

ORIGINAL ARTICLE

Revisiting the definition of estrogen receptor positivity in HER2-negative primary breast cancer

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Background: Although 1% has been used as cut-off for estrogen receptor (ER) positivity, several studies have reported that tumors with ER < 1% have characteristics similar to those with $1\% \le ER < 10\%$. We hypothesized that in patients with human epidermal growth factor 2 (HER2)-negative breast cancer, a cut-off of 10% is more useful than one of 1% in discriminating for both a better pathological complete response (pCR) rate to neoadjuvant chemotherapy and a better long-term outcome with adjuvant hormonal therapy. Our objectives were to identify a percentage of ER expression below which pCR was likely and to determine whether this cut-off value can identify patients who would benefit from adjuvant hormonal therapy.

Patients and methods: Patients with stage II or III HER2-negative primary breast cancer who received neoadjuvant chemotherapy followed by definitive surgery between June 1982 and June 2013 were included. Logistic regression models were used to assess the association between each variable and pCR. Cox models were used to analyze time to recurrence and overall survival. The recursive partitioning and regression trees method was used to calculate the cut-off value of ER expression.

Results: A total of 3055 patients were analyzed. Low percentage of ER was significantly associated with high pCR rate (OR = 0.99, 95% CI = 0.986–0.994, P < 0.001). The recommended cut-off of ER expression below which pCR was likely was 9.5%. Among patients with ER \ge 10% tumors, but not those with 1% \le ER < 10% tumors, adjuvant hormonal therapy was significantly associated with long time to recurrence (HR = 0.24, 95% CI = 0.16–0.36, P < 0.001) and overall survival (HR = 0.32, 95% CI = 0.2–0.5, P < 0.001).

Conclusion: Stage II or III HER2-negative primary breast cancer with ER < 10% behaves clinically like triple-negative breast cancer in terms of pCR and survival outcomes and patients with such tumors may have a limited benefit from adjuvant hormonal therapy. It may be more clinically relevant to define triple-negative breast cancer as HER2-negative breast cancer with <10%, rather than <1%, of ER and/or progesterone receptor expression.

Key words: breast cancer, ER positivity, triple-negative breast cancer, adjuvant hormonal therapy

Introduction

Approximately 12%–17% of breast cancers are triple-negative breast cancer (TNBC), which is defined as estrogen receptor (ER)-negative, progesterone receptor (PR)-negative, and human epidermal growth factor 2 (HER2)-negative disease [1]. With contemporary treatments, patients with TNBC generally have a better overall response to chemotherapy but—particularly among those with chemotherapy-insensitive disease—a poorer

prognosis than do patients with other breast cancer subtypes, such as breast cancers that are ER- and/or HER2-positive [1–6]. This suggests that TNBC is a heterogeneous disease and that patients with TNBC have chemotherapy responses and survival outcomes that differ from those of patients with other breast cancer subtypes. Compared with TNBC patients who do not achieve pathological complete response (pCR) to neoadjuvant chemotherapy (NACT), TNBC patients who do achieve pCR have a

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better prognosis; thus, pCR is a surrogate survival marker in this population [2, 6–10].

According to the American Society of Clinical Oncology/ College of American Pathologists (ASCO/CAP), breast cancers with <1% of ER or PR expression should be considered hormone receptor-negative tumors [11]. However, we found in a previous retrospective study that breast cancers with 1%-9% of ER expression had gene expression profiles similar to those of breast cancers with <1% of ER expression [12]. Another retrospective study including both HER2-negative and HER2-positive cases showed that the recurrence-free survival (RFS) and overall survival (OS) rates of patients with 1%-9% of ER expression and those of patients with <1% of ER expression did not differ significantly (P = 0.96 and 0.10, respectively) and that both groups' RFS and OS rates were significantly worse than those of patients with \geq 10% of ER expression (P < 0.0001 for both) [13]. These results suggest that breast cancers with >1% but <10% of ER expression and those with $\leq 1\%$ of ER expression have similar molecular features and clinical prognoses. Another retrospective study suggested that low percentage of ER expression as a continuous variable is associated with a high pCR rate [14]. However, those studies included both HER2-negative and HER2-positive diseases. In addition, no previous study has determined the optimal cut-off value of the ER expression level as a continuous variable in terms of the association with pCR rate or the difference in the survival benefit from adjuvant hormonal therapy based on the calculated cut-off value of ER. Some studies have reported that hormone receptor-positive tumors are less sensitive to systemic chemotherapies than hormone receptornegative tumors are, which suggests that the ER expression level is associated with sensitivity to NACT [15-17]. More recently, the St. Gallen International Expert Consensus 2015 reported that ER expression values between 1% and 9% were considered equivocal and that endocrine therapy alone cannot be relied upon for patients with these values [18]. Thus, whether the percentage of ER expression as a continuous variable affects the pCR rate after NACT in patients with HER2-negative primary breast cancer is unknown and the exact clinical definition of TNBC with consideration of the survival benefit from adjuvant hormonal therapy has not been fully investigated.

We hypothesized that in patients with HER2-negative primary breast cancer, a cut-off ER expression level of 10% is more useful than one of 1% for discriminating patients who are likely to have a better pCR rate to NACT and those who are likely to have a better long-term outcome with adjuvant hormonal therapy. In this retrospective chart review study, our primary objective was to identify the percentage of ER expression below which pCR was likely in patients with newly diagnosed stage II or III HER2-negative primary invasive breast cancer treated with NACT. Our secondary objective was to determine whether this cut-off value can identify patients who would benefit from adjuvant hormonal therapy.

Methods

Study population

This retrospective chart review study was approved by The University of Texas MD Anderson Cancer Center's Institutional Review Board

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(protocol number: PA14-0046), and a waiver of informed consent was granted based on the study's retrospective nature. We reviewed the Breast Medical Oncology management system database (protocol number: 2004-0541) to identify patients with newly diagnosed stage II or III HER2-negative primary invasive breast cancer who received NACT followed by definitive surgery between June 1982 and June 2013 at MD Anderson Cancer Center. We included only patients for whom the percentages of ER and PR expression levels were available.

Data collection

From the database, we extracted age, race, menopausal status, body mass index, clinical stage, percentage of ER, percentage of PR, histology, nuclear grade, NACT regimens [anthracycline alone (A), taxane alone (T), or anthracycline plus taxane (A + T)], treatment response (pCR or nonpCR), lymphovascular invasion (positive or negative), adjuvant hormonal therapy (yes or no), adjuvant chemotherapy (yes or no), and adjuvant radiation therapy (yes or no). Stage was assessed using American Joint Committee on Cancer (AJCC) classification. HER2 positivity was defined as a HER2/CEP17 fluorescence *in situ* hybridization (FISH) ratio of \geq 2.0 and/or an immunohistochemical (IHC) staining score of 3+ [19]. pCR was defined as no invasive carcinoma in the breast or no tumor in the axilla at the time of surgery.

Data were initially extracted from electronic medical records and entered into a prospectively maintained database. The dataset was assessed and cleaned by TF and TK independently. Follow-up information for patients in the Breast Medical Oncology management system database is obtained every 2 years by direct review of the medical records and linkage to the MD Anderson Tumor Registry, which mails annual followup letters to each patient registered at MD Anderson to confirm that the patient is alive and free of cancer. The MD Anderson Tumor Registry also checks the Social Security Death Index and the Texas Bureau of Vital Statistics for the status of patients who do not respond to the letters.

Pathological evaluation

For IHC staining, the 1D5 antibody was used until 2000. Since then, the 6F11 antibody has been used for IHC staining of formalin-fixed paraffinembedded (FFPE) tissue sections in our clinical IHC laboratory, which is certified under the provisions of the United States Clinical Laboratory Improvement Act (CLIA) and accredited by the CAP. The IHC protocol, whose use with 4-µm FFPE tissue sections, has been validated and includes a de-paraffinization step for 30 min at 72 °C and a rehydration with antigen retrieval carried out at 100 °C for 20 min with Citrate buffer. In the protocol, endogenous peroxidase is blocked with 3% peroxide for 5 min. The primary antibody is applied at a 1 : 35 dilution for 15 min. Post-primary antibody detection is carried out using a commercial polymer system (Bond Polymer Refine Detection, Leica), and stain development is achieved by incubation with DAB and DAB Enhancer (Leica). Hematoxylin is applied as a counterstain. A positive control (cervical tissue) is added to every slide. At MD Anderson, a multidisciplinary approach to grossing breast specimens, used for more than two decades, involves the use of whole and sectioned specimen radiography, the correlation of gross findings with imaging findings, and discussion among the pathologist, radiologist, and surgeon. For specimens obtained after neoadjuvant therapy, we follow protocol for the examination of specimens from patients with invasive carcinoma of the breast from College of American Pathologists (CAP) [20] and an international working group for standardization of specimen handing and reporting for breast specimens after neoadjuvant therapy [21].

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confirm that the patient is alive and free of cancer. The MD Anderson Tumor Registry also checks the Social Security Death Index and the Texas Bureau of Vital Statistics for the status of patients who do not respond to the letters.

Statistical analysis

Standard descriptive statistics and frequency tabulation were used to summarize data. The chi-square test and Fisher's exact test were used to evaluate the association between two categorical variables. The Kruskal–Wallis test was used to compare the distributions of continuous variables among different groups. Univariate and multivariate logistic regression models were used to investigate the association between each variable and pCR. The variables with *P* values of ≤ 0.2 in the univariate analysis were included in the selection of the full multivariate model. We obtained the reduced multivariate model using a backward selection approach, removing the least significant covariates from the full model one at a time, and *P* values of < 0.05 were used as the limit for inclusion in this analysis. ER and PR were treated as continuous and categorical variables separately in the multivariate analyses.

Kaplan-Meier curves for patients sorted by prognostic factors of interest were produced, and the log-rank test was used to assess the difference between the prognostic factor groups. OS was defined as the time from surgery to death or last follow-up; patients who were alive at the end of the study period were censored at the date of last follow-up. Time to recurrence (TTR) was defined as the time from surgery to recurrence or breast cancerspecific death; patients whose disease had not recurred or who had not died from breast cancer were censored at the date of last follow-up. Univariate and multicovariate Cox proportional hazard models were used to determine the effects of prognostic factors on survival distributions. The recursive partitioning and regression trees method [22] was used to select a percentage of ER expression below which pCR was likely. The package was downloaded via the Comprehensive R ArchiveNetwork (CRAN, https://cran.r-project. org/web/packages/rpart/). All tests were two-sided. P values <0.05 were considered statistically significant. All analyses were conducted using SAS 9.3 (SAS, Cary, NC), S-Plus 8.0 (TIBCO Software Inc., Palo Alto, CA), and R 2.14.2.

Results

Patients

A total of 3055 patients were included in the analysis (supplementary Figure S1, available at *Annals of Oncology* online). Among these 3055 patients, 1726 (56.5%) had stage II disease and 1329 (43.5%) had stage III disease; 932 (30.5%) had tumors with <1% of ER expression (ER < 1% tumors), 171 (5.6%) had tumors with \geq 1% but <10% of ER expression (1% \leq ER < 10% tumors), and 1952 (63.9%) had tumors with \geq 10% of ER expression (ER \geq 10% tumors). Most patients (2577; 84.4%) had ductal carcinoma, and most patients (2651; 86.8%) received A + T as NACT. Of the 171 patients with 1% \leq ER < 10% tumors, 43 (25.1%) had adjuvant hormonal therapy, and of the 1952 patients with ER \geq 10% tumors, 1906 (97.6%) had adjuvant hormonal therapy.

The patients' demographic and clinicopathological characteristics by ER expression level are given in Table 1. Compared with patients with ER < 1% or $1 \le \text{ER} < 10\%$ tumors, those with ER $\ge 10\%$ tumors were more likely to be white (P < 0.001), have nuclear grade I or II (P < 0.001), have non-lobular disease (P < 0.001), have lymphovascular invasion (P = 0.005), and have received adjuvant radiation (P < 0.001).

Univariate logistic regression analysis of pCR

In the univariate logistic regression analysis, high percentages of ER and PR expression as continuous variables were significantly associated with low pCR rates [ER: odds ratio (OR), 0.98; 95% confidence interval (CI), 0.978–0.983; P < 0.001; PR: OR, 0.976; 95% CI, 0.971–0.98; P < 0.001]. Consistent with this result, low ER and PR expression levels as categorical variables were significantly associated with high pCR rates. Ductal carcinoma, clinical stage II, high nuclear grade, and the A + T regimen were significantly associated with high pCR rates (supplementary Table S1, available at *Annals of Oncology* online).

Multivariate logistic regression analysis of pCR

The reduced multivariate logistic regression model using percentages of ER and PR expression as continuous variables is shown in Table 2. After adjustment for other covariates, low ER and PR expression levels remained significantly associated with high pCR rates (ER: OR, 0.99; 95% CI, 0.986–0.994; P < 0.001; PR: OR, 0.989; 95% CI, 0.984–0.995; P < 0.001). Ductal carcinoma, clinical stage II, high nuclear grade, and the A + T regimen also remained significantly associated with high pCR rates.

We also carried out multivariate logistic regression using ER expression levels as categorical variables. Compared with patients with ER \geq 10% tumors, patients with ER < 1% or 1% \leq ER < 10% tumors had a significantly higher probability of pCR (ER < 1%: OR, 2.15; 95% CI, 1.62–2.87; P < 0.001; 1% \leq ER < 10%: OR, 2.27; 95% CI, 1.48–3.47; P < 0.001). There was no significant difference of pCR rates between ER < 1% and 1% \leq ER < 10% groups in multivariate logistic regression (OR, 0.95; 95% CI, 0.64–1.40; P = 0.792).

Recommended cut-off value for ER positivity

Recursive partitioning and regression trees method without boundary values including all 3055 patients was used to find an optimal cut-off point for the continuous ER with respect to pCR. In this method, the best threshold to split the observations into two separated subgroups was investigated. The recursive partitioning and regression trees method revealed that the recommended cut-off of ER expression below which pCR was likely was 9.5%. Multivariate analysis revealed that the pCR probability of patients with ER < 10% tumors (26.6%) was significantly higher than that of patients with ER \geq 10% tumors (7.0%; OR, 2.17; 95% CI, 1.64–2.87; *P* < 0.0001). Using the same methods, we found that the recommended cut-off of ER expression with respect to TTR was also 9.5%.

Survival analysis

The median follow-up time was 3.9 years for OS. The TTR and OS curves of the ER < 1% and 1% \leq ER < 10% groups were overlapping. The TTR and OS rates of the patients with ER \geq 10% tumors were significantly higher than those of the patients with ER < 1% or 1% \leq ER < 10% tumors (log-rank *P* < 0.0001 for both TTR and OS) (Figure 1A and B). When 10% was used as the cut-off to categorize ER expression level, the TTR and OS of the patients with ER < 10% tumors were significantly lower than

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Table 1. Patient characteristics at diagnosis					
Variable	TOTAL (<i>n</i> = 3055)	ER<1% (<i>n</i> = 932)	1% ≤ ER < 10% (<i>n</i> = 171)	ER \geq 10% (<i>n</i> = 1952)	P value
Age, median (range), years	49 (19–83)	49 (22–83)	49 (19–77)	49 (22–83)	0.49
BMI, median (range), kg/m ²	27.7 (14.5–65.9)	27.8 (14.5–56.4)	29 (17.7–31.3)	27.5 (15.5–66)	0.03
Race/ethnicity					
White	1915 (62.7)	565 (60.6)	97 (56.7)	1253 (64.2)	< 0.001
Black	463 (15.2)	197 (21.1)	35 (20.5)	231 (11.8)	
Hispanic	483 (15.8)	128 (13.7)	25 (14.6)	330 (16.9)	
Asian/others	194 (6.4)	42 (4.5)	14 (8.2)	138 (7.1)	
Menopausal status					
Premenopausal	1461 (47.8)	438 (47.2)	84 (49.1)	939 (48.4)	0.82
Postmenopausal	1579 (51.7)	489 (52.8)	87 (50.9)	1003 (51.6)	
Unknown	15 (0.5)	0 (0)	0 (0)	0 (0)	
Histology					
Ductal	2577 (84.4)	851 (91.3)	156 (91.2)	1570 (80.4)	< 0.001
Lobular	231 (7.6)	15 (1.6)	2 (1.2)	214 (11)	
Others	215 (7.0)	40 (4.3)	7 (4.1)	168 (8.6)	
Unknown	32 (1.0)	26 (2.8)	6 (3.5)	0 (0)	
Nuclear grade					
1/11	1163 (38.1)	91 (9.8)	19 (11.1)	1053 (53.9)	< 0.001
	1803 (59.0)	813 (87.2)	148 (86.5)	842 (43.1)	
Unknown	89 (2.9)	28 (3)	4 (2.3)	57 (2.9)	
Clinical stage	, , ,		. ,	, , ,	
Stage II	1726 (56.5)	511 (54.8)	99 (57.9)	1116 (57.2)	0.46
Stage III	1329 (43.5)	421 (45.2)	72 (42.1)	836 (42.8)	
Neoadjuvant regimen					
Α	292 (95.6)	70 (7.5)	9 (5.3)	213 (10.9)	< 0.001
Т	112 (36.7)	36 (3.9)	3 (1.8)	73 (3.7)	
A+T	2651 (86.8)	826 (88.6)	159 (93)	1666 (85.3)	
ER, continuous, mean ± SD	51.88 ± 43.17	_	_	_	
PR, categorical	51100 = 15117				
PR < 1%	1245 (40.8)	818 (87.8)	110 (64.3)	317 (16.2)	<0.001
$1\% \le PR < 10\%$	326 (10.7)	61 (6.5)	41 (24)	224 (11.5)	<0.001
PR > 10%	1484 (48.6)	53 (5.7)	20 (11.7)	1411 (72.3)	
PR, continuous, mean \pm SD	31.6 ± 38.24	2.4 ± 11.3	4.6 ± 12.5	47.9 ± 38.5	< 0.001
LVI	J1.0 ± J0.24	2.4 - 11.5	4.0 ± 12.5	47.9 - 30.5	<0.001
Negative	2030 (66.4)	631 (67.7)	124 (72.5)	1275 (65.3)	0.005
Positive	915 (30)	254 (27.3)	36 (21.1)	625 (32)	0.005
Unknown	. ,			. ,	
	110 (3.6)	47 (5)	11 (6.4)	52 (2.7)	
Adjuvant chemotherapy	2504 (040)	770 (02 ()	155 (00 6)	1669 (85.5)	0.012
No	2594 (84.9)	770 (82.6)	155 (90.6)	· · ·	0.013
Yes	461 (15.1)	162 (17.4)	16 (9.4)	283 (14.5)	
Adjuvant hormonal therapy	1000 (00 4)	0.46 (00.0)	122 (74.0)	16 (2.4)	.0.001
No	1020 (33.4)	846 (90.8)	128 (74.9)	46 (2.4)	< 0.001
Yes	2035 (66.6)	86 (9.2)	43 (25.1)	1906 (97.6)	
Adjuvant radiation	(2)((2))	227 (25.4)	50 (20 2)	240 (170)	-0.001
No	636 (20.8)	237 (25.4)	50 (29.2)	349 (17.9)	< 0.001
Yes	2419 (79.2)	659 (74.6)	121 (70.8)	1603 (82.1)	
pCR				1016 (02)	
No	2626 (86)	687 (73.7)	123 (71.9)	1816 (93)	< 0.001
Yes	429 (14)	245 (26.3)	48 (28.1)	136 (7)	

All data are no. of patients (%) unless noted otherwise.

ER, estrogen receptor; BMI, body mass index; A, anthracycline; T, taxane; SD, standard deviation; PR, progesterone receptor; LVI, lymphovascular invasion; pCR, pathological complete response.

Table 2. Multivariate logistic regression analysis by pathological complete response (pCR) status

Variable	OR (95% CI)	P value	
Clinical stage			
Stage III	(reference)		
Stage II	1.94 (1.53–2.45)	< 0.001	
ER, continuous	0.990 (0.986-0.994)	< 0.001	
PR, continuous	0.989 (0.984–0.995)	< 0.001	
Race/ethnicity			
White	(reference)		
Black	1.18 (0.88–1.58)	0.28	
Hispanic	1.37 (1.00–1.85)	0.04	
Asian/Others	0.87 (0.52-1.45)	0.58	
Histology			
Ductal	(reference)		
Lobular	0.36 (0.14-0.90)	0.03	
Others	0.510 (0.226-1.000)	0.05	
Nuclear grade			
	(reference)		
1/11	0.45 (0.32-0.63)	< 0.001	
Neoadjuvant regimen			
A+T	(reference)		
А	0.30 (0.17-0.52)	< 0.001	
Т	0.26 (0.10-0.66)	0.004	

OR, odds ratio; Cl, confidential interval; ER, estrogen receptor; PR, progesterone receptor; A + T, anthracycline and taxane; A, anthracycline; T, taxane.

those of the patients with ER \geq 10% tumors (log-rank *P* < 0.0001 for both TTR and OS) (Figure 1C and D).

The univariate Cox proportional hazard models revealed that compared with ER < 1% or $1\% \leq \text{ER} < 10\%$ tumors, ER $\geq 10\%$ tumors were significantly associated with better TTR and OS (supplementary Table S2, available at *Annals of Oncology* online). Patients with ER < 1% or $1\% \leq \text{ER} < 10\%$ tumors did not have significantly different TTR (HR, 1.033; 95% CI, 0.78–1.37: P < 0.82) or OS (HR, 1.086; 95% CI, 0.8–1.48: P < 0.6). After adjustment for covariates including pCR status and adjuvant hormonal therapy, the percentage of ER expression was not significantly associated with TTR or OS (supplementary Table S3, available at *Annals of Oncology* online). Compared with PR $\geq 10\%$ tumors, PR < 1% tumors were significantly associated with worse TTP and OS, and $1\% \leq \text{PR} < 10\%$ tumors were not a significant factor for TTP or OS (supplementary Table S3, available at *Annals of Oncology* online).

Next, to explore the effect of the percentage of ER expression on survival outcomes, we removed adjuvant hormonal therapy from the multivariate Cox regression model. After the removal of adjuvant hormonal therapy and adjustment for other covariates, the multivariate Cox regression model revealed that compared with patients with ER \geq 10% tumors, those with ER < 1% or 1% \leq ER < 10% tumors had worse TTR (ER < 1%: HR, 1.64; 95% CI, 1.34–1.99; *P* < 0.001; 1% \leq ER < 10%: HR, 2.07; 95% CI, 1.4–2.62; *P* < 0.001) and OS (ER < 1%: HR, 2.07; 95% CI, 1.67–2.58; *P* < 0.001; 1% \leq ER < 10%: HR, 2.35; 95% CI, 1.66–3.32; *P* < 0.001).

Difference of survival benefit from adjuvant hormonal therapy

To explore the difference of survival benefit of adjuvant hormonal therapy between patients with $1\% \le ER < 10\%$ tumors and those with ER > 10% tumors, we included an interaction term between adjuvant hormonal therapy and ER category $(1\% \le ER < 10\%$ versus $ER \ge 10\%)$, which was statistically significant in the multivariate models (P < 0.001 for TTR and P = 0.048 for OS). We carried out additional subgroup analyses by these two ER categories. Multivariate analysis revealed that among the 171 patients with $1\% \le ER \le 10\%$ tumors, adjuvant hormonal therapy was not significantly associated with either TTR (HR, 0.88; 95% CI, 0.48–1.6; P = 0.67) or OS (HR, 0.65; 95% CI, 0.32–1.36; P = 0.25) (Table 3). However, among the 1952 patients with ER \geq 10% tumors, adjuvant hormonal therapy was significantly associated with better TTR (HR, 0.24; 95%) CI, 0.16–0.36; P<0.001) and OS (HR, 0.32; 95% CI, 0.2–0.5; *P* < 0.001) (Table 4).

Discussion

We demonstrated that for patients with newly diagnosed stage II or III HER2-negative primary invasive breast cancer treated with NACT, a low percentage of ER expression was significantly associated with a high pCR rate. We also demonstrated that the percentage of ER expression below which pCR was likely was 10%, which is the cut-off value previously recommended by the ASCO/CAP. In addition, we found adjuvant hormonal therapy to have a significant benefit in terms of TTR and OS only in patients with $ER \ge 10\%$ tumors. Although patients with $1\% \le ER < 10\%$ tumors are currently recommended to receive adjuvant hormonal therapy, such patients in the present study did not receive a significant survival benefit from adjuvant hormonal therapy. At MD Anderson Cancer Center, treatment decisions regarding both neoadjuvant and adjuvant systemic therapies follow Breast Medical Oncology department consensus based on National Comprehensive Cancer Network (NCCN) clinical practice guidelines. The consensus meeting is a monthly meeting, and the influence of the preference of the treating physician is minimal.

We found that 5.4% of the HER2-negative breast cancers in the present study had ER expression rates of 1%-9%. This is in line with previous studies that reported that \sim 5% of all breast cancers have ER expression rates of 1%-9%. Because of the low frequency of this category, conducting prospective trials targeting this cohort is not feasible. Until recently, only retrospective studies and post hoc reviews have informed the discussion of stratifying patients with breast cancer according to their likelihood of benefiting from hormonal therapy. However, our findings shed some light on whether patients with HER2-negative breast cancer can be grouped in this manner. St. Gallen International Expert Consensus 2005 recommended three categories of endocrine responsiveness: endocrine responsive, endocrine response uncertain, and endocrine non-responsive [23]. Following this statement, St. Gallen International Expert Consensus 2009 recommended adjuvant endocrine therapy for almost all patients whose tumors show evidence of endocrine responsiveness, which is

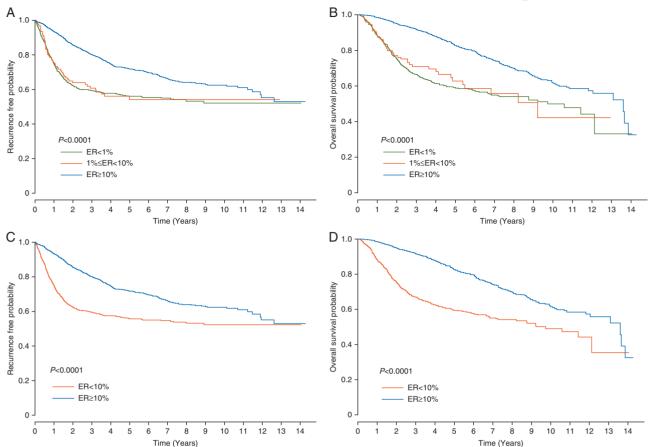


Figure 1. Time to recurrence (TTR) and overall survival (OS) analysis. (A) TTR and (B) OS curves for patients with tumors with <1% of ER expression (ER < 1%), patients with tumors with $\ge1\%$ but <10% of ER expression (1% \le ER < 10%), and patients with tumors with $\ge10\%$ of ER expression (ER $\ge 10\%$). (C) TTR and (D) OS curves for patients with tumors with <10% of ER expression (ER < 10%) and patients with tumors with tumors with $\ge10\%$ of ER expression (ER < 10%) and patients with tumors with $\le10\%$ of ER expression (ER < 10%) and patients with tumors with $\ge10\%$ of ER expression (ER < 10%) and patients with tumors with $\ge10\%$ of ER expression (ER < 10%) and patients with tumors with $\ge10\%$ of ER expression (ER < 10%).

Table 3. Subgroup analysis of patients with 1% \leq ER $<$ 10% expression (n = 171)						
	TTR		os			
Variable	HR (95% CI)	P value	HR (95% CI)	P value		
Clinical stage						
Stage III	(reference)		(reference)			
Stage II	0.38 (0.21–0.67)	< 0.001	0.22 (0.11–0.45)	< 0.001		
Pathological r	esponse					
pCR	(reference)		(reference)			
Non-pCR	3.79 (1.47–9.74)	0.006	4.55 (1.37–15.08)	0.01		
LVI						
Negative	(reference)		(reference)			
Positive	2.22 (1.24–3.97)	0.007	1.93 (1.00–3.68)	0.048		
Adjuvant horr	monal therapy					
No	(reference)		(reference)			
Yes	0.88 (0.48–1.60)	0.67	0.65 (0.32–1.36)	0.25		

Multivariate Cox regression analysis of time to recurrence (TTR) and overall survival (OS).

HR, hazard ratio; CI, confidential interval; pCR, pathological complete response; LVI, lymphovascular invasion.

now defined as the presence of any detectable ER expression [24]. Then, St. Gallen International Expert Consensus 2015 reported that ER values between 1% and 9% were considered equivocal and that endocrine therapy alone cannot be relied upon for patients with these values [18]. However, these categories have not been clearly defined.

One of the novelties of the current study is that we evaluated ER expression as a continuous variable. Also, we derived the cutoff value of ER expression to identify patients who would be more or less likely to achieve pCR after NACT and investigated the benefit from adjuvant hormonal therapy in terms of TTR and OS.

The current study had several potential limitations. First, this was a retrospective study and treatment was not randomly assigned. However, conducting a randomized controlled trial that includes patients with $1\% \leq ER < 10\%$ tumors would be extremely difficult, given the small number of patients with such tumors; moreover, the current recommendations for defining ER and PR positivity are based on retrospective analyses. Second, there were fewer patients with $1\% \leq ER < 10\%$ tumors than patients with $ER \geq 10\%$ tumors. This sample size of patients with $1\% \leq ER < 10\%$ tumors might have been too small to detect statistically significant differences. Despite the small sample size, our multivariate analysis revealed that the interaction term between

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	TTR		OS		
Variable	HR (95% CI)	P value	HR (95% CI)	P value	
Age	0.974 (0.965–0.983)	<0.001	-	_	
BMI	_	-	1.02 (1.01–1.04)	0.01	
Race					
White	(reference)		(reference)		
Black	1.00 (0.75-1.34)	0.98	-	-	
Hispanic	0.74 (0.56–0.98)	0.03	-	_	
Asian/Others	1.03 (0.70-1.52)	0.87	-	_	
Nuclear grade	-	-	(reference)		
	-	-	0.77 (0.6–0.98)	0.03	
1/11					
Histology					
Ductal	_	_	(reference)		
Lobular	_	_	0.78 (0.52–1.16)	0.22	
Others	_	_	0.57 (0.37–0.89)	0.01	
Clinical stage					
Stage III	(reference)		(reference)		
Stage II	0.51 (0.42–0.62)	< 0.001	0.51 (0.4–0.65)	<0.001	
PR, categorical					
$PR \ge 10\%$	(reference)		(reference)		
PR < 1%	1.77 (1.39–2.24)	< 0.001	1.80 (1.36–2.36)	< 0.001	
$1\% \le PR < 10\%$	1.34 (1.01–1.76)	0.04	1.44 (1.03–2.00)	0.03	
Pathological response	1.51 (1.61 1.76)	0.01	1.11(1.05 2.00)	0.05	
pCR	(reference)		(reference)		
Non-pCR	3.31 (1.75–6.25)	<0.001	5.61 (2.07–15.16)	< 0.001	
LVI	5.51 (1.75 0.25)	20.001	3.01 (2.07 13.10)	<0.001	
Negative	(reference)		(reference)		
Positive	1.72 (1.42–2.10)	<0.001	1.68 (1.32–2.13)	< 0.001	
Neoadjuvant regimens	1.72 (1.42-2.10)	<0.001	1.00 (1.32-2.13)	<0.001	
A+T	(reference)				
A	(reference) 1.17 (0.90–1.51)	0.24	_	_	
Т		0.24	-	-	
	1.93 (1.27–2.93)	0.002	-	-	
Adjuvant hormonal therapy	(10601000)		(referrer ce)		
No	(reference)	<0.001	(reference)	<0.001	
Yes	0.24 (0.16–0.36)	<0.001	0.32 (0.20–0.50)	<0.001	

Multivariate Cox regression analysis of time to recurrence (TTR) and overall survival (OS).

HR, hazard ratio; CI, confidential interval; BMI, body mass index; PR, progesterone receptor; pCR, pathological complete response; LVI, lymphovascular invasion; A + T, anthracycline and taxane; A, anthracycline; T, taxane.

adjuvant hormonal therapy and ER category ($1\% \le ER < 10\%$ versus $ER \ge 10\%$) was statistically significant for both TTR and OS, which supported this significant result. Third, some patients' diagnostic biopsies were carried out at outside centers, which could have contributed to low reproducibility of biomarker evaluation in our study sample. However, at MD Anderson Cancer Center, only dedicated breast pathologists review and evaluate all slides, including slides from outside centers. Also, cases are reviewed as part of MD Anderson's internal quality control program and its concordance rate in biomarker evaluation is >95%. Thus, the reproducibility problem of ER and PR expression in the current study must be small. Fourth, data on the biological features of tumors, such as PAM50 results, were not included in the study. ASCO guidelines state that clinicians may use the PAM50 risk of recurrence score to inform their decision

to use adjuvant systemic therapy in patients with ER/PR-positive, HER2-negative, node-negative breast cancer [25]. Fifth, owing to its retrospective nature, our analysis did not account for Ki-67 expression status, which may be associated with pCR and survival outcomes. However, the interlaboratory reproducibility of current methods of Ki-67 assessment is inadequate, and the role of Ki-67 in patients treated with NACT without endocrine therapy is still unclear [26, 27]. In addition, the NCCN Guidelines do not recommend routine testing of Ki-67. Given the lack of reproducibility and the recommendation of the NCCN, Ki67 staining is not routinely carried out as standard practice, and many patients in our study did not have Ki-67 data. Lastly, there is a heterogeneity of systemic treatments after definitive surgery because of the nature of retrospective study. This might affect the analysis of benefit from adjuvant hormonal therapy on the survival outcomes.

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One factor that may need to be taken into consideration when interpreting the findings of the present study is BRCA1 somatic mutation status and its association with poly ADP ribose polymerase (PARP) inhibitors and platinum agents. Approximately 60% or more of breast cancer patients with BRCA1 mutations have TNBC and ~35% of TNBC patients have BRCA1 mutation [28, 29]. Tumors with BRCA mutation tend to respond to PARP inhibitors and/or platinum agents [29-32]. In the era of developing platinum and PARP inhibitors for BRCA-mutated breast tumors, the comprehensive analysis with IHC and genomic analysis of breast tumors, especially low ER expression HER2negative breast tumors, would become more important. Something else that should be considered in the interpretation of the present study's findings is the fact that two different antibodies were used during the study period. Both the 1D5 and 6F11 antibodies have been demonstrated to produce results that correlate with clinical outcomes. However, the quality of antibody as well as the quality of IHC staining methods have improved. Stricter guidelines for the documentation of pre-analytical factors such as fixation time have been implemented, and more robust detection systems are available for archival tissues.

Current ASCO/CAP guidelines recommend that, regardless of HER2 status, breast cancers with <1% of ER expression should be considered ER-negative. However, we found that for patients with newly diagnosed stage II or III HER2-negative primary invasive breast cancer, a low percentage of ER expression was associated with a high pCR rate and that the cut-off percentage of ER expression below which pCR was likely was 10%. In addition, only patients with ER \geq 10% tumors had better survival with adjuvant hormonal therapy. Also, the cut-off of 1% is more reproducible than that of 10% in IHC staining, and the cut-off of 10% tends to be more affected by the number of cells on the slide and the quality of the slides. The quality of pathological assessment was maintained in the current study because only dedicated pathologists evaluated all slides.

In conclusion, stage II or III HER2-negative primary invasive breast cancer with <10% of ER expression behaves clinically like TNBC in terms of pCR and survival outcomes and patients with such tumors may have a limited benefit from adjuvant hormonal therapy. It may be more clinically relevant to define TNBC as HER2-negative breast cancer with <10%, rather than <1%, of ER and/or PR expression. Studies with other datasets to confirm our findings are warranted.

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