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Y-Box binding protein 1 expression in breast cancer and its correlation with hormone receptors and other prognostic markers

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Abstract:

INTRODUCTION: The present histologic and immunohistochemical prognostic markers of breast carcinoma do not effectively identify the subset of patients with poor prognosis. Y-Box binding protein 1 (YB1) is a novel biomarker which may identify and aid in targeted personalized therapy for such patients.

MATERIALS AND METHODS: The study was conducted on histopathology specimens of 74 patients of breast carcinoma who had undergone modified radical mastectomy. YB1 immunohistochemistry (IHC) was performed on manual tissue microarray blocks (each having 15 cores). The YB1 expression was quantified in terms of “immunoreactive score” which was correlated with clinical parameters, hormone receptor status, and Her2neu overexpression by IHC. The Her2neu status of the equivocal cases was further evaluated by fluorescent *in situ* hybridization (FISH).

RESULTS: YB1 was positive in 36/74 (48.6%) cases. On IHC and analysis by FISH, 25/74 (34%) cases had Her2neu overexpression. Estrogen receptor (ER) and progesterone receptor (PR) positivity was found in 42% and 36.5% cases, respectively. YB1 immunopositivity was negatively correlated with ER and PR expression, but showed a significant positive correlation with Her2neu expression. No correlation was found with other clinical parameters, tumor stage, and grade, except lymph node involvement, which showed a positive association with YB1 expression. Triple-negative breast carcinoma constituted 25.6% of the total cases, out of which 73.6% were YB1 positive.

CONCLUSION: This study found that YB1 has an association with Her2neu expression. It may in future provide a therapeutic target in Her2neu overexpressing tumors.

Key words:

Manual tissue microarray, triple-negative breast carcinoma, Y-Box binding protein 1

Introduction

Though clinically and morphologically similar cases of carcinoma breast may appear similar, genetically they form a divergent group. Hence, novel genetic markers are needed to prognosticate and stratify cases of breast cancer for appropriate therapy.

Y-Box binding protein 1 (YB1) is a nucleic acid-binding protein encoded by gene YBX1,

which was initially identified as a factor bound to the Y-box (an inverted CCAAT box).^[1,2] It is a pleiotropic, multifunctional protein with many functions including regulation of transcription, translation, cell survival, DNA replication and repair, drug resistance, and epithelial–mesenchymal transition.^[2] It contains a highly conserved cold shock domain, which is involved in binding nucleic acid including RNA as well as dsDNA.^[3]

Within the nucleus, YB1 binds to inverted CCAAT boxes known as YB-1-responsive

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elements to activate or repress transcription.^[2] YB1 has been shown to interact with the epidermal growth factor receptor (*EGFR*) enhancer and the *Her2* promoter and also induces transcription of drug-resistant genes including the multidrug resistance-1 (MDR1) and multi-drug resistance-related protein-1.^[4-6]

In the cytoplasm, YB1 is found in complexes with translationally inactive mRNAs.^[7] This has led to the notion that YB1 is involved in translational repression.

YB1 is expressed in various malignancies including breast carcinoma, colorectal carcinomas, ovarian serous adenocarcinoma, prostate adenocarcinoma, lung carcinoma, large B-cell lymphoma, multiple myeloma, osteosarcoma, embryonal rhabdomyosarcoma, glioblastoma multiforme, and melanoma.^[8] Little is known about the mechanism involved in the overexpression of YB1 in cancers. It was once postulated that it occurred due to gene amplification. However, gene mapping and comparative genomic hybridization failed to reveal any amplification at chromosome 1p34.^[9] Thus, the high expression of YB1 in breast cancer does not appear to be due to gain in copy number and could be due to transcriptional activation.^[10]

In all the studies done so far, YB1 expression correlated with a poor patient prognosis. Tumors expressing high YB-1 had significantly decreased disease-free and recurrence-free survival.

Studies have shown YB1 protein to be elevated in a range of 36%–41% of breast cancers.^[11,12] As in other malignancies, increased expression of YB1 in breast cancer is associated with a poorer prognosis in terms of decreased patient survival and increased recurrence rates.^[13] Nuclear localization of YB1 is thought to be associated with intrinsic MDR1 gene expression.^[5] Studies have further concluded that overexpression of YB1 may be an initiating event for early development of breast cancer by causing centrosome amplification, chromosomal instability, and aneuploidy, hence a potential target for therapy.^[8]

The *Her2* proto-oncogene is located on chromosome 17q21. *Her2* is found to be amplified in 20%–30% of breast cancer.^[14] *Her2* positivity is associated with a number of adverse tumor characteristics such as larger tumor size, higher nuclear grade, and S-phase fraction.^[15] YB1 directly induces transcription of *EGFR* and *Her2* in a S102-dependent manner.^[16]

Materials and Methods

This study was performed on 74 histologically confirmed cases of breast carcinoma from October 2015 to April 2017.

All these patients had undergone modified radical mastectomy. The study was cleared by the Institutional Ethics Committee and informed patient consent was taken. Clinical data of the patients were recorded from the histopathology requisition forms. All the specimens were fixed in 10% buffered formalin and embedded in paraffin wax. Sections stained with hematoxylin and eosin were used to identify the histological type of breast cancer and study other histological features such as presence of *in situ* component, lymph node status, and grading of the tumor. For grading, Nottingham modification of the Bloom Richardson system and for staging, American Joint Committee on Cancer staging were used.^[17,18]

Manual tissue microarray (TMA) without prefabricated recipient block using bone marrow biopsy needle was made as described by Chen and Zhou as it was found to be superior to the technique described by Kononen *et al.* in our department.^[19,20] We constructed six TMA blocks for 74 cases. Each TMA block comprised of tissues from 12 to 15 cases.

Immunohistochemistry procedure

3–4 µm-thick sections were taken from each microarray block for hematoxylin and eosin staining and ER, PR, *Her-2*, and YB1 detection by immunohistochemistry (IHC) on poly-l-lysine-coated slides. The rabbit polyclonal anti-YB1 protein antibody (orb 336555 biorbyt, UK) was used in a dilution of 1:200. Sections from human duodenal biopsy were taken as positive control for YB1 protein expression [Figure 1a].

Immunohistochemical analysis

The IHC staining was assessed as percentage of positive staining cells. For interpretation of ER, PR protein expression nuclear staining was assessed [Figure 2a and b]. >10% of cells showing positive nuclear staining for ER/PR were considered positive. For *Her-2*, 0–1+ was taken as negative and 3+ as positive (ASCO/CAP guidelines 2013) [Figure 2c, d and Table 1].^[21] Cases which were equivocal (2+) by IHC were further evaluated for *Her2* amplification by fluorescent *in situ* hybridization (FISH). For YB1, nuclear staining was assessed by “Immune-reactive score” [Table 2]. The result is the product of the two scores, ranging from 0 to 12, considering YB1 <6 as negative expression and YB1 ≥6 as positive expression [Figure 1].^[22]

Fluorescent *in situ* hybridization

ZytoLight SPEC ERBB2/CEN17 Dual Color Probe Kit was used for the detection of *Her2* gene amplification by FISH. The probe contained green-labeled polynucleotides (ZyGreen: Excitation at 503 nm and emission at 528 nm), which targets the ERBB2 gene and orange-labeled polynucleotides (ZyOrange: Excitation

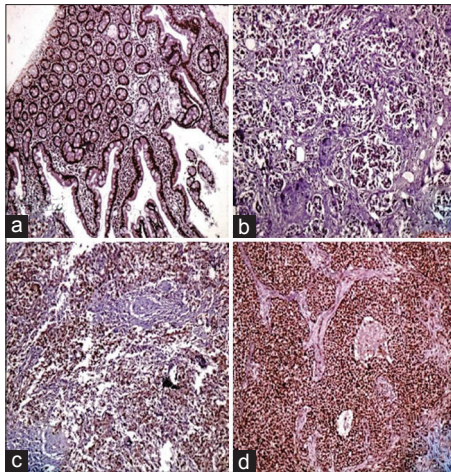


Figure 1: (a) Human duodenal biopsy (control) showing nuclear and cytoplasmic expression of Y-Box binding protein 1 antibody, ×100; (b) negative nuclear Y-Box binding protein 1 expression; score 3, ×100; (c) moderate positivity of nuclear Y-Box binding protein 1 expression; score 9, ×100; (d) strong positivity of nuclear Y-Box binding protein 1 expression; score 12, ×100

Table 1: Estrogen receptor, progesterone receptor, and Her2 scoring

	ER	PR	Her2
Negative	0	0	No staining or membrane staining which is incomplete and is faint/barely perceptible and within ≤10% of tumor cells
1+	<10% cells show positive staining	<10%	Incomplete membrane staining that is faint/barely perceptible and within >10% of tumor cells
2+	10%-75%	10%-75%	Circumferential membrane staining that is incomplete and/or weak/moderate and within >10% of tumor cells OR complete and circumferential membrane staining that is intense and within ≤10% of tumor cells
3+	>75%	>75%	Circumferential membrane staining that is complete, intense, and within >10% of tumor cells

ER = Estrogen receptor, PR = Progesterone receptor

Table 2: Y-Box binding protein 1 immuno-reactive scoring

Percentage of positive cells	Intensity of reaction
≤25% positive cells - 1	Negative staining - 0
26%-50% positive cells - 2	Weak staining - 1
51%-75% positive cells - 3	Moderate staining - 2
≥75% positive cells - 4	Strong staining - 3

at 547 nm and emission at 572 nm), which targets alpha-satellite sequences of the centromere of chromosome 17. Duplex formation of the fluorescence-labeled probes was visualized using Nikon Eclipse 80i fluorescence microscope (Nikon Corporation) employing red and

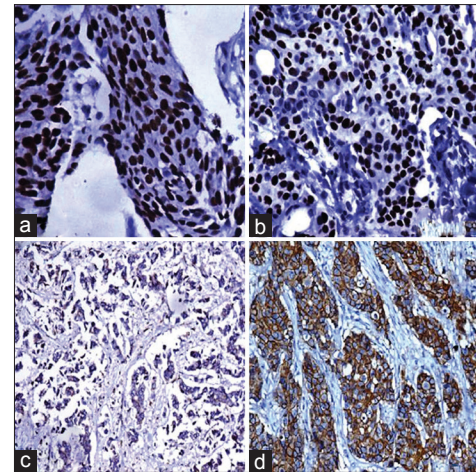


Figure 2: (a) Positive estrogen receptor expression (3+), ×400; (b) positive progesterone receptor expression (3+), ×400; (c) negative Her2 expression, ×100; (d) positive Her2 expression (3+), ×200

green filters. In interphases of normal cells or cells without aberrations of chromosome 17, two ERBB2 signals and two chromosome 17 signals appeared and were taken as normal.

Results

Y-Box binding protein 1 immunoexpression

YB1 expression was studied by IHC. The expression levels were then correlated with clinicomorphological parameters including age, side, menopausal status, grade, stage, lymph node status, presence/absence of *in situ* component, and hormone receptor status [Table 3].

In our study, YB1 showed cytoplasmic positivity in all the cases, whereas nuclear expression of YB1 was found to be positive in 48.6% cases and had a significant association with Her2 positivity and ER and PR negativity [Table 4]. For the purpose of our study, only nuclear expression was taken as positive. However, no association was found with tumor grade and stage. Lymph node metastasis had significant positive correlation with YB1 expression [Table 3].

Her2 gene amplification analysis by fluorescent *in situ* hybridization

Analysis of Her2 amplification by FISH was done on TMA slides. Her2 gene amplification was found in all three IHC equivocal (score 2+) cases. The cells having Her2/CEP17 ratio ≥2 or Her2/CEP17 ratio <2 with average Her2 copy number ≥6 signals/cell were taken as positive (as per ASCO/CAP guidelines) [Figure 3a and b].

Discussion

YB1 protein is a multifunctional protein having a role in transcription, translation, and DNA repair.

Table 3: Distribution of the clinicopathological parameters included in the study

Clinicopathological parameter	Percentage of cases		Association with YB1 expression (P)
	Present	Absent	
Histological type			
IDC-NOS	70/74 (95)		0.164
Invasive mucinous carcinoma	3		
IDC with medullary features	1		
Invasive lobular carcinoma	1		
<i>In situ</i> component	54/74 (72.3)	20/74 (27.7)	1
Tumor grade**			
Grade 1	13/65 (20)		0.333
Grade 2	44/65 (67.7)		
Grade 3	8/65 (12.3)		
Pathological (T) staging			
T1	3		0.28
T2	69		
T3	19		
T4	9		
Lymph node metastasis#	42/72 (58.33)	30/72 (41.67)	0.02
Molecular subtypes			
Luminal A	40		
Luminal B	19		
Her2 positive	15		
Triple negative	26		

**9/74 cases were postchemotherapy, hence not graded, #Data regarding lymph node status of two cases were not available. IDC-NOS = Invasive ductal carcinoma-not otherwise specified, YB1 = Y-Box binding protein 1

Table 4: Distribution of Y-Box binding protein 1 according to Her2, estrogen receptor, and progesterone receptor expression among study cases

	YB1 positive (n=36), n (%)	YB1 negative (n=38), n (%)	P
Her2 positive (n=25)	19 (76)	6	0.0012
Her2 negative (n=49)	17	32 (65.3)	
ER positive (n=31)	8 (25.8)	23 (74.2)	0.001
ER negative (n=43)	28 (65)	15 (35)	
PR positive (n=27)	5 (18.5)	22 (81.5)	0.0001
PR negative (n=47)	31 (66)	16 (34)	

ER = Estrogen receptor, PR = Progesterone receptor, YB1 = Y-Box binding protein 1

An increased YB1 expression was demonstrated in not only breast cancer but also various other tumors that include osteosarcoma, prostate cancer, glioblastoma, and ovarian malignancies.^[22] It induces cell proliferation, apoptosis, and control of tumor response to chemotherapy.^[22]

Previous studies on YB1 expression in breast cancer have supported its utility as an independent prognostic and predictive marker in breast cancer management.

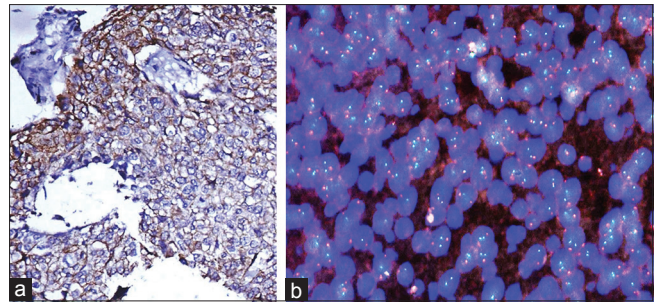


Figure 3: (a) A case equivocal on immunohistochemistry for Her2 (2+); (b) photomicrograph showing Her2 gene amplification by fluorescent *in situ* hybridization, with the cells having Her2/CEP17 ratio ≥2

Her2 in breast cancer is overexpressed either due to gene amplification or via transcriptional activation through increased tyrosine kinase activity and phosphorylation of itself and cellular substrates. Her2 and Her3 coexpression activates the downstream PI3k/Akt pathway which is considered to be a tumorigenic function of overexpressed Her2.^[23] In our study, Her-2 was positive in 33.7% of cases. Studies have reported a wider range of positivity from 10% to 34%.^[24] In our study, nuclear expression of YB1 was found to be positive in 48.6% of cases and had a significant association with Her2 positivity. However, no association was found with tumor grade, stage, histological subtype, presence of *in situ* component, or other clinical parameters such as age or menopausal status.

Studies have reported YB1 overexpression in 36%–41% of human breast cancers, while 80% of primary breast tumors showed increased copy numbers of chromosome 1.^[5,12] Wang *et al.* conducted their study on 1094 breast cancer patients and found YB1 overexpression in 398/1094 (36.38%) of cases.^[13] Bergmann *et al.* constructed TMA representing 4049 cases of invasive breast cancer with 20 years of follow-up and found YB1 expression in 41% of cases.^[8] In a similar study in 239 breast cancer cases, nuclear YB1 expression was found to be positive in 36.8% of cases.^[25]

A significant association was found between YB1 expression and Her2 receptor overexpression in our study ($P = 0.0012$). Wang *et al.* also found a significant association between YB1 expression and Her2 positivity.^[13]

Few other studies have suggested that YB1 acts as a promoter of Her2, and its expression leads to the clonal selection and expansion of Her2-positive cells.^[25,26] Similar results were obtained in another study, which reported that YB1 directly binds to the Her2 promoter in cells where Her2 gene was amplified. It thus allowed the cells with Her2 amplification to slip through the cell cycle check points and promote tumor growth.^[6] Her2

expression can thus be downregulated by silencing YB1 using SiRNA to knock down its expression or by mutating YB1 (Ser¹⁰²).^[6] Patients with Her2 receptor positivity and hormone receptor negativity have been found to have an unfavorable prognosis and higher risk of recurrence. YB1 expression acting via Her2-Akt-dependent pathway might be responsible for this poor prognosis.^[27] Hence, it has been suggested that the YB1-Her2 axis can be very important for personalized therapeutics in breast cancer by YB1-targeted drugs. Bergmann *et al.* reported that YB1 was expressed in 65% of Her2-positive breast cancer. In our study, we found YB1 to be positive in 76% of Her2-positive cases.^[8]

Lymph node status and tumor stage are important prognostic factors in breast cancer management. In our study, we found significant association between YB1 expression and lymph node status, suggesting the role of YB1 in prognostication.^[22]

In another study, YB1 was expressed in 30% of node-negative cases and based on the results, the authors suggested that patients with node-positive or node-negative cancer were more likely to die from the disease if YB1 was expressed.^[8] Xie *et al.* also found a significant positive correlation of YB1 with tumor grade and Her2 expression and a negative correlation with ER expression.^[25]

Our study showed both cytoplasmic and nuclear expression of YB1. For the purpose of analysis, only the nuclear expression of YB1 was considered positive. Studies in the past have also demonstrated both cytoplasmic and nuclear YB1 expression.^[28] Other workers have also found cytoplasmic YB1 expression in 100% of their cases.^[25] However, nuclear expression of YB1 was found only in 48.6% of our study cases. Different YB1 antibodies show different staining patterns that are determined by the accessibility of epitopes, and this depends on the nature of YB1 complexes. Hence, it is important to standardize the protocols and validate the staining with transcription of the protein if YB1 is to be used reproducibly as a prognostic guide for different cancers.^[11]

Studies have found YB1 expression in 70%–75% of triple-negative breast cancer (TNBC) and have found YB1 to be crucial in the genesis and maintenance of TNBC.^[25] In our study, TNBC constituted 25.6% of all cases and YB1 expression was seen in 73.6% of these cases. Targeting YB1 in TNBC may be a useful therapeutic option. Post chemoresponse or survival data were not available for the cases, and further studies with larger sample size and adequate follow-up data would strengthen the role of YB1 in the prognostication of Indian breast cancer patients.

Limitations of the study

This study could only study YB1 in relation to breast carcinoma though it has been associated with many other carcinomas. The other limitation was that follow-up of the patients was not available to validate the prognostic role of YB1 in breast carcinoma patients.

Conclusion

This study found that YB1 has an association with Her2neu expression in breast carcinoma. It may in future provide a therapeutic target in Her2neu overexpressing tumors.

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Conflicts of interest

There are no conflicts of interest.

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