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#### Cancer Risks Associated With BRCA1 and BRCA2 Pathogenic Variants

Li, Shuai; Silvestri, Valentina; Leslie, Goska; Rebbeck, Timothy R.; Neuhausen, Susan L.; Hopper, John L.; Nielsen, Henriette Roed; Lee, Andrew; Yang, Xin; McGuffog, Lesley; Parsons, Michael T.; Andrulis, Irene L.; Arnold, Norbert; Belotti, Muriel; Borg, Ake; Buecher, Bruno; Buys, Saundra S.; Caputo, Sandrine M.; Chung, Wendy K.; Colas, Chrystelle; Colonna, Sarah V.; Cook, Jackie; Daly, Mary B.; de la Hoya, Miguel; de Pauw, Antoine; Delhomelle, Hélène; Eason, Jacqueline; Engel, Christoph; Evans, D. Gareth; Faust, Ulrike; Fehm, Tanja N.; Fostira, Florentia; Fountzilas, George; Frone, Megan; Garcia-Barberan, Vanesa; Garre, Pilar; Gauthier-Villars, Marion; Gehrig, Andrea; Glendon, Gord; Goldgar, David E.; Golmard, Lisa; Greene, Mark H.; Hahnen, Eric; Hamann, Ute; Hanson, Helen; Hassan, Tiara; Hentschel, Julia; Horvath, Judit; Izatt, Louise; Janavicius, Ramunas; Jiao, Yue; John, Esther M.; Karlan, Beth Y.; Kim, Sung-Won; Konstantopoulou, Irene; Kwong, Ava; Laugé, Anthony; Lee, Jong Won; Lesueur, Fabienne; Mebirouk, Noura; Meindl, Alfons; Mouret-Fourme, Emmanuelle; Musgrave, Hannah; Ngeow Yuen Yie, Joanne; Niederacher, Dieter; Park, Sue K.; Pedersen, Inge Sokilde; Ramser, Juliane; Ramus, Susan J.; Rantala, Johanna; Rashid, Muhammad U.; Reichl, Florian; Ritter, Julia; Rump, Andreas; Santamariña, Marta; Saule, Claire; Schmidt, Gunnar; Schmutzler, Rita K.; Senter, Leigha; Shariff, Saba; Singer, Christian F.; Southey, Melissa C.; Stoppa-Lyonnet, Dominique; Sutter, Christian; Tan, Yen; Teo, Soo Hwang; Terry, Mary Beth; Thomassen, Mads; Tischkowitz, Marc; Toland, Amanda E.; Torres, Diana; Vega, Ana; Wagner, Sebastian A.; Wang-Gohrke, Shan; Wappenschmidt, Barbara; Weber, Bernhard H. F.; Yannoukakos, Drakoulis; Spurdle, Amanda B.; Easton, Douglas F.; Chenevix-Trench, Georgia; Ottini, Laura; Antoniou, Antonis

Published in: Journal of Clinical Oncology

DOI (link to publication from Publisher): 10.1200/JCO.21.02112

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Publication date: 2022

Document Version
Publisher's PDF, also known as Version of record

Link to publication from Aalborg University

# original report

## Cancer Risks Associated With *BRCA1* and *BRCA2* Pathogenic Variants

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**PURPOSE** To provide precise age-specific risk estimates of cancers other than female breast and ovarian cancers associated with pathogenic variants (PVs) in *BRCA1* and *BRCA2* for effective cancer risk management.

**METHODS** We used data from 3,184 *BRCA1* and 2,157 *BRCA2* families in the Consortium of Investigators of Modifiers of *BRCA1/2* to estimate age-specific relative (RR) and absolute risks for 22 first primary cancer types adjusting for family ascertainment.

**RESULTS** *BRCA1* PVs were associated with risks of male breast (RR = 4.30; 95% CI, 1.09 to 16.96), pancreatic (RR = 2.36; 95% CI, 1.51 to 3.68), and stomach (RR = 2.17; 95% CI, 1.25 to 3.77) cancers. Associations with colorectal and gallbladder cancers were also suggested. *BRCA2* PVs were associated with risks of male breast (RR = 44.0; 95% CI, 21.3 to 90.9), stomach (RR = 3.69; 95% CI, 2.40 to 5.67), pancreatic (RR = 3.34; 95% CI, 3.21 to 3.21

**CONCLUSION** In addition to female breast and ovarian cancers, *BRCA1* and *BRCA2* PVs are associated with increased risks of male breast, pancreatic, stomach, and prostate (only *BRCA2* PVs) cancers, but not with the risks of other previously suggested cancers. The estimated age-specific risks will refine cancer risk management in men and women with *BRCA1/2* PVs.

J Clin Oncol 40:1529-1541. © 2022 by American Society of Clinical Oncology

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#### **Data Supplement**

Author affiliations and support information (if applicable) appear at the end of this article.

Accepted on December 20, 2021 and published at ascopubs.org/journal/jco on January 25, 2022: DOI https://doi.org/10.1200/JC0.21.

#### INTRODUCTION

It is well established that pathogenic variants (PVs) in *BRCA1* and *BRCA2* (*BRCA1/2*) are associated with increased risks of breast and ovarian cancers in women for which reliable risk estimates are available. Accumulated evidence indicates that *BRCA1/2* PVs are also

associated with pancreatic cancer<sup>2-8</sup> and male breast cancer risks<sup>3,6,9-13</sup> and that *BRCA2* PVs are associated with prostate cancer risk, particularly aggressive prostate cancer, whereas the association between *BRCA1* PVs and prostate cancer risk is still debated.<sup>2,5,6,8,14-17</sup> Associations with risks for other cancers have also been



#### CONTEXT

#### **Key Objective**

The associations of pathogenic variants (PVs) in *BRCA1* and *BRCA2* with cancers other than female breast and ovarian cancers remain uncertain. Precise risk estimates are required to inform effective cancer risk management. This study investigates the associations between the risks of 22 cancers and *BRCA1/2* PVs using data from 5,341 families segregating *BRCA1* or *BRCA2* PVs.

#### **Knowledge Generated**

BRCA1 and BRCA2 PVs are associated with increased risks of male breast, pancreatic, and stomach cancers; male BRCA2 carriers are also at increased prostate cancer risk. No associations were found with risks of other cancers. The cumulative risks to age 80 years ranged from 0.4% for male breast cancer to approximately 2.5% for pancreatic cancer for BRCA1 carriers and from approximately 2.5% for pancreatic cancer to 27% for prostate cancer for BRCA2 carriers.

#### Relevance

The findings provide age-specific cancer risk estimates and will allow for improved cancer risk assessment of male and female carriers.

suggested, including colorectal, liver, and stomach cancers for *BRCA1/2* PVs; cervical, corpus uteri, kidney, and testis cancers for *BRCA1* PVs;<sup>3,4,6,8,18,19</sup> and bone, brain, blood, and gallbladder cancers and malignant melanoma for *BRCA2* PVs.<sup>2,5,6,8,20</sup> However, these associations are based on studies with relatively small sample sizes, resulting in imprecise cancer risk estimates.

The National Comprehensive Cancer Network and other guidelines recommend breast and ovarian cancer screening for BRCA1/2 carriers and prostate cancer screening particularly for BRCA2 carriers. Notably, National Comprehensive Cancer Network guidelines recently addressed testing and management for pancreatic cancer risk in BRCA1/2 carriers, but only in the presence of a positive family history of the disease. 21,22 Overall, current guidelines suggest that men and women with BRCA1/2 PVs should consider participation in investigational screening studies and receive education regarding signs and symptoms of cancers possibly associated with BRCA1/2 PVs.21 The availability of more precise risk estimates will aid translation into evidence-based clinical guidelines for the cancer risk management in BRCA1/2 carriers and may guide treatment options for patients with cancer.

To inform clinical management strategies and optimize guidelines for cancer risk management in female and male *BRCA1/2* carriers, we comprehensively assess the associations of *BRCA1/2* PVs with risks of 22 cancers, other than female breast and ovarian cancers.

#### **METHODS**

#### **Study Sample**

Data on 7,618 families with at least one family member having a *BRCA1* or *BRCA2* PV were obtained from 26 study groups in the Consortium of Investigators of Modifiers of *BRCA1*/2 (Data Supplement, online only).<sup>23</sup> Only families with a clearly PV identified were included.<sup>24</sup> The majority of families (7,281)

were ascertained through an index individual attending cancer family clinics, mainly because of having multiple affected relatives, and 337 families were ascertained through an index case with breast or ovarian cancer, unselected for family history. All index individuals were age  $\geq 18$  years. For each family member, data including familial relationship, BRCA1/2 PV status, sex, year of birth, and years or age at pedigree data collection, death, and cancer diagnoses were collected (Data Supplement). All participants provided written informed consent and participated in studies at the host institutions under ethically approved protocols.

#### Statistical Analysis

BRCA1 and BRCA2 families were analyzed separately. Complex segregation analysis,<sup>25</sup> which considered the observed phenotype and observed or inferred genotype information of all family members, was used to estimate relative risks (RRs) for 22 first primary cancer sites, excluding female breast and ovarian cancers (Table 1). This involved comparing the observed cancer incidences for carriers with the age-, country- and birth cohort-specific population incidences (Cancer Incidence in Five Continents<sup>26</sup>); thus, the estimated RRs were equivalent to standardized incidence ratios. Noncarriers were assumed to develop the cancers according to population incidences. Pedigree likelihoods were constructed and maximized using the pedigree analysis software MENDEL.<sup>27</sup>

Individuals were followed from birth until the age of the first primary cancer diagnosis, death, age at pedigree-data collection, risk-reducing mastectomy and/or salpingo-oophorectomy (if these occurred at least 1 year before breast or ovarian cancer diagnoses, respectively), or age 80 years, whichever occurred first. Missing year of birth and cancer diagnosis age were imputed (Data Supplement).

Each individual was assumed to be at risk of developing the cancer of interest, as well as breast or ovarian cancer. The RRs for female breast and ovarian cancers were assumed

**TABLE 1.** No. of First Primary Cancer Cases in the Informative *BRCA1* and *BRCA2* Families

	BRCA1 Families, No.						BRCA2 Families, No.							
			Males			Females				Males			Females	
Cancer Site	Total	Carriers (n = 1,508)	Noncarriers (n = 1,716)	Untested (n = 44,396)	Carriers (n = 7,376)	Noncarriers (n = 4,154)	Untested (n = 40,801)	Total	Carriers (n = 1,063)	Noncarriers (n = 1,064)	Untested (n = 30,032)	Carriers (n = 5,032)	Noncarriers (n = 2,371)	Untested (n = 28,094)
Bladder	123	6	6	79	1	5	26	72	5	1	48	4	2	12
Brain and CNS	186	5	1	105	1	1	73	156	0	1	82	2	3	68
Breast	9,389	17	3	26	3,648	271	5,424	7,143	82	4	133	2,612	205	4,107
Cervix uteri	187	0	0	0	34	20	133	125	0	0	0	26	10	89
Colon-rectum	726	20	14	360	20	13	299	490	12	8	240	3	10	217
Connective and soft tissue	20	1	0	7	1	0	11	11	0	0	4	1	0	6
Corpus uteri	120	0	0	0	5	4	111	50	0	0	0	3	3	44
Esophagus	88	1	1	64	1	0	21	69	1	1	52	0	0	15
Eye	10	1	0	5	0	0	4	11	1	1	5	1	1	2
Gallbladder and extrahepatic ducts	27	0	0	11	0	1	15	18	0	0	9	2	1	6
Head and neck	226	9	4	161	1	1	50	158	5	3	114	3	0	33
Kidney	117	3	2	82	2	0	28	76	3	2	50	1	1	19
Leukemia	198	2	3	101	2	1	89	150	3	1	76	3	1	66
Lung	746	13	6	567	2	5	153	504	6	7	376	4	0	111
Lymphoma	134	6	6	71	3	3	45	80	5	4	35	2	4	30
Melanoma	174	11	10	71	19	28	35	96	8	11	33	12	11	21
Multiple myeloma	14	1	0	5	1	2	5	10	0	0	8	0	0	2
Ovary	2,743	0	0	0	885	28	1,830	827	0	0	0	293	18	516
Pancreas	252	9	2	146	4	1	90	266	12	2	151	8	2	91
Prostate	686	34	64	588	0	0	0	685	71	31	583	0	0	0
Stomach	463	5	0	263	0	0	195	387	5	2	243	4	0	133
Testis	47	1	2	44	0	0	0	38	4	1	33	0	0	0
Thyroid	58	1	0	12	9	8	28	47	0	0	11	14	6	16
Affected by any cancer	16,500	144	123	2,762	4,577	391	8,503	11,354	221	80	2,277	2,976	275	5,525
Unaffected	83,451	1,364	1,593	41,634	2,799	3,763	32,298	56,300	842	984	27,755	2,055	2,095	22,569

TABLE 2. Primary Cancer RRs and 95% Cls for BRCA1 and BRCA2 Carriers From the Main Analysis

**BRCA1** Carriers **BRCA2** Carriers **Cancer Site** RR (95% CI) P RR (95% CI) Ρ Age, years Bladder 40-79 0.88 (0.33 to 2.36) .80 1.71 (0.75 to 3.89) .20 Brain and CNS 20-79 1.15 (0.52 to 2.55) .73 1.10 (0.42 to 2.87) .85 Male breast 30-79 4.30 (1.09 to 16.96) 04 44.03 (21.32 to 90.93) < .001 20-79 .18 Cervix uteri 1.45 (0.85 to 2.49) 1.61 (0.86 to 3.04) .14 Colon-rectum 30-79 1.48 (1.01 to 2.16) .04 1.30 (0.80 to 2.11) .29 Connective and soft tissue 30-79 0.80 (0.07 to 8.71) .86 0.17 (0 to 25.94) .49 .95 Corpus uteri 40-79 0.97 (0.35 to 2.70) 0 (0 to 3.2E+280) .94 40-79 Esophagus 0.96 (0.35 to 2.65) .93 0.85 (0.29 to 2.49) .77 .65 Eye 30-79 1.56 (0.23 to 10.77) 4.60 (1.00 to 21.16) .05 Gallbladder and extrahepatic ducts .01 40-79 3.34 (1.34 to 8.28) 2.28 (0.77 to 6.70) .14 40-79 .78 0.71 (0.18 to 2.86) Head and neck 1.13 (0.49 to 2.62) .63 .19 Kidney 40-79 1.84 (0.74 to 4.56) 0.26 (0.01 to 6.20) .41 Leukemia 20-79 0.90 (0.36 to 2.26) .82 0.91 (0.29 to 2.85) .87 40-79 .19 .68 Lung 1.37 (0.85 to 2.21) 1.13 (0.63 to 2.03) Lymphoma 20-79 1.03 (0.33 to 3.22) .96 0.97 (0.16 to 5.87) .97 40-79 56 0.93 (0.26 to 3.25) 91 Melanoma 0.64 (0.14 to 2.95) Multiple myeloma 30-79 3.06 (0.83 to 11.26) .09 0.84 (0.10 to 7.31) .87 30-79 < .001 3.34 (2.21 to 5.06) < .001 Pancreas 2.36 (1.51 to 3.68) 40-79 2.22 (1.63 to 3.03) Prostate 0.82 (0.54 to 1.27) .38 < .001 Stomach 30-79 2.17 (1.25 to 3.77) .01 3.69 (2.40 to 5.67) < .001 **Testis** 20-79 0.07 (0 to 1.63) .10 2.17 (0.82 to 5.70) .12 Thyroid 30-79 0.14 (0.01 to 1.55) 11 0.84 (0.22 to 3.24) .80

Abbreviation: RR, relative risk.

to be equal to previous estimates<sup>28</sup>; therefore, we only estimated the RR for the cancer of interest. We fitted models in which the RRs were assumed to be constant with age, birth cohort, sex, and study group and separate models with sex-specific RRs. For cancers with significant associations, we investigated whether the RRs varied by age. RRs from the best fitting models were used to estimate age-specific absolute risks on the basis of UK cancer incidences in year 2008-2012 (Data Supplement).

Because family ascertainment varied across study groups, we adjusted for the ascertainment of each family separately using an ascertainment-assumption-free approach. Pedigree likelihoods were computed conditional on any data that may be relevant to the ascertainment (Data Supplement). Non-informative families, in which no additional information beyond the data relevant to the ascertainment was available, were excluded from analysis. Since cancer family history was self-reported, we assessed the possibility of systematic underreporting of specific cancers at the individual study group level and excluded any study groups in which under-reporting was likely relative to the population incidences (Data Supplement).

Sensitivity analyses under alternative inclusion, censoring, or ascertainment assumptions were performed for cancers

that demonstrated associations: (1) stratifying by geographical region (Asian countries v others); (2) including study groups with possible cancer under-reporting; (3) excluding individuals with missing age at diagnosis; (4) individuals with risk-reducing bilateral mastectomy and/or salpingo-oophorectomy were still considered to be at risk of developing the other cancers, except breast and ovarian cancers; and (5) assuming the data relevant to the ascertainment for clinic-based families do not include the family history of cancer of interest. To account for population differences in melanoma skin pigmentation, we also conducted sensitivity analyses for melanoma by using (1) only families from Australia, Northern Europe, and North America; (2) only families in which probands self-identified as White European; and (3) only the families satisfying both (1) and (2).

All statistical tests were two-sided, and associations with a nominal P < .05 were considered statistically significant.

#### **RESULTS**

After ascertainment adjustment, 3,184 *BRCA1* families and 2,157 *BRCA2* families were informative for inclusion in the analysis, including 14,979 carriers, 9,296 noncarriers,

TABLE 3. Sex-Specific RRs and 95% CIs for BRCA1 and BRCA2 Carriers From the Main Analysis

		BRCA1 Carriers		BRCA2 Carriers				
Cancer Site	Male RR (95% CI)	Female RR (95% CI)	<i>P</i> for Difference <sup>a</sup>	Male RR (95% CI)	Female RR (95% CI)	<i>P</i> for Difference <sup>a</sup>		
Bladder	0.97 (0.34 to 2.78)	0.53 (0.05 to 5.95)	.61	1.26 (0.46 to 3.47)	4.07 (1.09 to 15.21)	.20		
Brain and CNS	0.72 (0.25 to 2.06)	2.56 (0.98 to 6.67)	.11	0.48 (0.10 to 2.25)	2.27 (0.83 to 6.21)	.09		
Colon-rectum	1.54 (0.98 to 2.42)	1.34 (0.66 to 2.73)	.74	1.57 (0.90 to 2.74)	0.89 (0.36 to 2.20)	.28		
Connective and soft tissue	0.08 (0 to 196.37)	1.61 (0.15 to 16.78)	.36	0 (0 to 3.5E+122)	1.33 (0 to 3,851.9)	.53		
Esophagus	0.88 (0.29 to 2.70)	1.63 (0.13 to 20.17)	.68	1.12 (0.37 to 3.42)	0.07 (0 to 3.18)	.13		
Eye	1.98 (0.15 to 25.33)	NA	NA	3.26 (0.29 to 36.23)	6.19 (0.71 to 54.34)	.70		
Gallbladder and extrahepatic ducts	3.75 (1.23 to 11.43)	2.52 (0.36 to 17.56)	.71	2.35 (0.59 to 9.35)	2.20 (0.49 to 9.92)	.95		
Head and neck	1.04 (0.41 to 2.64)	1.69 (0.29 to 9.93)	.65	0.71 (0.19 to 2.73)	0.83 (0 to 474.33)	.96		
Kidney	1.35 (0.36 to 5.06)	3.10 (0.74 to 12.93)	.41	0.19 (0.01 to 4.46)	3.13 (0.37 to 26.16)	.27		
Leukemia	1.03 (0.36 to 2.92)	NA	NA	0.77 (0.23 to 2.60)	1.85 (0.30 to 11.57)	.48		
Lung	1.36 (0.79 to 2.33)	1.43 (0.49 to 4.22)	.93	0.81 (0.39 to 1.69)	2.84 (1.23 to 6.60)	.05		
Lymphoma	0.69 (0.12 to 3.91)	1.56 (0.33 to 7.43)	.49	0.78 (0.09 to 6.37)	2.24 (0.13 to 39.84)	.64		
Melanoma	0.44 (0.04 to 5.44)	0.80 (0.13 to 5.06)	.70	NA	1.82 (0.43 to 7.71)	NA		
Multiple myeloma	3.60 (1.00 to 12.96)	2.04 (0.22 to 18.87)	.66	1.11 (0.13 to 9.46)	0.01 (0 to 19.48)	.52		
Pancreas	1.92 (1.12 to 3.28)	4.27 (2.01 to 9.05)	.11	2.96 (1.78 to 4.94)	4.34 (2.19 to 8.62)	.38		
Stomach	1.67 (0.86 to 3.27)	4.86 (2.13 to 11.08)	.08	2.76 (1.59 to 4.80)	6.89 (3.71 to 12.78)	.04		
Thyroid	0.05 (0 to 4,319.91)	0.14 (0.01 to 1.78)	.88	NA	1.01 (0.25 to 4.19)	.31		

Abbreviations: NA, No. of cancers too small to obtain a sex-specific estimate; RR, relative risk.

and 153,323 untested individuals (Data Supplement). 61.3% of probands had self-reported ethnicity data. Of those, 77.0%, 11.5%, 4.7%, 3.3%, and 1.2% self-identified as White European, Asian, Ashkenazi Jewish, Hispanic, and Black, respectively. Prostate, lung, colorectal, stomach, and pancreatic cancers were the most common cancers in the data set, aside from breast and ovarian (Table 1). The age at diagnosis for each cancer by PV status is shown in the Data Supplement. After excluding study groups in which there was potential cancer under-reporting (Data Supplement), the proportions of families included in the estimation of cancerspecific risks varied from approximately 15% for lymphoma and multiple myeloma to > 90% for pancreatic and male breast cancers (Data Supplement).

#### Cancer Associations With BRCA1 PVs

*BRCA1* PVs were associated with male breast (RR = 4.30; 95% CI, 1.09 to 16.96), gallbladder (RR = 3.34; 95% CI, 1.34 to 8.28), pancreatic (RR = 2.36; 95% CI, 1.51 to 3.68), stomach (RR = 2.17; 95% CI, 1.25 to 3.77), and colorectal (RR = 1.48; 95% CI, 1.01 to 2.16) cancers (Table 2). No association was found for prostate cancer (RR = 0.82; 95% CI, 0.54 to 1.27). No difference in the RR estimates by sex was observed for any of the 17 non–sex-specific cancers (all P > .07; Table 3).

A model with RRs stratified by age 65 years (Data Supplement) provided a significantly better fit for stomach cancer: RR = 3.50 (95% CI, 2.01 to 6.10) for age < 65 years and higher than 0.61 (95% CI, 0.16 to 2.30) for age  $\geq$  65 years (*P*-heterogeneity = .01). For male breast cancer, a model with RRs stratified by age decade provided a better fit than the model with an age-constant RR (P = .03), but this was mainly driven by the lack of cases in the age group of 50-59 years (Data Supplement).

#### Cancer Associations With BRCA2 PVs

*BRCA2* PVs were associated with increased risks of male breast (RR = 44.0; 95% CI, 21.3 to 90.9), stomach (RR = 3.69; 95% CI, 2.40 to 5.67), pancreatic (RR = 3.34; 95% CI, 2.21 to 5.06), and prostate (RR = 2.22; 95% CI, 1.63 to 3.03) cancers (Table 2). Female carriers had a higher risk of stomach cancer (RR = 6.89; 95% CI, 3.71 to 12.78) than male carriers (RR = 2.76; 95% CI, 1.59 to 4.80; P-heterogeneity = .04; Table 3).

A model with RRs stratified by age 65 years (Data Supplement) provided a significantly better fit for pancreatic cancer: RR = 4.92 (95% CI, 2.96 to 7.80) for age <65 years and higher than 1.77 (95% CI, 0.87 to 3.58) for age  $\geq65$  years (*P*-heterogeneity = .03). There was a suggestion that the prostate cancer RR was greater for

<sup>&</sup>lt;sup>a</sup>P value by comparing the model of the same RR between males and females with its nested model of sex-specific RR.

TABLE 4. Age-Specific Absolute Risks (%) and 95% Cls of Primary Cancers With Significant Associations for BRCA1 and BRCA2 Carriers<sup>a</sup>

Cancer Site	Sex	Age 50 Years	Age 60 Years	Age 70 Years	Age 80 Years
Absolute risk (95% CI) for BRCA1 carriers					
Breast	Male	0.02 (0.01 to 0.08)	0.07 (0.02 to 0.3)	0.2 (0.05 to 0.7)	0.4 (0.1 to 1.5)
Pancreas	Male	0.1 (0.07 to 0.2)	0.4 (0.3 to 0.7)	1.3 (0.8 to 2.0)	2.9 (1.9 to 4.5)
	Female	0.08 (0.05 to 0.1)	0.3 (0.2 to 0.5)	1.0 (0.6 to 1.5)	2.3 (1.5 to 3.6)
Stomach	Male	0.2 (0.1 to 0.3)	0.6 (0.3 to 1.0)	1.1 (0.6 to 2.2)	1.6 (0.7 to 4.0)
	Female	0.1 (0.06 to 0.2)	0.3 (0.2 to 0.5)	0.5 (0.3 to 0.9)	0.7 (0.3 to 1.7)
Absolute risk (95% CI) for BRCA2 carriers					
Breast	Male	0.2 (0.1 to 0.5)	0.7 (0.4 to 1.5)	1.8 (0.9 to 3.7)	3.8 (1.9 to 7.7)
Pancreas	Male	0.2 (0.1 to 0.3)	0.9 (0.5 to 1.4)	2.0 (1.2 to 3.3)	3.0 (1.7 to 5.4)
	Female	0.2 (0.09 to 0.2)	0.6 (0.4 to 1.0)	1.5 (0.9 to 2.5)	2.3 (1.3 to 4.2)
Prostate	Male	0.2 (0.2 to 0.3)	2.9 (2.1 to 3.9)	12.6 (9.4 to 16.7)	26.9 (20.5 to 34.7)
Stomach	Male	0.1 (0.08 to 0.2)	0.5 (0.3 to 0.8)	1.4 (0.8 to 2.3)	3.5 (2.1 to 6.1)
	Female	0.2 (0.1 to 0.4)	0.6 (0.3 to 1.0)	1.3 (0.7 to 2.5)	3.5 (1.9 to 6.4)

<sup>&</sup>lt;sup>a</sup>Absolute risks were calculated on the basis of UK cancer incidences in years 2008-2012 in the Cancer Incidence in Five Continents. <sup>26</sup>

age < 65 years (RR = 3.10; 95% CI, 2.00 to 4.79) than age  $\geq$  65 years (RR = 1.69; 95% CI, 1.09 to 2.62), but this model did not fit significantly better than the model with an age-constant RR (P=.06).

#### Sensitivity Analysis

The results are described in detail in the Data Supplement. There was no significant difference in the RR estimates by geographical region. The observed cancer associations were robust to all sensitivity analyses, except for colorectal and gallbladder cancers. No association was found for melanoma even when analyses were restricted to families from Australia, Northern Europe, and North America or families in which probands self-identified as White European.

#### **Absolute Risks**

RRs from the main analysis best-fitting models were used to calculate age-specific absolute cancer risks (Table 4 and Fig 1). By age 80 years, the male breast cancer risk for *BRCA1* and *BRCA2* carriers was 0.4% (95% CI, 0.1 to 1.5) and 3.8% (95% CI, 1.9 to 7.7), respectively; the pancreatic cancer risk varied between 2.3% and 3.0% for both male and female *BRCA1* and *BRCA2* carriers; the stomach cancer risks were 1.6% (95% CI, 0.7 to 4.0) for male and 0.7% (95% CI, 0.3 to 1.7) for female *BRCA1* carriers and approximately 3.5% for both male and female *BRCA2* carriers. The prostate cancer risk associated with *BRCA2* PVs was 26.9% (95% CI, 20.5 to 34.7) by age 80 years and 33.1% (95% CI, 25.5 to 42.2) by age 85 years.

#### **DISCUSSION**

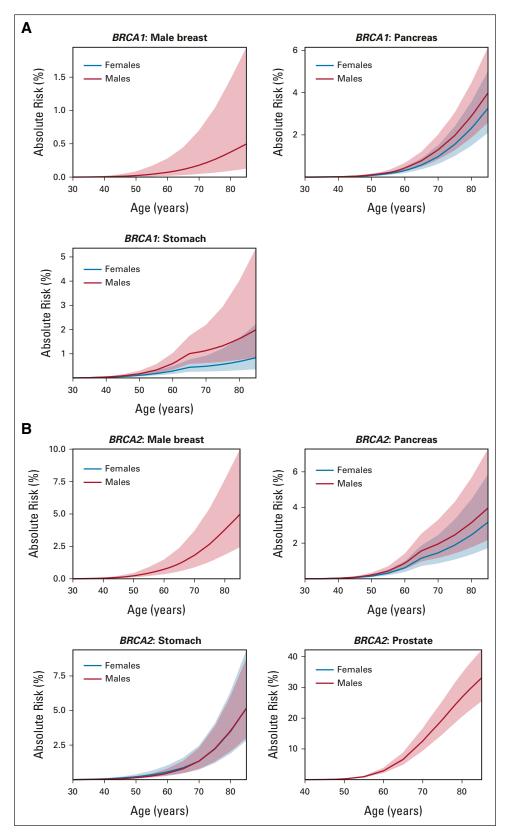
This study assessed the risks associated with *BRCA1/2* PVs for 22 first primary cancers, other than female breast and ovarian cancers, and further clarified the cancer spectrum associated with *BRCA1/2* PVs.

The associations of *BRCA1/2* PVs with the risks of male breast and pancreatic cancers were confirmed and refined, as well as the association of prostate cancer with *BRCA2* PVs, regardless of age and aggressiveness.

The lifetime male breast cancer risks were previously reported to be 2%-6% for BRCA1 and 7%-13% for BRCA2 carriers (Data Supplement).<sup>3,6,9-13</sup> We estimated these risks to be somewhat lower, 0.4% (95% CI, 0.1 to 1.5) and 3.8% (95% CI, 1.9 to 7.7), respectively. The pancreatic cancer associations were consistent with previously reported RRs of 2-3 and lifetime risks of 1%-4% for BRCA1 carriers<sup>3,4,6</sup> and RRs of 3-6 and lifetime risks of 3%-5% for BRCA2 carriers (Data Supplement).<sup>2,5-8</sup> Notably, the RR was higher for BRCA2 carriers age < 65 years.

Previous retrospective studies reported prostate cancer RRs of 2-6 and absolute risks of 17%-31% by age 80 years for BRCA2 carriers (Data Supplement). 2,5,6,8,14-17 Our estimated absolute risk by age 85 years was 33%, lower than the recently reported prospective estimate of 60% by Nyberg et al.<sup>32</sup> However, after adjusting for possible increased prostate-specific antigen screening effects in the prospective study, their estimate was 41% (95% CI, 22 to 59), consistent with our estimate. The present estimate is unlikely to be subject to increased screening biases since prostate cancer family history was retrospectively collected, and increased screening in relatives is unlikely to have taken place before the identification of BRCA2 PVs. The reported associations of BRCA1 PVs with prostate cancer risk are inconsistent, with RRs of 0.4-4, most not statistically significant. 3,4,6,8,14-18,32,33 This study confirms that BRCA1 PVs are not associated with overall prostate cancer

Among the suggested associations with other cancers, the association between *BRCA1/2* PVs and stomach cancer is



**FIG 1.** Age-specific absolute risks (%) and 95% CIs of primary cancers on the basis of UK cancer incidences in years 2008-2012 for (A) *BRCA1* and (B) *BRCA2* carriers. Solid lines are the age-specific absolute risk estimates, and ribbons are the relevant 95% CIs.

under considerable debate. 4.5 This study validated and further elucidated this association: there were associations with both *BRCA1* and *BRCA2* PVs, with RRs of 2.17 (3.50 for age < 65 years) and 3.69, respectively. Our estimates better refined the previously reported RRs of 2-7 for *BRCA1* carriers 3.6.8 and approximately 2.6 for *BRCA2* carriers (Data Supplement). 2.8 Notably, our findings showed that the stomach cancer RR for female *BRCA2* carriers was higher than the estimate for male carriers although this translated in similar absolute risks, given the higher incidence of male stomach cancer in the general population. However, we cannot exclude the possibility that the higher female RR may be due to the misclassification of some ovarian cancers as stomach cancers. 2.34

Data in the current study come from either epidemiologic studies or families undergoing PV screening collected at genetics centers. Although individual studies and clinical genetic centers, where possible, confirmed reported cancer diagnoses in families through medical records or registries as part of standard clinical practice, cancer confirmation information is not available centrally and it was not feasible to collect this at such a large scale. However, a key advantage of the present study is the large sample size, which results in RR estimates with greater precision. Only a small number of family-based studies reported cancer confirmation rates.<sup>2,4,5,8</sup> Our RR estimates for stomach cancer, which may be susceptible to a greater degree of misclassification bias than other cancers, <sup>2,34</sup> are not significantly different from the estimates from studies that reported cancer confirmation. However, the present RRs have similar or greater precision than the published estimates from studies with high cancer confirmation rates (Data Supplement).

In the present study, previously suggested associations of *BRCA1/2* PVs with risks of other genitourinary cancers and melanoma<sup>2,8</sup> were not replicated. Although associations of *BRCA1* PVs with colorectal and gallbladder cancers were observed, the results were not robust in the sensitivity analyses performed.

Increased risks of bone and liver cancer have also been reported for *BRCA1* or *BRCA2* carriers.<sup>4-6</sup> However, liver and bone are common metastatic sites for breast, prostate, or pancreatic cancers and could be the presenting cancer. Since no pathology confirmation data were available, we did not examine these associations in the main analysis. If we assume that the reported bone and liver cancers in the data set are indeed first primaries, the data suggest no association with *BRCA1* PVs, but that *BRCA2* carriers are at seven-fold increased risk of bone cancer and five-fold increased risk of liver cancer without significant differences between males and females (Data Supplement). However, no conclusion for these associations can be drawn without pathology confirmation.

Overall, the estimated age-specific relative and absolute risks suggest that, in addition to breast and ovarian

cancers, the clinical management of BRCA1/2 carriers should focus on cancer sites, which now show robust associations, such as prostate (BRCA2 carriers only), pancreatic, and possibly stomach cancers. Notably, although rare, pancreatic and stomach cancers are associated with poor prognosis and their incidences have been rising over time, and thus, our results highlight the importance of screening for upper gastrointestinal tract malignancies for BRCA1 and BRCA2 carriers, particularly for age < 65 years. On the other hand, some cancers previously taken into consideration for screening for BRCA1/2 carriers, like melanoma, may be reconsidered, to further optimize cancer prevention screening strategies and eventually reduce carriers' distress. Given that the cancer risk associations were found for both male and female carriers, the results also suggest that male relatives of known BRCA1/2 carriers should be informed about their individual cancer risk and encouraged to be tested.35,36 It has been shown that knowing the germline BRCA1/2 PV status can influence treatment options for patients with cancer, leading to improved prognosis. For example, poly (ADP-ribose) polymerase inhibitor therapies that have been used successfully in the treatment of BRCA-related breast and ovarian cancers<sup>37</sup> are now beginning to be used for pancreatic and prostate cancers, 38,39 and in the near future, they might also be used for stomach cancer.<sup>40</sup>

To avoid biases in the risk estimates related to the ascertainment of clinic-based families, on the basis of multiple affected family members, we used a conservative ascertainment adjustment approach by conditioning on the family histories of cancers of breast and ovary and the cancer site under investigation. When only family history of female breast and ovarian cancers was considered in the ascertainment, the RR estimates were somewhat higher for most cancers but with narrower CIs (Data Supplement). Therefore, conditioning on the family history of the cancer of interest is unlikely to have led to substantial underestimation of risk. A notable exception was male breast cancer, where much higher RR estimates were obtained. However, this estimate is most likely biased because male breast cancer family history has been an important factor in considering BRCA1/2 germline genetic testing since the discovery of BRCA1/2.

This study has several limitations. First, this is a retrospective family-based study, with self-reported cancer family history, which may be inaccurate. Second, 7%-40% of reported cancer cases had missing age at diagnosis, with stomach cancer having the largest proportion. To minimize these potential biases, we performed sensitivity analyses excluding any study groups in which underreporting was likely and any cases with missing age at diagnosis, and conclusions remained similar for most cancers. Third, we presented our results without any multiple testing adjustment. However, even using a false discovery rate adjustment, all the observed associations for

BRCA2 carriers and the pancreatic cancer association for BRCA1 carriers had false discovery rates < 0.05. Fourth, the ethnicity of the family proband was not systematically collected by all studies because of variations in local data collection protocols. Among those with recorded ethnicity, in Asia-based studies, 97.7% of probands were Asian and in the rest of the studies 86.1%, 5.2%, 3.7%, 1.3%, and 1.1% of probands were White European, Ashkenazi, Hispanic, Black, and Asian, respectively. Therefore, the power to investigate the associations by all ethnic groups was limited. However, we did not find evidence of heterogeneity in the RRs by geographical region (Asia  $\nu$  others). Whether our risk estimates are applicable to non-European populations requires further investigation. Fifth, we did not have data on other genetic and environmental

factors, so we were unable to investigate the modification effects of these factors; therefore, our risk estimates should be interpreted as the average risks across all potential genetic and environmental modifiers.

In conclusion, this study confirms that, aside from female breast and ovarian cancers, *BRCA1/2* PVs are associated with increased risks of breast cancer in men, and pancreatic and stomach cancers in both sexes, and that only *BRCA2* carriers are at elevated prostate cancer risk. *BRCA1/2* PVs were not associated with the risks of any other cancers previously suggested. The association results and estimated age-specific risks will improve the cancer risk management for men and women with *BRCA1/2* PVs.

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The content of this manuscript does not necessarily reflect the views or policies of the National Cancer Institute or any of the collaborating centers in the Breast Cancer Family Registry (BCFR), nor does mention of trade names, commercial products, or organizations imply endorsement by the US Government or the BCFR.

#### **EQUAL CONTRIBUTION**

S.L. and V.S. contributed equally to this work as joint first authors. L.O. and A.C.A. contributed equally to this work as joint senior authors.

#### PRIOR PRESENTATION

Presented at the Familial Aspects of Cancers: Research and Practice conference, Kingscliff, New South Wales, Australia, September 1, 2021.

#### **SUPPORT**

Supported by grants from Cancer Research UK (C12292/A20861, C12292/A11174, and PPRPGM-Nov20\100002), the Gray Foundation; by grants from Fondazione AIRC (Associazione Italiana Ricerca sul Cancro) under IG 2018 ID. 21389, LILT (Lega Italiana per la Lotta Contro i Tumori) under IG 2019, P.I. Ottini Laura, and Italian Ministry of Education, Universities and Research—Dipartimenti di Eccellenza-L. 232/2016; and by the Victorian Cancer Agency Early Career Research Fellowship (ECRF19020) and the Picchi Award for Excellence in Cancer Research from the Victorian Comprehensive Cancer Centre. G.C.-T. is a National Health and Medical Research Council (NHMRC) Research Fellow. A.B.S. and M.T.P. were supported by NHMRC Funding (APP177524). T.R.R. was supported by NIH R01CA207365. M. Tischkowitz was supported by the NIHR Cambridge Biomedical Research Centre (BRC-1215-20014). BCFR: UM1 CA164920 from the National Cancer Institute. BFBOCC: Lithuania (BFBOCC-LT): Research Council of

Lithuania grant SEN-18/2015. DEMOKRITOS: European Union (European Social Fund—ESF) and Greek national funds through the Operational Program "Education and Lifelong Learning" of the National Strategic Reference Framework (NSRF)—Research Funding Program of the General Secretariat for Research and Technology: SYN11\_10\_19 NBCA. Investing in knowledge society through the European Social Fund. DFKZ: German Cancer Research Center. EMBRACE: Cancer Research UK Grants C1287/A23382 and C1287/A26886. D.G.E. was supported by the Manchester NIHR Biomedical Research Centre (IS-BRC-1215-20007). Fiona Lalloo was supported by an NIHR grant to the Biomedical Research Centre, Manchester. The Investigators at The Institute of Cancer Research and The Royal Marsden NHS Foundation Trust were supported by an NIHR grant to the Biomedical Research Centre at The Institute of Cancer Research and The Royal Marsden NHS Foundation Trust. Ros Eeles and Elizabeth Bancroft were supported by Cancer Research UK Grant No. C5047/A8385. Ros Eeles was also supported by NIHR support to the Biomedical Research Centre at The Institute of Cancer Research and The Royal Marsden NHS Foundation Trust. FPGMX: A.V. was supported by the Spanish Health Research Foundation, Instituto de Salud Carlos III (ISCIII), partially supported by FEDER funds through Research Activity Intensification Program (contract Grant Nos.: INT15/ 00070, INT16/00154, INT17/00133, INT20/00071) and through Centro de Investigación Biomédica en Red de Enferemdades Raras CIBERER (ACCI 2016: ER17P1AC7112/2018), Autonomous Government of Galicia (Consolidation and structuring program: IN607B), and by the Fundación Mutua Madrileña, GC-HBOC: German Cancer Aid (Grant Nos. 110837 and 113049, R.K.S.). GEMO: Ligue Nationale Contre le Cancer; the Association "Le cancer du sein, parlons-en!" Award, the Canadian Institutes of Health Research for the "CIHR Team in Familial Risks of Breast Cancer" program, the Fondation ARC pour la recherche sur le cancer (Grant No. PJA 20151203365) and the French National Institute of Cancer (INCa grants AOR 01 082, 2001-2003, 2013-1-BCB-01-ICH-1, and SHS-E-SP 18-015). HCSC: Spanish Ministry of Health PI15/00059, PI16/01292, and CB-161200301 CIBERONC from ISCIII (Spain), partially supported by European Regional Development FEDER funds. M.d.I.H. was supported by a grant from the Spanish Ministry of Science and Innovation, Plan Nacional de I + D + I 2013-2016, ISCIII (PI20/00110), and FEDER from Regional Development European Funds (European Union). HRBCP: Hong Kong Sanatorium and Hospital, Dr Ellen Li Charitable Foundation, The Kerry Group Kuok Foundation, National Institute of Health1R 03CA130065, and North California Cancer Center. kConFab: The National Breast Cancer Foundation, and previously by the National Health and Medical Research Council (NHMRC), the Queensland Cancer Fund, the Cancer Councils of New South Wales, Victoria, Tasmania, and South Australia, and the Cancer Foundation of Western Australia. KOHBRA: the Korea Health Technology R&D Project through the Korea Health Industry Development Institute (KHIDI) and the National R&D Program for Cancer Control, Ministry of Health and Welfare, Republic of Korea (HI16C1127; 1020350; 1420190). NCI: the Intramural Research Program of the US National Cancer Institute, NIH, and by support services contract Nos. NO2-CP-11019-50, NO2-CP-21013-63, and NO2-CP-65504 with Westat Inc, Rockville, MD. OSUCCG: Ohio State University Comprehensive Cancer Center. SEABASS: Ministry of Science, Technology and Innovation, Ministry of Higher Education (UM.C/HIR/ MOHE/06) and Cancer Research Initiatives Foundation. SWE-BRCA: the Swedish Cancer Society. UKFOCR: Cancer Research UK. WCP: B.Y.K. was funded by the American Cancer Society Early Detection Professorship (SIOP-06-258-01-COUN) and the National Center for Advancing Translational Sciences (NCATS), Grant No. UL1TR000124.

### AUTHORS' DISCLOSURES OF POTENTIAL CONFLICTS OF INTEREST

Disclosures provided by the authors are available with this article at DOI https://doi.org/10.1200/JC0.21.02112.

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Financial support: Shuai Li, Rita K. Schmutzler, Melissa C. Southey, Shan Wang-Gohrke, Drakoulis Yannoukakos, Laura Ottini, Antonis C. Antoniou Administrative support: John L. Hopper, Shan Wang-Gohrke, Drakoulis Yannoukakos

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Manuscript writing: All authors
Final approval of manuscript: All authors
Accountable for all aspects of the work: All authors

#### **ACKNOWLEDGMENT**

We thank all the families and clinicians who contributed to the studies; GEMO study Collaborators; SWE-BRCA investigators; kConFab Investigators; Catherine M. Phelan for her contribution to CIMBA until she passed away on September 22, 2017; Sue Healey, in particular taking on the task of mutation classification with the late Olga Sinilnikova; Maggie Angelakos, Judi Maskiell, Gillian Dite, and Helen

Tsimiklis; members and participants in the New York site of the Breast Cancer Family Registry; members and participants in the Ontario Familial Breast Cancer Registry; Vilius Rudaitis and Laimonas Griškevičius; Drs Janis Eglitis, Anna Krilova, and Aivars Stengrevics; Yuan Chun Ding and Linda Steele for their work in participant enrollment and biospecimen and data management; Bent Eilertsen and Anne-Marie Gerdes for the recruitment and genetic counseling of participants; Alicia Barroso, Rosario Alonso, and Guillermo Pita; Ms JoEllen Weaver and Dr Betsy Bove; FPGMX: members of the Cancer Genetics group (IDIS): Ana Blanco, Miguel Aguado, Olivia Fuentes, and Ana Crujeiras; and IFE-Leipzig Research Centre for Civilization Diseases (Markus Loeffler, Joachim Thiery, Matthias Nüchter, and Ronny Baber). We thank all participants, clinicians, family doctors, researchers, and technicians for their contributions and commitment to the DKFZ study and the collaborating groups in Lahore, Pakistan (Noor Muhammad, Sidra Gull, Seerat Bajwa, Faiz Ali Khan, Humaira Naeemi, Saima Faisal, Asif Loya, and Mohammed Aasim Yusuf) and Bogota, Colombia (Diana Torres, Ignacio Briceno, and Fabian Gil). Genetic Modifiers of Cancer Risk in BRCA1/2 Mutation Carriers (GEMO) study is a study from the National Cancer Genetics Network UNICANCER Genetic Group, France. We would like to pay a tribute to Olga M. Sinilnikova, who with Dominique Stoppa-Lyonnet initiated and coordinated GEMO until she sadly passed away on June 30, 2014. The team in Lyon (Olga Sinilnikova, Mélanie Léoné, Laure Barjhoux, Carole Verny-Pierre, Sylvie Mazoyer, Francesca Damiola, and Valérie Sornin) managed the GEMO samples until the biological resource center was transferred to Paris in December 2015 (Noura Mebirouk, Fabienne Lesueur, and Dominique Stoppa-Lyonnet). We would like to thank all the GEMO collaborating groups for their contribution to this study: Coordinating Centre, Service de Génétique, Institut Curie, Paris, France: Muriel Belotti, Ophélie Bertrand, Anne-Marie Birot, Bruno Buecher, Sandrine Caputo, Chrystelle Colas, Anaïs Dupré, Emmanuelle Fourme, Marion Gauthier-Villars, Lisa Golmard, Marine Le Mentec, Virginie Moncoutier, Antoine de Pauw, Claire Saule, Dominique Stoppa-Lyonnet, and Inserm U900, Institut Curie, Paris, France: Fabienne Lesueur and Noura Mebirouk. Contributing Centers: Unité Mixte de Génétique Constitutionnelle des Cancers Fréquents, Hospices Civils de Lyon-Centre Léon Bérard, Lyon, France: Nadia Boutry-Kryza, Alain Calender, Sophie Giraud, and Mélanie Léone. Institut Gustave Roussy, Villejuif, France: Brigitte Bressac-de-Paillerets, Olivier Caron, Marine Guillaud-Bataille. Centre Jean Perrin, Clermont-Ferrand, France: Yves-Jean Bignon and Nancy Uhrhammer. Centre Léon Bérard, Lyon, France: Valérie Bonadona and Christine Lasset. Centre François Baclesse, Caen, France: Pascaline Berthet, Laurent Castera, and Dominique Vaur. Institut Paoli Calmettes, Marseille, France: Violaine Bourdon, Catherine Noguès, Tetsuro Noguchi, Cornel Popovici, Audrey Remenieras, and Hagay Sobol. CHU Arnaud-de-Villeneuve, Montpellier, France: Isabelle Coupier and Pascal Pujol. Centre Oscar Lambret, Lille, France: Claude Adenis, Aurélie Dumont, and Françoise Révillion. Centre Paul Strauss, Strasbourg, France: Danièle Muller. Institut Bergonié, Bordeaux, France: Emmanuelle Barouk-Simonet, Françoise Bonnet, Virginie Bubien, Michel Longy, and Nicolas Sevenet. Institut Claudius Regaud, Toulouse, France: Laurence Gladieff, Rosine Guimbaud, Viviane Feillel, and Christine Toulas. CHU Grenoble, France: Hélène Dreyfus, Christine Dominique Leroux, Magalie Peysselon, and Rebischung. CHU Dijon, France: Amandine Baurand, Geoffrey Bertolone, Fanny Coron, Laurence Faivre, Caroline Jacquot, and Sarab Lizard. CHU St-Etienne, France: Caroline Kientz, Marine Lebrun, and Fabienne Prieur. Hôtel Dieu Centre Hospitalier, Chambéry, France: Sandra Fert Ferrer. Centre Antoine Lacassagne, Nice, France: Véronique Mari. CHU Limoges, France: Laurence Vénat-Bouvet. CHU Nantes, France: Stéphane Bézieau and Capucine Delnatte. CHU Bretonneau, Tours and Centre Hospitalier de Bourges France: Isabelle Mortemousque. Groupe Hospitalier Pitié-Salpétrière, Paris, France: Florence Coulet and Mathilde Warcoin. CHU Vandoeuvre-les-Nancy, France: Myriam Bronner and Johanna Sokolowska, CHU Besancon, France: Marie-Agnès Collonge-Rame, Alexandre Damette. CHU Poitiers, Centre Hospitalier d'Angoulême and

Centre Hospitalier de Niort, France: Paul Gesta. Centre Hospitalier de La Rochelle: Hakima Lallaoui. CHU Nîmes Carémeau, France: Jean Chiesa. CHI Poissy, France: Denise Molina-Gomes. CHU Angers, France: Olivier Ingster; CHU de Martinique, France: Odile Bera; Mickaelle Rose; Ilse Coene and Brecht Crombez; Alicia Tosar and Paula Diague; Drs.Sofia Khan, Taru A. Muranen, Carl Blomqvist, Irja Erkkilä, and Virpi Palola; Hong Kong Sanatorium and Hospital; the Hungarian Breast and Ovarian Cancer Study Group members (Janos Papp, Aniko Bozsik, Timea Pocza, Zoltan Matrai, Miklos Kasler, Judit Franko, Maria Balogh, Gabriella Domokos, and Judit Ferenczi, Department of Molecular Genetics, National Institute of Oncology, Budapest, Hungary) and the clinicians and patients for their contributions to this study; the Oncogenetics Group (VHIO) and the High Risk and Cancer Prevention Unit of the University Hospital Vall d'Hebron, Miguel Servet Progam (CP10/00617), and the Cellex Foundation for providing research facilities and equipment; the ICO Hereditary Cancer Program team led by Dr Gabriel Capella; the ICO Hereditary Cancer Program team led by Dr Gabriel Capella; Dr Martine Dumont for sample management and skillful assistance; Ana Peixoto, Catarina Santos, and Pedro Pinto; members of the Center of Molecular Diagnosis, Oncogenetics Department and Molecular Oncology Research Center of Barretos Cancer Hospital; Heather Thorne, Eveline Niedermayr, all the kConFab research nurses and staff, the heads and staff of the Family Cancer Clinics, and the Clinical Follow-Up Study (which has received funding from the NHMRC, the National Breast Cancer Foundation, Cancer Australia, and the National Institute of Health [United States]) for their contributions to this resource, and the many families who contribute to kConFab; the KOBRA Study Group; Csilla Szabo (National Human Genome Research Institute, National Institutes of Health, Bethesda, MD); Lenka Foretova and Eva Machackova (Department of Cancer Epidemiology and Genetics, Masaryk Memorial Cancer Institute and MF MU, Brno, Czech Republic); and Michal Zikan, Petr Pohlreich, and Zdenek Kleibl (Oncogynecologic Center and Department of Biochemistry and Experimental Oncology, First Faculty of Medicine, Charles University, Prague, Czech Republic); Anne Lincoln and Lauren Jacobs; the participants in Hereditary Breast/Ovarian Cancer Study and Breast Imaging Study for their selfless contributions to our research; the NICCC National Familial Cancer Consultation Service team led by Sara Dishon, the laboratory team led by Dr Flavio Lejbkowicz, and the research field operations team led by Dr Mila Pinchev; the investigators of the Australia New Zealand NRG Oncology group; members and participants in the Ontario Cancer Genetics Network; Kevin Sweet, Caroline Craven, Julia Cooper, Amber Aielts, and Michelle O'Conor; Yip Cheng Har, Nur Aishah Mohd Taib, Phuah Sze Yee, Norhashimah Hassan, and all the research nurses, research assistants, and doctors involved in the MyBrCa Study for assistance in patient recruitment, data collection, and sample preparation; Philip Iau, Sng Jen-Hwei, and Sharifah Nor Akmal for contributing samples from the Singapore Breast Cancer Study and the HUKM-HKL Study, respectively; the Meirav Comprehensive breast cancer center team at the Sheba Medical Center; Christina Selkirk; Håkan Olsson, Helena Jernström, Karin Henriksson, Katja Harbst, Maria Soller, and Ulf Kristoffersson; from Gothenburg Sahlgrenska University Hospital: Anna Ofverholm, Margareta Nordling, Per Karlsson, and Zakaria Einbeigi; from Stockholm and Karolinska University Hospital: Anna von Wachenfeldt, Annelie Liljegren, Annika Lindblom, Brita Arver, and Gisela Barbany Bustinza; from Umeå University Hospital: Beatrice Melin, Christina Edwinsdotter Ardnor, and Monica Emanuelsson; from Uppsala University: Hans Ehrencrona, Maritta Hellström Pigg, and Richard Rosenquist; from Linköping University Hospital: Marie Stenmark-Askmalm, Sigrun Liedgren; Cecilia Zvocec, Qun Niu; Joyce Seldon and Lorna Kwan; Dr Robert Nussbaum, Beth Crawford, Kate Loranger, Julie Mak, Nicola Stewart, Robin Lee, Amie Blanco and Peggy Conrad and Salina Chan; Simon Gayther, Paul Pharoah, Carole Pye, Patricia Harrington, and Eva Wozniak; and Geoffrey Lindeman, Marion Harris, Martin Delatycki, Sarah Sawyer, Rebecca Driessen, and Ella Thompson for performing all DNA amplification.

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#### **AUTHORS' DISCLOSURES OF POTENTIAL CONFLICTS OF INTEREST**

#### Cancer Risks Associated With BRCA1 and BRCA2 Pathogenic Variants

The following represents disclosure information provided by authors of this manuscript. All relationships are considered compensated unless otherwise noted. Relationships are self-held unless noted. I = Immediate Family Member, Inst = My Institution. Relationships may not relate to the subject matter of this manuscript. For more information about ASCO's conflict of interest policy, please refer to www.asco.org/rwc or ascopubs.org/jco/authors/author-center.

Open Payments is a public database containing information reported by companies about payments made to US-licensed physicians (Open Payments).

Timothy R. Rebbeck Honoraria: AstraZeneca (I)

Consulting or Advisory Role: AstraZeneca (I)

Andrew Lee Employment: Illumina

Patents, Royalties, Other Intellectual Property: I am an inventor of the BOADICEA model that is licensed to Cambridge Enterprise (part of the University of Cambridge) for commercialization. I have received royalties from Cambridge Enterprise.

Norbert Arnold Honoraria: AstraZeneca

Ake Borg

Honoraria: Roche, AstraZeneca

Travel, Accommodations, Expenses: Roche, AstraZeneca

Sandrine Caputo

Research Funding: AstraZeneca/France (Inst)

Wendy K. Chung

Consulting or Advisory Role: Regeneron

Research Funding: Biogen

Chrystelle Colas Consulting or Advis

Consulting or Advisory Role: AstraZeneca/Merck, Pfizer

Miguel de la Hoya

Consulting or Advisory Role: AstraZeneca Research Funding: AstraZeneca

D. Gareth EvansHonoraria: AstraZeneca

Tanja N. Fehm

Consulting or Advisory Role: Roche, Novartis, Pfizer, AstraZeneca, Daiichi

Sankyo, MSD Oncology

Travel, Accommodations, Expenses: Roche

George Fountzilas

Stock and Other Ownership Interests: GENPREX Inc (I), ARIAD (I), Deciphera Pharmaceuticals Inc (I), Daiichi Sankyo, RFL Holdings, FORMYCON

Honoraria: AstraZeneca

Consulting or Advisory Role: Pfizer, Roche, Novartis Speakers' Bureau: Roche (I), LEO Pharma (I), Pfizer (I)

Travel, Accommodations, Expenses: Merck (I), Pfizer (I), K.A.M Oncology/

Hematology (I)

Megan Frone

Patents, Royalties, Other Intellectual Property: Receive royalties for being a

codeveloper of CancerGene Connect

Eric Hahnen

Consulting or Advisory Role: AstraZeneca

Helen Hanson Honoraria: Pfizer

Beth Y. Karlan

Consulting or Advisory Role: Roche Pharma AG, Merck, Mercy Bioanalytics,

GRAIL

Research Funding: GOG Foundation (Inst), NCI-NRG Oncology (Inst),

Genentech/Roche (Inst)

Patents, Royalties, Other Intellectual Property: US and EU patent on gene

signature

Other Relationship: Elsevier Irene Konstantopoulou

Speakers' Bureau: AstraZeneca Research Funding: AstraZeneca

Ava Kwong

Honoraria: Stryker, AstraZeneca Taiwan Limited, Roche Singapore, Pfizer,

AstraZeneca Hong Kong Ltd

Joanne Ngeow Yuen Yie Research Funding: AstraZeneca

Rita K. Schmutzler

Honoraria: AstraZeneca, Clovis, MSD/AstraZeneca Consulting or Advisory Role: AstraZeneca, MSD/AstraZeneca

Research Funding: Amgen

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Consulting or Advisory Role: AstraZeneca/Merck

Speakers' Bureau: AstraZeneca/Merck

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Honoraria: Novartis, AstraZeneca/MedImmune, Daiichi Sankyo Europe GmbH Consulting or Advisory Role: AstraZeneca/MedImmune, Daiichi-Sankyo, Gilead

Sciences, Sanofi/Aventis, Novartis

Speakers' Bureau: Novartis, AstraZeneca/MedImmune

Research Funding: Novartis, Sanofi, Myriad Genetics, Roche, AstraZeneca/

MedImmune

Travel, Accommodations, Expenses: Roche, Novartis

Dominique Stoppa-Lyonnet Honoraria: AstraZeneca/Merck (Inst)

Soo Hwang Teo

Speakers' Bureau: AstraZeneca, Pfizer, Roche

Sebastian A. Wagner

Consulting or Advisory Role: Bayer

Antonis C. Antoniou

Patents, Royalties, Other Intellectual Property: Inventor of the BOADICEA model, which has been licensed to Cambridge Enterprise for commercialization.

May receive royalties if commercialization is realized.

No other potential conflicts of interest were reported.