

ORIGINAL ARTICLE

STAT3, a Poor Survival Predictor, Is Associated with Lymph Node Metastasis from Breast Cancer

Yujuan Chen, Jing Wang, Xiaodong Wang, Xuejuan Liu, Hongjiang Li, Qing Lv, Jingqiang Zhu, Bing Wei, Ying Tang

Department of Thyroid and Breast Surgery, Western China Hospital of Sichuan University, Chengdu, China

Purpose: The aim of this study is to explore signal transducer and activator of transcription 3 (STAT3) expression in breast cancer and to analyze the detailed mechanism that STAT3 contributes to the progression of breast cancer. **Methods:** We retrospectively analyzed the clinicopathologic characteristics and overall survival (OS) of 140 breast cancer patients after curative surgery, and detected STAT3 expression, phosphorylated STAT3 (pSTAT3) expression, Ki-67 expression, vascular endothelial growth factor (VEGF)-C and -D expression in breast cancer tissues, and adjacent nontumor tissues. Survival analysis and relationship analysis were adopted for demonstrated the important mechanism of STAT3 contribution to progression of breast cancer. **Results:** STAT3 expression, pSTAT3 expression, Ki-67 expression, VEGF-C expression, and VEGF-D expression in breast cancer tissues

were significantly higher than those in adjacent nontumor tissues, respectively. With survival analysis, only number of lymph node metastasis (N stage) was identified as the independent predictors of the OS of breast cancer patients. Besides, we demonstrated there was the most prominent correlation between STAT3 expression and lymph node metastasis in breast cancer tissues by using the multinomial regression method. **Conclusion:** STAT3, a poor survival biomarker potential association with lymph node metastasis, was suitable for predication the OS of breast cancer patients after curative resection.

Key Words: Breast neoplasms, Neoplasm metastasis, Prognosis, STAT3 transcription factor

INTRODUCTION

Breast cancer, as an increasing incidence malignancy, is the most common lethal disease in females [1]. Most breast cancer patients have lymph node metastasis when they are initially diagnosed or operated on, which results in a poor prognosis [2]. Although there was no significant improvement on disease-free survival (DFS) and overall survival (OS), lymph node dissection was recently identified to reduce the local recurrence rate in breast cancer patients with node micrometastasis [3]. To date, sentinel lymph node biopsy (SLNB) has become the standard of care as a staging procedure for breast cancer patients with clinically node-negative disease, and regional nodal irradiation after a positive SLNB has been considered as a reasonable alternative to axillary lymph node dissection

(ALND) [4]. ALND remains a standard method of treating regional disease not only in patients with clinically positive nodes but also in some SLNB positive patients [5], while the latest researches demonstrated that most SLNB positive patients would be candidates to avoid ALND if they underwent breast-conserving therapy (BCT) [6,7]. Therefore, the accurate evaluation of lymph node metastasis is very important to the procedure of surgery, in addition to the predication the prognosis of breast cancer patients. Unfortunately, there is not an authoritative method for detection the nodal metastasis from breast cancer before surgery. On the other hand, we have to admit the fact that there is not a satisfactorily biomarker with the high specificity and sensitivity for screen the relapse of patients following the mastectomy. Lymphatic metastasis represents a series of sequential processes that include the dissemination and invasion of tumor cells from the primary tumor into the surrounding stromal tissue, penetration of the tumor cells across the lymphatic walls, implantation in the regional lymph nodes, and extravasation and proliferation in the parenchyma of target organs [7]. In general, vascular endothelial growth factor (VEGF)-C and -D have been defined

Correspondence to: Jing Wang

Department of Thyroid and Breast Surgery, Western China Hospital of Sichuan University, Guoxuexiang 37, Wuhou District, Chengdu, China
Tel: +86-28-8542-2114, Fax: +86-28-8542-2114
E-mail: wangjinghxy@sohu.com

Received: November 8, 2012 Accepted: February 7, 2013

as lymphangiogenic growth factors and play an important role in tumor lymphangiogenesis via activation of the VEGF receptor (VEGFR)-3, which is expressed in lymphatic endothelial cells [8,9]. However, these two biomarkers can not provide the information of dissemination and invasion of tumor cells, penetration of the tumor cells, and so on. These aforementioned defects make VEGF-C and -D not to be widely applicable to diagnosis the lymph node metastasis or to screen the nodal relapse in clinic.

Signal transducer and activator of transcription 3 (STAT3), an important member of the STAT family, is a latent transcription factor that has been shown to act as an oncogene in several malignant diseases [10]. It is well known that STAT3 is activated (phosphorylated STAT3, pSTAT3) in response to the binding of a large number of cytokines, hormones and growth factors to their receptors. Actually, STAT3 has been demonstrated to up-regulate genes encoding apoptosis inhibitors (Bcl-xL, Mcl-1, and survivin), cell-cycle regulators (cyclin D1 and c-Myc) and angiogenesis inducers (e.g., VEGF), which are important molecular cues for tumor progression [11-13]. Recently, several investigators reported that STAT3 aberrant expression were significantly associated with the status of lymph node metastasis in various malignant disease [14-19]. Theoretically, STAT3 has the ability to promote the lymphatic metastasis from malignancy. The detailed correlation between lymph node metastasis and STAT3 expression is not elucidated.

In view of aforementioned causalities, we designed this study for exploration the implication of the STAT3 aberrant expression in breast cancer. They are as follows: 1) to elucidate the STAT3 aberrant expression for predication the OS of breast cancer patients after curative resection; and 2) to initially analyze the correlation between the STAT3 aberrant expression and lymph node metastasis from breast cancer.

METHODS

Patients

A total of 140 patients who underwent potentially curative resection for breast cancer at the Department of Thyroid and Breast Surgery, West China Hospital of Sichuan University between June 2005 and December 2005 were eligible for this study. Eligibility criteria for this study included: 1) histologically proven primary breast cancer, 2) no history of mastectomy or other malignancy, 3) a lack of noncurative surgical factors except for distant metastasis (such as liver, lung, brain, or bone-marrow metastasis) and supraclavicular lymph node metastasis, 4) ALND performed, and 5) no patients died during the initial hospital stay or for 1 month after surgery, 6) no neoadjuvant therapy (including chemotherapy, radiotherapy, and

hormone therapy).

Surgical treatment

All patients were operated on according to the potentially curative mastectomy plus ALND method. Curative resection was defined as a complete lack of grossly visible tumor tissue and metastatic lymph nodes remaining after resection, with pathologically negative resection margins. Primary tumors were resected en bloc. The choice of surgical procedure of mastectomy (radical or modified radical mastectomy) was made by the attending surgeon's preference, and based mainly on the breast cancer treatment guidelines. Surgical specimens were evaluated as recommended by 7th American Joint Committee on Cancer (AJCC) TNM classification for breast cancer.

Immunohistochemistry

Paraffin sections (4 μ m thick) were deparaffinized and rehydrated. Antigen retrieval treatment was done at 95°C for 40 minutes in 0.01 mol/L sodium citrate buffer (pH 6.0), and endogenous peroxidases were blocked using 3% hydrogen peroxide for 30 minutes. Purchased antibodies were: rabbit anti-STAT3 (BA0621, 1:100 dilution; Boster, Wuhan, China), rabbit anti-pSTAT3 (AP0247, 1:150 dilution; Bioworld, Nanjing, China), rabbit anti-VEGF-C (BA0548, 1:100 dilution; Boster), rabbit anti-VEGF-D (BA1461, 1:50 dilution; Boster), rabbit anti-Ki-67 (BA1653, 1:150 dilution; Boster), rabbit anti-estrogen receptor (ER) (BA3599, 1:100 dilution; Boster), Wuhan, China), rabbit anti-progesterone receptor (PR) (ZA-0255, 1:150 dilution; Zhongshan, Beijing, China), and rabbit anti-human epidermal growth factor receptor 2 (HER2) (RM-2112, 1:50 dilution; Lab Vision\Neomarker, Kalamazoo, USA). All sections were incubated overnight with the primary antibody at 4°C. The sections were then treated with peroxidase using the labeled polymer method with Peroxidase (Boster) for 30 minutes. Antibody binding was visualized using the Avidin Biotin Complex Elite Kit and 3,3'-diaminobenzidine according to the manufacturer's instructions (Molecular Cancer Laboratory, West China Hospital of Sichuan University, Chengdu, China). Sections were then counterstained in hematoxylin. For general negative controls, the primary antibody was replaced with PBS.

Microscopic assessment of STAT3, pSTAT3, VEGF-C, VEGF-D, Ki-67, ER, PR, and HER2 expression

All sections were assessed blindly by two independent observers, and in cases of assessing disagreement a third independent assessment was performed. Staining for STAT3, VEGF-C, and VEGF-D was considered potentially positive if there was cytoplasmic staining. Staining for HER2 was con-

sidered potentially positive if there was membranous staining. The grade of staining intensity of STAT3, VEGF-C, VEGF-D, and HER2 was rated on a scale from 0 to 3, with 0, indicating no staining; 1, weak staining; 2, moderate staining; and 3, strong staining. The intensity scores of 2 and 3 were considered positive staining. Positive staining for pSTAT3, Ki-67, ER, and PR was defined as >25% nuclear staining with more than a moderate intensity for the tumor cell.

Evaluated variables

The following clinicopathological variables were evaluated: 1) age at surgery (<50, or ≥50 years); 2) depth of primary tumor invasion (T stage, according to 7th AJCC TNM classification) (T1, T2, T3, or T4); 3) number of lymph node metastasis (N stage, according to 7th AJCC TNM classification) (N0, N1, N2, or N3); 4) TNM classification (I, II, III, or IV); 5) World Health Organization classification of primary tumor (I, II, or III); 6) STAT3 expression (negative or positive); 7) pSTAT3 expression (negative or positive); 8) VEGF-C expression (negative or positive); 9) VEGF-D expression (negative or positive); 10) Ki-67 expression (negative or positive); 11) ER expression (negative or positive); 12) PR expression (negative or positive); and 13) HER2 expression (negative or positive).

Ethics statement

The study was approved by the Research Ethics Committee of West China Hospital of Sichuan University (2012-251), China. Informed consent was obtained from all patients before participating in the study.

Statistical analysis

Categorical variables were statistically compared a chi-square or Fisher's exact test. Continuous data were shown as mean ± SD and were statistically compared using the Mann-Whitney test. The median OS was determined by using the Kaplan-Meier method, and log-rank test was used to determine significance. Factors that were deemed of potential importance on univariate analyses ($p < 0.05$) were included in the multivariate analyses. Multivariate analysis of OS was performed by means of the Cox proportional hazards model, using the forward procedure for variable selection. Hazard ratios (HR) and 95% confidence interval (CI) were generated. Akaike information criterion (AIC) and Bayesian information criterion (BIC) were performed in the multinomial logistical regression model for evaluation the best clinicopathological variable for predication the lymph node metastasis from breast cancer. The smaller are AIC value and BIC value, the better is clinicopathological variable for predication the lymph node metastasis. In all statistical analyses, significance was defined as $p < 0.05$

and the statistical significance was two sided. The OS analysis of all patients was initially completed in March 2012. All statistical analyses were performed with statistical analysis program package SPSS version 18.0 (SPSS Inc., Chicago, USA).

Follow-up

After curative surgery, all patients were followed every 3 months for 2 year, then every half a year or until death. The median follow-up for the entire cohort was 54 months (range, 18-79 months). The follow-up of all patients who were included in this study was completed in February 2012. Serum values of CEA and CA 15-3, B ultrasonography, chest X-ray, and radionuclide bone scan were obtained with every visit.

RESULTS

Patient characteristics

The clinicopathological characteristics and biomarkers' expression of 140 breast cancer patients after curative resection are shown in Table 1. The 5-year survival rate (5-YSR) of all enrolled patients was 83.5%, and 111 patients were alive when the follow-up was completed.

Immunohistochemical staining for STAT3, pSTAT3, VEGF-C, VEGF-D, Ki-67, ER, PR, and HER2 in breast cancer or adjacent nontumor tissues

STAT3, VEGF-C, and VEGF-D expression was observed in the cytoplasm. Staining for pSTAT3 and Ki-67 appeared in the cell nucleus. STAT3 positive expression, pSTAT3 positive expression, VEGF-C positive expression, VEGF-D positive expression, Ki-67 positive expression, ER positive expression, PR positive expression, and HER2 positive expression were respectively detected in 87 (62.1%), 67 (47.9%), 58 (41.1%), 34 (24.3%), 87 (62.1%), 64 (45.7%), 72 (51.4%), and 41 (29.3%) of 140 breast cancer specimens, whereas 27 (19.3%), 14 (10.0%), 31 (22.1%), 16 (11.4%), 20 (14.3%), 13 (9.3%), 16 (11.4%), and 0 (0%) of 140 adjacent nontumor tissue specimens were assessed as STAT3 positive expression, pSTAT3 positive expression, VEGF-C positive expression, VEGF-D positive expression, Ki-67 positive expression, ER positive expression, PR positive expression, and HER2 positive expression, respectively. There were significantly statistical differences in the positive expression incidences of STAT3 ($p = 0.024$), pSTAT3 ($p = 0.015$), VEGF-C ($p = 0.032$), VEGF-D ($p = 0.040$), Ki-67 ($p = 0.011$), ER ($p = 0.004$), PR ($p = 0.025$), and HER2 ($p < 0.001$), between breast cancer and adjacent nontumor tissue specimens, respectively (Figures 1-4).

Table 1. Clinicopathologic characteristics of 140 breast cancer patients

Characteristic	No. (%)
Gender	
Male	0 (0)
Female	140 (100)
Age at surgery (yr), mean \pm SD (range)	48.79 \pm 10.49 (32-77)
< 50	88 (62.9)
\geq 50	52 (37.1)
T stage*	
T1	44 (31.4)
T2	80 (57.1)
T3	16 (11.4)
No. of metastatic LN, mean \pm SD (range)	2.30 \pm 0.36 (0-18)
N stage*	
N0	76 (54.3)
N1	36 (25.7)
N2	16 (11.4)
N3	12 (8.6)
TNM classification*	
I	30 (21.4)
II	78 (55.7)
III	32 (22.9)
WHO classification of primary tumor	
I	10 (7.1)
II	42 (30.0)
III	88 (62.9)
STAT3 expression	
Negative	53 (37.9)
Positive	87 (62.1)
pSTAT3 expression	
Negative	73 (52.1)
Positive	67 (47.9)
VEGF-C expression	
Negative	82 (58.6)
Positive	58 (41.4)
VEGF-D expression	
Negative	106 (75.7)
Positive	34 (24.3)
Ki-67 expression	
Negative	53 (37.9)
Positive	87 (62.1)
ER expression	
Negative	76 (54.3)
Positive	64 (45.7)
PR expression	
Negative	68 (48.6)
Positive	72 (51.4)
HER2 expression	
Negative	99 (70.7)
Positive	41 (29.3)

SD=standard deviation; LN=lymph node; WHO=World Health Organization; STAT3=signal transducer and activator of transcription 3; pSTAT3=phosphorylated STAT3; VEGF=vascular endothelial growth factor; ER=estrogen receptor; PR=progesterone receptor; HER2=human epidermal growth factor receptor 2. *According to the 7th AJCC (American Joint Committee On Cancer) TNM classification for breast cancer.

Univariate and multivariate survival analysis

With univariate analysis (Kaplan-Meier method), we found five clinicopathological variables and biomarkers' expression had significant associations with the OS of breast cancer patients after curative resection. They are as follows: N stage, TNM classification, STAT3 expression, pSTAT3 expression, Ki-67 expression, ER expression, and PR expression. All above seven variables were included in a multivariate Cox proportional hazards model (forward stepwise procedure) to adjust for the effects of covariates. In that model, only number of lymph node metastasis was identified as the independent predictors with the OS of breast cancer patients postoperatively (HR, 3.220; 95% CI, 2.257-4.594; $p < 0.001$) (Table 2, Figure 5).

Correlation analysis between STAT3 expression and other variables in 140 breast cancer patients

A univariate analysis (chi-square test) identified nine variables that had statistically significant associations with STAT3 expression from breast cancer: age at surgery, T stage, TNM classification, N stage, ER expression, pSTAT3 expression, VEGF-C expression, VEGF-D expression, and Ki-67 expression (Table 3). All the nine variables were included in a multinomial logistic regression to adjust for the effects of covariates to lymph node metastasis from breast cancer. In that model, AIC and BIC values were executed by using the multinomial logistic regression according to STAT3 expression of patients when the follow-up was over. We demonstrated that both AIC value and BIC value of N stage were the smallest (AIC value, 176.930; BIC value, 256.355) in the aforementioned nine relative factors of lymph node metastasis from breast cancer, which indicated that N stage was the most important predictor of STAT3 expression in breast cancer (Table 4, Figure 6).

DISCUSSION

Lymph node metastasis from breast cancer is a complex pathophysiological process of cancer progress, which is mainly associated with the biological habits of cancer cells. Usually, biological factors may directly affect the end-results of the malignant cells to selectively invade lymph nodes rather than visceral organs, just as certain tumor types metastasize to certain organs and not others [20]. Exact recognition of the lymph node metastatic status is considered as the important clue for contribution to not only choice the appropriate surgical procedure of lymph node dissection, but improvement the prognostic predication. To date, SLNB has become the standard of care as a staging procedure for breast cancer patients with clinically node-negative disease [5]. The current guide-

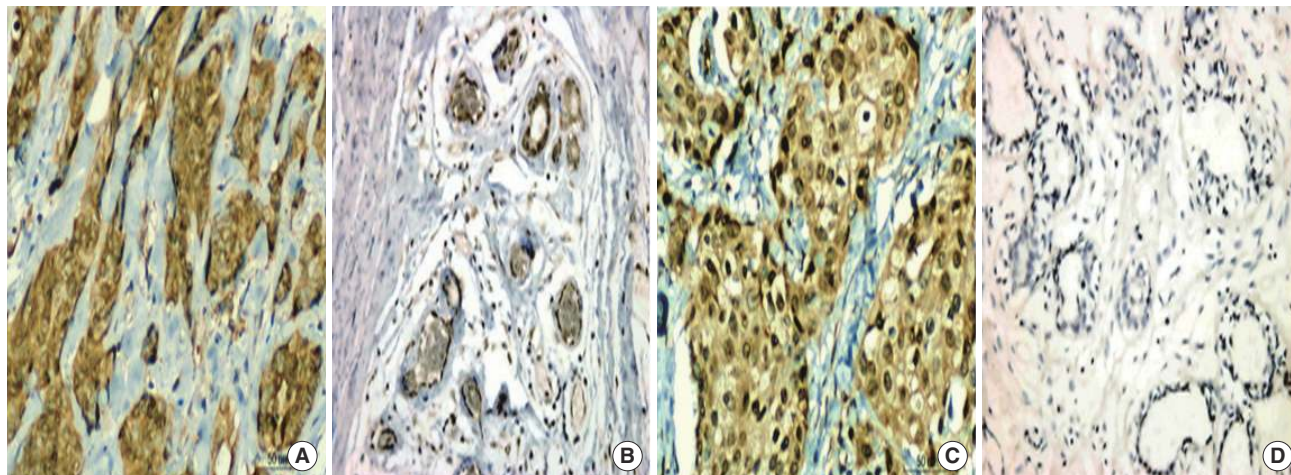


Figure 1. (A) Signal transducer and activator of transcription 3 (STAT3) expression in breast cancer tissue. (B) STAT3 expression in adjacent nontumor tissue. (C) pSTAT3 expression in breast cancer tissue. (D) pSTAT3 expression in adjacent nontumor tissue (Immunohistochemical staining, $\times 400$).

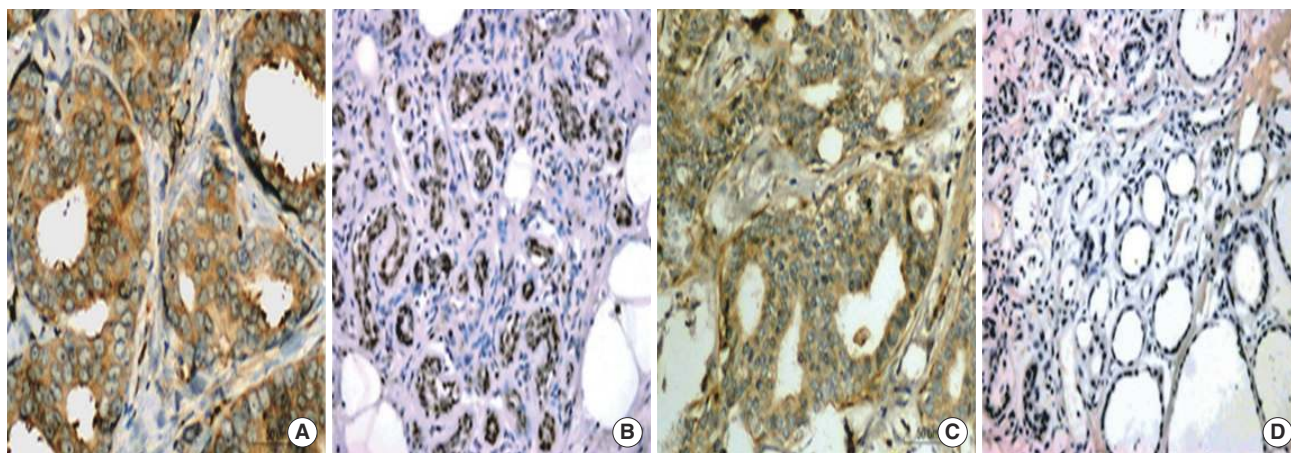


Figure 2. (A) Vascular endothelial growth factor (VEGF)-C expression in breast cancer tissue. (B) VEGF-C expression in adjacent nontumor tissue. (C) VEGF-D expression in breast cancer tissue. (D) VEGF-D expression in adjacent nontumor tissue (Immunohistochemical staining, $\times 400$).

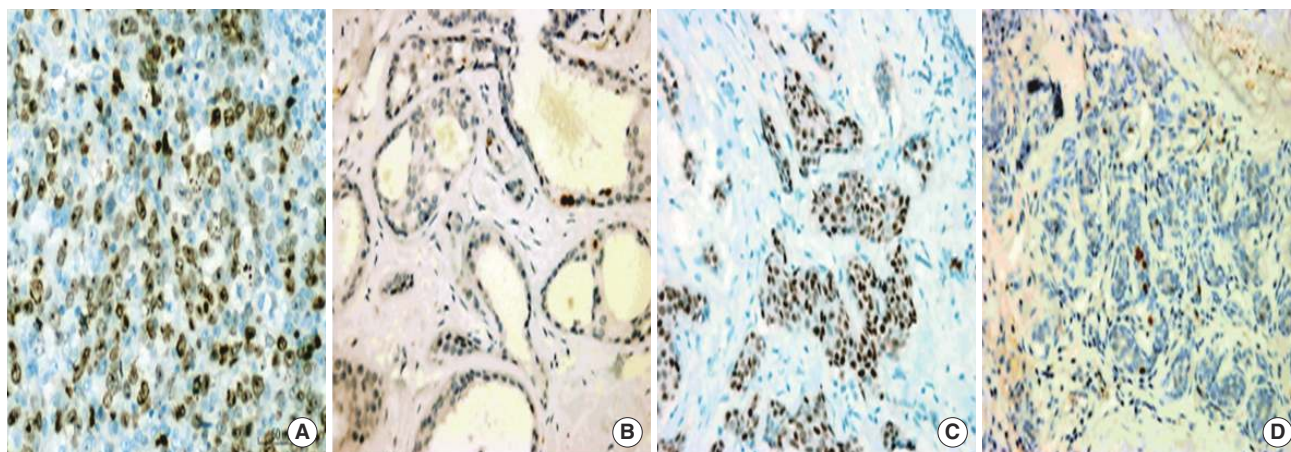


Figure 3. (A) Ki-67 expression in breast cancer tissue. (B) Ki-67 expression in adjacent nontumor tissue. (C) Estrogen receptor (ER) expression in breast cancer tissue. (D) ER expression in adjacent nontumor tissue (Immunohistochemical staining, $\times 400$).

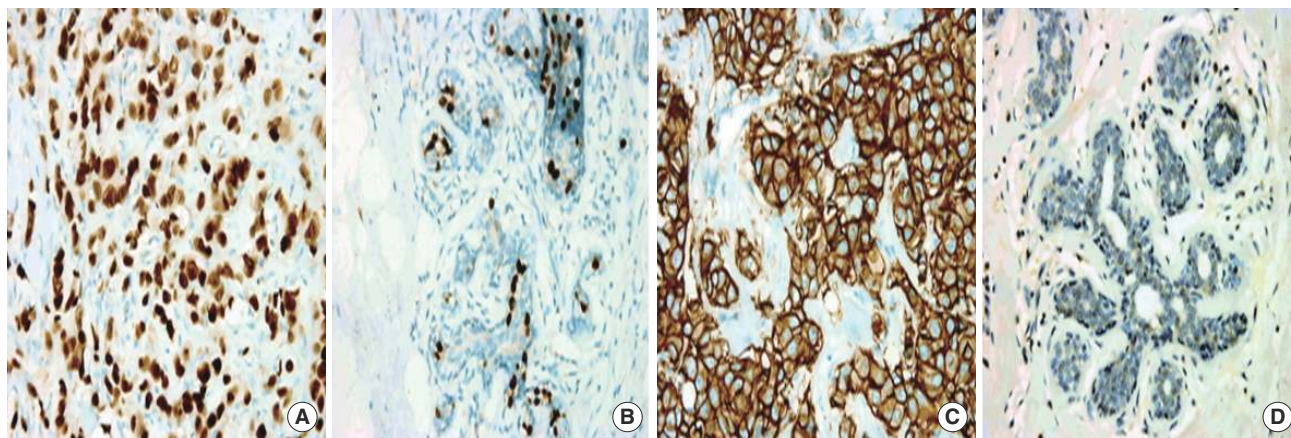


Figure 4. (A) Progesterone receptor (PR) expression in breast cancer tissue. (B) PR expression in adjacent nontumor tissue. (C) Human epidermal growth factor receptor 2 (HER2) expression in breast cancer tissue. (D) HER2 expression in adjacent nontumor tissue (Immunohistochemical staining, $\times 400$).

lines recommend completion ALND when lymph node involvement is identified by SLNB [4], although the need for completion ALND has been questioned [2]. Kim et al. [21] reported that approximately 50% of patients had additional nonsentinel axillary nodal metastases, which indicated that ALND might improve the prognosis of breast cancer patients. Furthermore, Park et al. [6] found that axillary recurrences were more frequent in a group of 287 patients who underwent SLNB alone compared with those who underwent completion ALND (2.0% vs. 0.4%, $p = 0.004$). Therefore, we have to believe that the exact diagnosis of nodal metastasis from breast cancer is a crucial factor for application the correctly surgical procedure and decrease the recurrence rate.

However, there is not an optimal method for exact detection the lymph node metastatic status before surgery. The insidiously nodal metastasis and micrometastasis of lymph nodes are so difficult to be detected with the general examinations. Biomarkers are high hopes for accurate prediction the lymph node metastasis from primary tumor. In theory, cancer cell metastasis should be considered to involve the cell proliferation, antiapoptosis, invasion enhancement, and lymphangiogenesis. According to these biological characteristics, STAT3, a cub of molecular regulator in various malignancies, is competent to evaluate the lymph node metastasis, rather than other conventional lymphangiogenic factors [10-13].

In many solid tumors, overexpression of human epidermal growth factor receptors (e.g., HER1/EGFR and HER2) correlates with poor prognosis. As a poor prognostic factor, EGFR overexpression is frequently shown in the breast cancer cells and tissues [22]. The key one of the down-stream molecules of EGFR which is identified to the oncogenesis in various organs is STAT3 [23]. Actually, STAT3 is constitutively activated by aberrant upstream tyrosine kinase activities in a broad

spectrum of cancer cell lines and human tumors. STAT3 is an oncogenic protein that is constitutively activated in many human cancers, including 30% to 60% of primary breast cancers [22]. The biological significance of STAT3 activation in breast cancer is not fully understood. STAT3 activation may modulate tumor invasiveness of breast cancer by regulating TIMP1 expression. Pytowski et al. [24] generated an *in vitro* system in which active STAT3 levels can be tightly controlled in breast cancer cells by using MCF-7 cells transfected with tetracycline-off constitutively active STAT3. Using immunohistochemistry and 142 primary breast tumors, they also found a significant association between the expression of the phosphorylated/active form of STAT3 (pSTAT3) and that of tissue inhibitor of matrix metalloproteinases-1 (TIMP-1) [24]. At present study, we demonstrated there were significantly statistical differences in the positive expression incidences of STAT3, and pSTAT3 between breast cancer and adjacent nontumor tissue specimens. In addition, we also found that both STAT3 expression and pSTAT3 expression were significantly associated with the OS of breast cancer patients after curative resection. Therefore, we did consider that STAT3 was a pivotal biomarker for predication the prognosis of breast cancer. As other investigators reported, STAT3 was significantly associated with the bad prognosis in various human malignancies [18, 25-27].

Lymph node metastasis was well known as the most important predictor of prognosis of breast cancer after surgery. Recently, several investigations demonstrated that STAT3 was involved in a series of molecular events of lymph node metastasis in different human cancers [11-15]. From the results of this study, we found that lymph node metastasis was the only independent predictor of the OS of breast cancer patients after surgery. Although we demonstrated that nine variables

Table 2. Univariate and multivariate survival analysis of 140 breast cancer patients after curative surgery

Variable	Mean OS (mo)	Chi-square value	Univariate <i>p</i> -value	HR value (95% CI)	Multivariate <i>p</i> -value
Age at surgery (yr)					
<50	105.17 ± 4.30	0.567	0.451	-	-
≥50	79.93 ± 3.60				
T stage*					
T1	108.19 ± 4.75	5.426	0.066	-	-
T2	81.57 ± 2.49				
T3	59.06 ± 5.66				
N stage*					
N0	115.75 ± 3.50	71.070	<0.001	3.220 (2.257-4.594)	<0.001
N1	79.87 ± 3.61				
N2	57.50 ± 4.73				
N3	38.74 ± 4.97				
TNM* classification					
I	114.00 ± 5.66	2.401	0.301	-	0.784
II	99.17 ± 5.68				
III	100.79 ± 4.47				
WHO classification					
I	114.00 ± 5.66	2.401	0.301	-	-
II	99.17 ± 5.68				
III	100.79 ± 4.47				
STAT3 expression					
Negative	110.58 ± 4.40	4.765	0.029	-	0.566
Positive	78.26 ± 2.60				
pSTAT3 expression					
Negative	111.95 ± 3.95	12.268	<0.001	-	0.916
Positive	74.57 ± 3.25				
VEGF-C expression					
Negative	105.50 ± 4.41	1.501	0.221	-	-
Positive	91.70 ± 4.32				
VEGF-D expression					
Negative	105.15 ± 4.06	2.835	0.092	-	-
Positive	73.07 ± 3.91				
Ki-67 expression					
Negative	111.44 ± 4.76	5.936	0.015	-	0.601
Positive	84.7 ± 3.43				
ER expression					
Negative	84.21 ± 3.57	5.885	0.015	-	0.591
Positive	109.91 ± 4.76				
PR expression					
Negative	89.02 ± 5.76	4.831	0.028	-	0.861
Positive	110.47 ± 3.76				
HER2 expression					
Negative	102.10 ± 4.66	0.038	0.845	-	-
Positive	95.33 ± 4.40				

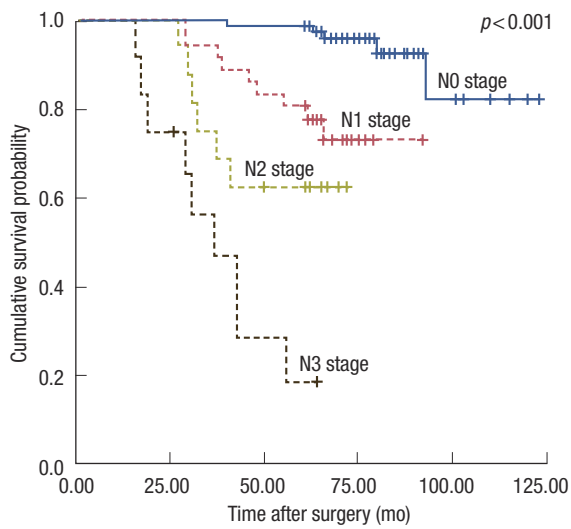
OS=overall survival; HR=hazard ratio; CI=confidence interval; WHO=World Health Organization; STAT3=signal transducer and activator of transcription 3; pSTAT3=phosphorylated STAT3; VEGF=vascular endothelial growth factor; ER=estrogen receptor; PR=progesterone receptor; HER2=human epidermal growth factor receptor 2.

*According to the 7th AJCC (American Joint Committee On Cancer) TNM classification for breast cancer.

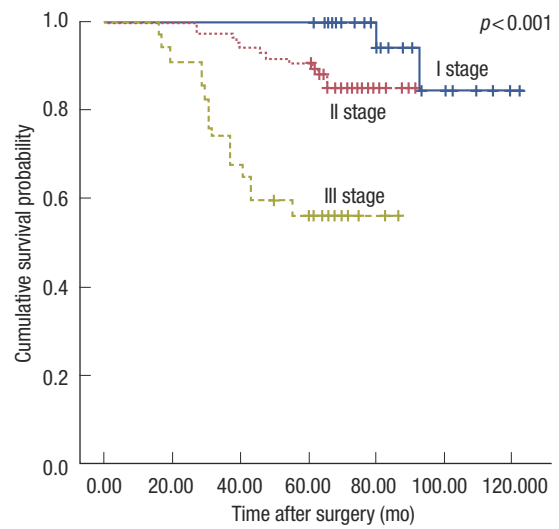
(including VEGF-C expression, VEGF-D expression, and Ki-67 expression) that had statistically significant associations with lymph node metastasis from breast cancer, the STAT3 expression was the most important predictor of lymph node metastasis from breast cancer by using the multinomial logis-

tical regression model analysis rather than other variables.

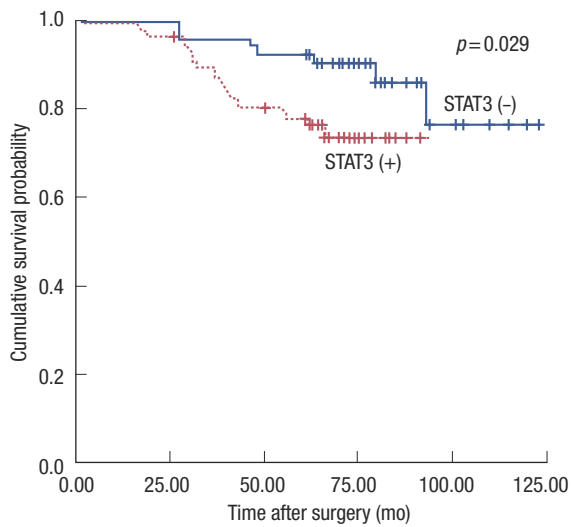
Notwithstanding a small scale study was performed, we concluded that STAT3 might improve the cancer cell lymphatic metastasis, which induced the bad prognosis of breast cancer patients. The anti-STAT3 monoclonal antibody might change



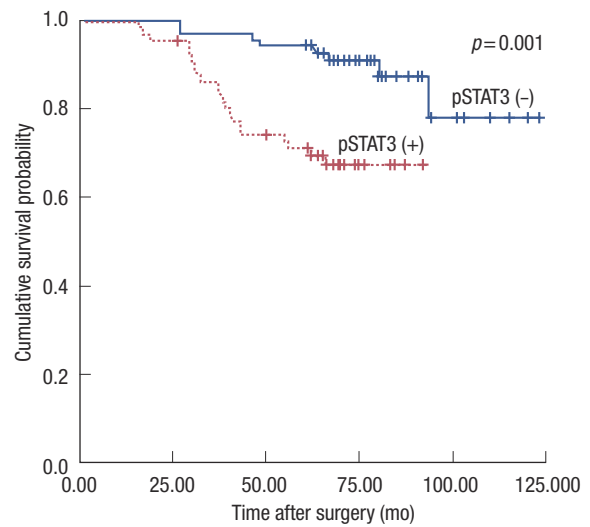
A



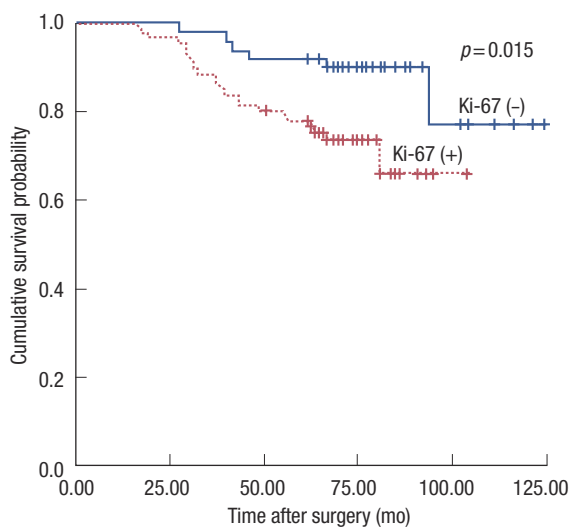
B



C



D



E

Figure 5. Survival curve for 140 breast cancer patients after curative surgery according to stage subgroup. (A) N stage (the AJCC (American Joint Committee On Cancer)) TNM classification of breast cancer. (B) TNM stage (the International Union against Cancer TNM classification of breast cancer). (C) Signal transducer and activator of transcription 3 (STAT3) expression. (D) Phosphorylated STAT3 (pSTAT3) expression. (E) Ki-67 expression.

Table 3. Univariate correlation analysis between STAT3 expression and other variables of 140 breast cancer patients

Variable	STAT3 immunohistochemical scores				Chi-square value	p-value
	0 score	1 score	2 scores	3 scores		
Age at surgery (yr)					18.968	<0.001
<50	36	20	18	14		
≥50	10	4	26	12		
T stage*					57.779	<0.001
T1	17	16	7	4		
T2	29	8	33	10		
T3	0	0	4	12		
N stage*					58.317	<0.001
N0	34	20	20	2		
N1	12	4	12	8		
N2	0	0	8	8		
N3	0	0	4	8		
TNM classification*					78.752	<0.001
I	15	14	1	0		
II	31	10	29	8		
III	0	0	14	18		
WHO classification					9.871	0.130
I	4	2	4	0		
II	19	8	11	4		
III	23	14	29	22		
pSTAT3 expression					89.164	<0.001
Negative	46	17	10	0		
Positive	0	7	34	26		
VEGF-C expression					26.677	<0.001
Negative	40	14	20	8		
Positive	6	10	24	18		
VEGF-D expression					25.014	<0.001
Negative	46	18	28	14		
Positive	0	6	16	12		
Ki-67 expression					17.256	0.001
Negative	27	11	10	5		
Positive	19	13	34	21		
ER expression					12.631	0.006
Negative	21	10	23	22		
Positive	25	14	21	4		
PR expression					7.367	0.061
Negative	23	6	23	16		
Positive	23	18	21	10		
HER2 expression					1.123	0.772
Negative	34	15	32	18		
Positive	12	9	12	8		

STAT3=signal transducer and activator of transcription 3; WHO=World Health Organization; pSTAT3=phosphorylated STAT3; VEGF=vascular endothelial growth factor; ER=estrogen receptor; PR=progesterone receptor; HER2=human epidermal growth factor receptor 2.

*According to the 7th AJCC (American Joint Committee On Cancer) TNM classification for breast cancer.

treatment status in the nodal metastasis from breast cancer, which is worthy of hoping for further improvement the prognosis of patients.

Table 4. Multivariate correlation analysis between STAT3 expression and other variables of 140 breast cancer patients

Variable	AIC value	BIC value	-2log likelihood value
Age at surgery	205.313	284.738	151.313
T stage*	182.652	262.077	128.652
N stage*	176.930	256.355	122.930
TNM classification*	184.234	263.659	130.234
pSTAT3 expression	199.218	278.642	145.218
VEGF-C expression	180.532	259.956	126.532
VEGF-D expression	186.057	265.481	132.057
Ki-67 expression	184.126	263.550	130.126
ER expression	191.351	270.776	137.351

STAT3=signal transducer and activator of transcription 3; AIC=Akaike information criterion; BIC=Bayesian information criterion; pSTAT3=phosphorylated STAT3; VEGF=vascular endothelial growth factor; ER=estrogen receptor.

*According to the 7th AJCC (American Joint Committee On Cancer) TNM classification for breast cancer.

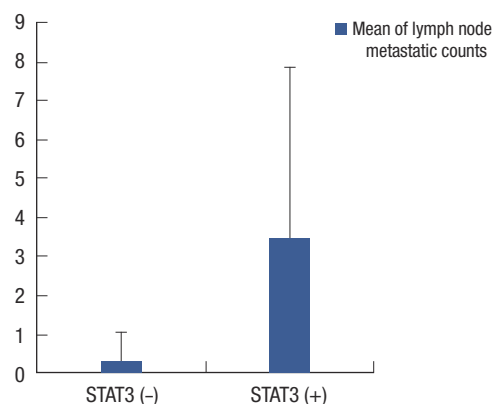


Figure 6. Comparison mean values of lymph node metastatic counts of 140 breast cancer patients according to signal transducer and activator of transcription 3 (STAT3) expression.

CONFLICT OF INTEREST

The authors declare that they have no competing interests.

REFERENCES

- Cowin P, Welch DR. Breast cancer progression: controversies and consensus in the molecular mechanisms of metastasis and EMT. *J Mammary Gland Biol Neoplasia* 2007;12:99-102.
- Naik AM, Fey J, Gemignani M, Heerdt A, Montgomery L, Petrek J, et al. The risk of axillary relapse after sentinel lymph node biopsy for breast cancer is comparable with that of axillary lymph node dissection: a follow-up study of 4008 procedures. *Ann Surg* 2004;240:462-8.
- Sola M, Alberro JA, Fraile M, Santesteban P, Ramos M, Fabregas R, et al. Complete axillary lymph node dissection versus clinical follow-up in breast cancer patients with sentinel node micrometastasis: final results from the multicenter clinical trial AATRM 048/13/2000. *Ann Surg Oncol* 2013;20:120-7.

4. Sanuki N, Takeda A, Amemiya A, Ofuchi T, Ono M, Ogata H, et al. Outcomes of clinically node-negative breast cancer without axillary dissection: can preserved axilla be safely treated with radiation after a positive sentinel node biopsy? *Clin Breast Cancer* 2013;13:69-76.
5. Noguchi M, Morioka E, Ohno Y, Nakano Y, Kosaka T. The changing role of axillary lymph node dissection for breast cancer. *Breast Cancer* 2013;20:41-6.
6. Park J, Fey JV, Naik AM, Borgen PI, Van Zee KJ, Cody HS 3rd. A declining rate of completion axillary dissection in sentinel lymph node-positive breast cancer patients is associated with the use of a multivariate nomogram. *Ann Surg* 2007; 245:462-468.
7. Glechner A, Wöckel A, Gartlehner G, Thaler K, Strobelberger M, Griebler U, et al. Sentinel lymph node dissection only versus complete axillary lymph node dissection in early invasive breast cancer: a systematic review and meta-analysis. *Eur J Cancer*. Epub 2012 Oct 17. DOI: <http://dx.doi.org/10.1016/j.ejca.2012.09.010>.
8. Mandriota SJ, Jussila L, Jeltsch M, Compagni A, Baetens D, Prevo R, et al. Vascular endothelial growth factor-C-mediated lymphangiogenesis promotes tumour metastasis. *EMBO J* 2001;20:672-82.
9. Van Trappen PO, Steele D, Lowe DG, Baithun S, Beasley N, Thiele W, et al. Expression of vascular endothelial growth factor (VEGF)-C and VEGF-D, and their receptor VEGFR-3, during different stages of cervical carcinogenesis. *J Pathol* 2003;201:544-54.
10. Couto JP, Daly L, Almeida A, Knauf JA, Fagin JA, Sobrinho-Simões M, et al. STAT3 negatively regulates thyroid tumorigenesis. *Proc Natl Acad Sci U S A* 2012;109:E2361-70.
11. Amin HM, McDonnell TJ, Ma Y, Lin Q, Fujio Y, Kunisada K, et al. Selective inhibition of STAT3 induces apoptosis and G(1) cell cycle arrest in ALK-positive anaplastic large cell lymphoma. *Oncogene* 2004;23: 5426-34.
12. Zhang F, Li C, Halfter H, Liu J. Delineating an oncostatin M-activated STAT3 signaling pathway that coordinates the expression of genes involved in cell cycle regulation and extracellular matrix deposition of MCF-7 cells. *Oncogene* 2003;22:894-905.
13. Niu G, Wright KL, Huang M, Song L, Haura E, Turkson J, et al. Constitutive Stat3 activity up-regulates VEGF expression and tumor angiogenesis. *Oncogene* 2002;21:2000-8.
14. Messina JL, Yu H, Riker AI, Munster PN, Jove RL, Daud AI. Activated stat-3 in melanoma. *Cancer Control* 2008;15:196-201.
15. Zhao X, Sun X, Li XL. Expression and clinical significance of STAT3, P-STAT3, and VEGF-C in small cell lung cancer. *Asian Pac J Cancer Prev* 2012;13:2873-7.
16. Cai QW, Li J, Li XQ, Wang JQ, Huang Y. Expression of STAT3, MMP-1 and TIMP-1 in gastric cancer and correlation with pathological features. *Mol Med Report* 2012;5:1438-42.
17. Jiang R, Jin Z, Liu Z, Sun L, Wang L, Li K. Correlation of activated STAT3 expression with clinicopathologic features in lung adenocarcinoma and squamous cell carcinoma. *Mol Diagn Ther* 2011;15:347-52.
18. Deng JY, Sun D, Liu XY, Pan Y, Liang H. STAT-3 correlates with lymph node metastasis and cell survival in gastric cancer. *World J Gastroenterol* 2010;16:5380-7.
19. Takemoto S, Ushijima K, Kawano K, Yamaguchi T, Terada A, Fujiyoshi N, et al. Expression of activated signal transducer and activator of transcription-3 predicts poor prognosis in cervical squamous-cell carcinoma. *Br J Cancer* 2009;101:967-72.
20. Fidler IJ. The pathogenesis of cancer metastasis: the 'seed and soil' hypothesis revisited. *Nat Rev Cancer* 2003;3:453-8.
21. Kim T, Giuliano AE, Lyman GH. Lymphatic mapping and sentinel lymph node biopsy in early-stage breast carcinoma: a metaanalysis. *Cancer* 2006; 106:4-16.
22. Hatake K, Tokudome N, Ito Y. Next generation molecular targeted agents for breast cancer: focus on EGFR and VEGFR pathways. *Breast Cancer* 2007;14:132-49.
23. Yue P, Zhang X, Paladino D, Sengupta B, Ahmad S, Holloway RW, et al. Hyperactive EGF receptor, Jaks and Stat3 signaling promote enhanced colony-forming ability, motility and migration of cisplatin-resistant ovarian cancer cells. *Oncogene* 2012;31:2309-22.
24. Pytowski B, Goldman J, Persaud K, Wu Y, Witte L, Hicklin DJ, et al. Complete and specific inhibition of adult lymphatic regeneration by a novel VEGFR-3 neutralizing antibody. *J Natl Cancer Inst* 2005;97:14-21.
25. Schoppmann SF, Jesch B, Friedrich J, Jomrich G, Maroske F, Birner P. Phosphorylation of signal transducer and activator of transcription 3 (STAT3) correlates with Her-2 status, carbonic anhydrase 9 expression and prognosis in esophageal cancer. *Clin Exp Metastasis* 2012;29:615-24.
26. Kim HS, Park YH, Lee J, Ahn JS, Kim J, Shim YM, et al. Clinical impact of phosphorylated signal transducer and activator of transcription 3, epidermal growth factor receptor, p53, and vascular endothelial growth factor receptor 1 expression in resected adenocarcinoma of lung by using tissue microarray. *Cancer* 2010;116:676-85.
27. Shah NG, Trivedi TI, Tankshali RA, Goswami JV, Jetly DH, Shukla SN, et al. Prognostic significance of molecular markers in oral squamous cell carcinoma: a multivariate analysis. *Head Neck* 2009;31:1544-56.