Prognostic Value of BCL2 in Women Patients with Invasive Breast Cancer

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Abstract

Background: Breast cancers are heterogeneous, making it essential to recognize several biomarkers for cancer outcome predictions especially in young women where the classical prediction parameters are not suitable. The goal from this study is to evaluate the impact of B cell lymphoma 2 (BCL2), P53 and Ki-67 proteins expression on survival in young women patients with invasive ductal carcinoma. Patients and methods: Samples and clinical data from 238 patients were collected between 2003 and 2017. They were selected according to 2 criteria: age <40 years old and most of them are affected by an Invasive Ductal Carcinoma. We evaluated BCL2, P53 and ki-67 expression by immunochemistry test, and then we assessed correlations of these biomarkers expression with patient's clinicopathological characteristics and survival. Results: Triple negative breast cancer group showed a high frequency among our cohort but we emphasize an almost equitable distribution among all molecular groups. Contrary to other studies which reported that luminal A was correlated with better prognosis, our analysis demonstrated that luminal A is correlated with the Scarff, Bloom and Richardson (SBR) grading 2 or SBR grading 3. To better investigate the prognosis, we analyze three biomarkers known by their impact on physiopathology behavior on breast cancer BCL2, ki-67 and P53. BCL2 is the more relevant one, it was correlated with molecular subtypes (p=0.0012) and SBR grading (p=0.0016). BCL2 seems to be the good prognostic biomarker related to survival (p=0.004) with a protective role among patients when endocrine therapy is not provided and Lymph Node (LN) involvement is positive (p=0.021, p=0.000 respectively). Conclusions: The classical prognostic parameters based mainly on the molecular classification in breast cancer seem insufficient in the case of young women. BCL2 protein expression analysis provides a better prognostic value. BCL2 should be clinically associated in current practice when young women specimens are diagnosticated.

Keywords: Breast cancer- BCL2- young women- prognosis

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Introduction

Breast cancer (BC) represents the most common disease in the world (WHO, 2013). It is the most frequent malignant neoplasm affecting Tunisian female patients with an incidence of 27.1/100,000 inhabitants (Maalej et al., 2004). BC is a heterogeneous disease with molecular subtypes that have biological distinctness and different behavior. This heterogeneity covers epidemiological risk factor, natural histories, biological etiology, clinical outcome, pathological characteristics and response to therapies (Peppercorn et al., 2008). Perou et al., (2000) identified four BC subtypes on the basis of gene-expression profiling of 39 invasive breast tumors and three normal breast specimens.

Molecular subtypes are defined by the immunohistochemical expression of estrogen receptor

(ER), progesterone receptor (PR), and human epidermal growth factor receptor 2 (HER2). Analyses of (ER), (PR) and (HER2) expressions have been routine practice for years. Endocrine therapy is considered for patients with hormone receptor positive (ER + and/or PR +) tumors. Furthermore, the ER negative (ER-), PR negative (PR-), and HER2 negative (HER2-) tumors, also known as triple-negative phenotype TNBC (ER-/PR-/HER2-) is characterized by expression of cytokeratin 5/6 (CK5/6) and/or the epidermal growth factor receptor (EGFR) (Nielsen et al., 2004; Change et al., 2008). For these patients, chemotherapy is the only available treatment. Classification and identification of new important biomarkers may help in prognostication and targeting to treatment to those most likely to benefit (Blows et al., 2010).

Many studies were focused to determine the impact

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E-Z Ayadi et al

of several biomarkers on survival and their inclusion in everyday clinical practice. Since 1979 studies were focused on the important role of the anti-proliferative biomarker P53, as the first tumor suppressor gene described (Gasco et al., 2002; Haldar et al., 1994). Then studies have demonstrated that P53 impact in clinical outcome remains controversial (Yang et al., 2013). While, BCL2 is an anti-apoptotic protein and its role is still also controversial. Some authors reported that BCL2 constitute a strong protein marker in BC (Kallel-Bayoudh et al., 2011), and is favorable prognostic factor in ER positive (Callagy et al., 2008). Other have shown BCL2 protein and gene expression to be a promising prognostic and predictive marker in human cancers (Von Minckwitz et al., 2008) especially node-negative BC (Ali et al., 2012; Paik et al., 2004).

BC among young women is known by its significant association with a poor prognosis (Anders et al., 2009) and generally correlated with worse pathological features including higher stage at presentation, grade 3, and estrogen receptor (ER) negative status (Gajdos et al., 2000). The distribution of molecular subtypes among women under 40 years old present less Luminal cancers and more HER2 positive and TNBC tumors than older women (Colleoni et al., 2002). Taking into account the heterogeneity of this disease, we choose to investigate the role of Ki-67, P53 and BCL2 on pathological and survival data in young Tunisian women patients reporting the correlation between biomarkers and each impact on survival.

Materials and Methods

Study patients

This is a retrospective cohort of young women with invasive BC. All patients are under 40 years old. They were studied between 2003 and 2017. Clinical features of patients were collected in collaboration with the anatomy pathology laboratories and oncologists. Recorded data provided age at diagnosis, BC grade determined by Elston-Ellis modification of Scarff-Bloom-Richardson grading system (SBR Grade I, II, III), tumor size (<2 cm, 2-5 cm, ≥ 5 cm), lymph node (LN) status (positive or negative) and vascular invasion (VI) status (positive or negative). In total, 238 patients from Tunisia accurately from Sfax, Sousse and Djerba regions were selected. Clinical analysis also provided data on whether or/ not the patient had been treated with adjuvant hormonal therapy or adjuvant chemotherapy.

Immunohistochemistry

Tissues samples from patient cohort were prepared in paraffin-embedded slides for protein investigation. Staining was done continuously and progressively according the specimen reception, for all samples, ER, PR, HER2, Ki-67, P53 and BCL2 were revealed using immunohitochemistry (IHC) tests. All markers were performed with specific antibody [primary antibody anti-human]: HER2 (Novocastra NCL-CB11), ER (Novocastra NCL-ER-6F11), PR (Novocastra NCL-PGR-312), Ki-67 (NovocastraNCL-L-KI-67-MM1), P53 (NovocastraNCL-L-P53-DO7), BCL2 (Novocastra NCL-L-BCL2). Secondary antibodies and streptavidine-HRP (Novocastra) were used for recognized antigen revelation. The enzymatic activation of the chromogen results in a visible reaction product at the antigen site. The specimen may then be counterstained using Hematoxyllin and cover slipped. Under a white light microscope, IHC expression interpretations from samples patients compared to positive controls tissues (using Colonic carcinoma for P53 and Tonsil lymphoma for BCL2 or Ki67 immunostaining) or negative controls (using normal breast showing any immunostained cell) were performed using conventional image analysis software.

Immunostaining scoring

Experienced pathologist (J.R) who had no knowledge of the patients' clinical status and outcomes evaluated immunohistochemistry (IHC) of ER, PR, HER2, Ki-67, P53 and BCL2 proteins expression. Hormone receptors (ER and PR) were considered positive when more than 1% of infiltrating tumor cell nuclei were stained (Hammond et al., 2010). Tumors were considered positive for HER2 if immunostaining was scored as 3+ according to Wolff criteria (Wolff et al., 2007; Wolff et al., 2013) and cancers with HER2 scored as 2+ (indeterminate) were assessed through fluorescent in situ hybridization (FISH) test. Immunostaining of Ki-67, P53 or BCL2 biomarkers was scored on the basis of the percentage of positive tumor cells. Therefore, positive cells were defined as cells with strongly and clearly brown immunostaining compared to controls. The scoring was graded according to the percentage of Ki-67, P53 and BCL2 -positive cells, the following scores were assigned: 0 (0%), 1 (1%), 2 (2%) 99 (99%), 100 (100%).

Breast Cancer subtyping

Based on ER, PR, HER2 and Ki-67 expression with IHC analysis, we classified samples in to 5 groups characterized by; LA group ER/PR +, HER2- and Ki-67 < 20%, LB like-group ER/PR +, HER2- and Ki-67 \geq 20% (Tashima et al., 2015), LB group ER/PR + and HER2+, HER2 group ER/PR- and HER2+ and TNBC group ER/PR - and HER2-.

Statistical analysis

Descriptive statistics for clinicopathological features were estimated using simple frequency. Bivariate analysis was performed by assessing the correlation between all the markers and prognosis using chi-square test for binary variables, Pearson-rank correlation for quantitative variables and ANOVA test for quantitative variation among class parameter. Survival analyzes were studied by Kaplan Meier Test. A p-value < 0.05 is considered as statistically significant. All these tests were done using the SPSS 13.0 software version or R-3.2.3 environment.

Results

Clinicopathological characteristics of young women patients

The clinicopathological characteristics are listed in Table 1. Among 238 young women affected with invasive BC, the mean of patients' age at diagnosis is 35 ± 4.2 years old ranged from 17-40 years old. Tumor size ranged from 0.8 cm to 16 cm with a mean size of $4.24 \text{ cm} \pm 2.7 \text{ (pT2)}$ and 6.2% of patients are affected with inflammatory BC. (Table 1). Informative BC cases about LN involvement and vascular invasion showed 64.1% and 13.7% positive cases respectively. Only 9.5% were grade I BC (SBR classification), and 90.5% were grade II or III (43.7% and 46.8% respectively).

Based on immunohistochemistry staining of ER, PR and HER2 positive rates were 60.5%, 60.1% and 37.1%

Table 1. Clinical Pathology Characteristics of YoungWomen Patients with Breast cancer

Clinicopathological Data	Characteristics	Frequency	Percent (%)
Tumor size	T1<2 cm	38	18.2
Total=209	2 <t2< 5="" cm<="" td=""><td>110</td><td>52.6</td></t2<>	110	52.6
	T3>5 cm	48	23
	T4 inflammatory	13	6.2
Lymphe Node status	Negative	69	35.9
Total=192	Positive	123	64.1
SBR grading	Ι	21	9.5
Total=222	II	97	43.7
	III	104	46.8
Vascular invasion	Negative	88	86.3
Total=102	Positive	14	13.7
ER status	Negative	94	39.5
Total=238	Positive	144	60.5
PR status	Negative	95	39.9
Total=238	Positive	143	60.1
HER2 status	Negative	149	62.9
Total=237	Positive	88	37.1
Ki-67 20% status	Negative	71	46.4
Total=153	Positive	82	53.6
MolecularSubtypes	Luminal A	38	18.72
Total=203	Luminal B Like	30	14.77
	Luminal B	53	26.1
	HER2+	35	17.24
	TNBC	47	23.15
Overall Survival	> 2years	75	54
Total=139	<2 Years	64	46
Metastasis	0	106	67.9
Total=156	1	50	32.1

ER, estrogen receptor; HER2, human epidermal growth factor receptor 2; PR, progesterone receptor;SBR grading, Scarff Bloom and Richardson grading; T, tumor; LA group, luminal A (hormonal receptor (+), her2 (-) and ki-67 <20%); LB group, luminal B (hormonal receptor (+) and her2 (+)); LB Like group, luminal B (hormonal receptor (+), her2 (-) and ki-67 \geq 20%); HER2 group, human epidermal growth factor receptor 2 (hormonal receptor (-) and her2 (+)); TNBC group, triple negative breast cancer (hormonal receptor (-) and her2 (-)); hormonal receptor=ER and PR.

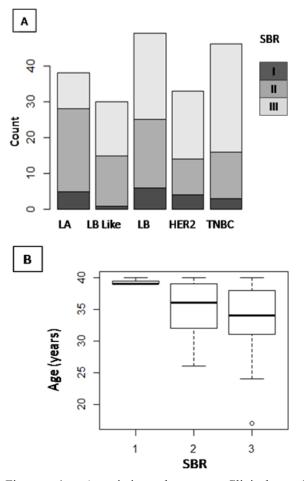


Figure Association between Clinical 1. and Histopathological Parameters of Young Women Breast Cancer Patients. (A) Barplots of SBR grading distribution among molecular subtypes (*p-value*=0.004). (B) Boxplots of patients age according to SBR grading (p-value=0.045). SBR grading, Scarff Bloom and Richardson grading; LA group, luminal A (hormonal receptor (+), her2 (-) and ki-67 <20%); LB group, luminal B (hormonal receptor (+) and her2 (+)); LB Like group, luminal B (hormonal receptor (+), her2 (-) and ki- $\overline{67} \ge \overline{20\%}$; HER2 group, human epidermal growth factor receptor 2 (hormonal receptor (-) and her2 (+)); TNBC group, triple negative breast cancer (hormonal receptor (-) and her2 (-)); hormonal receptor= estrogen receptor and progesterone receptor.

respectively. Ki-67 was assessed in 153 patients. Based on ER, PR, HER2 and Ki-67 rate, samples were classified in to 5 groups as shown in material and methods. Molecular classification showed that Luminal B and TNBC subtypes were the most frequent groups with 26.1% and 23.15% respectively followed by LA and HER2 subtypes (Table 1).

Correlation between clinical and histopathological parameters

Distribution of SBR grading among molecular subtypes showed that BC with grade I represents only 9.5% of tumor cases, while SBRI and SBRIII are the most common in studied cohort (Figure 1-A). Statistical analysis showed that there is significant difference of molecular subtype of BC among SBR grading (p=0.004). These results confirm that BC in young women presents

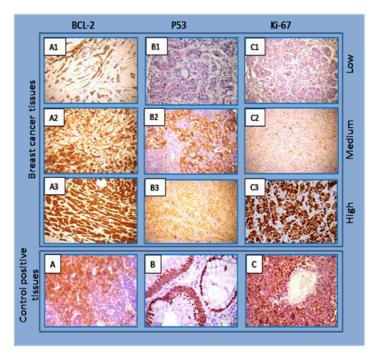


Figure 2. BCL-2, P53 and Ki-67 Expression Analysis by Immunohistochemical Staining in Breast Cancer Tissues. Positive controls revelations are presented by Tonsil lymphoma for BCL2 (A) or Ki67 (C) and Colonic carcinoma for P53 (B) immunostaining. Representative images for low, moderate and high BCL2 (A1-A3), P53 (B1-B3) and Ki-67 (C1-C3) showed biomarkers expression in young women breast cancer specimens. (Original magnification x100).

always an aggressive grade 46.8% (Table 1), therefore HER2+ and TNBC subtypes are the two groups associated with the worst prognosis with SBRIII grading. Moreover, age variation according to the SBR grading shows that the youngest have an SBRIII breast cancers, these data confirm that young age is often associated with the worst grade (p<0.05) (Figure 1-B). However, there is no relationship between age of patients and molecular

subtypes nor lymph node involvement and vascular invasion.

BCL2, P53 and Ki-67 immunodetection in breast cancer

BCL2, P53 and Ki-67 expression were evaluated successfully by IHC at least in 140 BC tissues. Based on the extent and intensity of tumor cells, immunostaining was scored as percentage of positive cells according

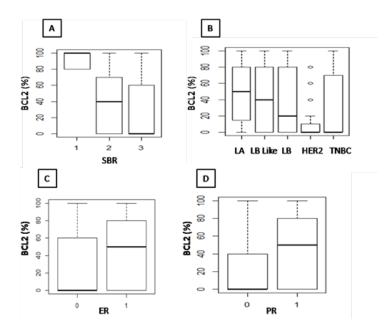


Figure 3. Relationship between BCl-2 Expression and Histopathological Parameters of Young Women Patients with Breast Cancer. Anova tests give significant variation of BCL2 rates among SBR grading (A) (p= 0.00168) showing high percentage with SBR I and molecular groups (B) (p= 0.0012) showing high percentage with LA and LB like groups mainly in comparison with HER2 group. Highly correlations were obtained for BCL2 expression levels with estrogen receptor (C) and progesterone receptor (D) (p-value < 0.001).

3560 Asian Pacific Journal of Cancer Prevention, Vol 19

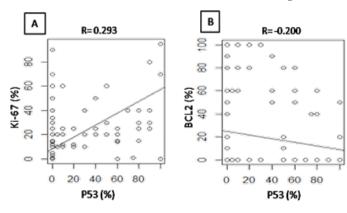


Figure 4. Biomarkers Associations. Plots of P53 levels according to ki-67 levels (A) showing highly significant of positive correlation (R = 0.293, p < 0.001) but less significant of negative correlation with BCL-2 level (B) (R = -0.200, p < 0.05). R, coefficient of correlation

to respective biomarker expression localization. An IHC score was generated as described in methods and quantification was performed of cytoplasmic expression for BCL2 (Figure 2, A1-A3) and nuclear expression for P53 (Figure 2, B1-B3) or ki-67 (Figure 2, C1-C3). Choosing this method for scoring is supported by IHC results obtained showing a measurable graduation

immunostaining revelation, indeed we observe clearly for all biomarkers an expression level ranging from low (% of positive cells \leq 30%), moderate (30 \leq % of positive cells \leq 70%) to strong (% of positive cells \geq 70%) (Figure 2). BCL2, P53 and Ki-67 immunodectection are compared and confirmed according to positive controls (Figure 2: A-C) and negative controls (data not shown).

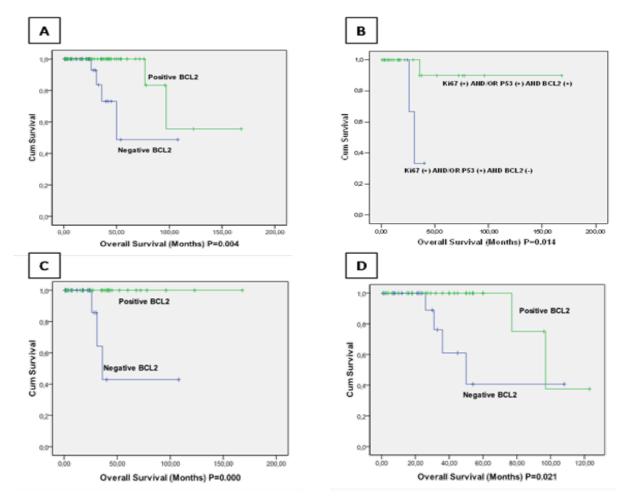


Figure 5. Survival analysis according to BCL2 expression. Graph (A) represents overall survival according to BCL2 expression. Graph (B) shows overall survival according to BCL2 protective impact in presence of Ki-67 or P53. Graph (C) indicates data of overall survival according to BCL2 presence in invasive breast cancer in case of positive node involvement. Graph (D) gives data of overall survival according to BCL2 presence in invasive breast cancer when endocrine therapy is not provided.

E-Z Ayadi et al

Correlation between all biomarkers and clinical data

Ki-67 is a marker of cell proliferation, apart from its tissues expression to perform molecular classification, we have investigated its status as a related biomarker among our cohort. Ki-67 expression rate is significantly related to hormonal receptors status (p=0.03) and LN involvement (p=0.045). However, Ki-67 levels were slightly progressive with significance limitation according to the SBR grading (p=0.068) and no significant variations were observed between ki-67 expression and tumor seize. For BCL2 investigations a significant variation of its rate among molecular groups (Figure 3-B, p=0.0012) showing high percentage with LA and LB Like groups mainly in comparison with HER2 group.

In addition, statistical analysis showed highly significant differences between the BC SBR grading according to BCL2 expressions, indeed its rates among SBR grading (Figure 3-A, p= 0.00168) showed high percentage in SBR I. Therefore, a similar highly significant correlations were observed between BCL2 levels and ER status (Figure 3-C, p= 0.000882) and PR status (Figure 3-D, p= 0.000528).

For P53 investigations, surprisingly no significant relationship was observed regarding P53 expression with clinical parameters. However relationship investigations between analyzed biomarkers showed that P53 was positively correlated to Ki-67 (Figure 4A, R= 0.293) (p<0.001) and negatively correlated to BCL2 (Figure 4B, R=-0.200) (p<0.05) these suggest that P53 can be indirectly related to a poor prognosis.

Biomarkers and disease outcome

A significant correlation between BCL2 and overall survival (p=0.004) (Figure 5-A). These data show that BCL2 is an important biomarker to evaluate prognosis. Patients expressing BCL2 have a better outcome than patients with negative BCL2. However, no significant results were found between survival and Ki-67 or P53 (data not shown).

To better investigate the role of BCL2 and its relation with the two important biomarkers Ki-67 and P53, we had analyzed survival according to BCL2 when one of the two biomarkers Ki-67 and P53 is at least expressed. Figure 5-B indicates that BCL2 is a strong biomarker with a protective effect and that its presence is significantly correlated with a better overall survival even if Ki-67 or/ and P53 are expressed (p=0.014).

Patients who have positive node involvement have more risk to recurrence, we wanted to explore if BCL2 can better evaluate the prognosis for this group of patients. Figure 5-C, demonstrated that this group of patients has a better outcome of survival if they express BCL2 (p=0.000). Surprisingly when we analyzed the group of patients who didn't receive endocrine therapy and focused on the impact of BCL2, we found that patients who expressed BCL2 have better survival than those who didn't express BCL2 (p=0.021) (Figure 5-D).

Discussion

Our results showed that 46% of our patients have a **3562** *Asian Pacific Journal of Cancer Prevention, Vol 19*

tumor size ranging from 2 to 5cm followed by T3 >5cm with a percentage of 25.7 (Table 1). This is probably due to the rapid proliferation features of BC in young patients compared to elderly patients (Maggard et al., 2003; Althuis et al., 2009) and that the breast tissue of younger women is denser and therefore it is more difficult to detect BC by physical examination and mammography (Schreer et al., 2009; Lin et al., 2009).

In our cohort, BC patients are most likely to present luminal B subtype followed by TNBC. These results disagree with others showing that younger breast cancer patients had a higher prevalence of luminal A and a lower prevalence of basal-like subtype compared with older patients (Tang et al., 2011; Farouk et al., 2016). However, our cohort molecular feature resemble to others who found that LB was the most common type among young women patients (Collins et al., 2012; Fourati et al., 2014). Data disagreement can be explained by differences in genetic backgrounds of populations studied, so comparable results are supported by population's consanguinity through different nations.

This study focuses on the different interactions between BCL2, P53 and Ki-67 in front of clinical data. Significant correlations have be found between Ki-67 and others biomarkers, assuming that Ki-67 is positively correlated with P53, high SBR grading and lymph node involvement. However, we didn't found any significant correlations between Ki-67 or P53 with overall survival (data not shown). Otherwise BCL2 was found positively correlated with ER and PR status and inversely correlated with P53 and high SBR grading (p<0.05 and p<0.01 respectively) (Figure 3 and Figure 4).

Recent studies have shown that BCL2 protein is a promising prognostic and predictive marker in BC, especially in hormone receptor-positive, LN negative BC (Kallel-Bayoudh et al., 2011; Ali et al., 2012; Paik et al., 2004). BCL2 protein expression in tumors was first studied in non-Hodgkin's lymphoma (Yang et al., 2013; Chan et al., 2011; Chin et al., 2005; Horwitz and Alpha-crystallin, 2003; Lambi et al., 2011). Many studies have demonstrated that BCL2 is a strong prognostic factor correlated with a better survival (Martinez-Arribas et al., 2007; Dawson et al., 2010; Callagy et al., 2006). Our results confirm this findings, we proved that BC young women expressing BCL2 have a better Overall Survival (p=0.004) (Figure 5-A).

Moreover, it has been reported that Ki-67 is an important biomarker which provides additional and independent predictive information regarding the response to chemotherapy and the prognosis in a group of patients receiving neoadjuvant treatment for BC (Fasching et al., 2011). So, this marker can be used to select patients who are unable to benefit from chemotherapy, such as those with HER2-negative and hormone receptor-positive tumors with low proliferation (Fasching et al., 2011; Inwald et al., 2013). In despite of some authors who have found that P53 protein was associated with high tumor proliferation rate, early disease recurrence and early death in node negative BC (Allred et al., 1993; Yamamoto et al., 2014). Fountzials and colleagues (2016) have proved that HER2 positive and P53 IHC positive

tumors were associated with increased risk for relapse in the pre-trastuzumab era, while the same phenotype conferred favorable disease free survival (DFS) in the post-trastuzumab era trials.

Keeping in mind these facts, we wanted to focus on supplementary effect that BCL2 can provide in the presence of the 2 prognostic biomarkers Ki-67 and P53. So we devised our patients on two groups; first one expresses at least one of the poor biomarkers and do not express BCL2 and the second one expresses at least one of the poor biomarkers and express BCL2. We analyzed survival in these two groups using Kaplan Meier test. Knowing that high Ki-67/BCL2 index is correlated with short disease free survival (Min et al., 2016). Our result confirmed that BCL2 do have a supplementary positive effect on survival with protective role (p=0.014) (Figure 5-B). Raising this result, we want to know better about this protective role. For this purpose, we studied the protective role of BCL2 among patients with positive node involvement and patients with endocrine therapy not provided. In our cohort 64% are node positive, this criterion is known to be associated with high risk of recurrence and bad survival (Kim et al., 2016), (Table 1). Hence, the protective role of BCL2 was confirmed inside the two groups presenting a better survival correlated with the presence of BCL2 (p=0.000, p=0.021 respectively) (Figure 5-C, D).

In conclusion, different molecular subtypes will lead to different prognosis and therapeutic option. Thus, molecular subtyping is essential for breast carcinoma management. But considering only the molecular group to establish a prognostic value is insufficient in young woman where the physiopathology seems to be different from their counterpart. Our results revealed that BCL2 is the best prognostic biomarker with a protective role. BCL2 expression analysis seems to be essential in medical practice to better evaluate BC prognosis in young women.

Abbreviations

ANOVA, analysis of variance; BCL2, B cell lymphoma 2; BC, Breast cancer; ER, estrogen receptor; HER2, growth factor receptor 2; LA, luminal A; LB, luminal B; LN, lymph node; PR, progesterone receptor; SBR grade (I, II, III), Scarff-Bloom-Richardson grading system; TNBC, triple negative breast cancer; VI, vascular invasion.

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Conflict of Interest

Boutheina Cherif has received research grants from ISECO and Hammadi Ayadi has received research grants from Era Wide UE FP7 Bioprotech Project (Ref N°26625).

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