RESEARCH ARTICLE





Mammographic density and breast cancer in women from high risk families

Teresa Ramón y Cajal¹, Isabel Chirivella², Josefa Miranda³, Alexandre Teule⁴, Ángel Izquierdo⁵, Judith Balmaña⁶, Ana Beatriz Sánchez-Heras⁷, Gemma Llort⁸, David Fisas¹, Virginia Lope^{9,10,11}, Elena Hernández-Agudo¹², María José Juan-Fita¹³, Isabel Tena¹⁴, Luis Robles¹⁵, Carmen Guillén-Ponce¹⁶, Pedro Pérez-Segura¹⁷, Mari Sol Luque-Molina¹⁸, Susana Hernando-Polo¹⁹, Mónica Salinas⁴, Joan Brunet^{5,20}, María Dolores Salas-Trejo³, Agustí Barnadas¹ and Marina Pollán^{9,10,11*}

Abstract

Introduction: Mammographic density (MD) is one of the strongest determinants of sporadic breast cancer (BC). In this study, we compared MD in *BRCA1/2* mutation carriers and non-carriers from *BRCA1/2* mutation-positive families and investigated the association between MD and BC among *BRCA1/2* mutation carriers per type of mutation and tumor subtype.

Methods: The study was carried out in 1039 female members of *BRCA1* and *BRCA2* mutation-positive families followed at 16 Spanish Genetic Counseling Units. Participants' density was scored retrospectively from available mammograms by a single blinded radiologist using a 5-category scale (<10 %, 10-25 %, 25-50 %, 50-75 %, >75 %). In BC cases, we selected mammograms taken prior to diagnosis or from the contralateral breast, whereas, in non-cases, the last screening mammogram was evaluated. MD distribution in carriers and non-carriers was compared using ordinal logistic models, and the association between MD and BC in *BRCA1/2* mutation carriers was studied using logistic regression. Huber-White robust estimators of variance were used to take into account correlations between family members. A similar multinomial model was used to explore this association by BC subtype.

Results: We identified and scored mammograms from 341 *BRCA1*, 350 *BRCA2* mutation carriers and 229 non-carriers. Compared to non-carriers, MD was significantly lower among *BRCA2* mutation carriers (odds ratio (OR) =0.71; *P*-value=0.04), but not among *BRCA1* carriers (OR=0.84; *P*-value=0.33). MD was associated with subsequent development BC (OR per category of MD=1.45; 95 % confidence interval=1.18-1.78, *P*-value<0.001), with no significant differences between *BRCA1* and *BRCA2* mutation carriers (*P*-value=0.48). Finally, no statistically significant differences were observed in the association of MD with specific BC subtypes.

Conclusions: Our study, the largest to date on this issue, confirms that MD is an independent risk factor for all BC subtypes in either *BRCA1* and *BRCA2* mutation carriers, and should be considered a phenotype risk marker in this context.

Introduction

Mammographic density (MD) is one of the strongest determinants of breast cancer (BC) in most ethnic groups. In general population studies, higher MD is consistently associated with a higher BC risk [1]. Individual variation of MD does not only depend on well-established

¹⁰Consortium for Biomedical Research in Epidemiology and Public Health (CIBERESP), Carlos III Institute of Health, Madrid, Spain

epidemiologic factors, such as age, parity, menopausal status, external hormonal manipulation or body mass index (BMI), but based on family studies [2–7], it also depends on the as yet unknown genetic background.

Knowledge of the genetic basis of MD could be a major milestone in BC risk prediction and prevention. Previous candidate gene analysis and linkage studies have shown either inconclusive or discordant results to date. In contrast, genome-wide association studies have started to identify some genetic variants that could explain part of the variation in MD [8]. Furthermore,



© 2015 Ramón y Cajal et al. This is an Open Access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/4.0), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly credited. The Creative Commons Public Domain Dedication waiver (http://creativecommons.org/publicdomain/zero/1.0/) applies to the data made available in this article, unless otherwise stated.

^{*} Correspondence: mpollan@isciii.es

⁹National Center for Epidemiology, Carlos III Institute of Health, Monforte de Lemos 5, 28029 Madrid, Spain

Full list of author information is available at the end of the article

some of the well-established BC susceptibility variants are also associated with MD variability [9, 10].

Germline mutations in *BRCA1* and *BRCA2* genes are the most frequent cause of strong genetic predisposition to breast and ovarian cancer. In the last decades, several studies have evaluated whether the mutational status of both genes is associated with differences in MD compared to the general population. Therefore, whereas earlier studies with limited numbers of patients [11–16] had conflicting results, two larger and well-conducted studies compared MD in either affected or healthy *BRCA1/2* mutation carriers [17] and in mutation carriers versus women with low to average risk [18]. In one of them, authors proved MD to be an independent risk factor for BC in carriers [17]. In contrast, findings from a recent study did not reveal any association between baseline MD and subsequent risk of BC among carriers [19].

In this paper, we aimed to compare MD in Spanish *BRCA1/2* mutation carriers and non-carriers and assess the potential impact of MD on subsequent BC risk among carriers according to the mutated gene and pathologic subtype.

Methods

A total of 1,039 women were enrolled from 509 breast and ovarian cancer families who had been counseled and tested for mutations in *BRCA1* and *BRCA2* genes, from January 1993 to December 2012, and followed up at 16 Cancer Genetic Units in Spain. The study was approved by national and regional Ethics Committees (see Acknowledgments for details) and conducted in compliance with the Helsinki Declaration. All women were older than 18 years and gave written informed consent for the study.

Eligible participants were women from mutation-positive families, including *BRCA1/2* mutation carriers and true mutation-negative women. Women diagnosed with BC were considered cases, whereas those individuals with no personal history of BC were regarded as non-cases or controls.

Structured questionnaires collecting baseline information were administered by investigators during either the highrisk breast screening or the post-test visit. Apart from demographic characteristics, data for other covariates were collected, such as the genetic condition, age, weight and height, reproductive history, menopausal status, the use of exogenous hormonal supplements and history of riskreducing surgery up to the time of the study mammogram. Controls' weight and height were measured by the investigator at the enrollment visit. Tumor subtype, weight and height at the time of the scored mammogram were extracted from the medical charts of BC patients.

Mammograms were either obtained from the radiological chart or provided by the participants. Due to the retrospective design of our study, we considered density measures from analog and digital mammographic films. For cancer patients, we aimed to collect the earliest mammogram obtained before diagnosis (within a 10-year period and identical menopausal status) and excluded the films from the affected breast. When unavailable, we used the mammogram of the unaffected contralateral breast taken at the time of cancer diagnosis. To make the age of controls at the time of the mammogram more comparable with those of women with BC, we attempted to obtain the latest available mammogram. For each participant, films from craniocaudal and mediolateral oblique views were requested. There were 119 women who either did not have a suitable mammogram (110) or the quality of the film was insufficient (9), rendering a total of 920 individuals for whom the mammogram was available and MD reading was feasible. There were 691 women carrying a deleterious mutation in the BRCA1 or BRCA2 (341 women with BRCA1 and 350 with BRCA2) genes and 229 women were relatives without a mutation.

Breast density was visually assessed by one experienced radiologist, with high intra-observer concordance [20]. The radiologist was blinded to the diagnosis, the carrier status and the referral centre of the mammogram. MD was classified using the Boyd semiquantitative scale into 6 categories, namely 0 %, <10 %, 10–25 %, 25–50 %, 50–75 % and >75 %. The radiologist provided a single reading per participant, considering the craniocaudal and mediolateral oblique views in both breasts when available. For those breast cancer cases without previous mammography, the two views of the contralateral breast were read.

The association between MD and the carrier status as well as other variables of interest was assessed in healthy women using an ordinal logistic regression model. This procedure results from fitting k-1 logistic regression models, dichotomizing MD into two groups, i.e., high versus low MD, using all possible cutoffs. This model, also known as the proportional odds model, assumes that odds ratios (ORs) remain constant irrespective of the cutoff chosen, but allows the intercepts to differ. The following variables were included as possible confounders: age at mammogram, BMI, menopausal status (premenopause/natural menopause/surgical menopause), parity (nulliparous/parous), type of image (analog/digital) and carrier status (non-carrier, BRCA1, BRCA2). The correlation between members of the same family was taken into account using Huber-White robust estimators of variance, considering individuals clustered inside families [21]. The Brant test was used to verify the proportional odds assumption.

The association between MD and subsequent BC development was evaluated using a logistic regression model, adjusted for age at mammogram, menopausal status (premenopause/natural menopause/surgical menopause), BMI, parity, age at first live birth, use of hormonal

replacement therapy (HRT) (never/ever), type of image and time elapsed from mammogram to either BC diagnosis (cases) or end of follow up (non-cases). The consistency of the association between MD and BC was explored performing subgroup analysis in models including MD as a continuous variable and an interaction term between MD and each of the explanatory variables already mentioned. Finally, possible differences in the effect of MD per BC subtype were assessed using a multinomial logistic model, adjusted for the same confounders and considering the following subtypes: 1) hormonal receptor positive (estrogen receptor (ER) or progesterone receptor (PR)) and human epidermal growth factor receptor 2 (HER2) negative, 2) HER2-positive, 3) triple-negative tumors, and 4) unknown subtype. Again, Huber-White robust estimators of variance were used in all instances to take into account the correlation between relatives [21]. Sensitivity analyses were carried out considering only those cases with mammograms taken 6 months or more before BC diagnosis. All statistical analyses were performed by using the STATA version 12.0 software program (Stata Corp, College Station, TX, USA).

Results

Baseline characteristics

Baseline characteristics of our population according to BRCA1/2 mutation carrier status and development of breast cancer are shown in Table 1. Carriers were younger at the scored mammogram (mean age was 41 years, regardless of the mutated gene and disease condition) compared with non-carriers (45 years; *P* value <0.001). In addition, carriers were more likely to have ever used hormone preventive treatment (1.4 % vs 0.4 %; *P* value = 0.06) and have undergone surgical menopause at the time of the selected mammogram (10.3 % vs 1.7 %; *P* value <0.001). There were 31 women, 21 *BRCA1* and 10 *BRCA2* mutation carriers, with a previous diagnosis of ovarian cancer, and four of them (2 *BRCA1* and 2 *BRCA2* mutation carriers) subsequently developed a breast tumor.

Among 691 *BRCA1/2* mutation carriers, 233 women developed invasive breast cancer (111 and 122 with *BRCA1* and *BRCA2* mutations, respectively), and 15 patients had ductal carcinoma in situ (3 *BRCA1* and 12 *BRCA2* mutation carriers, respectively), whereas 7 women had invasive breast cancer and the other 3 women developed ductal carcinoma in situ, among 229 non-carriers. Regarding pathologic subtypes, 54 cases, 2 of them among non-carriers, could not be classified due to lack of information on hormone receptor expression (11 cases) or HER2 status (54 cases). As expected, *BRCA2* mutation carriers were more likely to have hormone receptor positive tumors and no expression of HER2 (71 % versus 26 %; *P* value <0.001), whereas *BRCA1* mutation carriers had a higher proportion of triple-negative tumors (66 % versus 14 %; *P* value <0.001).

When we looked specifically at *BRCA1/2* mutation carriers, breast cancer patients had a lower BMI compared to controls, with 92 % of cases versus 85 % of controls having BMI <30 (*P* value = 0.03) (Table 1). No differences were seen in the age at the time of scored mammogram, but analog mammography was more frequent among cases (84 % versus 30 %). Finally, MD was higher in affected carriers, with 33 % of them in the two highest categories (MD >50 %) as opposed to 22 % of controls (*P* value = 0.001). Menopausal status, parity, age at first birth, oral contraceptives and hormonal replacement therapy were not significantly different between both groups. As expected, surgical menopause was more frequent among controls (14 % versus 4 %; *P* value <0.001).

Association of mammographic density with other variables

The association of all categories of mammographic density and explanatory variables is shown in Table 2. Given the small number of women with 0 % MD, the two first categories were combined in subsequent analyses. In all instances, ORs were adjusted for age at mammogram, BMI, menopausal status (premenopause/natural menopause/surgical menopause), type of mammogram, parity, carrier status and subsequent development of BC. We did not find a significant association between carrying a BRCA1/2 mutation and a lower MD (OR = 0.77; 95 % CI = 0.57, 1.05). However, when *BRCA1* and *BRCA2* mutation carriers were considered separately, BRCA2 mutation carriers had a statistically significant lower MD (OR = 0.71; 95 % CI = 0.51, 0.98: P value = 0.04), but no differences were observed for BRCA1 mutation carriers (P value = 0.332). Most risk factors considered, including age at mammography, BMI, menopausal status and parity were, as expected, significantly associated with MD. Moreover, MD was higher among women who subsequently developed a breast tumor (OR = 1.69; 95 % CI = 1.24, 2.31; P value = 0.001), Finally, neither age at first birth nor hormonal interventions such as HRT or oral contraceptives (OC) significantly modified MD in our study.

Effect of mammographic density in development of breast cancer in *BRCA1/2* mutation carriers

The distribution of MD in BC cases and non-cases for women who were non-carriers and *BRCA1* and *BRCA2* mutation-carriers is presented in Fig. 1. Cases tended to be more frequently classified in the higher MD categories. Given the limited number of BC cases among the noncarriers group, we did not evaluate the potential different effect of MD on cancer risk in non-carriers. Table 3 shows the association between MD and subsequent development of BC among *BRCA1/2* mutation carriers. After adjusting

Variable	BRCA1/2 mutatio	n carriers		Relatives without		
	Without BC	BC patients	P value ^a	Without BC	BC patients	P value ^a
	N = 443	N = 248		N = 219	N = 10	
Affected gene			0.183			
BRCA1	227 (51.2 %)	114 (46.0 %)				
BRCA2	216 (48.8 %)	134 (54.0 %)				
Age at mammography			0.909			0.907
Mean (SD)	41.6 (11.5)	41.5 (10.2)		45.6 (11.8)	45.2 (11.2)	
Body mass index			0.030			0.948
<18.5	6 (1.4 %)	7 (2.8 %)		11 (5.0 %)	0 (0.0 %)	
18.5–24.9	259 (58.5 %)	151 (60.9 %)		120 (54.8 %)	7 (70.0 %)	
25–29.9	110 (24.8 %)	69 (27.8 %)		56 (25.6 %)	2 (20.0 %)	
30-34.9	49 (11.1 %)	19 (7.7 %)		28 (12.8 %)	1 (10.0 %)	
>=35	19 (4.3 %)	2 (0.8 %)		4 (1.8 %)	0 (0.0 %)	
Unknown	0 (0.0 %)	0 (0.0 %)		11 (5.0 %)	0 (0.0 %)	
Menopausal status ^b			0.239			0.576
Premenopausal	310 (70.0 %)	184 (74.2 %)		147 (67.1 %)	7 (70.0 %)	
Postmenopausal	133 (30.0 %)	64 (25.8 %)		72 (32.9 %)	3 (30.0 %)	
Type of menopause ^{b,c}			< 0.001			0.154
Natural	73 (16.5 %)	53 (21.4 %)		69 (95.8 %)	2 (66.7 %)	
Surgery	60 (13.5 %)	11 (4.4 %)		3 (4.2 %)	1 (33.3 %)	
Time since menopause ^{b,c}			0.100			1.000
<=5 years	31 (7.0 %)	19 (7.7 %)		27 (12.3 %)	1 (10.0 %)	
6–10 years	34 (7.7 %)	8 (3.2 %)		17 (7.8 %)	1 (10.0 %)	
>10 years	63 (14.2 %)	35 (14.1 %)		28 (12.8 %)	1 (10.0 %)	
Unknown	5 (1.1 %)	2 (0.8 %)		0 (0.0 %)	0 (0.0 %)	
Nulliparous			0.151			0.264
No	302 (68.2 %)	182 (73.4 %)		167 (76.3 %)	6 (60.0 %)	
Yes	141 (31.8 %)	66 (26.6 %)		52 (23.7 %)	4 (40.0 %)	
Age at first birth, years ^d			0.110			0.464
>30	75 (16.9 %)	34 (13.7 %)		46 (27.5 %)	0 (0.0 %)	
26–30	111 (25.1 %)	64 (25.8 %)		60 (35.9 %)	3 (50.0 %)	
21-25	98 (22.1 %)	77 (31.0 %)		53 (31.7 %)	3 (50.0 %)	
<=20	18 (4.1 %)	7 (2.8 %)		8 (4.8 %)	0 (0.0 %)	
Oral contraceptives			0.487			0.508
No	156 (35.2 %)	81 (32.7 %)		58 (26.5 %)	4 (40.0 %)	
Yes	239 (54.0 %)	140 (56.5 %)		130 (59.4 %)	6 (60.0 %)	
Unknown	48 (10.8 %)	27 (10.9 %)		31 (14.2 %)	0 (0.0 %)	
Use of hormone replacement	nt therapy ^b		0.613			0.246
No	431 (97.3 %)	238 (96.0 %)		205 (93.6 %)	9 (90.0 %)	
Current use	3 (0.7 %)	3 (1.2 %)		5 (2.3 %)	1 (10.0 %)	
Past use	9 (2.0 %)	7 (2.8 %)		9 (4.1 %)	0 (0.0 %)	
Use of hormonal preventive	treatment ^b		0.349			
No	438 (98.9 %)	243 (98.0 %)		219 (99.5 %)	10 (100 %)	
Current use	5 (1.1 %)	5 (2.0 %)		1 (0.5 %)	0 (0.0 %)	

Table 1 Characteristics of the women from BRCA1/2 families included in the study

Type of mammogram			<0.001			0.020
Analog	132 (29.8 %)	208 (83.9 %)		62 (28.3 %)	6 (60.0 %)	
Digital	311 (70.2 %)	40 (16 1 %)		157 (71 7 %)	4 (40.0 %)	
Number of projections	311 (7012 70)	10 (1011 70)	< 0.001		. (1010 70)	0.069
One	9 (2.0 %)	6 (2.4 %)				
Тwo	34 (7.7 %)	74 (29.8 %)		11 (4.7 %)	3 (25.0 %)	
Three	2 (0.5 %)	3 (1.2 %)		1 (0.4 %)	0 (0.0 %)	
Four	398 (89.8 %)	165 (66.5 %)		223 (94.9 %)	9 (75.0 %)	
Mammographic density			0.001			0.316
0%	6 (1.4 %)	0 (0.0 %)		3 (1.4 %)	1 (10.0 %)	
<=10 %	99 (22.3 %)	35 (14.1 %)		44 (20.1 %)	1 (10.0 %)	
11-25 %	89 (20.1 %)	47 (19.0 %)		44 (20.1 %)	2 (20.0 %)	
26-50 %	151 (34.1 %)	85 (34.3 %)		72 (32.9 %)	3 (30.0 %)	
51-75 %	72 (16.3 %)	47 (19.0 %)		37 (16.9 %)	3 (30.0 %)	
>75 %	26 (5.9 %)	34 (13.7 %)		19 (8.7 %)	0 (0.0 %)	
Time of follow up ^e			< 0.001			0.924
<=1 year	207 (46.7 %)	162 (65.3 %)		103 (47.0 %)	4 (40.0 %)	
1-2 years	82 (18.5 %)	30 (12.1 %)		39 (17.8 %)	2 (20.0 %)	
2–5 years	94 (21.2 %)	30 (12.1 %)		48 (21.9 %)	3 (30.0 %)	
5–10 years	46 (10.4 %)	24 (9.7 %)		22 (10.0 %)	1 (10.0 %)	
>10 years	14 (3.2 %)	2 (0.8 %)		7 (3.2 %)	0 (0.0 %)	
Prophylatic mastectomy			0.003			
No	428 (96.6 %)	248 (100.0 %)				
Yes	15 (3.4 %)	0 (0.0 %)				
Breast cancer features						
Histology						
Ductal in situ		15 (6.0 %)			3 (30.0 %)	
Ductal		187 (75.4 %)			6 (60.0 %)	
Lobular		12 (4.8 %)			1 (10.0 %)	
Medular		14 (5.6 %)			0 (0.0 %)	
Others		18 (7.3 %)			0 (0.0 %)	
Unknown		2 (0.8 %)			0 (0.0 %)	
Type of breast cancer						
ER/PR+&HER2-		99 (39.9 %)			6 (60.0 %)	
HER2+		23 (9.3 %)			1 (10.0 %)	
Triple-negative		74 (29.8 %)			1 (10.0 %)	
Unknown		52 (21.0 %)			2 (20.0 %)	

Table 1 Characteristics of the women from BRCA1/2 families included in the study (Continued)

^aComparison between women who did and did not develop breast cancer. ^bAt time of mammography. ^cOnly postmenopausal women at time of mammography. ^dOnly parous women. ^eTime elapsed from mammographic exploration to breast cancer diagnosis (cases) or last contact (controls). *BC* breast cancer, *ER* estrogen receptor, *PR* progesterone receptor, *HER2* human epidermal growth factor receptor 2

for age at mammogram, BMI, menopause (premenopause/ natural menopause/surgical menopause), parity, age at first birth, HRT, type of mammogram and time of follow up, there was a clear association between MD and subsequent BC development among *BRCA1/2* mutation carriers, with a statistically significant dose-response trend (*P* value <0.001). The OR in women with density 50–75 % relative to women with density ≤ 10 % was 3.24 (95 % CI = 1.43, 7.35; *P* value = 0.005), while women in the highest MD category, namely >75 %, had an OR of 4.34 (95 % CI = 1.71, 11.1; *P* value = 0.002). The association between MD and BC tended to be stronger among *BRCA2* mutation carriers

		Categories	of mammogr	aphic density						
Variable	Number	<=10 %	11-25 %	26-50 %	51-75 %	>75 %	Odds ratio ^a	95 % Cl ^a	P value	
Carrier status										
Non-carrier	219	21 %	20 %	33 %	17 %	9 %	1.00			
BRCA1 or BRCA2	443	24 %	20 %	34 %	16 %	6 %	0.75	0.54, 1.04	0.085	
BRCA1	227	22 %	25 %	29 %	19 %	6 %	0.84	0.58, 1.23	0.376	
BRCA2	216	26 %	15 %	39 %	14 %	6 %	0.67	0.47, 0.96	0.028	
Age at mammography, ye	ars									
<35	188	10 %	14 %	36 %	27 %	13 %	1.00			
35–44	199	17 %	19 %	39 %	17 %	8 %	0.94	0.54, 1.62	0.826	
45-54	170	29 %	25 %	31 %	12 %	3 %	0.79	0.33, 1.90	0.603	
>=55	105	49 %	25 %	24 %	3 %	0 %	0.57	0.15, 2.20	0.415	
Linear trend (per year)							0.97	0.95, 0.99	< 0.001	
Body mass index										
<25	396	10 %	17 %	39 %	24 %	10 %	1.00			
25.0-29.9	166	36 %	26 %	28 %	7 %	3 %	0.28	0.19, 0.40	< 0.001	
>=30	100	51 %	24 %	23 %	2 %	0 %	0.16	0.10, 0.25	< 0.001	
Linear trend (per unit)							0.83	0.80, 0.86	< 0.001	
Menopausal status & type	of menopause	2								
Premenopausal	457	15 %	17 %	37 %	21 %	9 %	1.00			
Natural menopause	142	41 %	24 %	28 %	6 %	1 %	0.72	0.45, 1.16	0.179	
Surgical menopause	63	38 %	35 %	21 %	6 %	0 %	0.56	0.33, 0.96	0.034	
Nulliparous										
No	460	29 %	23 %	33 %	13 %	2 %	1.00			
Yes	202	9 %	14 %	35 %	24 %	17 %	1.87	1.28, 2.73	0.001	
Age at first birth, years (or	nly parous won	nen)								
>30	131	25 %	17 %	35 %	17 %	6 %	1.00			
25–29	170	25 %	25 %	34 %	15 %	1 %	1.17	0.72, 1.88	0.526	
20–24	151	35 %	24 %	30 %	9 %	2 %	1.10	0.67, 1.82	0.706	
<20	26	38 %	23 %	27 %	12 %	0 %	0.72	0.31, 1.66	0.436	
Linear trend (per year)							1.00	0.97, 1.05	0.745	
Hormonal replacement th	erapy									
No	636	23 %	20 %	34 %	17 %	7 %	1.00			
Yes	26	23 %	27 %	27 %	15 %	8 %	1.71	0.87, 3.38	0.120	
Hormonal preventive ther	ару									
No	656	23 %	20 %	34 %	17 %	7 %	1.00			
Yes	6	33 %	17 %	50 %	0 %	0 %	0.67	0.25, 1.82	0.434	
Oral contraception										
No	214	22 %	18 %	34 %	19 %	7 %	1.00			
Yes	369	22 %	21 %	33 %	17 %	7 %	0.93	0.67, 1.27	0.637	
Type of image										
Analog	194	26 %	21 %	34 %	13 %	6 %	1.00			
Digital	468	22 %	20 %	34 %	18 %	7 %	1.09	0.79, 1.50	0.588	

 Table 2 Association of mammographic density with carrier status and other characteristics in healthy female mutation carriers (controls) from BRCA1/BRCA2 families

 Table 2 Association of mammographic density with carrier status and other characteristics in healthy female mutation carriers

 (controls) from BRCA1/BRCA2 families (Continued)

 Number of projections

Three or Four	609	24 %	20 %	33 %	16 %	7 %	1.17	0.70, 1.94	0.544
One or two	53	15 %	17 %	40 %	21 %	8 %	1.00		
Number of projections									

^aAdjusted for age at mammogram, body mass index, menopause (premenopausal, natural menopause, surgical menopause), type of mammogram, parity (nulliparous versus parous), carrier status (non-carrier, *BRCA1*, *BRCA2*) and subsequent development of breast cancer (no, yes)

(OR per increase in one category of MD of 1.60 in BRCA2 and of 1.37 in BRCA1 carriers), but these differences were not statistically significant (P value for heterogeneity = 0.449). Sensitivity analysis including only the cases with a mammogram obtained at least 6 months before cancer diagnosis had comparable results (see the central columns in Table 3). Finally, given that most mammograms from BC cases were analog, the last columns present the association between MD and BC using only analog images (Table 3). ORs tended to be stronger in this case, particularly for BRCA2 mutation carriers. The analysis of digital images was hampered by the small number of BC cases available (40 cases: 6 MD <=10 %, 6 MD = 11-25 %, 17 MD = 26-50 %, 9 MD = 51-75 % and 2 MD >75 %) and no association between MD and BC was observed (ORs of 1.23, 1.94, 1.88 and 1.22, respectively; none of the ORs were statistically significant).

The association between MD (as an ordinal variable) and BC per category of the other explanatory variables among mutation carriers is presented in Fig. 2. The OR represents the risk excess associated with an increase in one category of MD. In general, a consistent effect of MD on BC risk was seen within all subgroup analyses. Even though the association seemed to be less marked among women older than 45 years at the time of mammography, those who were nulliparous, and those who had not taken OC, none of these differences were statistically significant. The smaller number of women who reported the use of HRT was too small to reach conclusions in this group. The association with the type of image was stronger for analog mammograms, but most cases had analog mammograms, whereas the opposite was true for non-cases. Finally, MD increased the risk of all pathologic subtypes BC subtypes in a similar way (P value for heterogeneity = 0.638).

Discussion

In this case-control study of women from high-risk breast cancer families, we confirmed that breast density was an independent risk factor for BC among *BRCA1* and *BRCA2* mutation carriers. Compared to women with breast density <10 %, those with densities >50 % had at



Mammographic Non-cases density		All cases included				Excluding cases without a mammogram taken at least 6 months before diagnosis				Including only analog images				
	Non-cases	BC cases	OR ^a	95 % Cl ^a	P value ^a	BC cases ^a	OR ^a	95 % Cl ^a	P value ^a	Non-cases	BC cases	OR ^b	95 % Cl ^b	P value ^b
BRCA1+BRCA2														
<=10 %	105	35	1.00			12	1.00			36	29	1.00		
1125 %	89	47	1.51	0.74, 3.06	0.255	18	1.76	0.70, 4.41	0.228	29	41	1.74	0.72, 4.19	0.216
26-50 %	151	85	1.85	0.90, 3.82	0.095	39	2.26	0.94, 5.44	0.070	45	68	1.77	0.75, 4.15	0.191
51-75 %	72	47	3.24	1.43, 7.35	0.005	21	3.67	1.37, 9.80	0.010	15	38	4.88	1.72, 13.9	0.003
>75 %	26	34	4.34	1.71, 11.1	0.002	17	8.94	2.86, 28.0	< 0.001	7	32	9.45	2.73, 32.7	< 0.001
Linear trend	443	248	1.45	1.18, 1.78	<0.001	107	1.64	1.28, 2.09	< 0.001	132	208	1.68	1.31, 2.15	< 0.001
BRCA1														
<=10 %	49	15	1.00			3	1.00			15	14	1.00		
11-25 %	56	24	2.18	0.85, 5.58	0.103	8	2.71	0.63, 11.7	0.183	15	20	1.78	0.50, 6.48	0.370
26-50 %	66	36	1.90	0.78, 4.67	0.160	14	3.06	0.73, 12.9	0.127	23	30	1.15	0.39, 3.39	0.798
51-75 %	42	22	2.53	0.89, 7.18	0.082	10	4.22	0.93, 19.2	0.063	9	21	3.68	0.75, 18.1	0.109
>75 %	14	17	5.42	1.79, 16.4	0.003	9	12.0	2.20, 65.1	0.004	3	16	7.49	1.57, 35.7	0.012
Linear trend	227	114	1.37	1.06, 1.76	0.015	44	1.65	1.17, 2.32	0.004	65	101	1.51	1.08, 2.12	0.017
BRCA2														
<=10 %	56	20	1.00			9	1.00			21	15	1.00		
11-25 %	33	23	1.29	0.45, 3.68	0.639	10	1.70	0.47, 6.19	0.419	14	21	2.21	0.76, 6.46	0.146
26-50 %	85	49	2.03	0.65, 6.34	0.222	25	3.36	0.89, 12.7	0.074	22	38	3.85	1.06, 13.9	0.040
51-75 %	30	25	5.37	1.35, 21.4	0.017	11	7.38	1.26, 43.1	0.027	6	17	11.2	1.82, 68.9	0.009
>75 %	12	17	4.82	1.10, 21.1	0.037	8	13.4	2.33, 77.3	0.004	4	16	18.8	3.06, 114.9	0.002
Linear trend	216	134	1.60	1.13, 2.28	0.008	63	1.95	1.30, 2.91	0.001	67	107	2.11	1.40, 3.16	< 0.001

Table 3 Association of mammographic density with subsequent breast cancer in *BRCA1/2* mutation carriers

^aOdds ratio (OR), 95 % CI and P values adjusted for age at mammogram, menopausal status premenopausal, natural menopause, surgical menopause), body mass index, parity (nulliparous, parous), age at first live birth, use of hormonal replacement treatment (yes, no), type of image (analog, digital) and time elapsed from mammogram to breast cancer diagnosis (cases) or end of follow up (non-cases). ^bOR, 95 % CI and P values adjusted for the above-mentioned variables except type of image

least threefold increased risk of developing a breast tumor. The association between MD and the risk of subsequent BC seemed to be stronger for *BRCA2* than for *BRCA1* mutation carriers, but the sample size was insufficient to detect statistical differences between them. Our results also confirmed an association between MD and all pathologic BC subgroups. Finally, we found that mutation carriers had lower MD estimations compared to non-carriers and this difference remained significant among *BRCA2* mutation carriers.

The role of MD as a risk predictor for BC has relevant implications in screening and preventive interventions, particularly in highly motivated women with a family history of the disease. Our results were consistent with early evidence for the association between BC risk and MD among women from high-risk families [6, 7] and later findings on *BRCA1* and *BRCA2* mutation carriers [17, 18], assessed by computerized methods for obtaining quantitative measures of MD. Initial case-control studies evaluating the association between family history, MD and subsequent risk of BC were completed by Martin et al. [7] in a sample of 2,322 subjects from three Canadian screening programs. In that population, MD explained 14 % of the association between family history and BC risk, being considered an intermediate marker for BC. Moreover, women with breast density >50 % had threefold higher risk of BC than those with <10 % density (95 % CI = 2.17, 4.15). Our OR for carriers with >50 % breast density are in agreement with those previously reported in our country among BC screening attendants, using the same semiquantitative visual scale [22]. In that study, the risk excess associated with higher MD was similar in women with and without a family history of BC, similar to the findings of the Nurses' Health Study [23].

Raising the question of the potential impact of MD on women with a germline mutation in *BRCA1* and *BRCA2* genes, Mitchell et al. evaluated the association between MD and BC risk among 342 women from the EMBRACE study, 206 of whom were *BRCA* mutation carriers [17]. In spite of some methodological differences and a limited



stratified by other explanatory variables. Odds ratios (*OR*) and 95 % CI correspond to an increase in one category of MD. Estimators adjusted for age at mammogram, menopausal status, body mass index (*BMI*), parity, age at first live birth, use of hormonal replacement therapy (*HRT*), type of image and time elapsed from mammogram to either breast cancer diagnosis (cases) or end of follow up (non-cases). *P* value reflects the statistical significance of the interaction term between MD and the corresponding variable in all cases except for BC subtype, for which the *P* value is for the test of heterogeneity of the MD effect according to the type of tumor. *HPT* hormone preventative treatment, *ER* estrogen receptor, *PR* progesterone receptor, *HER2* human epidermal growth factor receptor 2

sample of participants, their results were comparable to ours. Assessing MD with a semi-automatic tool, the authors estimated the OR for BC among carriers with ≥ 50 % breast density to be 2.27 (95 % CI = 0.70, 7.39) compared with carriers who had <10 % breast density, while using a qualitative scale (Wolfe's), the highest density group had a risk excess close to 3 (OR = 2.78) [17]. In contrast, a recent prospective nested case-control study investigating the same issue in a cohort of 462 BRCA mutation carriers enrolled in a BC screening program failed to confirm this association [19]. These negative results could be partially explained by facts such as the relatively small sample size and small number of cases, the high proportion of patients with a prior diagnosis and treatment of BC (19%), the older mean age at the baseline mammogram, and the lack of information on BMI, one of the most important established confounding factors for MD. This confounding effect is particularly relevant when MD is assessed using computer-assisted tools, given the strong correlation between BMI and the non-dense area [24, 25]. Due to the potential impact of cancer treatment on MD, and similar to the EMBRACE study, we restricted our study to include only women with first diagnosis of BC as cases. Indeed, Passaperuma et al. did not provide information relative to adjuvant cancer therapy, which could have acted as a confounding factor [19]. In fact, based on the effect of tamoxifen in reducing MD, research has been lately focused on evaluating the potential use of MD variation as a target for identifying women who would benefit from this drug either as preventive or adjuvant therapy [26–28].

Despite initial conflicting results from a small series of patients, two later studies investigating the effect of mutation status on MD patterns have reported similar results. The first study was based on 505 retrospectively obtained mammographic readings from 342 women and found no significant difference in MD between carriers and non-carriers, either as a group or when mutated genes were considered separately [17]. In fact, the mean density was slightly lower in carriers than non-carriers. A second study compared MD between 143 healthy carriers and 119 women determined to be at low to average risk of BC [18]. In agreement with our results, the authors also found a marginally lower, albeit not significant, MD among carriers after adjustment for age and BMI. In our study, based on a larger sample size, mutation carriers had lower breast density compared to non-carriers, although this effect was only statistically significant for *BRCA2* mutation carriers.

As regards BC subtypes, previous studies have addressed the association between MD and BC in the general population with inconsistent results [22, 29–32]. Therefore, whereas a few studies have suggested that MD is mainly associated with hormone receptor positive tumors [32], others have reported a similar [22, 30] or event stronger association with triple-negative tumors, especially for women aged <55 years [29]. In addition, and similar to our results in *BRCA1/2* mutation carriers, a recent study in screening participants in Spain proved that high MD was associated with all histologic subtypes of BC [22].

Our study had several limitations, mostly related to its retrospective nature. Indeed, the heterogeneity of mammograms due to the use of multiple mammographers, the inclusion of both digital and analog images and differences in methodology, such as the subjective visual assessment of breast density and the use of semiquantitative assessments, limit the comparability of our results with other studies. Regarding the use of digital and analog mammograms, density tends to be lower when estimated via digital mammography and most images for BC cases were analog (84 % analog versus 16 % digital). Owing to this, we were able to show a clear effect of MD on BC risk in general and also restricting the analysis to cases and controls with analog images, but no effect was observed using digital mammograms, probably due to the small number of BCs in this category and the fact that only eight of them were classified in the extreme categories. Future studies are needed to confirm these results in mutation carriers studied with digital mammograms. Last, although the plan in original design of our study was to use healthy true negative BRCA1/2 mutation carriers and phenocopies as controls for comparing the effect of MD on cancer risk between carriers and non-carriers, the smaller number of cases in this group did not allow us to accomplish this goal.

The strengths of our study include the fact that this is the largest cohort of women with a *BRCA1/2* mutation and MD data ever published. In spite of being a multicentric study, a single blinded radiologist, who is a highly reputed expert on mammographic reading, with a very high internal consistency [20], assessed all mammograms using a semiquantitative scale that has proved to be associated with subsequent development of BC [22]. Finally, restricting the analysis to BC patients with mammograms obtained 6 months before the diagnosis and the adjustment for all known confounders of MD provides compelling support for our results.

Conclusions

This is the largest study to date to assess the influence of MD on BC risk and is concordant with previously published results in that MD is an independent risk factor among *BRCA1/2* mutation carriers. Even though MD was not increased among mutation carriers compared to non-carriers, high MD increased the risk of all pathologic subtypes of BC. Prospective studies are needed to clarify the impact of density variations on BC risk and provide a more pro-active prevention strategy for women harboring *BRCA1/2* mutations.

Abbreviations

BC: breast cancer; BMI: body mass index; ER: estrogen receptor; HER2: human epidermal growth factor receptor 2; HPT: hormone preventative treatment; HRT: hormonal replacement treatment; MD: mammographic density; OC: oral contraceptives; OR: odds ratio; PR: progesterone receptor.

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

MP, TRyC, ICh and DST conceived the study and discussed their idea with the other authors, and all of them contributed to the design of the research proposals that provided financial support. JM, VL and DF helped design the questionnaire and were in charge of organizing mammographic readings. TRyC, ICh, AT, AI, JB, ABSH, GLI, EHA, MJJF, IT, LR, CGP, PPS, MSLM, SHP, MS, JB and AB recruited women from *BRCA1/2* families and collected the mammograms and information needed together with the pathologic records of BC cases. MP performed the statistical analysis and results were circulated and commented on by all the authors. MP, TRyC and ICh drafted the first version that was critically reviewed by the other authors, who made suggestions on how to improve it. All of them have read and approved the final version of the manuscript.

Acknowledgements

This work was supported by three research grants, PS09/01006, PS09/01024 and PS09/01721, from Spain's Health Research Fund (Fondo de Investigación Sanitaria); and a grant from the Spanish Federation of Breast Cancer Patients (Federación Española de Cáncer de Mama) (FECMA 485 EPY 1170-10). The authors also acknowledge the contribution of the Asociación Española contra el Cáncer (AECC), Carlos III Health Institute; RTICC; and the Catalan Health Institute and Autonomous Government of Catalonia (grant numbers: RD06/0020/1051, RD06/0020/1050, RD12/0036/0008, RD12/0036/0031, PI10/01422, PI13/00285, PIE13/00022 and SGR290). This project was approved by the following ethical boards: Comité de Ética de la Investigación y de Bienestar Animal CElyBA, Instituto de Salud Carlos III; Comité de Ética de la Investigación Clínica Hospital de la Santa Creu y Sant Pau, Barcelona; Comité de Ética de la Investigación Clínica Hospital Universitari de Girona Doctor Josep Trueta, Girona; Comité de Ética de la Fundación INCLIVA, Hospital Clínico Universitario, Valencia; Comité de Ética de la Investigación Clínica del Hospital Universitario 12 de Octubre, Madrid, and Comité de Ética de la Investigación Clínica del Hospital Ramón y Cajal, Madrid.

Author details

¹Medical Oncology Department, Hospital Santa Creu I Sant Pau, Barcelona, Spain. ²Medical Oncology Department, Hospital Clinico Universitario de Valencia, Valencia, Spain. ³Foundation General Directorate Public Health and Foundation for the Promotion of Health and Biomedical Research in the Valencian Region, FISABIO – Public Health, Valencia, Spain. ⁴Hereditary Cancer Program, Catalan Institue of Oncology-IDIBELL, Barcelona, Spain. ⁵Hereditary Cancer Program, Catalan Institute of Oncology-IDIBGI, Girona, Spain. ⁶Medical Oncology Deartment, Hospital Vall Hebron/Vall Hebron Institute of Oncology, Barcelona, Spain. ⁷Medical Oncology Department, Hospital Universitario de Elche, Elche, Spain. ⁸Genetic Counseling Unit, Corporació Sanitaria Parc tauli, Consorci Sanitari de Terrassa, Terrasa, Spain. ⁹National Center for Epidemiology, Carlos III Institute of Health, Monforte de Lemos 5, 28029 Madrid, Spain. ¹⁰Consortium for Biomedical Research in Epidemiology and Public Health (CIBERESP), Carlos III Institute of Health, Madrid, Spain. ¹¹Consortium Cancer Epidemiology Research Group, Oncology and Hematology Area, IIS Puerta de Hierro (IDIPHIM), Madrid, Spain. ¹²Breast Cancer Unit, Clinical Research Programme, Spanish National Cancer Center (CNIO), Madrid, Spain. ¹³Medical Oncology Department, Foundation of the Valencian Oncologic Institute, Valencia, Spain. ¹⁴Medical Oncology Department, Hospital Provincial de Castellón, Castellón, Spain. ¹⁵Medical Oncology Department, Hospital 12 de Octubre, Madrid, Spain. ¹⁶Medical Oncology Department, Hospital Universitario Ramón y Cajal, IRYCIS, Madrid, Spain. ¹⁷Medical Oncology Department, Hospital Clínico San Carlos Madrid Spain. ¹⁸Gynecological Department, Hospital Gregorio Marañón, Madrid, Spain. ¹⁹Medical Department, Hospital Fundación Alcorcón, Alcorcón, Spain. ²⁰Medical Sciences Department, School of Medicine, University of Girona, Girona, Spain.

Received: 15 December 2014 Accepted: 24 June 2015 Published online: 11 July 2015

References

- McCormack VA, dos Santos Silva I. Breast density and parenchymal patterns as markers of breast cancer risk: a meta-analysis. Cancer Epidemiol Biomarkers Prev. 2006;15:1159–69.
- Kataoka M, Antoniou A, Warren R, Leyland J, Brown J, Audley T, et al. Genetic models for the familial aggregation of mammographic breast density. Cancer Epidemiol Biomarkers Prev. 2009;18:1277–84.
- Haars G, van Noord PA, van Gils CH, Peeters PH, Grobbee DE. Heritable aspects of dysplastic breast glandular tissue (DY). Breast Cancer Res Treat. 2004;87:149–56.
- Kaprio J, Alanko A, Kivisaari L, Standertskjold-Nordenstam CG. Mammographic patterns in twin pairs discordant for breast cancer. Br J Radiol. 1987;60:459–62.
- Boyd NF, Dite GS, Stone J, Gunasekara A, English DR, McCredie MR, et al. Heritability of mammographic density, a risk factor for breast cancer. N Engl J Med. 2002;347:886–94.
- Boyd NF, Lockwood GA, Martin LJ, Knight JA, Jong RA, Fishell E, et al. Mammographic densities and risk of breast cancer among subjects with a family history of this disease. J Natl Cancer Inst. 1999;91:1404–8.
- Martin LJ, Melnichouk O, Guo H, Chiarelli AM, Hislop TG, Yaffe MJ, et al. Family history, mammographic density, and risk of breast cancer. Cancer Epidemiol Biomarkers Prev. 2010;19:456–63.
- Lindstrom S, Vachon CM, Li J, Varghese J, Thompson D, Warren R, et al. Common variants in ZNF365 are associated with both mammographic density and breast cancer risk. Nat Genet. 2011;43:185–7.
- Vachon CM, Scott CG, Fasching PA, Hall P, Tamimi RM, Li J, et al. Common breast cancer susceptibility variants in LSP1 and RAD51L1 are associated with mammographic density measures that predict breast cancer risk. Cancer Epidemiol Biomarkers Prev. 2012;21:1156–66.
- Fernandez-Navarro P, Pita G, Santamarina C, Moreno MP, Vidal C, Miranda-Garcia J, et al. Association analysis between breast cancer genetic variants and mammographic density in a large population-based study (Determinants of Density in Mammographies in Spain) identifies susceptibility loci in TOX3 gene. Eur J Cancer. 2013;49:474–81.
- 11. Chang J, Yang WT, Choo HF. Mammography in Asian patients with BRCA1 mutations. Lancet. 1999;353:2070–1.
- 12. Tilanus-Linthorst M, Verhoog L, Obdeijn IM, Bartels K, Menke-Pluymers M, Eggermont A, et al. A BRCA1/2 mutation, high breast density and

prominent pushing margins of a tumor independently contribute to a frequent false-negative mammography. Int J Cancer. 2002;102:91–5.

- Hamilton LJ, Evans AJ, Wilson AR, Scott N, Cornford EJ, Pinder SE, et al. Breast imaging findings in women with BRCA1- and BRCA2-associtaed breast carcinoma. Clin Radiol. 2004;59:895–902.
- Kaas R, Kroger R, Peterse JL, Hart AA, Muller SH. The correlation of mammographic-and histologic patterns of breast cancers in BRCA1 gene mutation carriers, compared to age-matched sporadic controls. Eur Radiol. 2006;16:2842–8.
- Huo Z, Giger ML, Olopade OI, Wolverton DE, Weber BL, Metz CE, et al. Computerized analysis of digitized mammograms of BRCA1 and BRCA2 gene mutation carriers. Radiology. 2002;225:519–26.
- Li H, Giger ML, Huo Z, Olopade OI, Lan L, Weber BL, et al. Computerized analysis of mammographic parenchymal patterns for assessing breast cancer risk: effect of ROI size and location. Med Phys. 2004;31:549–55.
- Mitchell G, Antoniou AC, Warren R, Peock S, Brown J, Davies R, et al. Mammographic density and breast cancer risk in BRCA1 and BRCA2 mutation carriers. Cancer Res. 2006;66:1866–72.
- Gierach GL, Loud JT, Chow CK, Prindiville SA, Eng-Wong J, Soballe PW, et al. Mammographic density does not differ between unaffected BRCA1/2 mutation carriers and women at low-to-average risk of breast cancer. Breast Cancer Res Treat. 2010;123:245–55.
- Passaperuma K, Warner E, Hill KA, Gunasekara A, Yaffe MJ. Is mammographic breast density a breast cancer risk factor in women with BRCA mutations? J Clin Oncol. 2010;28:3779–83.
- Garrido-Estepa M, Ruiz-Perales F, Miranda J, Ascunce N, Gonzalez-Roman I, Sanchez-Contador C, et al. Evaluation of mammographic density patterns: reproducibility and concordance among scales. BMC Cancer. 2010;10:485.
- Hardin JW, Hilbe JM. Generalized Linear Models and Extensions. College Station: Stata Press; 2007.
- Pollan M, Ascunce N, Ederra M, Murillo A, Erdozain N, Ales-Martinez J, et al. Mammographic density and risk of breast cancer according to tumor characteristics and mode of detection: a Spanish population-based case–control study. Breast Cancer Res. 2013;15:R9.
- Yaghjyan L, Coldiz GA, Rosner B, Tamimi RM. Mammographic breast density and breast cancer risk by menopausal status, postmenopausal hormone use and a family history of breast cancer. Cancer Causes Control. 2012;23:785–90.
- 24. Assi V, Warwick J, Cuzick J, Duffy SW. Clinical and epidemiological issues in mammographic density. Nat Rev Clin Oncol. 2012;9:33–40.
- Nguyen TL, Schmidt DF, Makalic E, Dite GS, Stone J, Apicella C, et al. Explaining variance in the cumulus mammographic measures that predict breast cancer risk: a twins and sisters study. Cancer Epidemiol Biomarkers Prev. 2013;22:2395–403.
- Cuzick J, Warwick J, Pinney E, Duffy SW, Cawthorn S, Howell A, et al. Tamoxifen-induced reduction in mammographic density and breast cancer risk reduction: a nested case–control study. J Natl Cancer Inst. 2011;103:744–52.
- Ko KL, Shin IS, You JY, Jung SY, Ro J, Lee ES. Adjuvant tamoxifen-induced mammographic breast density reduction as a predictor for recurrence in estrogen receptor-positive premenopausal breast cancer patients. Breast Cancer Res Treat. 2013;142:559–67.
- Kim J, Han W, Moon HG, Ahn S, Shin HC, You JM, et al. Breast density change as a predictive surrogate for response to adjuvant endocrine therapy in hormone receptor positive breast cancer. Breast Cancer Res. 2012;14:R102.
- Bertrand KA, Tamimi RM, Scott CG, Jensen MR, Pankratz V, Visscher D, et al. Mammographic density and risk of breast cancer by age and tumor characteristics. Breast Cancer Res. 2013;15:R104.
- Phipps AI, Buist DS, Malone KE, Barlow WE, Porter PL, Kerlikowske K, et al. Breast density, body mass index, and risk of tumor marker-defined subtypes of breast cancer. Ann Epidemiol. 2012;22:340–8.
- Yaghjyan L, Colditz GA, Collins LC, Schnitt SJ, Rosner B, Vachon C, et al. Mammographic breast density and subsequent risk of breast cancer in postmenopausal women according to tumor characteristics. J Natl Cancer Inst. 2011;103:1179–89.
- Conroy SM, Pagano I, Kolonel LN, Maskarinec G. Mammographic density and hormone receptor expression in breast cancer: the Multiethnic Cohort Study. Cancer Epidemiol. 2011;35:448–52.