

Prognostic impact of interleukin-6 and C-reactive protein on patients with breast cancer

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Abstract. The prognostic impacts of preoperative C-reactive protein (CRP) and interleukin (IL)-6 expression levels in patients with breast cancer remain controversial. A total of 55 female patients with invasive breast cancer were enrolled, and preoperative prognostic parameters including IL-6 and CRP were analyzed. Overall survival (OS) and recurrence-free survival (RFS) were estimated using the Kaplan-Meier method, and candidates' prognostic factors were examined using a Cox proportional hazard model. Using receiver operating characteristic curve analysis, IL-6 at 10.0 pg/ml and CRP at 0.12 mg/dl were determined as threshold values to predict OS and RFS, respectively. Patients with IL-6 ≥ 10.0 pg/ml had poorer OS compared with those with IL-6 < 10.0 pg/ml ($P=0.003$), and patients with CRP ≥ 0.12 mg/dl had poorer RFS compared with those with CRP < 0.12 mg/dl ($P<0.001$). Serum IL-6 level (hazard ratio, 13.230; 95% confidence interval, 1.285-136.214; $P=0.030$) and triple-negative subtype (hazard ratio, 11.739; 95% confidence interval, 1.415-97.362; $P=0.023$) were independent prognostic factors for OS, and CRP expression level was an independent prognostic factor for RFS in patients with breast cancer (hazard ratio, 18.571; 95% confidence interval, 2.240-153.949; $P=0.007$). In patients with invasive breast cancer, preoperative serum IL-6 and

triple-negative subtype may be independent prognostic factors for OS, while for RFS, preoperative CRP may be a more accurate prognostic factor compared with those currently established.

Introduction

Breast cancer is the most frequently occurring type of cancer with 1.7 million cases worldwide. In 2012 it was the leading cause of cancer-associated mortality, accounting for 521,900 cases in females (1). The identification of biomarkers for breast cancer, including estrogen receptor (ER), progesterone receptor (PgR) and the human epidermal receptor 2 (HER2), has enabled the prediction of patient prognosis and the establishment of novel therapeutic agents (2). Notably, clinicopathological surrogate definitions for molecular subtypes of breast cancer, including luminal A-like, luminal B-like with HER2 negative, luminal B-like with HER2 positive, HER2 positive with ER and PgR negative, and triple-negative have been used for treatment recommendations (3). However, other studies have reported controversial results for the use of molecular subtyping in the prediction of recurrence (4,5).

One of the hallmarks of cancer is angiogenesis in conjunction with systemic and local inflammation (6). Crosstalk between angiogenesis and inflammatory signaling pathways may also contribute to cancer progression. Interleukin (IL)-6 and C-reactive protein (CRP) are principal mediators and indicators of the inflammatory response. CRP, which was named for its capacity to precipitate C-polysaccharide of *Streptococcus pneumoniae*, is a sensitive and widely used inflammatory marker produced primarily in the liver, in response to IL-1, IL-6, tumor necrosis factor- α and IL-17 (7,8). CRP was reported to be associated with poor prognosis in nasopharyngeal (9), hepatocellular (10), pancreatic (11), colorectal (12), renal (13), urothelial (14) and prostate cancer (15), in addition to breast cancer (16,17). However, the association between CRP and the prognosis of patients with breast cancer remains controversial (18-20). The association between serum IL-6 expression levels and breast cancer has been reported in treatment prognosis (21-23) and resistance to chemotherapy (24), and other studies have reported an association between serum IL-6, CRP and breast cancer (25,26). Although previous studies (21-23) have investigated the

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Abbreviations: ER, estrogen receptor; PgR, progesterone receptor; HER2, the human epidermal receptor 2; IL, interleukin; CRP, C-reactive protein; VEGF, vascular endothelial growth factor; G-CSF, granulocyte colony-stimulating factor; sICAM, soluble intercellular adhesion molecule; BMI, body mass index; RBP, retinol binding protein; TTR, transthyretin; TF, transferrin; NLR, neutrophil-to-lymphocyte ratio; LMR, lymphocyte-to-monocyte ratio; PBMCs, Peripheral blood mononuclear cells; ROC, receiver operating characteristic; OS, overall survival; RFS, recurrence-free survival

Key words: IL-6, CRP, OS, RFS, breast cancer

association between IL-6 and prognosis in patients with metastasis, the prognostic impact of preoperative IL-6 levels remains to be elucidated.

Galectin-3, a β -galactoside binding lectin is one of the most highly investigated key factors promoting angiogenesis and inflammation in breast cancer (27-29). Galectin-3 has been reported to induce the secretion of angiogenic factors from blood vascular endothelial cells in vitro, including as IL-6, vascular endothelial growth factor (VEGF), granulocyte colony-stimulating factor (G-CSF), granulocyte macrophage colony-stimulating factor and soluble intercellular adhesion molecule (sICAM)-1 (30). However, the association between circulating galectin-3 levels and the prognosis of breast cancer has yet to be clarified. The present study aimed to evaluate the prognostic impact of preoperative CRP and IL-6 expression levels in patients with breast cancer, in conjunction with angiogenic, inflammatory, immunological and nutritional parameters.

Materials and methods

Patients. Sera from 64 female preoperative patients with invasive breast cancer were collected between March 2011 and May 2013 at the Department of Breast Surgery, Fukushima Medical University Hospital (Fukushima, Japan), prior to starting treatment. Among these patients, 55 underwent curative-intent surgery (either modified radical mastectomy or partial mastectomy followed by irradiation); the remaining 9 patients had distant metastasis, and were therefore excluded. Thus, 55 preoperative patients with breast cancer (median age, 52; range, 37-83 years) were ultimately enrolled in the present study. Of the 55 patients, 17 were preoperatively diagnosed as having T4 tumor (n=6) and/or axillary lymph node metastasis (n=22), and subsequently received neoadjuvant chemotherapy using fluorouracil, epirubicin and cyclophosphamide. Following surgery, cancer stage was determined pathologically according to the Tumor-Node-Metastasis (TNM) classification system of malignant tumors, published in the Union for International Cancer Control, 8th edition (31). A total of 29 patients received adjuvant chemotherapy, including docetaxel, paclitaxel or capecitabine, and subsequent treatment with neoadjuvant chemotherapy (n=17) due to a postoperative diagnosis of metastasis to the axillary lymph nodes (n=20), and/or having a triple-negative (n=8) or HER2 (n=4) subtype. Patient characteristics are summarized in Table I. The study protocol was approved by the Ethics Committee of Fukushima Medical University (approval no. 1095) and written informed consent was obtained from the enrolled patients.

Measurement of parameters. Using an ELISA (R&D Systems, Inc., Minneapolis, MN, USA) according to the manufacturer's protocol, patient sera were evaluated to determine the concentrations of galectin-3 (cat. no. DGAL30), IL-6 (cat. no. D6050), VEGF (cat. no. DVE00), sICAM-1 (cat. no. DCD540) and G-CSF (cat. no. DCS50). The nutritional status of each patient was using a combination of the body mass index (BMI) and serum concentrations of total protein, albumin, retinol binding protein (RBP), transthyretin (TTR) and transferrin (TF). These parameters were assessed at the Central Clinical Laboratory of Fukushima Medical

Table I. Patient characteristics.

Category	Patients, n (total 55)	%
Pathological T factor		
1a	0	0.0
1b	1	1.8
1c	21	38.2
2	25	45.5
3	2	3.6
4a	0	0.0
4b	6	10.9
Pathological N factor		
0	35	63.6
1a	13	23.6
1b	0	0.0
1c	0	0.0
2a	6	11.0
2b	0	0.0
3a	0	0.0
3b	1	1.8
3c	0	0.0
Tumor-Node-Metastasis stage		
IA	17	31.0
IB	0	0.0
IIA	18	32.7
IIB	7	12.7
IIIA	7	12.7
IIIB	5	9.1
IIIC	1	1.8
Pathological subtype		
Papillo-tubular	18	32.7
Solid-tubular	9	16.4
Scirrhou	25	45.4
Other	3	5.5
Histological grade		
1	24	43.6
2	17	30.9
3	11	20.0
Not graded	3	5.5
Molecular subtype		
Luminal A-like	19	34.5
Luminal B/HER-	9	16.4
Luminal B/HER+	15	27.3
HER2	4	7.3
Triple-negative	8	14.5
Surgical procedure		
Bp+SN	19	34.5
Bp+AX	8	14.5
Bt+SN	11	20.0
Bt+AX	17	31.0

Luminal B/HER-, Luminal B-like human epidermal receptor 2 negative; Luminal B/HER+, Luminal B-like human epidermal receptor 2 positive; Bp+SN, partial mastectomy with sentinel node biopsy; Bp+AX, partial mastectomy with axillary lymph node dissection; Bt+SN, modified radical mastectomy with sentinel node biopsy; Bt+AX, modified radical mastectomy with axillary lymph node dissection.

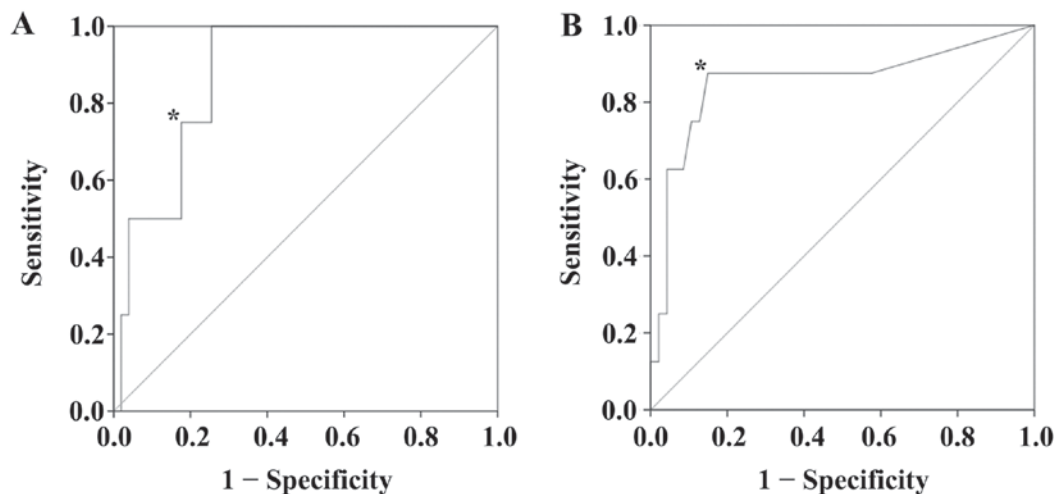


Figure 1. Receiver operating characteristic curve analysis. (A) The expression level of serum interleukin-6 was determined to be a biomarker for predicting the overall survival of patients with breast cancer ($P=0.013$). The AUC was 0.877, and 10.0 pg/ml was set as the cutoff value* (sensitivity=0.750, and specificity=0.824). (B) The CRP expression level was evaluated as a biomarker to predict RFS ($P=0.001$). The AUC was 0.854, and a level of 0.12 mg/dl was determined as the cutoff threshold* (sensitivity =0.875 and specificity =0.851). AUC, area under the curve; CRP, C-reactive protein; RFS, recurrence-free survival.

University Hospital. Indicators of inflammation, including CRP, white blood cell count, neutrophil, lymphocyte and monocyte counts, in addition to the neutrophil-to-lymphocyte ratio (NLR) and lymphocyte-to-monocyte ratio (LMR), were also verified.

The expression levels of the immunological cytokines IL-10, -12 and -17 were obtained. Peripheral blood mononuclear cells (PBMCs) were isolated using Ficoll-Hypaque columns (Pharmacia-Biotech, Uppsala, Sweden), and washed twice with RPMI-1640 medium (Wako Pure Chemical Industries, Ltd., Osaka, Japan). Isolated PBMCs were incubated at a concentration of 1×10^6 cells/ml in 1 ml RPMI-1640 (10% heat-inactivated fetal calf serum; Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA) for 24 h at 37°C and 5% CO_2 , with the following stimuli: i) 20 $\mu\text{g}/\text{ml}$ phytohemagglutinin for the IL-10 and IL-17 production assays; and ii) 0.01% *Staphylococcus aureus* Cowan strain 1 for the IL-12 assays. The supernatant was aliquoted and stored at -80°C until use. Supernatant samples were subsequently thawed and used to determine the concentrations of IL-10, IL-12, and IL-17 using an ELISA. Following thawing, samples were used only once, and not all blood samples were of sufficient volume for all measurements.

Statistical analysis. All statistical calculations were performed using SPSS® v.24 (IBM Corp., Armonk, NY, USA). Data are presented as frequencies or percentages for categorical variables, and means \pm standard deviation for continuous variables, unless otherwise indicated. For categorical clinical variables, the differences between two groups were evaluated using the χ^2 or the Fisher's exact test. The differences between continuous data were analyzed using the Mann-Whitney U test.

With regard to survival analysis, the mean observation period was 69.6 months (range, 55.3-81.4). The final assessment of disease status was made on December 12, 2017. Receiver operating characteristic (ROC) analysis was used to evaluate the prognostic value of the selected parameters.

Overall survival (OS) and recurrence-free survival (RFS) rates were determined using the Kaplan-Meier method, and the differences between the groups were assessed using the log-rank test. Prognostic factor candidates were subjected to univariate and multivariate analysis using a Cox proportional hazard model to identify independent predictors of prognosis; when $P < 0.1$ for univariate analysis, the candidate was also analyzed using multivariate analysis. $P < 0.05$ was considered to indicate a statistically significant difference.

Results

ROC curve analysis. Using ROC curve analysis, the serum expression level of IL-6 was determined to be a biomarker for the prediction of OS ($P=0.013$), with a cutoff value of 10.0 pg/ml (Fig. 1A), a sensitivity of 0.750 and a specificity of 0.824. Additionally, the serum expression level of CRP was evaluated as a biomarker to predict RFS ($P=0.001$), with a cutoff threshold of 0.12 mg/dl (Fig. 1B). At this cutoff value, the sensitivity was 0.875 and the specificity was 0.851.

Associations between serum expression levels of IL-6 and CRP, and patient characteristics. Table II displays patient characteristics according to serum IL-6 or CRP levels. There were no statistically significant differences in pathological T factor, pathological N factor, TNM stage, administration of neoadjuvant or adjuvant chemotherapy, ER, PgR, or HER2 expression levels, lymphatic invasion or microscopic vascular invasion associated with IL-6 and CRP expression levels between groups.

Serum VEGF levels in patients with $\text{CRP} \geq 0.12$ mg/dl (median, 472.0 ng/ml; range, 170.0-1,100.0 ng/ml) were significantly higher compared with those in patients with $\text{CRP} < 0.12$ mg/dl (median, 220.0 ng/ml; range, 74.0-1,460.7 ng/ml) ($P=0.011$). The Ki-67 labeling index of patients with $\text{CRP} \geq 0.12$ mg/dl (median, 35.3%; range, 3.1-80.0%) was significantly higher compared with that in patients with $\text{CRP} < 0.12$ mg/dl (median, 16.4%; range, 1.0-78.2%) ($P=0.035$). The

Table II. Patient characteristics according to IL-6 and CRP expression levels.

Category	IL-6 <10.0 pg/ml N=43	IL-6 ≥10.0 pg/ml N=12	P-value	CRP <0.12 pg/dl N=41	CRP ≥0.12 pg/dl N=14	P-value
pT			1.000			0.678
1+2	36	10		35	11	
3+4	7	2		6	3	
pN			0.319			0.755
0	29	6		26	8	
≥1	14	6		15	6	
Stage			0.477			0.736
I+II	33	8		31	10	
III+IV	10	4		10	4	
Neoadjuvant chemotherapy			0.735			0.742
-	29	9		29	9	
+	14	3		12	5	
Adjuvant chemotherapy			1.000			0.130
-	20	6		22	4	
+	23	6		19	10	
Estrogen receptor			0.689			0.477
-	8	3		8	4	
+	35	9		33	10	
Progesterone receptor			0.336			1.000
-	18	3		16	5	
+	25	9		25	9	
Human epidermal receptor 2			0.183			1.000
-	26	10		27	9	
+	17	2		14	5	
Lymphatic vessel invasion			0.320			0.346
-	24	4		23	5	
+	19	7		18	8	
Vascular invasion			1.000			1.000
-	27	7		26	8	
+	16	4		15	5	

IL-6, interleukin-6; CRP, C-reactive protein. One patient with IL-6 ≥ 10.0 pg/ml and CRP ≥ 0.12 mg/dl was not included in the evaluation of lymphatic or vascular invasion.

BMI of patients with CRP ≥0.12 mg/dl (median, 24.2 kg/m²; range, 16.4-29.6 kg/m²) was significantly higher compared with that in patients with CRP <0.12 mg/dl (median, 21.5 kg/m²; range, 20.8-40.3 kg/m²) (P=0.001; Fig. 2). When patients were categorized according to IL-6 expression level (cutoff, 10.0 pg/ml), there were no statistically significant differences among parameters. Serum expression levels of galectin-3, sICAM-1, RBP, TTR, TF, NLR and LMR, and the production of IL-12 and IL-17, were not significantly associated with the expression of IL-6 or CRP.

OS and RFS. As revealed in Fig. 3A and B, patients with IL-6 ≥10.0 pg/ml possessed poorer OS and RFS compared with those with IL-6 <10.0 pg/ml (P=0.010 and P=0.038, respectively). There were no statistically significant differences in OS between patients with different levels

of CRP (P=0.253; Fig. 3C), whereas patients with CRP ≥0.12 mg/dl exhibited a poorer RFS compared with those with CRP <0.12 mg/dl (P<0.001; Fig. 3D). Differences in the serum expression levels of galectin-3, VEGF, sICAM-1 and G-CSF did not significantly affect the prognosis of patients with breast cancer.

Cox proportional hazards model. The following prognostic factors were analyzed using a Cox proportional hazards model: pathological T factor (T1 + T2 vs. T3 + T4), pathological N factor (N0 vs. ≥ N1), TNM stage (I + II vs. III + IV), triple-negative subtype (no vs. yes), CRP (<0.12 mg/dl vs. ≥0.12 mg/dl) and serum IL-6 expression level (<10.0 pg/ml vs. ≥10.0 pg/ml). Triple-negative subtype (hazard ratio, 8.795; 95% confidence interval, 1.230-62.875; P=0.030) and serum IL-6 expression level (hazard ratio, 10.785; 95% confidence

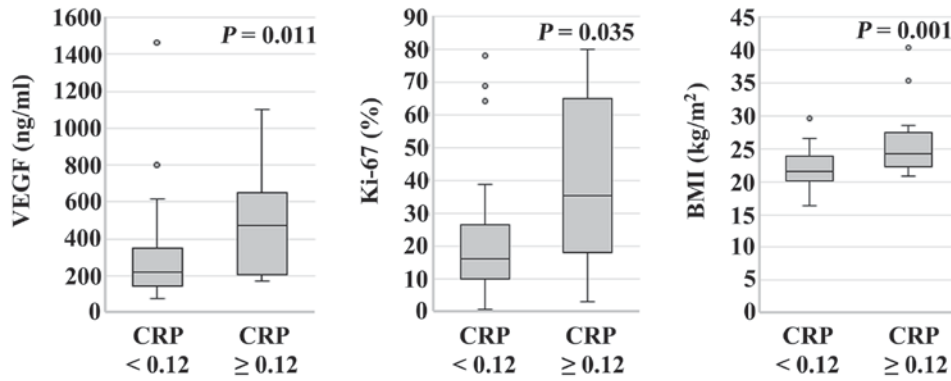


Figure 2. Association between CRP expression level and patient parameters. Serum VEGF levels in patients with CRP ≥ 0.12 mg/dl (median, 472.0 ng/ml; range, 170.0-1,100.0 ng/ml) were significantly higher compared with those in patients with CRP < 0.12 mg/dl (median, 220.0 ng/ml; range, 74.0-1,460.7 ng/ml) ($P=0.011$). Ki-67 labeling index in patients with CRP ≥ 0.12 mg/dl (median, 35.3%; range, 3.1-80.0%) was significantly higher compared with that of patients with CRP < 0.12 mg/dl (median, 16.4%; range, 1.0-78.2%) ($P=0.035$). BMI of patients with CRP ≥ 0.12 mg/dl (median, 24.2 kg/m²; range, 16.4-29.6 kg/m²) was significantly higher compared with that of patients with CRP < 0.12 mg/dl (median, 21.5 kg/m²; range, 20.8-40.3 kg/m²) ($P=0.001$). CRP, C-reactive protein; VEGF, vascular endothelial growth factor; BMI, body mass index.

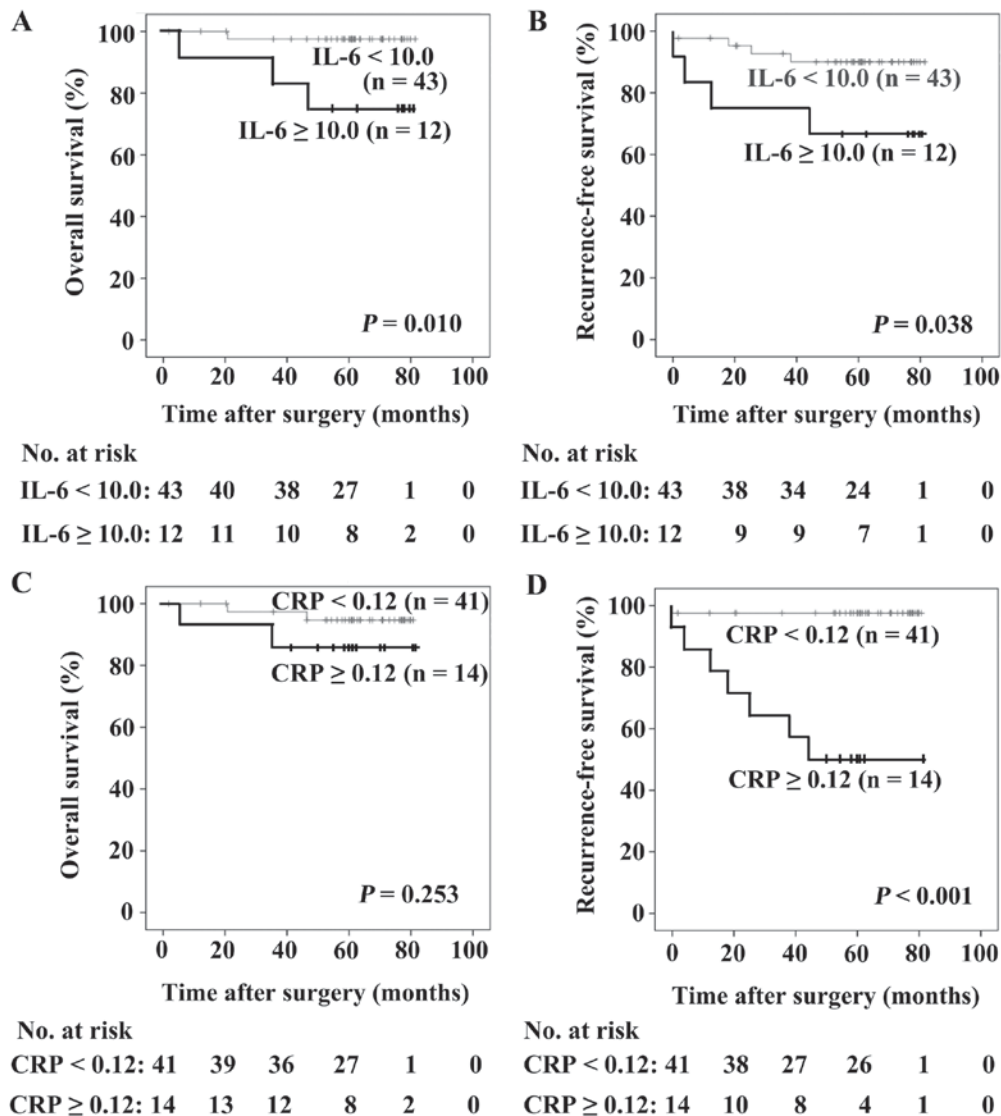


Figure 3. Patient survival calculated using the Kaplan-Meier method. (A) Patients with IL-6 ≥ 10.0 pg/ml exhibited poorer OS compared with those with IL-6 < 10.0 pg/ml ($P=0.010$). (B) Patients with IL-6 ≥ 10.0 pg/ml exhibited poorer RFS compared with those with IL-6 < 10.0 pg/ml ($P=0.038$). (C) There were no statistically significant differences in OS between patients with CRP < 0.12 mg/dl and with CRP ≥ 0.12 mg/dl ($P=0.253$). (D) Patients with CRP ≥ 0.12 mg/dl exhibited a poorer RFS compared with those with CRP < 0.12 mg/dl ($P<0.001$). CRP, C-reactive protein; IL-6, interleukin-6; OS, overall survival; RFS, recurrence-free survival.

Table III. Cox proportional hazards model.

A, Overall survival						
Category	Univariate analysis			Multivariate analysis		
	HR	95% CI	P-value	HR	95% CI	P-value
T	5.121	0.721-36.369	0.102	-	-	-
N	1.729	0.243-12.278	0.584	-	-	-
TNM stage	2.886	0.406-20.485	0.289	-	-	-
HG	7.198	0.652-79.503	0.107	-	-	-
TN	8.795	1.230-62.875	0.030	11.739	1.415-97.362	0.023
IL-6	10.785	1.122-103.693	0.039	13.230	1.285-136.214	0.030
CRP	2.970	0.41-21.100	0.277	-	-	-
B, Recurrence-free survival						
Category	Univariate analysis			Multivariate analysis		
	HR	95% CI	P-value	HR	95% CI	P-value
T	3.355	0.801-14.053	0.098	2.499	0.583-10.711	0.218
N	5.598	1.129-27.748	0.035	3.429	0.677-17.364	0.136
TNM stage	0.993	0.200-4.920	0.993	-	-	-
HG	3.172	0.708-14.206	0.131	-	-	-
TN	2.702	0.543-13.456	0.225	-	-	-
IL-6	3.637	0.733-18.040	0.114	-	-	-
CRP	23.865	2.930-194.408	0.003	18.571	2.240-153.949	0.007

HR, hazard ratio; CI, confidence interval; T, pathological T factor; N, pathological N factor; TNM, Tumor-Node-Metastasis; HG, histological grade; TN, triple-negative molecular subtype; IL-6, interleukin-6; CRP, C-reactive protein.

interval, 1.122-103.693; $P=0.039$) were significantly associated following univariate analysis. Triple-negative subtype (hazard ratio, 11.739; 95% confidence interval, 1.415-97.362; $P=0.023$) and serum IL-6 expression level were independent prognostic factors for the OS of patients with breast cancer. With regard to RFS, the pathological N factor (hazard ratio, 5.598; 95% confidence interval, 1.129-27.748; $P=0.035$) and CRP level (hazard ratio, 23.865; 95% confidence interval, 2.930-194.408; $P=0.003$) showed statistical significance following univariate analysis, and the CRP expression level was an independent prognostic factor for RFS in patients with breast cancer (hazard ratio, 18.571; 95% confidence interval, 2.240-153.949; $P=0.007$; Table III).

Discussion

The prognostic impact of preoperative serum IL-6 expression level has been reported previously, but based only on the results of Kaplan-Meier analysis (26). Furthermore, the prognostic impact of serum IL-6 expression level has only been observed in patients with metastatic disease (21-23). Although the prognostic impact of preoperative CRP has been reported (16,17), other studies have described controversial results (18-20). Thus, the present study aimed to evaluate the

prognostic impact of preoperative CRP and IL-6 expression levels in patients with breast cancer. It was revealed that preoperative expression levels of IL-6 and CRP effected OS and RFS, respectively.

IL-6 is secreted by various cell types, including breast cancer cells. The response to IL-6 in such cells depends on the expression of ERs (24). The ER-positive MCF-7 cell line does not secrete IL-6, whereas ER-negative MD-MBA-231 cells do (32,33). However, Fontanini *et al* (34) reported that expression levels of IL-6 correlated with ER expression. Ravishankaran and Karunanithi (25) reported that serum IL-6 expression levels correlated with the extent of tumor invasion, lymph node metastasis, distant metastasis and TNM staging. In the present study, IL-6 did not correlate with such tumor statuses.

In the present study, CRP expression level correlated with VEGF expression, Ki-67 labeling index and BMI. Expression levels of VEGF in breast cancer have been reported to correlate with poorer RFS (35). Although serum VEGF expression level was reportedly increased in patients with breast cancer, compared with healthy controls (36), numerous studies have suggested that it has no prognostic value (37,38). Highly proliferative tumors, including triple-negative breast cancer, show enhanced angiogenesis

and reportedly exhibit high levels of VEGF expression (39). Additionally, the Ki-67 labeling index has been associated with OS in breast cancer (40). In the present study, serum expression levels of VEGF and the Ki-67 labeling index had no significant association with OS or RFS; however, a significant correlation with higher CRP expression indicates that they may contribute to poor RFS. Furthermore, increased risk of breast cancer was associated with increased body weight and obesity in women, particularly in postmenopausal patients, and a significant link was observed between CRP expression levels and BMI (41). The suggested association between elevated CRP expression level, VEGF, Ki-67 and BMI warrants further investigation.

Limitations of the present study include its relatively small patient cohort and short observational period. Also, lifestyle factors that may influence IL-6 and/or CRP expression levels, including smoking, menopausal status, and other comorbidities (including diabetes and cardiovascular disease), were not taken into account.

For ROC curve analysis, 10.0 pg/ml IL-6 and 0.12 mg/dl CRP were determined as threshold values to predict OS and RFS, respectively. Patients with IL-6 ≥ 10.0 pg/ml exhibited poorer OS compared with those with IL-6 < 10.0 pg/ml, and patients with CRP ≥ 0.12 mg/dl exhibited poorer RFS compared with those with CRP < 0.12 mg/dl. In patients with breast cancer, serum IL-6 and CRP expression levels were independent prognostic factors for OS and RFS, respectively. Specifically, CRP is indicated to be a superior prognostic marker compared with established prognostic factors, including T factor, N factor and histological grade, supporting a possible association between inflammation and recurrence in breast cancer.

In conclusion, in patients with invasive breast cancer, preoperative serum IL-6 expression levels and triple-negative subtype may be independent prognostic factors for OS, while for RFS, preoperative CRP expression level may be a more accurate prognostic factor compared with those already established. Particular attention should be paid to patients with higher preoperative IL-6 and/or CRP expression levels during preoperative follow-up for breast cancer.

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Availability of data and materials

The datasets generated from the present study are available from the corresponding author upon reasonable request.

Authors' contribution

TS, MS and TO contributed to the conception, design, and integrity of this study. KG, YM, MN, KT and NA performed data acquisition, analysis, and interpretation. TS and MS drafted and critically revised the manuscript.

Ethics approval and consent to participate

The present study was approved by the Ethics Committee of Fukushima Medical University (approval no. 1095) and written informed consent was obtained from all enrolled patients. Additionally, all patient data were treated in accordance with the local privacy regulations.

Patient consent for publication

All patients provided written informed consent.

Competing interests

The authors declare that they have no competing interests.

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