

Supplementary Materials: Estrogen Receptor Status Oppositely Modifies Breast Cancer Prognosis in *BRCA1/BRCA2* Mutation Carriers Versus Non-Carriers

Michal Vocka, Martina Zimovjanova, Zuzana Bielcikova, Petra Tesarova, Lubos Petruzela, Martin Mateju, Ludmila Krizova, Jaroslav Kotlas, Jana Soukupova, Marketa Janatova, Petra Zemankova, Petra Kleiblova, Jan Novotny, Bohuslav Konopasek, Martina Chodacka, Milan Brychta, Marek Sochor, Denisa Smejkalova-Musilova, Vlastimila Cmejlova, Renata Kozevnikovova, Lenka Miskarova, Sonja Argalacsova, Lenka Stolarova, Klara Lhotova, Marianna Borecka and Zdenek Kleibl

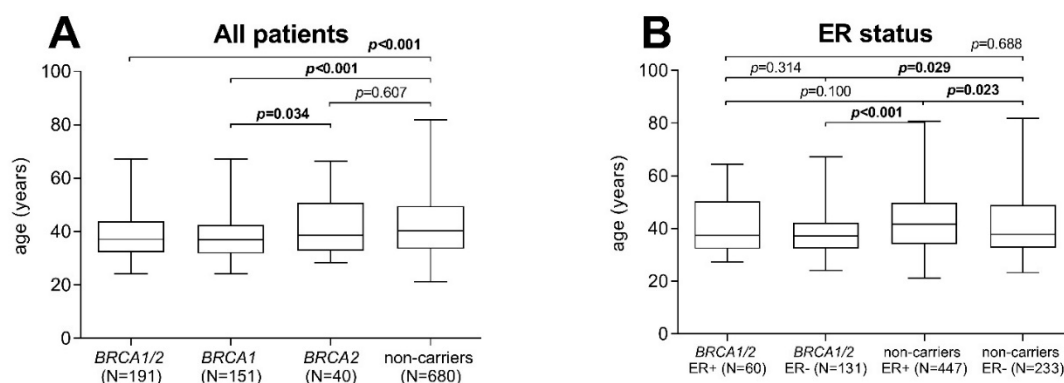


Figure S1. Differences in the age at diagnosis in mutation carriers and non-carriers in particular breast cancer subtypes; (A) distribution according to the mutation status; (B) distribution according to the mutation status and ER status.

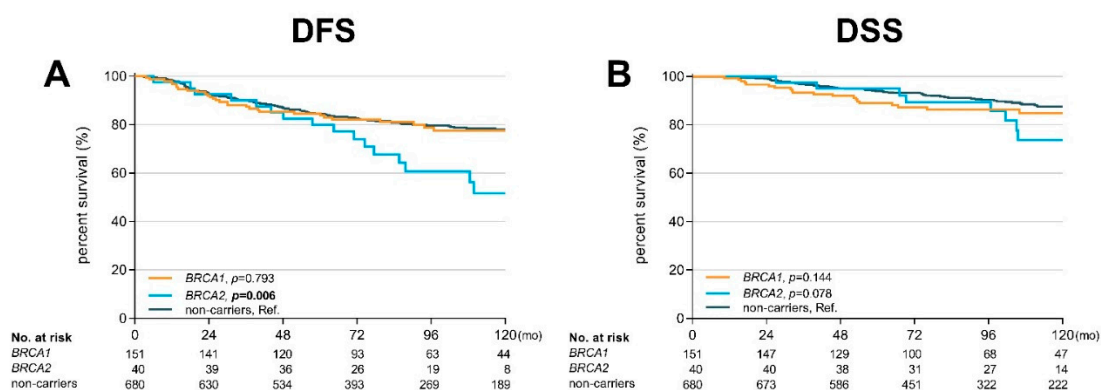


Figure S2. Kaplan–Meier plots showing DFS (A) and DSS (B) in *BRCA1* mutation carriers, *BRCA2* mutation carriers, and non-carriers.

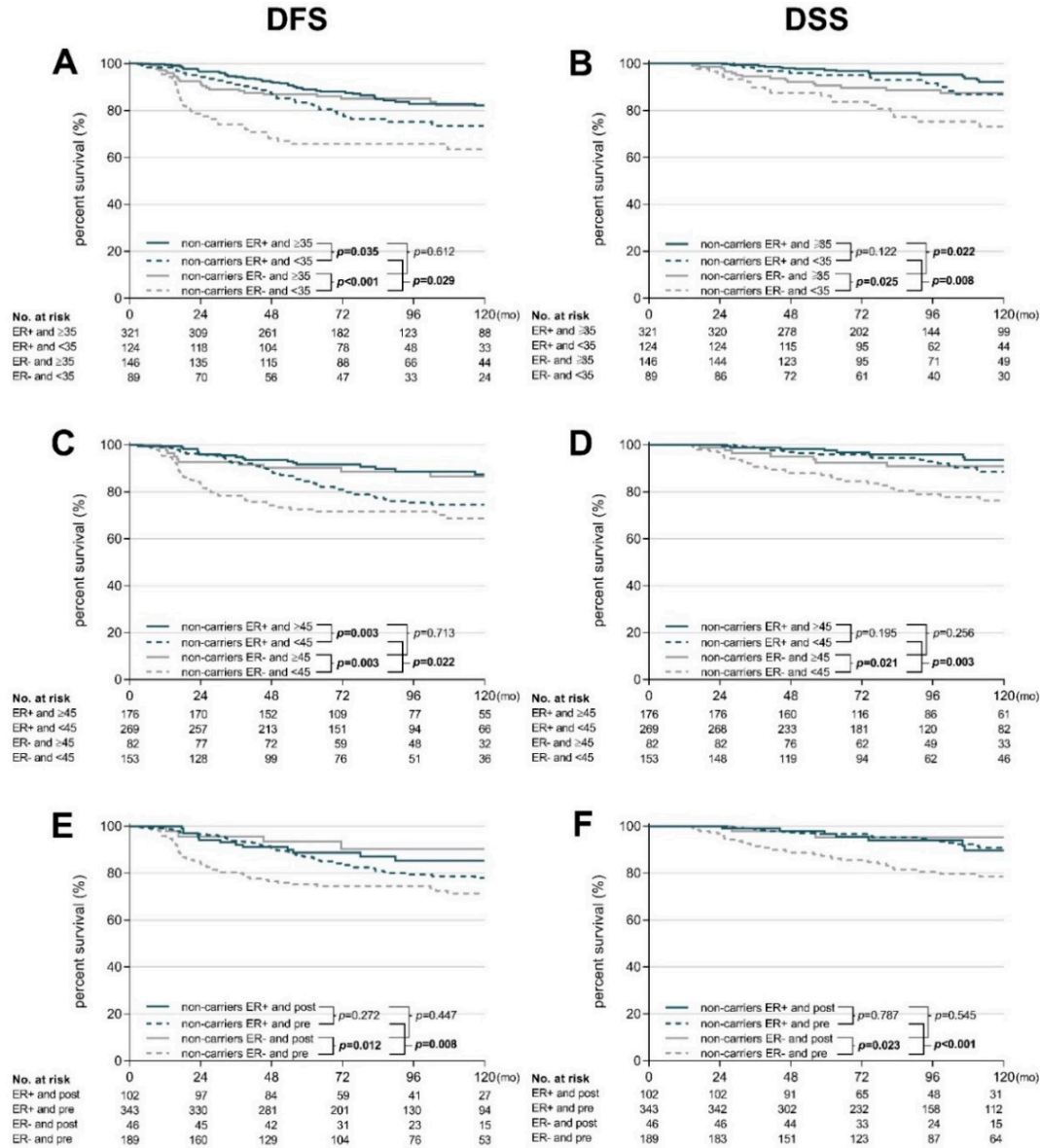


Figure S3. Kaplan–Meier plots of DFS and DSS in *BRCA1/2* mutations non-carriers considering a combined impact of ER-status and age of onset (**A**, **B** <35 vs. ≥35; **C**, **D** <45 vs. ≥45) or menopausal status (**E**, **F**) on survival. (**A**) ER-negative patients diagnosed at <35 years were at higher risk of recurrence (HR 2.61 [95%CI 1.49–4.56]) than ER-negative women diagnosed at ≥35 years. These high-risk ER-negative young patients (diagnosed at <35 years) were also at higher risk (HR 1.79 [95%CI 1.06–3.02]) than ER-positive young women diagnosed at the same age. (**B**) ER-negative patients diagnosed at <35 years were at higher risk of BC-related death than ER-negative women diagnosed at ≥35 years (HR 2.17 [95%CI 1.10–4.25]) and also than ER-positive young women diagnosed at the same age (HR 2.61 [95%CI 1.29–5.30]). (**C**) ER-negative patients diagnosed at <45 years were at higher risk of recurrence than ER-negative women diagnosed at ≥45 years (HR 2.29 [95%CI 1.32–3.98]) and also than ER-positive young women diagnosed at the same age (HR 1.63 [95%CI 1.07–2.47]). (**D**) ER-negative patients diagnosed at <45 years were at higher risk of BC-related death than ER-negative women diagnosed at ≥45 years (HR 2.22 [95%CI 1.13–4.35]) and also than ER-positive young women diagnosed at the same age (HR 3.00 [95%CI 1.67–5.42]). (**E**) Besides, ER-negative premenopausal patients were at higher risk than ER-negative postmenopausal patients (HR 2.31 [95%CI 1.20–4.42]) and ER-positive premenopausal patients (HR 1.71 [95%CI 1.15–2.53]). (**F**) ER-negative premenopausal patients were at higher risk than ER-negative postmenopausal patients (HR 2.53 [95%CI 1.14–5.66]) and ER-positive premenopausal patients (HR 3.36 [95%CI 1.93–5.87]).

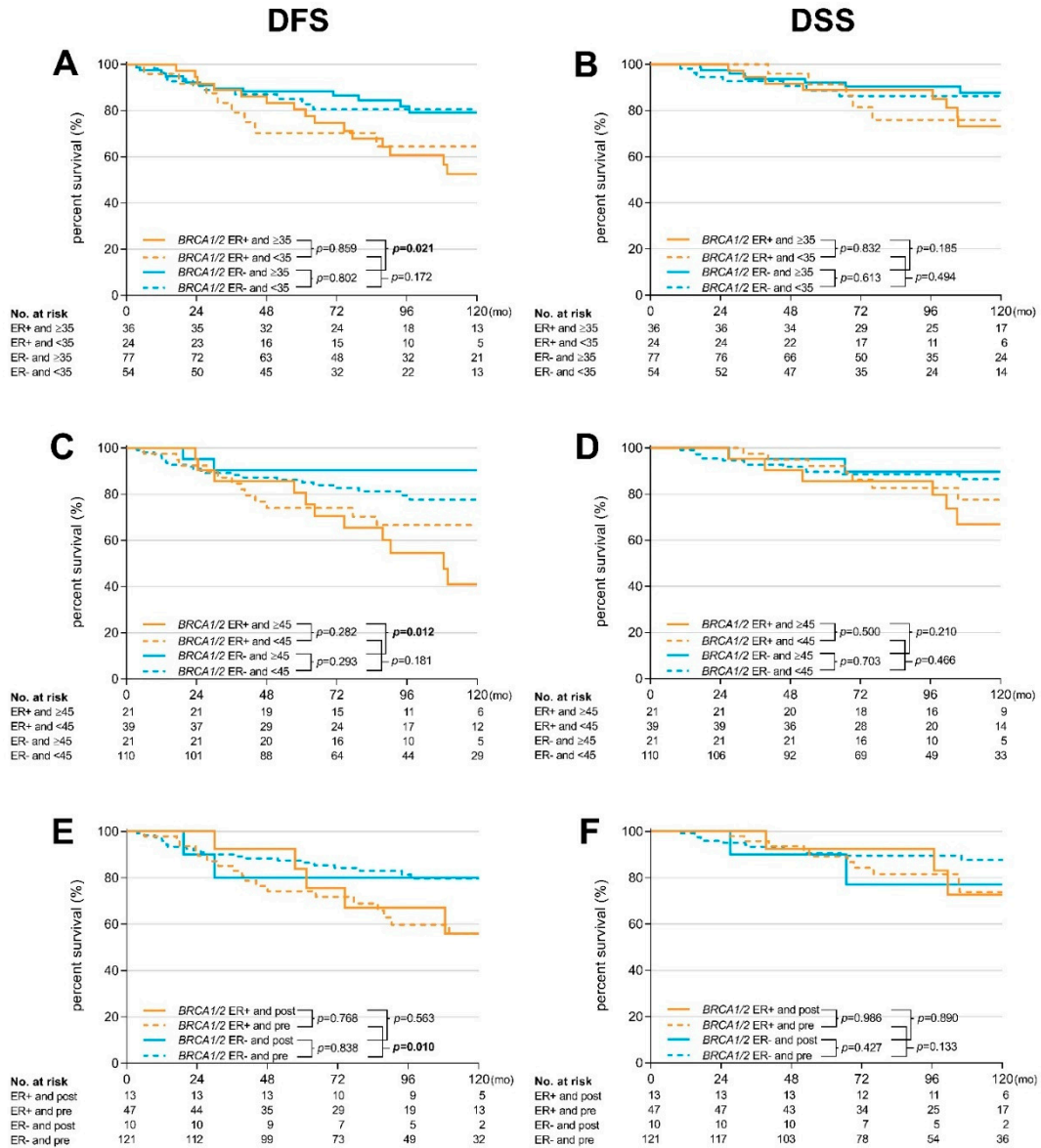


Figure S4. Kaplan–Meier plots of DFS and DSS in *BRCA1/2* mutation carriers considering a combined impact of ER-status and age of onset (A, B <35 vs. ≥35; C, D <45 vs. ≥45) or menopausal status (E, F) on survival. (A) Patients diagnosed at ≥35 years with ER-positive tumors showed a significantly increased risk of BC recurrence (HR 2.53 [95%CI 1.15–5.57]), compared with ER-negative mutation carriers of the same age. (B) Risk of BC-related death in ER-positive patients diagnosed at ≥35 years did not significantly differ from ER-negative women diagnosed at ≥35 years (HR 2.01 [95%CI 0.72–5.66]). (C) Increased risk of BC recurrence was shown for ER-positive patients ≥45 years (HR 4.03 [95%CI 1.36–12.00]), compared with ER-negative patients of the same age. (D) Risk of BC-related death in ER-positive patients diagnosed at ≥45 years did not significantly differ from ER-negative women diagnosed at ≥45 years (HR 2.44 [95%CI 0.60–9.85]). (E) ER-positive postmenopausal patients were at similar risk of BC recurrence as ER-negative postmenopausal patients (HR 1.57 [95%CI 0.34–7.15]). (F) Risk of BC-related death in ER-positive postmenopausal patients did not significantly differ from ER-negative postmenopausal patients (HR 0.88 [95%CI 0.14–5.46]).

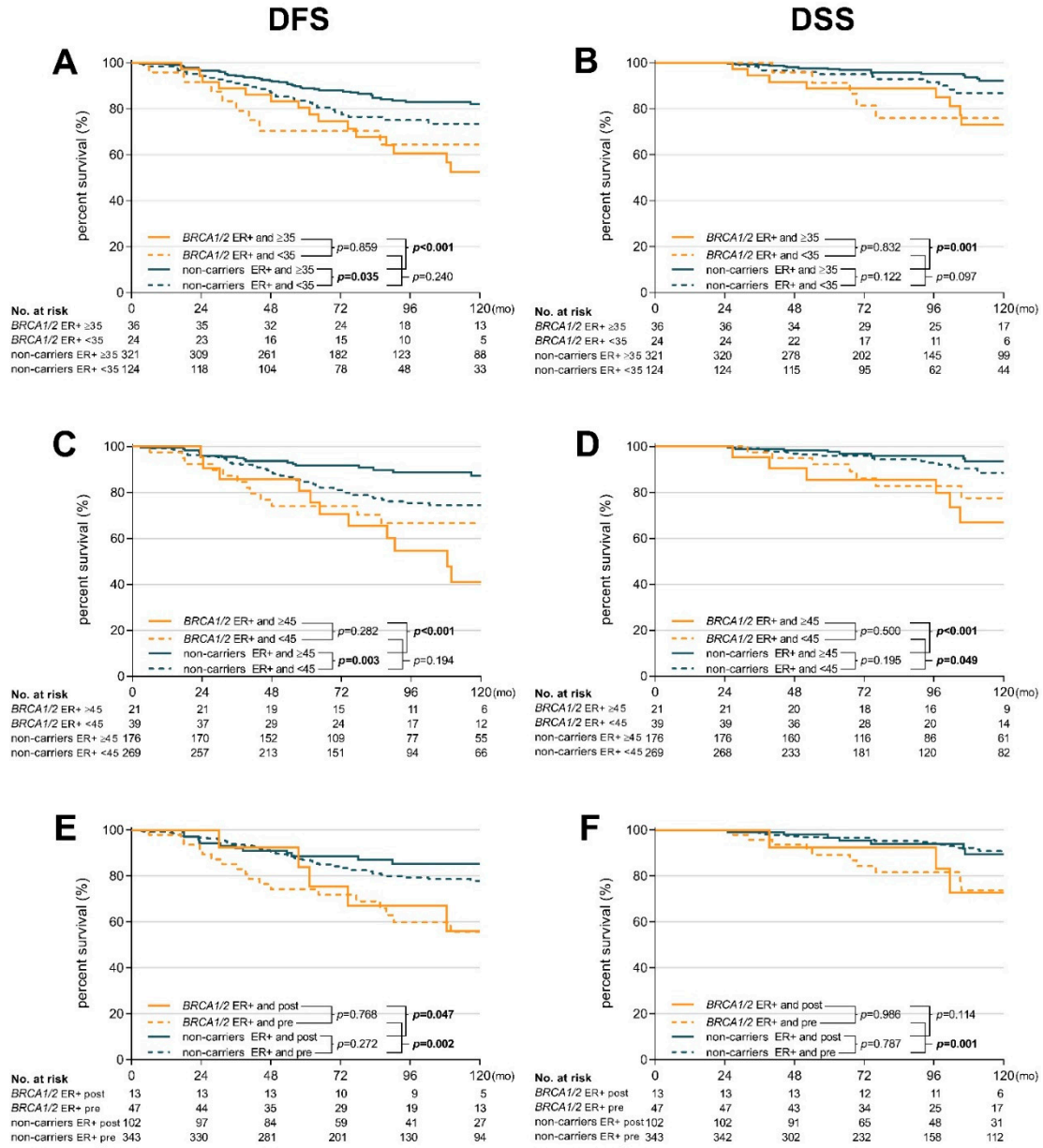


Figure S5. Cont.

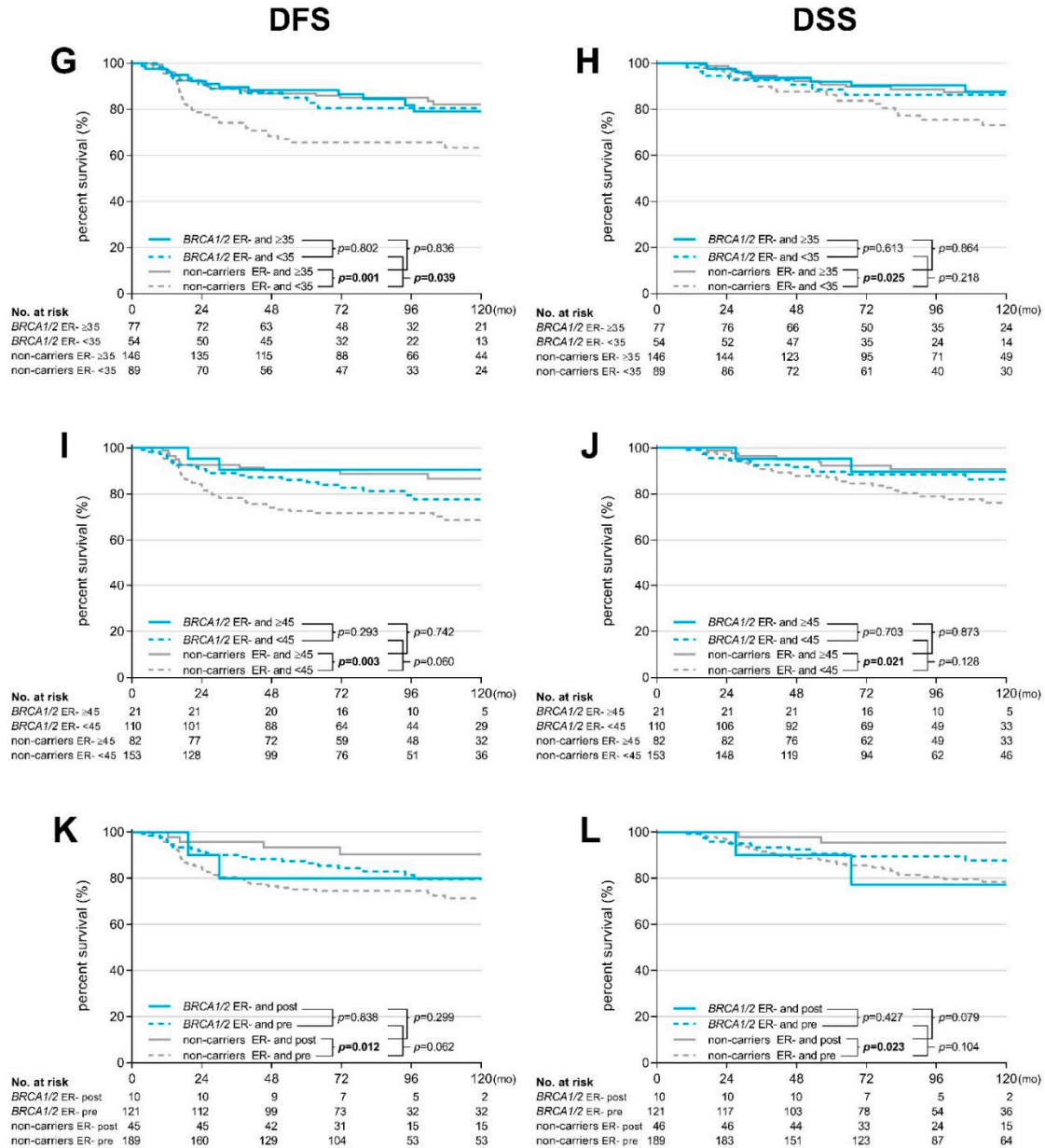


Figure S5. Kaplan-Meier plots of DFS and DSS in *BRCA1/2* mutation carriers and non-carriers considering a combined effect of ER-positivity (A–F) or ER-negativity (G–L) and age of disease onset or menopausal status on survival.

Table S1. List of pathogenic mutations identified in the *BRCA1* and *BRCA2* genes in analyzed breast cancer patients.

<i>BRCA1/2</i> Mutation Carriers					
Gene	Mutation	No.	Gene	Mutation	No.
<i>BRCA1</i>	c.5266dupC	73	<i>BRCA2</i>	c.5682C>G	4
	c.181T>G	15		c.3847_3848delGT	3
	c.3700_3704delGTAAA	14		c.5946_5946delT	3
	c.1687C>T	9		c.7913_7917delTTCCT	3
	c.213-12A>G	5		c.2808_2811delACAA	2
	c.66_67delAG	5		c.4284_4285insT	2
	c.2411_2412delAG	4		c.7595_7596insTT	2
	c.3756_3759delGTCT	4		c.8042_8043delCA	2
	c.1600C>T	2		c.8755-1G>A	2
	c.3016_3019delCATT	2		c.9097_9098insA	2
	c.4165_4166delAG	2		c.9435_9436delGT	2
	c.4956G>A	2		c.1296_1297delGA	1
	c.505C>T	2		c.1773_1776delTTAT	1
	c.5074+1G>T	2		c.1796_1800delCTTAT	1
	c.1016delA	1		c.1813_1814insA	1
	c.115T>C	1		c.3978_3979insTGCT	1
	c.1204delG	1		c.3G>A	1
	c.1916T>A	1		c.475+1G>T	1
	c.2073dupA	1		c.5238_5239insT	1
	c.2263G>T	2		c.5722_5723delCT	1
	c.2762delA	1		c.5763_5764insT	1
	c.315T>A	1		c.5763dupT	1
	c.3226delA	1		c.604_613del10	1
	c.3239T>A	1		c.6275_6276delTT	1
	c.3331C>T	1		c.6444_6445insT	1
	c.3607C>T	1		c.6444dupT	1
	c.3770_3771delAG	1		c.6447_6448dupTA	1
	c.4158_4162delCTCTC	1		c.6591_6592delTG	1
	c.4243delG	1		c.6754_6755dupT	1
	c.4539C>T	1		c.7151_7152delAA	1
	c.4689C>G	1		c.771_775delTCAAA	1
	c.4868C>G	1		c.8363G>A	1
	c.5075-2A>G	1		c.8535_8538delAGAG	1
	c.5152+1G>A	1		c.9403_9403delC	1
	c.5152+2dupT	1		c.994dupA	1
	c.5673delA	1			
CNVs					
	exon 5–14del	10			
	exon 1–17del	4			
	exon 13dup	1			
	exon 13–19del	1			
	exon 18–22del	1			
	exon 21–22del	1			
	exon 21–24del	1			

No. – number of carriers; CNVs – copy number variants. *Note:* Mutation (at coding sequence) were annotated according to the reference sequence NM_007294 (for *BRCA1*) and NM_000059 (for *BRCA2*).

Table S2. Clinicopathological characteristics of all *BRCA1/BRCA2* mutation carriers (together and separately) and non-carriers of mutations in cancer-susceptibility genes. (*N* = denotes for individuals with known variables).

	<i>BRCA1/2</i> Carriers (<i>N</i> = 234)			<i>BRCA1</i> Carriers (<i>N</i> = 183)			<i>BRCA2</i> Carriers (<i>N</i> = 51)			Non-Carriers (<i>N</i> = 899)		
	<i>N</i>	% *	<i>p</i>	<i>N</i>	% *	<i>p</i>	<i>N</i>	% *	<i>p</i>	<i>N</i>	% *	<i>p</i>
Median age at diagnosis year (25–75% percentile)		(<i>N</i> = 234)			(<i>N</i> = 183)			(<i>N</i> = 51)			(<i>N</i> = 899)	
	37.7	(32.9–44.0)	<0.001	37.3	(32.8–49.5)	<0.001	39.7	(34.0–50.1)	0.776	40.9	(33.8–49.5)	Ref.
Age diagnosis categories		(<i>N</i> = 234)			(<i>N</i> = 183)			(<i>N</i> = 51)			(<i>N</i> = 899)	
< 35 y	86	(36.8)	<0.001	71	(38.8)	<0.001	15	(29.4)	0.718	276	(30.7)	Ref.
35–44 y	96	(41.0)		78	(42.6)		18	(35.3)		270	(30.0)	
≥ 45 y	52	(22.2)		34	(18.6)		18	(35.3)		353	(39.3)	
Menopausal status		(<i>N</i> = 234)			(<i>N</i> = 183)			(<i>N</i> = 51)			(<i>N</i> = 889)	
pre	207	(88.5)	0.001	166	(90.7)	<0.001	41	(80.4)	0.838	712	(79.2)	Ref.
post	27	(11.5)		17	(9.3)		10	(19.6)		187	(20.8)	
Primary tumor (T)		(<i>N</i> = 227)			(<i>N</i> = 180)			(<i>N</i> = 47)			(<i>N</i> = 867)	
Tis	1	(0.4)	<0.001	1	(0.6)	<0.001	0	(0.0)	0.156	45	(5.2)	Ref.
T1 (<2 cm)	78	(34.4)		57	(31.7)		21	(44.7)		417	(48.1)	
T2 (2–5 cm)	110	(48.5)		94	(52.2)		16	(34.0)		303	(34.9)	
T3 (>5 cm)	26	(11.5)		19	(10.6)		7	(14.9)		81	(9.3)	
T4	12	(5.3)		9	(5.0)		3	(6.4)		21	(2.4)	
Regional lymphatic node (N)		(<i>N</i> = 227)			(<i>N</i> = 180)			(<i>N</i> = 47)			(<i>N</i> = 867)	
N0	121	(53.3)	0.088	101	(56.1)	0.120	20	(42.6)	0.068	526	(60.7)	Ref.
N1	92	(40.5)		67	(37.2)		25	(53.2)		301	(34.7)	
N2	9	(4.0)		7	(3.9)		2	(4.3)		33	(3.8)	
N3	5	(2.2)		5	(2.8)		0	(0.0)		7	(0.8)	
Tumor stage		(<i>N</i> = 228)			(<i>N</i> = 180)			(<i>N</i> = 48)			(<i>N</i> = 889)	
0 (TisN0)	1	(0.4)	<0.001	1	(0.6)	<0.001	0	(0.0)	0.024	44	(4.9)	Ref.
I (T1N0–1mi)	58	(25.4)		45	(25.0)		13	(25.5)		334	(37.6)	
II (T2–3N0, T1–2N1)	123	(53.9)		101	(56.1)		22	(43.1)		389	(43.8)	
III (T3N1, TXN2–3, T4NX)	45	(19.7)		33	(18.3)		12	(23.5)		99	(11.1)	
IV (M1)	1	(0.4)		0	(0.0)		1	(2.0)		23	(2.6)	
Tumor morphology		(<i>N</i> = 232)			(<i>N</i> = 182)			(<i>N</i> = 50)			(<i>N</i> = 891)	
ductal	195	(84.1)	0.012	151	(83.0)	0.001	44	(88.0)	0.390	738	(82.8)	Ref.

lobular	6	(2.6)		3	(1.6)		3	(6.0)		70	(7.9)	
medullar	18	(7.8)		18	(9.9)		0	(0.0)		43	(4.8)	
other	13	(5.6)		10	(5.5)		3	(6.0)		40	(4.5)	
Tumor grade		(N = 216)			(N = 173)			(N = 43)			(N = 832)	
low (1)	8	(3.7)	<0.001	5	(2.9)	<0.001	3	(7.0)	0.437	115	(13.8)	Ref.
intermediate (2)	70	(32.4)		49	(28.3)		21	(48.8)		383	(46.0)	
high (3)	138	(63.9)		119	(68.8)		19	(44.2)		334	(40.1)	
ER status		(N = 213)			(N = 169)			(N = 44)			(N = 827)	
positive	72	(33.8)	<0.001	45	(26.6)	<0.001	27	(61.4)	0.519	546	(66.0)	Ref.
PR status		(N = 207)			(N = 164)			(N = 43)			(N = 799)	
positive	69	(33.3)	<0.001	39	(23.8)	<0.001	30	(69.8)	0.419	502	(62.8)	Ref.
HER-2 status		(N = 193)			(N = 152)			(N = 41)			(N = 732)	
positive	13	(6.7)	<0.001	10	(6.6)	<0.001	3	(7.3)	0.012	556	(24.0)	Ref.
TNBC		(N = 193)			(N = 152)			(N = 41)			(N = 729)	
yes	115	(59.6)	<0.001	105	(69.1)	<0.001	10	(24.4)	0.547	145	(19.9)	Ref.
Surgery – primary tumor		(N = 231)			(N = 182)			(N = 49)			(N = 876)	
mastectomy	122	(52.8)	0.484	94	(51.6)	0.727	28	(57.1)	0.346	440	(50.2)	Ref.
breast-conserving surgery	109	(47.2)		88	(48.4)		21	(42.9)		436	(49.8)	
Surgery – lymphatic nodes		(N = 227)			(N = 181)			(N = 46)			(N = 859)	
axillary dissection	184	(81.1)	0.054	146	(80.7)	0.105	38	(82.6)	0.294	643	(74.9)	Ref.
sentinel node biopsy	43	(18.9)		35	(19.3)		8	(17.4)		216	(25.1)	
Radiotherapy		(N = 229)			(N = 181)			(N = 48)			(N = 869)	
yes	157	(68.6)	0.475	128	(70.7)	0.225	29	(60.4)	0.423	574	(66.1)	Ref.
Chemotherapy type		(N = 232)			(N = 182)			(N = 50)			(N = 873)	
Antra + Tax	122	(52.6)	<0.001	97	(53.3)	<0.001	25	(50.0)	0.214	311	(35.6)	Ref.
Antra	61	(26.3)		49	(26.9)		12	(24.0)		243	(27.8)	
Other	21	(9.1)		17	(9.3)		4	(8.0)		85	(9.7)	
No chemotherapy	28	(12.1)		19	(10.4)		9	(18.0)		234	(26.8)	
Endocrine therapy **		(N = 65)			(N = 38)			(N = 27)			(N = 490)	
TAM monotherapy	26	(40.0)	0.961	18	(47.4)	0.886	8	(29.6)	0.655	203	(41.4)	Ref.
AI monotherapy	8	(12.3)		4	(10.5)		4	(14.8)		66	(13.5)	
LHRH analogues + TAM	29	(44.6)		15	(39.5)		14	(51.9)		210	(42.9)	
LHRH analogues + AI	2	(3.1)		1	(2.6)		1	(3.7)		11	(2.2)	
Event during follow-up ***		(N = 233)			(N = 183)			(N = 51)			(N = 876)	

loco-regional recurrence	14	(6.0)	0.169	13	(7.1)	0.457	1	(2.0)	0.093	77	(8.8)	Ref.
distant metastasis	47	(20.2)	0.002	28	(15.3)	0.289	19	(38.0)	0.027	110	(12.6)	Ref.
second breast cancer	43	(18.5)	<0.001	34	(18.6)	<0.001	9	(18.0)	0.026	76	(8.7)	Ref.
second tumors	12	(5.2)	0.753	10	(5.5)	0.805	2	(4.0)	0.570	52	(5.9)	Ref.
Median of follow-up (years)	(N = 234)			(N = 183)			(N = 51)			(N = 899)		
median (25–75% percentil)	10.0 (6.3–16.0)		0.235	9.6 (6.14–15.44)		0.692	11.2 (7.7–17.3)		0.048	9.8 (5.9–15.8)		Ref.
Breast cancer-related death	(N = 234)			(N = 183)			(N = 51)			(N = 899)		
yes	32	(13.7)	0.507	22	(12.0)	0.429	13	(25.5)	0.028	128	(14.2)	Ref.

* % = percentage of known; ** N = number of patients with ER-positive BC; *** patient could be counted in more than one event; pre – premenopausal; post – postmenopausal; TNBC – triple-negative BC; Antra – anthracyclines; Tax – taxanes; TAM – tamoxifen; AI – aromatase inhibitor; LHRH - luteinizing hormone-releasing hormone; Ref. – reference.

Table S3. Analysis of 10-year DFS and DSS using the Mantel-Haenszel test comparing variables within subgroups of *BRCA1/2* mutation carriers and non-carriers, respectively.

Variable	Category	Disease Free Survival (DFS) Analysis											Disease Specific Survival (DSS) Analysis												
		<i>BRCA1/2</i> Carriers						Non-carriers					<i>BRCA1/2</i> Carriers						Non-Carriers						
		Pts No.	Ev No.	Ev %	HR	95% CI	<i>p</i>	Pts No.	Ev No.	Ev %	HR	95% CI	<i>p</i>	Pts No.	Ev No.	%	HR	95% CI	<i>p</i>	Pts No.	Ev No.	%	HR	95% CI	<i>p</i>
All Pts		191	46	24.1	-	-	-	680	128	18.8	-	-	-	191	28	14.7	-	-	-	680	64	9.4	-	-	-
Age at diagnosis	<35	78	18	23.1	0.98	0.55–1.78	0.960	213	60	28.2	2.21	1.51–3.22	<0.001	78	12	15.4	1.16	0.54–2.48	0.703	213	32	15.0	2.29	1.35–3.87	0.002
	≥35	113	28	24.8	Ref.			467	68	14.6	Ref.			113	16	14.2	Ref.			467	32	6.9	Ref.		
	<45	150	34	22.7	0.84	0.42–1.67	0.612	422	100	23.7	2.17	1.53–3.09	<0.001	150	21	14.0	0.88	0.36–2.13	0.779	422	49	11.6	1.98	1.20–3.26	0.008
	≥45	41	12	29.3	Ref.			258	28	10.9	Ref.			41	7	17.1	Ref.			258	15	5.8	Ref.		
Menopausal status	pre	168	39	23.2	0.80	0.34–1.90	0.611	532	119	22.4	1.71	1.15–2.56	0.009	168	23	13.7	0.62	0.20–1.90	0.406	532	55	10.3	1.53	0.88–2.64	0.129
	post	23	7	30.4	Ref.			148	18	12.2	Ref.			23	5	21.7	Ref.			148	9	6.1	Ref.		
Tumor size	T1	71	11	15.5	Ref.			355	45	12.7	Ref.			71	4	5.6	Ref.			355	18	5.1	Ref.		
	T2	84	20	23.8	1.63	0.80–2.29	0.176	244	52	21.3	1.94	1.28–2.92	0.002	84	12	14.3	2.53	0.95–6.76	0.063	244	29	11.9	2.76	1.53–4.98	0.001
	T3	24	9	37.5	3.45	1.22–9.77	0.020	62	23	37.1	7.25	3.52–14.93	<0.001	24	7	29.2	7.91	2.02–31.0	0.003	62	13	21.0	12.4	4.38–35.2	<0.001
	T4	12	6	50.0	13.18	2.83–61.48	0.001	19	8	42.1	19.71	4.91–79.13	<0.001	12	5	41.7	94.9	11.8–766	<0.001	19	4	21.1	31.7	3.96–253	0.001
Nodal status	N0	100	17	17.0	Ref.			387	51	13.2	Ref.			100	11	11.0	Ref.			387	19	4.9	Ref.		
	N1	78	25	32.1	2.35	1.27–4.38	0.007	256	64	25.0	2.17	1.49–3.16	<0.001	78	15	19.2	2.12	0.96–4.65	0.062	256	38	14.8	3.40	2.00–5.80	<0.001
	N2	8	2	25.0	1.62	0.29–9.12	0.587	30	9	30.0	5.19	1.77–15.21	0.003	8	1	12.5	1.26	0.13–11.8	0.842	30	4	13.3	6.09	1.16–32.1	0.033
	N3	5	2	40.0	5.56	0.56–55.54	0.144	7	4	57.1	141.7	12.93–1553	<0.001	5	1	20.0	2.94	0.17–51.1	0.459	7	3	42.9	-	-	-
Tumor stage	I	52	7	13.5	Ref.			282	28	9.9	Ref.			52	2	3.8	Ref.			282	8	2.8	Ref.		
	II	97	23	23.7	1.78	0.85–3.74	0.128	310	65	21.0	2.26	1.50–3.39	<0.001	97	15	15.5	2.86	1.06–7.71	0.038	310	37	11.9	3.61	2.01–6.48	<0.001
	III	42	16	38.1	3.25	1.41–7.46	0.006	88	35	39.8	10.35	5.53–19.38	<0.001	42	11	26.2	5.96	1.97–19.0	0.002	88	19	21.6	19.6	7.74–49.4	<0.001
Tumor grade	1	7	2	28.6	Ref.			91	6	6.6	Ref.			7	1	14.3	Ref.			91	1	1.1	Ref.		
	2	59	16	27.1	1.01	0.23–4.38	0.986	311	59	19.0	2.21	1.24–3.92	0.007	59	9	15.3	1.20	0.18–8.14	0.856	311	21	6.8	2.75	1.00–7.57	0.051
	3	125	28	22.4	0.89	0.20–4.01	0.880	278	63	22.7	2.55	1.50–4.33	0.001	125	18	14.4	1.16	0.18–7.59	0.874	278	42	15.1	3.44	1.74–6.79	<0.001
ER status	pos	60	23	38.3	2.33	1.25–4.33	0.008	445	74	16.6	0.66	0.46–0.95	0.027	60	13	21.7	1.82	0.83–4.00	0.136	445	28	6.3	0.37	0.22–0.62	<0.001
	neg	131	23	17.6	Ref.			235	54	23.0	Ref.			131	15	11.5	Ref.			235	36	15.3	Ref.		
PR status	pos	62	22	35.5	2.01	1.09–3.73	0.026	428	76	17.8	0.79	0.55–1.13	0.200	62	10	16.1	1.07	0.49–2.34	0.861	428	30	7.0	0.46	0.27–0.77	0.003
	neg	129	24	18.6	Ref.			252	52	20.6	Ref.			129	18	14.0	Ref.			252	34	13.5	Ref.		
HER-2 status	pos	13	4	30.8	1.57	0.47–5.25	0.464	164	34	20.7	1.21	0.86–2.08	0.369	13	2	15.4	1.18	0.25–5.50	0.832	164	19	11.6	1.46	0.81–2.61	0.205
	neg	178	42	23.6	Ref.			516	94	18.2	Ref.			178	26	14.6	Ref.			516	45	8.7	Ref.		
TNBC	yes	114	20	17.5	0.50	0.28–0.90	0.021	138	30	21.7	1.34	0.86–2.08	0.197	114	14	12.3	0.70	0.33–1.48	0.350	138	21	15.2	2.40	1.29–4.46	0.006
	no	77	26	33.8	Ref.			542	98	18.1	Ref.			77	14	18.2	Ref.			542	43	7.9	Ref.		

Pts No. – number of patients; Ev No. – number of events; Ev % - percentage of events; HR – hazard ratio; 95% CI – 95% confidential interval; pre – premenopausal; post – postmenopausal; pos – positive; neg – negative; Ref. – reference.

Table S4. Analysis of 10-year DFS and DSS using a univariate Cox regression model comparing variables within subgroups of *BRCA1/2* mutation carriers and non-carriers.

Variable	Category	Disease Free Survival (DFS) Analysis						Disease Specific Survival (DSS) Analysis					
		<i>BRCA1/2</i> Carriers			Non-Carriers			<i>BRCA1/2</i> Carriers			Non-Carriers		
		HR	95% CI	<i>p</i>	HR	95% CI	<i>p</i>	HR	95% CI	<i>p</i>	HR	95% CI	<i>p</i>
Age at diagnosis	continuous	1.01	0.98–1.04	0.564	0.96	0.95–0.98	<0.001	1.00	0.96–1.04	0.975	0.96	0.93–0.98	0.002
	<35 vs ≥35 (Ref.)	0.98	0.55–1.78	0.960	2.04	1.44–2.89	<0.001	1.16	0.55–2.45	0.703	2.13	1.30–3.47	0.003
	<45 vs ≥45 (Ref.)	0.84	0.44–1.63	0.612	2.44	1.60–3.71	<0.001	0.88	0.38–2.08	0.780	2.16	1.21–3.84	0.009
menopausal status	pre vs post (Ref.)	0.81	0.36–1.81	0.611	1.92	1.15–3.20	0.012	0.67	0.25–1.75	0.409	1.74	0.86–3.53	0.122
	T2–4 vs T1 (Ref.)	2.09	1.06–4.12	0.033	2.30	1.60–3.30	<0.001	3.92	1.36–11.3	0.011	3.18	1.84–5.49	<0.001
Tumor size	T3–4 vs T1–2 (Ref.)	2.43	1.31–4.51	0.005	2.82	1.88–4.23	<0.001	3.55	1.68–7.51	0.001	2.97	1.70–5.17	<0.001
	T4 vs T1–3 (Ref.)	3.00	1.27–7.08	0.012	2.76	1.35–5.65	0.005	4.34	1.64–11.4	0.003	2.73	0.99–7.50	0.052
Nodal status	N1–3 vs N0 (Ref.)	2.23	1.23–4.07	0.009	2.23	1.57–3.18	<0.001	1.97	0.92–4.21	0.080	3.43	2.01–5.87	<0.001
	N2–3 vs N0–1 (Ref.)	1.30	0.47–3.64	0.613	2.41	1.36–4.27	0.003	1.10	0.26–4.63	0.898	2.44	1.11–5.35	0.026
	N3 vs N0–2 (Ref.)	1.90	0.46–7.86	0.373	4.00	1.48–10.9	0.006	1.57	0.21–11.6	0.658	6.76	2.11–21.6	0.001
Tumor stage	II–III vs I (Ref.)	2.30	1.03–5.13	0.043	2.90	1.91–4.42	<0.001	5.26	1.25–22.2	0.024	5.55	2.65–11.7	<0.001
	III vs I–II (Ref.)	2.16	1.18–3.97	0.013	3.12	2.12–4.61	<0.001	2.58	1.21–5.50	0.015	3.22	1.88–5.51	<0.001
Tumor grade	2–3 vs 1 (Ref.)	0.94	0.23–3.88	0.933	3.44	1.52–7.82	0.003	1.20	0.16–8.81	0.860	9.97	1.38–71.9	0.023
	3 vs 1–2 (Ref.)	0.90	0.50–1.63	0.728	1.57	1.11–2.22	0.011	1.07	0.49–2.31	0.870	3.10	1.85–5.20	<0.001
ER status	pos vs neg (Ref.)	2.15	1.21–3.84	0.009	0.67	0.47–0.96	0.028	1.75	0.83–3.67	0.141	0.40	0.24–0.65	<0.001
PR status	pos vs neg (Ref.)	1.91	1.07–3.40	0.029	0.79	0.56–1.13	0.201	1.07	0.49–2.32	0.861	0.48	0.29–0.79	0.004
HER–2 status	pos vs neg (Ref.)	1.46	0.52–4.09	0.467	1.20	0.81–1.77	0.370	1.17	0.28–4.93	0.832	1.41	0.83–2.42	0.206
TNBC	yes vs no (Ref.)	0.51	0.28–0.91	0.024	1.31	0.87–1.97	0.198	0.70	0.34–1.48	0.352	2.05	1.22–3.45	0.007

Pts No. – number of patients; Ev No. – number of events; OR – odds ratio; 95% CI – 95% confidential interval; pre – premenopausal; post – postmenopausal; pos – positive; neg – negative; Ref. – reference.

Table S5. Analysis of 10-year DFS and DSS using multivariable Cox proportional-hazard models adjusted for age, menopausal status, stage, tumor grade and ER status.

	BRCA1/2 Carriers						Non-Carriers					
	DFS			DSS			DFS			DSS		
	HR	95%CI	p-Value	HR	95%CI	p-Value	HR	95%CI	p-Value	HR	95%CI	p-Value
model A												
Age (continuous)	1.01	0.97–1.06	0.552	0.99	0.94–1.04	0.661	0.96	0.93–0.98	0.001	0.95	0.92–0.99	0.014
Menopausal status (post vs pre)	1.04	0.32–3.38	0.948	0.41	0.10–1.77	0.234	0.60	0.27–1.32	0.205	0.44	0.14–1.34	0.148
Tumor stage (II–III vs I)	2.38	1.04–5.48	0.041	5.63	1.30–24.4	0.021	2.54	1.66–3.89	<0.001	4.56	2.16–9.63	<0.001
Tumor grade (3 vs 1–2)	1.20	0.64–2.27	0.571	1.36	0.59–3.17	0.470	1.10	0.75–1.63	0.629	1.92	1.09–3.38	0.025
ER–status (pos vs neg)	2.19	1.17–4.09	0.014	1.70	0.76–3.82	0.197	0.76	0.51–1.12	0.161	0.57	0.33–0.98	0.041
model B												
Age (<35 vs ≥35)	0.92	0.49–1.74	0.796	1.18	0.52–2.70	0.692	1.70	1.17–2.48	0.005	1.67	0.98–2.87	0.060
Menopausal status (post vs pre)	0.84	0.34–2.06	0.707	0.48	0.16–1.44	0.191	1.24	0.71–2.16	0.442	1.01	0.46–2.18	0.987
Tumor stage (II–III vs I)	2.33	1.02–5.35	0.045	5.75	1.34–24.8	0.019	2.61	1.71–4.01	<0.001	4.73	2.24–9.99	<0.001
Tumor grade (3 vs 1–2)	1.19	0.63–2.25	0.593	1.37	0.59–3.17	0.468	1.11	0.75–1.64	0.594	1.91	1.08–3.37	0.025
ER–status (pos vs neg)	2.19	1.17–4.11	0.015	1.70	0.76–3.82	0.200	0.76	0.51–1.12	0.169	0.57	0.33–0.97	0.039
model C												
Age (<45 vs ≥45)	0.92	0.38–2.23	0.847	1.14	0.34–3.83	0.830	2.41	1.30–4.48	0.005	1.86	0.81–4.27	0.141
menopausal status (post vs pre)	0.87	0.29–2.60	0.801	0.47	0.14–1.88	0.287	0.76	0.36–1.62	0.481	0.78	0.29–2.14	0.632
Tumor stage (II–III vs I)	2.33	1.02–5.32	0.046	5.76	1.34–24.8	0.019	2.53	1.65–3.88	0.000	4.57	2.16–9.67	<0.001
Tumor grade (3 vs 1–2)	1.18	0.63–2.23	0.602	1.38	0.60–3.19	0.452	1.12	0.76–1.65	0.578	1.93	1.09–3.41	0.023
ER–status (pos vs neg)	2.17	1.16–4.05	0.015	1.73	0.77–3.88	0.181	0.74	0.50–1.10	0.137	0.56	0.32–0.95	0.033
model D												
Age (continuous)	1.01	0.96–1.05	0.758	0.98	0.92–1.03	0.426	0.96	0.93–0.98	0.001	0.95	0.91–0.99	0.008
Menopausal status (post vs pre)	1.00	0.29–3.39	0.995	0.37	0.08–1.75	0.209	0.65	0.30–1.41	0.276	0.47	0.15–1.42	0.181
Tumor stage (III vs I–II)	1.93	1.01–3.67	0.047	2.50	1.11–5.63	0.026	2.99	2.00–4.47	0.000	2.78	1.60–4.85	<0.001
Tumor grade (3 vs 1–2)	1.10	0.57–2.11	0.775	1.15	0.48–2.72	0.755	1.02	0.69–1.52	0.918	1.81	1.01–3.24	0.047
ER–status (pos vs neg)	1.97	1.02–3.78	0.043	1.48	0.64–3.45	0.361	0.66	0.45–0.98	0.039	0.50	0.29–0.87	0.014
model E												
Age (<35 vs ≥35)	1.03	0.54–1.94	0.937	1.32	0.58–3.03	0.505	1.63	1.12–2.39	0.011	1.59	0.92–2.73	0.094
Menopausal status (post vs pre)	0.86	0.35–2.11	0.735	0.52	0.17–1.57	0.245	1.43	0.82–2.48	0.203	1.18	0.55–2.55	0.673
Tumor stage (III vs I–II)	1.93	1.01–3.68	0.047	2.56	1.14–5.75	0.023	2.90	1.94–4.34	0.000	2.70	1.55–4.71	<0.001
Tumor grade (3 vs 1–2)	1.08	0.56–2.08	0.811	1.15	0.49–2.72	0.752	1.06	0.71–1.57	0.773	1.88	1.05–3.37	0.033
ER–status (pos vs neg)	1.96	1.02–3.78	0.044	1.46	0.63–3.41	0.378	0.69	0.47–1.02	0.061	0.52	0.30–0.90	0.019
model F												
Age (<45 vs ≥45)	0.90	0.36–2.24	0.828	1.19	0.32–4.37	0.795	2.73	1.47–5.09	0.002	2.21	0.96–5.09	0.064
Menopausal status (post vs pre)	0.94	0.31–2.87	0.913	0.51	0.12–2.28	0.381	0.78	0.36–1.65	0.510	0.77	0.28–2.15	0.624
Tumor stage (III vs I–II)	1.94	1.01–3.69	0.045	2.50	1.11–5.65	0.027	3.10	2.07–4.63	0.000	2.82	1.62–4.90	<0.001
Tumor grade (3 vs 1–2)	1.09	0.57–2.08	0.794	1.19	0.51–2.79	0.691	1.06	0.71–1.57	0.782	1.90	1.06–3.40	0.031
ER–status (pos vs neg)	1.96	1.02–3.77	0.045	1.50	0.64–3.49	0.352	0.66	0.45–0.98	0.038	0.51	0.29–0.87	0.015

Please add footnote here to explain the bold text in the table.

Table S6. The indication criteria for genetic testing of hereditary BC/OC predisposition in the Czech Republic.

Familial Indication Criteria (IC):
<ul style="list-style-type: none"> • ≥ 3 first and/or second-degree relatives with BC (including the proband) • 2 first and/or second-degree relatives with BC (1 diagnosed at ≤ 50 years, or both diagnosed at ≤ 60 years, or one with male BC; including the proband) • BC and OC in first and/or second-degree relatives (including the proband), or BC ≤ 50 years and PaC (including the proband) • ≥ 2 first and/or second-degree relatives with OC and PaC (including the proband) • ≥ 2 first and/or second-degree relatives with OC (including the proband)
Personal (Individual) Indication Criteria:
<ul style="list-style-type: none"> • BC diagnosed at ≤ 45 years (or at ≤ 50 years, when family history is unknown or limited) <ul style="list-style-type: none"> • BC duplicity (first cancer diagnosed at ≤ 50 years or both cancers at ≤ 60 years) <ul style="list-style-type: none"> • “Triple negative” BC diagnosed at ≤ 60 years <ul style="list-style-type: none"> • Medullary BC • Male BC • Duplicity of BC and OC • Duplicity of BC and PaC • OC at any age

BC – breast cancer; OC – ovarian cancer (including the fallopian tube and primary peritoneal cancer); PaC – pancreatic cancer; close blood relatives include first-, second-, and/or third-degree relatives from the same family side. *Note:* The Czech national guidelines are based on NCCN guidelines (www.nccn.org) and since 2003 updated by the consensus of the expert panel constituted from the members of Czech Society for Oncology (www.linkos.cz) and Society of Medical Genetics and Genomics (www.slg.cz) [1–5]. The guidelines used for the enrollment of high-risk patients before 2003 were described by Pohlreich et al. [6,7].

Reference

- Bartonkova, H.; Foretová, L.; Helmichová, E.; Kalábová, R.; Kleibl, Z.; Konopásek, B.; Krutílková, V.; Machackova, E.; Novotny, J.; Petráková, K. Recommendations for care of patients with breast and ovarian cancer and healthy individuals with germline mutations in BRCA1 or BRCA2. *Klin. Onkol.* **2003**, *16*, 28–34.
- Plevová, P.; Novotný, J.; Petráková, K.; Palácová, M.; Kalábová, R.; Schneiderová, M.; Foretová, L.; Hereditary Breast and Ovarian Cancer Syndrome. *Klin. Onkol.* **2009**, *22*, 8–11.
- Pohlreich, P.; Kleibl, Z.; Kleiblová, P.; Janatová, M.; Soukupová, J.; Macháčková, E.; Házová, J.; Vašíčková, P.; Sťahlová, H.E.; Navrátilová, M.; et al. The Clinical Importance of a Genetic Analysis of Moderate-Risk Cancer Susceptibility Genes in Breast and Other Cancer Patients from the Czech Republic. *Klin. Onkol.* **2012**, *25 Suppl 1*, S59–S66.
- Janatova, M.; Borecka, M.; Soukupova, J.; Kleiblova, P.; Stribrna, J.; Vocka, M.; Zemánková, P.; Panczak, A.; Veselá, K.; Souček, P.; et al. PALB2 as Another Candidate Gene for Genetic Testing in Patients with Hereditary Breast Cancer in Czech Republic. *Klin. Onkol.* **2016**, *29 Suppl 1*, 31–34.
- Foretova, L.; Machackova, E.; Palacova, M.; Navratilova, M.; Svoboda, M.; Petrakova, K. Recommended Extension of Indication Criteria for Genetic Testing of BRCA1 and BRCA2 Mutations in Hereditary Breast and Ovarian Cancer Syndrome. *Klin. Onkol.* **2016**, *29 Suppl 1*, S9–S13.
- Pohlreich, P.; Stribrna, J.; Kleibl, Z.; Zikan, M.; Kalbacova, R.; Petruzelka, L.; Konopasek, B. Mutations of the BRCA1 gene in hereditary breast and ovarian cancer in the Czech Republic. *Med. Princ. Pract.* **2003**, *12*, 23–29.
- Pohlreich, P.; Stribrna, J.; Zikan, M.; Kleibl, Z.; Matous, B.; Novotny, J. Analysis of BRCA1 and BRCA2 mutations in high-risk patients from the Prague-area. *Eur. J. Cancer Suppl.* **2005**, *3*, 89.



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