

## ARTICLE

# Evaluation of the Dutch *BRCA1/2* clinical genetic center referral criteria in an unselected early breast cancer population

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In this study, we evaluated the diagnostic value of the Dutch Clinical Genetic Center (CGC) referral guidelines for *BRCA1/2* mutation testing in 903 early breast cancer patients, unselected for family history, diagnosed in a cancer hospital before the age of 50 years in 1974–2002; most prevalent Dutch pathogenic *BRCA1/2* mutations had been analyzed on coded DNA in a research setting. Forty-nine (5.4%) of the patients were proven to be *BRCA1/2* mutation carriers. We found that 78% and 69% of *BRCA1* and *BRCA2* mutation carriers identified met the criteria for referral to the CGC based on age, family history and synchronous multiple tumors; reflected by a combined sensitivity of 75.5% and specificity of 63.2%. More than half of the *BRCA1* mutation carriers, that is, 58% had a triple-negative tumor. The highest AUC was obtained by shifting the age at diagnosis threshold criterion from 40 to 35 years and by adding a 'triple-negative breast cancer' criterion with an age threshold of 45 years; the specificity increased to 71.2%, whereas the sensitivity remained the same; that is, a referral of fewer patients will lead to the identification of at least the same number of *BRCA1/2* mutation carriers. Two-thirds of the *BRCA1/2* mutation carriers identified in this research setting had been referred for counseling and testing. Our results indicate that, awaiting a possibly more extended mutation screening of all breast cancer patients, the triple-negative status of a breast cancer should be added to the CGC referral criteria.

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## INTRODUCTION

The risk of developing breast cancer in *BRCA1* and *BRCA2* mutation carriers is high, and is estimated to be between 27 and 80% up to 70 years of age.<sup>1–5</sup> *BRCA1/2* mutation carriers are more likely to develop breast cancer at a young age and have a higher risk of developing a second breast cancer in comparison with women without a mutation in the *BRCA1* or *BRCA2* gene.<sup>2,6</sup> Additionally, they also have a high risk of developing ovarian cancer, which is estimated to be between 5 and 60% up to 70 years of age.<sup>1–5</sup> Because of the high risks, it is important to identify *BRCA1/2* mutation carriers and offer them options to manage their risks; that is, more intensive screening or risk-reducing surgery (prophylactic mastectomy and/or salpingo-oophorectomy).<sup>7–10</sup> Additionally, it is important to identify breast cancer patients with a *BRCA1/2* mutation because of suggestions that targeted chemotherapy will be available for this patient group in the near future; for example, *BRCA1/2* mutation carriers have been shown to respond to treatment with PARP inhibitors.<sup>10–12</sup>

The prevalence of women carrying a *BRCA1/2* mutation in the general population is low,<sup>13</sup> and within breast cancer patients it was estimated to be around 1–2% for each gene;<sup>14</sup> testing every breast cancer patient for a mutation is debatable. Except for cost-effectiveness and feasibility reasons, there are also considerations

regarding the impact of the counseling and testing procedure.<sup>15</sup> In many Western countries, with no highly prevalent *BRCA1/2* founder mutations, the guidelines state that only patients should be tested with an *a priori* change of at least 10% of having a mutation.<sup>16,17</sup> The Clinical Genetic Center (CGC) referral criteria for breast cancer patients are based on family history of the patient and age at diagnosis of breast cancer.<sup>16</sup> Tumor type, that is, the receptor status of the tumor (estrogen receptor negative (ER–), progesterone receptor negative (PR–) and HER2/neu receptor negative (HER2–)), is suggested to improve the identification of *BRCA1* mutation carriers,<sup>18,19</sup> but is not yet included in most of the CGC referral criteria.<sup>16,17,20</sup> When a patient is referred to the CGC, DNA testing will be offered when the probability of being a carrier of a *BRCA1* and/or *BRCA2* mutation is estimated to be over 10% by the clinical geneticist.<sup>21–25</sup>

In this study, we were able to evaluate the performance of the current Dutch CGC referral criteria<sup>17</sup> for *BRCA1/2* mutation screening, used by clinical geneticists and oncologists in current practice, in an unselected breast cancer patient population diagnosed under the age of 50 years. Changes in sensitivity and specificity were explored by adding the tumor subtype to the criteria and shifting the age at diagnosis thresholds. Furthermore, we studied which part of the *BRCA1/2* mutation carriers identified in the research setting had actually been referred to the CGC.

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## METHODS

### Patient selection

Our study population, a retrospectively ascertained cohort, consists of a consecutive series of invasive early breast cancer patients unselected for a family history of cancer and diagnosed under the age of 50 years. Patients included in the study were treated for primary breast cancer between 1970 and 2003 in the Netherlands Cancer Institute, a cancer hospital in Amsterdam, the Netherlands. Patients with metastases at diagnosis or with earlier tumors were excluded. The breast cancer patients were identified through the Medical Registry of the hospital ( $N = 1893$ ). An overview of the inclusion of patients in the analyses is shown in Figure 1.

### Germline DNA collection and *BRCA1/2* testing (research setting)

For 1620 breast cancer patients (86%), we were able to gather germline DNA of sufficient quality and successfully perform *BRCA1* and *BRCA2* mutation analysis. For most of the patients formalin-fixed, paraffin-embedded tissue blocks containing normal tissue were used for DNA isolation (77%); for 23% of the patients, blood was collected instead. The methods for DNA isolation and mutation analysis have been described elsewhere.<sup>26</sup> In short, DNA was isolated using standard laboratory procedures. *BRCA1/2* mutation analysis included testing for 92 pathogenic variants using allelic discrimination or fragment length analyses; sequencing was used for confirmation of mutations. We estimated that with these methods we were able to capture about 61% of the *BRCA1/2* mutations prevalent in families in the Netherlands.<sup>26</sup> The *BRCA1/2* mutations have been submitted to <http://databases.lovd.nl/shared/genes/BRCA1> and <http://databases.lovd.nl/shared/genes/BRCA2>.

### Family history and clinicopathological data collection

For 903 breast cancer patients included in the cohort, both family history at the time of the breast cancer diagnosis and receptor expression data of the tumor were available. First- and second-degree family history data of the patients at the time of their breast cancer diagnosis, routinely recorded by the treating physician in the patient anamnesis, were collected through the Medical Registry of the hospital. Data regarding the immunohistochemical expression of the ER, PR and HER2 of the breast tumor were gathered from original pathology reports (72% ER/PR and 53% of HER2 status) or determined using

tissue microarray (28% ER/PR and 47% HER2; a tumor was scored negative following the clinically used cutoff points in the Netherlands, i.e., ER <10%, PR <10% and HER2: 0 or 1+).<sup>27,28</sup> Patients with a tumor without the expression of ER, PR and HER2 were considered to have a triple-negative tumor. The 903 patients with complete available data (56% of the total cohort) had a similar median age at diagnosis compared with the rest of the cohort (44 vs 43,  $P$ -value = 0.07), although the median year of diagnosis was somewhat higher (1994 vs 1991 (range for both: 1974–2002),  $P$ -value  $\leq 0.001$ ).

### Definition of the CGC referral criteria

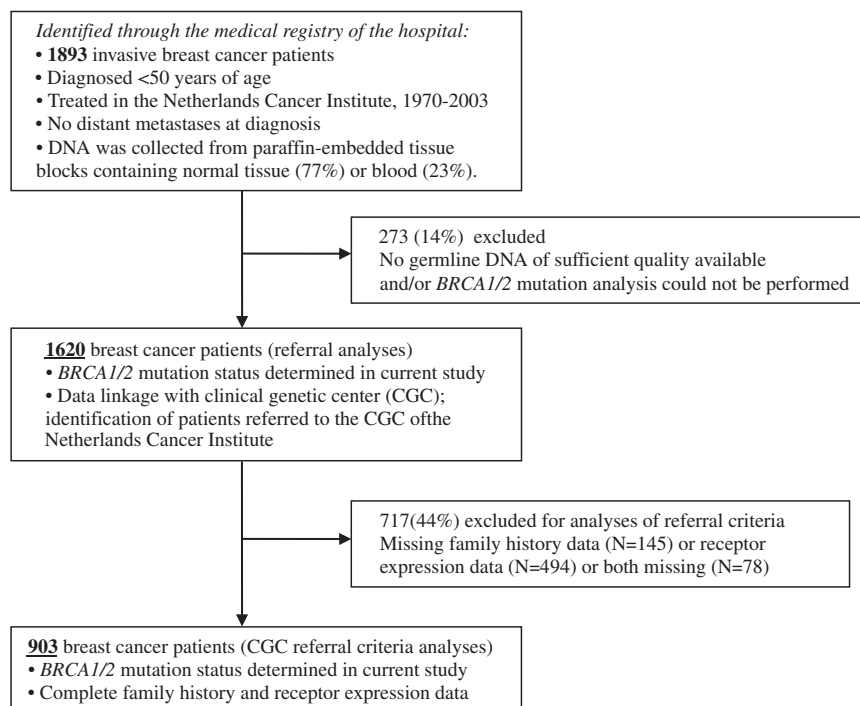
The CGC referral criteria used in this study were based on the Dutch situation.<sup>17</sup> The first criterion is breast cancer diagnosed under the age of 40 years. The second criterion is synchronous bilateral breast cancer (a primary tumor in both breasts) or multiple breast cancer (more primary tumors in one breast); we considered patients with two primary breast tumors diagnosed up to within 3 months apart as synchronous bilateral. The third criterion is when a patient has three or more relatives with cancer, of whom at least one had breast cancer. The family history data available in our hospital medical registry did not completely follow the Dutch CGC referral criteria;<sup>17</sup> therefore, we slightly adapted the family history criteria for the use in our study (Supplementary Table 1).

### Referral of patients to the CGC

In the 'Progeny' registry database of the CGC of the Netherlands Cancer Institute, information about all patients referred to the CGC is recorded. Through linkage with this database, we identified breast cancer patients from our cohort who were referred to this CGC until April 2012. For these patients, data regarding the date of referral, whether a patient was tested for *BRCA1/2* mutations in the CGC and the testing results, were collected from Progeny. Also, full pedigree family history data, recorded by the clinical geneticist for all referred patients and updated at every visit to the CGC by the patient and/or any relatives, were available.

### Coded data linkage

Using a coding procedure,<sup>29</sup> the clinical data and *BRCA1/2* mutation study results were anonymized before linkage. As discussed earlier,<sup>29</sup> no explicit



**Figure 1** Inclusion of patients in the study and in the analyses.

informed consent was asked and results were not reported back to the patients. The secondary use of long-term stored tissue samples and clinical data in this study was according to Dutch guidelines (Dutch codes of conduct: <http://www.federa.org/codes-conduct>) and approved by the review board of the Netherlands Cancer Institute.

### Statistical analyses

Differences in proportions of breast cancer patients with a family history and other clinicopathological characteristics were tested for an association with the *BRCA1* and *BRCA2* mutation status, as determined in our research cohort, using the Pearson's  $\chi^2$ -test.

*Analyses of the performance of the CGC referral criteria to identify women who proved to be BRCA1/2 mutation carriers in our research cohort.* The sensitivity, specificity, positive-predictive and negative-predictive values were calculated for the CGC referral criteria to identify the *BRCA1/2* mutation carriers in our research cohort (for the used methods see Tables 2 and 3). The discriminative performance between *BRCA1/2* mutation carriers and non-carriers was measured by the area under the receiver-operating characteristic curve (AUC) of the logistic regression model including the CGC referral criteria. Changes in sensitivity and specificity were explored by shifting the threshold of the age at diagnosis of breast cancer and by adding and shifting the threshold of the age at diagnosis of triple-negative breast cancer; the criteria regarding family history and bilateral/multiple breast cancer were kept unchanged. The AUC of the models including criteria with shifted thresholds were compared with the AUC of the original model including the CGC referral criteria using a  $\chi^2$ -test. A sensitivity analysis was performed by making the family history criterion less specific: all breast cancer cases with at least one family member with breast cancer at the time of diagnosis were taken into account instead of only cases with more than two relatives with breast and/or other cancer (at least one must have had breast cancer). Statistical analyses were performed using STATA (STATA version 11.2; STATA Corporation, College Station, TX, USA).

*Analyses of the CGC referral patterns of women who proved to be BRCA1/2 mutation carriers in our research cohort.* Proportions of patients who were referred and were not referred, related to the *BRCA1/2* mutation status as determined in our research cohort, were determined and patient characteristics (age at diagnosis of the tumor; family history) were summarized for the different groups. All the above analyses were stratified for calendar year of diagnosis of breast cancer: <1995 compared with >1994; the year of the discovery of the *BRCA1/2* genes and the start of testing in the Netherlands. Furthermore, for patients diagnosed after 1994, time of referral was used to distinguish patients who were referred to the CGC of the Netherlands Cancer Institute before or within 1 year after their breast cancer diagnosis, and those who were referred more than 1 year after their breast cancer diagnosis or were not referred (including 17 (13%) patients with an unknown referral date).

## RESULTS

Forty-nine (5.4%) women were proven to be *BRCA1/2* mutation carriers in our research cohort of 903 breast cancer patients diagnosed under the age of 50 years. Characteristics of the *BRCA1* mutation carriers, *BRCA2* mutation carriers and non-carriers in our cohort are shown in Table 1; 80.6% and 76.7% of *BRCA1* and *BRCA2* mutation carriers, respectively, had one or more family members with breast cancer at the time of breast cancer diagnosis compared with 46.0% of the non-carriers. In all, 58.3% of the *BRCA1* mutation carriers had a triple-negative tumor compared with only 13.2% and 7.7% of the non-carriers and *BRCA2* mutation carriers, respectively.

*Performance of the CGC referral criteria to identify women who proved to be BRCA1/2 mutation carriers in our research cohort.* Each CGC referral criterion, except the bilateral/multiple breast cancer criterion that only included small numbers, identified more than 40% of the

women who proved to be *BRCA1* or *BRCA2* mutation carriers in our research cohort (Table 1).

In all, 77.8% of the breast cancer patients with a *BRCA1* and 69.2% of the breast cancer patients with a *BRCA2* mutation fulfilled at least one of the CGC referral criteria, compared with 36.8% of the non-carriers. No patient in our cohort met all the criteria (Table 1).

The sensitivity and specificity of the CGC referral criteria to identify the *BRCA1/2* mutation carriers in our research cohort were 75.5% and 63.2%, respectively (Table 2). The corresponding AUC was 0.73 (Supplementary Figure 1). The sensitivity of each individual criterion was poor, whereas specificity was high (Table 2). The sensitivity and specificity of the CGC referral criteria to identify only the *BRCA1* mutation carriers were 77.8% and 62.7%, respectively (Supplementary Table 2).

In three different settings in which the triple-negative criterion was added to the CGC referral criteria and the age thresholds in the criteria shifted, the absolute increase of the specificity was up to 6.8%, whereas the sensitivity was equal or also increased (2.1%), compared with the original CGC referral criteria (Table 3). Accordingly, there was an increase in the AUC of 0.04 and 0.05 when the age at diagnosis threshold of all breast cancer cases was decreased to 35 years and the age at diagnosis threshold of the triple-negative breast cancer cases was added and set at 50 years (Supplementary Figure 1A;  $P$ -value = 0.21) or 45 years (Supplementary Figure 1B;  $P$ -value = 0.08), respectively. In the last setting, with the highest increase in discriminative performance compared with the original CGC referral criteria, only three extra mutation carriers were identified in addition to those identified by the original CGC referral criteria (Supplementary Figure 2). However, fewer non-carriers met the shifted criteria than the original CGC referral criteria ( $N$  difference = 68; Supplementary Figure 2). In other words, less patients will have to be tested to identify the same number of *BRCA1/2* mutation carriers, explaining the improved specificity and overall discriminative performance.

Referring to the Dutch NABON guidelines, under patients who are considered for referral, a 10% level of detection rate of *BRCA1/2* mutation carriers is desired.<sup>17</sup> In our research cohort, the percentage of *BRCA1/2* mutation carriers under patients who fulfilled the family history criterion and/or the bilateral/multiple breast cancer criterion was 13.7% (Supplementary Table 3A). Among patients who did not fulfill these criteria, only in patients diagnosed with breast cancer under the age of 30 years, the desired level of 10% detection rate of mutation carriers was reached. The proportion of *BRCA1/2* mutation carriers among patients with a triple-negative tumor was higher than 10% in all age groups except in the 45–50 years age groups (Supplementary Table 3A).

In the sensitivity analyses in which the family history criterion was made less stringent, that is, all women with at least one relative with breast cancer were considered to fulfill the criterion, overall the results did not change and confirmed the original results, although the effects found were less pronounced (Supplementary Tables 3B and 4).

*The CGC referral patterns of women who proved to be BRCA1/2 mutation carriers in our research cohort (N = 1620).* Of the breast cancer patients included in our cohort and diagnosed before 1995, that is, the discovery of the *BRCA1/2* genes (<1995), 14% had been referred to the CGC of the Netherlands Cancer Institute by April 2012 (Figure 2a). Women who were proven to be *BRCA1/2* mutation

**Table 1** Distribution of clinicopathological characteristics and the CGC referral criteria<sup>a,b</sup>

<i>BRCA1/2</i> mutation status <sup>c</sup>	Non-carriers		<i>BRCA1</i>		<i>P</i> -value <sup>d</sup>	<i>BRCA2</i>		<i>P</i> -value <sup>d</sup>
	<i>N</i> <sup>e</sup>	% <sup>e</sup>	<i>N</i> <sup>e</sup>	% <sup>e</sup>		<i>N</i> <sup>e</sup>	% <sup>e</sup>	
Total <i>N</i> =903	854	94.6	36	4.0		13	1.4	
Median age diagnosis (years, SD)	44	5.4	36	7.6	<0.001	39	4.5	0.002
Bilateral/multiple breast cancer	6	0.7	2	5.6	0.003	0	0	NA
Triple-negative tumor	113	13.2	21	58.3	<0.001	1	7.7	0.56
Family history (type <sup>f</sup> )								
No affected relatives	310	36.3	1	2.8	<0.001	1	7.7	0.06
Yes, breast cancer	393	46.0	29	80.6		10	76.9	
Yes, only other cancers	151	17.7	6	16.7		2	15.4	
Family history (number <sup>g</sup> )								
No affected relatives	310	36.3	1	2.8	<0.001	1	7.7	0.006
Yes, <3	373	43.7	18	50.0		5	38.5	
Yes, ≥3	171	20.0	17	47.2		7	53.9	
<i>CGC referral criteria<sup>a</sup></i>								
(A) Three or more family members with cancer <sup>h</sup>	146	17.1	16	44.4	<0.001	6	46.2	0.006
(B) Bilateral/multiple breast cancer	6	0.7	2	5.6	0.003	0	0	NA
(C) Breast cancer diagnosed <40 years	209	24.5	21	58.3	<0.001	7	53.8	0.02
At least one criterion met	314	36.8	28	77.8	<0.001	9	69.2	0.02
Number of <i>BRCA1/2</i> screening criteria met								
1	267	85.0	17	60.7	0.001	5	55.6	0.02
2	47	15.0	11	39.3		4	44.4	
3	0	0	0	0		0	0	
<i>Proportions BRCA1/2 mutation carriers under all patients, by age diagnosis (years)</i>								
<35	75	81.5	15	16.3	<0.001	2	2.2	0.02
35–40	134	92.4	6	4.1		5	3.5	
40–45	242	94.5	9	3.5		5	2.0	
45–50	403	98.3	6	1.5		1	0.2	
<i>Proportions BRCA1/2 mutation carriers under patients with triple-negative tumors, by age diagnosis (years)</i>								
<35	19	61.3	12	38.7	<0.001	0	0	0.12
35–40	16	80.0	3	15.0		1	5.0	
40–45	25	86.2	4	13.8		0	0	
45–50	53	96.4	2	3.6		0	0	

Abbreviations: CGC, clinical genetic centre; NA, not applicable.

<sup>a</sup>Adjusted criteria as described in Supplementary Table 1.

<sup>b</sup>Analyses among included breast cancer patients stratified by the *BRCA1/2* mutation status as determined in our research cohort.

<sup>c</sup>As determined in this research setting.

<sup>d</sup>*P*-value of the Pearson's  $\chi^2$ -test, *BRCA1* or *BRCA2* carriers compared with non-carriers.

<sup>e</sup>Unless otherwise specified.

<sup>f</sup>Type of cancer present in affected relatives.

<sup>g</sup>Number of affected relatives in the family.

<sup>h</sup>At least one must have had breast cancer.

carriers in our research cohort were more frequently referred compared with the non-carriers in our cohort (34% vs 13%, *P*-value <0.001). The referral time ranged from 4 months to more than 29 years after the breast cancer diagnosis (range referral years: 1995–2011).

Fifteen percent of all breast cancer patients included in our cohort and diagnosed after 1994 had been referred to the CGC of the Netherlands Cancer Institute before (2%) or within 1 year after diagnosis (14%) (Figure 2b). Patients who were referred before diagnosis (*N*=12) all had a family history of breast cancer at the time of diagnosis; most of them (*N*=11) even had three or more relatives affected with cancer. Furthermore, patients referred not before but within 1 year after diagnosis were more likely to have a family history of breast cancer at the time of diagnosis (86% vs 44%, *P*-value <0.001), to have more affected relatives with cancer (35% vs 20%, *P*-value = 0.001) and to be diagnosed with breast cancer under the age of 35 years (32% vs 9%, *P*-value <0.001) than patients who

were *not* referred to the CGC of the Netherlands Cancer Institute before or within a year after breast cancer diagnosis. However, of these patients who were not referred, 17% still had an extensive family history at the time of diagnosis (three or more relatives with cancer, of whom at least one had breast cancer) (Figure 2b).

Under the patients diagnosed with breast cancer after 1994, the proportion who had been referred to the CGC of the Netherlands Cancer Institute before diagnosis was higher among *BRCA1/2* mutation carriers than among the non-carriers identified in our research cohort (15% vs 1%, *P*-value <0.001). Also, the proportion of patients who were referred not before but within 1 year after diagnosis was higher among the carriers (26% vs 13%, *P*-value <0.001) (Figure 2b). These referred carriers all had a family history of breast cancer, of whom eight (50%) had more than three affected relatives with cancer, compared with 74% of the *BRCA1/2* mutation carriers who were not referred before or within 1 year after diagnosis (*P*-

**Table 2 Performance of each individual criterion and the CGC referral criteria<sup>a,b</sup>**

	No. false negatives		No. true positives		Total no. that met the criterion		% of <i>BRCA1/2</i> mutation carriers, positive-predictive value <sup>c</sup>		Sensitivity	Specificity
	Yes	No	No	Yes	Yes	No	value <sup>c</sup>	Negative-predictive value		
(A) Three or more family members with cancer <sup>d</sup>	27	708	146	22	168	13.1	96.3	44.9	82.9	
(B) Bilateral/multiple breast cancer	47	848	6	2	8	25.0	94.7	4.1	99.3	
(C) Breast cancer diagnosed <40 years	21	645	209	28	237	11.8	96.8	57.1	75.5	
<b>CGC referral criteria<sup>a</sup> – all above criteria combined</b>	<b>12</b>	<b>540</b>	<b>314</b>	<b>37</b>	<b>351</b>	<b>10.5</b>	<b>97.8</b>	<b>75.5</b>	<b>63.2</b>	
Triple-negative breast cancer (no age limitation)	27	741	113	22	135	16.3	96.5	44.9	86.8	

Abbreviations: negative-predictive value, no. of non-carriers that did not meet the CGC referral criteria/total no. that did not meet the CGC referral criteria; positive-predictive value, no. of *BRCA* mutation carriers that met the CGC referral criteria/total no. that met the CGC referral criteria; sensitivity, no. of *BRCA1/2* mutation carriers that met the CGC referral criteria/total no. of *BRCA1/2* mutation carriers; specificity, no. of non-carriers that did not meet the CGC referral criteria/total no. of non-carriers.

<sup>a</sup>Adjusted criteria (based on the Dutch situation) as described in Supplementary Table 1; bold represents the CGC referral criteria<sup>a</sup>.

<sup>b</sup>Analyses to identify women who proved to be *BRCA1/2* mutation carriers in our research cohort.

<sup>c</sup>Diagnostic yield: prior chance on detecting a mutation.

<sup>d</sup>At least one must have had breast cancer.

**Table 3 Performance of adaptations of the CGC referral criteria<sup>a,b</sup>**

Criteria shift		No. of false negatives		No. of true positives		Total no. that met the criterion	% of <i>BRCA1/2</i> mutation carriers, positive-predictive value <sup>c</sup>		Sensitivity	Specificity	AUC	P-value
Age diagnosis of all breast cancer	Age diagnosis of TN breast cancer	No. of false negatives	No. of true positives	No. of false positives	No. of true positives		value <sup>c</sup>	Negative-predictive value				
<25	x	25	701	153	24	177	13.6	96.6	49.0	82.1	0.66	NA
<25	<30	21	697	157	28	185	15.1	97.1	57.1	81.6	0.71	NA
<25	<35	19	687	167	30	197	15.2	97.3	61.2	80.4	0.73	NA
<25	<40	17	672	182	32	214	15.0	97.5	65.3	78.7	0.75	NA
<25	<45	14	648	206	35	241	14.5	97.9	71.4	75.9	0.77	NA
<25	<50	13	607	247	36	283	12.7	97.9	73.5	71.1	0.75	NA
<30	x	20	688	166	29	195	14.9	97.2	59.2	80.6	0.71	NA
<30	<35	18	678	176	36	212	17.0	97.4	66.7	79.4	0.73	NA
<30	<40	16	663	191	33	224	14.7	97.6	67.3	77.6	0.75	NA
<30	<45	13	639	215	36	251	14.3	98.0	73.5	74.8	0.78	NA
<30	<50	12	598	256	37	293	12.6	98.0	75.5	70.0	0.76	0.31
<35	x	17	647	207	32	239	13.4	97.4	65.3	75.8	0.72	NA
<35	<40	15	632	222	34	256	13.3	97.7	69.4	74.0	0.76	NA
<b>&lt;35</b>	<b>&lt;45</b>	<b>12</b>	<b>608</b>	<b>246</b>	<b>37</b>	<b>283</b>	<b>13.1</b>	<b>98.1</b>	<b>75.5</b>	<b>71.2</b>	<b>0.78</b>	<b>0.08</b>
<35	<50	11	567	287	38	325	11.7	98.1	77.6	66.4	0.77	0.21
<b>&lt;40</b>	<b>NA</b>	<b>12</b>	<b>540</b>	<b>314</b>	<b>37</b>	<b>351</b>	<b>10.5</b>	<b>97.8</b>	<b>75.5</b>	<b>63.2</b>	<b>0.73</b>	<b>Ref.</b>
<40	<45	9	516	338	40	378	10.6	98.3	81.6	60.4	0.79	NA
<40	<50	8	475	379	41	420	9.8	98.3	83.7	55.6	0.78	NA
<45	x	3	328	526	46	572	8.0	99.1	93.9	38.4	0.75	NA
<45	<50	2	287	567	47	614	7.7	99.3	95.9	33.6	0.80	NA
<50	x	0	0	854	49	903	5.4	0	100	0	0.50	NA

Abbreviations: NA, not applicable; negative-predictive value, no. of non-carriers that did not meet the CGC referral criteria/total no. that did not meet the CGC referral criteria; positive-predictive value, no. of *BRCA* mutation carriers that met the CGC referral criteria/total no. that met the CGC referral criteria; sensitivity, no. of *BRCA* mutation carriers that met the CGC referral criteria/total no. of *BRCA* mutation carriers; specificity, no. of non-carriers that did not meet the CGC referral criteria/total no. of non-carriers; x, no change in the criterion.

<sup>a</sup>Adjusted criteria (based on the Dutch situation) as described in Supplementary Table 1; bold represents the CGC referral criteria<sup>a</sup>; italics represents the adjusted criteria achieving an equal or higher sensitivity and higher specificity than the CGC referral criteria<sup>a</sup>; bold and italics represent the adjusted criteria achieving an equal or higher sensitivity and higher specificity than the CGC referral criteria, and with the best discriminatory performance (highest AUC).

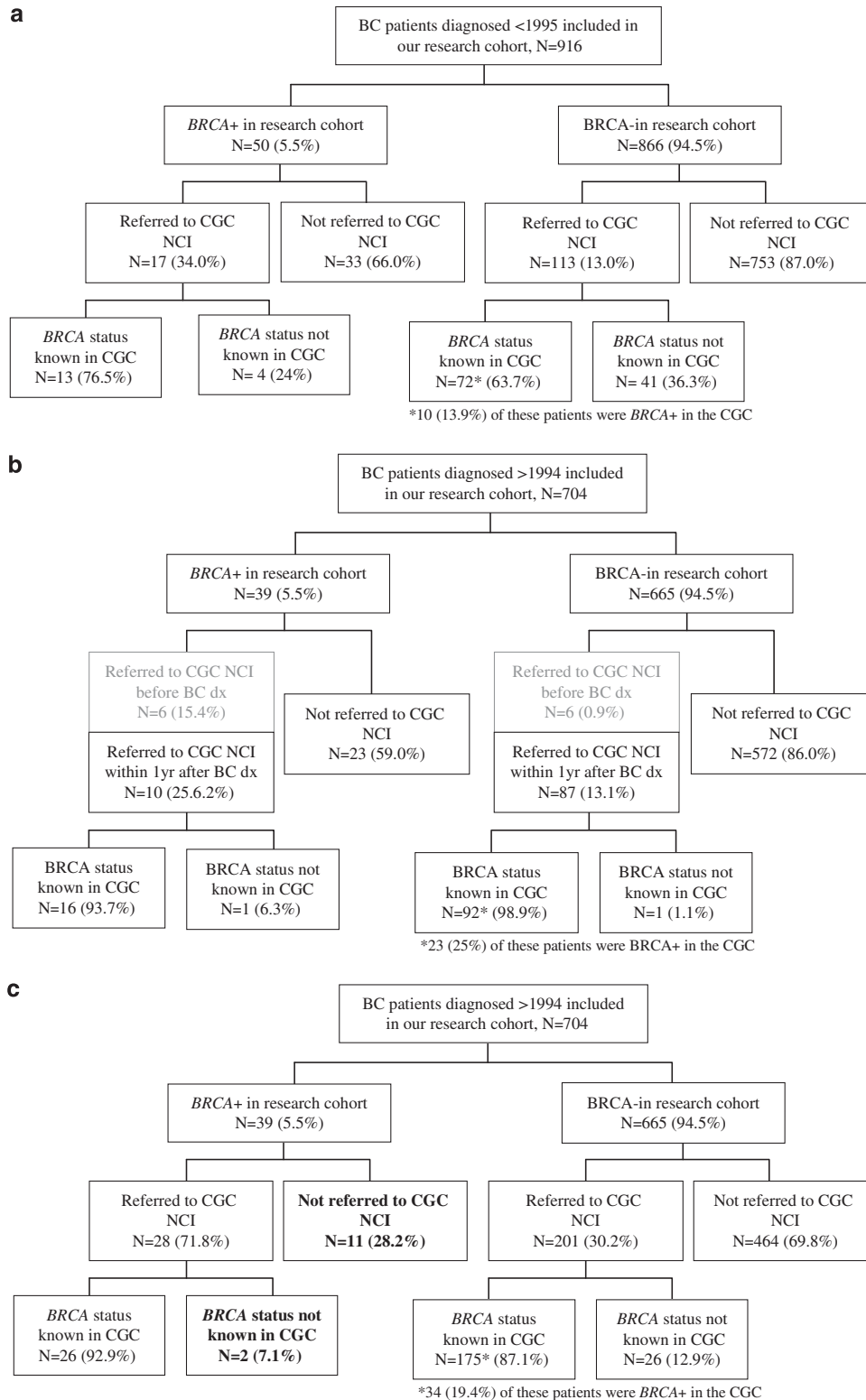
<sup>b</sup>Analyses to identify women who proved to be *BRCA1/2* mutation carriers in our research cohort when shifting the age at diagnosis of breast cancer and shifting the age at diagnosis of triple-negative breast cancer. No changes in the other criteria were made.

<sup>c</sup>Diagnostic yield: Prior chance on detecting a mutation.

value = 0.03; Figure 2a). Furthermore, nine (56%) of the referred *BRCA1/2* mutation carriers were diagnosed with breast cancer at a very young age (under 35 years) compared with six (26%) of the carriers who were not referred before or within 1 year after their diagnosis ( $P$ -value = 0.06).

Ten additional *BRCA1/2* mutation carriers diagnosed after 1994 were referred to the CGC in the later years after diagnosis. Eventually, 67% ( $N = 26$ , taking into account 2 with an unknown referral date) of the total group of women who proved to be *BRCA1/2* carriers in our research cohort were referred and were





**Figure 2** Overview of the women who proved to be *BRCA1/2* mutation carriers and non-carriers in our research cohort and referral of these patients to the CGC of the Netherlands Cancer Institute (NCI) ( $N = 1620$ ), stratified for the calendar year of diagnosis (**a–c**). Bold represents the number of patients who are carriers of a *BRCA1/2* mutation and were diagnosed after 1994, but who were not referred to the CGC of the NCI or the *BRCA* status was not known in the CGC of the NCI; BRCA–, *BRCA* mutation non-carrier as determined in our research cohort; BRCA+, *BRCA1/2* mutation carrier as determined in our research cohort. For 166 (18%) patients in panel a family history data was missing; for 57 (8%) patients in panels b and c family history data was missing. (a) Patients diagnosed <1995: all patients referred to the CGC by April 2012 are indicated as referred. (b) Patients diagnosed >1994: only the patients referred to the CGC before breast cancer diagnosis (gray), or within 1 year after their breast cancer diagnosis are indicated as referred (patients with unknown referral date ( $n = 17$ ; of which two BRCA+) are included in the group ‘not referred’). (c) Patients diagnosed >1994: all patients referred to the CGC by April 2012 are indicated as referred.

tested for *BRCA1/2* mutations in the CGC of the Netherlands Cancer Institute (Figure 2c). Thirteen *BRCA1/2* mutation carriers had not been referred or were not tested for *BRCA1/2* mutations by April 2012 (Figure 2c). Four of these carriers were diagnosed in 1995 or 1996, shortly after the discovery of the *BRCA1/2* genes; two died shortly (1 and 4 years) after the breast cancer diagnosis. Of the seven other carriers, only two had more than two family members with breast cancer at the time of diagnosis (Supplementary Table 5).

## DISCUSSION

Our study is one of the first studies to evaluate the performance of the CGC referral criteria to identify women who proved to be *BRCA1/2* mutation carriers in an *unselected* hospital-based cohort of breast cancer patients. Most earlier publications evaluated risk assessment models, which are based on extensive pedigree information and are most suitable *after* a patient has been referred to the CGC.<sup>30</sup> We calculated that the CGC referral criteria (based on the Dutch situation) had a sensitivity of 75.5% and specificity of 63.2% to identify the *BRCA1/2* mutation carriers. In the group of patients in our cohort who met the CGC referral criteria ( $N = 351$ ), there was a detection rate for *BRCA1/2* mutation carriers of 10.5%. An improved performance to identify *BRCA1/2* mutation carriers, that is, a better specificity (up to 6.8% absolute increase) and similar or better sensitivity (up to 2.1% increase), and accordingly an increase in detection rate up to 13.1%, was achieved by shifting the threshold of the age at diagnosis of breast cancer from 40 to 35 years and by adding an age threshold to 45 or 50 year for patients with a triple-negative breast cancer. In conclusion, to identify the same number or even more *BRCA1/2* mutation carriers, fewer patients need to be referred to the CGC.

It should be noted here that our results are most important for referral of *BRCA1* mutation carriers: they are more often diagnosed with triple-negative tumors compared with *BRCA2* mutation carriers (Table 1).<sup>31</sup> When performing our analyses on *BRCA1* mutation carriers only, we saw stronger and significant effects of the addition of triple-negative status to the CGC referral criteria on the sensitivity and specificity (Supplementary Table 2). We considered there was insufficient power to perform the analyses separately for *BRCA2* mutation carriers. Moreover, in clinical practice mutation testing is mostly performed for both genes simultaneously, and thus separation of the results is less relevant.

When interpreting the results from this study, it should be taken into account that our study is performed in a research setting; we estimated that we were able to screen for about 61% of the pathogenic *BRCA1/2* mutations (69% *BRCA1* mutations; 46% *BRCA2* mutations) known to be prevalent in Dutch families.<sup>26</sup> We screened for a higher percentage of *BRCA1* mutations than *BRCA2* mutations, and *BRCA2* mutation carriers are less prevalent in the Dutch population.<sup>26</sup> Consequently, we missed *BRCA1/2* mutation carriers in our cohort (see Figure 2) and we may assume results reported here are conservative and most important for the referral of *BRCA1* mutation carriers. In addition, it should be kept in mind that the difference in absolute numbers of *BRCA1/2* mutation carriers comparing the original and altered CGC referral criteria was small.

The earlier CGC referral criteria were solely based on the patients' family history of cancer.<sup>32</sup> In line with this, we noticed that *BRCA1/2* mutation carriers identified in our research cohort, who were diagnosed with breast cancer after 1994 and referred before or within 1 year after diagnosis, more often had a family history of breast cancer at the time of diagnosis compared with those who were referred later or not at all. Likely, the later referred patient developed

another cancer or had additional family members developing cancer. Thirteen (33%) *BRCA1/2* mutation carriers identified in our research cohort and diagnosed with breast cancer after 1994 were not referred to the CGC of the Netherlands Cancer Institute by April 2012. Most of them were not referred for logical reasons (these patients were diagnosed shortly after the discovery of the *BRCA1/2* genes or died shortly after the breast cancer diagnosis; Supplementary Table 5); in addition, a few, even though with a positive family history of breast cancer at the time of diagnosis, might not have wanted to be referred,<sup>33</sup> their treating physician lacked knowledge and/or awareness,<sup>34</sup> or even though diagnosed and treated at The Netherlands Cancer Institute, a specialized cancer center, were screened in another CGC.

The family history of a patient is still the most important criterion for referral of patients to the CGC<sup>17</sup> with the highest sensitivity to identify *BRCA1/2* mutation carriers (Tables 1 and 2).<sup>35</sup> Of note, non-carriers with an extensive family history of (breast) cancer are at increased risk of developing second cancers<sup>36</sup> and referral to the CGC of these patients remains important. As we did not have full pedigree information at the time of diagnosis available in our cohort, we probably missed information of some brothers/fathers affected by breast cancer or family members affected by ovarian cancer; however, we had information also about breast cancer patients non-referred to the CGC. It is difficult to predict what would be the effect of this on the results observed in this study, but in previously published Dutch data, the prevalence of breast cancer patients with ovarian cancer in the family was only 2%.<sup>37</sup> That same study reported 34% of breast cancer patients to have a family history of breast cancer to the extent of a CGC referral indication;<sup>37</sup> higher than the 19% in our study, this difference likely being explained by our younger population. Considering these figures, we can postulate that in our cohort the effect of the family history criterion might be somewhat underestimated. However, in a sensitivity analysis with a less specific family history criterion (48%; taking all breast cancer cases with at least one relative with breast cancer into consideration), we confirmed that decreasing the age at diagnosis threshold of all breast cancer cases and increasing the age at diagnosis threshold of the triple-negative breast cancer cases improves both the sensitivity and specificity of the criteria (Supplementary Table 4).

We also tried to get insight into the quality of the family history data we used in this study, by comparing family history information from the clinical report with the more extended family history data available from the subgroup of CGC referred patients. Not surprisingly, the former data reported fewer patients with a family history of cancer, although this difference was most pronounced for non-breast cancer (data not shown). The CGC data reflected for many patients the status *after* their diagnosis of breast cancer, or later since pedigrees are updated at every visit to the CGC by the patient and/or any relatives. Since for breast cancer patients it is mostly the treating physician who refers to the CGC, we feel that the data used in our analyses are more representative for the everyday situation in clinical practice. Although this family history information may not have been recorded consistently over the whole study period by clinicians, carrier status was unknown at diagnosis, and therefore it is unlikely that this led to differential bias.

Our results, certainly when taking into account previously published data,<sup>16,18,19,38–40</sup> indicate that histologic parameters, that is, the triple-negative status of the breast cancer, should be incorporated in the CGC referral criteria. Even though we evaluated specifically the Dutch guidelines, other Western countries are in a similar situation; of seven international guidelines, only two indicate triple-negative breast cancer as an indication for *BRCA1/2* testing.<sup>20</sup> Families in Western countries are becoming smaller and other factors to identify

*BRCA1/2* mutation carriers, such as the type of breast tumor, are becoming more important.<sup>41</sup> It has repeatedly been suggested that pathologic data should be incorporated in addition to family history in the prediction of the probability of carrying a mutation in one of the *BRCA1/2* genes.<sup>18,38</sup> Recent publications show that even testing all women younger than 50 years with triple-negative breast cancer, regardless of family history, is a cost-effective strategy and could reduce subsequent cancer risks,<sup>39,40</sup> and the NICE guidelines (UK) recently included the triple-negative criterion (<age 40 years).<sup>42</sup> Expanding the CGC referral criteria with this information will improve cancer risk management, for example, decisions of salpingo-oophorectomy or preventive contralateral surgery and/or treatment with PARP inhibitors, for these women and their families.

### CONFLICT OF INTEREST

The authors declare no conflict of interest.

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