

TABLE 3 Nomogram for breast cancer

| Decision factors | Decision goal |
|------------------|--|
| DCIS | To determine whether SLNB is required by examining possible microinvasion ⁴⁶ |
| PST | To determine the type of surgery required by examining possible pCR ⁴⁷ |
| IBTR | To determine whether RT or re-excision is necessary ⁴⁸ |
| SLNB | To determine whether ALND is required by predicting non-SLN metastasis ^{18,49-52} |

re-excision.²⁰ For a close margin (i.e., 2–5 mm), boost irradiation can be considered.

The 20-year follow-up data from the National Surgical Breast and Bowel Project (NSABP) B-06 trial showed that 39.2% of the patients who received wide local excision without radiotherapy developed ipsilateral breast tumor recurrence (IBTR), as compared with 14.3% of those who received postoperative radiotherapy. Some believe that IBTR does not influence overall survival and that it can be considered a marker of distant metastases rather than a cause; its presence therefore cannot change the intrinsic risk of distant disease.²¹ However, according to a meta-analysis performed by the Early Breast Cancer Trialists' Collaborative Group (EBCTCG), the impact of local radiation therapy (RT) to prevent local recurrence, either to the breast following BCT or to the chest wall after mastectomy, exhibited overall survival benefit in patients with greater than 10% risk of local recurrence, but it did not show any benefit in patients with less than 10% risk of local recurrence.²² An analysis of hazard ratios for distant metastases in patients who had undergone breast-conservation surgery with or without postoperative radiotherapy indicated that local recurrence might be a cause of distant metastases.²³ These results suggested that the group with a high risk for locoregional recurrence gained a survival benefit from local radiotherapy. In addition, local relapse could be a crucial psychological stressor for a patient even if her long-term survival was unaffected.

PST

In the case of sequential chemotherapeutic regimens such as doxorubicin and cyclophosphamide (AC) followed by a taxane, the pathological complete remission (pCR) rates are higher in patients who responded to the preceding regimen than in nonresponders. Furthermore, combining chemotherapy with an anti-HER2 treatment such as trastuzumab resulted in even higher pCR rates in HER2-positive cases.²⁴ A multidisciplinary team, which included an attending surgeon, a radiologist, a medical oncologist, and a pathologist, was indispensable in making appropriate decisions regarding

BCS after PST. The findings also led to the recommendation that long-term outcome data, particularly data related to local recurrence rates, and methodologies for assessing the response and success of treatment should be collected, analyzed, and clarified at each institution.²⁵

The large majority of the attendees agreed that neoadjuvant endocrine treatment (NAET) is an acceptable approach for certain patients, including those with low-grade, estrogen receptor (ER)-positive breast cancers and postmenopausal patients. Recent studies have suggested that NAET provides higher breast-conservation rates. Nevertheless, because of a lack of randomized clinical trial data, especially on local recurrence, this issue remains to be studied with respect to the tailoring of treatment using biomarkers.²⁶ Future studies are required to investigate the factors that are predictive of a shrinkage pattern in tumors that have responded to NAET and to determine their postoperative prognosis.

Hereditary Breast Cancer

There was some consensus that patients at higher risk for local recurrence or development of breast cancer in the contralateral breast due to genetic mutations (e.g., *BRCA1* or *BRCA2*) require a more aggressive surgery than BCS. Although this is a controversial topic, the risks of IBTR and of developing contralateral breast cancer may be higher in patients with *BRCA* abnormalities. Therefore, performing a bilateral mastectomy may be preferable to BCS. In addition, performing a bilateral mastectomy would avoid use of RT in a majority of patients. Fifteen years of follow-up data from postoperative radiotherapy in *BRCA* patients suggested that there is a higher risk of radiation toxicity in these patients. Taking these data together, bilateral mastectomy for this specific subgroup could result in reducing cancer recurrence in the affected breast, decreasing new breast cancer development in the unaffected breast, and avoiding the late toxicity of radiotherapy.²⁷⁻²⁹

BCS for DCIS

The Van Nuys Prognostic Index (VNPI), originally proposed and validated by Silverstein et al., is a scoring system for predicting the risk of IBTR in DCIS patients undergoing BCS. Three major factors—margin status, high histological grade, and young age—were recognized as significant risk factors for IBTR after resection of DCIS. The distribution of the opinions as to the proper margin needed for DCIS was similar to that for invasive ductal carcinoma (IDC).

Several retrospective studies have suggested that RT after BCS is useful in avoiding IBTR, especially in patients with high-risk DCIS.^{30,31} Tamoxifen in combination with RT has

also been reported to decrease IBTR in DCIS.³² Prospective trials of neoadjuvant therapies for DCIS using trastuzumab or lapatinib have recently been initiated. These trials may elucidate the effect of anti-HER2 treatments on the local management of HER-2-neu-overexpressing DCIS.

Dunne et al. performed a meta-analysis of 4,660 cases identified from Medline with regard to the margins required for DCIS and RT. They found that a negative margin significantly reduced the risk of IBTR compared with a close margin, and a 2-mm margin was superior to a margin less than 2 mm. However, they observed no significant differences in the IBTR rates with margins over 2 mm.³³ Fisher et al. demonstrated the benefit of tamoxifen in the treatment of DCIS in NSABP B-24, a randomized controlled trial.³⁴ Because these data suggest that BCS alone is insufficient to prevent IBTR after surgery for DCIS, there was consensus at the meeting that RT and/or endocrine therapy is necessary after BCS.

RADIATION THERAPY

RT as a Component of the Local Management of Breast Cancer

Postoperative RT reduces the risk of locoregional recurrence to approximately one-third of that without RT. Although the baseline risks have varied among existing reports, depending on the method of surgery and the pathological evaluation, the relative risk reduction related to RT was consistent.²²

For each group of patients who received BCS, there have been continual efforts to find a subgroup of patients who do not require RT.^{32,35-38} Unfortunately, such a subgroup had not yet been identified in a prospective trial. However, the eligibility criteria and systemic treatment used in early clinical trials were suboptimal in comparison with today's standards.³² A clinical trial in a selected group of patients, which included individuals over 70 years old with hormone-responsive tumors treated with a suitable resection margin and appropriate hormonal therapy, demonstrated that the absolute reduction in the risk of local recurrence due to RT, although significant, was small enough that omission of RT could be considered.³² It is suggested that the intrinsic subtype of breast cancer might be an independent predictive factor related to the benefit of postoperative RT.^{39,40} At the meeting it was indicated that these findings should be verified in prospective trials.

Trends in Postoperative Irradiation for the Conserved Breast

Both hypofractionated whole-breast RT and accelerated partial-breast irradiation (APBI) were increasingly used

after BCT. Hypofractionated whole-breast RT demonstrated equivalent tumor control and cosmetic results compared with conventional fractionation.^{41,42} In the consensus conference, we discussed hypofractionation as an option for certain patients, such as those who are margin free. However, APBI is still considered an experimental treatment.

Indication for Boost to the Tumor Bed after BCT

Although a large randomized clinical trial demonstrated a significant reduction of IBTR in patients with a negative margin, we were unable to reach a consensus on the indications for an RT boost. The most important issue to be resolved was the definition of a "positive" margin after BCS. This definition varied by country and region.⁴³ Therefore, it should be further examined whether patients with positive margins benefit from routine administration of boost irradiation after whole-breast radiation therapy. At the consensus conference, approximately half of the participants responded that boost irradiation is not necessary if the margin is greater than 5 mm.

In addition to the dose dependency of the ipsilateral tumor control, the European Organisation for Research and Treatment of Cancer (EORTC) 22881-10882 trial clearly demonstrated that younger patients receive a greater benefit from boost irradiation secondary to their greater baseline risk of IBTR. However, in this consensus conference, approximately half of the participants answered that young age alone is not a sufficient criterion for providing a boost, if the margin is widely clear. To resolve this issue, we must standardize the definition of a positive margin, clarify the relationship between the distances required for a clear margin, and understand the magnitude of the effect of boost irradiation.

Survival Benefit of Postoperative RT for Breast Cancer

Meta-analyses performed by EBCTCG demonstrated that a reduction in the risk of locoregional recurrence at 5-year postoperative follow-up could eventually lead to a reduction in death from all causes at 15-year postoperative follow-up.³⁰ This survival benefit was attributed to prevention of secondary dissemination from local recurrence. However, the benefit was substantial only if the absolute risk reduction of the locoregional recurrence at 5 years exceeded 10%.

Currently, patients with four or more positive lymph nodes are regarded as being at high risk for local recurrence. Postoperative RT to the supraclavicular lymph nodes and the chest wall and breast are recommended in this group after both breast-conserving surgery and mastectomy. Furthermore, meta-analyses of existing trials have

suggested that patients with one to three positive lymph nodes might also receive a survival benefit from postoperative RT, although a randomized clinical trial investigating this hypothesis is ongoing. Patients with negative axillary lymph nodes generally exhibit a low risk of local recurrence. These patients do not benefit from such RT and may have increased risks of radiation side-effects if RT is given. Of note, the number of positive axillary lymph nodes in this context is only a surrogate for the risk of isolated locoregional recurrence. The indication for postoperative RT should ultimately be based on the absolute risk of local recurrence.

Postoperative Radiation Therapy in Patients Receiving PST

Recently, PST has been offered not only to patients with advanced disease but also to patients with early-stage breast cancer. The expansion of this practice has unveiled a new clinical question: What is the optimal RT dose for patients who respond favorably to PST? Randomized trials are needed to answer this question. However, the general consensus was that, for all patients who receive PST and BCS, postoperative RT is recommended. Retrospective studies of patients who received a mastectomy after PST showed that RT significantly improved local control even in patients with pCR after PST.⁴⁴ These investigators also found that RT improved survival in patients at higher risk of locoregional recurrence after PST and mastectomy.⁴⁵ These results provide insight that the decision to offer RT should be based on both the pretreatment assessment and the final pathologic findings. Postoperative RT is recommended for patients initially diagnosed as having a high risk of locoregional recurrence, regardless of their response to PST.

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Prognostic significance of tumor/stromal caveolin-1 expression in breast cancer patients

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Caveolin-1 (Cav-1) has been extensively characterized in cancer biological research. However, the role of Cav-1 in the interaction between tumor and stromal cells remains unclear. In the present study, we examined Cav-1 expression in tumor cells and stromal cells in breast cancer tissue by immunohistochemical analysis and evaluated its prognostic value in a training cohort. Immunohistochemical analysis of Cav-1 expression was scored as (++) , (+) or (-) according to the proportion of positively stained tumor cells (T) and stromal cells (S). Correlation analysis between tumor/stromal Cav-1 expression and clinicopathological parameters revealed that only T(++) Cav-1 status was positively associated with tumor size and histological nodal status ($P = 0.019$ and 0.021 , respectively). Univariate analysis revealed that combined T(++)/S(-) status was significantly correlated with unfavorable prognostic outcomes ($P < 0.001$). Multivariate analysis demonstrated that this combined status is an independent prognostic factor for primary breast cancer ($P = 0.002$). Clinical outcomes in different subgroups of breast cancer patients were also strictly dependent on this combined status ($P < 0.05$). The prognostic value of T(++)/S(-) Cav-1 status was also validated in the testing cohort. Collectively, our data indicate that high Cav-1 expression in tumor cells and lack of this expression in stromal cells could help identify a particular subgroup of breast cancer patients with potentially poor survival. Further studies are required to understand the regulatory mechanism of Cav-1 in the tumor microenvironment. (*Cancer Sci* 2011; 102: 1590–1596)

Breast cancer is the most common female cancer. Late-onset diagnosis, axillary lymph node metastases, tumor size, pathological type and resistance to antitumor therapy indicate a poor prognosis for breast cancer patients. Although treatment strategies for breast cancer have recently made great progress, recurrence and death rates remain unacceptably high.⁽¹⁾ Therefore, molecular biomarkers for recurrence and progression of breast cancer must be explored to help clinicians identify new diagnostic and therapeutic techniques to detect and treat breast cancer.⁽²⁾

Caveolins (Cav) are a family of scaffolding proteins that coat 50–100 nm plasma membrane invaginations. The Cav family is composed of three isoforms: Cav-1, Cav-2 and Cav-3. The Cav-1 gene is located on chromosome 7 (locus 7q31.1) and includes three exons (30, 165 and 342 bp) and two introns (1.5 and 32 kb).⁽³⁾ Cav-1 expression depends on the type of tumor and its expression is downregulated in several human cancers such as sarcoma and lung cancer and might function as a tumor suppressor.^(4,5) However, upregulation of Cav-1 expression has been reported in esophageal and pancreatic cancers and is also correlated with histopathological grade and poor prognosis.^(6,7)

Cav-1 is mainly involved in vesicular transport, cholesterol homeostasis and signal transduction.⁽⁸⁾ Furthermore, it might facilitate DNA repair and stabilize the insulin receptor against

degradation. Cav-1 also plays a negative role in cell movement,⁽⁹⁾ cellular senescence⁽¹⁰⁾ and cell growth.⁽¹¹⁾ Endothelial cells from Cav-1-/- mice exhibit a diminished response to angiogenic growth factors.⁽¹²⁾ Furthermore, Cav-1 overexpression is sufficient to induce premature cellular senescence in fibroblasts.^(13,14) Cancer-associated fibroblasts (CAFs), which are derived from malignant or normal epithelial cells, promote tumor growth.⁽¹⁵⁾ *In vitro* studies have shown that both stromal and epithelial Cav-1 play a protective role against mammary hyperplasia and tumorigenesis in breast cancer.^(11,16,17) In addition, clinical studies have indicated that stromal loss of Cav-1 is a single independent predictor of early breast cancer recurrence and progression.^(18,19) However, the value of combined tumor/stromal Cav-1 expression on the outcome of breast cancer patients is largely unknown.

In the present study, we investigated the clinical significance of Cav-1 expression (including tumor and stromal expression) in a training cohort and the correlation between tumor/stromal Cav-1 expression and clinicopathological characteristics. In addition, effects of combined tumor/stromal Cav-1 expression on outcomes in breast cancer patients were investigated. In addition, the prognostic value of combined tumor/stromal Cav-1 expression was also clarified in a testing cohort. Intriguingly, our results indicated that a counter balance of Cav-1 levels in the tumor microenvironment and epithelial compartment were the most strongly influenced clinical outcomes.

Materials and Methods

Collection of tissue samples. Tissue specimens of the training cohort were collected from the Department of Breast Surgery, Kyoto University Hospital (Kyoto, Japan) between July 2000 and February 2006. Informed consent was obtained from all patients prior to specimen collection and all study protocols were approved by the Ethics Committee for Clinical Research, Kyoto University Hospital. The clinical stage was assessed by The Japanese Breast Cancer Society classification.⁽²⁰⁾ For analysis of survival and follow up, the date of surgery was used to represent the beginning of the follow-up period. All patients who died from diseases other than breast cancer or from unexpected events were excluded from the case collection. Follow ups were terminated in June 2010. The median follow up was 74 months (range, 3–119 months). Clinicopathological parameters of the training cohort are listed in Table 1. In addition, we validated the results using an independent testing cohort of 193 consecutive patients (Table 1) who underwent surgical resection of breast cancer at Osaka Red Cross Hospital (Osaka, Japan). The protocols used in the testing group were approved by the Ethics Committee of the Osaka Red Cross Hospital. Follow ups

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Table 1. Clinicopathological parameters of the training and testing cohort

| Parameter | Variable | n (100%) |
|---------------------------|-----------------|----------|
| Training cohort | | |
| Age | >50 | 75 (72) |
| | ≤50 | 29 (28) |
| Tumor size | ≤2 cm | 24 (23) |
| | 2–5 cm | 51 (49) |
| | >5 cm | 10 (10) |
| | Unknown | 17 (18) |
| Histological nodal status | Positive | 46 (44) |
| | Negative | 56 (54) |
| | Unknown | 2 (2) |
| ER | Positive | 80 (77) |
| | Negative | 24 (23) |
| PgR | Positive | 67 (64) |
| | Negative | 37 (36) |
| HER2† | Positive | 18 (18) |
| | Negative | 81 (82) |
| ER/PgR/HER2† | Triple negative | 12 (12) |
| | Others | 92 (88) |
| Graded† | Grade 1 | 12 (12) |
| | Grade 2 | 41 (39) |
| | Grade 3 | 44 (42) |
| | Unknown | 2 (7) |
| Recurrence | Yes | 22 (21) |
| | No | 72 (69) |
| | Unknown | 10 (10) |
| Death | Yes | 7 (7) |
| | No | 97 (93) |
| Testing cohort | | |
| Age | >50 | 147 (76) |
| | ≤50 | 45 (23) |
| | Unknown | 1 (1) |
| Tumor size | ≤2 cm | 85 (44) |
| | 2–5 cm | 97 (50) |
| | >5 cm | 11 (6) |
| Histological nodal status | Positive | 79 (41) |
| | Negative | 109 (56) |
| | Unknown | 5 (3) |
| ER | Positive | 139 (72) |
| | Negative | 50 (26) |
| | Unknown | 4 (2) |
| PgR | Positive | 95 (49) |
| | Negative | 95 (49) |
| | Unknown | 3 (2) |
| HER2 | Positive | 22 (11) |
| | Negative | 170 (88) |
| | Unknown | 1 (1) |
| ER/PgR/HER2 | Triple negative | 35 (18) |
| | Others | 154 (80) |
| | Unknown | 4 (2) |
| Grade | Grade 1 | 58 (30) |
| | Grade 2 | 50 (26) |
| | Grade 3 | 85 (44) |
| Recurrence | Yes | 28 (14) |
| | No | 165 (86) |
| Death | Yes | 15 (8) |
| | No | 178 (92) |

†Patients with ductal carcinoma in situ (DCIS) were excluded. ER, estrogen receptor; HER2, human epidermal growth factor receptor 2; PgR, progesterone receptor.

in the testing group were terminated in January 2011. The median follow up in the testing cohort was 42 months (range, 1–80 months).

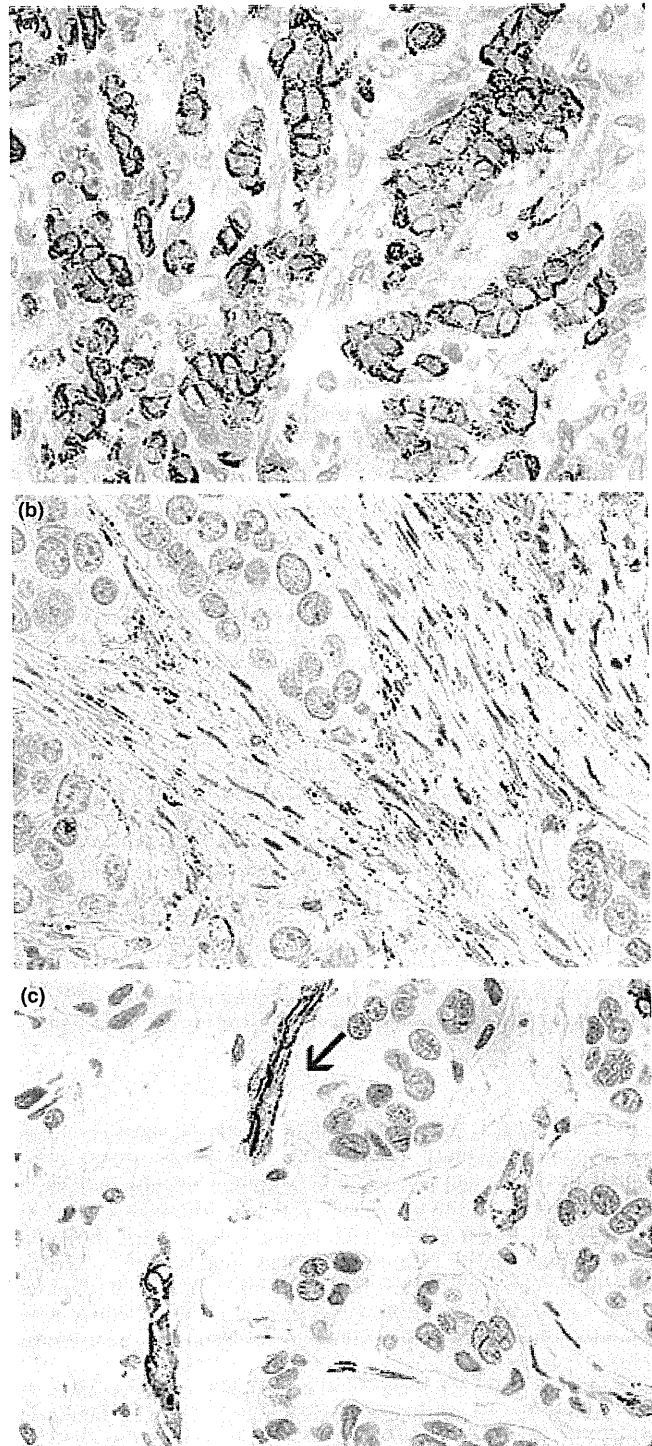


Fig. 1. Immunohistochemical analysis of caveolin-1 (Cav-1) expression (×400). (a) Tumor hot-spot expression. (b) Diffuse Cav-1 expression in stroma and (c) negative stromal Cav-1 expression. Endothelial cells indicate positive immunostaining for Cav-1 used as internal positive controls (arrows).

Immunohistochemical analysis. Immunohistochemical analysis was performed as described previously.^(18,19) In brief, slides were incubated with an anti-Cav-1 monoclonal antibody (1:800; Cell Signaling Technology, Danvers, MA, USA). The

signals were detected by Envision kit (Dako, Glostrup, Denmark) and the sections were counterstained with haematoxylin. Negative control sections were incubated with phosphate-buffered saline plus 1% bovine serum albumin instead of primary antibody. Endothelial cells were used as internal positive controls because these cells commonly express Cav-1 in cancerous regions. Results of the analyses were evaluated by two pathologists, who were independent and blinded to the clinical features of the study. We determined three hot-spots at $\times 400$ magnification, calculated the number of Cav-1-stained tumor cells (T) and stromal cells (S) and graded the cells as follows: negative expression (-), $\leq 5\%$; low expression (+), 5–50%; and high expression (++), $> 50\%$. For analyzing combined tumor/stromal Cav-1 expression, we determined tumors with a high tumor hot-spot grade that were stromal negative as T(++)/S(-).

Statistical analysis. The correlation between different types of Cav-1 expression was evaluated using Spearman's test. The correlation between Cav-1 and clinicopathological parameters was evaluated using the Kruskal–Wallis test. Disease-free survival was estimated using the Kaplan–Meier estimate and a comparison of stratified survival curves was performed using log-rank tests. Cox analysis was used to evaluate the correlation between Cav-1 and disease-free survival in the presence of various potential prognostic factors for disease-free survival. Differences were considered statistically significant at $P < 0.05$.^(18,19)

Results

Immunohistochemical analysis of Cav-1 expression in the training cohort. All patients were Japanese women and their clinicopathological characteristics are listed in Table 1. Twenty-two patients developed recurrence and seven of them died as a result. Distributions of recurrence and survival parameters are also indicated in Table 1. Table S1 summarizes the number of patients in each subgroup stratified by Cav-1 grade. T(-) Cav-1 expression was observed in 16%, T(+) Cav-1 expression was observed in 67% and T(++) Cav-1 expression was observed in 17% of breast cancer patients. Strong positive staining showed a prevalent membrane pattern associated with cytoplasm positive. For total stromal Cav-1 expression, 57% were S(+++) and 43% were S(-). In total, 72 patients were S(++/+) and 32 showed S(-) expression stratified by stromal hot-spot Cav-1 expression. For combined Cav-1 expression score grading, 5% of breast cancer patients were T(++)/S(-) and 16% were T(+)/S(-). Representative examples are illustrated in Figure 1.

Next, correlations between the different types of Cav-1 expression were analyzed. Stromal hot-spot Cav-1 expression was significantly correlated with total stromal Cav-1 expression ($R^2 = 0.517$, $P < 0.001$). However, stromal hot-spot Cav-1 expression was weakly correlated ($R^2 = 0.081$, $P = 0.003$) with tumor hot-spot expression. No significant difference was observed between total stromal and tumor hot-spot Cav-1 expression ($R^2 = 0.029$, $P = 0.083$). Because of the strong correlation between stromal hot-spot and total Cav-1 expression, we used the former as a representative for the following analysis.

Correlations between Cav-1 expression and clinicopathological parameters in the training cohort. Table 2 summarizes the correlation between Cav-1 expression and the clinicopathological parameters of breast cancer patients in the training cohort. T(+++) Cav-1 expression was positively associated with tumor size and histological nodal status ($P = 0.019$ and 0.021 , respectively). S(-) Cav-1 expression was independent of histological nodal and human epidermal growth factor receptor 2 (HER2) status ($P = 0.385$ and 0.055 , respectively). No significant correlations were found between stromal hot-spot and tumor hot-spot Cav-1 expression and age, tumor stage, grade, estrogen receptor (ER) status or progesterone receptor (PgR)

Table 2. Associations between caveolin-1 (Cav-1) expression and clinicopathological parameters in the training and testing cohorts

| Parameter | Variable | Tumor hot-spot (training cohort) | | Stromal hot-spot (training cohort) | | Tumor/stromal Cav-1 expression (training cohort) | | Tumor/stromal Cav-1 expression (testing cohort) | | P |
|-------------------------------|-----------|----------------------------------|-----------------|------------------------------------|-----------------|--|----------------|---|-----------------|--------|
| | | T(+/-) | T(++) | S(-) | S(++/++) | T(++)/S(-) | Others | T(++)/S(-) | Others | |
| Tumor size | ≤ 50 | 27.0 \pm 12.8 | 36.3 \pm 17.4 | 30.5 \pm 13.1 | 28.2 \pm 14.6 | 36.6 \pm 6.9 | 20.9 \pm 3.3 | 40.9 \pm 45.4 | 23.8 \pm 12.4 | <0.001 |
| Age (%) | > 50 | 24 (28) | 5 (28) | 10 (31) | 19 (26) | 1 (20) | 28 (28) | 13 (59) | 134 (79) | 0.129 |
| T status (%) | T0/T1 | 62 (72) | 13 (72) | 22 (69) | 53 (74) | 4 (80) | 71 (72) | 9 (41) | 36 (21) | 0.033 |
| | T2/T3/T4 | 24 (34) | 2 (12) | 5 (21) | 21 (33) | 0 (0) | 26 (31) | 5 (23) | 80 (47) | 0.002 |
| Histological nodal status (%) | Positive | 47 (66) | 14 (88) | 19 (79) | 42 (67) | 3 (100) | 58 (69) | 17 (77) | 91 (53) | 0.113 |
| | Negative | 34 (40) | 12 (71) | 16 (52) | 30 (42) | 5 (100) | 41 (42) | 5 (25) | 104 (62) | 0.208 |
| Grade† (%) | Grade 1 | 51 (60) | 5 (29) | 15 (48) | 41 (58) | 0 (0) | 56 (58) | 15 (75) | 64 (38) | 0.360 |
| | Grade 2 | 12 (15) | 0 (0) | 2 (7) | 10 (15) | 0 (0) | 12 (13) | 4 (18) | 54 (32) | 0.471 |
| | Grade 3 | 31 (39) | 10 (59) | 11 (37) | 30 (45) | 4 (80) | 37 (40) | 5 (23) | 45 (26) | 0.197 |
| ER (%) | Positive | 69 (80) | 7 (41) | 17 (56) | 27 (40) | 1 (20) | 43 (47) | 13 (59) | 72 (42) | 0.055 |
| | Negative | 17 (20) | 11 (61) | 21 (66) | 59 (82) | 3 (60) | 77 (78) | 13 (62) | 126 (75) | 0.817 |
| PgR (%) | Positive | 58 (67) | 9 (50) | 19 (59) | 48 (67) | 3 (60) | 64 (65) | 8 (38) | 42 (25) | 0.113 |
| | Negative | 28 (33) | 9 (50) | 13 (41) | 24 (33) | 2 (40) | 35 (35) | 11 (52) | 84 (50) | 0.201 |
| HER2† (%) | Positive | 15 (19) | 3 (17) | 9 (30) | 9 (13) | 2 (40) | 16 (17) | 3 (29) | 19 (33) | 0.667 |
| | Negative | 66 (81) | 15 (83) | 21 (70) | 60 (87) | 3 (60) | 78 (83) | 18 (71) | 152 (67) | 0.002 |

†Patients with ductal carcinoma in situ (DCIS) were excluded. ER, estrogen receptor; HER2, human epidermal growth factor receptor 2; PgR, progesterone receptor; S, stromal hot-spot; T, tumor hot-spot.

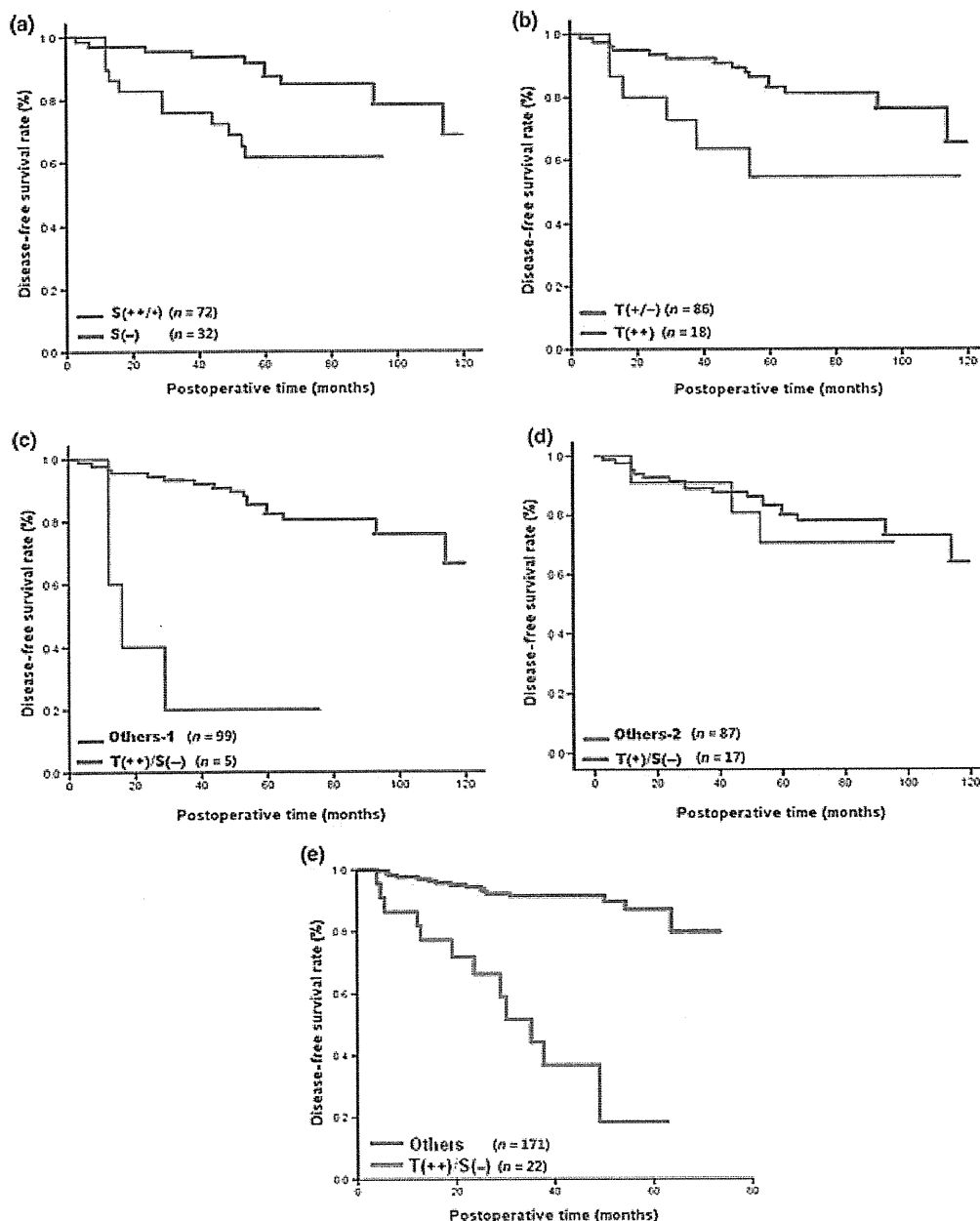


Fig. 2. Disease-free survival curves of the training cohort stratified by (a) stromal hot-spot caveolin-1 (Cav-1) expression status: S(-) vs S(++/+); $P = 0.009$, log-rank test); (b) tumor hot-spot Cav-1 expression status: T(++/+) vs T(+/-); $P = 0.019$, log-rank test); (c) combined Cav-1 expression: T(++)/S(-) vs Others 1 ($P < 0.001$, log-rank test); and (d) combined Cav-1 expression: T(+)/S(-) vs Others 2 ($P = 0.662$, log-rank test). Disease-free survival curves of the testing cohort stratified by (e) combined Cav-1 expression: T(++)/S(-) vs Others ($P < 0.001$, log-rank test).

status ($P > 0.05$). T(++)/S(-) Cav-1 expression was positively associated with tumor size and histological nodal status ($P < 0.001$ and 0.012 , respectively). No significant correlations were found between T(++)/S(-) Cav-1 expression and age, tumor stage, grade, ER, PgR or HER2 status ($P > 0.05$).

Prognostic value of Cav-1 expression in the training cohort. Figure 2a–d illustrates the Kaplan–Meier curves of disease-free survival for the training cohort constructed on the basis of the Cav-1 expression level. T(++), S(-) and T(++)/S(-) status correlated closely with poor disease-free survival ($P = 0.009$, 0.019 and < 0.001 , respectively). No major differences were observed between S(+) and S(++/+) or between T(+)

and T(-) Cav-1 expression on the predictive value of disease-free survival ($P > 0.05$; data not shown). T(+)/S(-) status also did not influence disease-free survival.

Kaplan–Meier analysis demonstrated a significant impact of certain clinicopathological prognostic factors such as HER2, tumor stage and ER on disease-free survival ($P = 0.015$, 0.010 and 0.010 , respectively). No significant correlations were found between disease-free survival and other clinicopathological factors, including PgR, histological nodal status, grade and age ($P > 0.05$; Table 3). Cox analysis was performed to evaluate whether the correlation between Cav-1 expression and disease-free survival was related to the correlation of Cav-1

Table 3. Univariate analyses of factors associated with recurrence in the training and testing cohorts

| Variable | Disease-free survival (<i>P</i>) |
|--|------------------------------------|
| Training cohort | |
| Tumor/stromal Cav-1 expression T(++)/S(-) vs others | <0.001 |
| Stromal hot-spot Cav-1 expression S(++/+) vs S(-) | 0.009 |
| Tumor stage (T0/T1 vs T2/T3/T4) | 0.010 |
| ER (positive vs negative) | 0.010 |
| HER2 (positive vs negative) | 0.015 |
| Tumor hot-spot Cav-1 expression T(++)/S(-) vs T(+/-) | 0.019 |
| PgR (positive vs negative) | 0.054 |
| Histological nodal (positive vs negative) | 0.069 |
| Age (>50 vs ≤50) | 0.411 |
| Testing cohort | |
| Tumor/stromal Cav-1 expression T(++)/S(-) vs others | <0.001 |
| Histological nodal status (positive vs negative) | <0.001 |
| Grade (1 vs 2 vs 3) | <0.001 |
| HER2 (positive vs negative) | <0.001 |
| Tumor stage (T0/T1 vs T2/T3/T4) | 0.008 |
| ER (positive vs negative) | 0.020 |
| PgR (positive vs negative) | 0.054 |
| Age (>50 vs ≤50) | 0.376 |

Cav-1, caveolin-1; ER, estrogen receptor; HER2, human epidermal growth factor receptor 2; PgR, progesterone receptor; S, stromal hot spot; T, tumor hot spot.

Table 4. Multivariate analysis of factors that might affect disease-free survival in the training and testing cohorts

| | Disease-free survival | | |
|-----------------------------------|-----------------------|--------------|----------|
| | HR | 95% CI | <i>P</i> |
| Training cohort | | | |
| Stromal hot-spot Cav-1 expression | 0.322 | 0.098–1.065 | 0.063 |
| ER | 1.092 | 0.306–3.901 | 0.893 |
| HER2 | 3.362 | 1.035–10.923 | 0.044 |
| Tumor stage | 7.772 | 0.985–61.288 | 0.052 |
| Histological nodal status | 1.536 | 0.900–2.619 | 0.115 |
| Grade | 3.807 | 1.018–7.472 | 0.021 |
| Tumor hot spot-Cav-1 expression | 0.370 | 0.108–1.269 | 0.114 |
| ER | 0.844 | 0.248–2.867 | 0.785 |
| HER2 | 4.778 | 1.337–17.081 | 0.016 |
| Tumor stage | 6.437 | 0.806–51.392 | 0.079 |
| Histological nodal status | 1.406 | 0.814–2.430 | 0.222 |
| Grade | 0.322 | 1.220–11.881 | 0.046 |
| Tumor/stromal Cav-1 expression | 0.041 | 0.006–0.297 | 0.002 |
| ER | 0.774 | 0.232–2.580 | 0.677 |
| HER2 | 3.665 | 1.037–12.954 | 0.044 |
| Tumor stage | 7.234 | 0.900–58.129 | 0.063 |
| Histological nodal status | 1.327 | 0.753–2.341 | 0.328 |
| Grade | 5.868 | 1.523–22.614 | 0.010 |
| Testing cohort | | | |
| Tumor/stromal Cav-1 expression | 0.249 | 0.107–0.582 | 0.001 |
| ER | 0.846 | 0.362–1.982 | 0.701 |
| HER2 | 2.141 | 0.875–5.238 | 0.096 |
| Tumor stage | 1.551 | 0.965–2.494 | 0.070 |
| Histological nodal status | 4.124 | 1.517–11.208 | 0.005 |
| Histological grade | 3.150 | 1.416–7.007 | 0.005 |

Cav-1, caveolin-1; CI, confidence interval; ER, estrogen receptor; HER2, human epidermal growth factor receptor 2; HR, hazard ratio.

expression with other prognostic factors. The results revealed that S(-) Cav-1 expression was not an independent prognostic factor for disease-free survival ($P = 0.063$). No significant correlation was found between T(++)/S(-) Cav-1 expression and disease-free survival ($P = 0.114$). Consistent with the univariate analysis results, the multivariate analysis revealed that combined T(++)/S(-) status was strongly associated with an unfavorable prognosis ($P = 0.002$; Table 4). We conducted further subgroup analysis stratified by histological nodal, ER and HER2 status, which were associated with poor clinical outcomes. As a result, S(-) Cav-1 showed clear trends for predicting disease-free survival in histological node+ patients ($P = 0.008$). However, T(+) Cav-1 expression did not influence disease-free survival in histological node+ patients ($P = 0.088$). Notably, histological node+ patients with T(++)/S(-) Cav-1 expression exhibited lower disease-free survival ($P = 0.001$). Furthermore, T(++)/S(-) Cav-1 expression also served as an important predictor of disease-free survival for ER+ and HER2+ patients ($P = 0.001$ and 0.045 , respectively). Moreover, patients with T(++)/S(-) Cav-1 expression who were in the ER-, HER2-, PgR (+ and -), tumor size (>5 and ≤5 cm), age (>50 and ≤50 years) and grade (Grade 2 and 3) subgroups had poorer disease-free survival (Table S2).

Validation of the prediction power of T(++)/S(-) Cav-1 status in the testing cohort. In the testing cohort, all patients were also Japanese women and their clinicopathological characteristics are listed in Table 1. T(++)/S(-) Cav-1 expression was observed in 11% and other expressions were observed in 89% breast cancer patients. T(++)/S(-) Cav-1 expression was significantly related to tumor size ($P < 0.001$), tumor stage ($P = 0.033$) and histological nodal status ($P = 0.002$). We could not find a statistically significant association between T(++)/S(-) Cav-1 expression and age, grade, ER, PgR or HER2 ($P > 0.05$; Table 2).

We then examined the association of T(++)/S(-) Cav-1 expression with the clinical outcome. Kaplan–Meier survival analysis showed patients with T(++)/S(-) Cav-1 expression had shorter disease-free survival than those with other expressions ($P < 0.001$, Fig. 2e). The significant impact of clinicopathological prognostic factors such as tumor stage, ER and HER2 on disease-free survival ($P = 0.008$, 0.020 and <0.001 , respectively) was also validated. These results were consistent with the above findings. In addition, a significant correlation between disease-free survival and grade or histological nodal status was observed in the testing cohort ($P < 0.001$ and <0.001 , respectively; Table 3).

Table 4 provides the multivariate analyses of factors related to patient disease-free survival. Cox analysis indicated that T(++)/S(-) Cav-1 status was an independent predictor of disease-free survival ($P = 0.001$), as were histological nodal status ($P = 0.005$) and grade ($P = 0.005$). Moreover, the role of T(++)/S(-) Cav-1 expression in disease-free survival in the ER (+ and -), HER2 (+ and -), PgR (+ and -), tumor size (>5 and ≤5 cm), age (>50 and ≤50 years) and grade (Grade 1 + 2 and 3) subgroups is shown in Table S2.

Discussion

The Cav-1 gene is colocalized at the D7S522 locus on human chromosome 7q31.1 and is commonly deleted in breast, colon, kidney, prostate, ovary, head and neck cancers. Thus, it seems feasible to propose that the Cav-1 gene might serve as a candidate tumor suppressor gene.⁽²¹⁾ In the present study, we focused on breast cancer patients to determine the correlation of tumor/stromal Cav-1 expression with clinicopathological parameters and survival. Cav-1 expression was evaluated semi-quantitatively based on the proportion of positively stained tumor and stromal cells. We found that tumor Cav-1 demonstrated a prevalent membrane pattern associated with cytoplasm positive and that T(-) Cav-1 expression was noted in 16% of

cases, which is consistent with previous reports.⁽²²⁾ The stromal Cav-1 expression rate and pattern in the present study are similar to a previous report.⁽²³⁾

According to previous reports, total tumor Cav-1 expression has no prognostic value in primary breast cancer patients.^(18,24) In the present study, we examined stromal hot-spot, total stromal and tumor hot-spot Cav-1 expression. First, we analyzed correlations between stromal hot-spot and total stromal and tumor Cav-1 expression and found a weak but significant correlation between stromal and tumor expression, indicating that Cav-1 expression could be regulated differently between tumor and stromal cells and that Cav-1 might influence different functions in those cells.^(25,26) T(+++) Cav-1 expression was positively associated with tumor size and histological nodal status. Previous reports revealed that tumor Cav-1 expression was negatively associated with HER2 status.⁽²⁷⁻²⁹⁾ Several studies have indicated that Cav-1 might function as a negative signal transduction regulator to HER2/neu and that it might play a negative regulatory role in mammary tumor development. In addition, activation of HER2/neu might downregulate Cav-1 expression *in vitro*.^(30,31) However, this finding was not supported by our results. Stromal Cav-1 expression showed no significant correlation with any of the clinicopathological parameters, which was inconsistent with a previous report,⁽¹⁸⁾ and therefore we focused on combined tumor/stromal Cav-1 expression. T(+++)/S(-) was observed frequently in large-size tumors and histological node+ cases, indicating that tumor/stromal Cav-1 expression is involved in breast cancer progression. Interestingly, survival analyses revealed that patients with T(+++)/S(-) Cav-1 expression had the shortest disease-free survival among various Cav-1 expression subgroups. Multivariate analysis confirmed an independent prognostic value of the combined status. Consistent with these results, T(+++)/S(-) Cav-1 expression was significantly related to tumor size, histological nodal status and disease-free survival in the testing cohort. Besides, the positive correlation between T(+++)/S(-) Cav-1 expression and tumor stage was also indicated in the testing cohort. A possible explanation for the discrepancy could be due to the difference in sample size. Furthermore, T(+++)/S(-) Cav-1 expression also impacted the clinical outcomes stratified by ER status, PgR status, HER2 status, tumor size, age, histological nodal status and grade in both the training and testing cohorts. Therefore, these results indicate that combined Cav-1 status has a more potent prognostic value than either stromal or tumor hot-spot alone. We believe that these results are important when considering breast cancer biology. Given the limited number of cases, prospective studies with long-term follow-up data are warranted.⁽³²⁾

Tumor Cav-1 expression with respect to tumorigenesis seems more complex than originally believed. Cav-1 loss-of-function induces ligand-independent hyperactivation of Ras-p42/44 MAPK and Smad signaling pathways as well as enhanced matrix metalloproteinase-2/9 secretion. Each of these pathways is likely to contribute to cell cycle progression, growth factor independence, cell invasiveness and epithelial-mesenchymal transition.⁽³³⁾ Despite extensive evidence supporting the role of Cav-1 as a tumor suppressor, several studies have suggested an alternative view of Cav-1 expression in tumors. In breast cancer, Cav-1 protects tumor cells from anoikis, promotes tumor cell survival and abrogates detachment-induced p53 activation.^(34,35) Furthermore, Cav-1 expression is upregulated in multidrug-resistant MCF-7 cells.^(36,37) A hypothesis has been proposed to

explain the divergent roles of Cav-1; even if an initial loss of Cav-1 is observed in breast cancer, re-expression of Cav-1 at later stages might correlate with more malignant characteristics.^(35,38)

Stromal Cav-1 plays a vital role in tumorigenesis. Loss of stromal Cav-1 is an independent predictor for therapeutic resistance and poor prognosis in primary breast cancers.^(12,18,19,39) Woodman *et al.*⁽¹²⁾ reported that endothelial cells from Cav-1-/- mice exhibit a disrupted response to angiogenic growth factors. Senescent human diploid fibroblasts exhibit increased levels of the Cav-1 protein.⁽¹⁰⁾ In addition, loss of Cav-1 in stromal cells of various organs directly leads to disorganised stromal compartments and dysfunctional organ systems.⁽⁴⁰⁾

Furthermore, recent studies have revealed a role played by Cav-1 in the interaction between tumor and stromal cells in breast cancer. During tumor formation, cancer cells and adjacent fibroblasts are metabolically coupled. A new model has been proposed in which glycolytic CAF promote tumor growth by secreting energy-rich metabolites that can be taken up by adjacent tumor cells.⁽⁴¹⁾ Loss of Cav-1 *in vitro* induces metabolic coupling between CAF and tumor cells and leads to the formation of a host-parasite relationship. Martinez-Outschoorn *et al.*⁽⁴²⁾ showed that Cav-1 expression is downregulated in fibroblasts co-cultured with MCF-7 cells and that it mediates autophagic/lysosomal degradation. Furthermore, autophagy induced by loss of Cav-1 in fibroblasts provides cancer cells with essential chemical building blocks.^(42,43) Loss of stromal Cav-1 fibroblasts protects adjacent cancer cells via decreased apoptosis, increased TP53-induced glycolysis and apoptosis regulator expression.⁽⁴⁴⁾ Furthermore, loss of Cav-1 induces oxidative stress in CAF, which is the root cause of mitochondrial dysfunction in CAF and promotes DNA damage. In the present study, the predictive value of T(+++)/S(-) was demonstrated in luminal-type cancers and HER2+ cancers. Its value was stratified by an intrinsic subtype and warrants an examination with a greater number of cases.

The regulatory mechanism of Cav-1 expression in breast cancer remains to be elucidated. Pro-autophagic stimuli such as hypoxia, oxidative stress and nuclear factor κB activation might cause the loss of Cav-1.⁽⁴⁵⁾ Conversely, multiple factors are present during Cav-1 upregulation.

In conclusion, we provide evidence that T(+++)/S(-) Cav-1 expression is closely associated with unfavorable prognostic outcomes in primary breast cancer patients. This particular subgroup seems to be engaged in rapid disease progression. Further studies involving analysis of molecular mechanisms of Cav-1 expression are required. The interaction between tumor and stromal cells and Cav-1 in the tumor microenvironment is also a key issue to investigate. Moreover, new therapies targeting Cav-1 expression might be a novel therapeutic approach, particularly for patients with T(+++)/S(-) Cav-1 status.

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Disclosure Statement

The authors have no conflict of interest.

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Supporting Information

Additional Supporting Information may be found in the online version of this article:

Table S1. The numbers of patients in each subgroup stratified by Cav-1 grade.

Table S2. Univariate analyses of Cav-1 expression associated with disease-free survival of the training and testing cohort.

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Docetaxel Followed by Fluorouracil/Epirubicin/Cyclophosphamide as Neoadjuvant Chemotherapy for Patients with Primary Breast Cancer

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Objective: This multicenter, open-label, single-arm, Phase II study assessed the efficacy of a neoadjuvant chemotherapy with docetaxel (75 mg/m² q3w) followed by 5-fluorouracil 500 mg/m², epirubicin 100 mg/m² and cyclophosphamide 500 mg/m² q3w in patients with early-stage breast cancer.

Methods: Women with resectable breast cancer (T1c–3 N0 M0 or T1–3 N1 M0) were enrolled. Before surgery, patients received four cycles of docetaxel followed by four cycles of 5-fluorouracil, epirubicin, and cyclophosphamide. The primary endpoint was the pathological complete response (pCR) rate defined for the breast alone, assessed by a central review committee. Secondary endpoints included clinical response and safety.

Results: One hundred and thirty-seven patients were enrolled. Of the 132 patients assessable for pathologic response, 23% (95% confidence interval, 16–31%) experienced a pathological complete response and 6% (95% confidence interval, 3–12%) had a near pathological complete response (few remaining cancer cells), resulting in a quasi-pathological complete response of 29% (95% confidence interval, 21–37%). Clinical response rate following the initial docetaxel regimen was 64%. The overall clinical response rate after completion of 5-fluorouracil, epirubicin, and cyclophosphamide was 79%; breast-conserving surgery was performed in 79% of patients. More patients with triple-negative disease (estrogen/progesterone receptors negative; human epidermal growth factor 2 negative) experienced a pathological complete response [14/29, (48%); 95% confidence interval, 29–68%] versus those with other molecular subtypes. The safety profile was acceptable.

Conclusions: Eight cycles of neoadjuvant chemotherapy—docetaxel followed by 5-fluorouracil, epirubicin, and cyclophosphamide—are tolerable and conferred high rates of pathological complete response and breast-conserving surgery. Patients with triple-negative disease were more likely to achieve pathological complete response versus other subtypes, suggesting that selecting appropriate neoadjuvant chemotherapy based on molecular subtype could be possible.

Key words: breast neoplasms – neoadjuvant therapy – FEC protocol – docetaxel

INTRODUCTION

Neoadjuvant chemotherapy has been widely used for patients with operable breast cancer to increase the chance of breast conservation (1–7). Furthermore, response to neoadjuvant treatment can provide important information on long-term survival outcomes. Pathological complete response (pCR) in the breast and axillary lymph nodes predicts a favorable prognosis, whereas a lack of pCR in the breast and node-positive status do not (6,7). This implies the possibility of tailoring subsequent treatment according to the response to initial treatment (7–12). In addition, correlative studies of tumor samples before and after treatment may provide information on markers that could predict response or resistance to treatment (13–16).

Results from the National Surgical Adjuvant Breast and Bowel Project (NSABP) Protocol B-18 trial demonstrated the impact of neoadjuvant chemotherapy in patients with operable early-stage breast cancer (17). The protocol-specified anthracycline-containing regimen—four cycles of doxorubicin and cyclophosphamide (AC)—resulted in an increased likelihood of breast-conserving surgery (BCS) compared with no neoadjuvant chemotherapy. The study established pCR as a prognostic marker for long-term disease-free survival (DFS) and demonstrated that there was no difference in survival if chemotherapy was administered before or after surgery. Subsequent studies, such as the Aberdeen trial, have demonstrated the benefit of the sequential addition of taxanes to neoadjuvant anthracycline regimens (5). The NSABP Protocol B-27 trial demonstrated that, compared with neoadjuvant AC alone, the addition of sequential docetaxel doubled the pCR rate, increased the clinical complete response rate (RR) and increased the proportion of patients with negative axillary nodes (7–18).

We previously conducted a Phase II study to evaluate the clinical and pathological response and safety of the FEC regimen (5-fluorouracil, epirubicin and cyclophosphamide) followed by docetaxel as neoadjuvant chemotherapy in Japanese women with early-stage breast cancer [Japan Breast Cancer Research Group (JBCRG) 01 trial]. The results of this study have been reported previously (19). Although the pCR rate was 16% and BCS was possible for 85% of patients, there were some safety concerns, with 18% of patients experiencing febrile neutropenia and 41% of patients experiencing Grade 1/2 peripheral edema (no Grade 3/4 events observed) following the docetaxel regimen (unpublished data). Disease progression occurred in 6% of patients after the completion of all planned treatment (unpublished data).

In an effort to achieve a higher pathological RR with an improved safety profile, we decided to evaluate the efficacy and safety of docetaxel followed by FEC (JBCRG 03 trial)—the reverse of the sequence of chemotherapy used in the JBCRG 01 trial (19). The clinical and pathological effects and the toxicity profile of this regimen are presented here, and the results of predictive marker analyses are discussed.

PATIENTS AND METHODS

PATIENT ELIGIBILITY

This was a multicenter, open-label, single-arm, Phase II study that recruited patients via central registration. Japanese women aged 20–59 years with histologically proven early-stage breast cancer (T1c–3 N0 M0 or T1–3 N1 M0) were enrolled. No prior chemotherapy, radiotherapy, hormonal therapy or immunotherapy was allowed. Other inclusion criteria were Eastern Cooperative Oncology Group performance status 0–1; white blood cell count 4000–12 000/mm³; neutrophil count \geq 2000/mm³; platelet count \geq 100 000/mm³; hemoglobin \geq 9.5 g/dl; serum bilirubin \leq 1.25 times upper limit of normal (ULN); creatinine \leq 1.5 times ULN and aspartate aminotransferase and alanine aminotransferase \leq 1.5 times ULN. Patients with congestive heart failure or left ventricular ejection fraction \leq 60% were excluded. Patients were also excluded if they had confirmed infection; serious concomitant illness such as severe cardiovascular disease, uncontrolled diabetes, malignant hypertension or hemorrhagic disease; active concomitant malignancy; brain metastasis; peripheral neuropathy; history of edema with severe drug allergy; or previous long-term corticosteroid therapy. Pregnant or lactating women were excluded. Mammography, ultrasonography, magnetic resonance imaging or computed tomography was used to assess the presence of tumors. Baseline evaluations included complete blood cell and platelet count, routine blood chemistry and liver function tests, chest X-ray, bone scan, electrocardiogram and echocardiogram.

The local ethics committee or institutional review board approved the study at each institution. All patients gave written informed consent to participate. The protocol was conducted in accordance with the Declaration of Helsinki and Good Clinical Practice.

TREATMENT

Four cycles of docetaxel (75 mg/m²) administered intravenously (i.v.) every 21 days were followed by four cycles of FEC (5-fluorouracil 500 mg/m², epirubicin 100 mg/m² and cyclophosphamide 500 mg/m²) administered i.v. on Day 1 every 21 days before surgery. Premedication was administered based upon each physician's decision to prevent edema, nausea and allergic reactions (e.g. dexamethasone 12 mg i.v. and/or granisetron 4 mg i.v. on Day 1, and oral dexamethasone 8 mg on Days 2 and 3 of docetaxel treatment; dexamethasone 24 mg i.v. on Day 1 and oral dexamethasone 8 mg on Days 2–6 with the FEC regimen). Administration of granulocyte colony-stimulating factor and antibiotics was left to the judgment of each investigator.

CLINICAL RESPONSE ASSESSMENT

Tumor assessments were performed within 4 weeks before docetaxel treatment, after completion of docetaxel treatment

and before surgery. Tumor response was assessed using the modified Response Evaluation Criteria in Solid Tumors guidelines (in which confirmatory scans/assessments were not required due to the timing of surgery), for patients who had measurable lesions.

CENTRAL PATHOLOGIC ASSESSMENT

Hematoxylin and eosin-stained slides were prepared from core needle biopsy and surgical specimens from the primary tumor. All surgical specimens were cut in 5 mm interval and all surfaces were microscopically examined in each institution. Pathological response of chemotherapy was assessed by a central review committee consisting of three pathologists who used criteria established by the Japanese Breast Cancer Society. pCR was defined as necrosis and/or disappearance of all tumor cells, and/or the replacement of cancer cells by granulation and/or fibrosis. If only ductal components remained, the pathological response was described as a pCR. Near pCR was defined as extremely high grade marked changes approaching a complete response, with only a few remaining isolated cancer cells (19). Quasi-pCR (QpCR) was the total of both pCR and near pCR. The central review committee evaluated the pathological responses independently from local pathologists. This committee was blinded to the local pathologists' reports. Patients who did not have surgery because of disease progression were considered not to have a pCR.

HORMONE RECEPTOR AND HUMAN EPIDERMAL GROWTH FACTOR 2 OVEREXPRESSION

Estrogen receptor (ER) and progesterone receptor (PgR) status was determined by immunohistochemistry (IHC) before docetaxel treatment at each participating institute. In general, tumors with more than 10% positively stained tumor cells were classified as positive for ER and PgR. The human epidermal growth factor 2 (HER2) status of the tumor was also determined at each institute by IHC or by fluorescence *in situ* hybridization (FISH) analysis. HER2-positive tumors were defined as those scoring 3+ with IHC staining or testing positive by FISH. HER2-negative tumors were defined as those scoring 0–1+ with IHC or scoring 2+ with IHC and testing negative by FISH.

SURGERY AND RADIOTHERAPY

Following chemotherapy and clinical assessment of response, patients underwent surgery. If the tumor was too large or invasive for BCS, a modified radical mastectomy was recommended. Careful pathological assessment of tumor margins was performed in accordance with the Japanese Breast Cancer Society criteria (20). Sentinel lymph node biopsy was performed to confirm disease stage or to avoid surgical axillary dissection. Autologous or heterologous reconstructive surgery was performed depending on the

patient's requirements and health status. All patients who underwent BCS were given standard radiotherapy to the remaining ipsilateral breast tissue after surgical recovery. For patients diagnosed as sentinel node negative and thus not requiring axillary dissection, radiotherapy to the axilla was allowed.

TOXICITY AND DOSE MODIFICATION

Toxicities were evaluated according to the National Cancer Institute Common Terminology Criteria for Adverse Events (version 3) throughout treatment with docetaxel and FEC before surgery. Treatment could be postponed for a maximum of 2 weeks only for severe toxicity. If the adverse event (AE) did not improve during this period, chemotherapy was discontinued and surgery was recommended. Dose reductions were permitted for docetaxel from 75 to 60 mg/m² and for epirubicin from 100 to 75 mg/m² in cases of febrile neutropenia or Grade 3/4 non-hematologic toxicities, except for nausea, vomiting and fatigue.

STATISTICAL METHODS

The primary endpoint was the pCR rate. Before the initiation of the current study, the pCR rate for non-taxane anthracycline regimens ranged from 12.8% (NSABP Protocol B-27) (18) to 15.4% (Aberdeen trial) (5). Previously, we had conducted JBCRG01 trial to evaluate the pCR rate defined for breast disease (19). Therefore, in order to detect improvement in the pCR rate in the same definition of our previous study, a sample of 119 patients was required according to binominal distribution, with a one-sided threshold pCR rate of 12%, an expected pCR rate of 22%, an α error of 5% and a β error of 10%. The target number of patients for recruitment was therefore 119, so assuming that 5% of patients would not be evaluable, we planned to enroll 130 patients. Secondary endpoints included safety, clinical RR, rate of BCS, DFS, overall survival and a subset analysis according to biomarkers. Pathological and clinical RRs were calculated with 95% confidence intervals (95% CIs), with each complete RR based on a binominal distribution. Pathological response was evaluated by hormone receptor status and HER2 status. A multiple logistic regression analysis was performed to examine which factors (menopausal status, tumor size, ER and PgR status, HER2 status and clinical response to docetaxel and FEC) were associated with pCR and QpCR.

RESULTS

PATIENTS CHARACTERISTICS AND TREATMENT

Enrollment took place from October 2005 through October 2006. One hundred and thirty-seven patients were enrolled. Two patients did not receive study treatment because of early withdrawal of consent; therefore, 135 patients were evaluable for safety and clinical response. These evaluable

Table 1. Patients' characteristics

| Characteristic | Value ^a |
|---|--------------------|
| Number of evaluable ^b patients | 135 |
| Age (years) | |
| Median | 46 |
| Range | 24–62 |
| Performance status, <i>n</i> (%) | |
| 0 | 133 (99) |
| 1 | 2 (1) |
| Menopausal status, <i>n</i> (%) | |
| Premenopausal | 94 (70) |
| Postmenopausal | 41 (30) |
| Clinical tumor stage, <i>n</i> (%) | |
| T1 | 13 (10) |
| T2 | 98 (73) |
| T3 | 24 (18) |
| Clinical nodal stage, <i>n</i> (%) | |
| N0 | 62 (46) |
| N1 | 73 (54) |
| ER status, <i>n</i> (%) | |
| Positive | 86 (64) |
| Negative | 46 (34) |
| Unknown | 3 (2) |
| PgR status, <i>n</i> (%) | |
| Positive | 63 (47) |
| Negative | 70 (52) |
| Unknown | 2 (1) |
| HER2 status, ^c <i>n</i> (%) | |
| 0 | 21 (16) |
| 1+ | 63 (47) |
| 2+ | 20 (15) |
| 3+ | 31 (23) |

ER, estrogen receptor; HER2, human epidermal growth factor receptor 2; PgR, progesterone receptor.

^aPercentages may not add up to 100% because of rounding.

^bNumber of patients evaluable for safety and clinical response.

^cEvaluated by immunohistochemistry.

patients included two patients aged 60 and 62 years (included because their age was not considered to influence the evaluation). Two patients were lost to follow-up before surgery, thus 133 patients were evaluable for surgical response. A total of 132 patients were evaluable for pathological response; one patient was excluded owing to lack of confirmation of invasive carcinoma (following the pathological central review) due to inadequate samples from core needle biopsy before study treatment.

The patient characteristics are summarized in Table 1. Thirty patients (22%) had triple-negative disease, defined as

ER-negative, PgR-negative and HER2-negative primary breast cancer, including one patient who was lost to follow-up before surgery.

Overall, 98 patients (73%) completed the planned eight cycles of treatment without dose reductions or study discontinuation. A total of 115 (85%) and 106 (82%) patients completed all four planned treatment cycles of docetaxel and FEC, respectively; dose reductions were necessary in 9 (7%) and 17 (13%) patients, respectively. The majority of the dose reductions were attributable to toxicities, particularly febrile neutropenia during treatment with FEC (10 versus 2 patients during docetaxel treatment). Dose reductions due to neutropenia were required by three patients each during the docetaxel and FEC regimens. Eleven (8%) and six patients (5%), respectively, discontinued treatment during docetaxel and FEC therapy because of toxicities (five patients discontinued during both regimens) or disease progression (six patients during docetaxel and one patient during FEC). The mean dose intensities were 24.2 and 30.3 mg/m²/week for docetaxel and epirubicin, respectively.

TOXICITIES

The incidence of treatment-related AEs is summarized in Table 2. Neutropenia was the most common Grade 3/4 treatment-related AE and was observed in 44% and 60% of patients during docetaxel and FEC therapy, respectively. Overall, 67% and 15% of patients experienced at least one episode of Grade 3/4 neutropenia or febrile neutropenia, respectively. For non-hematologic toxicities of any grade, rash, sensory neuropathy, edema, muscle pain and joint pain occurred more frequently during docetaxel treatment than with FEC. Conversely, the frequency of gastrointestinal symptoms, such as nausea, vomiting and anorexia, was higher with FEC than with docetaxel. The frequency of Grade 1/2 peripheral edema was similar during exposure to docetaxel (33%) and FEC (29%); no patient had Grade 3/4 edema. Grade 3/4 non-hematologic toxicities, including gastrointestinal disturbances, were infrequent during both docetaxel and FEC. No fatal AEs were reported.

CLINICAL RESPONSE TO TREATMENT

The overall clinical RR was 79% (106/135; 95% CI, 71–85%), with a clinical complete RR of 21% (29/135), a partial RR of 57% (77/135) and a disease progression rate of 5% (7/135). The clinical RR following the initial docetaxel regimen was 64%. The clinical responses to treatment with docetaxel followed by FEC according to response to initial docetaxel are shown in Table 3. Eight of the 135 patients (6%) progressed during docetaxel administration; 2 of 135 patients (1%) had disease progression during FEC. Of the 30 patients with triple-negative disease, 7 patients were observed to have disease progression following docetaxel treatment. One of the 17 patients with ER-positive, PgR-negative and HER2-negative tumors had disease

Table 2. Treatment-related adverse events

| Adverse event, <i>n</i> (%) | DOC (<i>n</i> = 135) | | FEC (<i>n</i> = 29) | | Overall (<i>n</i> = 35) | |
|-----------------------------------|-----------------------|-----------|----------------------|-----------|--------------------------|-----------|
| | All grades | Grade 3/4 | All grades | Grade 3/4 | All grades | Grade 3/4 |
| Non-hematologic toxicities | | | | | | |
| Infection with neutropenia | 6 (4) | 2 (1) | 3 (2) | 2 (2) | 9 (7) | 4 (3) |
| Fever | 15 (11) | 0 | 13 (10) | 1 (1) | 22 (16) | 1 (1) |
| Infection (other) | 3 (2) | 1 (1) | 2 (2) | 0 | 4 (3) | 1 (1) |
| Fatigue | 82 (61) | 0 | 84 (65) | 2 (2) | 98 (73) | 2 (1) |
| Nausea | 52 (39) | 1 (1) | 102 (79) | 3 (2) | 108 (80) | 4 (3) |
| Vomiting | 19 (14) | 1 (1) | 51 (40) | 3 (2) | 61 (45) | 4 (3) |
| Anorexia | 53 (39) | 1 (1) | 86 (67) | 2 (2) | 91 (67) | 2 (1) |
| Stomatitis | 50 (37) | 1 (1) | 51 (40) | 0 | 68 (50) | 1 (1) |
| Diarrhea | 39 (29) | 1 (1) | 20 (16) | 0 | 46 (34) | 1 (1) |
| Phlebitis | 2 (1) | 1 (1) | 2 (2) | 0 | 4 (3) | 1 (1) |
| Alanine aminotransferase | 36 (27) | 0 | 50 (39) | 2 (2) | 57 (42) | 2 (1) |
| Aspartate aminotransferase | 19 (14) | 0 | 34 (26) | 1 (1) | 40 (30) | 1 (1) |
| Nail changes | 2 (1) | 0 | 33 (26) | 1 (1) | 33 (24) | 1 (1) |
| Weight loss | 5 (4) | 0 | 6 (5) | 1 (1) | 8 (6) | 1 (1) |
| Creatinine | 4 (3) | 1 (1) | 6 (5) | 0 | 7 (5) | 1 (1) |
| Edema | 44 (33) | 0 | 37 (29) | 0 | 55 (41) | 0 |
| Hematologic toxicities | | | | | | |
| Neutropenia | 60 (44) | 59 (44) | 91 (71) | 77 (60) | 100 (74) | 91 (67) |
| Leukopenia | 69 (51) | 50 (37) | 101 (78) | 66 (51) | 108 (80) | 76 (56) |
| Thrombocytopenia | 13 (10) | 0 | 28 (22) | 2 (2) | 31 (23) | 1 (1) |
| Anemia | 66 (49) | 0 | 99 (77) | 1 (1) | 106 (79) | 1 (1) |
| Febrile neutropenia | 9 (7) | 9 (7) | 15 (12) | 15 (12) | 20 (15) | 20 (15) |

DOC, docetaxel; FEC, 5-fluorouracil, epirubicin and cyclophosphamide.

Table 3. Clinical response to DOC followed by FEC according to response to initial DOC treatment (*n* = 135)

| Clinical response, ^a <i>n</i> (%) | Total ^b | Responder | Non-responder |
|--|--------------------|-----------|---------------|
| Response to DOC | | | |
| Responder | 87 (64) | 79 (58) | 8 (6) |
| Non-responder | 48 (36) | 27 (20) | 21 (16) |

^aOverall response was confirmed after completion of chemotherapy in comparison with before docetaxel treatment.

^bPercent value of each column was calculated by dividing by the total number of the evaluable patients (*n* = 135).

progression; while of the 53 patients with ER-positive, PgR-positive, and HER2-negative tumors and of the 9 patients with ER-positive, PgR-positive, and HER2-positive tumors, no patient had disease progression during docetaxel treatment. Among those with triple-negative disease, the majority of patients with disease progression after initial

docetaxel were premenopausal [6/7 patients (86%)] and had solid-tubular carcinoma which characterized by solid cluster of cancer cells with expansive growth forming sharp borders [4/7 patients (57%)], as assessed using the Japanese Breast Cancer Society histological classification of breast tumors (21) (Table 4). Excluding the differences outlined above, there were no differences between patient and tumor characteristics for those with progressive disease versus non-progressive disease.

Twenty-seven of 48 non-responders to docetaxel (56%) had a response to FEC treatment; however, 8 of 87 responders to docetaxel (9%) showed no improvement in response with FEC treatment. Following chemotherapy, BCS was performed for 105 of 133 assessable patients (79%).

PATHOLOGICAL RESPONSE AND PREDICTIVE FACTORS TO TREATMENT

The primary endpoint—pCR rate—was 23% (95% CI, 16–31%). A near pCR rate of 6% (95% CI, 3–12%) resulted

Table 4. Clinical and pathologic characteristics of triple-negative breast cancer^a for patients with progressive disease versus patients without progressive disease, following initial docetaxel therapy

| Characteristic | Without PD | PD |
|--------------------------------|------------|---------|
| No. of evaluable patients | 23 | 7 |
| Age, years | | |
| Median | 43 | 46 |
| Range | (30–62) | (29–53) |
| Menopausal status, n (%) | | |
| Premenopausal | 15 (65) | 6 (86) |
| Postmenopausal | 8 (35) | 1 (14) |
| Tumor stage | | |
| T1 | 2 (9) | 0 |
| T2 | 14 (61) | 5 (71) |
| T3 | 7 (30) | 2 (29) |
| Nodal stage, n (%) | | |
| N0 | 13 (57) | 3 (43) |
| N1 | 10 (43) | 4 (57) |
| Tumor type, n (%) | | |
| Solid-tubular carcinoma | 6 (26) | 4 (57) |
| Papillotubular carcinoma | 5 (22) | 3 (43) |
| Scirrhous carcinoma | 3 (13) | 0 |
| Unspecified invasive carcinoma | 9 (39) | 0 |

PD, progressive disease.
^aTriple-negative tumors were defined as ER-negative, PgR-negative and HER2-negative primary breast cancer.

in a QpCR rate of 29% (95% CI, 21–37%) when combined with the pCR. Pathological response of each subset population according to their hormone receptor and HER2 status is summarized in Fig. 1A and B. Patients with triple-negative disease had the highest pCR rate of 48% (95% CI, 29–68%). Near pCR was not observed in triple-negative disease. Patients with HER2-positive, ER-negative and PgR-negative tumors had a pCR rate of 29% (95% CI, 8–58%) and a QpCR rate of 36% (95% CI, 13–65%); patients with HER2-positive and ER-positive and/or PgR-positive tumors had a pCR rate of 19% (95% CI, 4–46%) and a QpCR rate of 38% (95% CI, 15–65%). Patients with HER2-negative and ER-positive and/or PgR-positive tumors had the lowest pCR and QpCR rates (13%; 95% CI, 6–23% and 19%; 95% CI, 10–30%, respectively). One of the seven patients who experienced clinical disease progression with initial docetaxel treatment had a QpCR following FEC.

The relationship between tumor pathological feature and pCR rate is shown in Table 5. The only variable found to be significantly associated with a pCR after docetaxel treatment was ER status.

Survival outcomes will be reported when the 5-year follow-up has been completed for this study.

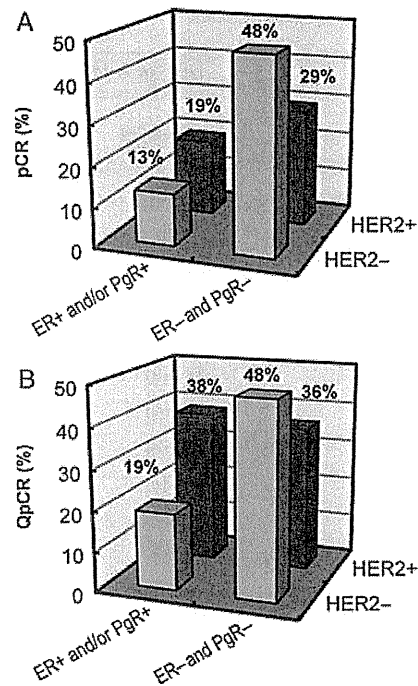


Figure 1. (A) Relationship between pCR versus HER2 and ER/PgR status following DOC and FEC ($n = 129$). (B) Relationship between QpCR versus HER2 and ER/PgR status following DOC and FEC ($n = 129$). Three patients were excluded from evaluable patients for pathologic response ($n = 132$) because of their unknown hormone receptor status. There were no near pCR case observed in triple-negative (ER-, PgR- and HER2-) diseases. DOC, docetaxel; ER, estrogen receptor; FEC, 5-fluorouracil, epirubicin, and cyclophosphamide; HER2, human epidermal growth factor receptor 2; pCR, pathologic complete response; PgR, progesterone receptor; QpCR, quasi-pathologic complete response.

DISCUSSION

This is the first report to evaluate the effectiveness of an initial docetaxel regimen for neoadjuvant therapy of Japanese patients with early-stage breast cancer. An additional component of the study was to analyze the data according to hormone receptor and HER2 status. Recently, Wildiers et al. (22) reviewed four adjuvant trials which had demonstrated the taxane-first regimens were favorable in terms of the relative drug dose intensity achieved. Also they mentioned larger non-randomized adjuvant studies for a series of 284 patients who first received three cycles of FEC followed by three cycles of docetaxel, the mean relative dose intensity was 91% for FEC and 76% for docetaxel, whereas in another series of 378 patients who received three cycles of docetaxel followed by four cycles of EC (epirubicin plus cyclophosphamide), a median docetaxel dose intensity of 100% was achieved. Therefore, they concluded such data suggest that the administration of a taxane first, followed by an anthracycline, may be preferable in line with the Norton–Simon hypothesis (23). In the JBCRG 01 study, the largest study to date to evaluate neoadjuvant chemotherapy in this patient population, the clinical and pathological responses

Table 5. Predictive variables for pCR before and following chemotherapy

| Variables | Before treatment | | | After DOC | | | After FEC following DOC | | |
|--|------------------|-----------|----------------|-----------|-----------|----------------|-------------------------|-----------|----------------|
| | OR | 95% CI | <i>P</i> value | OR | 95% CI | <i>P</i> value | OR | 95% CI | <i>P</i> value |
| Menopausal status: pre (versus post) | 1.5 | 0.94–2.40 | 0.0923 | 1.52 | 0.94–2.47 | 0.0867 | 1.42 | 0.87–2.31 | 0.1575 |
| Tumor size: ≥3 cm (versus <3 cm) | 1.51 | 0.94–2.41 | 0.0881 | 1.45 | 0.90–2.34 | 0.1266 | 1.56 | 0.96–2.52 | 0.0724 |
| ER: negative (versus positive) | 0.58 | 0.32–1.03 | 0.0650 | 0.51 | 0.28–0.95 | 0.0331 | 0.58 | 0.32–1.05 | 0.0709 |
| PgR: negative (versus positive) | 0.66 | 0.34–1.28 | 0.2211 | 0.72 | 0.37–0.95 | 0.3408 | 0.65 | 0.33–1.27 | 0.2083 |
| HER2: 3+ (versus <3+) | 1.32 | 0.76–2.28 | 0.3251 | 1.41 | 0.80–2.47 | 0.2360 | 1.39 | 0.80–2.41 | 0.2445 |
| Clinical response to DOC | | | | | | | | | |
| Response (versus no response) | — | — | — | 0.64 | 0.38–1.07 | 0.0875 | — | — | — |
| Clinical response to FEC following DOC | | | | | | | | | |
| Response (versus no response) | — | — | — | — | — | — | 0.58 | 0.29–1.14 | 0.1160 |

CI, confidence interval; OR, odds ratio; pCR, pathologic complete response.

and safety of FEC followed by docetaxel were investigated (19). The eligibility criteria, treatment dose and distribution of patient characteristics (menopausal status, tumor stage, hormone receptor status and HER2 status) studied in the JBCRG 01 trial were similar to those investigated in the present JBCRG 03 study (19). The incidences of Grade 3/4 neutropenia and febrile neutropenia observed in the current study were similar to those reported in the JBCRG 01 trial (19). However, the rate of Grade 1/2 edema during docetaxel treatment was lower in the present study (33%) than in the JBCRG 01 study (41%), suggesting that docetaxel might be better tolerated when given up front than when administered after completion of prior chemotherapy. Further studies are warranted to assess quality of life and the incidence of edema in order to confirm the effect of administering docetaxel as the initial therapy.

Many different neoadjuvant chemotherapy schedules and dose regimens are used in clinical practice. The NSABP Protocol B-18 trial, which compared AC treatment before and after surgery, reported no difference in DFS between the two approaches (17). However, the rate of BCS was greater with neoadjuvant AC chemotherapy, and the prognosis of patients who obtained a pCR was also better with this treatment regimen (17). Several other regimens have been evaluated in an effort to increase the pCR rate. The addition of a taxane to an anthracycline-containing regimen has been shown to improve the pCR and clinical RRs (5,18). Furthermore, excellent results have been reported by the MD Anderson Cancer Center using a regimen of paclitaxel plus trastuzumab followed by FEC plus trastuzumab in patients with operable breast cancer and HER2 overexpression (24). However, few studies have evaluated initial taxane therapy followed by an anthracycline-containing regimen in this indication (24). Thus, it was decided to evaluate such a reverse regimen and to analyze the findings according to molecular subtypes. Importantly, the primary endpoint—pCR rate—

achieved in the present study was 23% (95% CI, 16–31%), far exceeding our estimate of 12% (19). Even though the pCR rate here cannot be directly compared with the results from the JBCRG 01 trial (pCR rate: 12%, QpCR rate: 25%), the pCR rate from this study is a favorable result considering the similar patient characteristics in both trials (19).

The overall clinical RR of 79% was similar to that reported in the JBCRG 01 trial (74%) (19). Furthermore, the clinical RR following the initial docetaxel regimen was 64%, similar to the clinical response following the initial FEC regimen in the JBCRG 01 trial (61%) (19). The clinical RR following the initial docetaxel regimen, however, is lower in this study than those reported in other studies (71.7–85%) (25,26). It could be hypothesized that the clinical response might be influenced by the lower dose of docetaxel used in this study (75 mg/m²) compared with the 100 mg/m² dose used in previous studies (25,26).

The rate of BCS observed in our study (79%) was similar to that reported in the JBCRG 01 trial (85%) (19). Unfortunately, the overall disease progression rate (5%) was not lowered by the use of docetaxel followed by FEC in this study, and was similar to that seen in the JBCRG 01 trial (6%) (19).

Although 7 of the 29 patients with triple-negative disease had disease progression during the initial docetaxel regimen, 14 of the 22 patients without disease progression (64%) achieved a QpCR. This QpCR rate is markedly higher compared with previous findings (27).

Our results indicate that if patients with triple-negative disease who experienced disease progression following initial docetaxel therapy were excluded, the pCR rate for this group of patients would have been higher. We thus compared the clinical and pathological characteristics between patients with triple-negative disease who experienced disease progression following the initial docetaxel regimen with those who did not have disease progression. However, no

significant differences in patient or tumor characteristics were seen between these patient groups. It was noted, however, that six of seven premenopausal patients (86%) and four of seven patients (57%) with solid-tubular carcinoma had disease progression following docetaxel therapy. Given the high incidence of disease progression among patients with triple-negative disease who had solid-tubular subtype tumors, this phenotype could be used in future studies to predict which patients are more likely to experience progressive disease following docetaxel therapy. Accordingly, the identification of patients with hormone receptor-positive and HER2-negative disease would also enable the selection of patients who are more likely to benefit from neoadjuvant chemotherapy. Thus, studying patients' molecular subtypes, and selecting appropriate chemotherapy regimens accordingly, has the potential to provide superior results to those of the JBCRG 03 trial.

Recently, it has been shown that basal-like breast cancer defined by five biomarkers [epidermal growth factor receptor (EGFR), cytokeratin 5/6 (CK5/6), ER, PgR and HER2 status] provides a more specific definition of basal-like breast cancer that predicts survival better than the triple-negative phenotype (27,28). In patients treated with anthracycline-based chemotherapy, tumors found to be positive for the basal markers corresponded to a cohort of patients with a significantly worse outcome (29). Thus in future trials, it may be beneficial to assess EGFR and CK5/6 status in patients with triple-negative disease to help predict patient survival.

Interestingly, the pCR rate (27%) following neoadjuvant chemotherapy in patients with HER2-negative breast cancer was higher in this study than in the JBCRG 01 study (14%), suggesting that this subpopulation may benefit from initial docetaxel treatment. Conversely, a lower QpCR rate was observed in HER2-positive patients (37%) in this study than in the JBCRG 01 trial (52.8%). This suggests that initial anthracyclines may be required for HER2-positive disease. A study by Buzdar et al. (24) reported that a high pCR rate of 60% was observed in patients with HER2-positive disease treated with the combination of paclitaxel plus trastuzumab followed by FEC plus trastuzumab, indicating that the HER2-positive population in the current study may have benefited further from concomitant trastuzumab therapy. These findings demonstrate the benefit of selecting the most effective chemotherapy regimen according to each patient's molecular subtype and initial response to neoadjuvant treatment.

One limitation of the study was that HER2-positive patients were not treated with trastuzumab, which has been shown to improve outcomes in patients with HER2-overexpressing breast cancer (24). Further studies investigating optimal treatment regimens for different molecular subtypes should include concurrent trastuzumab for patients with the HER2-positive phenotype.

In conclusion, docetaxel followed by FEC as neoadjuvant chemotherapy is a tolerable and effective regimen for

patients with early-stage breast cancer. In addition, a high pCR rate made this regimen particularly promising in patients with triple-negative breast cancer. In the future, selection of a neoadjuvant chemotherapy regimen for operable breast cancer may be possible based on molecular subtype.

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Conflict of interest statement

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