

Fig. 1. In patient with positive node (n = 84), the most radioactive positive node was within third rank of radioisotope count.

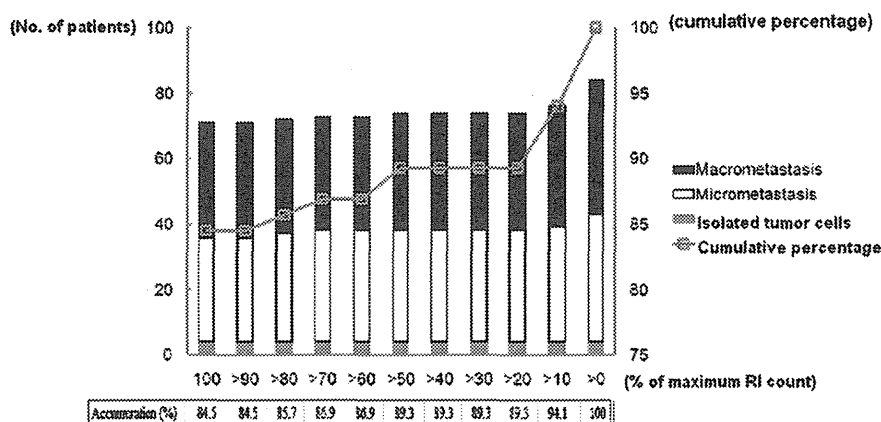


Fig. 2. Histological diagnosis according to percentage of the maximum radioisotope count.

and the TNM classification requires 6 or more lymph nodes to be excised for pN staging, the boundary between SNB and ALND can be defined at the number 6. Another guideline known as the '10% rule' was introduced in 2000 by Martin et al.⁴ According to this rule, any node with 10% or more of the *ex vivo* RI count of the hottest node should be removed as a SLN. When the '10% rule' was used in this study, the false-negative rate was 5.9% and slightly high. The reported false-negative rates were 1.7% in Chung et al.⁶ and 5.8% in Martin et al.⁴ The accepted false-negative rate for SNB was 5%⁷ and a record of less than 5% false-negatives at our institute since the introduction of SNB in 2000.

Although 22.9% of positive nodes were not stained but detected by RI only, all the hottest positive nodes were stained by indigo carmine. According to the current procedure of SNB, all the nodes that were stained and/or react to the gamma probe were excised by surgeon. Therefore the number of excised node by RI was apt to larger. And dye method is inferior to RI or combination method in detective rate, generally.^{10,11} However, Narui et al. reported that a 4 node sampling method using only patent blue was reliable,¹² and in our study 2 nodes were found only through the dye method not due to accumulate RI. Metastatic disease can injure the lymphatic system and technical errors can occur during SNB. Recognition of

the method's limitations and careful intraoperative palpitation are necessary.

In summary, the validity of these stopping rules in SNB under the use of indigo carmine blue dye was analyzed. Though the '10% rule' resulted in slightly high false-negative rate with the RI count method, a positive sentinel lymph node was identified in 100% of cases within the first 3 sentinel nodes when indigo carmine and RI was used. Under the use of indigo carmine, terminating SNB at 3 sampled nodes was the validity procedure to minimize the false-negative and complication rates.

Conflict of interest statement

None declared.

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Tumor-infiltrating lymphocytes are correlated with response to neoadjuvant chemotherapy in triple-negative breast cancer

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Received: 18 January 2011 / Accepted: 25 April 2011 / Published online: 12 May 2011
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Abstract The purpose of the present study was to identify histological surrogate predictive markers of pathological complete response (pCR) to neoadjuvant chemotherapy (NAC) in triple-negative breast cancer (TNBC). Among 474 patients who received NAC and subsequent surgical therapy for stage II–III invasive breast carcinoma between 1999 and 2007, 102 (22%) had TNBC, and 92 core needle biopsy (CNB)

specimens obtained before NAC were available. As controls, CNB specimens from 42 tumors of the hormone receptor-negative and HER2-positive (HR–/HER2+) subtype and 46 tumors of the hormone receptor-positive and HER2-negative (HR+/HER2–) subtype were also included. Histopathological examination including tumor-infiltrating lymphocytes (TIL) and tumor cell apoptosis, and immunohistochemical studies for basal markers were performed, and the correlation of these data with pathological therapeutic effect was analyzed. The rates of pCR at the primary site were higher for TNBC (32%) and the HR–/HER2+ subtype (21%) than for the HR+/HER2– subtype (7%) ($P = 0.006$). Expression of basal markers and p53, histological grade 3, high TIL scores, and apoptosis were more frequent in TNBC and the HR–/HER2+ subtype than in the HR+/HER2– subtype ($P = 0.002$ for TIL and $P < 0.001$ for others). In TNBC, the pCR rates of tumors showing a high TIL score and of those showing a high apoptosis score were 37 and 47%, respectively, and significantly higher or tended to be higher than those of the tumors showing a low TIL score and of the tumors showing a low apoptosis score (16 and 27%, respectively, $P = 0.05$ and 0.10). In a total of 180 breast cancers, the pCR rates of the tumors showing a high TIL score (34%) and of those showing a high apoptosis score (35%) were significantly higher than those of the tumors showing a low TIL score (10%) and those of the tumors showing a low apoptosis score (19%) ($P = 0.0001$ and 0.04 , respectively). Histological grade and basal marker expression were not correlated with pCR. Although the whole analysis was exploratory, the degree of TIL correlated with immune response appear to play a substantial role in the response to NAC in TNBC.

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Keywords Triple-negative breast cancer · Neoadjuvant
chemotherapy · Pathological complete response ·
Tumor-infiltrating lymphocytes · Tumor cell apoptosis

Introduction

The heterogeneous nature of breast cancer has been demonstrated by gene expression profiling using the DNA microarray technique [1–3]. Genetically, invasive breast cancers have been classified into distinct intrinsic subtypes comprising luminal A, luminal B, ERBB2 (HER2), basal-like, and normal breast subtypes [1–3], which demonstrate characteristic immunohistochemical features and clinical behavior [4–8]. Both basal-like and normal breast subtypes are immunohistochemically characterized by lack of expression of the estrogen receptor (ER), progesterone receptor (PgR), and HER2, and thus are also categorized as triple-negative breast cancer (TNBC). TNBC, which accounts for 10–15% of all breast cancers, tends to show visceral metastasis and aggressive clinical behavior [9].

TNBC is unresponsive to specific targeted therapies such as trastuzumab for HER2-positive breast cancer, or hormonal therapy for hormone-receptor-positive breast cancer. In cases of operable TNBC, only systemic chemotherapy has been shown to be effective in an adjuvant or neoadjuvant setting. Although patients with TNBC are more likely to achieve a pathological complete response (pCR) after neoadjuvant chemotherapy (NAC) than patients with the luminal subtypes, and pCR is correlated with an excellent clinical outcome, TNBC patients with residual disease after NAC have a poor prognosis [10, 11]. However, the factor that determines sensitivity to chemotherapy in patients with TNBC is uncertain.

TNBC itself may show heterogeneous characteristics including basal-like and normal breast subtypes, as judged from gene expression profiles [1–3]. Accordingly, it is important to investigate the pathological factors associated with response to chemotherapy in patients with TNBC.

The aim of the present study was to identify the factors that predict pCR after NAC in patients with TNBC by examination of histological parameters including histological grade and type, the presence of tumor-infiltrating lymphocytes (TIL), and tumor cell apoptosis, as well as immunohistochemical parameters including basal-like markers and p53.

Materials and methods

Patients and tissue samples

Among 474 patients who received NAC and subsequent surgical therapy for stage II–III invasive breast carcinoma between 1999 and 2007, 102 (22%) had TNBC. Originally, we planned to compare 100 TNBCs with 100 non-TNBCs as controls on the basis of matching for age (± 5 years) and clinical stage (II and III). In the 100 control cases, we planned to include 50 cases of the HR–/HER2+ subtype

(HER2 positive and ER/PgR negative in routine immunohistochemistry) and 50 cases of the HR+/HER2– subtype (ER and/or PgR positive but HER2 negative in routine immunohistochemistry). From these patients, sufficient CNB specimens before NAC were available from 92 tumors of TNBC, 42 tumors of the HR–/HER2+ subtype, and 46 tumors of the HR+/HER2– subtype. Clinical characteristics of all patients were obtained from the medical records. All patients received neoadjuvant anthracycline-based regimens (adriamycin 60 mg/m² plus cyclophosphamide 600 mg/m² (AC) or cyclophosphamide 600 mg/m² plus epirubicin 100 mg/m²/5-fluorouracil 600 mg/m² (CEF)) alone, taxane-based regimens (weekly paclitaxel 80 mg/m², or triweekly docetaxel 75 mg/m²) alone, or anthracycline and taxane sequentially or concurrently (adriamycin 50 mg/m² plus docetaxel 60 mg/m² (AT), AC or CEF followed by weekly paclitaxel or triweekly docetaxel). Trastuzumab was not used for the 42 patients with tumors of HR–/HER2+ subtype, because the use of trastuzumab for neoadjuvant therapy of primary breast cancer was not approved in Japan. The patients have been followed up for 64.8 months on an average (7.2–138.2 months). All specimens were formalin-fixed and paraffin-embedded, and 4- μ m-thick sections were prepared for hematoxylin and eosin staining and immunohistochemistry (IHC) and were reviewed by two observers including an experienced pathologist (T.H.). The present study was approved by the Institutional Review Board of the National Cancer Center.

Histopathological evaluation

Pathological therapeutic effect was assessed for resected primary tumors after NAC. Pathological complete response (pCR) was defined as the absence of all invasive disease in the breast tumor according to the National Surgical Adjuvant Breast and Bowel Project (NSABP) B-18 protocol [12]. In addition, we defined quasi-pCR (QpCR) as the absence of invasive tumor or only focal residual invasive carcinoma cells in the primary site [13]. In Japan, Breast Cancer Research Group (JBCRG) 01 study, QpCR after NAC was shown to be correlated with better patient prognosis in comparison with non-QpCR [13]. Furthermore, we took into consideration both the pCR in the primary tumor and no residual tumor in axillary lymph nodes as another classification for histopathological therapeutic effect [14, 15].

Histopathological assessment of predictive factors was made for CNB specimens. Histopathological parameters examined included histological grade [16], histological type [17], presence of tumor-infiltrating lymphocytes (TIL), apoptosis, and correlation of these parameters with intrinsic subtypes and pCR. Histological grade was assigned on the basis of the criteria of Elston and Ellis.

For the evaluation of TIL, both areas of stroma infiltrated by lymphocytes (proportional score) and intensity of lymphatic infiltration (intensity score) were taken into consideration. Proportional scores were defined as 3, 2, 1, and 0 if the area of stroma with lymphoplasmacytic infiltration around invasive tumor cell nests were >50 , >10 – 50 , $\leq 10\%$, and absent, respectively. Intensity scores were defined as 2, 1, and 0, if the intensity of lymphatic infiltration was marked, mild, and absent, respectively (Fig. 1). Lymphocyte infiltration surrounding non-invasive tumor cells was not taken into account. The proportional and intensity scores were summed for each tumor, and the TIL score was classified as high if the sum was 3–5, whereas the TIL score was classified as low if the sum was 0–2. As criteria for apoptosis, scores were defined as 2, 1, and 0 if apoptotic cells (arrows in Fig. 2) were >10 per 10 high-power fields (HPFs) using $40\times$ objective lens, 5–9 per 10 HPFs, and less than 5 per 10 HPFs, respectively.

Immunohistochemistry (IHC)

IHC was performed for CNB specimens using the following primary antibodies: anti-ER (clone 1D5; Dako), anti-PgR (clone PgR636; Dako), anti-HER2 (polyclonal, HercepTest II, Dako), anti-p53 (clone DO-7; Dako), anti-cytokeratin (CK) 5/6 (clone D5/16 B4; Dako), anti-CK14 (NCL-LL002, Novocastra), and anti-EGFR (pharmDX, clone 2-18C9, Dako).

Because ER, PgR, and HER2 tests had been performed by various antibodies and methods, these tests were re-tested again according to standardized antibodies and

methods in the present study. The sections were deparaffinized, subjected to antigen retrieval by incubating in target retrieval solution, high pH (Dako) for 40 min at 95°C for ER and PgR, in sodium citrate buffer (pH 6.0) with a microwave oven for 15 min at 97°C for CK14, in sodium citrate buffer (pH 6.0) with a water bath for 15 min at 98°C for CK5/6, or by autoclaving in sodium citrate buffer (pH 6.0) for 20 min at 121°C for p53, then allowed to cool at room temperature. Endogenous peroxidase and non-specific staining were blocked in 2% normal swine serum (Dako). The slides were incubated with primary antibodies at 4°C overnight and then reacted with a dextran polymer reagent combined with secondary antibodies and peroxidase (Envision Plus, Dako) for 2 h at room temperature. Specific antigen–antibody reactions were visualized using 0.2% diaminobenzidine tetrahydrochloride and hydrogen peroxide. Counterstaining was performed using Mayer's hematoxylin. For the HER2 and EGFR kits, immunohistochemistry was performed in accordance with the protocol recommended by the manufacturer.

ER and PgR were judged as positive if the Allred score was ≥ 3 and as negative if the Allred score was ≤ 2 [18]. HER2 protein overexpression was judged as positive when the score was 3+, equivocal when the score was 2+, and negative when the score was 0 or 1+ in accordance with the ASCO/CAP recommendation [19]. TNBC was defined as negative for ER, PgR, and HER2, while the HR+/HER2– subtype was defined as positive for ER or PgR and negative for HER2, and the HR–/HER2+ subtype was defined as negative for ER and PgR, and positive for HER2. The basal-like subtype was defined as CK5/

Fig. 1 Histopathological features of tumor-infiltrating lymphocytes (TILs). **a** High TIL score (proportional score 3+ intensity score 2); **b** High TIL score (proportional score 2+ intensity score 2); **c** Low TIL score (proportional score 1+ intensity score 2); **d** Low TIL score (proportional score 0, intensity score 0). Original magnification: $400\times$

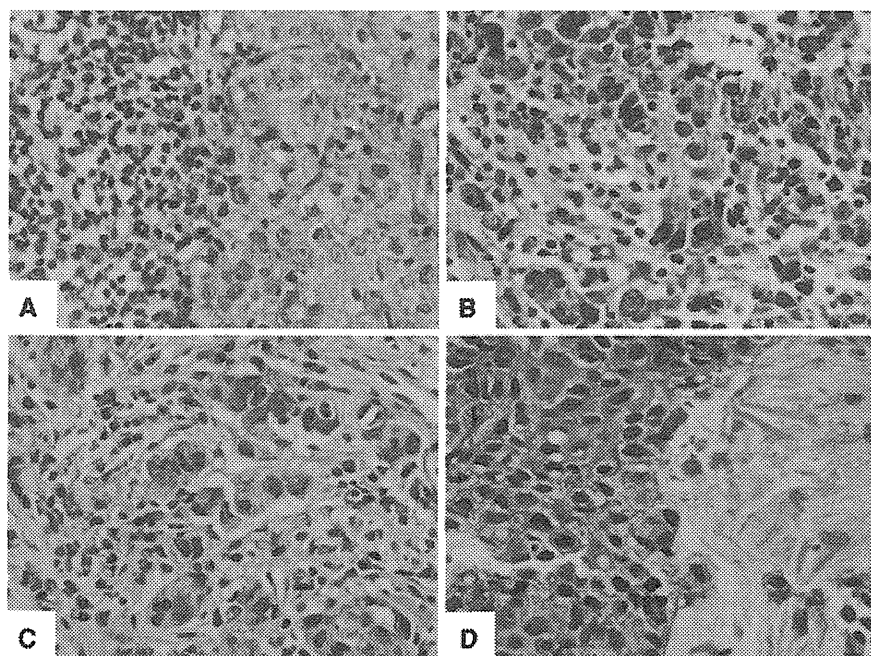
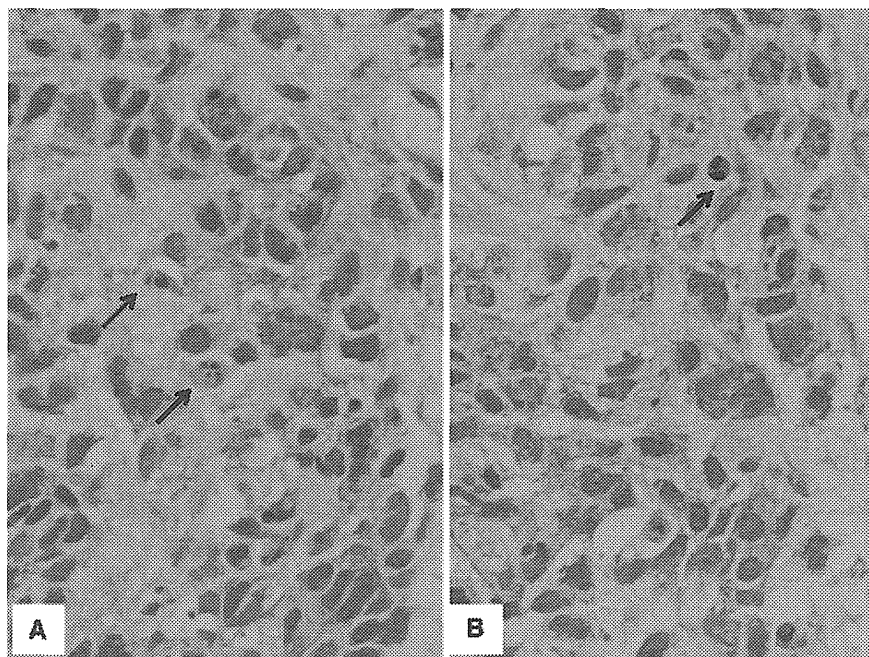


Fig. 2 Histopathological features of breast carcinoma with apoptosis (a, b) (arrows: apoptosis) Original magnification: 400 \times



6 > 1%, CK14 > 1%, or EGFR > 1%. For reference, data based on the criteria CK5/6 > 10%, CK14 > 10%, or EGFR > 10% were also acquired. p53 was scored using the Allred score and was regarded as positive when ≥ 5 .

Statistical analyses

Statistical analyses were performed using SPSS software. Patients' characteristics were compared between subgroups using the chi-squared test or Fisher's exact test for categorical variables, and Kruskal–Wallis test for continuous variables. Association of pathological parameters, including a basal-like subtype, with pCR, QpCR, or pCR and no residual axillary tumor were evaluated using the chi-squared test or Fisher's exact test. Predictive ratio of pCR, QpCR, or pCR plus residual axillary metastasis by clinicopathological parameters were analyzed using the univariate and multivariate logistic regression models. Survival curves of patients were drawn using Kaplan–Meier method, and statistical difference between survival curves were calculated by using the log-rank test. In all analyses, differences were considered significant at $P < 0.05$.

Results

We confirmed immunohistochemically that all 92 tumors were TNBC, 42 of 50 were of the HR–/HER2+ subtype, and 46 of 50 were of the HR+/HER2– subtype. A total of

180 specimens were investigated in this study. The characteristics of the patients are presented in Tables 1 and 2.

Clinicopathological characteristics and subtypes

In tumors with the TNBC and HR–/HER2+ subtype, the frequencies of the basal-like subtype were 59% (54 of 92) and 43% (18 of 42), respectively, compared with only 7% (3 of 46) in the HR+/HER2– subtype. Therefore, the incidence of the basal-like subtype was significantly higher in TNBC or in the HR–/HER2+ subtype than in the HR+/HER2– subtype ($P < 0.001$). Similarly, the frequency of p53 expression was significantly higher in TNBC (63%, 58 of 92) and the HR–/HER2+ subtype (62%, 26 of 42) than in the HR+/HER2– subtype (26%, 12 of 46) ($P < 0.001$). Tumors of histological grade 3 were more frequent in TNBC (89%, 82 of 92) and the HR–/HER2+ subtype (81%, 34 of 42) than in the HR+/HER2– subtype (13%, 6 of 46) ($P < 0.001$).

The incidence of high TIL score (score 3–5) was also higher in TNBC (73%, 67 of 92) and the HR–/HER2+ subtype (55%, 23 of 42) than in the HR+/HER2– subtype (17%, 8 of 46) ($P = 0.002$). An apoptosis score of 2 was also more frequent in TNBC (21%, 19 of 92) and the HR–/HER2+ subtype (48%, 20 of 42) than in the HR+/HER2– subtype (2%, 1 of 46) ($P < 0.001$). The incidences of a basal-like subtype, p53 expression, a high TIL score, and an apoptosis score of 2 did not differ between TNBC and the HR–/HER2+ subtype.

All six metaplastic carcinomas were TNBC [17].

Table 1 Evaluation of clinicopathological parameters in three subtypes of primary breast cancer

	TNBC (<i>n</i> = 92) No. of patients (%)	HR-/HER2+ (<i>n</i> = 42) No. of patients (%)	HR+/HER2- (<i>n</i> = 46) No. of patients (%)	<i>P</i> value
Age				
Median (range)	52 (23-76)	55 (31-71)	55 (31-71)	0.36
<i>T</i>				
1	2 (2)	0 (0)	0 (0)	0.37
2	48 (53)	17 (41)	26 (56)	
3	27 (29)	16 (38)	11 (24)	
4	15 (16)	9 (21)	9 (20)	
<i>N</i>				
0	45 (49)	24 (57)	24 (52)	0.96
1	35 (38)	14 (33)	18 (39)	
2	10 (11)	3 (7)	3 (7)	
3	2 (2)	1 (3)	1 (2)	
Stage				
II	56 (61)	25 (60)	28 (61)	0.99
III	36 (39)	17 (40)	18 (39)	
ER				
Positive	0 (0)	0 (0)	46 (100)	
Negative	92 (100)	42 (100)	0 (0)	
PgR				
Positive	0 (0)	0 (0)	32 (70)	
Negative	92 (100)	42 (100)	14 (30)	
HER2				
Positive	0 (0)	42 (100)	46 (0)	
Negative	92 (100)	0 (0)	0 (100)	
Basal marker				
Positive	54 (59)	18 (43)	3 (7)	<0.001
Negative	38 (41)	24 (57)	43 (93)	
p53				
Positive	58 (63)	26 (62)	12 (26)	<0.001
Negative	34 (37)	16 (38)	34 (74)	
Grade				
1	1 (1)	0 (0)	4 (9)	<0.001
2	9 (10)	8 (19)	36 (78)	
3	82 (89)	34 (81)	6 (13)	
TIL				
Low (0/1/2)	25 (4/8/13) (27)	19 (7/6/6) (45)	38 (25/8/5) (83)	0.002
High (3/4/5)	67 (22/24/21) (73)	23 (8/11/4) (55)	8 (6/2/0) (17)	
Apoptosis				
0	22 (24)	8 (19)	29 (63)	<0.001
1	51 (55)	14 (33)	16 (35)	
2	19 (21)	20 (48)	1 (2)	
pCR (NSABP B-18)				
Yes	29 (32)	9 (21)	3 (7)	0.004
No	63 (68)	33 (79)	43 (93)	
QpCR (JBCRG 01)				
Yes	35 (38)	17 (40)	3 (7)	<0.001
No	57 (62)	25 (60)	43 (93)	
pCR (primary and lymph nodes)				
Yes	26 (28)	6 (14)	3 (7)	0.006
No	66 (72)	36 (86)	43 (93)	

ER estrogen receptor, *HR* hormone receptors, *pCR* pathological complete response, *PgR* progesterone receptor, *TIL* tumor infiltrating lymphocytes, *TNBC* triple negative breast cancer

Table 2 Correlation between therapeutic effect of primary breast cancer to neoadjuvant chemotherapy (NAC) and infiltrating lymphocytes (TIL)

Subtype of breast cancer and response to NAC	No. of patients (%)			P
	Total	TIL score		
		0–2	3–5	
A. TNBC				
pCR (NSABP B-18)				
Yes	29 (32)	4 (16)	25 (37)	0.05
No	63 (68)	21 (84)	42 (63)	
QpCR (JBCRG)				
Yes	35 (38)	4 (16)	31 (46)	0.008
No	57 (62)	21 (84)	36 (54)	
pCR (primary + lymph nodes)				
Yes	26 (28)	4 (16)	22 (33)	0.11
No	66 (72)	21 (84)	45 (67)	
B. HR–/HER2+ subtype				
pCR (NSABP B-18)				
Yes	9 (21)	2 (11)	7 (30)	0.12
No	33 (79)	17 (89)	16 (70)	
QpCR (JBCRG)				
Yes	17 (40)	5 (26)	12 (52)	0.09
No	25 (60)	14 (74)	11 (48)	
pCR (primary + lymph nodes)				
Yes	6 (14)	1 (5)	5 (22)	0.13
No	36 (86)	18 (95)	18 (78)	
C. HR+/HER2– subtype				
pCR (NSABP B-18)				
Yes	3 (7)	2 (5)	1 (13)	0.44
No	43 (93)	36 (95)	7 (87)	
QpCR (JBCRG)				
Yes	3 (7)	2 (5)	1 (13)	0.44
No	43 (93)	36 (95)	7 (87)	
pCR (primary + lymph nodes)				
Yes	3 (7)	2 (5)	1 (13)	0.44
No	43 (93)	36 (95)	7 (87)	
D. Total (TNBC+ HR–/HER2+ HR+/HER2–)				
pCR (NSABP B-18)				
Yes	41 (23)	8 (10)	33 (34)	0.0001
No	139 (77)	74 (90)	65 (66)	
QpCR (JBCRG)				
Yes	55 (31)	11 (13)	44 (45)	< 0.0001
No	125 (69)	71 (87)	54 (55)	
pCR (primary + lymph nodes)				
Yes	35 (19)	7 (9)	28 (29)	0.0007
No	145 (81)	75 (91)	70 (71)	

HR hormone receptors, TNBC triple-negative breast cancer, TIL tumor-infiltrating lymphocyte, pCR pathologically complete response, QpCR quasi-pCR, NAC neoadjuvant chemotherapy

Clinicopathological characteristics and pCR

The pCR rate according to NSABP B-18 classification was significantly higher in TNBC (32%) and HR–/HER2+ subtype (21%) than in HR+/HER2– subtype (7%) ($P = 0.004$). Likewise, the QpCR rate according to

JBCRG 01 classification was significantly higher in TNBC (38%) and HR–/HER2+ subtype (40%) than in HR+/HER2– subtype (7%) ($P < 0.001$). Furthermore, the rate of pCR in both primary site and lymph nodes was significantly higher in TNBC (28%) than in HR–/HER2+ (14%) and HR+/HER2– (7%) subtypes ($P = 0.006$) (Table 1).

Table 3 Correlation between apoptosis of tumor cells and therapeutic effect of primary breast cancer to neoadjuvant chemotherapy (NAC)

Subtype of breast cancer and response to NAC	No. of patients (%)			<i>P</i>
	Total	Apoptosis		
		Score 0, 1	Score 2	
A. TNBC				
pCR (NSABP B-18)				
Yes	29 (32)	20 (27)	9 (47)	0.10
No	63 (68)	53 (73)	10 (53)	
QpCR (JBCRG)				
Yes	35 (38)	26 (36)	9 (47)	0.35
No	57 (62)	47 (64)	10 (53)	
pCR (primary + lymph nodes)				
Yes	26 (28)	17 (23)	9 (47)	0.04
No	66 (72)	56 (77)	10 (53)	
B. HR–/HER2+ subtype				
pCR (NSABP B-18)				
Yes	9 (21)	4 (18)	5 (25)	0.71
No	33 (79)	18 (82)	15 (75)	
QpCR (JBCRG)				
Yes	17 (40)	7 (32)	10 (50)	0.23
No	25 (60)	15 (68)	10 (50)	
pCR (primary + lymph nodes)				
Yes	6 (14)	2 (9)	4 (20)	0.40
No	36 (86)	20 (91)	16 (80)	
C. HR+/HER2– subtype				
pCR (NSABP B-18)				
Yes	3 (7)	3 (7)	0 (0)	1.00
No	43 (93)	42 (93)	1 (100)	
QpCR (JBCRG)				
Yes	3 (7)	3 (7)	0 (0)	1.00
No	43 (93)	42 (93)	1 (100)	
pCR (primary + lymph nodes)				
Yes	3 (7)	3 (7)	0 (0)	1.00
No	43 (93)	42 (93)	1 (100)	
D. Total (TNBC+ HR–/HER2+ HR+/HER2–)				
pCR (NSABP B-18)				
Yes	41 (23)	27 (19)	14 (35)	0.04
No	139 (77)	113 (81)	26 (65)	
QpCR (JBCRG)				
Yes	55 (31)	36 (26)	19 (47)	0.008
No	125 (69)	104 (74)	21 (53)	
pCR (primary + lymph nodes)				
Yes	35 (19)	22 (16)	13 (32)	0.02
No	145 (81)	118 (84)	27 (68)	

HR hormone receptors, *TNBC* triple-negative breast cancer, *pCR* pathologically complete response, *QpCR* quasi-pCR, *NAC* neoadjuvant chemotherapy

The association between pCR and TIL scores stratified by tumor subtype is shown in Table 2. In patients with TNBC, the pCR rate was significantly higher in those with tumors showing high TIL scores (3–5) (37%, 25 of 67) than in those with tumor showing low TIL scores (0–2) (16%, 4 of 25) ($P = 0.05$). Likewise, the QpCR rate was

significantly higher in those with tumors showing the high TIL scores (46%, 31 of 67) than in those with the low TIL scores (16%, 4 of 25, $P = 0.008$). Furthermore, the rate of pCR in both primary tumor and axillary lymph nodes tended to be higher in the patients with tumors showing the high TIL scores (35%, 22 of 67) than in those with tumors

showing the low TIL scores (16%, 4 of 25). A similar tendency of correlation was seen for tumors of HR-/HER2+ subtype (Table 2), although there was no statistic significance. There was no correlation between TIL and therapeutic effect in HR+/HER2- subtype tumors. In a total of 180 cases including all TNBC, HR-/HER2+, and HR+/HER2- subtypes studied, TIL was significantly correlated with pCR, QpCR, and the pCR in both the primary site and lymph nodes ($P = 0.0001$, $P < 0.0001$, and $P = 0.0007$, respectively, Table 2).

In the patients with TNBC, the pCR rate tended to be higher in those with tumors showing an apoptosis score of 2 (47%, 9 of 19) than in those with an apoptosis score 0 or 1 (27%, 20 of 73, $P = 0.10$) (Table 3). Furthermore, the rate of pCR in both primary tumor and axillary nodes was significantly higher in the tumors showing an apoptosis score 2 (47%, 9 of 19) than in those with an apoptosis score 0 or 1 (23%, 17 of 73, $P = 0.04$). A similar tendency of correlation was seen for tumors of HR-/HER2+ subtype (Table 3), although there was no statistic significance between an apoptosis score and these pCRs (Table 3). There was no statistically significant correlation between apoptosis score and therapeutic effect in HR+/HER2- subtype tumors. In a total of 180 cases including these three subtypes, apoptosis

was significantly correlated with pCR, QpCR, and the pCR in both the primary site and axillary lymph nodes ($P = 0.04$, 0.008, and 0.02, respectively) (Table 3).

The pCR rate did not differ significantly between p53-negative tumors (13 of 34, 38%) and p53-positive tumors (15 of 57, 26%) in patients with TNBC. In the HR-/HER2+ subtype, however, seven of nine patients who achieved pCR had p53-positive tumors. There was no correlation between pCR and p53 in the HR+/HER2- subtype.

The pCR rate did not differ between patients with tumors of the basal-like subtype and those with tumors of the non-basal-like subtype (Table 4). Same tendencies of relationship with p53 status or with basal-like subtype were seen for the classification of QpCR and for the pCR of both the primary site and axillary lymph nodes (data not shown).

When all 180 cases were combined, T, N, and grade were correlated or tended to be correlated with pCR (Table 4). QpCR, and the pCR of both primary site and axillary lymph nodes also showed similar tendency (data not shown). Age was not correlated with therapeutic effect.

A univariate regression model analysis showed that the high TIL score was significantly correlated with QpCR (relative ratio (RR) 4.52, 95% reliable range (95%RR) 1.40–14.59) and nearly significantly correlated with pCR in

Table 4 Correlation of clinicopathological parameters with pathological complete response (pCR) of primary breast cancer to neoadjuvant chemotherapy

	All	No. of pCR/No. of patients (%)						
		<i>P</i> value	TNBC	<i>P</i> value	HR-/HER2+	<i>P</i> value	HR+/HER2-	<i>P</i> value
Age								
≤50	14/64 (22)	0.80	11/40 (28)	0.46	3/12 (25)	0.72	0/12 (0)	0.39
>50	27/116 (23)		18/52 (35)		6/30 (20)		3/34 (9)	
T								
1, 2	26/93 (28)	0.09	18/50 (36)	0.31	6/17 (35)	0.07	2/26 (8)	0.60
3, 4	15/87 (17)		11/42 (26)		3/25 (12)		1/20 (5)	
N								
Positive	14/87 (16)	0.03	11/47 (23)	0.09	2/18 (11)	0.15	1/22 (5)	0.53
Negative	27/93 (29)		18/45 (40)		7/24 (29)		2/24 (8)	
Stage								
II	31/109 (28)	0.03	21/56 (38)	0.12	8/25 (32)	0.05	2/28 (7)	0.66
III	10/71 (14)		8/36 (22)		1/17 (6)		1/18 (6)	
Grade								
1, 2	7/58 (12)	0.02	3/10 (30)	0.91	1/8 (13)	0.44	3/40 (8)	0.65
3	34/122 (29)		26/82 (32)		8/34 (24)		0/6 (0)	
Basal-like								
Positive	23/75 (31)	0.03	19/54 (35)	0.36	4/18 (22)	0.60	0/3 (0)	0.81
Negative	18/105 (17)		10/38 (26)		5/24 (21)		3/43 (7)	
p53								
Positive	23/95 (24)	0.52	15/57 (26)	0.23	7/26 (27)	0.24	1/12 (8)	0.61
Negative	17/84 (20)		13/34 (38)		2/16 (13)		2/34 (6)	

HR hormone receptors, pCR pathological complete response

Table 5 Logistic analysis for prediction of pathological therapeutic effect to neoadjuvant chemotherapy to TNBC

	Relative ratio (95% reliable range)	P value
A. Univariate		
1. pCR (NSABP B-18)		
TIL (score 3-5 vs. 0-2)	3.12 (0.96–10.15)	0.058
Apoptosis (2 vs. 0, 1)	2.38 (0.85–6.73)	0.10
2. QpCR (JBCRG)		
TIL (score 3-5 vs. 0-2)	4.52 (1.40–14.59)	0.012
Apoptosis (2 vs. 0, 1)	1.63 (0.59–4.51)	0.35
3. pCR (primary + lymph node)		
TIL (score 3-5 vs. 0-2)	2.57 (0.79–8.39)	0.12
Apoptosis (2 vs. 0, 1)	2.97 (1.04–8.49)	0.043
B. Multivariate		
1. pCR (NSABP B-18)		
TIL (score 3-5 vs. 0-2)	2.78 (0.84–9.18)	0.09
Apoptosis (2 vs. 0, 1)	2.01 (0.70–5.81)	0.20
2. QpCR (JBCRG)		
TIL (score 3-5 vs. 0-2)	4.34 (1.33–14.21)	0.015
Apoptosis (2 vs. 0, 1)	1.27 (0.44–3.65)	0.66
3. pCR (primary + lymph node)		
TIL (score 3-5 vs. 0-2)	2.17 (0.65–7.28)	0.21
Apoptosis (2 vs. 0, 1)	2.60 (0.89–7.58)	0.08

pCR pathological complete response, TIL tumor-infiltrating lymphocyte, TNBC triple-negative breast cancer

N, T, grade, basal-like, p53, and histological type were not significant as predictor of pCR

92 TNBCs (relative ratio 3.12, 95%RR 0.96–10.15) ($P = 0.012$ and 0.058 , respectively) (Table 5). Apoptosis was significantly correlated with pCR (primary + lymph node) in 92 TNBCs (RR 2.97, 95%RR 1.04–8.49) ($P = 0.043$). Other parameters, including T, N, grade, basal-like subtype, p53 and histological type, were not significant predictors of pCR. TIL and apoptosis showed no mutual correlation. When these two parameters were subjected to multivariate analysis, only TIL was shown to be a significant independent factor for QpCR (RR 4.34, 95%RR 1.33–14.21, $P = 0.015$), but apoptosis was not significant (Table 5).

Survival analyses

In 92 patients with TNBC, disease-free survival (DFS) curves differed significantly between pCR and non-pCR groups (5-year DFS rate 93% vs. 66%, $P = 0.019$), between QpCR and non-QpCR groups (5-year DFS rate 91% vs. 64%, $P = 0.010$), and between the group of pCR in both primary tumor and axillary lymph nodes and others (5-year DFS rate 92% vs. 68%, $P = 0.043$) (Fig. 3). In TNBC, patients with a high TIL score tumor showed

slightly higher 5-year DFS rate than patients with a low TIL score tumor (77% vs. 70%), but the difference was not significant statistically ($P = 0.58$) (Fig. 4).

Discussion

Breast cancer has been shown to be a heterogeneous disease, and each intrinsic subtype of breast cancer differs in terms of gene expression and molecular features [1–5]. Previous studies reported differences between breast cancer subtypes in the pCR rate after primary chemotherapy [8, 10]: Rouzier et al. reported that the pCR rate after anthracycline and taxane chemotherapy in patients with luminal subtypes was 6%, while patients with both the basal-like and erbB2+ (HER2) subtypes had a pCR rate of 45%, based on classification using a “breast intrinsic” gene set [8]. Carey et al. also reported differences in the chemosensitivity of breast cancer subtypes when classified by immunohistochemistry: pCR rates after treatment with anthracycline either alone or in combination with taxane were 27, 36, and 7% for TNBC, and the HER2 and luminal subtypes, respectively [10]. In the present study, we confirmed that the pCR rate, QpCR rate, and the pCR rate in both the primary site and lymph nodes were significantly higher in patients with TNBC and tumors of the HR–/HER2+ subtype than in those with tumors of the HR+/HER2– subtype.

The proportions of cases showing a high TIL score (3, 4 or 5) and high apoptosis (score 2) were larger in TNBC and the HR–/HER2+ subtype than in the HR+/HER2– subtype. In addition, both TIL score and apoptosis were significantly associated with a response to NAC in TNBC, while in the HR–/HER2+ subtype and the HR+/HER2– subtype, these parameters were not significantly associated with pCR or QpCR. Because we used statistical tests on multiple related hypotheses, i.e., pCR, QpCR, and pCR in both the primary tumor and axillary lymph nodes, the data acquired should be considered exploratory. Nonetheless, these results suggest that patients with a high immune response to TNBC were more likely to show pCR, and that the immune component played a substantial role in the response of TNBC to NAC.

Although conflicting results have been reported [20, 21], earlier studies revealed a relationship between high lymphocyte infiltration and good prognosis in patients with breast cancer [22–25]. However, breast cancer subtypes were not taken into consideration in these studies. Kreike et al. demonstrated that a large amount of lymphocytic infiltrate was a significant indicator of longer distant metastasis-free survival in patients with TNBC [26]. In several studies, changes in TIL score or in the percentage in a certain subset of T cells were shown to be correlated

Fig. 3 Disease-free survival curves for patients with primary triple-negative breast cancer (TNBC) after neoadjuvant chemotherapy. **a** Survival curves for (a) patient group that showed pCR (NSABP B-18) and (b) patient group that showed non-pCR. Curves for two groups are significantly different (5-year DFS rate 93% vs. 66%, $P = 0.019$). **b** Survival curves for (a) patient group that showed QpCR (JBCRG) and (b) patient group that showed non-QpCR. Curves for two groups are significantly different (5-year DFS rate 91% vs. 64%, $P = 0.010$). **c** Survival curves for (a) patient group that showed pCR and (b) patient group that showed non-pCR in both primary tumor and axillary lymph nodes and others. Curves for two groups are significantly different (5-year DFS rate 92% vs. 68%, $P = 0.043$)

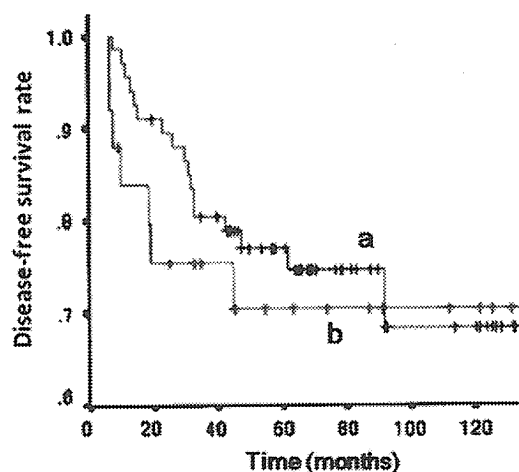
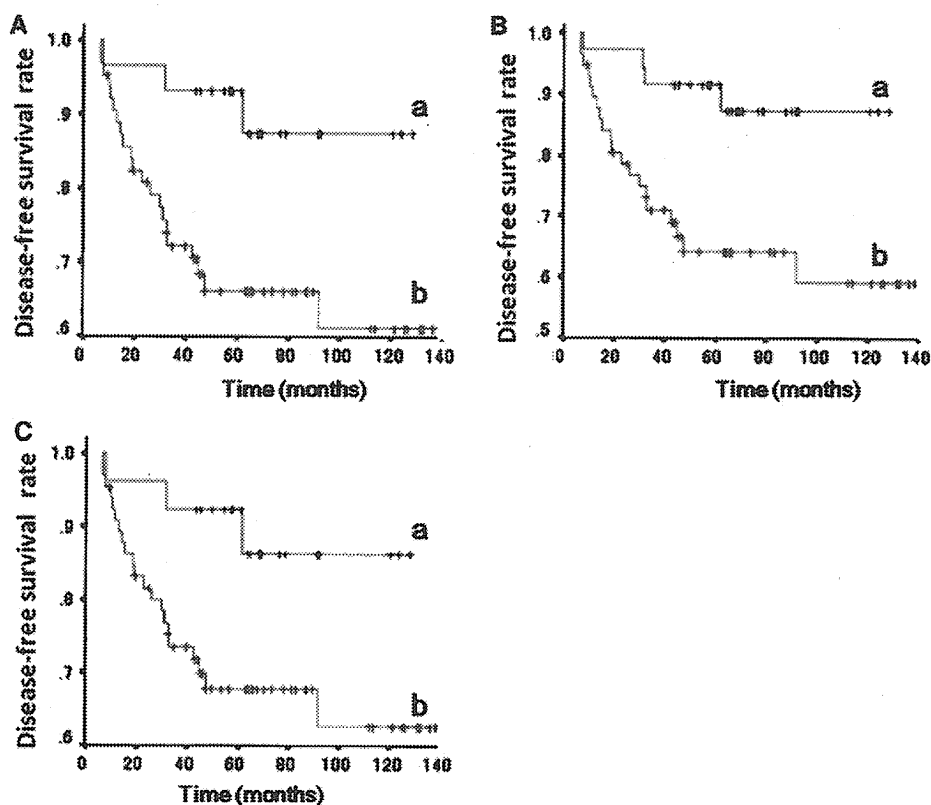


Fig. 4 Disease-free survival curves for patients with primary triple-negative breast cancer (TNBC) after neoadjuvant chemotherapy, stratified by the score of tumor infiltrating lymphocytes (TIL). **a** High TIL score group ($n = 67$). **b** Low TIL score group ($n = 25$). Although the 5-year disease-free survival rate was slightly higher in the high TIL score group (77%) than in the low TIL score group (70%), these two curves did not differ significantly ($P = 0.58$)

with pCR to neoadjuvant chemotherapy of breast cancer [27, 28].

It is also possible that gene expression associated with chemosensitivity and prognosis differs among breast cancer

subtypes. Teschendorff et al. also reported that a high level of gene expression representing an immune response was correlated with the better prognosis of patients with ER-negative breast cancer [29]. In fact, Rouzier et al. demonstrated that the genes predictive of pCR differed between the basal-like subtype and the HER2 subtype [8]. Furthermore, Desmedt et al. revealed that the gene expression modules associated with clinical outcome were different between the ER-/HER2- and HER2+ tumors: immune response genes only in the former and both tumor invasion and immune response genes in the latter [5]. Their results were consistent with those of the present study, which demonstrated a significant correlation between the presence of TIL and pCR/QpCR rate in TNBC, but the correlation was only marginal in the HR-/HER2+ subtype. Therefore, the molecular mechanisms determining chemosensitivity may differ between the basal-like and HR-/HER2+ subtypes.

We demonstrated a tendency of correlation between apoptosis and response to NAC in TNBC. Although Desmedt et al. examined the gene expression module associated with apoptosis, there was no association between expression of this gene set and prognosis in any of the breast cancer subtypes examined [5]. Because apoptosis has been defined as programmed cell death, and is usually unaccompanied by inflammation and cytokine release, apoptosis has been believed to be independent of TIL. In

the present study, there was no significant relationship between the presence of TIL and tumor cell apoptosis in TNBC. However, recent studies demonstrated that tumor cell death induced by chemotherapy can promote cytotoxic T-lymphocyte response that confers permanent antitumor immunity [30, 31]. We used histological examination only to identify apoptotic cancer cells. However, it would be more informative to add other techniques, such as the TUNEL method or immunohistochemistry, to identify apoptosis from multiple angles.

We revealed no correlation between the expression of basal-like markers and response to NAC in all of the breast subtypes examined. Although the significance of basal-like markers for clinical outcome is controversial [32–34], a lack of association between basal-like markers and chemosensitivity or prognosis has been demonstrated when breast cancers are divided into subtypes on the basis of ER and HER2 positivity [33, 34]. Nuclear p53 has been shown to be frequent in TNBC [35], but the significance of p53 as a predictive marker for pCR is also controversial [36]. In the present study we were unable to demonstrate any significant impact of p53 as such a marker.

It is unknown whether TILs cause susceptibility to chemotherapy, or they are simply a possible marker of chemosensitivity. There are reports that showed TILs are a predictor of response to neoadjuvant chemotherapy in breast cancer [37, 38]. Hornychova et al. reported that the infiltration of CD3⁺ T-lymphocytes and CD83⁺ dendritic cells were correlated with the effectiveness of primary chemotherapy, evaluated as pCR [38]. Denkert et al. showed that T-cell-related markers CD3D and CXCL9 expression were significantly associated with pCR [37]. Several studies suggested possible mechanisms of tumor-immune interaction in response to chemotherapy. pCR to neoadjuvant chemotherapy was shown to be associated with an immunologic profile combining the absence of immunosuppressive Foxp3⁺ regulatory T cells and the presence of a high number of CD8⁺ T cells and cytotoxic cells [28]. These reports suggest subsets of TILs caused susceptibility to chemotherapy.

In conclusion, we have demonstrated that the various breast cancer subtypes classified by ER, PgR, and HER2 status have different pathological characteristics and predictive factors for response to chemotherapy. TNBC with a high score for TIL and apoptosis is more likely to respond to chemotherapy. Therefore, in patients with TNBC, the immune response appears to influence on the response to chemotherapy. Further examination is warranted to elucidate the mechanism involved in the immune response component of chemosensitivity.

Acknowledgments We thank Mrs. Sachiko Miura and Mrs. Chizu Kina for excellent technical assistance. This study was supported in

part by grants from the Ministry of Health, Labor, and Welfare, Japan, the Ministry of Education, Culture, Sports, Science, and Technology, Japan, and the Princess Takamatsu Cancer Research Fund, Japan.

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Comparative study of the value of dual tracer PET/CT in evaluating breast cancer

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(Received April 9, 2012/Revised May 22, 2012/Accepted May 22, 2012/Accepted manuscript online May 28, 2012/Article first published online July 4, 2012)

The present study was conducted to assess the relationship between tumor uptake and pathologic findings using dual-tracer PET/computed tomography (CT) in patients with breast cancer. Seventy-four patients with breast cancer (mean age 54 years) who underwent ¹¹C-choline and 2-[¹⁸F]fluoro-2-deoxy-D-glucose (¹⁸F-FDG) PET/CT prior to surgery on the same day were enrolled in the present study. Images were reviewed by a board-certified radiologist and two nuclear medicine specialists who were unaware of any clinical information and a consensus was reached. Uptake patterns and measurements of dual tracers were compared with the pathologic findings of resected specimens as the reference standard. Mean (\pm SD) tumor size was 5.9 ± 3.2 cm. All primary tumors were identified on ¹⁸F-FDG PET/CT and ¹¹C-choline PET/CT. However, ¹⁸F-FDG PET/CT demonstrated focal uptake of the primary tumor with ($n = 38$; 51%) or without ($n = 36$; 49%) diffuse background breast uptake. Of the pathologic findings, multiple logistic regression analysis revealed an independent association between fibrocystic change and diffuse background breast uptake (odds ratio [OR] 8.57; 95% confidence interval [CI] 2.86–25.66; $P < 0.0001$). Tumors with higher histological grade, nuclear grade, structural grade, nuclear atypia, and mitosis had significantly higher maximum standardized uptake values (SUV_{max}) and tumor-to-background ratios (TBR) for both tracers. Multiple logistic regression analysis revealed that only the degree of mitosis was independently associated with a high SUV_{max} (OR 7.45; 95%CI 2.21–25.11; $P = 0.001$) and a high TBR (OR 5.41; 95%CI 1.13–25.96; $P = 0.035$) of ¹¹C-choline PET/CT. In conclusion, ¹¹C-choline may improve tumor delineation and reflect tumor aggressiveness on PET/CT in patients with breast cancer. (*Cancer Sci* 2012; 103: 1701–1707)

Positron emission tomography/computed tomography (PET/CT) with the glucose analog 2-[¹⁸F]fluoro-2-deoxy-D-glucose (¹⁸F-FDG) is recognized as an important tool in initial tumor evaluation, including staging, in the evaluation of treatment response, and in the assessment of recurrent disease for breast cancer.^(1,2) It has been reported that PET/CT adds incremental diagnostic confidence to PET in 60% of patients and in >50% of regions with increased ¹⁸F-FDG uptake.⁽³⁾ Tatsumi *et al.*⁽⁴⁾ concluded that PET/CT was preferable in evaluating breast cancer lesions in view of the level of diagnostic confidence that it allows. Regardless of the exact type of PET/CT fusion technique, ¹⁸F-FDG uptake in non-malignant conditions often leads to high background uptake on breast imaging.⁽⁵⁾

Histological changes are the cause of considerable variations and false-positive findings on breast imaging. Fibrocystic changes (FCC) are the most common of these conditions that can affect the assessment of imaging features on mammography^(6,7) and MRI.^(8,9) Similarly, there is evidence in the literature that ¹⁸F-FDG PET and accelerated glucose metabolism as

a result of FCC lead to false-positive findings and difficulty in determining the boundary of specificity.⁽¹⁰⁾

Choline is an essential component of the cell membrane and choline uptake is upregulated by choline kinase- α , which catalyzes the phosphorylation of choline.^(11,12) In mammary epithelial cells, levels of phosphocholine metabolites increase due to overexpression of choline kinase- α , which is regulated by the mitogen-activated protein kinase (MAPK) pathway.^(11–13) Recent clinical studies in patients with breast carcinoma undergoing molecular-targeted therapy suggest that ¹¹C-choline uptake is 10-fold higher in aggressive breast carcinoma phenotypes and that the uptake of ¹¹C-choline on PET is correlated with tumor grade.⁽¹³⁾ Thus, ¹¹C-choline is considered a promising radiotracer for the evaluation of breast cancer in the clinical setting prior to treatment.

Although both data from ¹⁸F-FDG and ¹¹C-choline PET/CT allow more precise evaluation of the primary breast cancer, direct comparisons of these two tracers in breast cancer have not been made. In the present study, we sought to confirm and extend previous findings of ¹¹C-choline PET/CT studies by investigating the association between histological findings and the results of ¹⁸F-FDG PET/CT investigations in patients with breast cancer.

Materials and Methods

Patients. Seventy-four patients (mean age 54 years; range 25–89 years) with breast carcinoma were enrolled in the present retrospective dual PET/CT study between March 2008 and March 2010. Patients were eligible for inclusion in the study if they met the following criteria: (i) performance status 0 or 1; (ii) no concomitant malignancy; (iii) histologically proven breast carcinoma diagnosed by biopsy at least 1 month before; and (iv) no history of hormone therapy. All patients were required to provide written informed consent. A regimen of 5-fluorouracil, epirubicin, and cyclophosphamide (FEC) plus paclitaxel was used as neoadjuvant chemotherapy in 32 patients (43%). As a rule, hormone therapy was introduced after completion of imaging studies if needed. Our institutional review board (National Cancer Center Hospital, Tokyo, Japan) approved the present study, which complied with the Health Insurance Portability and Accountability Act. The clinical records of all patients were available for review. All patients received surgery after imaging studies.

Phantom study. A phantom study of PET/CT was performed prior to the clinical study at two institutions to clarify the optimum conditions for data acquisition and to ensure quality control.⁽¹⁴⁾ Studies were performed with a whole-body PET/CT

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scanner (Aquiduo PCA-7000B; Toshiba Medical Systems, Tochigi, Japan). The CT component of the scanner has a 16-row detector. We used an NEMA image quality (IQ) phantom (NU 2-2001) for cross calibration, because this type of phantom is used in many institutions and data regarding the estimation of the optimum time are available. The radioactivity concentration of the background was set at 2.6 ± 0.2 kBq/mL ^{18}F -FDG, similar to that in clinical settings. The radioactivity concentration of the hot portion was fourfold greater than that of the background. Data were collected over a period of 2–5 min in the dynamic acquisition mode and for 30 min in the static acquisition mode. The data acquired, including normalization data, cross-calibration data, blank scan data, and transmission data, were assessed for visual inspection, phantom noise equivalent count ($\text{NEC}_{\text{phantom}}$), percentage contrast ($Q_{\text{H}10\text{ mm}}$) and percentage background variability ($N_{10\text{ mm}}$). The preferred parameters pertinent to the clinical condition were $\text{NEC}_{\text{phantom}} > 10.4$ (counts), $N_{10\text{ mm}} < 6.2\%$, and $Q_{\text{H}10\text{ mm}}/N_{10\text{ mm}} > 1.9\%$. After a review of the data analyses, the optimum conditions for the PET/CT were determined as follows: data acquisition, 180 s for one bed; field-of-view, 500 mm; iteration, 4; subset, 14; matrix size, 128×128 ; filter, Gaussian 8 mm in full width at half maximum; reconstruction, ordered-subsets expectation maximization (OSEM).

Data acquisition. ^{11}C -Choline was synthesized using a commercially available module, as described by Hara *et al.*⁽¹⁵⁾ Prior to the ^{11}C -choline PET/CT study, patients fasted for at least 6 h. Immediately after they had evacuated their bladder, patients were placed in a supine, arm-up position. For the PET/CT, low-dose CT data were first acquired at 120 kVp using an autoexposure control system (beam pitch 0.875 or 1 and 1.5 or 2 mm \times 16-row mode). Data acquisition was performed for each patient from the top of the skull to the mid-thigh. Patients maintained normal shallow respiration during the three-dimensional acquisition of CT scans. No iodinated contrast material was administered. Acquisition of emission scans from the head to the mid-thigh was started 5 min after intravenous administration of a mean ^{11}C -choline dose of 475 MBq (range 469–491 MBq). The ^{18}F -FDG PET/CT study was performed 1 h after the ^{11}C -choline PET/CT study in all patients. Patients received an intravenous injection of 311 MBq (range 197–397 MBq) ^{18}F -FDG with an uptake phase at 64 ± 5 min.

Image interpretation. Dedicated software (Vox-base SP1000 workstation; J-MAC Systems, Sapporo, Japan) was used to review all PET, CT, and coregistered PET/CT images in all standard planes. Images were analyzed visually and quantitatively by two independent reviewers, who recorded their findings after reaching a consensus. A region of interest (ROI) was outlined within areas of increased uptake and measured on each slice. When the lesion was extensively heterogeneous, the ROI was set so as to cover all the components of the lesion. The diffuse pattern of breast was assigned to the breast that shows homogeneous accumulation greater than aortic blood except for the primary lesion. For quantitative interpretations, the standardized uptake value (SUV) was determined according to the standard formula, with activity in the ROI recorded as Bq/mL per injected dose (Bq) per weight (kg), but time decay correction for whole-body image acquisition was not performed. The maximum SUV (SUV_{max}) was recorded using the maximum pixel activity within the ROI. The tumor-to-background ratio (TBR) was calculated with reference to uptake in the contralateral breast.

Pathologic analysis. All patients underwent surgery. Each tumor was staged according to the TNM classification of the International Union against Cancer.⁽¹⁶⁾ Resected specimens were fixed in 10% buffered formalin and embedded in paraffin wax. Then, 4- μm sections were obtained in a plane perpendic-

ular to the long axis of the breast. Paraffin-embedded microslides were stained with H&E. Tissue grading, nuclear grading, and structural grading were done using the grading system of Elston and Ellis.⁽¹⁷⁾ Estrogen receptor (ER) and progesterone receptor status was evaluated using the H-scoring system of McCarty *et al.*⁽¹⁸⁾ Human epidermal growth factor-2 (HER-2/neu) was evaluated by immunostaining with 4B5 primary antibody. Evaluation of the primary lesion was based on the following pathologic findings: FCC, differentiation, subtype, location, diameter of the invasive component, diameter of the non-invasive component, ratio of the invasive component in the tumor (%), tissue grading, nuclear grading, structural grading, nuclear atypia, mitosis, necrosis, fat invasion, cutaneous invasion, muscular invasion, ER status, progesterone receptor status, and HER-2/neu status. In the present study, “non-invasive component” referred to ductal carcinoma *in situ* (DCIS).

Statistical analysis. The Chi-squared test or Fisher’s exact probability test were used to compare pathologic findings associated with PET/CT findings. In addition, the Wald test and 95% confidence intervals (CI) were used to evaluate the statistical significance of individual variables. To determine relationships of SUV and TBR between the two tracers, we used Spearman rank correlation. Comparisons of mean values between groups were made using Student’s *t*-test or analysis of variance (ANOVA) with Bonferroni’s adjustment for multiple comparisons. Parsimonious univariate and multivariate logistic regression models were used to measure independent associations with PET/CT findings. Statistical tests used a two-sided significance level of 0.05. Statistical analyses were performed using PASW Statistics 19 (IBM, Tokyo, Japan).

Results

In all, 74 patients completed the study procedures. The demographic data for all patients are given in Table 1. There were 66 patients (89%) with invasive tumors, 60 of which were ductal carcinoma and six lobar carcinoma. Eight patients (11%) had non-invasive ductal carcinoma.

All primary tumors were identified on ^{18}F -FDG PET/CT and ^{11}C -choline PET/CT (Fig. 1). The SUV_{max} of ^{11}C -choline PET/CT was significantly lower than that of ^{18}F -FDG PET/CT ($P = 0.002$; Table 2). Conversely, the TBR of ^{11}C -choline PET/CT was significantly higher than that of ^{18}F -FDG PET/CT ($P < 0.0001$; Table 2). Using ^{18}F -FDG PET/CT, focal uptake of the primary tumor with ($n = 38$ [51%]; Fig. 2) or without ($n = 36$ [49%]) diffuse background breast uptake was demonstrated. Conversely, ^{11}C -choline PET/CT showed only focal uptake of the primary tumor in all patients. There were

Table 1. Patient demographics

Age (years)	54 \pm 13 (24–78)
Tumor side	
Right	44 (59)
Left	30 (41)
Tumor size (cm)	5.9 \pm 3.2 (1.8–12.0)
Main location	
Medial upper quadrant	8 (11)
Medial lower quadrant	8 (11)
Lateral upper quadrant	46 (62)
Lateral lower quadrant	6 (8)
Central	2 (3)
Invasive tumor	66 (89)
Non-invasive tumor	8 (11)

Data are given as the mean \pm SD, with the range in parentheses, or as the number of patients in each group with percentages in parentheses.

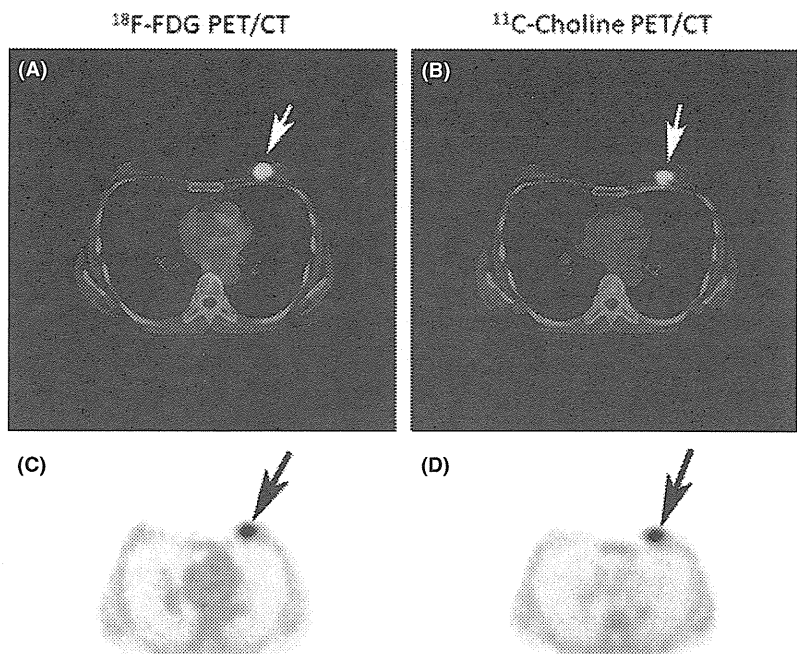


Fig. 1. Results for a 51-year-old woman with invasive ductal carcinoma of the left breast. (a,c) ^{18}F -fluorodeoxyglucose (FDG) PET/computed tomography (CT) images (fusion image: a; PET alone: c) reveal a focal hypermetabolic focus in the primary tumor (arrows). The maximum standardized uptake value (SUV_{max}) was 5.5 and the tumor-to-background ratio (TBR) was 47.0. (b,d) Transverse ^{11}C -choline PET/CT images (fusion image: b; PET alone: d) also reveal a focal hypermetabolic focus in the primary tumor (arrows). The SUV_{max} was 5.0 and the TBR was 137.5. On microscopy, the tumor contained 100% invasive component.

Table 2. Computed tomography (CT)/PET measurements and pathologic components with or without diffuse background breast uptake on ^{18}F -fluorodeoxyglucose PET/CT

	Total	With diffuse uptake	Without diffuse uptake	<i>P</i> -value
^{11}C -Choline uptake of tumor				
SUV_{max} (g/mL)	3.7 ± 2.9	3.6 ± 3.5	3.8 ± 2.0	0.789
TBR	8.0 ± 6.0	7.7 ± 9.9	8.3 ± 9.8	0.709
^{18}F -FDG uptake of tumor				
SUV_{max} (g/mL)	4.4 ± 3.1	4.6 ± 3.6	4.2 ± 2.5	0.571
TBR	3.7 ± 2.7	3.2 ± 2.4	4.5 ± 2.9	0.016
Diameter of invasive tumor (cm)	4.1 ± 3.5	4.2 ± 4.0	4.0 ± 3.0	0.800
Diameter of non-invasive tumor (cm)	1.8 ± 2.3	2.6 ± 2.9	0.9 ± 1.0	0.002
% Invasive component	66.0 ± 36.5	54.0 ± 41.4	78.6 ± 25.4	0.003

FDG, fluorodeoxyglucose; SUV_{max} , maximum standardized uptake value; TBR, tumor-to-background ratio.

significant differences between patients with or without diffuse background breast uptake on ^{18}F -FDG PET/CT for TBR of ^{18}F -FDG (Table 2). There was no interaction between ^{11}C -choline uptake and background breast uptake patterns on ^{18}F -FDG PET/CT. There were significant differences for the diameter of the non-invasive component and the percentage invasive component between patients with and without diffuse background breast uptake on ^{18}F -FDG PET/CT (Table 2).

The pathologic findings and background breast uptake patterns on ^{18}F -FDG PET/CT are listed in Table 3. Patients with

diffuse background breast uptake had significantly different values for percentage invasive component, FCC, necrosis, and triple negative tumor compared with patients without diffuse background breast uptake. There were no significant differences between the two groups in histologic grade, nuclear grade, structural grade, nuclear atypia, mitosis, fat invasion, or cutaneous invasion. Nor were there any significant differences in hormone receptor status between the two groups, specifically HER-2/neu, ER, and progesterone receptors. Only FCC showed an independent association with diffuse background breast uptake on multiple logistic regression analysis (OR 8.57; 95% CI 2.86–25.66; $P < 0.0001$).

There was a modest correlation between the diameter of the invasive tumor and SUV_{max} ($P < 0.0001$) or TBR ($P = 0.006$) on ^{18}F -FDG PET/CT (Table 4). Similar trends were found between the diameter of the invasive tumor and SUV_{max} ($P < 0.0001$) and TBR ($P < 0.0001$) on ^{11}C -choline PET/CT (Table 5). The TBR on ^{11}C -choline PET/CT also showed a modest correlation with the percentage invasive component ($P = 0.047$). The diameter of the non-invasive tumor was not correlated with SUV_{max} or TBR on either ^{18}F -FDG or ^{11}C -choline PET/CT.

Pathologic characteristics and tracer uptake are summarized in Table 6. Tumors with a higher histologic grade, nuclear grade, structural grade, nuclear atypia, and mitosis showed significantly higher SUV_{max} and TBR for both ^{18}F -FDG and ^{11}C -choline PET/CT. Tumors without expression of hormone receptors, including ER and progesterone receptors, and triple negative tumors showed significantly higher SUV_{max} and TBR for both ^{18}F -FDG and ^{11}C -choline PET/CT. Tumors expressing FCC and fat invasion were more likely to have high SUV_{max} and TBR on ^{11}C -choline PET/CT, but these differences were not identified in the TBR of ^{18}F -FDG PET/CT. In addition, tumors with necrosis and cutaneous invasion were found to have greater SUV_{max} and TBR only on ^{11}C -choline PET/CT. There was no significant association between the SUV_{max} or TBR and the percentage of invasive component or the HER-2/neu status for both tracers. After adjusting for age and tumor size, multiple logistic regression analysis revealed that the degree of mitosis was independently associated with high

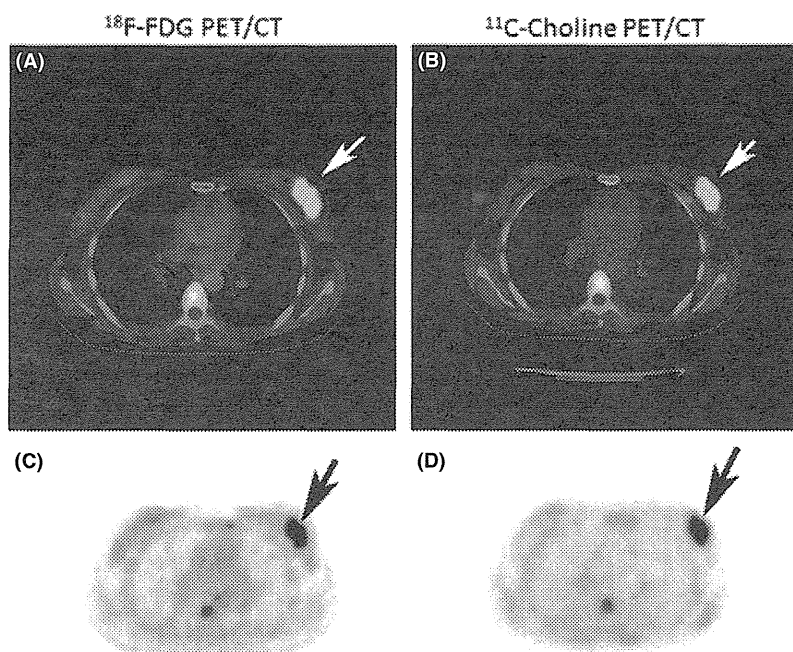


Fig. 2. Results for a 47-year-old woman with invasive scirrhous carcinoma of the left breast. (a,c) ^{18}F -fluorodeoxyglucose (FDG) PET/computed tomography (CT) images (fusion image: a; PET alone: c) reveal a focal hypermetabolic focus (arrows) of the primary tumor with diffuse background breast uptake. The maximum standardized uptake value (SUV_{max}) was 5.4 and the tumor-to-background ratio (TBR) was 35.7. (b,d) Transverse ^{11}C -choline PET/CT images (fusion image: b; PET alone: d) reveal only a focal hypermetabolic focus in the primary tumor (arrows). The SUV_{max} was 5.0 and the TBR was 125.0. On microscopy, the tumor contained 15% invasive component. Diffuse fibrocystic changes were found in the background breast.

SUV_{max} (OR 7.45; 95% CI 2.21–25.11; $P = 0.001$) and high TBR (OR 5.41; 95% CI; 1.13–25.96; $P = 0.035$) of ^{11}C -choline PET/CT.

Discussion

The present study examined the association between dual-tracer uptake and histological background in breast cancer. Despite positive correlations for SUV_{max} or TBR with ^{18}F -FDG and ^{11}C -choline, mitosis was found to be correlated with ^{11}C -choline uptake only, which reflects tumor aggressiveness reported in the previous study of patients with breast cancer.⁽¹⁹⁾ The results also reveal that diffuse background breast uptake on ^{18}F -FDG PET/CT depends on FCC and this pattern of uptake was not identified in any patients on ^{11}C -choline PET/CT. Our findings suggest that ^{11}C -choline may be feasible for the imaging of breast cancer particularly for patients with underlying FCC in whom mammography and ^{18}F -FDG PET/CT are limited.

Our observation of a positive correlation between mitosis and ^{11}C -choline uptake supports results reported in previous studies.^(13,19) This phenomenon was not affected by the underlying histological background because comparative correlation coefficients of SUV_{max} and TBR were similar on ^{11}C -choline PET/CT. Furthermore, the association between mitosis and ^{18}F -FDG uptake was not observed, regardless of positive correlation between ^{18}F -FDG and ^{11}C -choline uptake. This discrepancy in terms of mitosis and tracer uptake in our patients is presumably caused by differences in the degree of tracer uptake.

The present study demonstrated that there were significant differences in the diameter of the non-invasive component and the percentage invasive component between patients with and without diffuse background breast uptake on ^{18}F -FDG PET/CT. However, the SUV_{max} and TBR of both tracers were similar between patients with or without diffuse background breast uptake on ^{18}F -FDG PET/CT. These results suggest that the non-invasive component of breast cancer, which refers to the DCIS component in the present study, cannot be depicted by both tracers. These findings are consistent with that of another study that suggested DCIS could not be precisely visualized by PET.⁽¹⁾ Neubauer *et al.*⁽²⁰⁾ suggested that the DCIS component

could be detected by dynamic contrast-enhanced MRI, but the specificity was unfavorable because of an overlap in kinetic curve appearance. A major limitation of previous studies, as well as the present study, is that whole-body PET/CT scanners were used to evaluate primary lesion of the breast.

Fibrocystic changes are the most common diffuse benign condition of the breast related to changes in responses to estrogen and progesterone. The histology of FCC varies considerably and includes cysts, apocrine metaplasia, fibrosis, calcification, ductal hyperplasia, adenosis, and fibroadenomatous changes.^(21,22) Because of its diverse appearances and kinetic features, FCC is major cause of false-positive findings on MRI.^(23–25) As for PET studies, Yutani *et al.*⁽²⁶⁾ have previously explored the ^{18}F -FDG uptake of FCC in 38 patients with breast cancer, providing evidence that diffuse ^{18}F -FDG uptake caused by accompanying FCC obscures uptake by the primary tumor. Palmedo *et al.*⁽²⁷⁾ have confirmed that FCC is a major cause of reduced specificity in the detection of primary breast cancers on ^{18}F -FDG PET. Furthermore, Kole *et al.*⁽²⁸⁾ compared the detectability of primary lesions between ^{18}F -FDG PET and ^{11}C -tyrosine PET in patients with breast cancer and concluded that the visual assessment and delineation of the primary tumor were complicated only on ^{18}F -FDG PET when the contralateral breast tissue served as the control because FCC is a bilateral disease. As far as we were aware, the present study is the first that has been designed to evaluate the primary lesion of breast cancer using the dual tracers of ^{18}F -FDG and ^{11}C -choline. However, considering the high incidence of FCC, PET tracers including ^{11}C -tyrosine and ^{11}C -choline in addition to ^{18}F -FDG are more likely to fulfill specificity expectations.

The exact mechanism of ^{11}C -choline uptake by tumor cells is largely unknown; however, ^{11}C -choline has been proposed as a marker of the extracellular receptor kinase/MAPK pathway, exhibits significant uptake in tumor tissues, and is regarded as a favorable tracer for breast cancer.⁽¹³⁾ ^{11}C -Choline uptake may occur via a choline-specific transporter protein that is overexpressed in the cell membranes of breast cancer. ^{11}C -Choline is phosphorylated by choline kinase, which is upregulated in tumor cells for the synthesis of phosphatidylcholine, and is retained within tumor cells.^(11,12) Phosphatidylcholine is an essential

Table 3. Pathologic characteristics and background breast uptake on ¹⁸F-fludeoxyglucose PET/computed tomography

	No. patients		P-value
	With diffuse uptake	Without diffuse uptake	
Invasive component			
>30%	20	32	0.001
<30%	18	4	
Fibrocystic change			
Present	24	6	<0.0001
Absent	14	30	
Histologic grade			
1 or 2	16	22	0.102
3	22	14	
Nuclear grade			
1 or 2	16	22	0.102
3	22	14	
Structural grade			
1 or 2	14	18	0.253
3	24	18	
Nuclear atypia			
1 or 2	16	20	0.247
3	22	16	
Mitosis			
1 or 2	26	20	0.254
3	12	16	
Necrosis			
Present	22	8	0.002
Absent	16	28	
Fat invasion			
Present	22	24	0.437
Absent	16	12	
Cutaneous invasion			
Present	4	4	0.163
Absent	32	32	
HER-2/neu receptor			
Positive	22	14	0.102
Negative	16	22	
Estrogen receptor			
Positive	28	21	0.163
Negative	10	15	
Progesterone receptor			
Positive	28	21	0.163
Negative	10	15	
Triple negative			
Yes	4	11	0.032
No	34	25	

component of cell membranes and is involved in the modulation of transmembrane signaling by carcinogenesis. Therefore, ¹¹C-choline metabolism is accelerated in cell proliferation and is enhanced with increasing tumor grade of breast cancer. In the present study, tumors with higher histologic grade, nuclear grade, structural grade, nuclear atypia, and mitosis showed significantly higher SUV_{max} and TBR for ¹¹C-choline PET/CT. These results are in accord with those of previous *in vivo* and *in vitro* studies.^(19,29)

¹¹C-Choline PET/CT has been introduced as feasible method for the evaluation of breast cancer. In the present study, tumors without ER or progesterone receptors and triple negative tumors showed greater uptake of ¹¹C-choline compared with control groups. This suggests that ¹¹C-choline uptake reflects tumor aggressiveness. In a study of 32 patients with pathologically proven breast cancer expressing ER, no association was found between ¹¹C-choline uptake and hormone

Table 4. Relationship between ¹⁸F-fludeoxyglucose uptake and invasive or non-invasive tumor components

	¹⁸ F-FDG			
	SUV _{max}	P-value	TBR	P-value
Diameter of invasive tumor	0.381	<0.0001	0.318	0.006
Diameter of non-invasive tumor	-0.058	0.625	-0.14	0.234
% Invasive component	0.126	0.286	0.189	0.089

FDG, fludeoxyglucose; SUV_{max}, maximum standardized uptake value; TBR, tumor-to-background ratio.

Table 5. Relationship between ¹¹C-choline uptake and invasive or non-invasive tumor components

	¹¹ C-Choline			
	SUV _{max}	P-value	TBR	P-value
Diameter of invasive tumor	0.425	<0.0001	0.537	<0.0001
Diameter of non-invasive tumor	0.038	0.745	-0.066	0.575
% Invasive component	0.125	0.29	0.232	0.047

SUV_{max}, maximum standardized uptake value; TBR, tumor-to-background ratio.

receptor status.⁽¹⁹⁾ The apparent discrepancy between the present study and those of the previous study⁽¹⁹⁾ may be due, in large part, to differences in the patient populations studied.

In the present study, tumors exhibiting fat invasion were more likely to have a high SUV_{max} and TBR on ¹¹C-choline PET/CT, but these differences were not identified in the TBR of ¹⁸F-FDG PET/CT. This appeared to be associated with diffuse ¹⁸F-FDG uptake of breast caused by accompanying FCC, which may obscure tumor delineation. The presence of necrosis or cutaneous invasion was also found to have an association with SUV_{max} and TBR on ¹¹C-choline PET/CT. Overall, our results are consistent with those reported in *in vivo* and *in vitro* studies, in which ¹¹C-choline uptake was found to reflect tumor aggressiveness of breast cancer.^(13,19)

The present study design had limitations. First, the present study was designed to assess tumor uptake of dual tracers prior to surgery. The results from a breast cancer patient population of will not fully explain the detectability of advanced or recurrent disease. Second, the present study was an observational study and not a clinical trial, which raises the possibility of confounding factors affecting the results. Third, although ¹¹C-choline is clearly a possible PET tracer for tumor localization in patients with breast cancer, its short half-life restricts its practical application. However, ¹⁸F-choline is a tracer with a longer half-life than that of ¹¹C-choline, and so ¹⁸F-choline may improve the accuracy of tumor localization. Additional comparative studies regarding detectability and pathologic correlation are needed to validate the findings of the present study. Although we found that ¹¹C-choline uptake reflected tumor aggressiveness in patients with breast cancer, we did not have any data regarding nodal status and follow-up management of the patients. Further studies are needed to clarify the relationship between ¹¹C-choline uptake and patient outcome with a long follow-up period.

In conclusion, the results of the present study suggest that ¹¹C-choline PET/CT allows for the evaluation of tumor aggressiveness and improves delineation of primary tumors compared with ¹⁸F-FDG PET/CT in patients with breast cancer. The results demonstrate the advantages and potential of ¹¹C-choline, but clinical evaluation with a long follow-up