BRCA1 (185deIAG) Mutation among Egyptian Breast Cancer

Female Patients

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ABSTRACT: This study was conducted to estimate the frequency of BRCA1 (185delAG) mutation among Egyptian female patients with breast cancer. Forty selected female patients with breast cancer, 80 of their female relatives and 10 healthy females as a control group were included in this study. Result: The age of onset of breast cancer was below 40 years in 25 (62.5%) patients and above 40 years in 15 (37.5%) patients. There were significant differences among the patients regarding the age at menarche before 13 years (p=0.011, p<0.05), onset of breast cancer (p=0.000, p<0.001), parity (p=0.000, p<0.001), first delivery before 30 years of age (p=0.04, p<0.05), breast feeding (p=0.002, p<0.05), and positive family history (p=0.000, p<0.001). The frequency of BRCA1 (185delAG) mutation was found among 10% of the patients' group. Eight percent of patients with early onset below 40 years and 13.5% of patients with onset after 40 years were heterozygotes for the mutation. Three percent of patients with unilateral breast cancer, 40% of patients with bilateral breast cancer and 50% of patients with breast ovarian cancer were carrying the mutation. Our results indicated that breast ovarian cancer and bilateral breast cancer patients were likely to have BRCA1 (185delAG) mutation than in unilateral breast cancer.

INTRODUCTION

Breast cancer (BC) is the most common	BC is futile and disfiguring, making early
malignancy in women, accounting for 31%	detection has a high priority in medical
of all female cancers, and responsible for	management of the disease. ⁽²⁾
15% of cancer deaths in women. ⁽¹⁾ One	The risk factors of BC include genetic,
million females world wide are diagnosed	environmental, and hormonal. Genetic risk
with BC every year. Treatment of advanced	factors contribute to about 5%-10% of all

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cases, 90%-95% of them result from somatic mutation and about 5%-10% are inherited as the result of germ line mutation in autosomal dominant BC susceptibility genes.^(3,4)

Many genes have been found to increase susceptibility to cancer and are also associated with familial breast cancer. These genes include Breast Cancer Antiestrogen resistance-1 (BRCA1), Breast Cancer Anti-estrogen resistance-2 (BRCA2), Ataxia Telangiectasia mutant gene (ATM), Phosphate and Tensin homology (PTEN), and Tumor Protein (P53). Several other less frequently predisposing genes are also involved but to lesser extent.(3,4)

BRCA1 gene (chr 17q21) is a tumor suppressor gene that encodes tumor suppressor protein which acts as a negative regulator for tumor growth. It accounts for 45% of inherited BC and 90% of inherited breast-ovarian cancer (BOC) in highly affected families.⁽⁵⁾ Different types of mutation have been found in the BRAC1 gene and predispose to development of cancer. The most common of them is the frame shift mutation at position 185 in exon2, involving deletion of adenine and guanine (185 del AG mutation).⁽⁶⁻⁸⁾ The identification and study of this mutation could facilitate early diagnosis and proper counseling for BC. Therefore, the present study aimed at identifying the 185 del AG mutation (exon 2) of BRCA1 gene among Egyptian females with early BC and BOC and their female relatives. This will allow diagnosis proper early and aenetic counseling for susceptible cases in the families.

PATIENTS AND METHODS

The present study included 40 female patients with BC collected from the Oncology Unit, Medical Research Institute, Alexandria University. Two close female relatives for each patient (80 females) and ten healthy age matched females without any family history of any cancer, as a control group, were also included.

Female patients participating in the study were selected according to (The following criteria):

- Early onset breast cancer (below 40 years).
- 2- Bilateral breast cancer.
- 3- Breast-ovarian cancer.
- Positive family history of breast cancer.

The patients and control group were subjected to the following:

- 1- Detailed history:
- Personal history including age, age of menarche, marriage, age of first full term pregnancy, lactation, and menopause.
- Family history of breast cancer or any other cancer (for patients and relatives).
- 2- Molecular studies to detect BRCA1 (185 del AG) mutation as follows:
- a- DNA isolation and purification from peripheral blood using standard

method.⁽⁹⁾

Forward:

b- Polymerase chain reaction (PCR) amplification: according to Lahad *et al.*, (1977).⁽¹⁰⁾ The sequence of the primers used were as follows:

GAAGTTGTCATTTTATAAACCTTT-3' Reverse:

TGTCTTTTCTTCCCTAGTATGT-3'

The PCR mix was performed in a final volume of 25 μ L containing 100 ng (3 μ l) DNA, 10 pmole (3 μ l) of each primer, 12.5 μ l of the ready made PCR master mix (Promega Chemical Co.) and 3.5 μ l of nuclease free water was added to adjust the volume to 25 μ l.

- c- PCR conditions as follows: 1 cycle at 95°C for 5 minutes followed by 33 cycles at 94°C for 1 minute, 58°C for 1 minute and 72°C for 1 minute, 1 cycle at 72°C for 10 minutes.
- d- Agarose gel electrophoresis: was
 carried out on a 3% agarose gel and
 visualized with ethidium bromide.⁽¹¹⁾

5'-

5'-

- e- Single Strand Conformational Polymorphism (SSCP) Analysis: (12,13) Ten µl of PCR product was diluted 1:1 in formamide buffer (95% formamide, 20 mM EDTA pH 8.0, 0.05 xylene cyanol, and 0.05% bromophenol blue), kept at 85°C for 5 minutes and then cooled quickly in ice. The samples were loaded onto a non-denaturating polyacrylamide and run at room temperature for 4 hours using constant voltage (150V) and 1 X TBE buffer. Detection was carried out using silver stain and ethidium bromide.
- 3- Statistical analysis: Statistical differences between patients and controls, also among patients were determined with the Fisher's Exact Test.

RESULTS

This study included 40 female patients with BC or BOC, 80 of their close female relatives and 10 healthy adult females as a control group.

Characteristics of cases

The age of menarche was below 13 years in 28 patients (70%) and over 13 years in 12 patients (30%) with a mean age of 13.20±1.3 years .The age of menopause was more than 45 years in 8 patients (20%) with a mean age of 49.10±3.4 years. The remaining 32 patients (80%) did not experience menopause as they developed cancer at young age. The majority of the patients, 37 cases (92.5%), were married. Nullipara constituted 6 patients (15%). Among 34 patients (85%); 23 (67.6%) had their 1st child below 30 years of age while 11 (32.4%) were over 30 years. Breast feeding was practiced in 30 patients (75%).

Table 1 shows the characteristics of the patients and the controls .There was no significant differences between patients and controls in age at menarache (P=0.138, p >0.05), marital status (P=1.00, P>0.05), parity (P=0.653, P>0.05), age at first delivery (P=0.402, P>0.05), and breast feeding (P=0.707, P>0.05). However, there

was a significant difference among patients regarding, age at menarche before 13 years (P=0.011, P<0.05), onset of BC (P=0.000, P<0.05), marital status (P=0.000, P<0.05) parity (P=0.000, P<0.05) first delivery before 30 years of age (P=.040, P<0.05), and breast feeding (P=0.002, P<0.05).

Classification of cases

The female patients were classified into 2 groups according to the age of onset of the disease. The first group included 25 (62.5%) patients with early onset BC before 40 years. Among this group 2 patients (8%) were with bilateral BC, 22 patients (88%) with unilateral BC and one patient (4%) with BOC. The second group included 15 patients (37.5%) with late onset BC above 40 years of age. Three of them (20%) were with bilateral BC, 11 patients (73.3%) with unilateral BC and one patient (6.7%) with BOC. Positive family history with BC was detected in 2 patients (8%) in the first group and in13 patients (86.6%) in the second group. Positive family history was significantly higher in the second group patients (P=0.000, P<0.05). (Table 2)

Molecular study

DNA was amplified using intronic specific primers spanning exon 2 (262bp). Similar amplified PCR products for all samples subjected to the present study were detected (Fig. 1).

The SSCP analysis revealed 4 patients (10%) carring 185 del AG mutation in BRCA1 gene. (Fig. 2 and 3). No mutation was detected in the patient's relatives or in control groups.

The main findings of the four patients with BRCA1 mutation are summarized in (Table 3).

		Control		Patients	
		%	No.	%	No.
- Age at menarche					
< 13 y		40	4	70	28
≥ 13 y	p* = 0.138	60	6	30	12
	p** = 0.011				
- Onset of cancer					
Premenopausal	-	-	-	80	32
Postmenopausal		-	-	20	8
	p** = 0.00				
- Marital status					
Married		90	9	92.5	37
Not married	p* = 1.000	10	1	7.5	3
	p** = 0.000				
- Parity					
Parous women		80	8	85	34
Multiparous women	p* = 0.653	20	2	15	6
	p** = 0.000				
- Age at 1 st delivery			(n = 8)		(n = 34)
< 30 y		87.5	7	67.6	23
\geq 30 y	p* = 0.402	12.5	1	32.4	11
	p** = 0.040				
- Breast feeding					
Yes		70	7	75	30
No	p* = 0.707	30	3	25	10
	p* = 0.002				

Table (1): Characteristics of patients with breast cancer and control

p*: p value between patients and control.

p**: p value among patients.

	Age o	Total	
	> 40 y	□ 40 y	
	No. %	No. %	
	15 (37.5%)	25 (62.5%)	
Positive family history for BC*	13 (86.6%)	2 (8%)	15 (37.5%)
Positive family history for another cancer	1 (6.7%)	-	1 (2.5%)
Unilateral BC	11 (73.3%)	22 (88%)	33 (82.5%)
Bilateral BC	3 (20%)	2 (8%)	5 (12.5%)
Breast ovarian cancer	1 (6.7%)	1 (4%)	2 (5%)

Table (2): Distribution of patients according to age of onset of the disease

* Significant difference in positive family history of BC between the 2 groups of patients (p < 0.001).

Breast ovarian cancer (one case)	Bilateral breast cancer (2 cases)		Unilateral breast cancer (one case)	
47 y	47 y	35 y	32 y	Age of onset
> 13 y	< 13 y	< 13 y	< 13 y	Age of menarche

Table (3): Characteristics of the	patients with BRCA1	(185 del AG) mutation
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47 y	47 y	35 y	32 y	Age of onset
> 13 y	< 13 y	< 13 y	< 13 y	Age of menarche
Yes	Yes	Yes	Yes	Parity
> 30 y	> 30 y	< 30 y	< 30 y	Age at 1 st delivery
Yes	Yes	Yes	Yes	Lactation
+ve	-ve	-ve	+ve	Family history of BC
+ve	-ve	-ve	-ve	Family history of another cancer



Agarose gel (3%) electrophoresis of PCR products for BRCA1 exon2 Gene (lanes 2-11). Lane 1 represents 50 KP ladder mdecular weight marker

Figure (1): Agarose gel (3%) electrophoresis of PCR products for BRCA1 exon2 Gene (lanes 2-8). Lane 1 represents 50bp ladder molecular weight marker.



SSCP analysis of PCR products for BRCA1 exon2 gene Lanes (1.7) show band variation, lane (10.11) control PCR product (negative for the mutation)

Figure (2): Single strand conformational polymorphism (SSCP) analysis of PCR products for BRCA1 exon2 gene (using ethidium bromide stain). Lanes (1,7) show band variation: upper bands represent the single stranded DNA; the lower bands represent the reannealed double strand DNA. Lanes (10,11) control PCR product (negative for the mutation). Lanes (2,3,4,5,6,8,9) represent normal PCR products for patients without the muation.



SSCP analysis of PCR products for BRCA1 exon2 gene Lanes (1.2) show band variation, lane 5 control PCR product (negative for the mutation)

Figure (3): Single strand conformational polymorphism (SSCP) analysis of PCR product for BRCA1 exon2 gene (using silver staining). Lanes (1,2) show band variation: upper bands represent the single stranded DNA and lower bands represent the reannealed double stranded DNA. Lane 5 control PCR product (negative for the mutation). Lanes (3,4) represent PCR products for patients without the mutation.

DISCUSSION

Multiple menstrual and reproductive events; menarche, pregnancy, breast feeding; may alter BC risk. In the present study there was a significant increase in the patients with menarche before 13 years of age (P<0.05) and the mean age of menarche of patients was 13.2 years and menopause at 49 years. These results were similar to those reported by Knudson et al⁽¹⁴⁾ who found that both early age at menarche (before 13 years) and late age at menopause (more than 45 years) were found to be associated with increased risk of BC especially in susceptible women. This may be related to a higher life time exposure to the hormones estrogen and progesterone.

No association between BC and marital status of the patients was detected in the present study as there was no significant difference between patients and controls (P>0.05). This is consistent with other studies that confirmed the absence of significant association between BC and marital status which may indicate that marital status alone without parity has no role in hormone level change in the body.⁽¹⁵⁻¹⁷⁾ However, the married women were significantly higher in the patients group (P<0.05).

Single and nulliparous women were reported to have an increased risk of BC, about 1.4 times the risk of parous women. This low risk of BC in parous women might result from the protective effect of age at first pregnancy and child birth.⁽¹⁶⁻¹⁸⁾ The finding of the present study was contrary to that as parous women were significantly higher in the patients group (P<0.05). These parous women may be exposed to a strong risk factors that lead to the BC inspite of the protective effect of parity.

Late age at first child birth increases the life time incidence of BC.^(19,20) It was reported that being older than 30 years at first delivery is a risk factor for BC.⁽²¹⁾ The highest risk was in those who have a first child after the age of 35 years, they appear to be at even a higher risk than that of nulliparous women.⁽²²⁾ Moreover, epidemiological studies have consistently shown that women who undergo an early first full term pregnancy have a significantly reduced life time risk of BC, this association is independent of parity, i.e., number of live birth.^(19,20) In the present study there was no significant difference in age at first delivery between parous

patients and controls, moreover, the parous patients who gave birth to their first child before 30 years of age were significantly higher (P<0.05) than those who gave birth after 30 years of age .

It has been suggested that breast feeding may protect against BC and increasing years of nursing experience may decrease the BC risk.⁽²³⁾ In the present study no significant difference (p >0.05) between patients and controls was detected but the patients who had experienced breast feeding were significantly higher (p< 0.05) among patients group.

Our results indicated that parity, age at first delivery, and breast feeding had no effect on the BC risk.

In the preset study, the frequency of BRCA1 (185delAG) mutation was 10% among the patients. This is similar to that reported by Santrosa *et al.*,⁽²⁴⁾ who found the mutation in 10% of Italian females with familial BC, but lower than that reported by

Lahad *et al.*, $(19\%)^{(10)}$ in Ashkenazi Jewish. However our result was higher than that reported by Peelen *et al.*, $(1.2\%)^{(25)}$ This variation in the frequency may be attributed to the ethnic differences.

The age is an important risk factor for breast cancer as the risk increases steadily with age, so occurrence of breast cancer in young age group gives strong implication for the presence of inherited genetic predisposition for breast cancer.⁽¹⁰⁾ In the present study, two patients with 185delAG mutation were below 40 years (2/25; 8%). Several studies found frequencies of 3.5% ,6.7% , 9%, and 20%⁽²⁶⁻²⁹⁾ in the patients below 40 years of age . In the present study the other 2 patients carried the mutation were above 40 years (2/15; 13.5%) at the age of onset of the disease. Other studies reported frequencies of $1.9\%^{(26)}$ and $2\%^{(27)}$ in patients above 40 years of age . Although the differences between the results of many studies conducted in different populations, most of these studies concluded that BRCA1 185delAG mutation is a strong candidate for early onset breast cancer than in late onset breast cancer. This was inconsistent with our results as the frequency of the mutation was higher in the old age group . This may be attributed to a possible impact of gene-environment interaction which delays the onset of the BC in the old age group .

The occurrence of multilocular cancer is usually associated with inherited mutation of one of cancer predisposing genes, so occurrence of bilateral BC or BOV is suggestive for the presence of dominantly inherited mutation of one of the breast cancer susceptibility genes.⁽³⁰⁾ In the present study 2 out of 5 (40%) patients with bilateral breast cancer, 1 out of 2 (50%) patients with BOC and 1 out of 33 (3.0%) patients with unilateral breast cancer were found to have the mutation. Steinmann *et al.*,⁽³¹⁾ found that the frequency of the mutation was not different between the

unilateral and bilateral BC patients and explained the development of bilateral BC familial aggregation additional of to susceptibility factors modifying the penetrance of BRCA1 mutation. Other studies reported a mutation frequency of $20\%^{(32)}$ to $100\%^{(33)}$ in BOC. In the present unequal number of study, although patients with unilateral, bilateral BC and BOC, are present, our results indicate that BRCA1 (185delAG) frequency is higher in bilateral BC and BOC than in unilateral BC Egyptian female patients.

A positive family history of breast cancer usually reflects genetic susceptibility and it can be considered as one the strongest risk factors for the disease.^(34,35) In the present study, 2 out of 15 (13.5%) patients with positive family history of BC had BRCA1 gene mutation while 2 out of 25 (8%) patients with negative family history had the mutation. (Result) Our finding is higher than that reported by Friedman *et al.*,⁽³⁶⁾ (7.4%) and Guran *et al.*,⁽³⁷⁾ (5.8%) among patients with positive family history of BC. However it was lower than that reported by Kumer *et al.*,⁽³⁸⁾ (21%) and Schubert *et al.*,⁽³⁹⁾ (40%). Our results indicated that a considerable proportion of the familial risk of BC is attributable to genes other than BRCA1(185delAG) mutation.

In conclusion, our results indicated that BRCA1 (185deIAG) mutation has a role in breast caner but a considerable proportion of the early breast cancer and familial breast cancer may be due to genes other than BRCA1 (185deIAG) mutation. Also, bilateral breast cancer and breast ovarian cancer patients were likely of having the mutation than unilateral breast cancer patients.

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