Biomarkers, Genomics, Proteomics, and Gene Regulation

PTEN, PIK3CA, p-AKT, and p-p70S6K Status

Association with Trastuzumab Response and Survival in Patients with HER2-Positive Metastatic Breast Cancer

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Phosphatase and tensin homolog (PTEN) is a key modulator of trastuzumab sensitivity in HER2-overexpressing breast cancer. Because PTEN opposes the downstream signaling of phosphoinositide 3-kinase (PI3K), we investigated the role of PTEN and other components of the PI3K pathway in trastuzumab resistance. We analyzed the status of PTEN, p-AKT-Ser473, and p-p70S6K-Thr389 using immunohistochemistry. PIK3CA mutation status was analyzed by direct sequencing. Primary tumor tissue was available from 137 patients with HER2-overexpressing metastatic breast cancer who had received trastuzumabbased chemotherapy. We observed that each of the four biomarkers alone did not significantly correlate with trastuzumab response, whereas PTEN loss alone significantly correlated with shorter survival times (P = 0.023). PI3K pathway activation, defined as PTEN loss and/or PIK3CA mutation, was associated with a poor response to trastuzumab (P = 0.047) and a shorter survival time (P = 0.015). PTEN loss was significantly associated with a poor response to trastuzumab (P = 0.028) and shorter survival time (P =0.008) in patients who had received first-line trastuzumab and in patients with estrogen receptor- (P =0.029) and progesterone receptor-negative tumors (P = 0.033). p-AKT-Ser473 and p-p7086K-Thr389 each had a limited correlation with trastuzumab response. When these markers were combined with PTEN loss, an increased correlation with patient outcome was observed. In conclusion, PI3K pathway activation plays a pivotal role in trastuzumab resistance. Our findings may facilitate the evaluation of tumor response to trastuzumab-based and targeted therapies. (*Am J Pathol 2010, 177:1647–1656;* DOI: 10.2353/ajpath.2010.090885)

Human epidermal growth factor receptor 2 (HER2) is overexpressed in 20% to 25% of invasive breast cancers. Patients with HER2-overexpressing tumors experience a shorter time to relapse and shorter overall survival than patients with tumors of normal HER2 levels.^{1,2} HER2 overexpression can lead to activation of many downstream signaling molecules, including Ras-Gap, Src, phosphoinositide 3-kinase (PI3K)/AKT, and many other signaling events.^{3,4} Trastuzumab (Herceptin; Genentech, CA), a humanized monoclonal antibody that directly targets the extracellular domain of HER2, has a remarkable therapeutic efficacy in treating patients with HER2-expressing metastatic breast cancer (MBC)⁵ and patients with HER2-positive early-stage disease in adjuvant settings.6,7 Trastuzumab treatment, when combined with chemotherapy, resulted in a significant improvement in patients' response rate, time to progression, and duration

F.J.E. and H.G. contributed equally to this study.

Accepted for publication June 16, 2010.

Supplemental material for this article can be found on *http://ajp. amjpathol.org.*

Supported in part by NIH grants P30-CA 16672 (MDACC), M.D. Anderson Cancer Center Breast SPORE P50 CA116199 (project 4), and the Breast Cancer Research Foundation (F.J.E); RO1-CA112567, DOD Center of Excellence grant subproject W81XWH-06-2-0033, DOD Synergistic Award W81XWH-08-1-0712, and Susan G. Komen Breast Cancer Foundation Promise grant KG091020 (D.Y.). D.Y. is the Hubert L. & Olive Stringer Distinguished Chair in Basic Science at M.D. Anderson Cancer Center.

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of response.⁸ The underlying mechanisms of trastuzumab's antitumor activities include, but are not limited to, inducing antibody-dependent cellular cytotoxicity,⁹ inhibiting HER2 extracellular domain cleavage,¹⁰ activating phosphatase and tensin homolog (PTEN),¹¹ and inhibiting PI3K/AKT survival signaling.¹² However, the overall response rate to trastuzumab is low, and almost half of patients with HER2-positive breast cancer exhibit an initial resistance to trastuzumab-based therapy.^{11,13} Despite the large amounts of preclinical and clinical data, the causes of trastuzumab resistance are still poorly understood.¹⁴

The PI3K pathway, downstream of HER2, plays a central role in regulating a number of cellular processes, such as apoptosis, migration, angiogenesis, cell proliferation, and glucose metabolism, and it is involved in trastuzumab resistance.^{15,16} PI3K phosphorylates phosphatidylinositols on the cell membrane, generating phosphatidylinositol-3,4,5-trisphosphate (PIP3) from phosphatidylinositol-4,5-bisphosphate (PIP2). Then, at the cell membrane, PIP3 recruits protein kinases and activates protein kinase B (PKB)/AKT.¹⁷ In breast cancer cells, HER2 overexpression can lead to activation of the PI3K/AKT pathway.¹⁸ The activation of AKT and its downstream signaling have been demonstrated to inhibit cell cycle arrest and block trastuzumab-mediated apoptosis.¹² AKT phosphorylation and AKT kinase activities were found to be increased in trastuzumab-resistant cells, derived from BT474 HER2-overexpressing breast cancer cells, when compared with parental cells.¹⁹ These data provide insight into the trastuzumab-resistance mechanism of PI3K/AKT signaling.¹⁵

Aberrations in the components of the PI3K pathway have been reported in most solid tumors, including breast cancer.¹⁶ PTEN is a tumor suppressor that dephosphorylates the D3 position of PIP3 and inhibits the PI3K/AKT pathway.²⁰ Loss of PTEN function as a result of mutation, deletion, or promoter methylation has been reported in nearly 50% of breast cancers.¹¹ In addition, the gene encoding one of the PI3K catalytic subunits, p110 α (PIK3CA), has been found to be mutated in about 25% of breast cancers.^{21,22,23} Most of the reported mutations are localized to hotspots in exons 9 and 20 of the PIK3CA gene, which result in increased PI3K pathway signaling.^{22,24} We previously discovered that PTEN activation is a novel mechanism of trastuzumab antitumor function, and PTEN loss confers trastuzumab resistance in HER2overexpressing breast cancer cells.¹¹ PTEN loss significantly predicted poor response to trastuzumab-based therapy in a small cohort of HER2-positive patients with MBC.¹¹ Later, it was reported that both low PTEN levels and PI3K-activating PIK3CA mutations contribute to trastuzumab resistance in HER2-overexpressing breast cancer.25,26 PTEN loss or PIK3CA mutations, which indicate activation of the PI3K pathway, are considered as markers for poor response to trastuzumab in patients with HER2-overexpressing breast cancer.²⁵ On the other hand, some studies found no correlation between PTEN expression and trastuzumab response or survival in patients with HER2-positive breast cancer.27,28 These contradictory findings prompted us to further investigate the association between PTEN status and clinical outcomes in a large cohort of patients with MBC who were treated with trastuzumab-based therapy. We tested the hypothesis that a comprehensive assessment of PI3K pathway activation status provides biomarkers that can identify patients who may not benefit from trastuzumab-based therapy.

Materials and Methods

Patients and Tissue Samples

HER2-positive tumor samples were obtained from 137 eligible patients at the University of Texas M. D. Anderson Cancer Center under an institutional review board-approved protocol. All of the patients, who subsequently developed MBC, were treated with trastuzumab-based therapy between 1998 and 2006. Tumors that scored 3+ on immunohistochemical (IHC) analysis of HER2 and/or had an amplified *HER2* gene detected by fluorescence *in situ* hybridization (FISH) were defined as HER2 positive. PTEN, p-AKT, and p-p70S6K status, determined by IHC analysis, were evaluated on serial slides and scored on all tissue samples, except for one sample on which p-AKT scoring could not be obtained.

IHC Analysis

IHC staining was performed on 4-µm slices of formalinfixed paraffin-embedded (FFPE) tissue sections with primary antibodies against PTEN (138G6, 1:50; Cell Signaling Technology, Danvers, MA), p-AKT (Ser473, 736E11, IHC specific, Cell Signaling; 1:50), and p-p70S6K (Thr389, 1A5, Cell Signaling, 1:700). To validate the specificity of PTEN antibodies, Western blotting was performed first using both BT474 control cells and BT474 PTEN-deficient cell lines (by shRNA knockdown). Then mouse xenograft tumors derived from above cell lines were embedded in paraffin and used as negative and positive control for IHC staining. After comparing four PTEN antibodies from different companies (Cell Signaling; R&D Systems, Minneapolis, MN; Cascade Bioscience, Winchester, MA; and Dako, Carpinteria, CA), we found that the antibody from Cell Signaling is the most specific in our studies (data not shown). The p-AKT and p-p70S6K stainings were validated by Western blotting and other studies, as previously described.^{29,30} After deparaffinization and rehydration, slides were subjected to heat-induced epitope retrieval in 10 mmol/L citrate buffer (pH 6.0) for p-p70S6K staining or 1 mmol/L EDTA (pH 8.0) for PTEN and p-AKT staining in a hot water bath (95°C) for 20 minutes. Endogenous peroxidase activity was blocked for 15 minutes in 0.3% hydrogen peroxide. After blocking with 1% goat serum for 1 hour at room temperature, the sections were incubated with primary antibodies for at least 18 hours at 4°C overnight. Immunodetection was performed using the LSAB2 kit (Dako), 3-3'-diaminobenzidine was used for color development, and hematoxylin was used for counterstaining. Positive control slides (from BT474 xenograft tumors) and negative control slides (without primary antibody) were included within each batch for staining.

Evaluation of IHC Staining

The PTEN, p-AKT, and p-p70S6K expression levels were evaluated based on the staining intensity (SI) and percentage of positive cells (PP) within the whole tissue section. PTEN was scored semiquantitatively using the immunoreactive score (IRS), which was calculated as follows: IRS = SI × PP. SI was defined as 0 = negative, 1 = weak, 2 = moderate, and 3 = strong. PP was scored as 0 = 1%; 1 = 1%-10%; 2 = 11%-50%; 3 = 51%-80%; and 4 = >80% positive cells. PTEN loss was defined as an IRS of 3 or less.¹¹ The p-AKT and p-p70S6K stainings were evaluated by an H score,³¹ which was calculated by multiplying the PP by the corresponding SI (1 = weak, 2 = moderate, and 3 = strong), giving a maximum score of 300 (100% × 3). H scores >50 were considered positive.

PIK3CA PCR, Sequencing, and Mutational Analysis

We isolated genomic DNA from FFPE tumor samples for PIK3CA mutation screening from 90 patients. The other 47 tumors are very small in size and did not allow analysis of PIK3CA mutations. PIK3CA mutation screening was performed by using the QIAamp DNA FFPE tissue kit (Qiagen, Valencia, CA) according to the kit protocol. First, the FFPE slides were stained with hematoxylin and eosin and checked by a pathologist to identify the tumor region. After deparaffinization, most of the tumor was cut out from the slides by a razor blade. For a few samples, dissection was aided by laser capture microscopy owing to the dispersion of the tumor cells. For laser capture, we used a PALM MicroBeam Microscope for Laser Micromanipulation (Carl Zeiss, Inc., Thornwood, NY) with HV-D30 Hitachi camera capture system (Tokyo, Japan). The microscope contains built-in class IIIB and class IV lasers. The equipment was run using PALM Robo Software 4.0.0.10 (Carl Zeiss), and all laser capture was directed by a pathologist.

The isolated genomic DNA was diluted to 3 ng/ μ l with TE buffer (10 mmol/L Tris-HCl, 0.1 mmol/L EDTA [pH 8.0]) and sent to a DNA sequencing facility for sequencing of exons 9 and 20 of PIK3CA. For each sample, we submitted 6 ng of genomic DNA. The DNA was amplified using the primer pair 5'-GTTTTCCCAGTCACGACGTATT-GAAAATGTATTTGCTTTTTC and 5'-GGAAACAGCTAT-GACCATAACATGCTGAGATCAGCCAAAT for exon 9 and the primer pair 5'-GTTTTCCCAGTCACGACGAT-TAACATCATTTGCTCCAAAC and 5'-AGGAAACAGCTA-TGACCATGCTGTTTAATTGTGTGGAAGAT for exon 20. The primers also contained M13 forward and reverse tails to facilitate sequencing. After amplification, the DNA seguence was determined on a Mega BASE 4500 (GE Health care, Waukesha, WI) using dye-primer chemistry, as previously described.³² The observed mutations were confirmed by two independent amplification and sequencing reactions. The success rate was 60% (55/90).

Evaluation of Trastuzumab Response and Statistical Analyses

Response to trastuzumab-based therapy was evaluated every 8 to 12 weeks using the response evaluation criteria in solid tumors (RECIST) with modifications.³³ A complete response (CR) was defined as the complete disappearance of all radiographically or visually apparent tumors for at least 4 weeks. A partial response (PR) was defined as a \geq 30% decrease in the maximum diameter of measurable metastatic lesions for at least 4 weeks, with no new lesions appearing. Stable disease (SD) was defined as no change in the size of measurable lesions \geq 20% and no appearance of new lesions. Progressive disease (PD) was defined as the appearance of any new lesion or an increase in the size of the existing lesions \geq 20%. In this study, response was defined as a CR, PR, or SD longer than 6 months (clinical benefit = CR + PR +SD). Overall survival (OS) was defined as the time from initiation of trastuzumab treatment to the patient's death or the final day of follow-up.

The relationship between the expression of the biomarkers and response to trastuzumab and clinicopathologic features was analyzed using Fisher's exact test and the χ^2 test. Survival analyses were performed using Kaplan–Meier curves with log-rank tests. A *P* value <0.05 (two-sided) was considered to be statistically significant. All statistical analyses were performed using the SPSS 16.0 program (SPSS, Chicago, IL).

Results

Demographic and Pathological Characteristics of the Study Population

The study population comprised 137 patients with HER2positive MBC who received trastuzumab treatment. The patients' clinical characteristics are summarized in Table 1. The median age at first diagnosis was 44 years (range, 20-73 years). Of the 137 patients, 123 (89.8%) were initially diagnosed with invasive ductal carcinoma. Most patients were white (104, 75.9%); 54 patients (39.4%) had estrogen receptor (ER)-positive tumors, and 44 (32.1%) had progesterone receptor (PR)-positive tumors. The median disease-free survival time before developing metastatic disease was 18 months (range, 0-169 months), and most patients (51.8%) had multiple distant metastases. Among the 137 patients, the three most frequent first metastatic sites were the liver (65, 47.4%), bone (60, 43.8%), and lungs (52, 38.0%). During followup, 45 patients (32.8%) developed brain metastases, of which 31 (31/45 = 68.9%) were diagnosed with brain metastasis after initiation of Trastuzumab-based therapy.

The patients' demographic characteristics had no relationship with response to trastuzumab or overall survival after trastuzumab treatment, except for race. Black patients had the shortest OS after starting trastuzumab (P = 0.014) compared with other groups. Information on the patients' trastuzumab-based therapy and patients' clinical outcomes were summarized in Table 2. Of the

Table 1.Clinicopathological Characteristics of 137 Metastatic
Breast Cancer Patients Treated with Trastuzumab-
Based Therapy

* *	
Characteristic	n (%)
Age at diagnosis (median 44 years, range 20–73)	
<50 years	89 (65)
≥50 years	48 (35)
Nuclear grade	
	0
II	22 (16)
111	106 (77.4)
Unknown	9 (6.6)
Histologic type	· · ·
Invasive ductal carcinoma	123 (89.8)
Invasive mixed ducal/lobular carcinoma	8 (5.8)
Invasive lobular carcinoma	2 (1.5)
Inflammatory carcinoma	1 (0.7)
Invasive micropapillary carcinoma	1 (0.7)
Breast cancer, NOS	2 (1.5)
Race/ethnicity	
White	104 (75.9)
Black	8 (5.8)
Hispanic	17 (12.4)
Asian	6 (4.4)
Others	2 (1.5)
ER status	FO (40 0)
Positive	56 (40.9)
Negative	78 (56.9)
Unknown PR status	3 (2.2)
Positive	45 (32.8)
Negative	45 (32.8) 85 (62.0)
Unknown	7 (5.1)
First site of metastasis	7 (0.1)
Liver	65 (47.4)
Bone	60 (43.8)
Lung	52 (38)
Brain	9 (6.6)
Brain metastasis	0 (0.0)
Total cases from time of original	45 (32.8)
diagnosis	(02.0)
2.2.3.10010	

ER indicates estrogen receptor; PR, progesterone receptor.

137 patients, 87 (63.5%) used trastuzumab treatment as first-line therapy, and 91 (66.4%) had received no prior chemotherapy. Eighty-two patients (59.9%) responded to trastuzumab-based therapy (CR + PR + SD), with a median OS of 30 months (range, 2–84 months). However, 55 patients (40.1%) were resistant to trastuzumab (PD), with a median OS of only 13 months (range, 1–49, months; P < 0.001 compared with CR + PR + SD group).

PTEN and PIK3CA Status and Their Relationship with the Patients' Clinicopathologic Characteristics

To determine PTEN protein expression in breast tumors, we first validated the specificity and sensitivity of PTEN antibodies for IHC (see *Materials and Methods* and Supplemental Figure S1, A and B at *http://ajp.amjpathol.org*). As shown in Figure 1A, PTEN expression in tumors was heterogeneous and PTEN had both cytoplasmic and nuclear localizations. A strong PTEN staining in fibroblasts cells was used as internal positive controls along with paraffin sections containing both control.shRNA and

 Table 2.
 Trastuzumab-Based Therapy Information and Clinical Outcomes of the 137 Metastatic Breast Cancer Patients

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Variable	Description
Ttzm-based therapy	
Ttzm single agent	16 (11.7%)
Ttzm + navelbine	30 (21.9%)
Ttzm + taxane-based	76 (55.5%)
chemotherapy	70 (00.070)
Ttzm $+$ other therapy	15 (10.9%)
Lines of Ttzm-based therapy	10 (10.070)
First-line	87 (63.5%)
Second-line	26 (19.0%)
Other	24 (17.5%)
Prior chemotherapy	21(11.070)
Without	91 (66.4%)
With	39 (28.5%)
Unknown	7 (5.1%)
Response to Ttzm-based	. (0.1.70)
therapy	
CR + PR + SD	82 (59.9%)
PD	55 (40.1%)
Outcome	
Dead	97 (70.8%)
Alive	40 (29.2%)
Median time of overall	23m (range 1-84 m)
survival from Ttzm-based	(S)
therapy start date	
Patients responsive (CR +	30m (range 2–84 m)
PR + SD to therapy	
Patients resistant (PD) to	13m (range 1–49 m)
therapy	

Ttmz indicates trastuzumab; CR, complete response; PR, partial response; SD, stable disease; PD, progressive disease.

PTEN.shRNA xenograft tumors as positive and negative controls (Figure 1B). Based on our previous studies, we used an IRS score \leq 3 as the criterion for PTEN loss or reduced PTEN expression.¹¹ Using this cut-off, we observed loss of PTEN expression in 71 tumors (51.8%) in the 137 patients' HER2-positive MBC (Table 3). The patients' clinico-pathological characteristics were not significantly associated with PTEN status (data not shown).

We performed *PIK3CA* mutation analysis on 90 of the 137 samples. The other 47 tumors are very small in size and did not allow analysis of *PIK3CA* mutations. The success rate (as measured by getting reliable sequence at E542, E545, and H1047) was 60% (55/90). We also confirmed the findings by PCR analyses. We found that 12 tumors (21.8%) had *PIK3CA* mutations among the 55 patients with HER2-overexpressing breast cancer (Table 3). The most common mutation was an *H1047R* mutation in exon 20 (7/12, 12.7%). Four of the 12 tumors had an *E545K* mutation, and one had an *E542K* mutation. *PIK3CA* mutation status had no association with any of the clinic-pathological parameters (P > 0.05, data not shown).

Relationship Between PTEN and PIK3CA Mutation Status and Patients' Clinical Outcomes after Trastuzumab-Based Therapy

We found that among the 137 patients, those who had tumors with PTEN loss (IRS \leq 3) were more likely to be resistant to trastuzumab-based therapy than those who

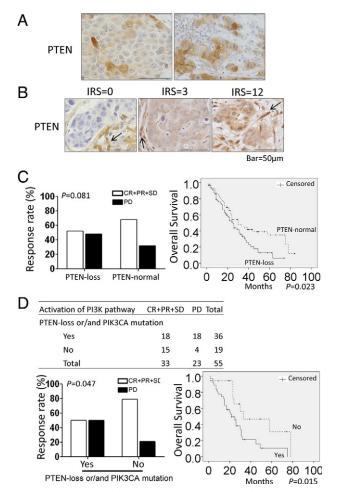


Figure 1. Activation of the PI3K pathway was more significantly associated with trastuzumab resistance than PTEN status alone. **A:** Heterogeneous expression of the PTEN protein in invasive ductal carcinoma lesions examined by IHC in HER2-positive MBC tumors. **B:** Representative PTEN IHC staining (IRS = 0, 3, 12). **Arrows** point to fibroblasts and endothelial cells that showed high expression of PTEN, which served as an internal positive control. **C:** Poor response (P = 0.081) and significantly shorter OS (P = 0.023) after trastuzumab-based therapy in patients with PTEN loss alone (PTEN loss defined as IRS ≤ 3). **D:** Significantly poorer response (P = 0.047) and shorter OS (P = 0.015) after trastuzumab-based therapy in patients with PTSN loss alone. Scale bar = $50 \ \mu m$.

had PTEN-normal tumors, but PTEN loss itself was not significantly correlated with response (P = 0.081; Table 3; Figure 1C). The Kaplan–Meier curve demonstrated a significantly shorter survival for patients with PTEN-deficient tumors (P = 0.023; Figure 1C). *PIK3CA* mutation was not significantly associated with either a response to trastuzumab-based therapy (P = 1.000; Table 3) or an improved OS time (P = 0.172); patients with the *PIK3CA* mutation seemed to have a slight trend toward shorter OS times (Table 3; also see Supplemental Figure S2 at *http://ajp.amjpathol.org*).

Because both PTEN loss and *PIK3CA* mutations lead to PI3K pathway activation and because activated PI3K signaling were reported to contribute to trastuzumab resistance, we analyzed trastuzumab response based on a combination of PTEN and *PIK3CA* mutation status. Patients having tumors with PI3K pathway activation (defined as PTEN loss and/or *PIK3CA* mutation) were more

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Variables	CR + PR + SD	PD	Total (%)	<i>P</i> Value
PTEN PTEN loss PTEN normal Total PIK3CA mutation	37 45	34 21	71 (51.8) 66 (48.2) 137 (100)	0.081
+ _ Total	7 26	5 17	12 (21.8) 43 (78.2) 55 (100)	1.000

 Table 3.
 Associations between PTEN and PIK3CA Status and Response to Trastuzumab-Based Therapy

CR indicates complete response; PR, partial response; SD, stable disease; PD, progressive disease.

resistant to trastuzumab-based therapy (PD) and received less clinical benefit (CR + PR + SD) than patients having tumors without PI3K pathway activation (P = 0.047; Figure 1D). In addition, patients having tumors with an activated PI3K pathway had a significantly shorter OS (P = 0.015) than patients whose tumors do not have PI3K pathway activation (Figure 1D).

Correlation between p-AKT and p-p70S6K Status and Their Relationship with Patients' Clinical Outcomes after Trastuzumab-Based Therapy

Because PTEN loss and PIK3CA mutations are involved in trastuzumab resistance and both can lead to activation of downstream signaling pathways, including increased p-AKT and p-p70S6K, we hypothesized that activation of signals downstream of the PI3KCA mutation and PTEN loss, such as increased p-AKT and p-p70S6K, would also correlate with trastuzumab resistance and serve as molecular markers for trastuzumab response. IHC staining showed that p-AKT had both cytoplasmic and nuclear localization, whereas p-p70S6K mainly had nuclear staining (Figure 2A). In some cases, tumors also displayed variable heterogeneous p-AKT and p-p70S6K staining similar to those of PTEN. We used the H score to quantify the staining, and positivity of p-AKT (p-AKT+) and p-p70S6K (p-p70S6K+) were defined as H \geq 50. We detected p-AKT+ and p-p70S6K+ in 72 (52.9%) and 93 (67.9%) tumors, respectively. Notably, p-AKT and p-p70S6K status were significantly correlated (P = 0.001; see Supplemental Table S1 at http://ajp.amjpathol.org). However, p-AKT and p-p70S6K status did not correlate with any of the patients' clinicopathologic characteristics described in Table 1 (data not shown).

The status of p-AKT and p-p70S6K alone showed no correlation with either trastuzumab response (P = 0.298, P = 0.136; see Supplemental Table S2 at *http://ajp. amjpathol.org*) or OS (P = 0.526, P = 0.319; see Supplemental Figure S3A and S3B at *http://ajp.amjpathol.org*). Combining p-AKT status and p-p70S6K status also had no correlation with either trastuzumab response or OS (data not shown).

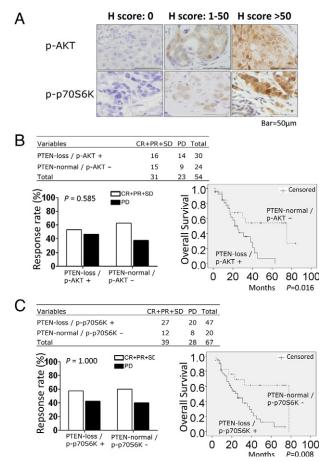


Figure 2. Combing either p-AKT+ or p-p70S6K+ with PTEN loss correlated with trastuzumab resistance. **A:** Representative IHC staining patterns of p-AKT and p-p70S6K in HER2-positive MBC tumors. **B:** Combining p-AKT+ and PTEN loss alone. Patients with PTEN-loss/p-AKT+ tumors showed worse responses and significantly shorter survival times (P = 0.016) after trastuzumab-based therapy. **C:** Combining p-p70S6K+ and PTEN loss alone. Patients with PTEN-loss/p-p70S6K+ and PTEN loss alone. Patients with PTEN-loss/p-p70S6K+ tumors showed worse responses and significantly shorter of trastuzumab resistance than PTEN loss alone. Patients with PTEN-loss/p-p70S6K+ tumors showed worse responses and significantly shorter OS (P = 0.008) after trastuzumab-based therapy. Scale bar = 50 μ m.

p-AKT or p-p70S6K Combined with PTEN Status Correlated Well with Patients' Clinical Outcomes after Trastuzumab-Based Therapy

We next examined whether combining either p-AKT+ or p-p70S6K+ with PTEN loss correlated with trastuzumab resistance. The combination of PTEN loss with p-AKT+ was not significantly correlated with trastuzumab response (P = 0.585; Figure 2B). However, we observed a trend that patients with PTEN-loss/p-AKT+ tumors were more resistant to trastuzumab-based therapy than those with PTEN-normal/p-AKT- tumors. Moreover, the OS time after trastuzumab-based therapy was significantly shorter among those patients who had PTEN-loss/p-AKT+ tumors than those with PTEN-normal/p-AKT- tumors (P = 0.016; Figure 2B). Combined p-AKT+ and PTEN loss might be a more powerful marker of trastuzumab resistance than PTEN loss alone (P = 0.023; Figure 1B). Similarly, combined PTEN and p-p70S6K status showed no correlation with trastuzumab response (P = 1.000; Figure 2C), whereas patients who had PTENloss/p-p70S6K+ tumors had a significantly shorter OS after trastuzumab-based therapy than those with PTENnormal/p-p70S6K- tumors (P = 0.008; Figure 2C). Thus, combined p-p70S6K+ and PTEN loss might also be a more powerful marker of trastuzumab resistance than PTEN loss alone (P = 0.023; Figure 2C).

Correlation of PTEN Status with Patients' Clinical Response to Trastuzumab-Based Therapy in Different Therapeutic Subgroups

In the whole cohort, PTEN loss alone was not significantly associated with resistance to trastuzumab-based therapy (P = 0.081). The patients had varying treatment backgrounds, and we postulated that the predictive effect of PTEN status for trastuzumab response might be confounded by these different backgrounds. To examine this, we performed extended analyses of PTEN status and patients' clinical response to trastuzumab-based therapy after adjusting for the patients' treatment backgrounds.

First, we found that the rate of PTEN loss in tumors did not differ among different therapeutic subgroups of patients (see Supplemental Table S3 at http://ajp.amjpathol. org). Next, we performed a correlation analysis of PTEN status and trastuzumab response in different therapeutic subgroups. Among patients using trastuzumab as firstline therapy, PTEN loss alone was significantly associated with resistance to trastuzumab (P = 0.028) and more significantly correlated with shorter OS (P = 0.008) than in the whole cohort (P = 0.023). However, if the analysis was confined to the subgroup using trastuzumab as "other-line therapy" (\geq second-line therapy), there were no significant differences in trastuzumab response or OS between patients with PTEN-loss tumors and those with PTEN-normal tumors (P = 1.000, P = 0.863; Table 4). We also compared the whole cohort with the subgroup who received no prior che-

Table 4.Associations of PTEN Status with Clinical
Outcomes after Trastuzumab-Based Therapy in
Different Therapeutic Subgroups

	Re	spoi	Overall	
Therapeutic Subgroups	CR+PF +SD	{		Survival P Value (Log Rank)
Ttzm as first-line therapy PTEN loss PTEN normal	24 31	22 10	0.028	0.008
Ttzm as other-line therapy PTEN loss PTEN normal	13 14	12 11	1.000	0.863
Without prior chemotherapy PTEN loss PTEN normal	21 35	22 13	0.030	0.013
With prior chemotherapy PTEN loss PTEN normal	12 9	11 7	1.000	0.922

Ttzm indicates trastuzumab; CR, complete response; PR, partial response; SD, stable disease; PD, progressive disease.

Subgroup Variable	PTEN Loss	PTEN Normal	Total	<i>P</i> Value
ER-/PR- patients Other patients (ER-/PR+ or ER+/PR- or ER+/PR+)	42 27	27 36	69 63	
Total	69	63	132	0.039

Table 5.Correlation between PTEN Status and ER/PR
Expression Status

ER indicates estrogen receptor; PR, progesterone receptor.

motherapy before trastuzumab-based treatment and found that PTEN loss was significantly associated with trastuzumab resistance (P = 0.030) and shorter OS (P = 0.013) in patients who received no prior chemotherapy before trastuzumab-based treatment. However, *PIK3CA* mutations had no correlation with trastuzumab response in any subgroup (see Supplemental Table S4 at *http://ajp.amjpathol.org*).

Associations of PTEN Status with Patients' Clinical Outcomes after Trastuzumab-Based Therapy in Different ER and PR Expression Subgroups

Previous studies reported that the HER2 signaling pathway can cross talk with ER,⁴ and ER may play a role in resistance to HER2 inhibitors.³⁴ To assess whether hormone receptor status involves in trastuzumab response, we performed trastuzumab response analyses based on hormone receptor status in patients' breast tumors. We found that neither ER status nor PR status was correlated with clinical outcomes in the whole cohort, but ER- patients had a trend toward PD after trastuzumab-based therapy (P = 0.099, data not shown). Interestingly, PTEN loss was significantly associated with ER-/PR- status in this cohort (P = 0.039; Table 5). When we classified patients based on their ER-/PR- status, PTEN loss was found to be significantly associated with trastuzumab resistance (P = 0.029) and shorter OS (P = 0.033) in the ER-/PR- subgroup (Table 6).

Discussion

To our knowledge, this is the analysis of PI3K pathway activation (PTEN loss, *PI3K* mutation, pAKT+, and p-p70S6K+) in the largest cohort of HER2-overexpressing breast cancer patients who received trastuzumab-based therapy for metastatic disease. Our results demonstrated that the status of the four biomarkers (PTEN, *PIK3CA*, p-AKT, and p-p70S6K) did not significantly correlate with trastuzumab response when analyzed individually. Only PTEN loss was significantly correlated with shorter OS after trastuzumab-based therapy. PI3K pathway activation, defined as PTEN loss and/or *PIK3CA* mutation, significantly contributed to both worse response to trastuzumab and shorter OS. Although p-AKT and

 Table 6.
 Association of PTEN Status with Clinical Outcomes after Trastuzumab-Based Therapy in Different ER/ PR Expression Subgroups

	Response			
Subgroup Variable	CR+PR +SD	PD	P Value (Fisher Exact)	Overall Survival <i>P</i> Value (Log Rank)
ER-/PR- patients PTEN loss PTEN normal Other patients (ER-/PR+ or ER+/PR- or ER+/PR+) PTEN loss	18 19 17	24 8	0.029	0.033
PTEN loss PTEN normal	24	10 12	0.795	0.348

ER indicates estrogen receptor; PR, progesterone receptor; CR, complete response; PR, partial response; SD, stable disease; PD, progressive disease.

p-p70S6K status had limited correlation with trastuzumab response and OS in this cohort, when combined with PTEN status, PTEN loss/p-AKT+ and PTEN loss/pp70S6K+ were significantly related to shorter OS, with more power than PTEN loss alone. Remarkably, PTEN loss was significantly associated with trastuzumab resistance in patients who received trastuzumab as their firstline therapy or those who did not receive prior chemotherapy and in patients whose tumors were ER-/PR-. These data suggest that PI3K pathway activation through PTEN loss or PIK3CA mutation plays a pivotal role in resistance to trastuzumab-based therapy in patients with HER2-overexpressing MBC, particularly in patients who received trastuzumab as first-line therapy, those who had not received prior chemotherapy, and those with ER-/ PR- tumors).

Trastuzumab resistance is becoming increasingly important in the clinic, as understanding mechanisms of resistance may lead to novel therapeutic approaches to overcome resistance. Although trastuzumab may decrease the risk of recurrence by 50% in patients with HER2-positive early-stage breast cancer, approximately 15% of patients develop metastatic disease despite adjuvant trastuzumab therapy.35 Identifying the most relevant and effective predictive biomarkers of trastuzumab response is essential for selecting subgroups of patients who can benefit from trastuzumab-based therapy and patients who may need additional or alternative therapies. Multiple genetic and molecular alterations have been found to contribute to trastuzumab resistance, among which PI3K pathway constitutive activation is one of the most important.11,25,36

In this study, PTEN-loss and *PIK3CA* mutations were found in 51.8% and 21.8%, respectively, of the patients with HER2-positive MBC, which is consistent with previous reports.^{11,37,38} Taking into consideration PTEN-loss and *PIK3CA* mutation together, we found that patients having tumors without PI3K pathway activation received more benefit from trastuzumab-based therapy than patients with PI3K pathway activation tumors; patients having tumors without PI3K pathway activation had longer OS and better responses to trastuzumab. This observation indicates that evaluating both PTEN loss and *PIK3CA* mutations, or PTEN loss and PI3K downstream signaling events (p-AKT or p-p70S6K), will better determine which patients are more likely to benefit from trastuzumabbased therapy in the metastatic setting. Our results also indicate that the PI3K pathway plays a pivotal role in conferring trastuzumab resistance, which is consistent with the results of previous studies.^{11,25} On the basis of the critical contribution of the PI3K pathway in human cancer, it is rational that combining trastuzumab with PI3K/AKT inhibitors could overcome trastuzumab resistance.³⁹

Previous studies showed that PTEN loss led to increased PI3K/AKT signaling and that HER2-overexpressing breast cancer cell lines with low PTEN expression were more resistant to trastuzumab.11,25,26,36 Additionally, an unbiased large-scale RNA interference screen on genes involved in trastuzumab resistance in breast cancer identified PTEN as the only modulator of trastuzumab sensitivity.²⁵ In our study, although PTEN loss alone was borderline significant with trastuzumab response (P =0.081), there was a trend toward lower response rates and significantly shorter OS (P = 0.023) after trastuzumab-based therapy in patients with tumors of low PTEN expression. However, PIK3CA mutation status did not correlate with either response or survival after trastuzumab treatment, suggesting that the role of PTEN loss may be more important than PIK3CA mutations in conferring trastuzumab resistance.

Our data indicated that the relationship between PTEN status and trastuzumab response and OS might be confounded by the patients' treatment backgrounds, as these were highly varied. Therefore, it was necessary to consider the patients' treatment background when analyzing the role of PTEN status in trastuzumab resistance. Remarkably, in patients who were treated with trastuzumab as first-line therapy or who did not receive prior chemotherapies, PTEN loss alone was highly associated with a worse response to trastuzumab-based therapy (P = 0.028; P = 0.030). Even more significantly, PTEN loss strongly correlated with shorter OS (P = 0.008; P =0.013) in these subgroups of patients. On the other hand, PIK3CA mutations had no significant correlation with either response or OS in any subgroup. These results suggest that i) prior treatments might alter PTEN expression levels or functions; ii) prior therapies might affect the activation level of PI3K pathway, indirectly altering the PTEN status; or iii) prior therapies might modify alternative signaling pathways related to PTEN expression. These data support the notion that PTEN loss plays a central role in trastuzumab resistance, while PIK3CA mutations may not be as critical, which is different from what has been previously reported.²⁵ In addition, the mixed treatment backgrounds of the patients might be the reason for the different results in our study compared to other clinical studies on PTEN conferring trastuzumab resistance. In our study, only 55 patients' HER2-positive MBCs were analyzed for PIK3CA mutations. Therefore, interpretation of our results of PIK3CA mutations and trastuzumab resistance might be limited by the relatively small population size, and a larger cohort is needed to verify our results in the future. Our data indicated that patients with PTEN-normal tumors received the greatest benefit from first-line trastuzumab-based therapy.

PTEN loss or PIK3CA mutations direct PI3K-dependent tumorigenesis largely through activation of AKT kinase. Patients with PTEN-loss/p-AKT+ tumors or PTEN-loss/pp70S6K+ tumors had significantly shorter OS than those with PTEN-normal/p-AKT- tumors or PTEN-normal/pp70S6K- tumors, suggesting the importance of an activated PI3K pathway in trastuzumab resistance. However, the two downstream biomarkers (p-AKT, p-p70S6K) alone had limited value as indicators of trastuzumab resistance as noted in previous studies.^{27,28} Recently, it was reported that PI3K may promote cancer progression through an AKT-independent pathway, such as phosphoinositide-dependent kinase 1 pathway.⁴⁰ In our cohort, there was no correlation between PTEN loss/PIK3CA mutations and p-AKT/p-p70S6K (data not shown), suggesting that PIK3CA mutations might not be sufficient to induce AKT and p70S6K phosphorylations. It is also possible that other upstream signaling events are contributing to p-AKT/p-p70S6K. Additionally, only five cases of PIK3CA mutations were seen among 29 cases of breast cancer patients with low PTEN levels, suggesting that *PIK3CA* mutation and PTEN loss may have distinct roles in trastuzumab response. Further studies are needed to address whether the phosphoinositide-dependent kinase 1 pathway or other pathways take part in trastuzumab resistance in addition to PTEN loss and PIK3CA mutations.

Concomitant up-regulation of PI3K/AKT signaling in HER2-overexpressing breast cancers can suppress ER and PR transcriptional activity.^{22,41,42} For patients who have HER2-positive and hormone receptor-negative breast tumors, finding a biomarker to predict trastuzumab resistance is even more important. In the present study, we found a correlation between hormone receptor negative status and PTEN deficiency (P = 0.039). Moreover, in the ER-/PR- group, PTEN expression alone was significantly correlated with both poor response (P =0.029) and shorter OS after trastuzumab-based therapy (P = 0.033), whereas in the ER+/PR+ group, it was not. Our limited data are consistent with findings from other studies that loss of heterozygosity at the PTEN locus contributed to PR reduction and that PTEN functions might be different in ER+ versus ER- tumors.^{22,29,41,42} The data also suggest that abnormal expression of ER or PR might be involved in trastuzumab response. Further studies are needed to investigate whether ER and PR signaling deregulation is involved in mediating PTEN loss-induced trastuzumab resistance.

In conclusion, we have demonstrated that PI3K pathway activation (defined as PTEN loss, *PIK3CA* mutation, p-AKT+, and p-p70S6K+) contributes to trastuzumab resistance and short OS in patients with HER2-positive MBC treated by trastuzumab-based therapies. Remarkably, PTEN loss itself may serve as a predictor of resistance to trastuzumab in patients who receive trastuzumab as first-line therapy and in patients with negative hormone receptor status. Further studies are necessary to dissect the detailed mechanism of PTEN loss-mediated trastuzumab resistance and to help select the best treatment for individual patients.

Acknowledgments

We thank Sumaiyah Rehman, Dr. Shau-Hsuan Li, and Dr. Jing Lu for their valuable comments on this manuscript. We also thank Michael Worley and Alyson Todd (Department of Scientific Publication, MDACC) for their professional editing of the manuscript.

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