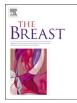
The Breast 55 (2021) 98-104



Contents lists available at ScienceDirect

The Breast



journal homepage: www.elsevier.com/brst

Original article

The effect of family history on screening procedures and prognosis in breast cancer patients - Results of a large population-based casecontrol study



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ARTICLE INFO

Article history: Received 14 August 2020 Received in revised form 6 December 2020 Accepted 18 December 2020 Available online 24 December 2020

Keywords: Breast cancer Screening Familial risk Prognosis Mammography

ABSTRACT

Background: The potential benefit of additional breast cancer screening examinations in moderate risk patients (patients with a history of breast cancer in one or two family members) remains unclear. *Methods:* A large population-based case—control study on breast cancer in postmenopausal women in Germany recruited 2002–2005 (3813 cases and 7341 age-matched controls) was used to assess the association of family history with breast cancer risk. Analysis of family history, participation in screening procedures, and tumor size regarding prognosis in patients was based on follow-up data until 2015. *Results:* A first degree family history of breast cancer was associated with higher breast cancer risk (OR 1.39, p < 0.001). Patients with a first degree family history of breast cancer were more likely to have had >10 mammograms (MG) (42.7% vs. 24.9%, p < 0.001) and showed a higher rate of imaging-detected tumors (MG or ultrasound) (45.8% vs. 31.9%, p < 0.001). A smaller tumor size at initial diagnosis (below 2 cm) was more likely in patients with a positive family history (OR 1.45, p < 0.001) and a higher number of MG (\geq 10 MG: OR 2.29). After accounting for tumor characteristics, mammogram regularity (HR 0.72, p < 0.001) and imaging-assisted tumor detection (HR 0.66, p < 0.001) were associated with better overall survival but not with a positive family history.

Discussion: Patients with a positive family history had a higher rate of imaging detected tumors with smaller size at initial diagnosis compared to patients without affected family members. Screening was associated with improved survival after a breast cancer diagnosis, irrespective of a positive family history. © 2020 The Authors. Published by Elsevier Ltd. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/).

1. Introduction

There is evidence that public breast cancer screening programs regardless of preexisting risk factors are beneficial in terms of reduced breast cancer mortality for women aged 50–69 years [1]. Besides that, a benefit of additional screening visits has also been documented in high-risk patients (e.g. BRCA 1 or BRCA 2 gene mutations) [2]. However, there is still a high rate of opportunistic screening procedures regardless of screening programs, especially

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in patients with family history of breast cancer [3]. Although the potential benefit of additional screening examinations in moderate risk patients (patients with a history of breast cancer in one or two family members) remains unclear [4], participation in these additional exams is supported by German statutory health insurances even if only one first-degree family member has a history of breast cancer [5]. Therefore, many women with a first- or second-degree relative with breast cancer are participating more frequently in breast cancer screening procedures outside mammography screening programs that offer screening every 2–3 years.

In this analysis, we first investigated the association of family history of breast cancer with breast cancer risk. We then examined in breast cancer patients the relationships between family history

https://doi.org/10.1016/j.breast.2020.12.008

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of breast cancer, frequency of screening procedures in a moderate risk situation (e.g. one or more affected family members), and tumor characteristic. Lastly, we assessed whether more frequent screening due to a positive family history was associated with overall and breast cancer specific survival after accounting for clinical and tumor characteristics.

2. Materials and methods

2.1. Study population and data source

The cohort consisted of 3813 cases and 7341 controls of the population-based case-control MARIE study (Mamma Carcinoma Risk factor Investigation) [6]. Patients were recruited in two regions of Germany; Hamburg and Rhine-Neckar-Karlsruhe if they were aged 50-74 years at diagnosis and had a histologically confirmed diagnosis of primary invasive (stage I to IV) or in situ breast cancer between January 1, 2001 and September 30, 2005. Two populationbased controls without a history of breast cancer matched by age and study region were recruited for each case. Information on prediagnostic lifestyle factors, socio-economic status, medical history, and information on specific medications, regimen, and duration of use was collected by a standardized face-to-face interview at recruitment [6]. The histological characteristics of the primary breast cancer were extracted from pathology reports. Treatment and clinical course were abstracted from medical records to verify clinical events either self-reported in the interview or reported by treating physicians during a first follow-up in 2009 and a second follow-up in June 2015 (in total > 90% self-reported events verified) resulting in a cohort study with a follow-up time of >10 years [7].

All study participants gave written informed consent. Ethics approval was obtained from the ethics committee of the University of Heidelberg, the Hamburg Medical Council, and the Medical Board of the State of Rhineland-Pfalz. The study was conducted in accordance with the Declaration of Helsinki.

Information on family history of breast cancer was collected by face-to-face interview at recruitment. Participants were asked whether family members were affected by breast cancer and other cancers, and to provide the information separately for mother, sister and daughter (first degree) and for second degree relatives.

2.2. Outcome assessment

Study participants were prospectively followed until June 30, 2015. Vital status was assessed via information from population registries. Causes of death were derived from death certificates obtained through the local/regional health offices and coded according to the 10th revision of the International Classification of Diseases (ICD-10-GM). The primary endpoints were overall survival (OS: including death from any cause), breast cancer-specific survival (BCS (non-breast cancer-related deaths were censored), and recurrences (including ipsilateral, contralateral, local/regional invasive recurrence, and distant recurrence). Participants without an event of interest were censored at the date of last contact or on June 30, 2015, whichever came first.

2.3. Statistical analysis

Case-control data were reported as mean \pm standard deviation (SD) or frequency (percent) for breast cancer cases and controls. Pearson's Chi²-tests and ANOVA were used for comparisons between groups.

To estimate the association between a positive 1st degree and 2nd degree family history and overall breast cancer risk, univariate and multivariable logistic regression models adjusted for diagnosis/ interview age (years), number of relatives with BC (1, \geq 2 or missing) age at first birth (\leq 21, 22–24, 25–28, 29+ years and missing), mammography ever (no/yes or missing) as well as benign breast disease in the past (y/n) were performed to estimate odds ratios (ORs) and 95% confidence intervals (95% Cl). Logistic regression analyses in cases only were performed to estimate associations between the frequency of participation in screening procedures and the tumor size or nodal status, respectively. Having a small tumor (below 2 cm versus above) or being node-negative (pN0 versus other) was the dependent variable. The categorial variable for family history served as adjusting variable.

The Kaplan-Meier method has been used to compare overall survival (OS) and breast cancer specific survival (BCS) with respect to the number of pre-diagnosis performed mammograms and to family history of breast cancer. The log-rank test has been applied to determine differences between the groups. Multivariable Cox proportional hazards regression analysis adjusted for tumor size, nodal status, grading, hormone receptor status, MG regularity, tumor detection mode and family history of breast cancer was conducted to estimate the association of the participation frequency in screening procedures, tumor size and nodal status with OS. To investigate if a positive family history is a potential effect modifier of the association of the frequent participation in screening procedures and OS, an interaction term was included in the Cox model.

All statistical analyses were two-sided at significance level $\alpha=0.05.$

All analyses were performed on all-available-cases basis. To keep as many observations as possible in the analyses, missing values in categorical variables were assigned an own category.

3. Results

3.1. Patient characteristics

There were no differences between 3813 cases and 7341 agematched controls with regard to age (mean age 62 y), menopausal status, age at first delivery, body mass index, education levels and use of contraceptive pills (Table 1). There was a high rate of childbirth in both groups (88.5% vs. 85.9%). There was a higher proportion of cases than controls who never breastfed (never breastfed: cases: 37.6% vs. controls 33.4%, p < 0.001, Table 1). On time of study inclusion, there was a higher rate of cases who were currently using hormone replacement therapy compared to controls (46.5%, vs. 33.3%, p < 0.001). There was no difference in having clinical examinations like breast palpation, but there was a higher proportion of cases who had regular MG compared to controls (61.5% vs. 53.6% in controls, p < 0.001), more than 10 performed MG before diagnosis of breast cancer (>10 MG, cases 28.9%, controls 20.7%, p < 0.001) and a history of benign breast diseases (41.9% vs. 34.2%, p < 0.001, Table 1).

3.2. Association of family history of breast cancer with breast cancer risk

Results of univariate logistic regression model analyses showed a higher proportion of cases with family history of breast cancer of a first degree relative (18.2% vs. 12.2%; OR 1.60, 95% confidence interval (Cl) 1.44–1.79, p < 0.001) and a second degree relative only (14.1% vs. 11.4%, OR 1.28, 95% CI: 1.14–1.44, p < 0.001) compared to controls (data not shown)).

There were no differences in breast cancer subtypes between cases with and cases without positive family history of breast cancer (p = 0.073). In this group of postmenopausal breast cancer patients with HR + disease, 18.5% (433/2338) of patients had a positive family history of breast cancer compared with 16.93% (107/

Table 1

≥3

39 (0.3%)

100

Characteristics of the MARIE study population.

	TOTAL	CONTROL	CASES	P-VALUE		
	N = 11,154	N = 7341	N = 3813			
Age, years						
Mean \pm SD	62 <u>±</u> 6.03)	62 (±6.02)	62 (±6.05)	0.274		
Age at Menopause, yea	rs					
Mean \pm SD	49 (±4.9)	49 (±5.0)	49 (±4.8)	< 0.001		
Age at first birth, years						
Mean \pm SD	24 (±4.6)	24 (±4.6)	24 (±4.7)	0.301		
BMI kg/m2	4000 (44 70)	2206 (44 0%)	1000 (44 5%)	0.110		
<22.5 22.5-<25	4982 (44.7%)	. ,	1696 (44.5%)	0.110		
22.3-<25	3453 (31.0%) 2274 (20.4%)	• • •	1209 (31.7%) 780 (20.5%)			
>30	439 (3.9%)	311 (4.2%)	128 (3.4%)			
missing	6	6	0			
Education						
low	6360 (57.0%)	4180 (56.9%)	2180 (57.2%)	0.881		
medium	3134 (28.1%)	2074 (28.3%)	1060 (27.8%)			
high	1658 (14.9%)	1087 (14.8%)	571 (15.0%)			
missing	2	0	2			
Pregnancy ever	1005 (10 400	0.45 (4.4 500)	500 (4 4 4 4)	0.001		
No	1385 (12.4%)	847 (11.5%)	538 (14.1%)	<0.001		
yes	9769 (87.6%)	6494 (88.5%) 1	3275 (85.9%)			
missing Age at first birth	0	1	0			
no pregnancy	1844 (16.5%)	1162 (15.8%)	682 (17.9%)	0.015		
<21 y	2697 (24.2%)	1797 (24,5%)		5.010		
22-24y	2504 (22,5%)					
25-28y	2383 (21.4%)		774 (20.3%)			
29 + y	1722 (15.4%)	1108 (15.1%)	614 (16.1%)			
missing	4	3	19			
Ever breastfed						
no	3885 (34.8%)	2452 (33.4%)	1433 (37.6%)	<0.001		
yes	7268 (65.2%)	4889 (66.6%)	2379 (62.4%)			
missing Use of contraceptive pi	1	0	1			
no	3985 (36.1%)	2610 (35.9%)	1375 (36.4%)	0.629		
yes	7066 (63.9%)	. ,	2406 (63.6%)	01020		
missing	103	71	32			
Use of hormone replac	ement therapy	/				
never	4356 (39.4%)	3071 (42.2%)	1285 (33.9%)	< 0.001		
past	2524 (22.8%)		742 (19.6%)			
current	• •	2425 (33.3%)	1759 (46.5%)			
missing	90	63	27			
Cinical examination/pa no	222 (2%)	105 (1.4%)	117 (3.1%)	<0.001		
yes		7229 (98.5%)	3694 (96.9%)	<0.001		
missing	9	7	2			
Mammogram regularit			-			
no	3583 (32.1%)	2572 (35.0%)	1011 (26.5%)	< 0.001		
yes	6281 (56.3%)	3935 (53.6%)	2346 (61.5%)			
no mammograms at all						
missing	45 (0.4%)	24 (0.3%)	21 (0.5%)			
Number of mammogra		910 (119)	125 (11 10)	<0.001		
0 1-4	1245 (11.2%)	3259 (44.4%)	435 (11.4%) 1293 (33.9%)	<0.001		
5-10		1745 (23.8%)				
>10		1527 (20.7%)	. ,			
Missing	0	0	0			
Benign breast diseases						
no	7024 (63.2%)	4815 (65.8%)	2209 (58.1%)	< 0.001		
yes	4096 (36.8%)	. ,	1597 (41.9%)			
missing	34	25	25			
first degree relative wi			2042 (01 223)	.0.001		
no	,	6067 (87,8%) 842 (12,2%)	2942 (81,8%)	<0.001		
yes missing	1499 (14,3%) 646	843 (12,2%) 431	656 (18,2%) 215			
second degree relative			21J			
No	-	6506 (88.6%)	3275 (85.9%)	< 0.001		
Yes	1373 (12.3%)	• • •	538 (14.1%)			
missing	0	0	0			
first and second degree relatives with breast cancer						
Number of relatives						
1	2591 (23.2%)	• • •	1061 (27.8%)	<0.001		
2	242 (2.2%) 39 (0.3%)	131 (1.8%) 17 (0.2%)	111 (2.9%) 22 (0.6%)			

Table 1	(continued)	
I able I	(continued)	

	TOTAL N = 11,154	$\begin{array}{l} \text{CONTROL} \\ \text{N} = 7341 \end{array}$	CASES N = 3813	P-VALUE
0	7636 (74.3%)	5232 (77.2%)	2404 (70.7%)	
missing	646	431	215	

632) in the HER2 positive subgroup and 15.58% (62/398) in the triple-negative subgroup.

Multivariate analysis showed a significantly higher breast cancer risk for participants with a positive family history of breast cancer (1 affected family member: OR 1.39, 95% CI: 1.26–1.54, p < 0.001, 2 or more affected family members: OR 1.75, 95% CI: 1.45–2.11, p < 0.001) and for women with a history of benign breast diseases (OR 1.40, 95% CI: 1.29–1.52, p < 0.001, Table 2). Reduced breast cancer risk was shown for patients with lower age at first birth (25–28 years: OR 0.80, 95% CI 0.71–0.91: p = 0.001) and also for patients who ever had a MG (OR 0.83, 95% CI: 0.73–0.94, p = 0.004, Table 2). There was no significant association for age (Table 2).

3.3. Relationship between family history of breast cancer and performed screening procedures as well as tumor detection

Among breast cancer patients, those with a first degree positive family history received a higher number of MG than those without positive family history of breast cancer (>10 MG: 42.7% vs. 24.9%, p < 0.001, Table 3). Also patients with a second degree positive family history of breast cancer received a higher number of MG (>10 MG: 34.0% vs. 27.2%, p = 0.009, Table 3) compared to patients without positive family history of breast cancer.

Breast cancer was imaging-detected by MG or ultrasound in 35.5% of all cases with a higher rate of imaging-detected cancers in patients with positive family history of breast cancer compared to patients without positive family history (first degree family history 48.6% vs. 33.1%, p < 0.001; second degree family history 42.7% vs. 34.3%, p = 0.001, Table 3).

3.4. Family history of breast cancer and screening effect

Patients with a positive family history (OR 1.45, 95% CI: 1.27–1.66, p < 0.001) and patients who received a higher number of MG (\geq 10 MG: OR 2.29, 95% CI: 1.88–2.81, p < 0.001) were more likely to be diagnosed with smaller tumor size at initial diagnosis (tumor size below and above 2 cm) compared to patients without positive family history and patients without any performed MG (Suppl. Table 1). These patients were also more likely to be diagnosed with node-negative than node-positive disease (with positive family history (OR 1.38, 95% CI 1.216–1.568, p < 0.001) and with higher number of MG (\geq 10 MG: OR 1.58, 95% CI 1.329–1.873 p < 0.001)) (*data not shown*).

Kaplan-Meier estimates showed significantly improved overall survival rates (5-year-cumulative risk of death 5 vs. 20%, p < 0.001, Fig. 1) and breast cancer specific survival rates in patients with a higher number of performed MG (5-year-cumulative risk of death 5 vs. 15%, p < 0.001). Patients with relatives with breast cancer also had improved overall survival rates (5-year-cumulative risk of death 5 vs. 12%, p = 0.006, Fig. 2) and breast cancer specific survival rates (5-year-cumulative risk of death 3 vs. 6%, p = 0.025).

Results of multivariate cox-proportional hazard regression analyses showed that prognostic factors like small tumor size, nodenegative disease, lower grading (G1, G2), positive hormone receptor status and older age were associated with improved overall survival. Even after adjustment for tumor characteristics,

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Table 2

Univariate and multivariate logistic regression analysis of breast cancer risk.

	UNIVARIATE		MULTIVARIATE	
Variable	OR (95% CI)	p-value	OR (95% CI)	p-value
Age, years	1.00 (0.98-1.01)	0.274	1.00 (0.99-1.01)	0.138
Number of relatives with breast	cancer			
Ref.: no relatives with BC			1.00	
1	1.41 (1.28-1.56)	< 0.001	1.39 (1.26-1.54)	< 0.001
≥ 2	1.77 (1.47-2.12)	<0.001	1.75 (1.45-2.11)	< 0.001
missing	1.06 (0.89-1.25)	0.522	1.05 (0.89-1.25)	0.53558
Mammography ever				
Ref.: no			1.00	
yes	0.96 (0.85-1.09)	0.550	0.83(0.73-0.94)	0.004
Missing	1.24 (0.21-7.55)	0.813	1.17 (0.19-7.25)	0.866
Age at first birth				
Ref.: no pregnancy			1.00	
≤21 y	0.85 (0.75-0.97)	0.012	0.86 (0.75-0.98)	0.019
22-24y	0.86 (0.76-0.98)	0.022	0.86 (0.76-0.97)	0.017
25-28y	0.82 (0.72-0.93)	0.002	0.80 (0.71-0.91)	0.001
29 + y	0.94 (0.82-1.08)	0.410	0.93 (0.81-1.07)	0.324
Missing	0.57 (0.06-5.48)	0.624	0.58 (0.06-5.57)	0.635
Benign breast disease				
Ref.: no			1.00	
yes	1.39 (1.28–1.51)	<0.001	1.40 (1.29–1.52)	< 0.001
missing	0.79 (0.37-1.68)	0.534	0.81 (0.38-1.75)	0.598

Table 3

Breast cancer detection in association with family history of breast cancer.

Number of Mammograms	Cases with	Cases without	p-value
	1st degree positive relative	1st degree positive relative	<0.001
0	38 (5.8%)	374 (12.7%)	
1-4	171 (26.0%)	1040 (35.3%)	
5–9	161 (24.5%)	769 (26.1%)	
10+	280 (42.7%)	733 (24.9%)	
missing	6(1,0%)	26 (0.9%)	
Number of Mammograms	Cases with 2nd degree positive relative only	Cases without 1st and 2nd degree positive relative	0.009
0	46 (8.5%)	389 (11.9%)	
1-4	169 (31.4%)	1124 (34.3%)	
5–9	136 (25.3%)	843 (25.7%)	
10+	183 (34.0%)	890 (27.2%)	
missing	4 (0.8%)	29 (0.9%)	
	Cases with	Cases without	
Detection of breast cancer by	1st degree positive relative	1st degree positive relative	p < 0.001
Palpation	334 (50.9%)	1959 (66.6%)	-
Mammogram/ultrasound	319 (48.6%)	979 (33.1%)	
missing	3	10	
Detection of breast cancer by	Cases with 2nd degree positive relative only	Cases without 1st and 2nd degree positive relative	p = 0.001
Palpation	306 (56.9%)	2139 (65.3%)	
Mammogram/ultrasound	230 (42.7%)	1122 (34.3%)	
missing	2	14	

mammogram regularity (HR 0.72, p < 0.001) and imaging-assisted tumor detection (HR 0.66, p = 0.002, Table 4) were associated with improved overall survival in multivariate analysis. After adjusting for screening and tumor characteristics, a positive family history itself was not associated with improved survival (Table 4). There was no significant interaction between family history and screening frequency with respect to overall survival.

4. Conclusion

Our analysis showed that family history of breast cancer was associated with higher breast cancer risk and resulted in a higher participation frequency in breast cancer-screening procedures like MG and ultrasound. Even in a moderate risk situation according to family history of breast cancer (e.g. one affected family member), patients participated more often in screening procedures and had a higher number of performed MG before diagnosis. Consecutively, there was also a higher rate of imaging-detected tumors in this moderate risk cohort, which resulted in smaller tumor size, less affected lymph nodes and better prognosis. However, there was no direct association between a positive family history of breast cancer and survival after adjusting for screening and tumor characteristics in the multivariate analysis.

Screening for breast cancer aims to reduce mortality from this cancer, as well as the morbidity associated with advanced stages of the disease, through early detection in asymptomatic women. There is still controversy about the benefit of breast cancer screening programs and concerns regarding overdiagnosis. Recently published data of 323,719 women participating in the German mammography screening program between 2003 and 2014 showed an increase of early stage breast cancer and a decrease in breast cancer mortality [8]. Also an independent panel of experts evaluated the screening benefit in 2015 and showed that women 50–69 years of age who were invited to attend mammographic screening had, on average, a 40% reduction in the risk of death from breast cancer irrespective of preexisting risk factors [1].

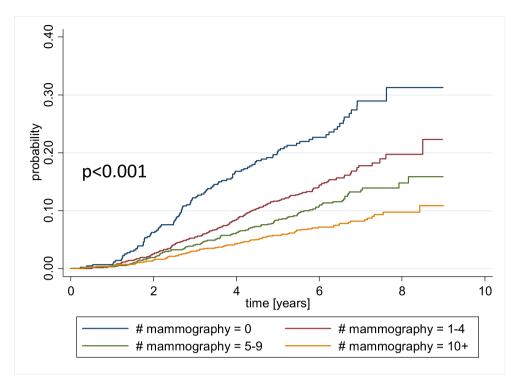


Fig. 1. Kaplan-Meier curves for 10-years cumulative risk of death, stratified by pre-diagnosis performed mammograms.

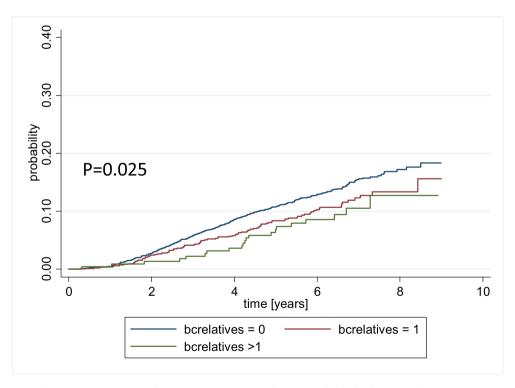


Fig. 2. Kaplan-Meier curves for 10-years cumulative risk of death, stratified by family history of breast cancer.

The German Consortium for Hereditary Breast and Ovarian Cancer (GC-HBOC) established since 1996 a panel of clinical criteria for genetic testing of individuals in a clinical setting, based on familial history of breast and ovarian cancer [9]. A "high risk" is defined as a lifetime risk of breast cancer of more than 30%, whereas other guidelines define a risk of more than 20% as elevated

[10]. A benefit of additional screening visits has been documented in "high-risk" patients (e.g. BRCA 1, BRCA 2 gene mutations) [2]. But it remains unclear how to advise patients, who are not "high-risk" patients but still have a family history of breast cancer, especially in a first or second degree relative. As a consequence of this, there is a high rate of opportunistic screening procedures regardless of

Table 4

Overall survival and risk factors (multivariate Cox-proportional hazard regression analysis).

Overall survivalTumor size pT1 (<2 cm)	Variable	HR (95% CI)	p-value
$\begin{array}{c c c c c c c c c c c c c c c c c c c $		Overall survival	
pT2 (2-5 cm)1.30 (1.04-1.63)0.023pT3+4 (>5 cm,)2.40 (1.77-3.25)<0.001	Tumor size		
pT3+4 (>5 cm,)2.40 (1.77–3.25)<0.001missing2.21 (1.52–3.20)<0.001Nodal statuspN01.00pN1 (1–3)1.72 (1.30–2.13)<0.001	pT1 (<2 cm)	1.00	
missing2.21 (1.52-3.20)<0.001Nodal status $pN0$ 1.00 $pN1$ (1-3)1.72 (1.30-2.13)<0.001	pT2 (2–5 cm)	1.30 (1.04-1.63)	0.023
Nodal statuspN01.00pN1 (1−3)1.72 (1.30−2.13)qN2 (≥4)2.88 (2.34−3.72)Q0012.88 (2.34−3.72)missing7.54 (4.06−13.98)Grading (1.00) G21.99 (1.34−2.95)G32.81 (1.86−4.25)G47.65 (1.72−34.11)Q40.001G47.65 (1.72−34.11)Q640.001Hormone Receptor StatusER/PR positive1.00ER + or PR+1.60 (1.28−2.01)<0.001	pT3+4 (>5 cm,)	2.40 (1.77-3.25)	< 0.001
$\begin{tabular}{ c c c c } \hline 1.00 & 1.00 & 0.001 \\ \hline $pN1$ (1-3)$ & 1.72 (1.30-2.13)$ & <0.001 \\ \hline $pN2$ (≥ 4)$ & 2.88 (2.34-3.72)$ & <0.001 \\ \hline $missing$ & 7.54 (4.06-13.98)$ & <0.001 \\ \hline $Grading$ & & & & & & & & & & & & & & & & & & &$	missing	2.21 (1.52-3.20)	< 0.001
Image: Picture interval and interval an	Nodal status		
$\begin{array}{cccc} {\sf pN2} (\geq 4) & 2.88 (2.34-3.72) & <0.001 \\ {\sf missing} & 7.54 (4.06-13.98) & <0.001 \\ {\sf Grading} & & & & & \\ {\sf G1} & 1.00 & & & \\ {\sf G2} & 1.99 (1.34-2.95) & 0.001 \\ {\sf G3} & 2.81 (1.86-4.25) & <0.001 \\ {\sf G4} & 7.65 (1.72-34.11) & 0.008 \\ {\sf missing} & 3.21 (1.61-6.41) & 0.001 \\ {\sf Hormone} {\sf Receptor Status} & & & \\ {\sf ER/PR positive} & 1.00 & & & \\ {\sf ER/PR positive} & 1.00 & & & \\ {\sf ER/PR negative} & 1.81 (1.44-2.27) & <0.001 \\ {\sf missing} & 0.79 (0.19-3.33) & 0.745 \\ {\sf Age} & 1.03 (1.02-1.04) & <0.001 \\ {\sf Mammogram regularity} & & & \\ {\sf No} & 1.00 & & & \\ {\sf yes} & 0.72 (0.59-0.89) & <0.001 \\ {\sf missing} & 1.16 (0.91-1.48) & 0.236 \\ {\sf Detection tumor by} & & & \\ {\sf Palpation (patient/doctor)} & 1.00 & & \\ {\sf Mammogram/Ultrasound} & 0.66 (0.50-0.85) & 0.002 \\ {\sf missing} & 0.38 (0.05-2.91) & 0.353 \\ \end{array}$	pN0	1.00	
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1 0.95 (0.75–1.20) 0.654	-		
≥2 0.84 (0.54−1.31) 0.447			
missing 1.41 (1.03–1.94) 0.032	missing	1.41 (1.03–1.94)	0.032

screening programs in patients with a moderate risk situation with regard to family history of breast cancer [11]. Even if it is well-known that there is a higher risk for patients in terms of developing breast cancer even if they have just one affected family member [12], the effect of additional screening procedures in this moderate risk cohort remains unclear [3,13].

In our analysis, we could identify a group of patients with moderate risk with regard to family history of breast cancer who were not eligible according to the guidelines for genetic counselling [14,15]. In this group, number of MG before diagnosis and imagingassisted tumor detection was also associated with improved overall survival. There was no difference in the magnitude of association of MG regularity, number of MG and imaging-assisted tumor detection with improved overall survival in patients with a moderate family history and those without a family history of breast cancer. A limitation of our study is the retrospective nature of the collected information. We could not exactly determine the age at breast cancer diagnosis in affected family members in our cohort. Thus, we were not able to differentiate between familial and hereditary breast cancer by the reported family history. Although we found a beneficial screening effect also in patients with a family history of breast cancer, based on the available data we cannot assess which screening method was superior or should be offered. It also remains unclear why patients with a family history of breast cancer in our cohort had a higher rate of performed screening procedures compared to patients without a family history of breast cancer. One reason might be a higher awareness of developing breast cancer by the patients themselves or by their physicians. Also, based on our data, we cannot conclude at what age screening procedures in a moderate risk group should start, although the median age of included patients in our study was 62 years and many patients had more than 10 performed MG before the diagnosis of breast cancer. This is in line with other data, showing low benefit of intensified screening procedures in high-risk patients without BRCA 1 or BRCA 2 gene mutations before the age of 50 [13].

The main strength of this large population-based patient sample with long-term follow-up is the availability of detailed information about family history of breast cancer and participation in screening procedures. To our knowledge, this is the first observational study that was able to show that patients with moderate breast cancer risk are more likely to have received a mammography screening, which is associated with a clinical benefit.

Our analysis showed an association between positive family history and participation in screening procedures, higher number of performed MG, higher rate of imaging-detected tumors as well as better prognostic factors in screen-detected breast cancer. This could be valued as positive screening effect which could support the role of screening even in a cohort of women with elevated breast cancer risk defined by family history of breast cancer. However, our results only show that ever mammography screening is beneficial with respect to survival and this holds whether or not the women had a positive family history. Regular mammography screening (which is only ever MG here) is associated with improved survival because of the more favourable tumor characteristics of screen-detected tumors.

Conclusion

Women with a family history of breast cancer are at higher risk of developing breast cancer. Additional screening procedures showed a benefit in terms of smaller tumor size, less affected lymph nodes at time of diagnosis and better prognosis in this group of breast cancer patients. However, it remains unclear when to start additional procedures and which screening method should be favored.

Funding source

This sub-analysis of the MARIE trial did not receive any specific grant from funding agencies in the public, commercial, or not-forprofit sectors.

Ethical approvements

Ethics approval was obtained from the ethics committee of the University of Heidelberg, the Hamburg Medical Council, and the Medical Board of the State of Rhineland-Pfalz. The study was conducted in accordance with the Declaration of Helsinki.

Declaration of competing interest

All authors declare that they have no conflict of interest.

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