

Poor outcome of patients with pulmonary adenocarcinoma showing decreased E-cadherin combined with increased S100A4 expression

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Abstract. A loss or reduced expression of E-cadherin, the main cell-to-cell adhesion molecule, correlates with distant metastasis in various cancers. Recent studies have reported a close correlation between the expression of E-cadherin and that of S100A4, calcium-binding protein. In this study, we investigated the expression of E-cadherin and S100A4 status in relation to the clinicopathological parameters of pulmonary adenocarcinoma. We finely and quantitatively examined the expression of E-cadherin and S100A4 using real-time polymerase chain reaction (PCR) in a total of 92 pulmonary adenocarcinomas obtained by surgical resection. All of the pulmonary adenocarcinomas showed significant expression of E-cadherin and S100A4. Real-time PCR showed lower E-cadherin expression in 21 adenocarcinomas, while 71 adenocarcinomas expressed a higher expression of E-cadherin. Of 21 adenocarcinomas with lower-expressing E-cadherin, 12 showed a higher expression of S100A4. These 12 cases significantly showed a poorer prognosis than others ($p=0.047$, Kaplan-Meier, log-rank test) and significantly showed more frequent venous involvement than others ($p=0.042$, χ^2 test). These results suggested that reduced E-cadherin expression combined with higher S100A4 expression is related to a poor prognosis through hematogenous metastasis in pulmonary adenocarcinoma.

Introduction

E-cadherin is the main cell-to-cell adhesion molecule that participates in homophilic, calcium-dependent interactions to

form the epithelial adherence junction (1,2). A loss or the reduced expression of E-cadherin is correlated with distant metastasis in various advanced cancers, including lung cancer (3,4). Thus, E-cadherin is assumed to act as a main invasion-suppressor molecule (5,6). Recent studies have reported the close correlations between S100A4 and E-cadherin expression in stomach and gallbladder cancers (7,8).

The S100 family of calcium-binding proteins are involved in a variety of physiological functions, such as cellular proliferation, adhesion and motility (9). S100A4, a small member of the S100 family, formerly known as p9Ka or mts1, has been characterized as a 'metastasis-related gene' (10,11). The elevated level of S100A4 protein correlated with the metastatic potential of breast neoplasm in rodent models (12-14). The induced expression of S100A4 increased the metastatic potentials in several rodent models of mammary carcinogenesis (15,16). A high incidence of pulmonary metastasis of mammary carcinomas has been observed in S100A4 transgenic mice (15,17). S100A4 overexpression has been associated with a poor prognosis in a variety of human cancers, such as stomach, colon, breast and gallbladder (7,18-20).

In this study, we finely and quantitatively examined the gene expression of E-cadherin and S100A4 by real-time PCR in a total of 92 pulmonary adenocarcinomas obtained by complete surgical resection. We also evaluated the clinical prognosis with long-term observation periods of 5-15 years in the 92 cases with pulmonary adenocarcinoma. We here discuss the relationship between the expression of E-cadherin and S100A4 and clinicopathological features in the 92 cases with pulmonary adenocarcinoma.

Patients and methods

Patients. Ninety-two pulmonary adenocarcinoma specimens were obtained at surgical resection from October 1989 to November 1999 with the patients' informed consents. All cases underwent complete resection (lobectomy or pneumonectomy and dissection of mediastinal lymph nodes) of pulmonary adenocarcinoma. Tissues were frozen and stored at -80°C until analyses. Surgical specimens were also processed for routine histopathological analysis. The pathological stages

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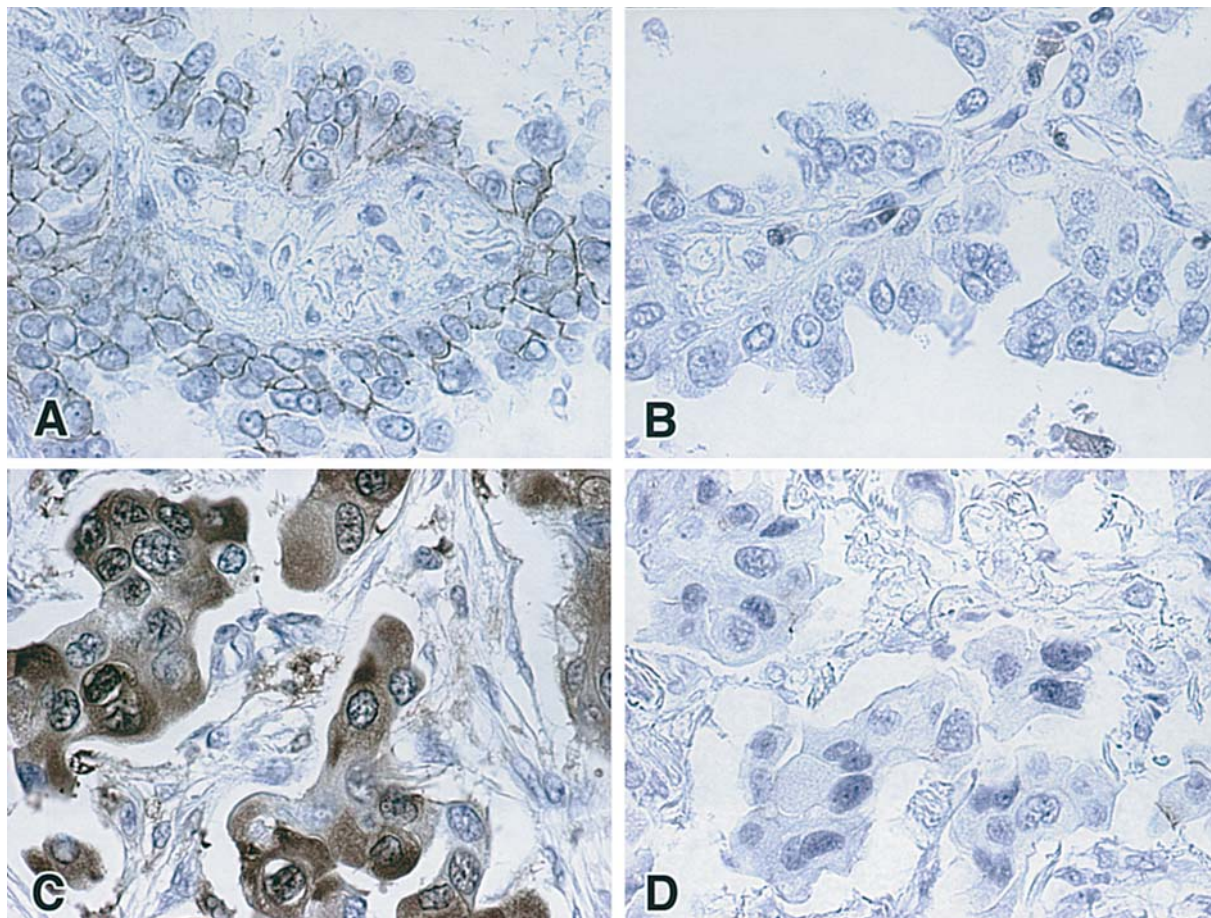


Figure 1. Protein localization of E-cadherin and S100A4. The cancerous nests with well-differentiated papillary features showed E-cadherin expression in the plasma membrane (A) but no apparent S100A4 expression (B). The poorly-differentiated cancerous nests revealed strong S100A4 expression in the cytoplasm and nucleus (C) but no apparent E-cadherin expression (D).

were estimated according to the World Health Organization (WHO) classification (21).

The patients consisted of 48 men and 44 women with a mean age of 63.0 ± 9.47 years (mean \pm standard deviation). The pathological stages of the pulmonary adenocarcinoma were as follows: stage I, 52 (56.5%) patients; stage II, 10 (10.9%) patients; stage III, 30 (32.6%) patients. Tumor status was T1 in 40 patients, T2 in 42, T3 in 7 and T4 in 3. Fifty-four patients had no lymph node metastasis (N0), whereas 38 had regional or mediastinal lymph node metastasis (11 had N1, 26 had N2 and 1 had N3 disease).

Quantitative evaluation of E-cadherin and S100A4. Total cellular RNA was prepared from the frozen specimens by standard acid guanidine isothiocyanate-phenol-chloroform extraction procedures. After heat-denaturation of total RNA specimens (1 μ g), reverse transcription was performed [10 mM DTT (Invitrogen Corp., Carlsbad, CA, USA), 0.2 mM dNTPs (Toyobo Co., Osaka, Japan) 100 pmol of Primer, Random PD (N6), (Roche Diagnostics Co., Indianapolis, IN, USA) and 200 units of SuperscriptTM II RNase H⁻ Reverse Transcriptase (Invitrogen) at 42°C, 60 min] (22,23). Real-time quantitative PCR for E-cadherin and S100A4 mRNA was performed according to the manufacturer's recommendation and our previous reports (24-26). The primers of E-cadherin

(Hs00170423) and S100A4 (Hs00243201) were purchased from TaqMan Gene Expression Assays. We used TaqMan Universal PCR Master Mix (PE Applied Biosystems, Foster City, CA) for the real-time PCR. For the internal controls, β -actin-probe-primer mixture for β -actin mRNA was used (human ACTB, 4310881E, PE Applied Biosystems). Real-time PCR assays were run on an ABI PRISM 7000 Sequence Detection System with the following protocol: initial denaturation for 2 min at 50°C and 10 min at 95°C, and amplification for 50 cycles of 15 sec at 95°C and 60 sec at 60°C. All of the samples were run in duplicate.

After determining the threshold cycle (C_t), which was defined as the PCR cycle number where the fluorescent intensity exceeded the threshold, the amount of E-cadherin and S100A4 mRNA in the sample specimen was calculated from the C_t of the sample and the RNA standard curve. Then, the relationship between the C_t and initial standard copy number was expressed as a logarithmic formula. The obtained copy of E-cadherin and S100A4 was then standardized with the β -actin mRNA quantity as the endogenous control using the following equation: Result = Log (E-cadherin or S100A4 RNA copy number in surgical samples)/(β -actin RNA copy number in surgical samples) $\times (6.1 \times 10^9)$.

Gene levels are estimated with standard expression in the control cell lines (defined as 1.00). We used G361 (human

skin malignant melanoma cell line) as standard for E-cadherin. The human skin malignant melanoma cell line, A375, was used as the standard for the evaluation of the S100A4.

Immunohistochemistry. The formalin-fixed and paraffin-embedded tissue samples were processed for immunohistochemistry. The sections were incubated with mouse monoclonal anti-E-cadherin antibodies (Takara, Ootsu Japan) and rabbit polyclonal anti-S100A4 antibodies (Dako, Glostrup, Denmark) at room temperature for 60 min. The E-cadherin immune complex was amplified with the HRP-labeled anti-mouse immunoglobulin (secondary antibody) after antigen retrieval (autoclaved in 1 mM EDTA for 10 min at 121°C) and the blocking of endogenous peroxidase activity with 0.3% hydrogen peroxide for 5 min. The S100A4 immune complex was amplified with the HRP-labeled anti-rabbit immunoglobulin (secondary antibody) after antigen retrieval (microwaved in 0.01 M citrate buffer for 20 min at 95°C) and blocking of endogenous peroxidase activity with 0.3% hydrogen peroxide for 5 min. The amplified immune products were visualized using a 3,3'-diaminobenzidine tetrahydrochloride (DAB) reaction.

Statistical analysis. The χ^2 test was applied for comparisons between group frequencies. Differences in survival between subgroups of patients were compared using the log-rank test, and survival curves were plotted according to the method of Kaplan-Meier. Data are shown as means \pm standard deviation.

Results

Expression of S100A4 and E-cadherin in pulmonary adenocarcinomas. All of the pulmonary adenocarcinomas showed significant expression of E-cadherin and S100A4 by real-time PCR. The expression levels of E-cadherin in neoplastic tissue varied from 0.600 to 15.369 (5.549 ± 3.243). In the 21 extraneoplastic tissues, the expression levels of E-cadherin mRNA varied from 0.822 to 4.559 (2.069 ± 0.891). The expression levels of S100A4 in neoplastic tissue varied from 0.027 to 0.730 (0.155 ± 0.117). In the 21 extraneoplastic tissues, the expression levels of S100A4 varied from 0.132 to 0.893 (0.403 ± 0.187). No inverse relation was apparent between E-cadherin and S100A4 expression.

We confirmed the cellular localization of E-cadherin and S100A4 protein by immunohistochemistry. E-cadherin expression was observed in the plasma membrane of tumor cells of adenocarcinoma. The cancerous cells with well-differentiated papillary features showed E-cadherin expression (Fig. 1A) but no S100A4 expression (Fig. 1B). The poorly differentiated cancerous cells revealed strong S100A4 expression (Fig. 1C) but no E-cadherin expression (Fig. 1D).

Clinicopathological implications of S100A4 and E-cadherin expression. The post-surgical observation period lasted from 0.2 to 149.6 months (54.3 ± 41.6 month). Twenty-one cases were categorized as the lower E-cadherin expressing (<3.000) group of 92 pulmonary adenocarcinoma cases.

There was no significant difference in prognosis between the lower E-cadherin expressing group and the higher E-cadherin expressing group ($p=0.114$, Kaplan-Meier, log-rank

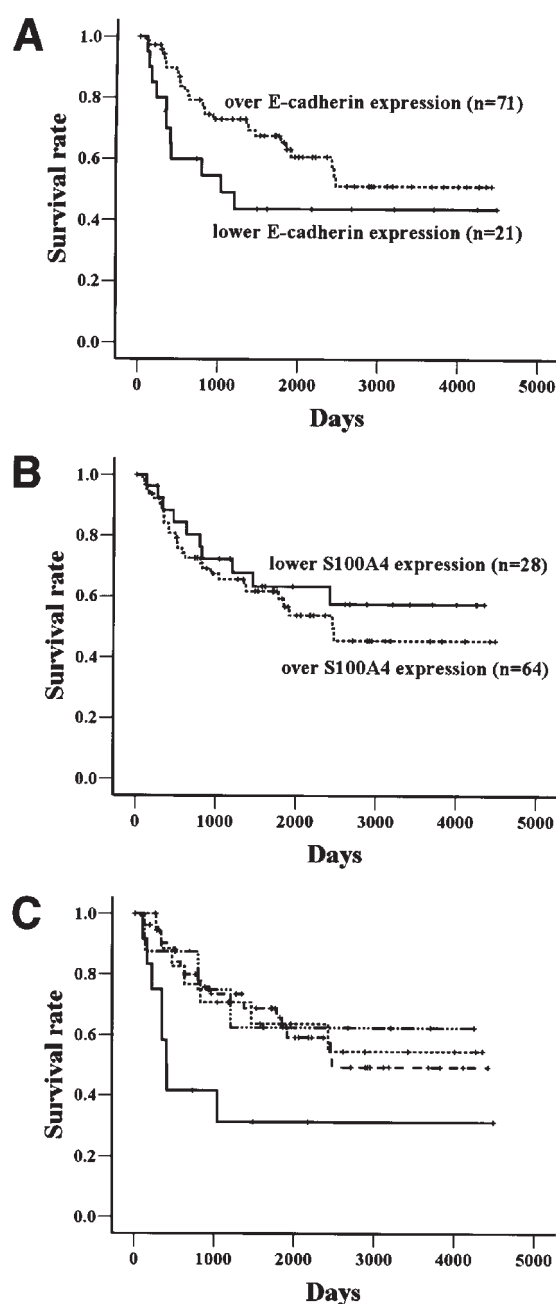


Figure 2. Survival curves of the cases with pulmonary adenocarcinoma. (A) There was no significant difference of prognosis between the 21 cases with lower E-cadherin expression and the other 71 cases ($p=0.114$, Kaplan-Meier, log-rank test). (B) There was also no significant difference of prognosis between the 28 cases with lower S100A4 expression and the other 64 cases ($p=0.453$, Kaplan-Meier, log-rank test). (C) Survival curves of four groups subdivided according to the expression status of E-cadherin and S100A4. The cases with lower-expressing E-cadherin and overexpressing S100A4 tumors (continuous line, $n=12$) showed a significantly poorer prognosis than those with tumors of the other three groups; those with lower-expressing E-cadherin and lower-expressing S100A4 (dashed and dotted line, $n=9$), overexpressing E-cadherin and lower-expressing S100A4 (dotted line, $n=19$), and overexpressing E-cadherin and overexpressing S100A4 (dashed line, $n=52$) ($p=0.047$, Kaplan-Meier, log-rank test).

test) (Fig. 2A). The sixty-four pulmonary adenocarcinomas showed higher levels of S100A4 (>0.1). There was also no significant difference in prognosis between the lower S100A4 expressing group and the higher S100A4 expressing group ($p=0.453$, Kaplan-Meier, log-rank test) (Fig. 2B).

Table I. Gene expression of E-cadherin and S100A4, and venous involvements.

Gene expression	Venous involvements		p-value
	(+) (%)	(-) (%)	
E-cadherin/S100A4			0.042 ^a
Higher/higher	29 (31.5)	23 (25.0)	
Higher/lower	9 (9.8)	10 (10.9)	
Lower/higher	10 (10.9)	2 (2.2)	
Lower/lower	2 (2.2)	7 (7.6)	
Total	50 (54.3)	42 (45.7)	

^aSignificance estimated with χ^2 test.

Table II. Gene expression of E-cadherin and S100A4, and metastatic organs.

Metastatic organ	E-cadherin overexpression			S100A4 overexpression		
	(+)	(-)	p-value	(+)	(-)	p-value
Total						
(+)	30	7	0.464	29	8	0.132
(-)	41	14		35	20	
Lung						
(+)	10	1	0.247	9	2	0.347
(-)	61	20		55	26	
Brain						
(+)	13	3	0.669	12	4	0.603
(-)	58	18		52	24	
Bone						
(+)	11	4	0.698	12	3	0.337
(-)	60	17		52	25	
Liver						
(+)	1	0	0.584	0	1	0.128
(-)	70	21		64	27	
Adrenal gland						
(+)	0	1	0.064	1	0	0.506
(-)	71	20		63	28	

Twelve of the 21 adenocarcinomas with lower-expressing E-cadherin showed higher expression of S100A4. These lower E-cadherin and higher S100A4 expressing cases significantly showed a poorer prognosis ($p=0.047$, Kaplan-Meier, log-rank test) (Fig. 2C). These 12 cases also showed significantly enhanced venous involvement ($p=0.042$, χ^2 test, Table I).

Table III. Univariate analysis of the associations between E-cadherin gene expression and clinicopathological features in pulmonary adenocarcinoma.

Variable	E-cadherin overexpression		p-value
	(+) (%)	(-) (%)	
Stage			0.827
I	41 (44.6)	11 (12.0)	
II	8 (8.7)	2 (2.2)	
III	22 (23.9)	8 (8.7)	
T factor			0.745
T1	33 (35.9)	7 (7.6)	
T2	31 (33.7)	11 (12.0)	
T3	5 (5.4)	2 (2.2)	
T4	2 (2.2)	1 (1.1)	
N factor			0.503
N (+)	28 (30.4)	10 (10.9)	
N (-)	43 (46.7)	11 (12.0)	
M factor			0.464
M0	41 (44.6)	14 (15.2)	
M1	30 (32.6)	7 (7.6)	
Total	71 (74.0)	21 (26.0)	

Table IV. Univariate analysis of the associations between S100A4 gene expression and clinicopathological features in pulmonary adenocarcinoma.

Variable	S100A4 overexpression		p-value
	(+) (%)	(-) (%)	
Stage			0.342
I	33 (35.9)	19 (20.7)	
II	8 (8.7)	2 (2.2)	
III	23 (25.0)	7 (7.6)	
T factor			0.321
T1	29 (31.5)	11 (12.0)	
T2	26 (28.3)	16 (17.4)	
T3	6 (6.5)	1 (1.1)	
T4	3 (3.3)	0 (0)	
N factor			0.238
N (+)	29 (31.5)	9 (9.8)	
N (-)	35 (38.0)	19 (20.7)	
M factor			0.132
M0	29 (31.5)	8 (8.7)	
M1	35 (38.0)	20 (21.7)	
Total	64 (69.6)	28 (30.4)	

However, no significant correlation was noted between metastatic organs and gene expression patterns of E-cadherin or S100A4 (Table II). Univariate analysis showed no significant correlations between E-cadherin or S100A4 expression levels and clinicopathological features in pulmonary adenocarcinoma (Tables III and IV).

Discussion

We finely and quantitatively examined the gene expression of E-cadherin and S100A4 in 92 surgical cases with pulmonary adenocarcinoma by real-time PCR. Twelve lower E-cadherin expressing pulmonary adenocarcinomas showed a higher expression of S100A4 and a significantly poorer prognosis and enhanced venous involvement. These results suggested that reduced E-cadherin expression combined with higher S100A4 expression is related to a poorer prognosis through hematogenous metastasis in pulmonary adenocarcinoma.

E-cadherin is a factor for cell-cell adhesion and maintaining epithelial structure. The decrease or loss of E-cadherin contributed to the reduction of cell-to-cell adhesiveness, followed by the loss of cell polarity and the destruction of structures. It may induce cells to dissociate from their primary tumors and invade the surrounding tissue or metastasize to distant organs (3). The relationship of S100A4 expression with the down-regulation of E-cadherin was first reported in mouse mammary tumor cells (27). Subsequent studies further supported the evidence that there was an inverse correlation between the expression of E-cadherin and S100A4 in several human cancers; stomach, gallbladder and malignant melanoma (7,20,28). Kimura *et al* reported that S100A4 plays a role in progression and metastasis in non-small cell lung cancer by immunohistochemistry (4). Our results revealed no inverse correlation of S100A4 and E-cadherin expression in pulmonary adenocarcinoma at the gene levels but showed that reduced E-cadherin expression combined with increased S100A4 expression was linked to a poorer prognosis in pulmonary adenocarcinoma according to real time-PCR assay.

Significant relations of S100A4 to lymph node metastasis were reported in several types of cancers, such as stomach, breast, colon and mouth (7,18,19,29). Our results showed no apparent relation between S100A4 and nodal metastases in pulmonary adenocarcinoma, while adenocarcinoma with decreased E-cadherin and increased S100A4 showed no increased lymphatic invasion. The critical roles of S100A4 expression may be tumor-type dependent. S100A4 molecules may modulate the cell cycle, cell motility, cell adhesion and invasive properties (30-32). In conclusion, reduced E-cadherin expression combined with increased S100A4 expression is related to a poorer prognosis through hematogenous metastasis in pulmonary adenocarcinoma.

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