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B Cell Lymphoma-2 (Bcl-2) in Serum Increased with Breast Cancer in Egyptian Women

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Abstract

Breast cancer is the most commonly diagnosed female-specific cancer, and shows an increasing trend in diagnosed cases worldwide. An estimated one in eight women will develop breast cancer in her lifetime. BC is estimated to be the most common female cancer in Egypt, and the incidence rate among Egyptian women is 48.8/10⁵. Evasion of cell death is now acknowledged as a hallmark of cancer, required to overcome the counterbalancing effects of cell death on enhanced cell proliferation. B cell lymphoma-2 (Bcl-2) gene involves in cell survival mechanism rather than cell proliferation and it prolongs cell life by preventing apoptosis via activating different signaling routes which are induced by various agents. We aimed in this study to investigate whether the Bcl-2 activities are associated with breast cancer in Egyptian women, and correlated to different clinicopathological features of the disease. The study was conducted on forty five females; thirty women with different stages of breast cancer, and fifteen normal healthy females were included as a control group. All patients under study were subjected to full history taking and clinical examination. Fresh blood samples were obtained from all subjects, serum separation was done for measurement of Bcl-2 levels by ELISA technique. Our results showed that there were significantly higher serum Bcl-2 levels in breast cancer patients before surgery than in normal healthy controls. The results obtained also revealed that the mean value of Bcl-2 for the breast cancer patients was significantly higher than the control group. We almost can say that the Bcl-2 values would increase with advanced stages of breast cancer.

Keywords: B cell lymphoma-2 (Bcl-2), Breast cancer, Egyptian women.

Introduction

Cancer can be viewed as the result of a succession of genetic changes during which a normal cell is transformed into a malignant one while evasion of cell death is one of the essential changes in a cell that cause this malignant transformation⁽¹⁾. As early as the 1970's, Kerr et al had linked apoptosis to the elimination of potentially malignant cells, hyperplasia and tumor progression^(2, 3). Hence, reduced apoptosis or its resistance plays a vital role in carcinogenesis. There are many ways a malignant cell can acquire reduction in apoptosis or apoptosis resistance. Generally, the mechanisms by which evasion of apoptosis occurs can be broadly divided into: 1) disrupted balance of pro-apoptotic and anti-apoptotic proteins (Bcl-2), 2) reduced caspase function and 3) impaired death receptor signaling⁽⁴⁾.

Bcl-2 was the first protein of this family to be identified more than 20 years ago and it is encoded by the Bcl-2 gene, which derives its name from B-cell lymphoma 2, the second member of a range of proteins found in human B-cell lymphomas with the t (14; 18) chromosomal translocation. All the Bcl-2 members are located on the outer mitochondrial membrane. They are dimmers which are responsible for membrane permeability either in the form of an ion channel or through the creation of pores⁽⁵⁾.

Apoptosis within the breast occurs with each hormonal cycle throughout a woman's reproductive lifespan⁽⁶⁾. Some investigators demonstrated that Bcl-2 family proteins are key regulators of cell survival during lactation and are critical for inducing cell death of the milk producing epithelium once lactation ceases⁽⁷⁾. Less is known regarding the Bcl-2 family in lobular (menopausal) involution of the breast. However, increasing

evidence suggests that the extent to which lobular involution occurs is directly proportional to a woman's risk of developing post-menopausal breast cancer. Nearly 70% of all breast cancers occur in post-menopausal women. It is therefore important to understand the molecular mechanisms contributing to lobular involution. Given the importance of Bcl-2 proteins in other stages of breast development, these observations support the exploration of Bcl-2 family proteins in lobular involution⁽⁶⁾. However, Bcl-2 and Bcl-xL over-expression in combination with oncogene-induced proliferation resulted in lumen filling, a morphological characteristic of DCIS, an early pre-malignant state, and these findings demonstrated that pro-proliferative and anti-apoptotic signals cooperate in early MEC transformation and that anti-apoptotic Bcl-2 proteins contribute to this process⁽⁸⁾.

Resistance to antigrowth stimuli is the major hallmark of cancer cells which is due to the overexpression of antiapoptotic Bcl-2 proteins. The overexpression of Bcl-2 family proteins also contributes to the resistance and recurrence of cancer⁽⁹⁾. Bcl-2 family proteins also mediate resistance to targeted breast cancer therapies. The anti-HER2 antibody, trastuzumab, is clinically approved for use in 20% of all breast cancers that exhibit *HER2* gene amplification. However, many *HER2*-amplified tumors display innate trastuzumab resistance, while others rapidly acquire trastuzumab resistance. Trastuzumab-resistant *HER2*+ breast cancer cell lines frequently upregulate Bcl-2 and decrease Bax as a means of enhancing cell survival⁽¹⁰⁾.

Materials and Methods

This study was conducted on thirty women with different stages of confirmed breast cancer recruited from Clinical and Experimental Surgery Department, Medical Research Institute, Alexandria, Egypt. Fifteen ages matched normal healthy individuals were also included as a control group. All individuals under study were subjected to the following: Full History taking, clinical examination, radiological investigations, and histopathological examination. Venous blood sample was collected from each participant into serum separator tube. Serum separation was done by centrifugation at 4000-5000 rpm/min for 5 minutes. Analysis of circulating levels of Bcl-2 in serum samples was done by ELISA technique according to Bender MedSystems GmbH (eBioscience), 2015 manufacturer ⁽¹¹⁾. An anti-human Bcl-2 coating antibody is adsorbed onto microwells. Human Bcl-2 present in the sample or standard binds to antibodies adsorbed to the microwells. Absorbance of each microwell was read on a spectrophotometer using 450 nm as the primary wave length. The plate reader was blanked by using the blank wells. The absorbance of both the samples and the standards were determined. The average absorbance values were calculated for each set of duplicate standards

and samples. A standard curve was created by plotting the mean absorbance for each standard concentration on the ordinate against the human Bcl-2 concentration on the abscissa. A best fit curve was drawn through the points of the graph. The concentration of circulating human Bcl-2 was determined for each sample by found the mean absorbance value on the ordinate and extends a horizontal line to the standard curve. At the point of intersection, a vertical line was extended to the abscissa and read the corresponding human Bcl-2 concentration. Data were fed to the computer using IBM SPSS software package version 20.0. Quantitative data were described using mean and standard deviation for normally distributed data while abnormally distributed data was expressed using range.

Results:

Clinicopathological features of the patients:

Age

The age range in breast cancer patient group was 21-75 years with a mean±SD of 51.3±12.86, while it was 26-62 years in control group with a mean ±SD of 47.8±9.30. There is no significant difference between the mean age values of the two studied groups (Table 1).

Table (1): Age distribution of studied groups.

Age (years)	Breast cancer (n=30)	Control (n=15)
Range	21-75	26-62
Mean	51.3	47.8
Std. Deviation (SD)	12.86	9.30

Menopausal status

The menopausal status of all subjects under study is summarized in table (2). In breast cancer group 13 out of 30 patients were premenopausal

(43.3 %) while 17 were post menopausal (56.7 %). In control group 10 out of 15 females were premenopausal (66.7 %) while 5 were post menopausal (33.3 %).

Table (2): Menopausal status among studied groups.

Menopausal status	Breast cancer (n=30)		Control (n=15)	
	No.	%	No.	%
Pre-menopausal	13	43.3	10	66.7
Post-menopausal	17	56.7	5	33.3

Family History for breast cancer

The family history of all subjects under study is summarized in table (3). In breast cancer patient group 14 patients had positive family history

to breast cancer (46.7 %) where are 16 had negative family history (53.3 %), while in control group 2 females had family history to breast cancer (13.3 %) and 13 had no family history to the disease (86.7 %).

Table (3): Family History for breast cancer among studied groups

Family History for breast cancer	Breast cancer (n=30)		Control (n=15)	
	No.	%	No.	%
Positive	14	46.7	2	13.3
Negative	16	53.3	13	86.7

Hormone Replacement Therapy (HRT) use

In breast cancer patients, 13 out of 30 HRT used (43.3 %) and 17 out of 30 not used (56.7 %) (Table 4).

Table (4): HRT used by breast cancer patient group.

HRT	Breast cancer (n=30)	
	No.	%
Used	13	43.3
Not used	17	56.7

Histopathological tumor type

According to the results of histopathological examination; in breast cancer patients, 17 patients had DCIS type (56.7 %) and 13 had IDC (43.3 %) (Table 5).

Table (5): Distribution of the breast cancer patients according to histopathological type.

Histological tumor type	Breast cancer (n=30)	
	No.	%
Ductal carcinoma in situ (DCIS)	17	56.7
Infiltrated Ductal Carcinoma (IDC)	13	43.3
Other types	0	0
Unknown	0	0

Tumor grade

In patients group, 15 cases were found to be grade I (50%), 5 cases were grade II (16.7 %), 6 cases were grade III (20 %), and 4 cases unknown grade (13.3 %), while neither of them were at grade IV (Table 6).

Table (6): Distribution of the breast cancer patient cases according to tumor grade.

Tumor grade	Breast cancer (n=30)	
	No.	%
I	15	50
II	5	16.7
III	6	20
IV	0	0
Unknown	4	13.3

Estrogen Receptor/ Progesteron Receptor (ER/PR) expression

Immunohistochemical (IHC) analysis revealed that 13 patients had positive ER/PR (43%), 13 had negative ER/PR (43.3 %), and 4 were unknown results (13.3 %) (Table 7).

Table (7): ER/PR expression in breast cancer patient group.

ER/PR expression	Breast cancer (n=30)	
	No.	%
Positive	13	43.3
Negative	13	43.3
Unknown	4	13.3

Human Epidermal Growth Factor Receptor 2 (HER-2) expression

Immunohistochemical (IHC) analysis revealed that 10 patients had positive HER-2 (33.3%), 16 had negative HER-2 (53.3 %), and 4 were unknown results (13.3 %) (Table 8).

Table (8): HER-2 expression in breast cancer patient group.

HER-2 expression	Breast cancer (n=30)	
	No.	%
Positive	10	33.3
Negative	16	53.3
Unknown	4	13.3

Bcl-2 levels:

The mean ±SD for Bcl-2 values were 2.01±1.91 and 1.17±0.17 in breast cancer and control groups, respectively. The mean Bcl-2 level was significantly (p=0.025) increased in breast cancer patients when compared to the control group (Table 9).

Table (9): Bcl-2 levels of studied groups.

Bcl-2 levels (ng/ml)	Breast cancer (n=30)	Control (n=15)	p-value*
Range	1.0-10.0	0.75-1.4	0.025
Mean	2.0050	1.1733	
Std. Deviation (SD)	1.91	0.17	

* = significant at p ≤ 0.05 level

Bcl-2 levels and clinicopathological features of breast cancer patients:

Table (10) summarizes Bcl-2 levels and clinicopathological parameters in breast cancer patients. Our data revealed that there was a significant difference in the mean values of **Bcl-2** in the different clinicopathological parameters.

Statistical analysis of results indicated that there was a significant increase in the mean value Bcl-2 in patients ≥50 years than <50 years (p= <0.005), in

post menopausal than premenopausal (p= 0.006), in –ve family history than +ve (p= <0.005) , in not used HRT than HRT used (p= <0.005) and in IDC patients than DCIS patients (p= <0.005). In addition, it was found that there was a significant increase in the mean values of Bcl-2 for grade III than grade I&II (p= 0.05). We almost can say that the Bcl-2 values would increase with advanced stages of breast cancer. Also, it was observed that there was a significant increase in mean values of Bcl-2 in patients with –ve ER/PR and Her-2

expression than +ve expression patients (p=<0.005).

Table (10): Bcl-2 levels and clinicopathological features of breast cancer patients.

	No. of cases	Bcl-2 (ng/ml)	
		Mean ±SD	P- value
Age (year)			
≥50	18	10.09±32.51	<0.005*
<50	12	1.43±0.41	
Menopausal status			
Pre	13	1.4 ± 0.4	0.006*
Post	17	10.62±33.4	
Family history			
Positive	14	2.54±2.72	<0.005*
Negative	16	10.2±34.6	
HRT use			
Used	13	2.42±2.86	<0.005*
Not used	17	9.84±33.55	
Tumor type			
DCIS	16	2.33±2.59	<0.005*
IDC	14	11.53±37.0	
Tumor grade			
I	15	1.49±0.45	0.05*
II	5	2.39±2.65	
III	6	25.12±56.29	
ER/PR			
Positive	13	2.08±2.42	< 0.005*
Negative	13	12.85±38.24	
HER-2			
Positive	10	2.57±2.51	< 0.005*
Negative	16	10.31±34.59	

* = significant at p ≤ 0.05 level

Discussion

Evasion of cell death is now acknowledged as a hallmark of cancer, required to overcome the counterbalancing effects of cell death on enhanced cell proliferation ⁽¹²⁾. Bcl-2 gene involves in cell survival mechanism rather than cell proliferation and it prolongs cell life by preventing apoptosis via activating different signaling routes which are induced by various agents ⁽¹³⁾.

In the present work, there were significant higher serum Bcl-2 levels in breast cancer patients

before surgery than in normal healthy controls. The mean±SD value for the breast cancer group was 2.01±1.92 ng/ml, and 1.17±0.17 ng/ml for the control group. Statistical analysis of results indicated that there was a significant increase in the mean value Bcl-2 in patients ≥50 years than <50 years (p= <0.005), in post menopausal than premenopausal (p= 0.006), in –ve family history than +ve (p= <0.005) , in not used HRT than HRT used (p= <0.005) and in IDC patients than DCIS patients (p= <0.005). In addition, it was found that there was a significant increase in the mean values

of Bcl-2 for grade III than grade I&II ($p= 0.05$). We almost can say that the Bcl-2 values would increase with advanced grades of breast cancer. Also, it was observed that there was a significant increase in mean values of Bcl-2 in patients with -ve ER/PR and Her-2 expression than +ve expression patients ($p= <0.005$) .

Our results agreed with those of Mahdy et al. (2011)⁽¹⁴⁾ and Kallel-Bayouhd et al. (2011)⁽¹⁵⁾ who reported high levels of Bcl-2 in breast cancer patients before surgery compared with normal controls. The increase of Bcl-2 in cancer cells points to a potentially critical role of this anti-apoptotic protein in breast cancer progression. Overexpression of Bcl-2 protein may serve as a determinant of an advantageous cell survival in breast tumor cells, ultimately leading to tumor progression and metastases⁽¹⁶⁾. In addition, Liu et al, (2014)⁽¹⁷⁾ found that the downregulation of Bcl-2 in human breast cancer cells has been associated with a good prognosis.

Similar results obtained by Honma et al, (2015)⁽¹⁸⁾ demonstrated that Bcl-2 is positivity and significantly correlated with poor clinical outcome in patients with ER-negative and PR-negative or triple-negative tumors. Bcl-2 is positivity independently predicted recurrence/mortality in hormone receptor-negative or triple-negative cases, but not in ER-positive and/or PR positive cases. It seems that the anti-apoptotic effect of Bcl-2, which usually correlates with poor clinical outcome or resistance to therapy in tumors other than breast cancer, is evident only in cases without hormone receptors and without adjuvant therapy. They suggested that the anti-apoptotic nature of Bcl-2 is clearly exhibited under such conditions. The prognostic value of Bcl-2 is more evident in postmenopausal women.

Tawfik et al. (2012)⁽¹⁹⁾ also found that positivity of Bcl-2 was in correlation with more aggressive tumor histology and longer overall survival in non-triple-negative group whereas bcl-2 positivity was found to be related with shorter survival in triple-negative group, interestingly.

Furthermore, Sezgin et al, (2014)⁽²⁰⁾ showed that Bcl-2 expression both in Triple Negative breast cancer (TNBC) and non-TNBC patients, and analysis of expression of Bcl-2 may be meaningful for deciding treatment strategies for TNBC. They suggested that treatment strategies targeting Bcl-2 seem to be promising for this aggressive disease with no specific treatment.

Berrak et al., (2016)⁽¹³⁾ reported that the increased expression level of Bcl-2 was found correlated with drug resistance mechanism in various cancer cells including breast cancer cases.

In partial agreement with our study, Merino et al, (2015)⁽²¹⁾ demonstrated that approximately 75% of primary breast cancers express high levels of BCL-2, with a predominance in ER-positive tumors: BCL-2 is overexpressed in nearly 85% of ER-positive tumors, 50% of HER2- positive tumors, 41% of TNBCs and 19% of basal-like tumors. These findings appear to be consistent with gene expression profiling studies, where Bcl-2 is predominantly expressed in ER-positive tumors. Interestingly, the frequency of BCL-2-positive tumors is lower (31%) in BRCA1- associated cancers, compared with cancers without BRCA1 mutations, probably attributable to their triple-negative status.

However, other studies^(22, 23) reported that Bcl-2 expression was positively correlated with ER and PR expression, but negatively correlated with HER2 expression, grade, and tumor size,

confirming Bcl-2's association with favorable prognostic factors or differentiated markers.

Furthermore, Williams et al, (2015) ⁽²⁴⁾ reported that Bcl-2 expression frequently correlates with ER expression levels in ER+ breast cancers. Several studies have indicated that estrogen may up-regulate Bcl-2, allowing for evasion of apoptosis ⁽²⁵⁻²⁷⁾.

According to our study, we can conclude Breast cancer patients showed high serum levels of Bcl-2 as compared to normal controls, and a clear increase with the advanced grade of the tumor. Serum Bcl-2 can be considered a biomarker for breast cancer patients, and could be used for monitoring the effect of chemotherapy on these patients. Our study may be applied for prevention and early intervention strategies in Egyptian population.

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