

ER α and ER β co-expression: An indicator of aggressive tumors and hormonal sensitivity

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Abstract. The estrogen receptors (ERs) ER α and ER β are important factors in breast cancer progression. Nevertheless, the molecular interplay between ER α and ER β and its clinical significance in breast cancer is controversial. The establishment of a clear association is required; therefore, the current study analyzed the expression patterns of ER α and ER β in 32 breast tumor tissues using reverse transcription-quantitative polymerase chain reaction. Furthermore, human epidermal growth factor receptor 2 (HER2) and the Ki-67 status were detected by immunohistochemistry. The results revealed that the ER α and ER β expression rates recorded were 68 and 65%, respectively. The ER α :ER β ratio exhibited a decline along with disease progression. ER α and ER β were found to be negatively correlated with HER2 status but positively correlated with Ki-67. Co-expression of ER α and ER β was associated with breast cancer aggressiveness, including higher histological grade and positive nodal status, which commonly occur following the menopause. In addition, in cases where ER β was coexpressed with ER α , HER2 expression was frequently found to be negative, whereas the Ki-67 index was upregulated. These data suggest that ER α and ER β co-expression may be an indicator of tumor aggressiveness and the sensitivity of hormonal therapy via the downregulation of HER2.

Introduction

Estrogen regulates numerous physiological processes including normal cell growth, the central nervous and skeletal systems

and the development and regulation of tissue-specific genes in the genital tract (1-3). Estrogen also affects the pathological process of numerous hormone-dependent diseases including breast, endometrial and ovarian cancer (1). The biological actions of estrogen are mediated by the binding to one of two specific estrogen receptors (ERs), ER α or ER β , which belong to the nuclear receptors superfamily (3). The binding of estrogen to its receptor leads to a conformational change in the structure of the ER and to the formation of estrogen receptor dimers that bind estrogen response elements as homo- or hetero-dimers within the regulatory sequences of estrogen-dependent genes (4). This conformational change, occurring as a result of ligand binding, facilitates the association of coactivator receptors and stabilizes the estrogen receptor complex with estrogen response elements (4). This promotes gene transcription and supports the stimulation of cell growth in various tissues (5-7). ER α was the first estrogen receptor to be isolated and cloned in 1980 from a cell line of human breast cancer (MCF7) (8,9). In the presence of estradiol (E2), this receptor may induce cell proliferation via the regulation of certain genes including Myc, cyclinD and Wnt11 (10). These genes have several effects, including interfering with cadherins, stimulating the cell cycle, promoting the transition from G1 to S phase and altering apoptosis (10,11). This leads to an increase in cell division, which may cause errors in replication and promote cancer development (11).

In the 1990s, ER β was the second estrogen receptor discovered and identified in the rat prostate and ovaries, encoded by 485 amino acids (1996) (12). In the same year, ER β was also isolated from human tissues, in this case encoded by 477 amino acids (13). The role of ER β in breast cancer remains to be established (14,15). The majority of previous studies have shown that ER β functions as a negative modulator of ER α and is associated with a good prognosis and prolonged disease-free survival (16,17).

Previous studies have revealed that the expression of ER β is correlated with a poor prognosis, including accelerated cell proliferation and distant metastasis (18,19). However, Speirs *et al* (20) have identified that co-expression of ER α and ER β is associated with high-grade tumors and metastases. In addition, Grober *et al* (21) demonstrated that ER β is able

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to interfere with ER α in the regulation of target genes. It is therefore necessary to understand the role of ER β in breast cancer and to elucidate the nature of its association with ER α . In this context, the present study was conducted to investigate the expression of estrogen receptors ER α and ER β in a series of breast cancer tumors. This was achieved by comparing the results with clinical and pathological parameters, as well as the expression of the oncoprotein human epidermal growth factor receptor 2 (HER2) and the proliferation index Ki-67. The objective of this study was also to analyze the ER α and ER β subgroups according to the aforementioned parameters.

Materials and methods

Patients and samples. The tissue of malignant mammary tumors was excised during tumorectomy from 32 females (mean age, 58.5 years; range, 32-85 years) was analyzed. Healthy tissues collected from patients during the tumorectomy were used as controls. The patients all had invasive ductal carcinoma and did not receive radiotherapy or chemotherapy prior to surgery. The samples were subject to a histological examination by a pathology specialist to determine the presence of malignant cells. Each diagnosed sample was divided into two portions: One portion was immediately processed for immunohistochemistry and the other portion was frozen and maintained at -80°C until RNA extraction. All pathological, clinical and personal data were anonymized and separated from any personal identifiers. All the procedures followed were examined and approved by the Saleh Azaiez Oncology Institute (Tunis, Tunisia).

Total RNA isolation and reverse transcription. Total RNA was extracted from breast specimens using a mechanical stirrer in the presence of a lysis buffer [4.5 M guanidine-HCl, 50 ml Tris-HCl, 30% Triton X-100 (w/v) pH 6.6 (25°C)], prior to the use of total RNA isolation and high pure RNA isolation kits (Roche Diagnostics, Basel, Switzerland). Equal amounts of total RNA (1 μ g) were reverse transcribed. cDNA synthesis was carried out using the PrimeScript™ 1st strand cDNA Synthesis kit (Takara Biotechnology Co., Ltd., Dalian, China).

Primers and quantitative polymerase chain reaction (qPCR). All PCR reactions were performed using an ABI Prism 7700 Sequence Detection system (Applied Biosystems; Thermo Fisher Scientific, Inc., Waltham, MA, USA). cDNA was amplified using the SYBR1-Green PCR Core Reagents kit (Applied Biosystems; Thermo Fisher Scientific, Inc.). The reaction mixes used for qPCR were as follows: 10 μ l SYBR-Green (Applied Biosystems; Thermo Fisher Scientific, Inc.), 6 μ l water; 1 μ l forward primer; 1 μ l reverse primer; and 2 μ l cDNA. The primers used to amplify ER α were as follows: forward, 5'-TGCCAAGGAGACTCGCTA-3'; reverse, 5'-TCAACATTCTCCCTCCTC-3'. For ER β , the primer sequences were forward, 5'-TGTTACGAAGTGGGAATGTGA-3' and reverse, 5'-TCTTGTTCTGGACAGGGATG-3' (40 cycles of: 94°C for 30 sec, 55°C for 30 sec, 72°C for 30 sec for both ER α and ER β). 18S was used as an endogenous control. The primer sequences used to amplify 18S were forward, 5'-GTAACCCGTGTAACCCCAT-3' and reverse, 5'-CCATCCAATCGGTAGTAGCG-3' (40 cycles of: 94°C for 30 sec, 56°C for 30 sec and

72°C for 30 sec). Relative mRNA levels were calculated based on Cq values and corrected for the 18S expression according to the equation $2^{-\Delta\Delta Cq}$ (22). Relative mRNA levels in the control tissue were equated to 1 and other values were expressed relative to this.

All primer pairs were initially validated by testing them for equal amplification efficiencies. The amplification efficiency was close to 2 under these conditions. Experiments were performed in triplicate for each data point.

Immunohistochemical staining. The immunohistochemistry expression of the oncoprotein Her2/neu and the proliferation index Ki-67 were tested on the same set of tumors. The primary antibodies used were as follows: Mouse anti-human Her2 (#CB11) and mouse anti-human Ki-67 (#MM1) (Novocastra; Leica Biosystems GmbH, Wetzlar, Germany).

A total of 32 tissue samples were fixed for 24 h at room temperature in 0.1 M phosphate buffered 10% formaldehyde, dehydrated in a graded ethanol series of increasing concentrations (70, 85, 90 and 100% for 10 min each), impregnated with xylene and embedded in paraffin. Sections (4 μ m thick) were processed following the NovoLink™ Polymer Detection systems (Novocastra; Leica Biosystems GmbH) method. Sections were deparaffinized by an overnight incubation at 59°C, and subsequently placed in a xylene bath for 15 min at room temperature. Sections were subsequently hydrated, incubated for 30 min in 1% hydrogen peroxide to block endogenous activity, and then antigen retrieval was performed by incubating the sections in a 0.01 M citrate buffer (Epitope Retrieval Solution pH 6.0; Leica Microsystems GmbH, Wetzlar, Germany) for 30 min at 98°C. Subsequently, the primary antibodies were applied for 1 h at 4°C, with a dilution of 1:40 for Her2 and 1:200 for Ki-67. The sections were then incubated at room temperature with Post Primary Block for 30 min to block non-specific polymer binding. The sections were incubated with a NovoLink™ Polymer for 30 min at room temperature, followed by incubations with 3,3'-diaminobenzidine (DAB) working solution for 5 min at room temperature to develop peroxidase activity. The slides were counterstained with hematoxylin and mounted. Staining specificity was checked using negative controls. Negative controls were obtained by replacing the primary antibody with an antibody of the same unrelated isotope during the immunohistochemistry technique or by omission of the primary antibody. Primary breast tissues were incubated in blocking peptides (Santa Cruz Biotechnology, Inc., Dallas, TX, USA) instead of primary antibodies.

The Ki-67 assessment and the Her2 status tests were performed by two experienced breast pathologists. The percentage of positively stained cells obtained is an average following counting of the stained cells and the total number of cells were counted in four high-magnification fields using a light microscope (magnification, x400; Media Cybernetics, Inc., Rockville, MD, USA). The staining of Ki-67 was scored for the percentage of positive cells (0, 0-5%; 1, 6-25%; 2, 26-50%; 3, >50%). The optimal cutoff value was identified as 1 for the low Ki-67 expression level and >1 for high expression (23). The scoring of Her2 was performed on a 0-3 scale (24). Positive (3+) was defined as intense complete membranous staining in >30% of the tumor cell population; borderline (2+) was defined as moderate membranous

Table I. Association between mRNA ER α and ER β levels and standard clinicopathological factors and molecular settings.

Characteristics	Total population	ER α		P-value	ER β		P-value
		Negative (%)	Positive (%)		Negative (%)	Positive (%)	
Total	32 (100%)	10 (31.28)	22 (68.75)		11 (34.37)	21 (65.62)	>0.05
Age							
≤50 years	12 (37.5)	4 (40)	8 (36.36)		5 (45.45)	7 (33.33)	
>50 years	20 (62.5)	6 (60)	14 (63.63)	0.13	6 (54.54)	14 (66.66)	0.06
Grade							
SBRI	7 (21.87)	4 (40)	3 (13.63)		4 (36.36)	3 (14.28)	
SBRII	17 (53.12)	3 (30)	14 (63.63)		5 (45.45)	12 (57.1)	
SBRIII	8 (25)	3 (30)	5 (22.72)	0.01	2 (18.18)	6 (28.57)	0.01
Lymph node status							
Positive	17 (53.12)	6 (60)	11 (50)		4 (36.36)	13 (61.90)	
Negative	15 (46.87)	4 (40)	11 (50)	>0.05	7 (63.63)	8 (38.09)	0.20
Tumor size							
≤30 mm	22 (68.75)	6 (60)	16 (72.72)		8 (72.72)	14 (66.66)	
>30 mm	10 (31.25)	4 (40)	6 (27.27)	0.006	3 (27.27)	7 (33.33)	0.06
HER2 status							
Positive	8 (25)	3 (30)	5 (22.72)		6 (72.72)	2 (57.14)	
Negative	24 (75)	7 (70)	17 (77.27)	0.0009	5 (27.27)	19 (42.85)	7.905 ⁻⁷
Ki-67							
≤5%	9 (28.12)	3 (44)	6 (27.27)		3 (27.27)	6 (28.57)	
>5%	23 (71.87)	7 (56)	16 (72.72)	0.0066	8 (72.72)	15 (71.42)	0.01

P-values calculated using the χ^2 test in R. SBR, Scarff-Bloom-Richardson; HER2, human epidermal growth factor receptor 2; ER, estrogen receptor.

staining in >10% of tumor cells; 1+ was defined as either weak or barely perceptible membranous staining in >10% of the tumor cells. Furthermore, a chromogenic *in situ* hybridization (CISH) analysis was performed, as described previously (25) for Her2/neu gene amplification in all 2+ cases, as defined by IHC. Scores of 0, 1+ and 2+ according to IHC but negative following CISH were considered as negative for the Her2/neu expression, whereas 3+ scores and 2+ cases defined as positive by CISH were considered as positive for Her2/neu expression.

Statistical analysis. The analysis of the results involving the expression of the estrogen receptors ER α and ER β together with the various histological and clinical parameters was performed using the χ^2 test with R (i386 3.2.1) software. Data are presented as the mean \pm standard deviation. P<0.05 was considered to indicate a statistically significant difference.

Results

Expression of mRNA ER α and ER β in malignant human breast tissues and their association with clinical parameters. In total, 32 breast tumor samples were analyzed and compared for the expression of the two ER isoforms by reverse transcription-qPCR. The expression of estrogen receptors in breast cancer tumors had a positivity of ~68% for ER α and 65% for ER β . As presented in Table I, no significant difference

was identified between the levels of expression of the two ER isoforms (P>0.05).

In order to determine whether the expression of each ER isoform in the breast tumors was associated with clinical parameters, associations between tumor grade and size, lymph node metastasis and menopausal status were examined (Table I). When compared with tumor grade, there was a significant association between ER α and ER β expression and tumor grade (P=0.01; Table I). Analysis of the mRNA expression of the receptors ER α and ER β revealed a significant increase of ER α in tumors that were grade 2 compared with grades 1 and 3. However, the highest ER β expression was found in the Scarff-Bloom-Richardson (SBR)3 grade (Fig. 1). According to the histopathological grade progress, the ER α :ER β ratio declined from SBR1 to SBR2 and SBR3 (Fig. 2).

Tumor size was correlated with ER α (P=0.006) expression and did not correlate with ER β expression (P=0.06; Table I). In addition, there was no significant difference in ER α and ER β expression according to lymph node metastasis status (P>0.05; Table I). As presented in Fig. 3, the expression of the two ER isoforms differed with menopausal status. The amount of mRNA ER α expression was 1.5X higher in premenopausal breast tumors compared with postmenopausal breast tumors. Inversely, mRNA expression of ER β increased between the premenopausal and postmenopausal status (Fig. 3). Furthermore, a shift in the ER α :ER β ratio was noted and this was

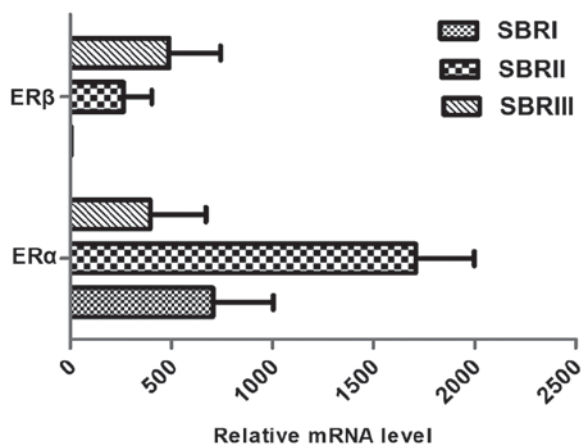


Figure 1. ER α and ER β mRNA expression level according to SBR grade. ER, estrogen receptor; SBR, Scarff-Bloom-Richardson.



Figure 2. ER α :ER β ratio according to SBR grade. ER, estrogen receptor; SBR, Scarff-Bloom-Richardson.

revealed to decline in the postmenopausal vs. premenopausal status group (7.66 to 3.36) (Fig. 4).

HER2/neu and Ki-67 in ER-positive vs. ER-negative cases. Differences in the HER2 and Ki-67 status between ER-positive and ER-negative cases were further analyzed. Immunohistochemical analysis was performed to assess the expression of HER2 and Ki-67 in breast cancer samples. Fig. 5 reveals membrane localization of the HER2 protein in malignant breast cells. Ki-67 was primarily localized to the nucleus of breast neoplastic cells (Fig. 5). As summarized in Table I, a marked negative association between the two estrogen receptors and the oncoprotein HER2/neu was observed. In total, 17/22 ER α positive cases were negative for HER2. Similarly, 19/21 ER β positive cases were negative for HER2. This negative association was more significant for ER β ($P=7.905 \times 10^{-7}$) than ER α ($P=0.0009$; Table I).

For Ki-67, there was a discrepancy in the prognostic importance of this factor between the ER-positive and ER-negative cases. A high Ki-67 index of $\geq 5\%$ was associated with the ER-positive subgroup ($P=0.006$ for ER α and $P=0.01$ for ER β ; Table I).

Association between ER α and ER β breast cancer subgroups and clinical information. The expression of ER subtypes within

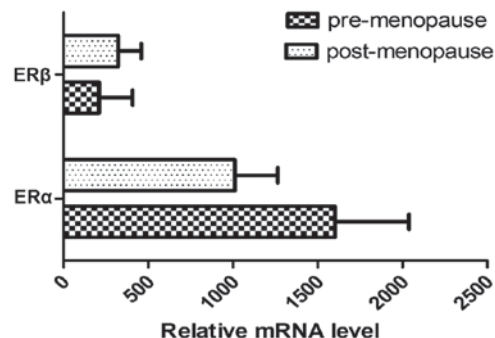


Figure 3. mRNA expression level of ER α and ER β according to menopausal status. ER, estrogen receptor; SBR, Scarff-Bloom-Richardson.

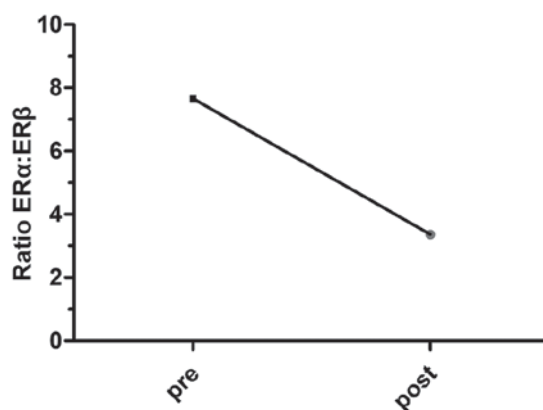


Figure 4. ER α :ER β ratio according to menopausal status. ER, estrogen receptor; pre, premenopausal; post, postmenopausal.

the tumor group was further analyzed. According to positive or negative expression of the hormone receptors ER α and ER β , ER α + and ER β + was significantly the most expressed ER subgroup in patients with breast cancer ($P=0.01$; Fig. 6). A total of 43.75% of the malignant breast samples co-expressed the two ER subtypes, compared with only 25 and 21.87% of the breast tumors which expressed either (ER α +, ER β - or ER α -, ER β +, respectively). A small number of breast cancer samples were in the ER α -, ER β - subgroup (9.37%; Fig. 6).

The distribution of ER α and ER β expression groups according to the menopausal status exhibited a significant difference for the ER α +, ER β + subgroup ($P=0.05$). As presented in Table II, 71.42% of the ER α and ER β co-expression cases occurred following the menopause stage, whereas only 28.57% occur prior to menopause. The other ER subgroups studied were distributed approximately homogeneously prior to and following the menopause phase. When compared with tumor grade, there was a significant association between the ER α +, ER β + subgroup and SBR grade ($P=0.005$). This was not observed for the other ER subgroups, but there was a non-significant association between the ER α +, ER β -subgroups and the primary stages of cancer ($P=0.08$). Similarly, depending on the nodal status, there was an association between the ER α +, ER β + subgroup with the infiltration of the lymphatic ganglion ($P=0.05$; Table II).

Subset analyses of HER2/neu and Ki-67 in ER α , ER β breast cancer subgroups. As presented in Table II, the complex

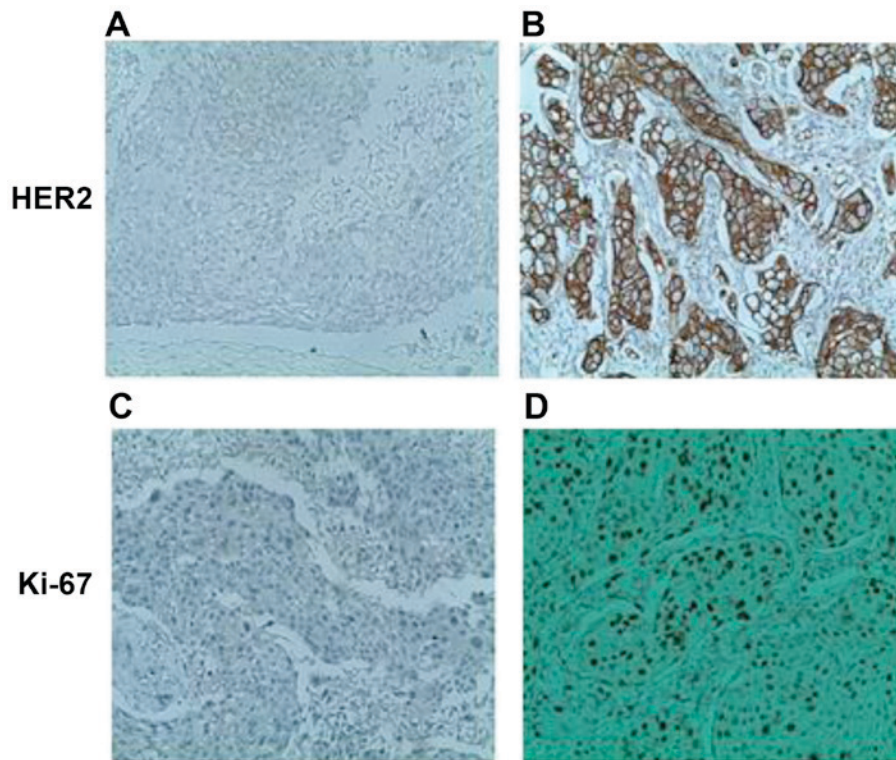


Figure 5. Immunohistochemical staining with anti-HER2 and anti-Ki-67 antibodies in breast tumors. (A) Absence of overexpression of HER2 oncoprotein (HER2 score, 0; original magnification, x250). (B) Presence of overexpression of HER2 oncoprotein (HER2 score, 3; original magnification, x400). (C) Proliferation index estimated at 2% and (D) Ki-67 proliferation index estimated at 70% (original magnification, x250).

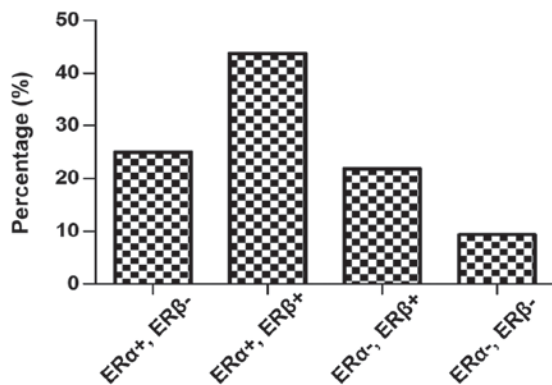


Figure 6. ERα and ERβ breast cancer subgroups. ER, estrogen receptor.

associations between various pathological variables were considered in the current study. Using this analysis, it is possible to visualize the association of biological factors (ERα, ERβ, HER2 and Ki-67) with ER subgroups and study their associations with conventional pathological factors. Breast tumors with an ERα+, ERβ+ profile were characterized by negative HER2 status. A statistically significant relationship between the co-expression of estrogen receptors and HER2 negative was identified in the tumor samples ($P=3.21 \times 10^{-5}$). A statistically significant negative association was also observed between the ERα-, ERβ+ profile and HER2 expression ($P=0.03$; Table II).

Using a Ki-67 cut-off value of 5% (Table II), only the ERα+, ERβ+ breast cancer subgroup was significantly associated with a Ki-67 index $>5\%$ ($P=0.008$).

Discussion

Previous studies have been conducted to decipher the role of ERs in breast carcinogenesis (16,18,19). Nonetheless, the role of the ERβ isoform in this malignancy and its association with ERα remains to be elucidated and the results of earlier studies are somewhat contradictory (26,27). The results of the current study indicate that ERα and ERβ expression was retained in a majority of breast cancer cases (68 and 65%, respectively). The associations between ERα and ERβ expression and menopausal status revealed that each isoform was more significantly expressed in postmenopausal patients than in premenopausal ones (Table I), primarily for ERβ which was ~2 times higher in post-menopausal cases ($P=0.06$). These data are concordant with previous studies demonstrating that the two ER isoforms are frequently positive in postmenopausal patients (28-30). However, considering the relative amount of mRNA, the ERα expression was found to be higher in premenopausal phases (1.5-fold) compared with postmenopausal breast tumors in the present study. Inversely, the mRNA expression level of ERβ was often higher in postmenopausal patients. Consequently, the ERα:ERβ ratio decreases from 7.66 to 3.36 (Fig. 4), which translates into the ERβ activity increasing in malignant breast tumors of post-menopausal patients. Furthermore, the expression levels of ERα and ERβ were revealed to be associated with smaller tumors and significantly associated with the SBR histopathological grade ($P=0.01$; Table I). In fact, ERβ was more highly expressed than ERα in the SBRIII grade. In addition, the ERα mRNA level reached a maximum in the SBRII grade and decreased in SBR grade III. However, the ERβ mRNA level was associated with the advancement of

Table II. Association between ER α and ER β breast cancer subgroups with clinicopathological and molecular parameters.

Clinicopathological and molecular parameters	ER α +, ER β - (n=8)	ER α +, ER β + (n=14)	ER α -, ER β + (n=7)	ER α -, ER β - (n=3)
	n (%)	n (%)	n (%)	n (%)
Menopausal status				
Premenopausal	4 (50)	4 (28.57)	3 (42.85)	1 (33.33)
Postmenopausal	4 (50)	10 (71.42)	4 (57.14)	2 (66.66)
P-value	>0.05	0.05	>0.05	>0.05
Histological grade				
SBRI	2 (25)	1 (7.14)	2 (28.57)	2 (66.66)
SBR II	5 (62.5)	9 (64.28)	3 (42.85)	0
SBR III	1 (12.5)	4 (28.57)	2 (28.57)	1 (33.33)
P-value	0.08	0.01	0.80	0.22
Lymph node status				
Positive	2 (25)	10 (71.42)	4 (57.14)	2 (66.66)
Negative	6 (75)	4 (28.57)	3 (42.85)	1 (33.33)
P-value	0.13	0.05	>0.05	>0.05
HER2/neu				
Positive	4 (50)	1 (7.14)	1 (14.28)	2 (66.66)
Negative	4 (50)	13 (92.85)	6 (85.71)	1 (33.33)
P-value	>0.05	3.22×10^{-5}	0.03	>0.05
Ki-67				
$\leq 5\%$	6 (75)	3 (21.42)	3 (42.85)	1 (33.33)
$> 5\%$	2 (25)	11 (78.57)	4 (57.14)	2 (66.66)
P-value	0.10	0.01	>0.05	>0.05

P-values calculated using the χ^2 test in R. SBR, Scarff-Bloom-Richardson; HER2, human epidermal growth factor receptor 2; ER, estrogen receptor.

the SBR grade. ER β expression reached its maximum in the SBR III grade and was higher than that of ER α . Consequently, the ER α :ER β ratio was inversely associated with SBR grade advancement, and the ratio tended to 0 at the SBR III grade (Fig. 2).

These results demonstrate a slight association between ER β expression (but not ER α) and node-positive breast cancer ($P=0.06$), which is corroborated by the findings of Hou *et al* (31), which demonstrated that ER β exerts stimulatory effects on breast cancer development and metastasis. Furthermore, Yan *et al* (23) revealed that the expression of ER β 2 is also correlated with high-grade tumors, distant metastasis and breast cancer mortality (23).

Estrogen-ER signaling is important in normal mammary gland development and breast carcinogenesis (32). Due to the crosstalk between the ER and epidermal growth factor receptor (EGFR)/HER2 signaling pathways and/or their downstream effectors, the majority of patients develop resistance to endocrine therapy (predominantly tamoxifen) (33). In fact, the relation of ER α and ER β with HER2 is of particular interest. In the current study, the HER2-negative status was associated with positive ER α and ER β expression. This result is concordant with previous data that demonstrated a significant negative correlation between the expression of the hormone receptor and HER2/neu amplification (24,34,35).

The combination of HER2 overexpression and a high Ki-67 index has been suggested to be a prognostic molecular marker for breast cancer (36). The present study revealed that positive ER receptors (α and β) were associated with a high Ki-67 index of $\geq 5\%$ ($P=0.006$ and $P=0.01$ respectively) for breast cancer (Table I). These results are concordant with previous reports (18,37), and imply that ER α and ER β may have a role in breast cancer development, metastasis and proliferation.

ER β may be expressed alone (ER α -, ER β +) or co-expressed with ER α (ER α +, ER β); therefore, a comparison of the previously cited clinical and molecular parameters referring to the (ER α , ER β) breast cancer subgroups is required. The present study identified that ER α and ER β co-expression was retained in the majority of breast cancer cases ($P=0.01$; Fig. 6), hypothesizing an interaction between the two nuclear receptors. In this context, Järvinen *et al* (38) and Leung *et al* (39) reported that positive ER β expression is associated with positive ER α in breast cancer tumors. Given that ER α and ER β were coexpressed in the majority of breast tumors, this expression may occur in the form of their heterodimers. ER α and ER β heterodimers may also serve a significant role in breast cancer, but this remains to be established (40).

The present study demonstrated that the co-expression of ER α and ER β occurs frequently following menopause, which may be associated with the *in situ* synthesis of steroid

hormones (the epithelium of the mammary gland) from the steroid precursors dehydroepiandrosterone (DHEA) and DHEA-sulfate. Through an intracrinology mechanism, E2 may be synthesized locally by the aromatization of the androgen by aromatase (41). Depending on the presence of ER α and ER β in certain cells, the receptors form functional homo- or heterodimers on promoter elements (40). The results of the current study reveal that the progression of breast cancer may be dependent on the expression levels of estrogenic receptors, particularly the co-expression of the ER α and ER β isoforms (ER α +, ER β + subgroup). Indeed, ER α and ER β co-expression is associated with high-grade tumors (grade II and III; $P=0.005$) and potentially lymph node infiltration ($P=0.05$). This finding is concordant with the findings of Speirs *et al* (20), which demonstrated that tumors that expressed ER α and ER β were node-positive and tended to be of a higher grade. This implies that ER α and ER β may cooperate to generate a tumor phenotype with a higher metastatic potential. The present study hypothesizes that there is a synergic effect between ER α and ER β , which exerts stimulative effects on breast cancer development and metastasis.

The potential stimulative effect exerted by the co-expression of the ER α and ER β receptors is supported by the significantly high proliferation rate estimated by the Ki-67 index in the (ER α +, ER β +) breast cancer subgroup. Notably, it has been reported that an increase in the proliferation rate occurs during the progression towards invasive ductal carcinoma, implying a significant role for Ki-67 in breast tumorigenesis (42). However, the (ER α +, ER β -) subgroup is associated with primary cancer grade (SBRI and II), non-infiltrated lymph nodes and a lower proliferation index (<5%). This reveals that the expression of ER α alone may be a marker of non-aggressive tumors, as has been suggested by a previous study (28). Our results demonstrate a negative association between the (ER α -, ER β +) subgroup and HER2. Such results are corroborated by a study by Marotti *et al* (34), which demonstrated that the co-expression of ER α and ER β was associated with a negative status of HER2. In addition, Lindberg *et al* (35) revealed that ER β is able to increase phosphatase and tensin homolog levels and decrease HER2/HER3 signaling, thereby reducing protein kinase B signaling. The co-expression of ER β and ER α (ER α +, ER β + subgroup) was negatively associated with HER2 expression in the present study. These findings are consistent with a previous report, which concluded that ER β is significantly associated with ER α expression and inversely associated with HER2 over-expression (34). As these two receptor systems (ER isoforms and HER2) have the capacity to activate one another (30), the present study hypothesizes a synergistic effect between ER α and ER β to abrogate HER2 activation as a part of the crosstalk between ER and growth factor receptors. In this context, ER-induced signaling pathways, demonstrated through *in vitro* cellular models, were found to induce EGFR ligands, such as transforming growth factor α (43) and lead to downregulation of EGFR (44) and HER2 (45). This growth factor receptor, HER2/neu, is not typically over-expressed in normal or benign breast lesions (46). However, a significantly lower level of HER2/neu expression in invasive carcinoma has been previously reported (47). The results of the current study imply that the molecular phenotype defined by the presence

of ER α , ER β and the absence of HER2 may be a precursor for the development of a more aggressive and malignant invasive ductal carcinoma. Furthermore, the coexpression of ER α and ER β may be a marker of hormonal sensitivity in association with downregulation of HER2 expression.

In conclusion, the results of the current study suggest an important role for ER β in breast cancer development, proliferation and metastasis, particularly when coexpressed with ER α . The ER α +, ER β + subgroup is associated with high tumor grade, metastasis and a high proliferation index. Therefore, the co-expression of ER α and ER β may be an indicator of aggressive tumors. In addition, the ER α +, ER β + subgroup is significantly associated with an HER2-negative status, and this may indicate a sensitivity towards hormonal therapy due to the downregulation of HER2 expression. Given that the ER α -ER β balance is influenced by the tumor microenvironment, including cytokines and growth factors, decrypting signaling pathways elicited by those components and their crosstalk with ER signaling may be investigated further in future studies.

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