kpzx329

Table 3.4.57 PAOLA1: Summary of subgroup analysis of AESI G1-2: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=25		mab			Placebo + be (N=13		nab			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		С	Number (%) of patients with events	(95	n time % CI) ns) [a]	- Hazard ratio [b]	95% CI [b]	2-sided p-value [b]	
FIGO Stage (Disease state)												
III	182	11 (6.0)	NE (NE,	NE)	89	3 (3.4)	NE (NE, N	E) 1.77	0.55, 7.82	0.3575
IV	73	3 (4.1)	NE (NE,	NE)	42	1 (2.4)	NE (NE, N	E) 1.74	0.22, 35.13	0.6176
Interaction p-value												0.9898
Region												
Europe	245	13 (5.3)	NE (NE,	NE)	125	4 (3.2)	NE (NE, N	E) 1.64	0.58, 5.81	0.3692
Japan	10	1 (10.0)	NE (NE,	NE)	6	0	NE (NE, N	E) NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Basel	ine										
(0) Normal activity	190	10 (5.3)	NE (NE,	NE)	100	3 (3.0)	NE (NE, N	E) 1.73	0.53, 7.74	0.3812
(1) Restricted activity	61	4 (6.6)	NE (NE,	NE)	30	1 (3.3)	NE (NE, N	E) 1.98	0.29, 38.62	0.5166
Interaction p-value												0.9196
Baseline CA-125 value												
<=ULN	228	13 (5.7)	,	NE,	NE)	117	4 (3.4)	NE (NE, N	,	0.58, 5.86	0.3609
>ULN	27	1 (3.7)	NE (NE,	NE)	14	0	NE (NE, N	E) NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	14 (5.5)	NE (NE,	NE)	131	4 (3.1)	NE (NE, N	E) 1.78	0.64, 6.29	0.2835
Interaction p-value												NC
Cytoreductive surgery outco	me											
No residue	166	10 (6.0)	NE (NE,	NE)	80	4 (5.0)	NE (NE, N	E) 1.18	0.39, 4.30	0.7790
Residue	79	3 (3.8)	NE (NE,	NE)	43	0	NE (NE, N	E) NC	NC	NC
Interaction p-value												NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeace 11AUG2022:11:30 kpzx329

Table 3.4.57 PAOLA1: Summary of subgroup analysis of AESI G1-2: Thrombocytopenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		Placebo + b (N=1				
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]		2-sided p-value [b]
Timing of cytoreductive s	surgery							
Upfront	146	5 (3.4)	NE (NE, NE)	78 1 (1.3)	NE (NE, NE)	2.61	0.42, 49.93	0.3345
Interval	99	8 (8.1)	NE (NE, NE)	45 3 (6.7)	NE (NE, NE)	1.22	0.35, 5.55	0.7695
Interaction p-value								0.5400
Myriad tumour BRCA mutat:	ion statu	ıs						
tBRCAm	158	9 (5.7)	NE (NE, NE)	77 4 (5.2)	NE (NE, NE)	1.06	0.34, 3.89	0.9287
Non-tBRCAm	97	5 (5.2)	NE (NE, NE)	54 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC
Status somatic BRCA mutat	tions							
sBRCAm	25	2 (8.0)	NE (NE, NE)	9 0	NE (NE, NE)	NC	NC	NC
gBRCAm	69	4 (5.8)	NE (NE, NE)	36 4 (11.1)	NE (NE, NE)	NC	NC	NC
Non-BRCAm	43	3 (7.0)	NE (NE, NE)	23 0	NE (NE, NE)	NC	NC	NC
Interaction p-value								NC

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeace 11AUG2022:11:30 kpzx329

Table 3.4.58 PAOLA1: Summary of subgroup analysis of AESI G1-2: Nausea Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + N			Placebo + be (N=1)			•		
Subgroup	n	Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median t (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IV	RS)								
NED [PDS]	92	52 (56.5)	3.7 (0.8, NE)	48	11 (22.9)	NE (NE,	NE)	3.29	1.79, 6.66	<0.0001*
NED/CR [IDS]	74	47 (63.5)	1.4 (0.3, 5.0)	38	13 (34.2)	NE (NE,	NE)	2.37	1.32, 4.57	0.0031*
NED/CR [Chemo]	40	21 (52.5)	0.6 (0.3, NE)	20	2 (10.0)	NE (NE,	NE)	7.94	2.33, 49.67	0.0002*
PR	49	22 (44.9)	NE (NE, NE)	25	4 (16.0)	NE (NE,	NE)	3.51	1.34, 12.01	0.0084*
Interaction p-value										0.4170
Screening laboratory tBR0	CA statu	ıs (IVRS)								
tBRCAm	150	85 (56.7)	3.7 (0.8, NE)	65	12 (18.5)	NE (NE,	NE)	4.10	2.33, 7.91	<0.0001*
non-tBRCAm	105	57 (54.3)	3.3 (0.3, NE)	66	18 (27.3)	NE (NE,	NE)	2.62	1.58, 4.59	0.0001*
Interaction p-value										0.2713
First line treatment out	come (eC	RF)								
NED [PDS]	89	51 (57.3)	2.7 (0.5, NE)	47	10 (21.3)	NE (NE,	NE)	3.64	1.93, 7.60	<0.0001*
NED/CR [IDS]	74	48 (64.9)	1.3 (0.3, 4.3)	32	10 (31.3)	NE (NE,	NE)	2.80	1.48, 5.88	0.0010*
NED/CR [Chemo]	39	19 (48.7)	NE (NE, NE)	17	2 (11.8)	NE (NE,	NE)	6.15	1.79, 38.62	0.0021*
PR	50	23 (46.0)	NE (NE, NE)	34	7 (20.6)	NE (NE,	NE)	2.61	1.18, 6.57	0.0171*
Interaction p-value										0.7038
Screening laboratory tBR0	CA statu	ıs (eCRF)								
tBRCAm	147	83 (56.5)	3.7 (1.1, NE)	67	13 (19.4)	NE (NE,	NE)	3.85	2.23, 7.25	<0.0001*
non-tBRCAm	108	59 (54.6)	2.7 (0.3, NE)	64	17 (26.6)	NE (NE,	NE)	2.74	1.63, 4.85	<0.0001*
Interaction p-value										0.3983
Age group										
<65 years	185	108 (58.4)	2.3 (0.8,10.3)	98	21 (21.4)	NE (NE,	NE)	3.71	2.37, 6.09	<0.0001*
>=65 years	70	34 (48.6)	NE (NE, NE)	33	9 (27.3)	NE (NE,	NE)	2.21	1.11, 4.91	0.0234*
Interaction p-value										0.2554

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacf 11AUG2022:11:30 kpzx329

Table 3.4.58 PAOLA1: Summary of subgroup analysis of AESI G1-2: Nausea Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + N=2			Placebo + be (N=1					
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	Median (95% C	I)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)										
III	182	98 (53.8)	5.4 (0.9, NE)	89	25 (28.1)	NE (NE	, NE)	2.44	1.60, 3.87	<0.0001*
IV	73	44 (60.3)	2.1 (0.4, NE)	42	5 (11.9)	NE (NE	, NE)	7.29	3.18, 21.06	<0.0001*
Interaction p-value										0.0233*
Region										
Europe	245	139 (56.7)	2.9 (0.9,16.0)	125	30 (24.0)	NE (NE	, NE)	3.15	2.15, 4.76	<0.0001*
Japan	10	3 (30.0)	NE (NE, NE)	6	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value										NC
ECOG performance status at	Basel	ine								
(0) Normal activity	190	111 (58.4)	2.1 (0.8, 8.0)	100	19 (19.0)	NE (NE	, NE)	4.18	2.64, 7.03	<0.0001*
(1) Restricted activity	61	30 (49.2)	NE (NE, NE)	30	11 (36.7)	NE (NE	, NE)	1.66	0.86, 3.47	0.1351
Interaction p-value										0.0375*
Baseline CA-125 value										
<=ULN	228	127 (55.7)	3.3 (0.9, NE)	117	28 (23.9)	NE (NE	, NE)	3.07	2.07, 4.71	<0.0001*
>ULN	27	15 (55.6)	4.1 (0.1, NE)	14	2 (14.3)	NE (NE	, NE)	5.45	1.54, 34.60	0.0061*
Interaction p-value										0.4334
Histological grade										
High grade	255	142 (55.7)	3.4 (1.1, NE)	131	30 (22.9)	NE (NE	, NE)	3.22	2.21, 4.87	<0.0001*
Interaction p-value										NC
Cytoreductive surgery outco	ome									
No residue	166	100 (60.2)	1.9 (0.6, 7.8)	80	21 (26.3)	NE (NE	, NE)	3.10	1.98, 5.10	<0.0001*
Residue	79	38 (48.1)	NE (NE, NE)	43	9 (20.9)	NE (NE	, NE)	2.92	1.48, 6.45	0.0014*
Interaction p-value										0.8952

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacf 11AUG2022:11:30 kpzx329

Table 3.4.58 PAOLA1: Summary of subgroup analysis of AESI G1-2: Nausea Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + k (N=2			Placebo + be (N=1)		mab						
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events	nts (95% CI) ratio		95% CI	[b]	2-sided p-value [b]			
Timing of cytoreductive s	surgery												
Upfront	146	76 (52.1)	7.9 (0.8, NE)	78	14 (17.9)	NE (NE, N	E) 3.	.92	2.29,	7.23	<0.0001*	
Interval	99	62 (62.6)	2.0 (0.4, 6.1)	45	16 (35.6)	NE (NE, N	E) 2.	. 24	1.33,	4.02	0.0020*	
Interaction p-value												0.1677	
Myriad tumour BRCA mutati	ion stat	us											
tBRCAm	158	90 (57.0)	2.9 (0.8, NE)	77	15 (19.5)	NE (NE, N	E) 3.	. 87	2.31,	6.97	<0.0001*	
Non-tBRCAm	97	52 (53.6)	3.4 (0.4, NE)	54	15 (27.8)	NE (NE, N	E) 2.	. 56	1.48,	4.71	0.0006*	
Interaction p-value												0.3053	
Status somatic BRCA mutat	cions												
sBRCAm	25	16 (64.0)	0.8 (0.1, NE)	9	2 (22.2)	NE (NE, N	E) 3.	.92	1.11,	24.81	0.0311*	
gBRCAm	69	43 (62.3)	2.3 (0.5, 6.1)	36	8 (22.2)	NE (NE, N	E) 3.	. 88	1.92,	8.92	<0.0001*	
Non-BRCAm	43	23 (53.5)	2.7 (0.3, NE)	23	7 (30.4)	NE (NE, N	E) 2.	. 37	1.07,	5.98	0.0327*	
Interaction p-value												0.6746	

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacf 11AUG2022:11:30 kpzx329

Table 3.4.59 PAOLA1: Summary of subgroup analysis of AESI G1-2: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ımab			Placebo + be (N=13		ımab					
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI)	C	Number (%) of patients with events	(95	an ti 5% CI ths))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV)	RS)												
NED [PDS]	92	23 (25.0)	NE (NE,	NE)	48	5 (10.4)	NE (NE,	NE)	2.56	1.06,	7.63	0.0364*
NED/CR [IDS]	74	16 (21.6)	NE (NE,	NE)	38	6 (15.8)	NE (NE,	NE)	1.41	0.58,	3.92	0.4641
NED/CR [Chemo]	40	4 (10.0)	NE (NE,	NE)	20	1 (5.0)	NE (NE,	NE)	2.06	0.31,	40.33	0.4887
PR	49	9 (18.4)	NE (NE,	NE)	25	2 (8.0)	NE (NE,	NE)	2.52	0.65,	16.53	0.1962
Interaction p-value														0.8336
Screening laboratory tBR0	CA statu	s (IVRS)												
tBRCAm	150	32 (21.3)	NE (NE,	NE)	65	5 (7.7)	NE (NE,	NE)	2.93	1.25,	8.57	0.0114*
non-tBRCAm	105	20 (19.0)	NE (NE,	NE)	66	9 (13.6)	NE (NE,	NE)	1.47	0.69,	3.40	0.3235
Interaction p-value														0.2642
First line treatment out	come (eCl	RF)												
NED [PDS]	89	21 (23.6)	NE (NE,	NE)	47	5 (10.6)	NE (NE,	NE)	2.37	0.97,	7.10	0.0604
NED/CR [IDS]	74	15 (20.3)	NE (NE,	NE)	32	4 (12.5)	NE (NE,	NE)	1.71	0.62,	5.99	0.3176
NED/CR [Chemo]	39	5 (12.8)	NE (NE,	NE)	17	2 (11.8)	NE (NE,	NE)	1.14	0.25,	7.99	0.8703
PR	50	9 (18.0)	NE (NE,	NE)	34	3 (8.8)	NE (NE,	NE)	2.11	0.63,	9.53	0.2357
Interaction p-value														0.8964
Screening laboratory tBR0	CA statu	s (eCRF)												
tBRCAm	147	31 (21.1)	NE (NE,	NE)	67	5 (7.5)	NE (NE,	NE)	2.98	1.27,	8.74	0.0102*
non-tBRCAm	108	21 (19.4)	NE (NE,	NE)	64	9 (14.1)	NE (NE,	NE)	1.46	0.69,	3.36	0.3311
Interaction p-value														0.2447
Age group														
<65 years	185	41 (22.2)	NE (NE,	NE)	98	8 (8.2)	NE (NE,	NE)	2.93	1.45,	6.74	0.0019*
>=65 years	70	11 (15.7)	NE (NE,	NE)	33	6 (18.2)	NE (NE,	NE)	0.86	0.33,	2.50	0.7681
Interaction p-value														0.0596

[a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacg 11AUG2022:11:30 kpzx329

Table 3.4.59 PAOLA1: Summary of subgroup analysis of AESI G1-2: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		mab	,		Placebo + be (N=13		ab	:		
Subgroup		Number (%) of patients with events	(95	Median time (95% CI) (months) [a] Number (%) Median time (95% CI) (months) [a] (months) [a]	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]				
FIGO Stage (Disease state)												
III	182	39 (21.4)	NE (NE,	NE)	89	10 (11.2)	NE (NE, NE)	2.02	1.05, 4.28	0.0342*
IV	73	13 (17.8)	NE (NE,	NE)	42	4 (9.5)	NE (NE, NE)	1.95	0.69, 6.94	0.2167
Interaction p-value												0.9593
Region												
Europe	245	52 (21.2)	NE (NE,	NE)	125	13 (10.4)	NE (NE, NE)	2.17	1.22, 4.16	0.0074*
Japan	10	0	NE (NE,	NE)	6	1 (16.7)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Basel	ine										
(0) Normal activity	190	39 (20.5)	NE (NE,	NE)	100	11 (11.0)	NE (NE, NE)	1.94	1.03, 3.99	0.0393*
(1) Restricted activity	61	13 (21.3)	NE (NE,	NE)	30	3 (10.0)	NE (NE, NE)	2.36	0.76, 10.30	0.1451
Interaction p-value												0.7858
Baseline CA-125 value												
<=ULN	228	41 (18.0)	NE (NE,	NE)	117	13 (11.1)	NE (NE, NE)	1.68	0.93, 3.27	0.0876
>ULN	27	11 (40.7)	NE (NE,	NE)	14	1 (7.1)	NE (NE, NE)	6.85	1.33,125.10	0.0172*
Interaction p-value												0.1355
Histological grade												
High grade	255	52 (20.4)	NE (NE,	NE)	131	14 (10.7)	NE (NE, NE)	2.01	1.15, 3.78	0.0134*
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	38 (22.9)	NE (NE,	NE)	80	9 (11.3)	NE (NE, NE)	2.17	1.10, 4.78	0.0248*
Residue	79	12 (15.2)	NE (NE,	NE)	43	5 (11.6)	NE (NE, NE)	1.34	0.50, 4.20	0.5783
Interaction p-value												0.4618

Table 3.4.59 PAOLA1: Summary of subgroup analysis of AESI G1-2: Vomiting Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		Placebo + b (N=1			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) Median time Hazard of patients (95% CI) ratio n with events (months) [a] [b] 95		ratio	2-sided p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	29 (19.9)	NE (NE, NE)	78 7 (9.0)	NE (NE, NE)	2.35 1.09, 5.83	0.0281*
Interval	99	21 (21.2)	NE (NE, NE)	45 7 (15.6)	NE (NE, NE)	1.41 0.63, 3.58	0.4185
Interaction p-value							0.4006
Myriad tumour BRCA mutat:	ion stat	us					
tBRCAm	158	34 (21.5)	NE (NE, NE)	77 6 (7.8)	NE (NE, NE)	2.93 1.32, 7.75	0.0064*
Non-tBRCAm	97	18 (18.6)	NE (NE, NE)	54 8 (14.8)	NE (NE, NE)	1.31 0.59, 3.20	0.5142
Interaction p-value							0.1884
Status somatic BRCA muta	tions						
sBRCAm	25	7 (28.0)	NE (NE, NE)	9 1 (11.1)	NE (NE, NE)	2.77 0.49, 51.85	0.2807
gBRCAm	69	15 (21.7)	NE (NE, NE)	36 3 (8.3)	NE (NE, NE)	2.86 0.94, 12.36	0.0642
Non-BRCAm	43	5 (11.6)	NE (NE, NE)	23 5 (21.7)	NE (NE, NE)	0.52 0.14, 1.85	0.2994
Interaction p-value							0.1165

Table 3.4.60 PAOLA1: Summary of subgroup analysis of AESI G1-2: Fatigue and Asthenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + 1		:-		Placebo + b (N=1		ab	,			2-sided p-value [b]
Subgroup	n	Number (%) of patients with events	Median ti (95% CI) (months) [)		Number (%) of patients with events	(95%	n time (CI) (S) [a]	Hazard ratio [b]	95% CI	I [b]	
First line treatment out	come (IV	RS)										
NED [PDS]	92	47 (51.1)	13.2 (2.9,	NE)	48	17 (35.4)	NE (NE, NE)	1.72	1.01,	3.07	0.0477*
NED/CR [IDS]	74	45 (60.8)	9.1 (2.2,2	22.1)	38	15 (39.5)	NE (NE, NE)	1.73	0.99,	3.20	0.0565
NED/CR [Chemo]	40	17 (42.5)	NE (NE,	NE)	20	5 (25.0)	NE (NE, NE)	1.99	0.79,	6.07	0.1512
PR	49	24 (49.0)	14.9 (4.2,	NE)	25	7 (28.0)	NE (NE, NE)	2.01	0.91,	5.05	0.0854
Interaction p-value												0.9843
Screening laboratory tBR0	CA statu	ıs (IVRS)										
tBRCAm	150	78 (52.0)	13.4 (5.4,	NE)	65	23 (35.4)	NE (NE, NE)	1.63	1.04,	2.65	0.0322*
non-tBRCAm	105	55 (52.4)	13.0 (3.5,	NE)	66	21 (31.8)	NE (NE, NE)	2.00	1.23,	3.38	0.0047*
Interaction p-value												0.5549
First line treatment out	come (eC	PRF)										
NED [PDS]	89	46 (51.7)	13.0 (2.9,	NE)	47	18 (38.3)	NE (NE, NE)	1.58	0.93,	2.80	0.0889
NED/CR [IDS]	74	44 (59.5)	6.2 (2.1,	NE)	32	9 (28.1)	NE (NE, NE)	2.77	1.42,	6.07	0.0020*
NED/CR [Chemo]	39	16 (41.0)	NE (NE,	NE)	17	4 (23.5)	NE (NE, NE)	2.04	0.75,	7.11	0.1747
PR	50	26 (52.0)	15.3 (4.2,	NE)	34	13 (38.2)	NE (NE, NE)	1.34	0.70,	2.68	0.3858
Interaction p-value												0.4705
Screening laboratory tBR0	CA statu	ıs (eCRF)										
tBRCAm	147	76 (51.7)	14.9 (5.6,	NE)	67	23 (34.3)	NE (NE, NE)	1.68	1.07,	2.74	0.0230*
non-tBRCAm	108	57 (52.8)	11.3 (3.5,	NE)	64	21 (32.8)	NE (NE, NE)	1.95	1.20,	3.28	0.0064*
Interaction p-value												0.6725
Age group												
<65 years	185	100 (54.1)	11.1 (4.9,	NE)	98	30 (30.6)	NE (NE, NE)	2.12	1.43,	3.25	0.0001*
>=65 years	70	33 (47.1)	NE (NE,	NE)	33	14 (42.4)	NE (NE, NE)	1.16	0.63,	2.23	0.6426
Interaction p-value												0.1179

Table 3.4.60 PAOLA1: Summary of subgroup analysis of AESI G1-2: Fatigue and Asthenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		_	bevacizumab 255)			Placebo + be (N=1						
Subgroup		Number (%) of patients with events)		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	[b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	93 (51.1)	15.8 (6.0,	NE)	89	31 (34.8)	NE (NE,	NE)	1.65	1.11,	2.52	0.0119*
IV	73	40 (54.8)	11.3 (2.1,	NE)	42	13 (31.0)	NE (NE,	NE)	2.20	1.21,	4.28	0.0089*
Interaction p-value												0.4485
Region												
Europe	245	133 (54.3)	11.3 (5.1,	NE)	125	44 (35.2)	NE (NE,	NE)	1.80	1.29,	2.56	0.0004*
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC		NC
Interaction p-value												NC
ECOG performance status at	Basel	ine										
(0) Normal activity	190	101 (53.2)	11.0 (3.7,	NE)	100	32 (32.0)	NE (NE,	NE)	1.96	1.34,	2.97	0.0005*
(1) Restricted activity	61	28 (45.9)	NE (NE,	NE)	30	11 (36.7)	NE (NE,	NE)	1.34	0.68,	2.80	0.4057
Interaction p-value												0.3549
Baseline CA-125 value												
<=ULN			15.3 (7.1,		117	39 (33.3)	NE (NE,	NE)	1.76	1.24,		0.0014*
>ULN	27	16 (59.3)	2.1 (0.5,	NE)	14	5 (35.7)	NE (NE,	NE)	2.18	0.85,	6.67	0.1064
Interaction p-value												0.6906
Histological grade												
High grade	255	133 (52.2)	13.4 (6.0,	NE)	131	44 (33.6)	NE (NE,	NE)	1.80	1.29,	2.56	0.0004*
Interaction p-value												NC
Cytoreductive surgery outco	me											
No residue	166	91 (54.8)	9.1 (3.5,	NE)	80	27 (33.8)	NE (NE,	NE)	2.00	1.32,	3.12	0.0009*
Residue	79	38 (48.1)	NE (NE,	NE)	43	13 (30.2)	NE (NE,	NE)	1.68	0.92,	3.28	0.0926
Interaction p-value												0.6623

Table 3.4.60 PAOLA1: Summary of subgroup analysis of AESI G1-2: Fatigue and Asthenia Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + 1 (N=2				oevacizumab 131)		:			
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) of patients with events)	Hazard ratio [b] 95% CI [b]		2-sided p-value [b]	
Timing of cytoreductive s	surgery										
Upfront	146	72 (49.3)	NE (NE, NE)	78	23 (29.5)	NE (NE,	NE)	1.96	1.25,	3.21	0.0030*
Interval	99	57 (57.6)	9.8 (2.4, NE)	45	17 (37.8)	NE (NE,	NE)	1.75	1.04,	3.10	0.0344*
Interaction p-value											0.7501
Myriad tumour BRCA mutati	on stat	us									
tBRCAm	158	83 (52.5)	13.4 (5.4, NE)	77	26 (33.8)	NE (NE,	NE)	1.73	1.13,	2.74	0.0106*
Non-tBRCAm	97	50 (51.5)	13.0 (3.5, NE)	54	18 (33.3)	NE (NE,	NE)	1.89	1.13,	3.33	0.0153*
Interaction p-value											0.8029
Status somatic BRCA mutat	ions										
sBRCAm	25	12 (48.0)	NE (NE, NE)	9	0	NE (NE,	NE)	NC	NC		NC
gBRCAm	69	45 (65.2)	3.5 (1.4,11.8)	36	16 (44.4)	NE (NE,	NE)	1.84	1.06,	3.36	0.0292*
Non-BRCAm	43	23 (53.5)	11.0 (2.0, NE)	23	12 (52.2)	10.4 (2.9,	NE)	1.15	0.58,	2.38	0.6996
Interaction p-value											0.3066

Table 3.4.61 PAOLA1: Summary of subgroup analysis of AESI G1-2: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	:	Olaparib + b (N=2		ımab				bevacizumab 131)		;			
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI)		Number (%) of patients with events	Median t: (95% CI (months))	Hazard ratio [b]	95% CI	[b]	2-sided p-value [b]
First line treatment out	come (IV	RS)											
NED [PDS]	92	41 (44.6)	NE (NE,	NE)	48	22 (45.8)	NE (NE,	NE)	1.07	0.65,	1.83	0.7854
NED/CR [IDS]	74	25 (33.8)	NE (NE,	NE)	38	16 (42.1)	NE (NE,	NE)	0.75	0.41,	1.44	0.3849
NED/CR [Chemo]	40	15 (37.5)	NE (NE,	NE)	20	11 (55.0)	6.9 (1.4,	NE)	0.60	0.28,	1.35	0.2100
PR	49	17 (34.7)	NE (NE,	NE)	25	10 (40.0)	NE (NE,	NE)	0.82	0.38,	1.85	0.6149
Interaction p-value													0.6405
Screening laboratory tBR0	CA statu	s (IVRS)											
tBRCAm	150	54 (36.0)	NE (NE,	NE)	65	26 (40.0)	NE (NE,	NE)	0.94	0.60,	1.52	0.7974
non-tBRCAm	105	44 (41.9)	NE (NE,	NE)	66	33 (50.0)	10.4 (4.1,	NE)	0.79	0.50,	1.25	0.3032
Interaction p-value													0.5924
First line treatment out	come (eCl	RF)											
NED [PDS]	89	40 (44.9)	NE (NE,	NE)	47	21 (44.7)	NE (NE,	NE)	1.12	0.67,	1.93	0.6822
NED/CR [IDS]	74	23 (31.1)	NE (NE,	NE)	32	12 (37.5)	NE (NE,	NE)	0.80	0.40,	1.65	0.5261
NED/CR [Chemo]	39	17 (43.6)	NE (NE,	NE)	17	8 (47.1)	NE (NE,	NE)	0.97	0.43,	2.37	0.9345
PR	50	17 (34.0)	NE (NE,	NE)	34	17 (50.0)	9.2 (3.4,	NE)	0.57	0.29,	1.12	0.0993
Interaction p-value													0.4654
Screening laboratory tBR0	CA statu	s (eCRF)											
tBRCAm	147	53 (36.1)	NE (NE,	NE)	67	26 (38.8)	NE (NE,	NE)	0.98	0.62,	1.59	0.9361
non-tBRCAm	108	45 (41.7)	NE (NE,	NE)	64	33 (51.6)	7.6 (3.6,	NE)	0.74	0.48,	1.17	0.2008
Interaction p-value													0.4027
Age group													
<65 years	185	69 (37.3)	NE (NE,	NE)	98	44 (44.9)	NE (NE,	NE)	0.81	0.56,	1.20	0.2907
>=65 years	70	29 (41.4)	NE (NE,	NE)	33	15 (45.5)	NE (NE,	NE)	0.92	0.50,	1.76	0.7911
Interaction p-value													0.7443

Table 3.4.61 PAOLA1: Summary of subgroup analysis of AESI G1-2: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b		ımab			Placebo + b						2-sided p-value [b]
Subgroup		Number (%) of patients with events	Media (95 (mont	% CI))		Number (%) of patients with events	Median ti (95% CI) (months)		Hazard ratio [b]	95% CI	5% CI [b]	
FIGO Stage (Disease state)													
III	182	68 (37.4)	NE (NE,	NE)	89	41 (46.1)	NE (NE,	NE)	0.80	0.55,	1.19	0.2622
IV	73	30 (41.1)	NE (NE,	NE)	42	18 (42.9)	NE (NE,	NE)	0.96	0.54,	1.75	0.8794
Interaction p-value													0.6165
Region													
Europe	245	95 (38.8)	NE (NE,	NE)	125	53 (42.4)	NE (NE,	NE)	0.92	0.66,	1.30	0.6278
Japan	10	3 (30.0)	NE (NE,	NE)	6	6 (100)	3.5 (0.3,	NE)	0.19	0.04,	0.73	0.0154*
Interaction p-value													0.0254*
ECOG performance status at	Basel	ine											
(0) Normal activity	190	73 (38.4)	NE (NE,	NE)	100	45 (45.0)	NE (NE,	NE)	0.82	0.57,	1.19	0.2883
(1) Restricted activity	61	24 (39.3)	NE (NE,	NE)	30	14 (46.7)	NE (NE,	NE)	0.92	0.48,	1.83	0.8142
Interaction p-value													0.7479
Baseline CA-125 value													
<=ULN	228	89 (39.0)	NE (NE,	NE)	117	53 (45.3)	NE (NE,	NE)	0.85	0.61,		0.3542
>ULN	27	9 (33.3)	NE (NE,	NE)	14	6 (42.9)	NE (NE,	NE)	0.78	0.28,	2.33	0.6397
Interaction p-value													0.8755
Histological grade													
High grade	255	98 (38.4)	NE (NE,	NE)	131	59 (45.0)	NE (NE,	NE)	0.84	0.61,	1.17	0.3060
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	64 (38.6)	NE (NE,	NE)	80	34 (42.5)	NE (NE,	NE)	0.94	0.63,	1.44	0.7810
Residue	79	30 (38.0)	NE (NE,	NE)	43	21 (48.8)	NE (NE,	NE)	0.71	0.41,	1.26	0.2419
Interaction p-value													0.4346

Table 3.4.61 PAOLA1: Summary of subgroup analysis of AESI G1-2: Hypertension Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		Placebo + k (N=1	pevacizumab 131)		
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) Median time Hazard of patients (95% CI) ratio n with events (months) [a] [b]		ratio	2-sided p-value [b]
Timing of cytoreductive s	surgery						
Upfront	146	60 (41.1)	NE (NE, NE)	78 37 (47.4)	NE (NE, NE)	0.88 0.59, 1.34	0.5480
Interval	99	34 (34.3)	NE (NE, NE)	45 18 (40.0)	NE (NE, NE)	0.83 0.48, 1.51	0.5384
Interaction p-value							0.8786
Myriad tumour BRCA mutat:	ion stat	us					
tBRCAm	158	59 (37.3)	NE (NE, NE)	77 31 (40.3)	NE (NE, NE)	0.97 0.63, 1.52	0.8982
Non-tBRCAm	97	39 (40.2)	NE (NE, NE)	54 28 (51.9)	9.0 (3.6, NE)	0.70 0.43, 1.15	0.1565
Interaction p-value							0.3255
Status somatic BRCA muta	tions						
sBRCAm	25	7 (28.0)	NE (NE, NE)	9 1 (11.1)	NE (NE, NE)	2.78 0.49, 51.95	0.2797
gBRCAm	69	24 (34.8)	NE (NE, NE)	36 16 (44.4)	NE (NE, NE)	0.78 0.42, 1.49	0.4431
Non-BRCAm	43	15 (34.9)	NE (NE, NE)	23 13 (56.5)	7.6 (2.2, NE)	0.48 0.23, 1.03	0.0579
Interaction p-value							0.1856

Table 3.4.62 PAOLA1: Summary of subgroup analysis of AESI G1-2: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	:	Olaparib + b (N=2		ab		Placebo + be (N=1			:		2-sided p-value [b]
Subgroup	(Number (%) of patients with events	Median (95% (month)	CI)		Number (%) of patients with events	Median t (95% CI (months)	[)	Hazard ratio [b]	95% CI [b]	
First line treatment out	come (IVI	RS)									
NED [PDS]	92	5 (5.4)	NE (NE, NE)	48	9 (18.8)	NE (NE	NE)	0.25	0.08, 0.73	0.0115*
NED/CR [IDS]	74	6 (8.1)	NE (NE, NE)	38	3 (7.9)	NE (NE	NE)	1.03	0.27, 4.86	0.9720
NED/CR [Chemo]	40	3 (7.5)	NE (NE, NE)	20	3 (15.0)	NE (NE	NE)	0.44	0.08, 2.36	0.3170
PR	49	4 (8.2)	NE (NE, NE)	25	4 (16.0)	NE (NE	NE)	0.49	0.12, 2.07	0.3181
Interaction p-value											0.4634
Screening laboratory tBR0	CA status	s (IVRS)									
tBRCAm	150	13 (8.7)	NE (NE, NE)	65	9 (13.8)	NE (NE	NE)	0.58	0.25, 1.40	0.2174
non-tBRCAm	105	5 (4.8)	NE (NE, NE)	66	10 (15.2)	NE (NE	NE)	0.29	0.09, 0.82	0.0191*
Interaction p-value											0.3176
First line treatment out	come (eCI	RF)									
NED [PDS]	89	4 (4.5)	NE (NE, NE)	47	8 (17.0)	NE (NE	NE)	0.23	0.06, 0.74	0.0135*
NED/CR [IDS]	74	8 (10.8)	NE (NE, NE)	32	2 (6.3)	NE (NE	NE)	1.75	0.44, 11.62	0.4535
NED/CR [Chemo]	39	1 (2.6)	NE (NE, NE)	17	2 (11.8)	NE (NE	NE)	0.20	0.01, 2.07	0.1704
PR	50	4 (8.0)	NE (NE, NE)	34	6 (17.6)	NE (NE	NE)	0.41	0.11, 1.45	0.1655
Interaction p-value											0.1444
Screening laboratory tBR0	CA status	s (eCRF)									
tBRCAm	147	12 (8.2)	NE (NE, NE)	67	9 (13.4)	NE (NE	NE)	0.56	0.24, 1.38	0.1994
non-tBRCAm	108	6 (5.6)	NE (NE, NE)	64	10 (15.6)	NE (NE	NE)	0.33	0.11, 0.89	0.0283*
Interaction p-value											0.4301
Age group											
<65 years	185	13 (7.0)	NE (NE, NE)	98	17 (17.3)	NE (NE	NE)	0.37	0.17, 0.75	0.0062*
>=65 years	70	5 (7.1)	NE (NE, NE)	33	2 (6.1)	NE (NE	NE)	1.17	0.25, 8.18	0.8479
Interaction p-value											0.1829

Table 3.4.62 PAOLA1: Summary of subgroup analysis of AESI G1-2: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2				Placebo + be (N=1)			:			2-sided p-value [b]
Subgroup		Number (%) of patients with events	Median t (95% CI (months)	:)		Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI	95% CI [b]	
FIGO Stage (Disease state)												
III	182	13 (7.1)	NE (NE,	NE)	89	12 (13.5)	NE (NE,	NE)	0.49	0.22,	1.08	0.0776
IV	73	5 (6.8)	NE (NE,	NE)	42	7 (16.7)	NE (NE,	NE)	0.39	0.11,	1.21	0.1008
Interaction p-value												0.7386
Region												
Europe	245	18 (7.3)	NE (NE,	NE)	125	16 (12.8)	NE (NE,	NE)	0.54	0.27,	1.06	0.0735
Japan	10	0	NE (NE,	NE)	6	3 (50.0)	NE (NE,	NE)	NC	NC		NC
Interaction p-value												NC
ECOG performance status at	Basel	ine										
(0) Normal activity	190	14 (7.4)	NE (NE,	NE)	100	13 (13.0)	NE (NE,	NE)	0.53	0.25,	1.15	0.1062
(1) Restricted activity	61	4 (6.6)	NE (NE,	NE)	30	6 (20.0)	NE (NE,	NE)	0.29	0.07,	1.01	0.0520
Interaction p-value												0.4113
Baseline CA-125 value												
<=ULN	228	16 (7.0)	NE (NE,	NE)	117	17 (14.5)	NE (NE,	NE)	0.45	0.22,	0.89	0.0225*
>ULN	27	2 (7.4)	NE (NE,	NE)	14	2 (14.3)	NE (NE,	NE)	0.47	0.06,	3.92	0.4561
Interaction p-value												0.9635
Histological grade												
High grade	255	18 (7.1)	NE (NE,	NE)	131	19 (14.5)	NE (NE,	NE)	0.45	0.23,	0.86	0.0164*
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	13 (7.8)	NE (NE,	NE)	80	11 (13.8)	NE (NE,	NE)	0.53	0.24,	1.22	0.1316
Residue	79	4 (5.1)	NE (NE,	NE)	43	7 (16.3)	NE (NE,	NE)	0.28	0.07,	0.92	0.0356*
Interaction p-value												0.3748

Table 3.4.62 PAOLA1: Summary of subgroup analysis of AESI G1-2: Proteinuria Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2			Placebo + b (N=1				
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]		Number (%) Median time Hazard of patients (95% CI) ratio n with events (months) [a] [b]		95% CI [b	2-sided o] p-value[b]	
Timing of cytoreductive s	surgery								
Upfront	146	7 (4.8)	NE (NE, N	TE) 78	14 (17.9)	NE (NE, NE)	0.23	0.09, 0.	.56 0.0011*
Interval	99	10 (10.1)	NE (NE, N	IE) 45	4 (8.9)	NE (NE, NE)	1.14	0.38, 4.	16 0.8230
Interaction p-value									0.0273*
Myriad tumour BRCA mutat:	ion stat	us							
tBRCAm	158	14 (8.9)	NE (NE, N	ΙΕ) 77	10 (13.0)	NE (NE, NE)	0.63	0.28, 1.	.46 0.2719
Non-tBRCAm	97	4 (4.1)	NE (NE, N	ΙΕ) 54	9 (16.7)	NE (NE, NE)	0.23	0.06, 0.	71 0.0099*
Interaction p-value									0.1569
Status somatic BRCA mutat	cions								
sBRCAm	25	2 (8.0)	NE (NE, N	ΙΕ) 9	2 (22.2)	NE (NE, NE)	0.30	0.04, 2.	.46 0.2365
gBRCAm	69	6 (8.7)	NE (NE, N	ΙΕ) 36	7 (19.4)	NE (NE, NE)	0.42	0.13, 1.	26 0.1197
Non-BRCAm	43	2 (4.7)	NE (NE, N	ΙΕ) 23	2 (8.7)	NE (NE, NE)	0.53	0.06, 4.	41 0.5276
Interaction p-value									0.9172

Table 3.4.63 PAOLA1: Summary of subgroup analysis of AESI G1-2: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + k (N=2		nab	•		Placebo + be (N=1)					
Subgroup	c	Number (%) of patients with events	Median (95% (month	CI)		C	Number (%) of patients with events	Median (95% C (months)	I)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	LS)										
NED [PDS]	92	1 (1.1)	NE (NE,	NE)	48	3 (6.3)	NE (NE	, NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE	, NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE	, NE)	NC	NC	NC
PR	49	1 (2.0)	NE (NE,	NE)	25	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR	CA status	(IVRS)										
tBRCAm	150	2 (1.3)	NE (NE,	NE)	65	2 (3.1)	NE (NE	, NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	1 (1.5)	NE (NE	, NE)	NC	NC	NC
Interaction p-value												NC
First line treatment out	come (eCR	PF)										
NED [PDS]	89	1 (1.1)	NE (NE,	NE)	47	3 (6.4)	NE (NE	, NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE	, NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE	, NE)	NC	NC	NC
PR	50	1 (2.0)	NE (NE,	NE)	34	0	NE (NE	, NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR	CA status	(eCRF)										
tBRCAm	147	2 (1.4)	NE (NE,	NE)	67	2 (3.0)	NE (NE	, NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	1 (1.6)	NE (NE	, NE)	NC	NC	NC
Interaction p-value												NC
Age group												
<65 years	185	2 (1.1)	NE (NE,	NE)	98	2 (2.0)	NE (NE	, NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	1 (3.0)	NE (NE	, NE)	NC	NC	NC
Interaction p-value												NC

Table 3.4.63 PAOLA1: Summary of subgroup analysis of AESI G1-2: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2		b		Placebo + be (N=1)					
Subgroup	C	Number (%) of patients with events	Median (95% (months	CI)	(Number (%) of patients with events	Median ti (95% CI (months))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)											
III	182	1 (0.5)	NE (N	E, NE)	89	3 (3.4)	NE (NE,	NE)	NC	NC	NC
IV	73	1 (1.4)	NE (N	E, NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Region											
Europe	245	2 (0.8)	NE (N	E, NE)	125	3 (2.4)	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (N	E, NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
ECOG performance status at	Baseli	ne									
(0) Normal activity	190	1 (0.5)	NE (N	E, NE)	100	2 (2.0)	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	1 (1.6)	NE (N	E, NE)	30	1 (3.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Baseline CA-125 value											
<=ULN	228	1 (0.4)	NE (N	E, NE)	117	3 (2.6)	NE (NE,	NE)	NC	NC	NC
>ULN	27	1 (3.7)	NE (N	E, NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Histological grade											
High grade	255	2 (0.8)	NE (N	E, NE)	131	3 (2.3)	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC
Cytoreductive surgery outco	ome										
No residue	166	1 (0.6)	NE (N	E, NE)	80	3 (3.8)	NE (NE,	NE)	NC	NC	NC
Residue	79	1 (1.3)	NE (N	E, NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value											NC

Table 3.4.63 PAOLA1: Summary of subgroup analysis of AESI G1-2: Wound healing complications Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		b		Placebo + be (N=1)		nab				
Subgroup	C	Number (%) of patients with events	Median (95% ((months	CI)	(Number (%) of patients with events	(95	n time % CI) ns) [a]	— Haz rat [k	io	% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	1 (0.7)	NE (N	E, NE)	78	3 (3.8)	NE (NE, N	E) N	C NC		NC
Interval	99	1 (1.0)	NE (N	E, NE)	45	0	NE (NE, N	E) N	C NC		NC
Interaction p-value												NC
Myriad tumour BRCA mutat:	ion statu	ıs										
tBRCAm	158	1 (0.6)	NE (N	E, NE)	77	2 (2.6)	NE (NE, N	E) N	C NC		NC
Non-tBRCAm	97	1 (1.0)	NE (N	E, NE)	54	1 (1.9)	NE (NE, N	E) N	C NC		NC
Interaction p-value												NC
Status somatic BRCA mutat	cions											
sBRCAm	25	0	NE (N	E, NE)	9	0	NE (NE, N	E) N	C NC		NC
gBRCAm	69	0	NE (N	E, NE)	36	0	NE (NE, N	E) N	C NC		NC
Non-BRCAm	43	0	NE (N	E, NE)	23	0	NE (NE, N	E) N	C NC		NC
Interaction p-value												NC

Table 3.4.64 PAOLA1: Summary of subgroup analysis of AESI G1-2: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		ımab	;		Placebo + be (N=13		ımab				
Subgroup		Number (%) of patients with events		an ti % CI; hs) [)	C	Number (%) of patients with events	(95	an tir 5% CI) :hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IV)	RS)											
NED [PDS]	92	10 (10.9)	NE (NE,	NE)	48	6 (12.5)	NE (NE,	NE)	0.82	0.30, 2.41	0.7030
NED/CR [IDS]	74	9 (12.2)	NE (NE,	NE)	38	3 (7.9)	NE (NE,	NE)	1.55	0.46, 7.00	0.4950
NED/CR [Chemo]	40	4 (10.0)	NE (NE,	NE)	20	1 (5.0)	NE (NE,	NE)	1.95	0.29, 38.08	0.5258
PR	49	5 (10.2)	NE (NE,	NE)	25	1 (4.0)	NE (NE,	NE)	2.70	0.44, 51.74	0.3152
Interaction p-value													0.6834
Screening laboratory tBRO	CA statu	s (IVRS)											
tBRCAm	150	17 (11.3)	NE (NE,	NE)	65	9 (13.8)	NE (NE,	NE)	0.78	0.36, 1.84	0.5616
non-tBRCAm	105	11 (10.5)	NE (NE,	NE)	66	2 (3.0)	NE (NE,	NE)	3.49	0.94, 22.57	0.0637
Interaction p-value													0.0639
First line treatment outo	come (eCl	RF)											
NED [PDS]	89	9 (10.1)	NE (NE,	NE)	47	6 (12.8)	NE (NE,	NE)	0.75	0.27, 2.23	0.5826
NED/CR [IDS]	74	9 (12.2)	NE (NE,	NE)	32	2 (6.3)	NE (NE,	NE)	1.99	0.51, 13.05	0.3457
NED/CR [Chemo]	39	4 (10.3)	NE (NE,	NE)	17	1 (5.9)	NE (NE,	NE)	1.75	0.26, 34.24	0.5971
PR	50	5 (10.0)	NE (NE,	NE)	34	2 (5.9)	NE (NE,	NE)	1.71	0.37, 11.95	0.5062
Interaction p-value													0.6706
Screening laboratory tBRO	CA statu	s (eCRF)											
tBRCAm	147	16 (10.9)	NE (NE,	NE)	67	9 (13.4)	NE (NE,	NE)	0.78	0.35, 1.83	0.5471
non-tBRCAm	108	12 (11.1)	NE (NE,	NE)	64	2 (3.1)	NE (NE,	NE)	3.62	0.99, 23.23	0.0529
Interaction p-value													0.0545
Age group													
<65 years	185	20 (10.8)	NE (NE,	NE)	98	8 (8.2)	NE (NE,	NE)	1.30	0.59, 3.13	0.5263
>=65 years	70	8 (11.4)	NE (NE,	NE)	33	3 (9.1)	NE (NE,	NE)	1.27	0.37, 5.80	0.7197
Interaction p-value													0.9783

Table 3.4.64 PAOLA1: Summary of subgroup analysis of AESI G1-2: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		mab	,		Placebo + be (N=1)		ab	;		
Subgroup		Number (%) of patients with events	Media (95° (mont)	% CI)			Number (%) of patients with events	(958	n time (CI) (s) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	22 (12.1)	NE (NE,	NE)	89	10 (11.2)	NE (NE, NE)	1.04	0.51, 2.30	0.9148
IV	73	6 (8.2)	NE (NE,	NE)	42	1 (2.4)	NE (NE, NE)	3.55	0.61, 67.08	0.1777
Interaction p-value												0.2360
Region												
Europe	245	28 (11.4)	NE (NE,	NE)	125	11 (8.8)	NE (NE, NE)	1.28	0.66, 2.69	0.4788
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Basel:	ine										
(0) Normal activity	190	24 (12.6)	NE (NE,	NE)	100	9 (9.0)	NE (NE, NE)	1.39	0.67, 3.15	0.3924
(1) Restricted activity	61	4 (6.6)	NE (NE,	NE)	30	1 (3.3)	NE (NE, NE)	1.96	0.29, 38.24	0.5230
Interaction p-value												0.7654
Baseline CA-125 value												
<=ULN	228	23 (10.1)	NE (NE,	NE)	117	11 (9.4)	NE (NE, NE)		0.52, 2.23	0.9003
>ULN	27	5 (18.5)	NE (NE,	NE)	14	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	28 (11.0)	NE (NE,	NE)	131	11 (8.4)	NE (NE, NE)	1.29	0.66, 2.71	0.4641
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	19 (11.4)	NE (NE,	NE)	80	8 (10.0)	NE (NE, NE)	1.12	0.51, 2.71	0.7905
Residue	79	8 (10.1)	NE (NE,	NE)	43	3 (7.0)	NE (NE, NE)	1.45	0.42, 6.62	0.5734
Interaction p-value												0.7422

Table 3.4.64 PAOLA1: Summary of subgroup analysis of AESI G1-2: Haemorrhage Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2			oevacizumab 131)		
Subgroup		Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]	2-sided p-value [b]
Timing of cytoreductive :	surgery						
Upfront	146	15 (10.3)	NE (NE, NE)	78 7 (9.0)	NE (NE, NE)	1.12 0.47, 2.92	0.8099
Interval	99	12 (12.1)	NE (NE, NE)	45 4 (8.9)	NE (NE, NE)	1.36 0.47, 4.87	0.5827
Interaction p-value							0.7850
Myriad tumour BRCA mutat:	ion stat	us					
tBRCAm	158	18 (11.4)	NE (NE, NE)	77 10 (13.0)	NE (NE, NE)	0.84 0.40, 1.89	0.6636
Non-tBRCAm	97	10 (10.3)	NE (NE, NE)	54 1 (1.9)	NE (NE, NE)	5.71 1.09,104.84	0.0368*
Interaction p-value							0.0426*
Status somatic BRCA muta	tions						
sBRCAm	25	2 (8.0)	NE (NE, NE)	9 0	NE (NE, NE)	NC NC	NC
gBRCAm	69	10 (14.5)	NE (NE, NE)	36 6 (16.7)	NE (NE, NE)	0.85 0.32, 2.51	0.7624
Non-BRCAm	43	5 (11.6)	NE (NE, NE)	23 1 (4.3)	NE (NE, NE)	2.66 0.43, 50.91	0.3239
Interaction p-value							0.3144

Table 3.4.65 PAOLA1: Summary of subgroup analysis of AESI G1-2: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + be (N=1)		nab	•			
Subgroup	c	Number (%) of patients with events	(95	an ti 5% CI .hs))	C	Number (%) of patients with events	Media (95 (mont)	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	RS)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(IVRS)											
tBRCAm	150	2 (1.3)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	RF)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	s (eCRF)											
tBRCAm	147	2 (1.4)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	0	NE (NE,	NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	2 (2.9)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

Table 3.4.65 PAOLA1: Summary of subgroup analysis of AESI G1-2: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		ımab			Placebo + be (N=1		nab				
Subgroup	c	Number (%) of patients with events	Media (95 (mont	% CI))	(Number (%) of patients with events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	2 (1.1)	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	2 (0.8)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	ne											
(0) Normal activity	190	1 (0.5)	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	1 (1.6)	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	2 (0.9)	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	2 (0.8)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	2 (1.2)	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

Table 3.4.65 PAOLA1: Summary of subgroup analysis of AESI G1-2: Arterial thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2				Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	Median ti (95% CI (months))	(Number (%) of patients with events	(95	an ti % CI) hs) [Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	0	NE (NE,	NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	2 (2.0)	NE (NE,	NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Myriad tumour BRCA mutat:	ion statu	ıs										
tBRCAm	158	2 (1.3)	NE (NE,	NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Status somatic BRCA muta	tions											
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE,	NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC

Table 3.4.66 PAOLA1: Summary of subgroup analysis of AESI G1-2: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		ımab			Placebo + bo (N=1		mab	•			
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI))	C	Number (%) of patients with events		an ti 5% CI :hs))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)											
NED [PDS]	92	3 (3.3)	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	4 (5.4)	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	1 (2.0)	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(IVRS)											
tBRCAm	150	5 (3.3)	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	3 (2.9)	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	2 (2.2)	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	6 (8.1)	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(eCRF)											
tBRCAm	147	5 (3.4)	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	3 (2.8)	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	4 (2.2)	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	4 (5.7)	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

Table 3.4.66 PAOLA1: Summary of subgroup analysis of AESI G1-2: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		mab]	Placebo + be (N=13		ab			
Subgroup	c	Number (%) of patients with events	(95%	n time k CI) ns) [a]		0	Number (%) f patients ith events	Mediar (95% (month	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	6 (3.3)	NE (NE, N	IE)	89	1 (1.1)	NE (NE, NE)	NC	NC	NC
IV	73	2 (2.7)	NE (NE, N	IE)	42	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	8 (3.3)	NE (NE, N	IE) 1	125	1 (0.8)	NE (NE, NE)	NC	NC	NC
Japan	10	0	NE (NE, N	IE)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	6 (3.2)	NE (NE, N	IE)]	100	1 (1.0)	NE (NE, NE)	NC	NC	NC
(1) Restricted activity	61	2 (3.3)	NE (NE, N	IE)	30	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	8 (3.5)	NE (NE, N	IE)]	117	1 (0.9)	NE (NE, NE)	NC	NC	NC
>ULN	27	0	NE (NE, N	IE)	14	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	8 (3.1)	NE (NE, N	IE) 1	131	1 (0.8)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	8 (4.8)	NE (NE, N	IE)	80	1 (1.3)	NE (NE, NE)	NC	NC	NC
Residue	79	0	NE (NE, N	IE)	43	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC

Table 3.4.66 PAOLA1: Summary of subgroup analysis of AESI G1-2: Venous thromboembolic events Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		Placebo + k (N=1			
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [k	2-sided b] p-value[b]
Timing of cytoreductive s	surgery						
Upfront	146	2 (1.4)	NE (NE, NE)	78 1 (1.3)	NE (NE, NE)	NC NC	NC
Interval	99	6 (6.1)	NE (NE, NE)	45 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Myriad tumour BRCA mutat:	ion statu	ıs					
tBRCAm	158	6 (3.8)	NE (NE, NE)	77 0	NE (NE, NE)	NC NC	NC
Non-tBRCAm	97	2 (2.1)	NE (NE, NE)	54 1 (1.9)	NE (NE, NE)	NC NC	NC
Interaction p-value							NC
Status somatic BRCA muta	tions						
sBRCAm	25	1 (4.0)	NE (NE, NE)	9 0	NE (NE, NE)	NC NC	NC
gBRCAm	69	3 (4.3)	NE (NE, NE)	36 0	NE (NE, NE)	NC NC	NC
Non-BRCAm	43	2 (4.7)	NE (NE, NE)	23 0	NE (NE, NE)	NC NC	NC
Interaction p-value							NC

Table 3.4.67 PAOLA1: Summary of subgroup analysis of AESI G1-2: Posterior Reversible Encephalopathy Syndrome (PRES) Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		ımab			Placebo + bo (N=1		mab				
Subgroup	C	Number (%) of patients with events		an ti % CI :hs))	C	Number (%) of patients with events	(95	an ti 5% CI) ths) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	.S)											
NED [PDS]	92	0	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(IVRS)											
tBRCAm	150	0	NE (NE,	NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	1 (1.5)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	0	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(eCRF)											
tBRCAm	147	0	NE (NE,	NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	1 (1.6)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	0	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

Table 3.4.67 PAOLA1: Summary of subgroup analysis of AESI G1-2: Posterior Reversible Encephalopathy Syndrome (PRES) Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + be (N=1)		nab			
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI ths))	C	Number (%) of patients with events	(95	n time % CI) ns) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	0	NE (NE,	NE)	89	1 (1.1)	NE (NE, NE) NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	0	NE (NE,	NE)	125	1 (0.8)	NE (NE, NE) NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	0	NE (NE,	NE)	100	0	NE (NE, NE) NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	1 (3.3)	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	0	NE (NE,	NE)	117	1 (0.9)	NE (NE, NE) NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	0	NE (NE,	NE)	131	1 (0.8)	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	0	NE (NE,	NE)	80	1 (1.3)	NE (NE, NE) NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC

Table 3.4.67 PAOLA1: Summary of subgroup analysis of AESI G1-2: Posterior Reversible Encephalopathy Syndrome (PRES) Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		ımab			Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]		C	Number (%) f patients ith events		an ti 5% CI :hs))	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]	
Timing of cytoreductive s	surgery												
Upfront	146	0	NE (NE,	NE)	78	1 (1.3)	NE (NE,	NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Myriad tumour BRCA mutat:	ion statu	ıs											
tBRCAm	158	0	NE (NE,	NE)	77	0	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	1 (1.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Status somatic BRCA mutat	cions												
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE,	NE)	36	0	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

Table 3.4.68 PAOLA1: Summary of subgroup analysis of AESI G1-2: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2		umab			Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI :hs))	C	Number (%) of patients with events	Media (95 (mont	% CI))	Hazard ratio [b] 95% CI [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	S)											
NED [PDS]	92	0	NE (NE,	NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	1 (2.6)	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(IVRS)											
tBRCAm	150	0	NE (NE,	NE)	65	1 (1.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	F)											
NED [PDS]	89	0	NE (NE,	NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	1 (2.9)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR	CA status	(eCRF)											
tBRCAm	147	0	NE (NE,	NE)	67	1 (1.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	0	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

Table 3.4.68 PAOLA1: Summary of subgroup analysis of AESI G1-2: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		umab			Placebo + be (N=1		nab			
Subgroup	C	Number (%) of patients with events	(95	an ti 5% CI ths))	C	Number (%) of patients with events	(95	n time k CI) ns) [a]	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)												
III	182	0	NE (NE,	NE)	89	1 (1.1)	NE (NE, NE) NC	NC	NC
IV	73	0	NE (NE,	NE)	42	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	0	NE (NE,	NE)	125	1 (0.8)	NE (NE, NE) NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	0	NE (NE,	NE)	100	1 (1.0)	NE (NE, NE) NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	0	NE (NE,	NE)	117	0	NE (NE, NE) NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	1 (7.1)	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	0	NE (NE,	NE)	131	1 (0.8)	NE (NE, NE) NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	0	NE (NE,	NE)	80	0	NE (NE, NE) NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	1 (2.3)	NE (NE, NE) NC	NC	NC
Interaction p-value												NC

Table 3.4.68 PAOLA1: Summary of subgroup analysis of AESI G1-2: Non-GI fistula or abscess Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2!		ımab			Placebo + be (N=1		ımab				
Subgroup	C	Number (%) of patients with events	(95	an time 5% CI) ths) [a]		C	Number (%) of patients with events	(95	an ti 5% CI; ths) [)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery												
Upfront	146	0	NE (NE,	NE)	78	0	NE (NE,	NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	1 (2.2)	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Myriad tumour BRCA mutat:	ion statu	ıs											
tBRCAm	158	0	NE (NE,	NE)	77	1 (1.3)	NE (NE,	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Status somatic BRCA muta	tions												
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE,	NE)	NC	NC	NC
gBRCAm	69	0	NE (NE,	NE)	36	1 (2.8)	NE (NE,	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

Table 3.4.69 PAOLA1: Summary of subgroup analysis of AESI G1-2: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2		ımab	;		Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI)	C	Number (%) of patients with events	Medi (95 (mont	% CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	LS)											
NED [PDS]	92	2 (2.2)	NE (NE,	NE)	48	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	1 (2.5)	NE (NE,	NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	0	NE (NE,	NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(IVRS)											
tBRCAm	150	3 (2.0)	NE (NE,	NE)	65	1 (1.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	0	NE (NE,	NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
First line treatment out	come (eCR	RF)											
NED [PDS]	89	2 (2.2)	NE (NE,	NE)	47	1 (2.1)	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	0	NE (NE,	NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	1 (2.6)	NE (NE,	NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE,	NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Screening laboratory tBR0	CA status	(eCRF)											
tBRCAm	147	3 (2.0)	NE (NE,	NE)	67	1 (1.5)	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	0	NE (NE,	NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Age group													
<65 years	185	3 (1.6)	NE (NE,	NE)	98	1 (1.0)	NE (NE,	NE)	NC	NC	NC
>=65 years	70	0	NE (NE,	NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

Table 3.4.69 PAOLA1: Summary of subgroup analysis of AESI G1-2: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	ı	Olaparib + b (N=2!		ımab			Placebo + be (N=13		ıb	•		
Subgroup	Number (%) of patients n with events		Median time (95% CI) (months) [a]		C	Number (%) of patients with events	Median (95% (month)	CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]	
FIGO Stage (Disease state)												
III	182	2 (1.1)	NE (NE,	NE)	89	1 (1.1)	NE (NE, NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE,	NE)	42	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Region												
Europe	245	3 (1.2)	NE (NE,	NE)	125	1 (0.8)	NE (NE, NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
ECOG performance status at	Baseli	.ne										
(0) Normal activity	190	2 (1.1)	NE (NE,	NE)	100	1 (1.0)	NE (NE, NE)	NC	NC	NC
(1) Restricted activity	61	1 (1.6)	NE (NE,	NE)	30	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Baseline CA-125 value												
<=ULN	228	3 (1.3)	NE (NE,	NE)	117	1 (0.9)	NE (NE, NE)	NC	NC	NC
>ULN	27	0	NE (NE,	NE)	14	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Histological grade												
High grade	255	3 (1.2)	NE (NE,	NE)	131	1 (0.8)	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC
Cytoreductive surgery outco	ome											
No residue	166	2 (1.2)	NE (NE,	NE)	80	1 (1.3)	NE (NE, NE)	NC	NC	NC
Residue	79	1 (1.3)	NE (NE,	NE)	43	0	NE (NE, NE)	NC	NC	NC
Interaction p-value												NC

Table 3.4.69 PAOLA1: Summary of subgroup analysis of AESI G1-2: Secondary cancer Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2				Placebo + be (N=1)		mab				
Subgroup	C	Number (%) of patients with events	Median tin (95% CI) (months) [95% CI)		Number (%) of patients with events	(95	an time % CI) hs) [a]		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
Timing of cytoreductive s	surgery											
Upfront	146	3 (2.1)	NE (NE,	NE)	78	1 (1.3)	NE (NE, 1	NE)	NC	NC	NC
Interval	99	0	NE (NE,	NE)	45	0	NE (NE, 1	NE)	NC	NC	NC
Interaction p-value												NC
Myriad tumour BRCA mutat:	ion statu	ıs										
tBRCAm	158	3 (1.9)	NE (NE,	NE)	77	1 (1.3)	NE (NE, 1	NE)	NC	NC	NC
Non-tBRCAm	97	0	NE (NE,	NE)	54	0	NE (NE, 1	NE)	NC	NC	NC
Interaction p-value												NC
Status somatic BRCA muta	tions											
sBRCAm	25	0	NE (NE,	NE)	9	0	NE (NE, 1	NE)	NC	NC	NC
gBRCAm	69	1 (1.4)	NE (NE,	NE)	36	0	NE (NE, 1	NE)	NC	NC	NC
Non-BRCAm	43	0	NE (NE,	NE)	23	0	NE (NE, 1	NE)	NC	NC	NC
Interaction p-value												NC

Table 3.4.70 PAOLA1: Summary of subgroup analysis of AESI G1-2: Pneumonitis Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	(Olaparib + b (N=2)		ab		Placebo + bo (N=1		mab				
Subgroup	C	Number (%) of patients with events	Mediar (95% (month	CI)		Number (%) of patients with events	Media (95 (mont	% CI)	Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
First line treatment out	come (IVR	S)										
NED [PDS]	92	0	NE (NE, NE)	48	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	2 (2.7)	NE (NE, NE)	38	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	40	0	NE (NE, NE)	20	0	NE (NE,	NE)	NC	NC	NC
PR	49	1 (2.0)	NE (NE, NE)	25	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR0	CA status	(IVRS)										
tBRCAm	150	1 (0.7)	NE (NE, NE)	65	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	105	2 (1.9)	NE (NE, NE)	66	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
First line treatment out	come (eCR	F)										
NED [PDS]	89	0	NE (NE, NE)	47	0	NE (NE,	NE)	NC	NC	NC
NED/CR [IDS]	74	3 (4.1)	NE (NE, NE)	32	0	NE (NE,	NE)	NC	NC	NC
NED/CR [Chemo]	39	0	NE (NE, NE)	17	0	NE (NE,	NE)	NC	NC	NC
PR	50	0	NE (NE, NE)	34	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Screening laboratory tBR0	CA status	(eCRF)										
tBRCAm	147	1 (0.7)	NE (NE, NE)	67	0	NE (NE,	NE)	NC	NC	NC
non-tBRCAm	108	2 (1.9)	NE (NE, NE)	64	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC
Age group												
<65 years	185	1 (0.5)	NE (NE, NE)	98	0	NE (NE,	NE)	NC	NC	NC
>=65 years	70	2 (2.9)	NE (NE, NE)	33	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value												NC

Table 3.4.70 PAOLA1: Summary of subgroup analysis of AESI G1-2: Pneumonitis Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

		Olaparib + b (N=2!		ımab	•		Placebo + be (N=1)		nab	•			
Subgroup	C	Number (%) of patients with events	Media (95 (mont	% CI)		(Number (%) of patients with events	Media (95 (mont	% CI)		Hazard ratio [b]	95% CI [b]	2-sided p-value [b]
FIGO Stage (Disease state)													
III	182	2 (1.1)	NE (NE,	NE)	89	0	NE (NE,	NE)	NC	NC	NC
IV	73	1 (1.4)	NE (NE,	NE)	42	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Region													
Europe	245	3 (1.2)	NE (NE,	NE)	125	0	NE (NE,	NE)	NC	NC	NC
Japan	10	0	NE (NE,	NE)	6	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
ECOG performance status at	Baseli	.ne											
(0) Normal activity	190	3 (1.6)	NE (NE,	NE)	100	0	NE (NE,	NE)	NC	NC	NC
(1) Restricted activity	61	0	NE (NE,	NE)	30	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Baseline CA-125 value													
<=ULN	228	2 (0.9)	NE (NE,	NE)	117	0	NE (NE,	NE)	NC	NC	NC
>ULN	27	1 (3.7)	NE (NE,	NE)	14	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Histological grade													
High grade	255	3 (1.2)	NE (NE,	NE)	131	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC
Cytoreductive surgery outco	ome												
No residue	166	3 (1.8)	NE (NE,	NE)	80	0	NE (NE,	NE)	NC	NC	NC
Residue	79	0	NE (NE,	NE)	43	0	NE (NE,	NE)	NC	NC	NC
Interaction p-value													NC

Table 3.4.70 PAOLA1: Summary of subgroup analysis of AESI G1-2: Pneumonitis Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

	1	Olaparib + b (N=2!		Placebo + b (N=1				
Subgroup	C	Number (%) of patients with events	Median time (95% CI) (months) [a]	Number (%) of patients n with events	Median time (95% CI) (months) [a]	Hazard ratio [b] 95% CI [b]		2-sided p-value [b]
Timing of cytoreductive s	surgery							
Upfront	146	0	NE (NE, NE)	78 0	NE (NE, NE)	NC I	NC	NC
Interval	99	3 (3.0)	NE (NE, NE)	45 0	NE (NE, NE)	NC I	NC	NC
Interaction p-value								NC
Myriad tumour BRCA mutati	ion statu	ıs						
tBRCAm	158	2 (1.3)	NE (NE, NE)	77 0	NE (NE, NE)	NC I	NC	NC
Non-tBRCAm	97	1 (1.0)	NE (NE, NE)	54 0	NE (NE, NE)	NC I	NC	NC
Interaction p-value								NC
Status somatic BRCA mutat	cions							
sBRCAm	25	1 (4.0)	NE (NE, NE)	9 0	NE (NE, NE)	NC I	NC	NC
gBRCAm	69	0	NE (NE, NE)	36 0	NE (NE, NE)	NC I	NC	NC
Non-BRCAm	43	1 (2.3)	NE (NE, NE)	23 0	NE (NE, NE)	NC I	NC	NC
Interaction p-value								NC

[[]a] Median time to event based on the Kaplan Meier estimate. NE = not estimable as median (or 95% CLs) not reached.

[b] HR, 95% profile likelihood CIs and likelihood-ratio test p-values estimated from Cox PH model. Efron method for handling ties. If >=10 patients for all subgroup levels, >=10 events for 1 subgroup level, >0 events for all subgroup levels/treatments, results from model including treatment, subgroup, treatment by subgroup interaction. For subgroups with >2 levels, levels with insufficient data were dropped and interaction model re-fitted. If interaction model conditions not met but 1 level had >=10 events, and >0 events for both treatments, results from model for 1 level including treatment only. * p-value <0.05. HR <1 favours olaparib. root/cdar/d081/_iemt/ar/iemt_payer_paola_germany_s3/tlf/prod/program/ttesubae.sas ettesubaeacr 11AUG2022:11:30 kpzx329

Figure 3.5.1 PAOLA1: Kaplan-Meier plot of AESI: Nausea for FIGO Stage (Disease state) = III Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

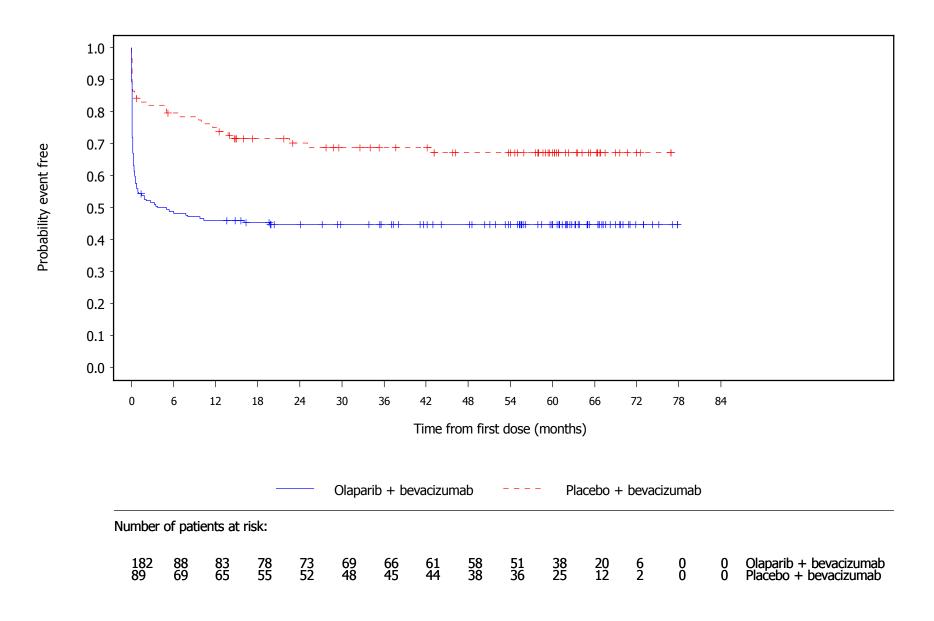


Figure 3.5.2 PAOLA1: Kaplan-Meier plot of AESI: Nausea for FIGO Stage (Disease state) = IV Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

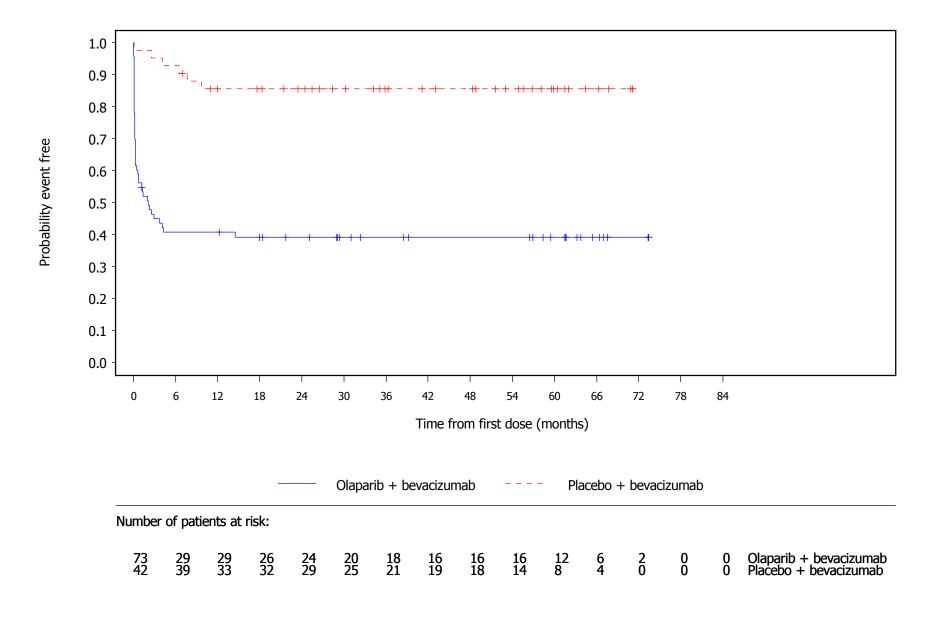


Figure 3.5.3 PAOLA1: Kaplan-Meier plot of AESI: Nausea for ECOG performance status at Baseline = (0) Normal activity Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

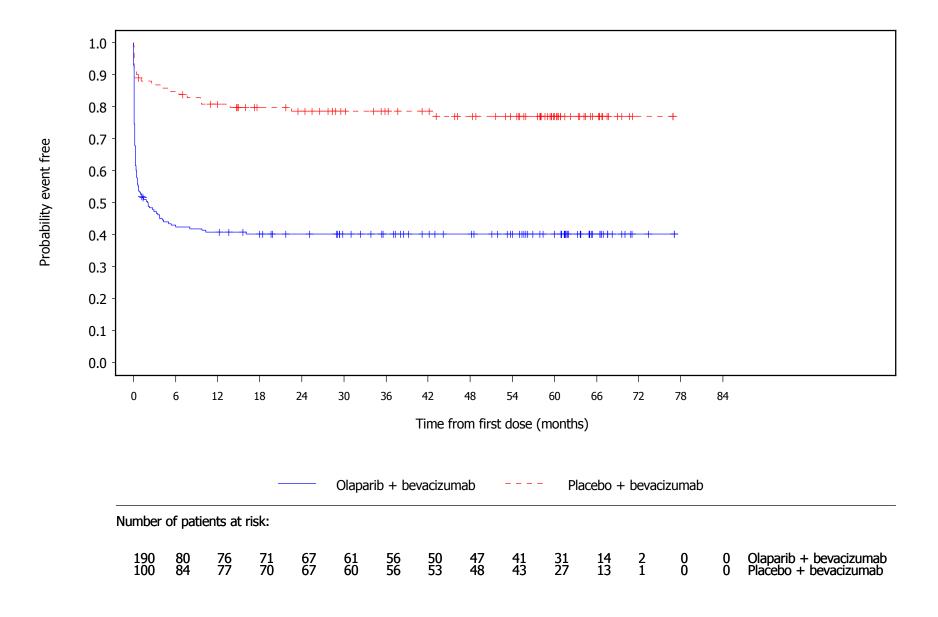


Figure 3.5.4 PAOLA1: Kaplan-Meier plot of AESI: Nausea for ECOG performance status at Baseline = (1) Restricted activity Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

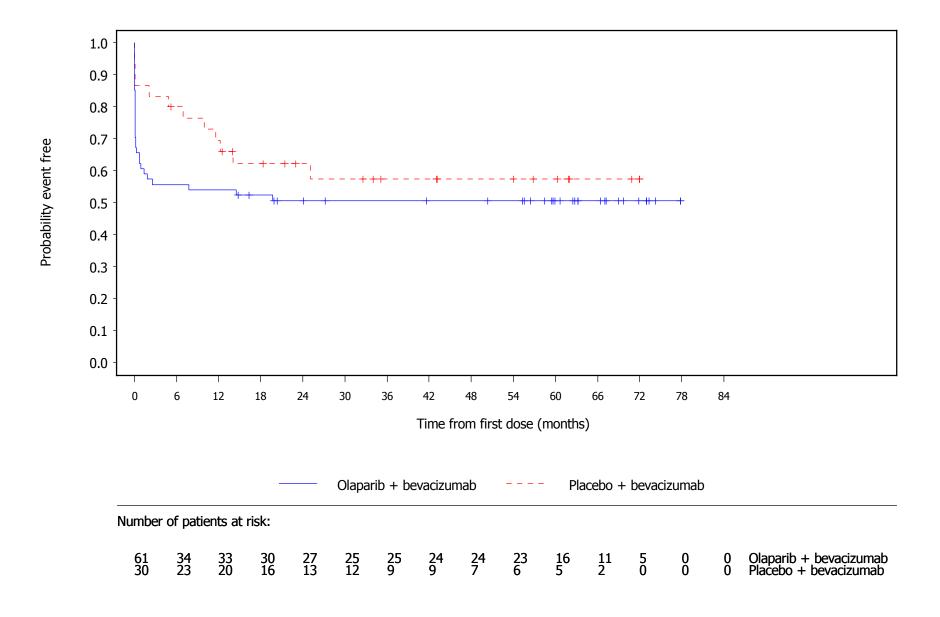


Figure 3.5.5 PAOLA1: Kaplan-Meier plot of AESI: Hypertension for Region = Europe Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

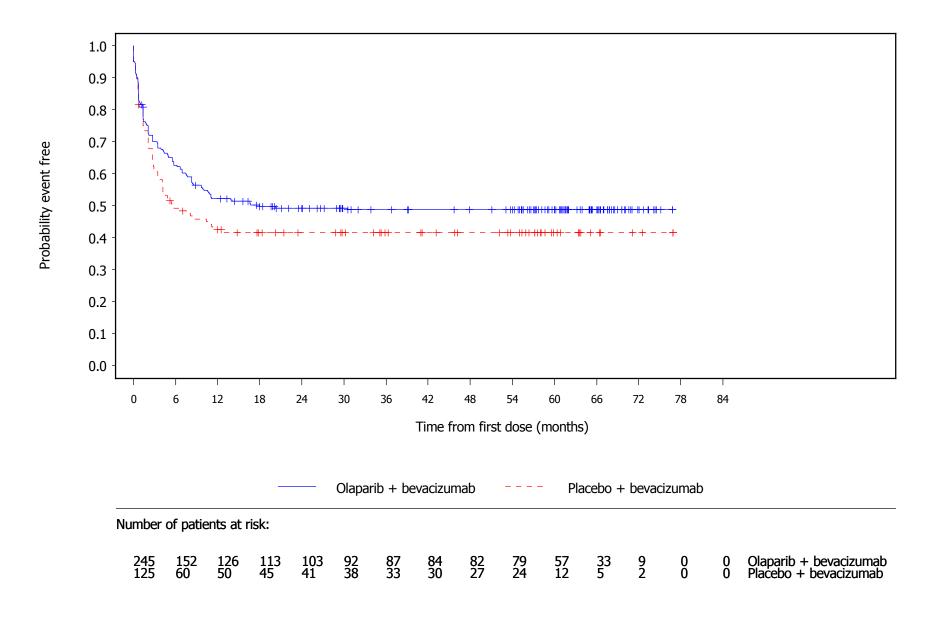


Figure 3.5.6 PAOLA1: Kaplan-Meier plot of AESI: Hypertension for Region = Japan Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

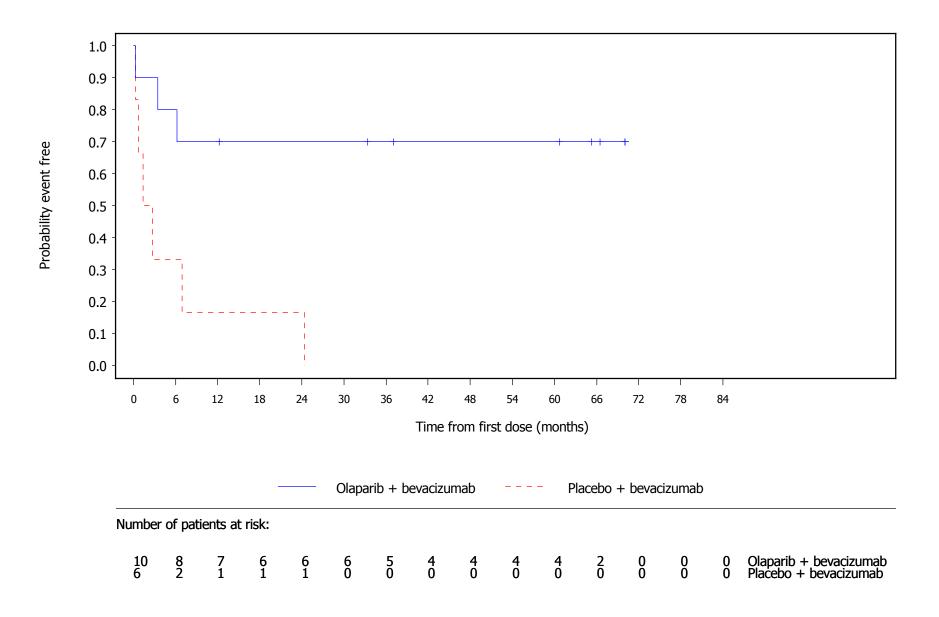


Figure 3.5.7 PAOLA1: Kaplan-Meier plot of AESI: Proteinuria for Timing of cytoreductive surgery = Upfront Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

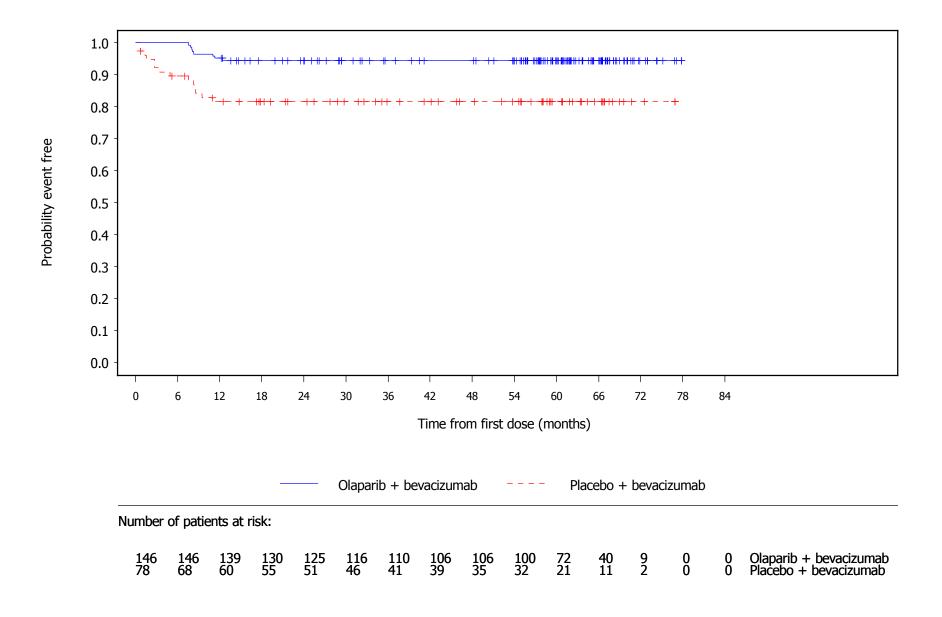


Figure 3.5.8 PAOLA1: Kaplan-Meier plot of AESI: Proteinuria for Timing of cytoreductive surgery = Interval Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

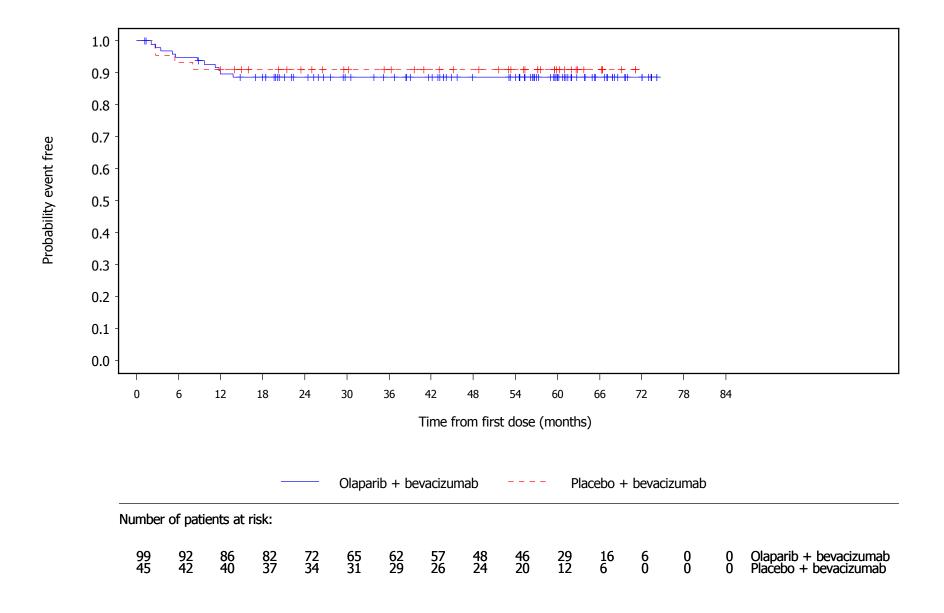


Figure 3.5.9 PAOLA1: Kaplan-Meier plot of AESI G1-2: Nausea for FIGO Stage (Disease state) = III Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

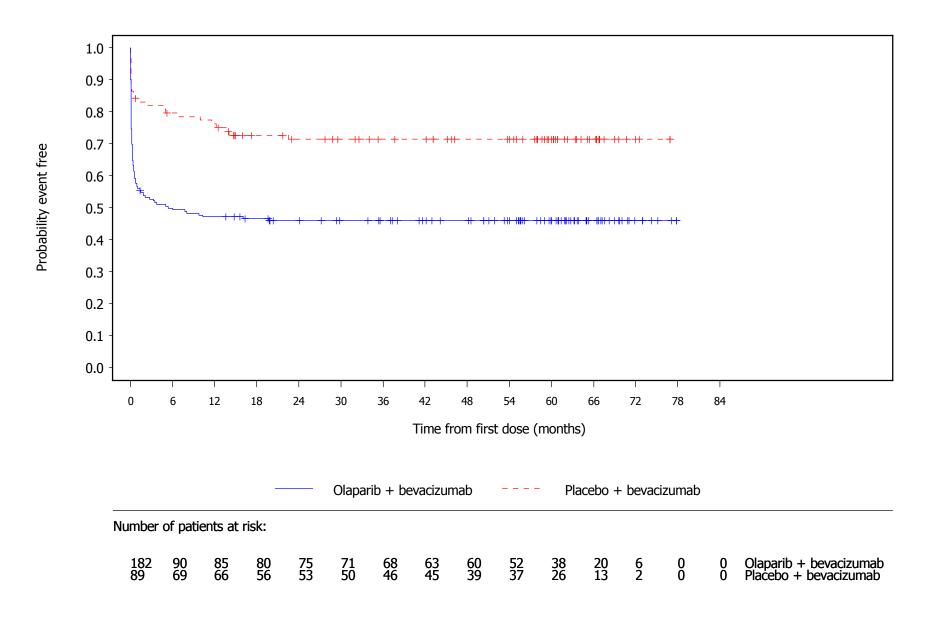


Figure 3.5.10 PAOLA1: Kaplan-Meier plot of AESI G1-2: Nausea for FIGO Stage (Disease state) = IV Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

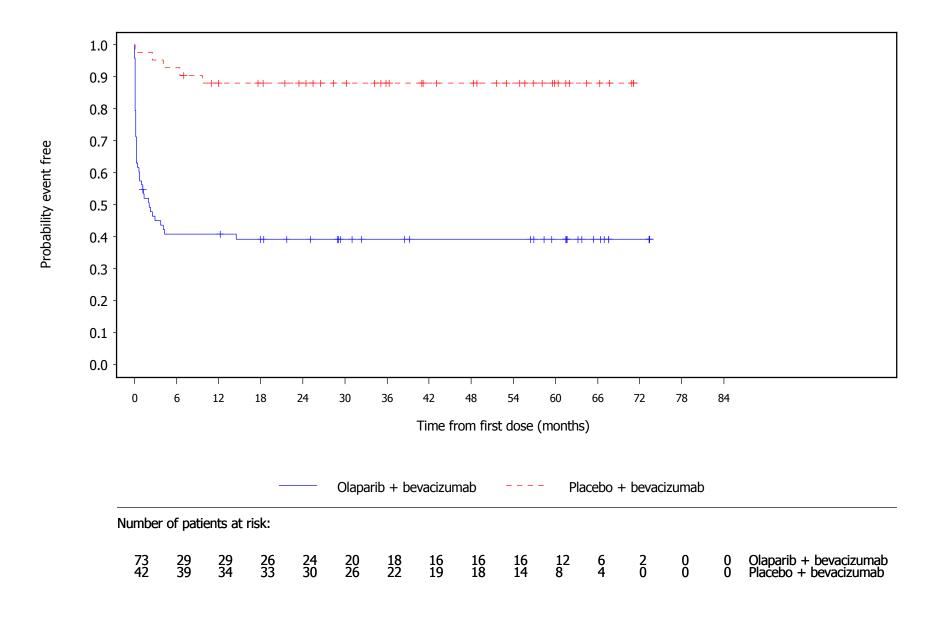


Figure 3.5.11 PAOLA1: Kaplan-Meier plot of AESI G1-2: Nausea for ECOG performance status at Baseline = (0) Normal activity Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

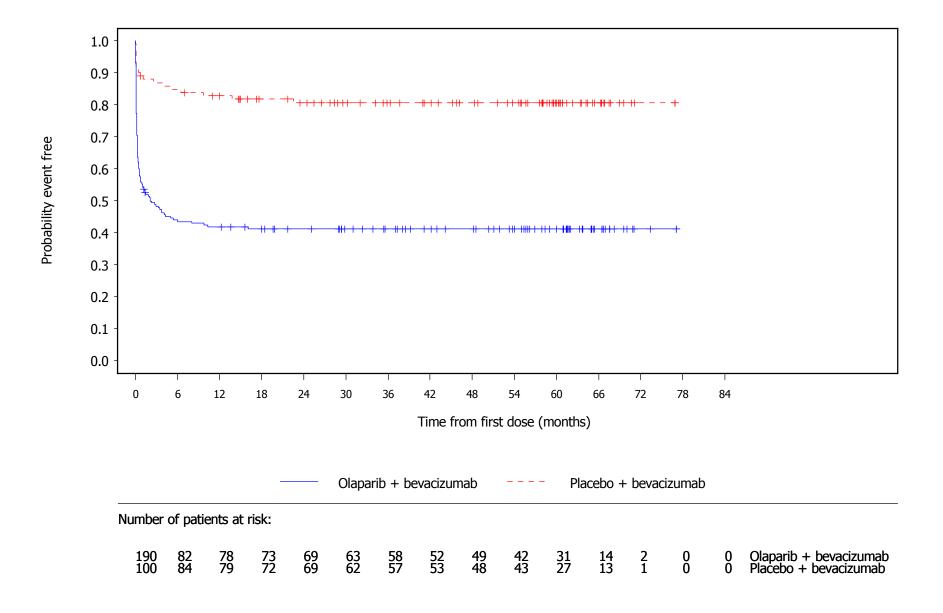


Figure 3.5.12 PAOLA1: Kaplan-Meier plot of AESI G1-2: Nausea for ECOG performance status at Baseline = (1) Restricted activity Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

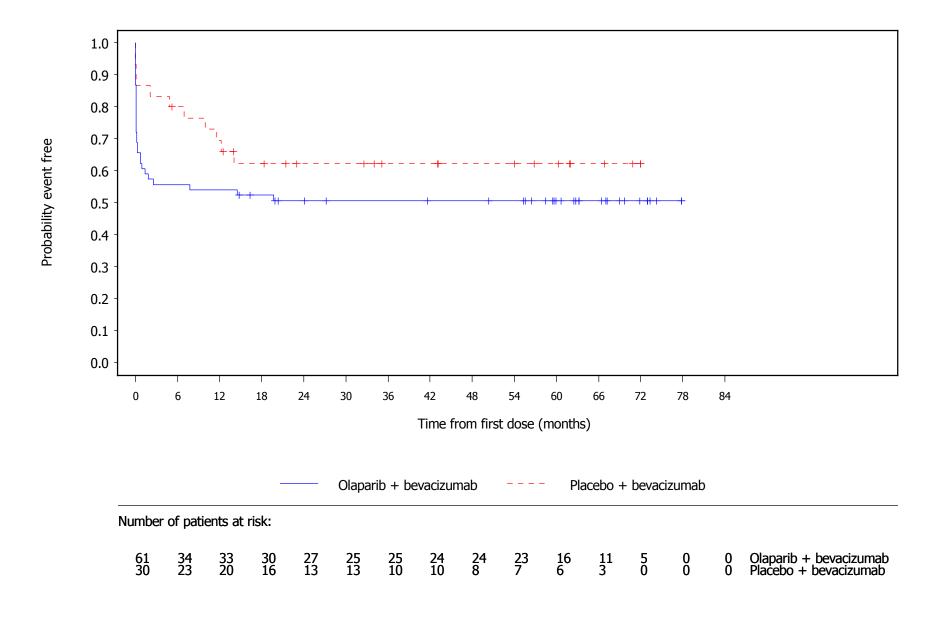


Figure 3.5.13 PAOLA1: Kaplan-Meier plot of AESI G1-2: Hypertension for Region = Europe Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

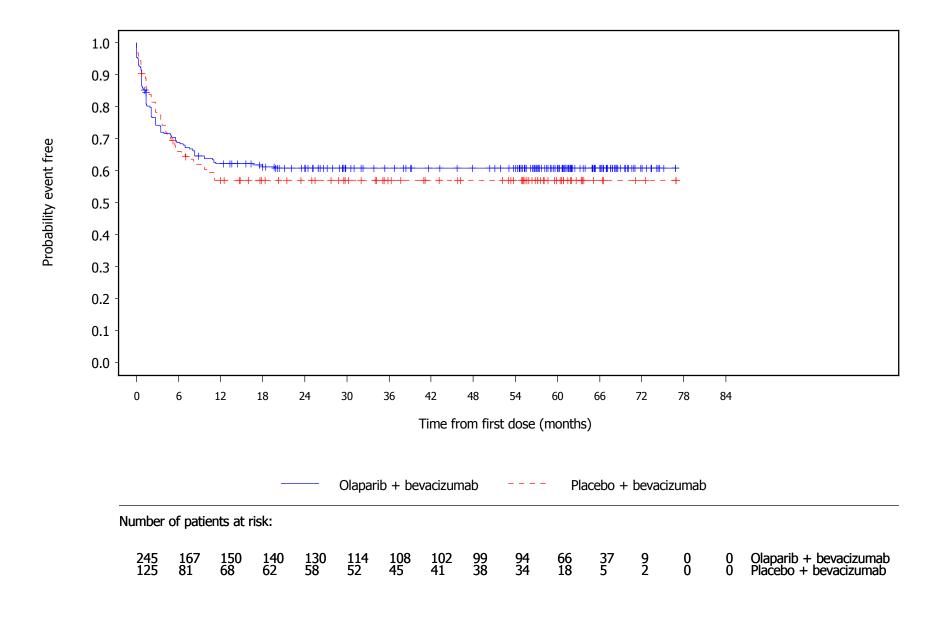


Figure 3.5.14 PAOLA1: Kaplan-Meier plot of AESI G1-2: Hypertension for Region = Japan Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

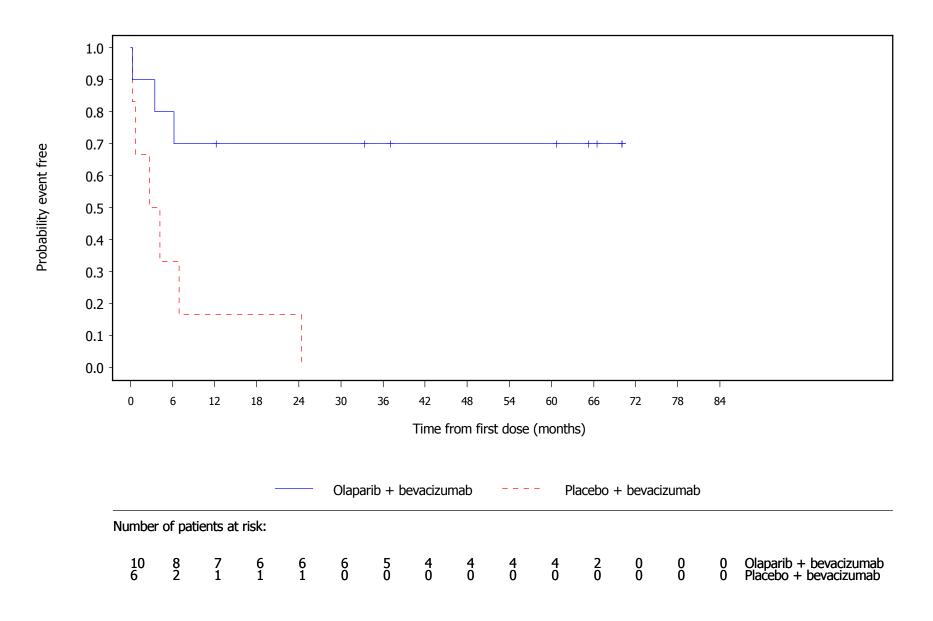


Figure 3.5.15 PAOLA1: Kaplan-Meier plot of AESI G1-2: Proteinuria for Timing of cytoreductive surgery = Upfront Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

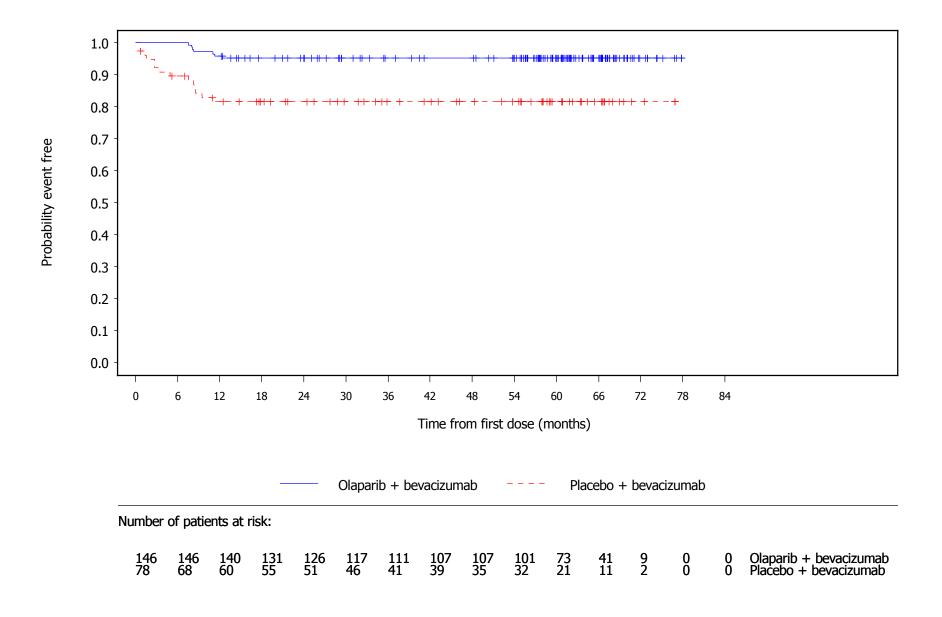


Figure 3.5.16 PAOLA1: Kaplan-Meier plot of AESI G1-2: Proteinuria for Timing of cytoreductive surgery = Interval Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

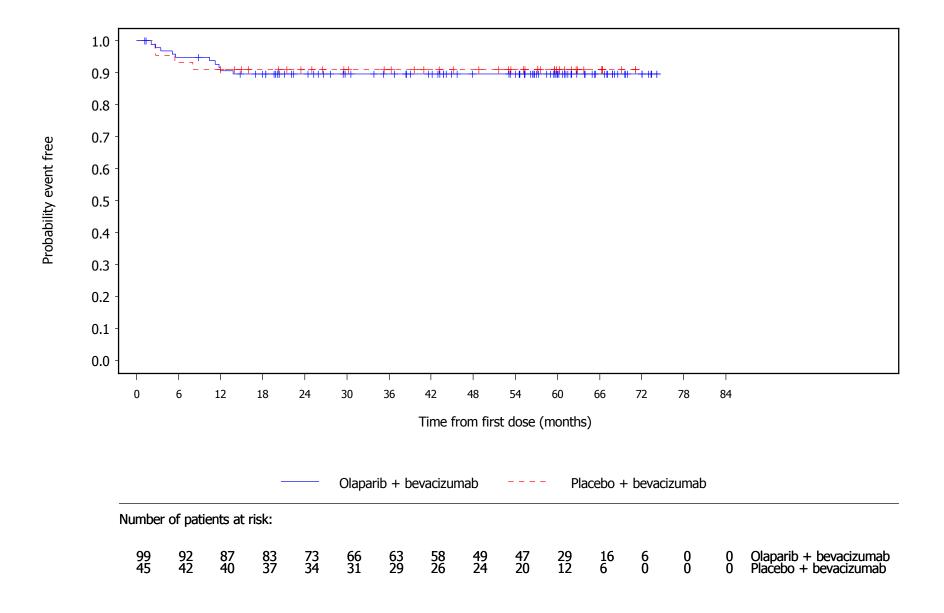


Figure 3.5.17 PAOLA1: Kaplan-Meier plot of AESI G1-2: Haemorrhage for Myriad tumour BRCA mutation status = tBRCAm Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

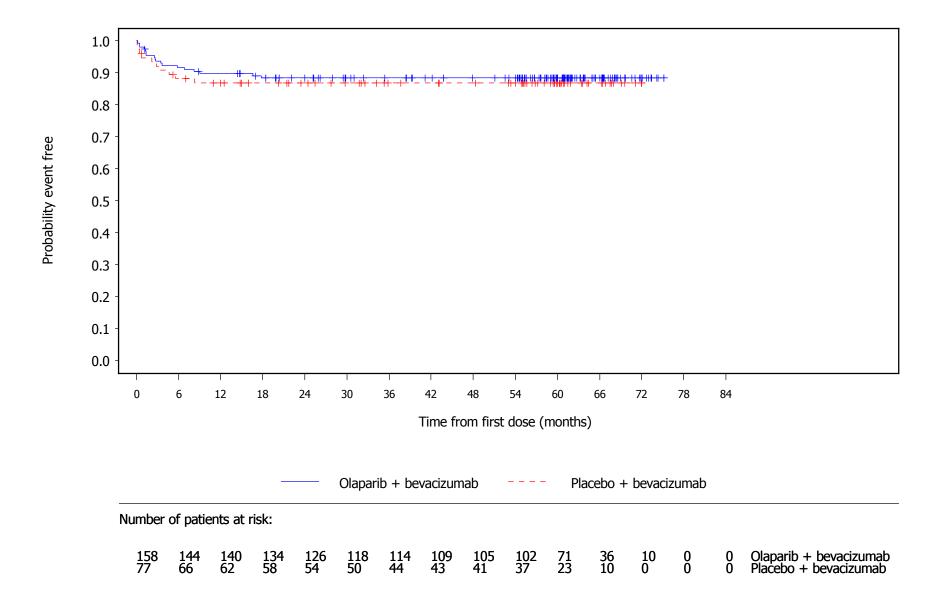


Figure 3.5.18 PAOLA1: Kaplan-Meier plot of AESI G1-2: Haemorrhage for Myriad tumour BRCA mutation status = Non-tBRCAm Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

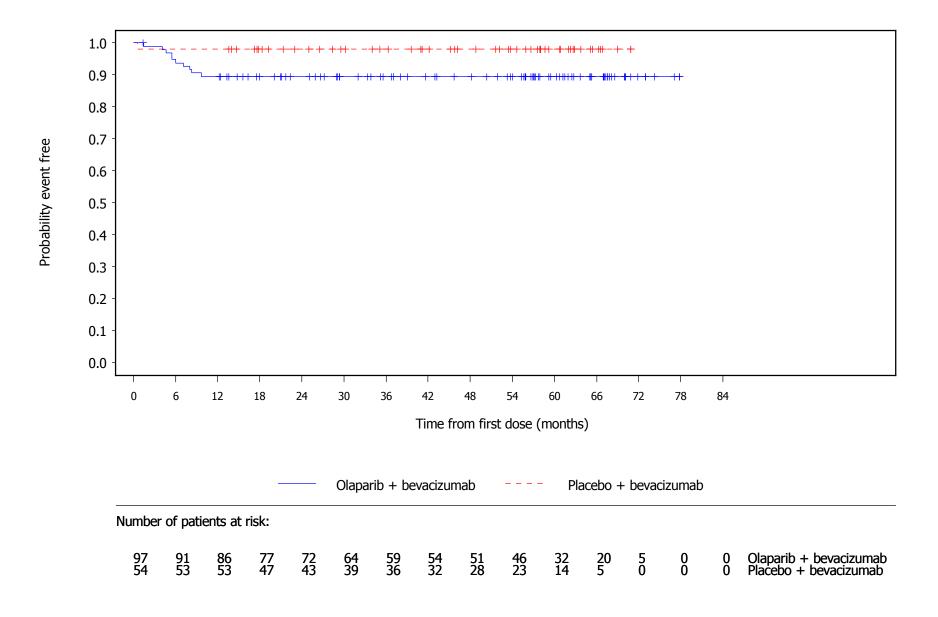


Table 3.6.1 PAOLA1: Summary of analysis of adverse events of special interest (total, and by grouped or preferred term)

(odds ratio, relative risk and risk difference)

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

									ŗ	Treatme	nt effe	ect			
	beva (N	parib + cizumab J=255)	bev	lacebo + vacizumab (N=131)		Odd	s Ratio			Relati	ive Ris	k	Risk D	ifferen	ce
) q w	Jumber %) of patients with events [a]	n	Number (%) of patients with events [a]		Estima (95% C		2- sided p- value		Estimat (95% CI		2- sided p- value	Estima (95% C		2- sided p- value
AESI: Anaemia [b][e][h]	255 1	.02(40.0)	131	12(9.2)	6.61(3.60,	13.19)	<0.0001 *	4.37(2.62,	8.11)	<0.0001 *	0.31(0.23,	0.38)	<0.0001 *
AESI: Neutropenia [b][e][h]	255	53(20.8)	131	22(16.8)	1.30(0.76,	2.29)	0.3437	1.24(0.80,	1.99)	0.3437	0.04(-0.04,	0.12)	0.3437
AESI: Thrombocytopeni a [b][e][h]	255	18(7.1)	131	7(5.3)	1.35(0.57,	3.54)	0.5105	1.32(0.59,	3.32)	0.5105	0.02(-0.04,	0.07)	0.5105
AESI: Nausea [b][e][h]	255 1	44(56.5)	131	34(26.0)	3.70(2.35,	5.94)	<0.0001 *	2.18(1.63,	3.03)	<0.0001 *	0.31(0.21,	0.40)	<0.0001
AESI: Vomiting [b][e][h]	255	55(21.6)	131	18(13.7)	1.73(0.98,	3.16)	0.0575	1.57(0.99,	2.64)	0.0575	0.08(-0.00,	0.15)	0.0575
AESI: Fatigue and Asthenia [b][e][h]	255 1	42(55.7)	131	47(35.9)	2.25(1.46,	3.48)	0.0002	1.55(1.22,	2.03)	0.0002	0.20(0.09,	0.30)	0.0002
AESI: Hypertension [b][e][h]	255 1	.27(49.8)	131	78(59.5)	0.67(0.44,	1.03)	0.0688	0.84(0.70,	1.01)	0.0688	-0.10(-0.20,	0.01)	0.0688
AESI: Proteinuria [b][e][h]	255	20(7.8)	131	19(14.5)	0.50(0.26,	0.98)	0.0445	0.54(0.30,	0.98)	0.0445	-0.07(-0.14,	-0.00)	0.0445

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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Table 3.6.1 PAOLA1: Summary of analysis of adverse events of special interest (total, and by grouped or preferred term)

(odds ratio, relative risk and risk difference)

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

									Т	reatme	nt effe	ect			
	bev	aparib + racizumab N=255)	ber	lacebo + vacizumab (N=131)		Odds	Ratio			Relati	ve Ris	k	Risk	Differer	ice
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]	(9	stimat 95% CI	.)	2- sided p- value		Estimat (95% CI	.)	2- sided p- value	Estin (95%	CI)	2- sided p- value
AESI: GI perforations, abscesses and fistulae [c][g][i]	255	3(1.2)	131	0	3.65(0.35,4	91.54)	0.3214		N	C			NC	
AESI: Wound healing complications [b][e][h]	255	2(0.8)	131	3(2.3)	0.34(0.04,	2.06)	0.2314	0.34(0.05,	2.04)	0.2314	-0.02(-0.05	, 0.01)	0.2314
AESI: Haemorrhage [b][e][h]	255	30(11.8)	131	12(9.2)	1.32(0.67,	2.77)	0.4307	1.28(0.70,	2.53)	0.4307	0.03(-0.04	, 0.09)	0.4307
AESI: Arterial thromboembolic events [b][e][h]	255	3(1.2)	131	4(3.1)	0.38(0.07,	1.74)	0.2055	0.39(0.08,	1.72)	0.2055	-0.02(-0.06	, 0.01)	0.2055
AESI: Venous thromboembolic events [b][e][h]	255	10(3.9)	131	1(0.8)	5.31(0	.9996,	97.88)	0.0501	5.14(0).9996,	93.73	0.0501	0.03(-0.00	, 0.06)	0.0501
AESI: Posterior Reversible Encephalopathy Syndrome (PRES) [c][g][i]	255	0	131	1(0.8)	0.17(0.00,	3.21)	0.2352		N	rC			NC	

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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Table 3.6.1 PAOLA1: Summary of analysis of adverse events of special interest (total, and by grouped or preferred term)

(odds ratio, relative risk and risk difference)

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

									Tre	eatment e	ffect			
	ber	aparib + vacizumab (N=255)		lacebo + vacizumab (N=131)		Odds	Ratio		R	elative 1	Risk	Risk D	ifferen	ıce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with		Estimat		2- sided p- value		timate 5% CI)	2- sided p- value	Estima (95% C		2- sided p- value
AESI: Congestive heart failure [d][g][i]	255	0	131	0		1	IÇ.		,	NC		· · · · · · · · · · · · · · · · · · ·	NC	
AESI: Non-GI fistula or abscess [c][g][i]	255	0	131	2(1.5)	0.10	0.00,	1.26)	0.0769		NC		1	NC	
AESI: MDS/AML [b][e][h]	255	4(1.6)	131	3(2.3)	0.68	0.15,	3.49)	0.6211	0.68(0	.15, 3.4	3) 0.6211	-0.01(-0.04,	0.02)	0.6211
AESI: Myelodysplastic syndrome and Acute myeloid leukaemia [b][e][h]	255	4(1.6)	131	4(3.1)	0.51	0.12,	2.17)	0.3452	0.51(0	.12, 2.3	4) 0.3452	-0.01(-0.06,	0.02)	0.3452
AESI: Secondary cancer [b][e][h]	255	15(5.9)	131	4(3.1)	1.98	0.70,	7.07)	0.2062	1.93(0	.72, 6.6	5) 0.2062	0.03(-0.02,	0.07)	0.2062
AESI: Pneumonitis [c][g][i]	255	3(1.2)	131	0	3.65	0.35,4	191.54)	0.3214		NC]	NC	

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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Table 3.6.2 PAOLA1: Summary of analysis of serious adverse events of special interest (odds ratio, relative risk and risk difference)

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

								Treatment effe	ect			
	be	aparib + vacizumab (N=255)	be ⁻	lacebo + vacizumab (N=131)	00	dds Ratio		Relative Ris	k	Risk Di	fferen	ce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]	Esti (95%		2- sided p- value	Estimate (95% CI)	2- sided p- value	Estimat (95% CI		2- sided p- value
Serious AESI: Anaemia [b][e][h]	255	13(5.1)	131	1(0.8)	6.98(1.3	7,127.51)	0.0151	6.68(1.35,120.60)	0.0151	0.04(0.01,	0.08)	0.0151
Serious AESI: Neutropenia [c][g][i]	255	2(0.8)	131	0	2.59(0.2	1,358.70)	0.4990	NC		И	С	
Serious AESI: Thrombocytopeni a [c][g][i]	255	4(1.6)	131	0	4.71(0.5	0,625.36)	0.2078	NC		N	С	
Serious AESI: Vomiting [d][g][i]	255	0	131	0		NC		NC		N	С	
Serious AESI: Hypertension [b][e][h]	255	22(8.6)	131	16(12.2)	0.68(0.3	4, 1.36)	0.2698	0.71(0.39, 1.32)	0.2698	-0.04(-0.11,	0.03)	0.2698
Serious AESI: Proteinuria [c][g][i]	255	1(0.4)	131	0	1.55(0.0	8,226.84)	0.7818	NC		N	С	
Serious AESI: GI perforations, abscesses and fistulae [c][g][i]	255	2(0.8)	131	0	2.59(0.2	1,358.70)	0.4990	NC		N	С	

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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Table 3.6.2 PAOLA1: Summary of analysis of serious adverse events of special interest (odds ratio, relative risk and risk difference)

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

									Tr	reatme	nt effe	ect			
	Olapar: bevaciz (N=25	umab	bev	acebo + acizumab N=131)		Odds	Ratio		:	Relati	lve Ris	k	Risk D	ifferen	ce
	with	of ents		Number (%) of patients with events [a]		Estimat (95% Cl		2- sided p- value		stimat 95% CI		2- sided p- value	Estima (95% C		2- sided p- value
Serious AESI: Wound healing complications [c][g][i]	255 0		131	1(0.8)	0.17(0.00,	3.21)	0.2352		Ŋ	īĊ			NC	
Serious AESI: Haemorrhage [c][e][h]	255 2(0.8)	131	1(0.8)	0.86(0.11,	9.45)	0.8835	1.03(0.10,	21.97)	0.9823	0.00(-0.03,	0.02)	0.9823
Serious AESI: Arterial thromboembolic events [c][e][h]	255 1(0.4)	131	3(2.3)	0.22(0.02,	1.33)	0.0989	0.17(0.01,	1.32)	0.0916	-0.02(-0.05,	0.003)	0.0916
Serious AESI: Venous thromboembolic events [c][g][i]	255 2(0.8)	131	0	2.59(0.21,3	358.70)	0.4990		N	īC		:	NC	
Serious AESI: Non-GI fistula or abscess [d][g][i]	255 0		131	0		Ŋ	IC			N	rc			NC	
Serious AESI: MDS/AML [b][e][h]	255 3(1.2)	131	3(2.3)	0.51(0.09,	2.78)	0.4150	0.51(0.10,	2.74)	0.4150	-0.01(-0.05,	0.01)	0.4150

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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Table 3.6.2 PAOLA1: Summary of analysis of serious adverse events of special interest (odds ratio, relative risk and risk difference) Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

									Tre	eatme	nt effe	ect			
		aparib + vacizumab (N=255)		lacebo + vacizumab (N=131)		Odds	Ratio		R	Relati	ve Ris	k	Risk	Differe	nce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estimat (95% Cl	[)	2- sided p- value	(9	timat 5% CI)	2- sided p- value	Estir (95%	CI)	2- sided p- value
Serious AESI: Myelodysplastic syndrome and Acute myeloid leukaemia [b][e][h]	255	3(1.2)	131	4(3.1)	0.38(0.07,	1.74)	0.2055	0.39(0	.08,	1.72)	0.2055	-0.02(-0.06	0.01)	0.2055
Serious AESI: Secondary cancer [b][e][h]	255	15(5.9)	131	4(3.1)	1.98(0.70,	7.07)	0.2062	1.93(0	.72,	6.65)	0.2062	0.03(-0.02	0.07	0.2062
Serious AESI: Pneumonitis [c][g][i]	255	2(0.8)	131	0	2.59(0.21,3	358.70)	0.4990		N	С			NC	

Includes adverse events of special interest with an onset date on or after the date of first dose and up to the date of DCO.

MedDRA version 25.0.

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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									Tre	atment effe	ect			
	be ⁻	aparib + vacizumab (N=255)	ber	lacebo + vacizumab (N=131)		Odds	Ratio		Re	elative Ris	k	Risk Di	fferen	ce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estimat		2- sided p- value		imate % CI)	2- sided p- value	Estimat (95% CI		2- sided p- value
AESI G>=3: Anaemia [b][e][h]	255	47(18.4)	131	1(0.8)	29.37(6.31,5	523.04)	<0.0001	24.15(5.	39,425.07)	<0.0001	0.18(0.13,	0.23)	<0.0001
AESI G>=3: Neutropenia [b][e][h]	255	21(8.2)	131	4(3.1)	2.85(1.06,	9.93)	0.0377	2.70(1.	05, 9.10)	0.0377	0.05(0.003,	0.10)	0.0377
AESI G>=3: Thrombocytopeni a [b][e][h]	255	5(2.0)	131	4(3.1)	0.63(0.17,	2.60)	0.5090	0.64(0.	17, 2.56)	0.5090	-0.01(-0.05,	0.02)	0.5090
AESI G>=3: Nausea [b][e][h]	255	9(3.5)	131	4(3.1)	1.16(0.37,	4.35)	0.8046	1.16(0.	38, 4.20)	0.8046	0.00(-0.04,	0.04)	0.8046
AESI G>=3: Vomiting [b][e][h]	255	4(1.6)	131	6(4.6)	0.33(0.08,	1.18)	0.0883	0.34(0.	09, 1.18)	0.0883	-0.03(-0.08,0	.004)	0.0883
AESI G>=3: Fatigue and Asthenia [b][e][h]	255	17(6.7)	131	3(2.3)	3.05(1.0005,	13.22	0.0499	2.91(1.0	0004, 12.31) 0.0499	0.04(0.00002	,	0.0499
AESI G>=3: Hypertension [b][e][h]	255	50(19.6)	131	42(32.1)	0.52(0.32,	0.84)	0.0074	0.61(0.	43, 0.87)	0.0074	-0.12(-0.22,	-0.03)	0.0074

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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Table 3.6.3 PAOLA1: Summary of analysis of severe adverse events of special interest with max. CTCAE grade >=3 (odds ratio, relative risk and risk difference)

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

								Treatm	ent effe	ct			
	beva	parib + cizumab =255)	bev	lacebo + vacizumab (N=131)	Odds	Ratio		Relat	ive Risl	ς	Risk D	ifferen	
	(p w n e	umber %) of atients ith vents [a]	n	Number (%) of patients with events [a]	Estimat (95% CI	e g	2- sided p- value	Estima (95% C	I)	2- sided p- value	Estimat (95% C	I)	2- sided p- value
AESI G>=3: Proteinuria [c][g][i]	255	3(1.2)	131	0	3.65(0.35,4	91.54) (0.3214		NC		Ţ	1C	
AESI G>=3: GI perforations, abscesses and fistulae [c][g][i]	255	3(1.2)	131	0	3.65(0.35,4	91.54) (0.3214		NC		1	VC	
AESI G>=3: Wound healing complications [d][g][i]	255	0	131	0	N	С			NC		1	IC	
AESI G>=3: Haemorrhage [c][e][h]	255	2(0.8)	131	1(0.8)	0.86(0.11,	9.45) 0	0.8835	1.03(0.10,	21.97)	0.9823	0.00(-0.03,	0.02)	0.9823
AESI G>=3: Arterial thromboembolic events [b][e][h]	255	1(0.4)	131	4(3.1)	0.13(0.01,		0.0333	0.13(0.01,	0.86)	0.0333	-0.03(-0.07,	-0.00)	0.0333
AESI G>=3: Venous thromboembolic events [c][g][i]	255	3(1.2)	131	0	3.65(0.35,4	91.54) (0.3214		NC		1	1C	

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c].

[e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h].

Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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								Treatmo	ent effe	ect			
	ber	aparib + vacizumab (N=255)	be ⁻	lacebo + vacizumab (N=131)	Odd	s Ratio		Relat	ive Ris	k	Risk D	ifferen	nce
	n	Number (%) of patients with	n	Number (%) of patients with	Estima (95%)		2- sided p- value	Estima (95% C		2- sided p- value	Estimat (95% CI		2- sided p- value
AESI G>=3: Congestive heart failure [d][g][i]	255	0	131	0		NC			NC		Ν	ĪĊ	
AESI G>=3: Non-GI fistula or abscess [c][g][i]	255	0	131	1(0.8)	0.17(0.00	3.21)	0.2352		NC		P	IC	
AESI G>=3: MDS/AML [b][e][h]	255	4(1.6)	131	3(2.3)	0.68(0.15	3.49)	0.6211	0.68(0.15,	3.43)	0.6211	-0.01(-0.04,	0.02)	0.6211
AESI G>=3: Myelodysplastic syndrome and Acute myeloid leukaemia [b][e][h]	255	4(1.6)	131	4(3.1)	0.51(0.12	2.17)	0.3452	0.51(0.12,	2.14)	0.3452	-0.01(-0.06,	0.02)	0.3452
AESI G>=3: Secondary cancer [b][e][h]	255	11(4.3)	131	3(2.3)	1.92(0.59	8.61)	0.2958	1.88(0.60,	8.22)	0.2958	0.02(-0.02,	0.06)	0.2958

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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Table 3.6.4 PAOLA1: Summary of analysis of non-severe adverse events of special interest with max. CTCAE grade 1 or 2 (odds ratio, relative risk and risk difference)

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

										Treatme	nt effe	ect			
	be	laparib + vacizumab (N=255)	be	lacebo + vacizumab (N=131)		Odds	s Ratio			Relat	ive Ris	k	Risk Di	fferen	ce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estima (95% C		2- sided p- value		Estimat (95% C]		2- sided p- value	Estimat (95% CI		2- sided p- value
AESI G1-2: Anaemia [b][e][h]	255	86(33.7)	131	11(8.4)	5.55(2.95,	11.41)	<0.0001	4.02(2.34,	7.73)	<0.0001	0.25(0.18,	0.33)	<0.0001
AESI G1-2: Neutropenia [b][e][h]	255	39(15.3)	131	19(14.5)	1.06(0.59,	1.96)	0.8366	1.05(0.65,	1.79)	0.8366	0.01(-0.07,	0.08)	0.8366
AESI G1-2: Thrombocytopeni a [b][e][h]	255	14(5.5)	131	4(3.1)	1.84(0.65,	6.61)	0.2659	1.80(0.66,	6.25)	0.2659	0.02(-0.02,	0.06)	0.2659
AESI G1-2: Nausea [b][e][h]	255	142(55.7)	131	30(22.9)	4.23(2.65,	6.90)	<0.0001	2.43(1.78,	3.47)	<0.0001	0.33(0.23,	0.42)	<0.0001
AESI G1-2: Vomiting [b][e][h]	255	52(20.4)	131	14(10.7)	2.14(1.17,	4.17)	0.0133	1.91(1.14,	3.46)	0.0133	0.10(0.02,	0.17)	0.0133
AESI G1-2: Fatigue and Asthenia [b][e][h]	255	133(52.2)	131	44(33.6)	2.16(1.40,	3.36)	0.0005	1.55(1.20,	2.06)	0.0005	0.19(0.08,	0.28)	0.0005
AESI G1-2: Hypertension [b][e][h]	255	98(38.4)	131	59(45.0)	0.76(0.50,	1.17)	0.2119	0.85(0.67,	1.10)	0.2119	-0.07(-0.17,	0.04)	0.2119

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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Table 3.6.4 PAOLA1: Summary of analysis of non-severe adverse events of special interest with max. CTCAE grade 1 or 2 (odds ratio, relative risk and risk difference)

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

									,	Treatme	ent effe	ect			
	ber	aparib + vacizumab (N=255)	ber	lacebo + vacizumab (N=131)		Odds	Ratio			Relat	ive Ris	k	Risk D	ifferen	ce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estimat (95% CI		2- sided p- value		Estima (95% C		2- sided p- value	Estimat (95% C)		2- sided p- value
AESI G1-2: Proteinuria [b][e][h]	255	18(7.1)	131	19(14.5)	0.45(0.22,	0.89)	0.0218	0.49(0.26,	0.90)	0.0218	-0.07(-0.15,	-0.01)	0.0218
AESI G1-2: Wound healing complications [b][e][h]	255	2(0.8)	131	3(2.3)	0.34(0.04,	2.06)	0.2314	0.34(0.05,	2.04)	0.2314	-0.02(-0.05,	0.01)	0.2314
AESI G1-2: Haemorrhage [b][e][h]	255	28(11.0)	131	11(8.4)	1.35(0.66,	2.91)	0.4188	1.31(0.69,	2.67)	0.4188	0.03(-0.04,	0.08)	0.4188
AESI G1-2: Arterial thromboembolic events [c][g][i]	255	2(0.8)	131	0	2.59(0.21,3	358.70)	0.4990		1	NC		I	IC	
AESI G1-2: Venous thromboembolic events [b][e][h]	255	8(3.1)	131	1(0.8)	4.21(0.76,	78.53)	0.1092	4.11(0.77,	75.81)	0.1092	0.02(-0.01,	0.05)	0.1092
AESI G1-2: Posterior Reversible Encephalopathy Syndrome (PRES) [c][g][i]	255	0	131	1(0.8)	0.17(0.00,	3.21)	0.2352		1	NC		7	IC	

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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Table 3.6.4 PAOLA1: Summary of analysis of non-severe adverse events of special interest with max. CTCAE grade 1 or 2 (odds ratio, relative risk and risk difference)

Safety Analysis Set, HRD[42] positive, DCO 22Mar2022

								Treatment effe	ect		
		laparib + vacizumab (N=255)		lacebo + vacizumab (N=131)		Odds Ratio		Relative Ris	k	Risk Differe	nce
	n	Number (%) of patients with events [a]	n	Number (%) of patients with events [a]		Estimate (95% CI)	2- sided p- value	Estimate (95% CI)	2- sided p- value	Estimate (95% CI)	2- sided p- value
AESI G1-2: Non-GI fistula or abscess [c][g][i]	255	0	131	1(0.8)	0.17(0.00, 3.21)	0.2352	NC		NC	
AESI G1-2: Secondary cancer [c][e][h]	255	3(1.2)	131	1(0.8)	1.21(0.20, 12.53)	0.8470	1.54(0.20, 30.97)	0.6971	0.00(-0.02, 0.03	0.6971
AESI G1-2: Pneumonitis [c][g][i]	255	3(1.2)	131	0	3.65(0.35,491.54)	0.3214	NC		NC	

NC=not calculable. CI=confidence interval. PL=profile likelihood. LR=likelihood ratio test. All models include treatment only.

[a] Patients with multiple events in the same category are counted only once in that category.

Patients with events in more than one category are counted once in each of those categories.

[b] Odds ratio (OR), 95% PL CI, LR p-value via logistic regression. [c] As [b] but with Firth method. [d] OR NC via [b] or [c]. [e] Relative risk (RR), 95% PL CI, LR p-value via log-binomial regression. [f] RR, 95% CI, p-value via modified poisson regression. [g] RR NC via [e] or [f]. [h] Risk difference (RD), 95% PL CI, LR p-value via binomial regression. [i] RD NC via [h]. Odds ratio and relative risk <1, and risk difference <0 favours olaparib. * p<0.05.

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Table 4.1 PAOLAl Appendix: Summary of Myelodysplastic syndrome/Acute myeloid leukaemia - adverse events of special interests by preferred term

Safety Analysis Set, HRD[42] positive, DCO 22MAR2022

AESI term/ MedDRA Preferred term	Number (%) of patients	
	Olaparib+ bevacizumab (N=255)	Placebo + bevacizumab (N=131)
MDS/AML	4 (1.6)	3 (2.3)
Acute myeloid leukaemia	4 (1.6)	1 (0.8)
Myelodysplastic syndrome	1 (0.4)	3 (2.3)
Myelodysplastic syndrome and Acute myeloid leukaemia	4 (1.6)	4 (3.1)
Acute leukaemia	0	1 (0.8)
Acute myeloid leukaemia	4 (1.6)	1 (0.8)
Myelodysplastic syndrome	1 (0.4)	3 (2.3)