# Reduced expression of the *claudin-7* gene correlates with venous invasion and liver metastasis in colorectal cancer

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Abstract. Claudins, members of a large family of adherent junction proteins, regulate the integrity and function of tight junctions and influence tumorigenesis. Studies have suggested that altered levels of different claudins are related to carcinoma-cell invasion and disease progression. This study examined the relationship between the relative expression of claudin genes and clinicopathological factors, especially invasion and metastasis, in patients with colorectal cancer. We studied surgical specimens of cancer tissue and adjacent normal mucosa from 205 patients with untreated colorectal carcinoma. The relative expression levels of claudin-1, -3, -4 and -7 mRNA in cancer and in normal adjacent mucosa were measured by quantitative real-time, reverse-transcription polymerase chain reaction. The relative expression levels of the claudin-1, -3 and -4 genes were higher in cancer than in normal adjacent mucosa, whereas the relative expression of the claudin-7 gene was similar. An analysis of the relationship between the clinicopathological features and gene expression showed that reduced expression of claudin-7 correlated with venous invasion and liver metastasis. There was also a correlation between claudin-3 and -4 gene expression. Our results suggested that a reduced expression of the claudin-7 gene might lead to venous invasion and liver metastasis in colorectal cancer. Reduced expression of the claudin-7 gene may thus be a useful predictor of liver metastasis in patients with colorectal cancer.

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#### Introduction

In simple epithelium, tight junctions are positioned at the boundaries of apical and basolateral plasma membranes. These junctions are thought to play an important role in the paracellular barrier and cell polarity (1-4). Several lines of evidence indicate that the granular cell layer of stratified epithelium of the skin possesses tight junctions that are crucial for barrier function (1,2,5,6). The tight junctions consist of membrane and peripheral proteins. Claudins are membrane proteins composed of four transmembrane domains and two extracellular loops, through which they bind to corresponding claudins in cell-to-cell contact. Claudin-1, -3, -4 and -7 are four representative members of the 24-claudin multigene family (4), associated with cancer. An enhanced expression of claudin-1 has been reported in colorectal cancer (7). Ovarian epithelial cells that express claudin-3 and -4 show increased invasiveness in vitro (8). Claudin-4 is a potent inhibitor of the invasiveness and phenotype of pancreatic cancer cells (9). The loss of claudin-7 expression has been observed in ductal carcinoma of the breast and squamous cell carcinoma of the head and neck (10,11). Usami et al (12) reported that a reduced expression of claudin-7 correlates with tumor invasion and metastasis in squamous cell carcinoma of the esophagus. However, whether the expression of claudin-1, -3, -4 and -7 is associated with the malignant potential of colorectal cancer remains to be clarified.

In this study, we measured the expression levels of the *claudin-1*, -3, -4 and -7 genes in 205 pairs of cancer tissue and adjacent normal mucosa obtained from patients with colorectal cancer. To evaluate the clinical significance of the claudins, we examined the correlation between the relative expression of these genes and the clinicopathological features.

# Materials and methods

Patients and samples. We studied surgical specimens of cancer tissue and adjacent normal mucosa obtained from 205 patients

Table I. PCR primers and conditions.

Gene	Primer	Temperature (C)	Product size (bp)
Claudin-1	5'-CCAGTTAGAAGAGGTAGTGTG-3' 5'-GAGAGGAAGGCAGTGAATC-3'	60	168
Claudin-3	5'-ACCACCACCACCAAC-3' 5'-GGGCTTCCTGGCTTCTGG-3'	65	113
Claudin-4	5'-TGCCTTGCTCACCGAAACCC-3' 5'-CCTCTAAACCCGTCCATCCACTC-3'	64.5	95
Claudin-7	5'-GGAGACGACAAAGTGAAGAAG-3' 5'-GCCATACCAGGAGCAAGC-3'	60	99
$\beta$ -actin	5'-AGTTGCGTTACACCCTTTCTTGAC-3' 5'-GCTCGCTCCAACCGACTGC-3'	60	171

with untreated colorectal carcinoma. The patients underwent surgery at the Yokohama City Medical Center, Gastroenterological Center and at the Kanagawa Cancer Center between 2002 and 2006. Informed consent was obtained from each patient and the Ethics Committees of the Yokohama City Medical Center and Kanagawa Cancer Center approved the protocol before initiation of the study. Each tissue sample was embedded in O.C.T. compound (Sakura Finetechnical Co., Ltd., Tokyo) and immediately stored at -80°C until use. No patient had any other malignancies. The histopathological features of specimens stained with hematoxylin and eosin were examined and sections that consisted of >80% carcinoma cells were used to prepare total RNA.

Quantitative real-time, reverse-transcription polymerase chain reaction (PCR). Total RNA isolated from colorectal cancer and adjacent normal mucosa was prepared with the use of Trizol (Gibco, Life Tech, Gaithersburg, MD). Complemetary DNA (cDNA) was synthesized from 2 µg of total RNA with an iScript cDNA Synthesis kit (Bio-Rad Laboratories, Hercules, CA). After synthesis, the cDNA was diluted 1:4 with water and stored at -20°C until use. Quantitative real-time PCR was performed with an iQ SYBR-Green Supermix (Bio-Rad Laboratories). PCR reactions were carried out in a total volume of 15 µl containing cDNA derived from 75 ng of RNA, 0.27  $\mu$ M of each primer, 7.5  $\mu$ l of iQ SYBR-Green Supermix containing dATP, dCTP, dGTP and dTTP at a concentration of 400  $\mu$ M each and 50 units/ml of iTag DNA polymerase. The PCR consisted of 10 min at 94°C, followed by 50 cycles of denaturation of the cDNA for 30 sec at 94°C, annealing for 30 sec at an appropriate temperature (Table I) and a primer extension for 1 min at 72°C followed by 72°C for 10 min. The PCR primer sequences of MMP2, MMP9, MT-MMP, RECK and \(\beta\)-actin, used as an internal control, are shown in Table I.

Statistical analysis. Gene expression levels of colorectal cancer were compared with those of normal adjacent mucosa with the use of the Wilcoxon test. The relationship between gene expression and potential explanatory variables, including age, gender, tumor size, histological type, depth of invasion, lymph node metastasis, location, lymphatic invasion, venous invasion and liver metastasis, were evaluated with the  $\chi^2$  test.

Associations between variables were assessed using the Mann-Whitney U test. Correlation coefficients between the different variables were calculated by simple regression analysis. Each statistical analysis was performed using Statview J 5.0 software (Abacus, CA). Two-sided P-values were calculated and a difference was considered significant at P-value <0.05.

## Results

Comparison of claudin-1, -3, -4 and -7 mRNA expression between colorectal cancer tissue and adjacent normal mucosa. Claudin-1, -3 and -4 gene expression levels were higher in cancer than in normal adjacent mucosa (P<0.001, P=0.001 and P<0.001) (Fig. 1A, B and C). The claudin-7 gene expression level of cancer did not differ significantly from that of normal adjacent mucosa (P=0.524) (Fig. 1D).

Relationship of claudin-1, -3, -4 and -7 gene expression levels to clinicopathological features. Expression levels of the claudin-1, -3, -4 and -7 genes were categorized as low or high according to their median values. The relationship between the expression of these genes and clinicopathological features was then examined. The expression levels of the claudin-1, -3, -4 and -7 genes were unrelated to age, gender, tumor size, lymph node metastasis and lymphatic invasion. There were correlations between claudin-1 expression and histological type (P=0.047) and between claudin-4 expression and tumor location (P=0.039). Moreover, a reduced expression of the claudin-7 gene correlated with venous invasion (P=0.029) and liver metastasis (P=0.022) (Table II).

Associations of claudin-1, -3, -4 and -7 gene expression with lymph node metastasis in patients with colorectal cancer. There was no significant association between the expression level of any gene and the presence or absence of lymph node metastasis (Fig. 2).

Associations of claudin-1, -3, -4 and -7 gene expression with venous invasion in patients with colorectal cancer. Claudin-3 and claudin-7 gene expression levels were higher in the absence than in the presence of venous invasion (P=0.043, P=0.001) (Fig. 3).

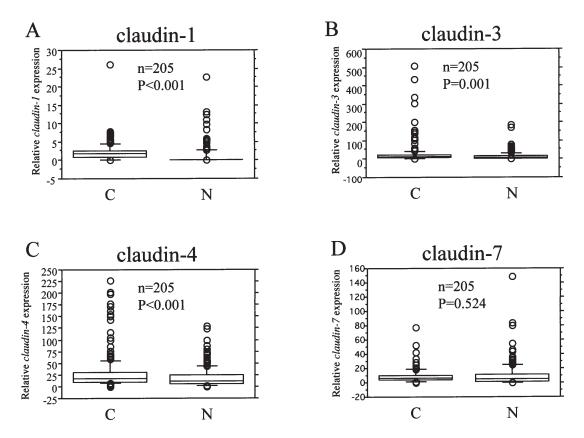


Figure 1. A comparison of *claudin-1*, -3, -4 and -7 mRNA expression levels between colorectal cancer tissue and adjacent normal mucosa. The *claudin-1*, -3 and -4 gene expression levels were higher in cancer than in normal adjacent mucosa (P<0.001, P=0.001, P<0.001). *Claudin-7* gene expression levels did not differ significantly between cancer and normal adjacent mucosa.

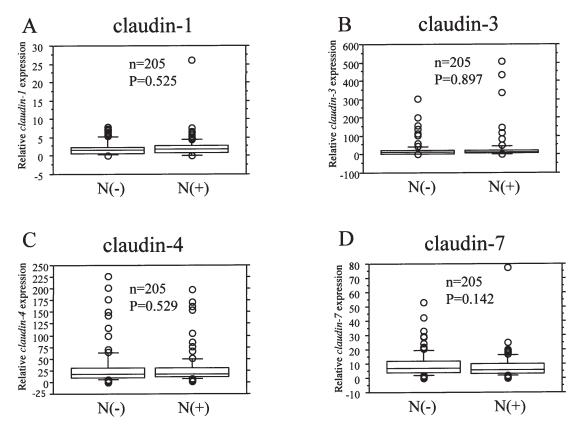


Figure 2. Associations of *claudin-1*, -3, -4 and -7 gene expression with lymph node metastasis in 205 patients with colorectal cancer. Box boundaries, the 25th and 75th percentiles of the observed values; capped bars, the 10th and 90th percentiles; solid line, median. P-values were calculated by the Mann-Whitney U test. The expression level of none of the genes examined correlated with the presence or absence of lymph node metastasis.

Table II. Relationship between the expression of claudin-1, -3, -4 or -7 genes and clinicopathological features.

Voisibloofootoonio	claudin-1	claudin-1 expression		claudin-3	claudin-3 expression		claudin-4 expression	xpression		claudin-7 expression	expression	
v anables/categones	low (n=102)	high (n=103)	P-value	low (n=102)	high (n=103)	P-value	low (n=102)	high (n=103)	P-value	low (n=102)	high (n=103)	P-value
Age .	65.6±11.3	66.0±10.3	0.775	65.6±11.1	66.0±10.5	0.805	65.7±11.2	65.8±10.4	0.917	65.1±11.0	66.5±10.6	0.344
Gender Male Female	58	54 49	0.524	51	61	0.160	50	62	0.108	50	62	0.108
Size  ≤5 cm  >5 cm	8 4	57 46	0.826	57 45	58 54 54	0.951	56 46	5 65	0.731	2 54 8 4	61 42	0.365
Histological type Well differentiated Moderately differentiated Poorly differentiated	28 54 20	33 62 8	0.047	26 60 16	35 56 12	0.362	29 60 13	32 56 15	0.809	28 60 14	33 56 14	0.762
Depth of invasion T1 T2 T3 T4	10 44 41 7	9 50 39 5	0.846	o 14 4 8	10 53 36 4	0.294	7 52 36 7	21 24 44 8	0.320	10 38 46 8	9 5 8 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	0.085
Lymph node metastasis Absent Present	50	45 58	0.930	46	49 54	0.722	51	44 59	0.296	42	53	0.140
Location Colon Rectum	61	51	0.139	62 40	50 53	0.784	96	46	0.039	56 46	56 47	0.940
Lymphatic invasion Absent Present	99	95	0.843	95	67 36	0.924	75 27	59 44	0.145	70 32	64 39	0.829
Venous invasion Absent Present	40	37	0.237	40	37	0.626	35	42 61	0.340	28	49 54	0.029
Liver metastasis Absent Present	70	934	0.802	934	70	0.802	72 30	67	0.396	59	80	0.022

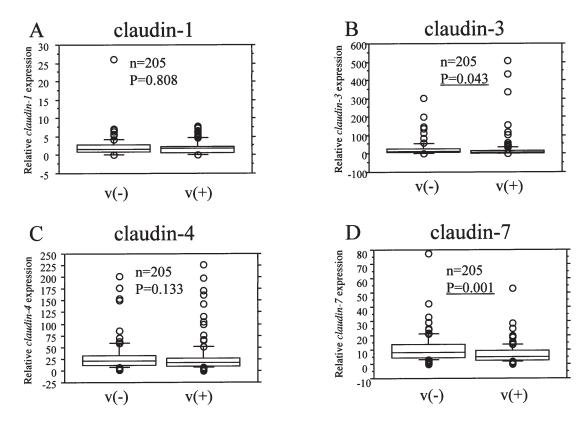


Figure 3. Associations of *claudin-1*, -3, -4 and -7 gene expression levels with venous invasion in 205 patients with colorectal cancer. Box boundaries, the 25th and 75th percentiles of the observed values; capped bars, the 10th and 90th percentiles; solid line, median. P-values were calculated by the Mann-Whitney U test. *Claudin-3* and -7 gene expression levels were higher in the absence than in the presence of venous invasion (P=0.043, P=0.001).

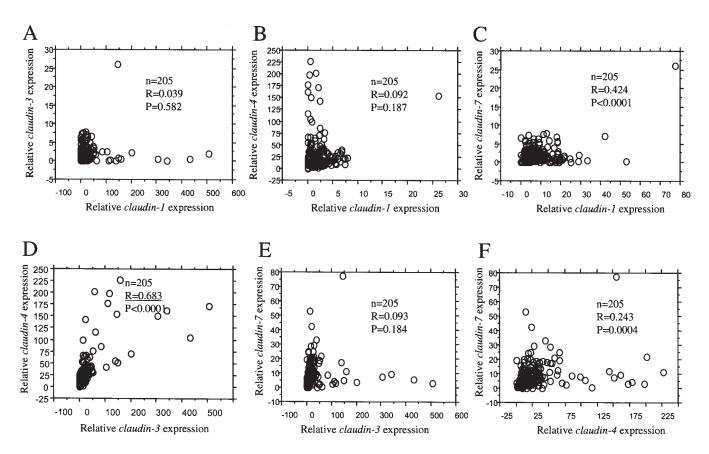


Figure 4. Correlation among *claudin-1*, -3, -4 and -7 gene expression levels in colorectal cancers. Each gene expression level is relative to that of the  $\beta$ -actin gene. The expression of the *claudin-3* gene correlated with that of the *claudin-4* gene (R=0.683).

Correlations among claudin-1, -3, -4 and -7 gene expression. Correlations between gene expression levels are shown in Fig. 4. The expression of the *claudin-3* gene correlated with that of the *claudin-4* gene (R=0.683).

#### Discussion

Cell-to-cell adhesiveness is generally reduced in various human cancers. The dissociation of cancer cells from primary cancer nests is a crucial step in metastasis. The suppression of cell-to-cell adhesiveness may trigger the release of cancer cells from primary cancer nests and increase tumor invasiveness (13). In this study, we examined the expression levels of the *claudin-1*, -3, -4 and -7 genes in colorectal cancer and the relationship of such levels to clinicopathological variables.

We compared the mRNA expression of each *claudin* gene between colorectal cancer tissue and adjacent normal mucosa. Dhawan et al (14) reported that the expression of claudin-1 is higher in human primary colon carcinoma and metastasis than in normal colorectal tissue. Pan et al (15) found that the expression of claudin-3 and -4 is significantly higher in human endometrial carcinoma than in normal endometrial tissue at the protein and mRNA levels. As for claudin-7, Kominsky et al (10) reported that this gene is down-regulated in breast cancers as compared with normal breast tissue. However, Sobel et al (16) found no significant difference in the expression of claudin-7 between human invasive cervical carcinoma and normal cervical tissue. In our study, expression levels of the claudin-1, -3 and -4 genes were higher in cancer than in normal adjacent mucosa, whereas the expression level of the *claudin-7* gene cancer did not differ significantly between cancer and normal adjacent mucosa.

We then examined the relationship between claudin gene expression levels and clinicopathological features. Sheehan et al (17) reported that a decreased expression of claudin-1 correlates with high tumor grade and biochemical disease recurrence in prostate carcinomas. Resnick et al (18) showed that a low expression level of claudin-1 is associated with a higher tumor grade and recurrence in patients with colorectal cancer. In our study, claudin-1 expression was associated with the histological type. As for claudin-3 and -4, Sheehan et al (17) reported that the expression of claudin-3 correlates with advanced-stage tumors and recurrence, whereas the expression of claudin-4 correlates with only advanced-stage tumors. Pan et al (15) found a slight though insignificant trend towards positive associations of claudin-3 and -4 levels with tumor grade and disease stage in patients with endometrial carcinoma. Our study found no significant relationship between the expression level of the *claudin-3* gene and any clinicopathological feature. The expression of the claudin-4 gene correlated with only tumor location. As for claudin-7, Kominsky et al (10) reported that the loss of claudin-7 expression is associated with nodal metastasis in primary breast carcinomas. Sauer et al (19) found that a reduced expression of claudin-7 correlates with metastatic disease in breast carcinoma. Usami et al (12) demonstrated that a reduced expression of claudin-7 correlates with metastasis in squamous cell carcinoma of the esophagus. In our study, a reduced expression of the claudin-7 gene correlated with venous invasion and liver metastasis in colorectal cancer.

When expression levels of the *claudin-1*, -3, -4 and -7 genes were contrasted with the presence or absence of lymph node metastasis, no correlation was noted for any gene. We also examined potential correlations of gene expression levels with the presence or absence of venous invasion. Sauer *et al* (19) reported that a reduced expression of claudin-7 correlates with metastatic disease. Usami *et al* (12) found that a reduced expression of claudin-7 correlates with tumor invasion in squamous cell carcinoma of the esophagus. In our study, *claudin-3* and -7 gene expression levels were higher in the absence than in the presence of venous invasion. This finding suggested that reduced *claudin-3* or -7 gene expression levels might contribute to venous invasion in colorectal cancer.

We then examined correlations among *claudin-1*, *-3*, *-4* and *-7* gene expression in colorectal cancers. Expression of the *claudin-3* gene was found to correlate with that of the *claudin-4* gene.

In conclusion, our results show that a reduced expression of the *claudin-7* gene correlates with venous invasion and liver metastasis in colorectal cancer. Reduced levels or the absence of claudin-7 expression may thus be a novel marker or predictor of metastasis.

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