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ORIGINAL ARTICLE

Basic Study

Loss of stromal caveolin-1 expression in colorectal cancer predicts poor survival

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Abstract

AIM: To investigate the clinicopathological significance and prognostic value of caveolin-1 (CAV-1) in both tumor and stromal cells in colorectal cancer (CRC).

METHODS: A total of 178 patients with CRC were included in this study. The correlation between CAV-1

expression and clinicopathologic features and survival was studied.

RESULTS: CAV-1 expression was detected in tumor and stromal cells. The expression of stromal CAV-1 was closely associated with histological type (P = 0.022), pathologic tumor-node-metastasis stage (P = 0.047), pathologic N stage (P = 0.035) and recurrence (P = 0.000). However, tumor cell CAV-1 did not show any correlation with clinical parameters. Additionally, the loss of stromal CAV-1 expression was associated with shorter disease-free survival (P = 0.000) and overall survival (P = 0.000). Multivariate analysis revealed that the loss of stromal CAV-1 expression was an independent prognostic factor for both overall survival (P = 0.014) and disease-free survival (P = 0.006).

CONCLUSION: The loss of stromal CAV-1 expression in CRC was associated with poor prognosis and could be a prognostic factor for CRC patients.

Key words: CAV-1; Stroma; Colorectal cancer; Prognosis

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Core tip: Caveolin-1 (CAV-1), an essential structural protein of the endocytic caveolae plasma membrane, plays a major role in modulating tumorigenic processes. Recent studies have revealed that the loss of stromal CAV-1 results in an activated tumor microenvironment and is significantly related to tumor recurrence and a poor prognosis for many tumors. However, the association between CAV-1 and colorectal cancer remains unknown. In our study, we observed CAV-1 expression in both tumor and stromal cells. Our results demonstrate that the loss of stromal CAV-1 is an independent predictor of poor overall survival and disease-free survival, whereas CAV-1 expression in tumor cells has no prognostic value.



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INTRODUCTION

Colorectal cancer (CRC) is the third most common cancer and the third leading cause of cancer mortality in the United States, with an estimated 142820 new cases and 50830 deaths per year^[1]. Over the past ten years, the rate of death from CRC has declined by 3%, but 30% to 40% of CRC patients will develop distant metastases, and 50% will die of CRC recurrence^[2].

The tumor microenvironment has recently been recognized to play an important role in determining tumor initiation and progression^[3-6]. The tumor microenvironment is composed of immune cells, stromal cells (including cancer-associated fibroblasts (CAFs), adipocytes, and bone marrow-derived progenitors), and the vasculature^[5,7]. Under normal physiological conditions, the stroma serves as a critical barrier to prevent malignant transformation. However, during neoplastic tumorigenesis, the stroma facilitates tumor progression and metastasis in response to molecular signals derived from carcinoma cells and other host cell types.

Caveolin-1 (CAV-1), an essential structural protein of endocytic caveolae plasma membrane invaginations, is present in most mammalian cells, such as adipocytes, endothelial cells, pneumocytes, fibroblasts, and smooth muscle cells^[8,9]. CAV-1 plays a major role in modulating tumorigenic processes through its various functions, such as gene regulation, membrane trafficking, and signal transduction^[10-12]. However, CAV-1 shows a compartment-dependent role in tumors. Furthermore, the role of CAV-1 is controversial in epithelial tumor cells. Several authors have proposed that high CAV-1 expression in tumor cells predicts poor survival^[13,14]. In contrast, El-Gendi et al^[15] reported that CAV-1 expression in tumor cells has no prognostic value. Recent studies have revealed that the loss of CAV-1 in the tumor stroma results in an activated tumor microenvironment and is significantly related to early tumor recurrence, metastasis, and poor clinical outcome in breast cancer, prostate cancer, and other solid tumors^[16-19].

However, the association between CAV-1 and colorectal cancer remains unknown. Thus, to explore the relationship between CAV-1 expression and colorectal cancer, we analyzed CAV-1 expression in CRC stromal cells and tumor cells in a series of human CRC tissue sections.

MATERIALS AND METHODS

Patients and specimens

Colorectal tissue samples were obtained from the pathology department at the First Affiliated Hospital of Sun Yat-sen University between May 2004 and November 2009. The patients included in this study fulfilled the following criteria: (1) pathologically confirmed diagnosis of colorectal adenocarcinoma; (2) no treatment prior to curative excision of the primary tumor; (3) no previous malignancy or second primary tumor; (4) no severe coincident diseases; and (5) the availability of clinical information and follow-up data. The tumors were classified according to the American Joint Committee on Cancer staging manual (7th edition).

All patients were followed every 3 mo for the first year, every 6 mo from the second to fifth years and every 12 mo after 5 years. The study protocol was approved by the Ethics and Scientific Committee of the First Affiliated Hospital of Sun Yat-sen University and conforms to the Declaration of Helsinki. All patients and their families provided written informed consent prior to surgery for their information to be stored in the hospital database and used for research.

Immunohistochemistry

Tissue section immunohistochemistry was performed according to the instructions provided by the manufacturer. Briefly, 4 µm formalin-fixed, paraffinembedded tissue sections were deparaffinized and rehydrated through a graded alcohol series. Antigen retrieval was performed in 10 mmol/L sodium citrate (pH 6.0) for 5 min using a pressure cooker and the sections were allowed to cool for 45 min at room temperature. After blocking with 3% hydrogen peroxide for 10 min to inactivate endogenous peroxidase, the slides were washed three times with phosphate-buffered saline and incubated with 10% goat serum for 30 min at room temperature. The sections were then incubated with an anti-CAV-1 antibody (rabbit monoclonal, dilution 1:100, Cell Signaling Technology, CST) overnight at 4 °C, and the slides were then incubated with an HRPconjugated sheep anti-rabbit secondary antibody (GTVision; Shanghai, China) for 30 min at room temperature. Immunoreactivity was revealed using 3,3'-diaminobenzidine and counterstained with Mayer's hematoxylin. A known positive tissue sample (breast cancer slide) was used as a positive control.

Evaluation of immunostaining

The staining of stromal CAV-1 was scored semiquantitatively as negative (0, no staining), weak (1, either diffuse weak staining or strong staining in less than 30% of the stromal cells) or strong (2, strong

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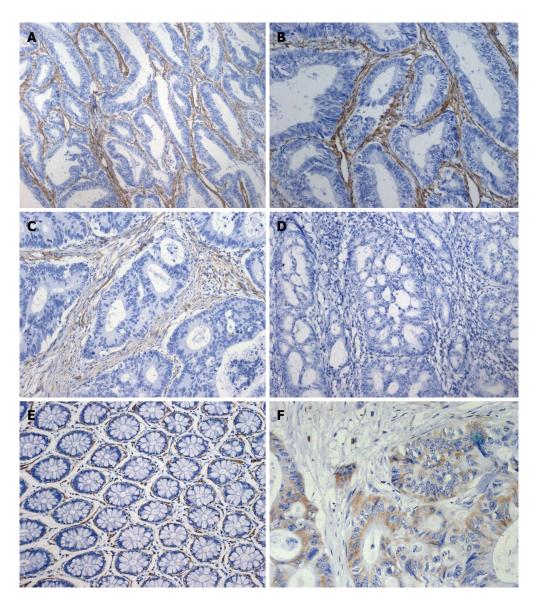


Figure 1 Immunohistochemical staining for caveolin-1. High caveolin-1 (CAV-1) expression in colorectal cancer stroma (A: × 200; B: × 400). Panel C (× 200) shows moderate stromal CAV-1 expression, and panel D (× 200) shows negative stromal CAV-1 expression. Panel E (× 200) depicts normal colorectal tissue. Panel F (× 400) demonstrates CAV-1 expression in colorectal tumor cells.

staining of 30% or more of the stromal cells)^[20]. We also evaluated CAV-1 expression in tumor cells. Any expression in tumor cells was considered to be positive CAV-1 staining^[20]. The immunostaining results were evaluated by two independent pathologists.

Statistical analysis

All statistical calculations were carried out using SPSS 17.0 statistical software. The associations between CAV-1 expression and various clinical parameters were evaluated using the Mann-Whitney test, Spearman's test or χ^2 test. The Kaplan-Meier test was used to evaluate disease-free survival (DFS) and overall survival (OS), and the survival curve was compared using the Log rank test. Multivariate Cox regression models were applied to evaluate CAV-1 expression and other prognostic factors with respect

to DFS and OS. All P-values are two-sided and were considered statistically significant at the level of < 0.05.

RESULTS

The present study included 178 matched colorectal cancer tissues and 30 paraneoplastic normal tissues, which were more than 5 cm from the primary tumor sites. Of the 178 CRC patients, 103 were male (57.9%) and 75 (42.1%) were female. The median age was 54 years (range: 24-85 years). The median follow-up period was 50 (range: 9-127) months for all patients.

CAV-1 expression was observed in both the tumor and stromal cells. Additionally, adipocytes, endothelial cells, and perineurial cells showed strong CAV-1 expression and served as internal positive controls.



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clinicopathologic parameters <i>n</i> (%)						
Variable		P-value				
	0	1	2			
Age (yr)				0.095 ¹		
≤ 50	22 (33.8)	19 (29.2)	24 (36.9)			
> 50	27 (23.9)	31 (27.4)	55 (48.7)			
Gender				0.404^{1}		
Male	26 (25.2)	29 (28.2)	48 (46.6)			
Female	23 (30.7)	21 (28.0)	31 (41.3)			
Tumor size (cm)				0.688^{1}		
< 5	28 (27.7)	30 (29.7)	43 (42.6)			
≥ 5	21 (27.3)	20 (26.0)	36 (46.8)			
Tumor location				0.378^{1}		
Colon	32 (30.8)	28 (26.9)	44 (42.3)			
Rectum	17 (23.3)	22 (30.1)	34 (46.6)			
Histological type				0.022 ^{2,a}		
Well	1 (12.5)	4 (50.0)	3 (37.5)			
Moderate	31 (23.8)	35 (26.9)	64 (49.2)			
Poor	17 (42.5)	11 (27.5)	12 (30.0)			
Pathologic T stage				0.118^{2}		
T1	1 (50.0)	0 (0)	1 (50.0)			
T2	0 (0)	5 (35.7)	9 (64.3)			
T3	45 (29.2)	44 (28.6)	65 (42.2)			
T4	3 (37.5)	1 (12.5)	4 (50.0)			
Pathologic N stage	()	~ /	()	0.035 ^{2,a}		
N0	10 (16.4)	20 (32.8)	31 (50.8)			
N1	23 (29.1)	23 (29.1)	33 (41.8)			
N2	16 (42.1)	7 (18.4)	15 (39.5)			
Pathologic TNM stage		~ /	()	0.047 ^{2,a}		
I	0 (0)	4 (50.0)	4 (50.0)			
П	10 (18.9)	16 (30.2)	27 (50.9)			
Ш	39 (33.3)	30 (25.6)	48 (41.0)			
Tumor cell Cav-1				0.697^{1}		
Negative	26.7 (61.8)	29.1 (63.8)	76 (44.2)			
Positive	3 (50.0)	0 (0)	3 (50.0)			
Recurrence	()	. (*)	()	< 0.001 ^{1,a}		
No		28 (24.1)	68 (58.6)			
Yes		21 (36.2)	10 (17.2)			
		()	()			

Table 1 Association of stromal caveolin-1 expression with clinicopathologic parameters n (%)

¹Mann-Whitney test; ²Spearman's test; ^aP < 0.05, stromal CAV-1 vs clinical variables. CAV-1: Caveolin-1; TNM: Tumor-node-metastasis.

Analyses for CAV-1 expression in the stroma revealed positive expression in 129 patients (72.5%) and negative expression in 49 patients (27.5%). Of the 129 positive cases, 50 (28.1%) had a score of 1, and 79 (44.4%) had a score of 2. Representative examples are shown in Figure 1. The correlations between stromal CAV-1 and clinical variables are listed in Table 1. The expression of stromal CAV-1 was closely associated with histological type (P = 0.022), pathologic tumor-node-metastasis (TNM) stage (P = 0.047), pathologic N stage (P = 0.035) and recurrence (P < 0.001). However, there was no significant correlation between the expression level of CAV-1 and age, gender, tumor size, tumor location, pathologic T stage, or tumor cell CAV-1.

Interestingly, with regard to CAV-1 expression in tumor cells, we only observed 6 tissues that were positive for expression in our study. However, as presented in Table 2, tumor cell CAV-1 did not showed any correlation with clinical parameters.

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Table 2 Association between tumor cell caveolin-1 expression and clinicopathologic parameters n (%)

Variable	Cav	Cav-1 <i>P</i> -value	
	Negative	Positive	
Age (yr)			1.000
≤ 50	63 (96.9)	2 (3.1)	
> 50	109 (96.5)	4 (3.5)	
Gender			0.698
Male	100 (97.1)	3 (2.9)	
Female	72 (96.0)	3 (4.0)	
Tumor size (cm)			0.237
< 5	96 (95.0)	5 (5.0)	
≥ 5	76 (98.7)	1 (1.3)	
Tumor location			0.694
Colon	101 (97.1)	3 (2.9)	
Rectum	71 (95.9)	3 (4.1)	
Histological type			0.498
Well	8 (100)	0 (0)	
Moderate	124 (95.4)	6 (4.6)	
Poor	40 (100.0)	0 (0)	
Pathologic T stage			0.073
T1	2 (100.0)	0 (0)	
T2	14 (100)	0 (0)	
T3	150 (97.4)	4 (2.6)	
T4	6 (75.0)	2 (25.0)	
Pathologic N stage			0.304
N0	57 (93.4)	4 (6.6)	
N1	77 (97.5)	2 (2.5)	
N2	38 (100)	0 (0)	
Pathologic TNM stage			0.158
Ι	8 (100)	0 (0)	
Π	49 (92.5)	4 (7.5)	
Ш	115 (98.3)	2 (1.7)	
Stromal Cav-1			0.208
0	46 (93.9)	3 (6.1)	
1	50 (100)	0 (0)	
2	76 (96.2)	3 (3.8)	
Recurrence			0.180
No	110 (94.8)	6 (5.2)	
Yes	58 (100)	0 (0)	

CAV-1: Caveolin-1; TNM: Tumor-node-metastasis.

Survival analysis

Next, we evaluated the prognostic value of CAV-1 expression in colorectal cancer. Figure 2 shows that the patients who were positive for stromal CAV-1 expression had significantly longer overall survival and disease-free survival (P < 0.001 for both) than the patients who were negative. Of note, the patients with high levels of stromal CAV-1 (score = 2) had a good prognosis, with 89.8% of the patients surviving the follow-up period. Similarly, 77.4% of the patients with moderate stromal CAV-1 staining (score = 1) survived. In contrast, 46.9% of the patients who were negative for stromal CAV-1 expression (score = 0) survived. The 5-year survival rates showed similar patterns. CRC patients with high stromal CAV-1 had a good 5-year survival rate (92%), whereas CRC patients with moderate or absent stromal CAV-1 expression had progressively worse 5-year survival rates (61% and 46%, respectively). We also analyzed the prognostic significance of CAV-1 expression

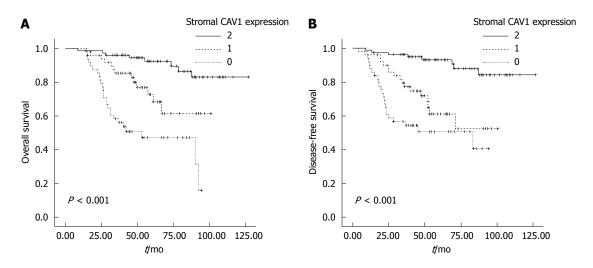


Figure 2 Kaplan-Meier curves of overall survival and disease-free survival. Patients with high stromal caveolin-1 expression have good overall survival (A) and disease-free survival (B).

Table 3 Univariate and multivariate analyses of prognostic factors in colorectal cancer with respect to overall survival and disease-free survival					
Parameter	Univariate analysis <i>P</i> -value	Multivariate analysis HR (95%CI)	<i>P</i> -value		
OS					
Age (> 50)	0.185	-	1.632		
Gender	0.794	-	1.431		
Tumor location	0.384	-	1.414		
Tumor cell Cav-1	0.216	-	0.742		
Stromal Cav-1	< 0.001 ^a	0.578 (0.373-0.895)	0.014^{a}		
Tumor size	0.443	-	0.382		
Histological type	0.533	-	0.666		
Pathological T stage	0.067	-	0.347		
Pathological N stage	< 0.001 ^a	-	0.868		
TNM	0.025 ^a	-	0.378		
Recurrence DFS	< 0.001 ^a	2.065 (1.121-3.803)	< 0.001 ^a		
Age (> 50)	0.165	-	0.218		
Gender	0.664	-	0.833		
Tumor location	0.061	-	0.325		
Tumor cell Cav-1	0.189	-	0.705		
Stromal Cav-1	< 0.001 ^a	0.546 (0.353-0.844)	0.006 ^a		
Tumor size	0.603	-	0.511		
Histological type	0.478	-	0.553		
Pathological T stage	0.171	-	0.888		
Pathological N stage	0.002 ^a	-	0.510		
TNM	0.184	0.451 (0.233-0.874)	0.018^{a}		
Recurrence	< 0.001 ^a	1.863 (1.106-2.434)	$< 0.001^{a}$		

 aP < 0.05, univariate analysis vs multivariate analysis. CAV-1: Caveolin-1; TNM: Tumor-node-metastasis; OS: Overall survival; DFS: Disease-free survival.

in tumor cells, but no prognostic significance for either OS or DFS was found (P = 0.216 and 0.189, respectively).

A univariate analysis of possible prognostic indicators identified stromal CAV-1, pathologic N stage, pathologic TNM stage and recurrence as indicators of OS, and stromal CAV-1, pathologic N stage and recurrence as indicators of DFS (Table 3). In a multivariable analysis, stromal CAV-1 expression was a significant independent prognostic factor for OS and DFS (P = 0.006, HR = 0.546, 95%CI: 0.353-0.844; P = 0.014, HR = 0.578, 95%CI: 0.373-0.895, respectively). Moreover, the results also showed that recurrence was an independent prognostic factor for OS and that TNM and recurrence were independent prognostic factors for DFS (Table 3).

DISCUSSION

In this study, we evaluated the expression of CAV-1 in the stroma and tumor cells of CRC. Our results demonstrated that the loss of CAV-1 expression in the stroma is a strong and independent predictor of poor OS and DFS, whereas CAV-1 expression in tumor cells had no prognostic value. Moreover, stromal CAV-1 expression was closely associated with histological type (P = 0.022), pathologic TNM stage (P = 0.047), pathologic N stage (P = 0.035) and recurrence (P < 0.001). However, CAV-1 expression in tumor cells did not show any correlation with the clinical parameters.

Tumors, which are composed of tumor cells and stromal cells, grow within a complex tumor microenvironment. CAFs were recently demonstrated to promote tumor initiation, prevent cancer cell apoptosis, induce cancer cell proliferation, and stimulate tumor angiogenesis by secreting a large amount of growth factors, extracellular matrix components, and matrix metalloproteinases^[4,6]. Although the underlying mechanisms are not fully elucidated, the loss of CAV-1 in the stroma plays a major role.

Witkiewicz *et al*^[16] reported that the absence of stromal CAV-1 expression predicts early tumor recurrence and poor clinical outcome in human breast cancer. Di Vizio *et al*^[21] found that the loss of stromal CAV-1 expression was closely related

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to disease progression and metastasis. Jia et al^[18] indicated that the down-regulation CAV-1 expression in the stroma predicts lymph node metastases, early tumor recurrence, and poor prognosis in esophageal squamous cell carcinoma. Consistent with previous studies, we found that the loss of CAV-1 expression in the stroma is a strong and independent predictor of a poor prognosis in CRC. In our cohort, the patients with high levels of stromal CAV-1 had a better prognosis, with 89.8% surviving the followup period. Similarly, 77.4% of the patients with moderate stromal CAV-1 staining survived. In contrast, 46.9% of the patients who were negative for stromal CAV-1 expression survived. We found a similar pattern for the 5-year survival rates. CRC patients with high stromal CAV-1 had a good (92%) 5-year survival rate. In contrast, CRC patients with moderate levels or absent CAV-1 expression in the stroma had progressively worse 5-year survival rates (61% and 46%, respectively). These results suggest that stromal CAV-1 expression could be used as an important prognostic factor for CRC patients.

In accordance with previous studies, we also found that the absence of CAV-1 expression in the stroma was significantly associated with early tumor recurrence^[17,20]. Patients who had lower stromal CAV-1 expression were more likely to experience tumor recurrence. This finding highlights the role of the tumor stroma in determining disease recurrence in CRC patients. Therefore, new drugs that target the tumor stroma may have unexpected effects in CRC therapies.

We also found that the loss of stromal CAV-1 expression was significantly associated with lymph node invasion. Thus, the stromal CAV-1 status may serve as a predictor of lymph node invasion. CAV-1 expression in the stroma of a biopsy sample may also be useful in deciding whether endoscopic surgery should be performed for patients with early CRC. Moreover, considering that the loss of CAV-1 expression in the stroma may predict a high risk of lymph node invasion, it would also be informative when choosing a postoperative therapy for patients with no lymph node invasion.

The role of CAV-1 in tumor cells is controversial. Steffens *et al*^[14] reported that low CAV-1 expression in renal cell carcinoma predicts a good clinical outcome compared with patients with high CAV-1 expression. However, El-Gendi *et al*^[15] reported that CAV-1 expression in tumor cells has no prognostic value. Additionally, Friedrich *et al*^[22] reported that CAV-1 deficiency in Apc (min/+) mice facilitates colorectal initiation. Luo *et al*^[23] suggested that CAV-1 could be a biomarker of early-onset CRC. In our study, we also examined CAV-1 expression in epithelial tumor cells in the same patients. However, we only found 6 patients with positive tumor cell expression, and our results revealed no significant

correlation between tumor cell CAV-1 and clinical characteristics or prognosis. The discrepancy could be explained by the tumor specificity of CAV-1 expression. Additionally, variation in scoring methods could explain some of the observed discrepancies.

Although the role of CAV-1 in the tumor stroma is not fully understood, recent findings support the hypothesis that CAV-1 plays a crucial role in the stroma. Koleske et $al^{[24]}$ were the first to find that CAV-1 functions as a tumor suppressor in fibroblasts. The reduction of CAV-1 levels by the constitutive activation of oncogenes, such as c-Myc, v-Src, H-Ras (G12V), and Neu/ErB2, can significantly promote tumor growth^[25-27]. The tumor suppressor p53 can also transcriptionally regulate stromal CAV-1 expression and p53 inactivation induces CAV-1 downregulation and promotes tumor progression^[28]. In a xenograft model, CAV-1-deficient cancer-associated fibroblasts were found to promote both tumor growth and angiogenesis^[29]. Several studies have found that the loss of CAV-1 in fibroblasts is sufficient to induce the conversion of benign stromal fibroblasts to tumorassociated fibroblasts via the TGF- β pathway^[30,31]. In turn, CAFs promote tumor initiation, progression, and prevent cancer cell apoptosis. Recently, Pavlides et al^[32] proposed a new tumor metabolism model, the Reverse Warburg Effect. In this model, the loss of CAV-1 causes stromal CAFs to undergo autophagy and aerobic glycolysis. As a consequence, the CAFs secrete energy-rich metabolites and chemical building blocks to adjacent tumor cells to prompt growth. The results of the present study support this hypothesis and indicate that the loss of CAV-1 expression in CAFs plays an important role in supporting tumor growth.

There are several limitations to this study. This research is a retrospective study, and there are limitations to any retrospective data collection, which is prone to bias. The relatively small sample size and the reliability of immunohistochemical techniques are other limitations.

In conclusion, our results suggest that the loss of CAV-1 expression in the stroma could be a predictor of poor clinical outcomes. As such, stromal CAV-1 levels could be used as a valuable biomarker for stratifying CRC patients into high-risk and lowrisk groups at diagnosis; this information could be used to provide a more personalized approach to postoperative therapy. Further studies are needed to elucidate the mechanisms of stromal CAV-1 reduction and the tumor-stroma cross-talk that is crucial for tumor growth and metastasis.

COMMENTS

Background

Caveolin-1 (CAV-1), an essential structural protein of the endocytic caveolae plasma membrane, plays a major role in modulating tumorigenic processes. Recent studies have revealed that the loss of stromal CAV-1 results in an

activated tumor microenvironment and is significantly related to early tumor recurrence, metastasis, and poor prognosis in breast cancer, prostate cancer, and other solid tumors. However, the association between CAV-1 and colorectal cancer (CRC) remains unknown.

Research frontiers

Tumors are composed of tumor cells and stromal cells. Stromal cells have recently been recognized to play an important role in determining tumor initiation and progression. Under normal physiological conditions, the stroma serves as a critical barrier to prevent malignant transformation. However, during neoplastic tumorigenesis, the stroma facilitates tumor progression and metastasis in response to molecular signals derived from carcinoma cells and other host cell types. CAV-1, an essential structural protein of endocytic caveolae plasma membrane invaginations, is present in most mammalian cells, such as adipocytes, endothelial cells, fibroblasts, and smooth muscle cells. CAV-1 plays a major role in modulating tumorigenic processes. Recent studies have revealed that the loss of CAV-1 in the tumor stroma results in an activated tumor microenvironment and is significantly related to early tumor recurrence, metastasis, and poor clinical outcome in breast cancer, prostate cancer and other solid tumors.

Innovations and breakthroughs

This study is the first to report CAV-1 expression in both tumor and stromal cells of colorectal cancer. Moreover, our results demonstrate that the loss of CAV-1 expression in the stroma is a strong and independent predictor of poor overall survival and disease-free survival, whereas CAV-1 expression in tumor cells has no prognostic value. Additionally, CAV-1 expression in the stroma was closely associated with histological type, pTNM stage, pN stage and recurrence. Conversely, CAV-1 expression in tumor cells did not show any correlation with clinical parameters.

Applications

The results suggest that stromal CAV-1 levels could be used as a valuable biomarker for stratifying CRC patients into high-risk and low-risk groups at diagnosis; this information could be used to provide a more personalized approach to postoperative therapy.

Terminology

The stroma is a component of tumors. During neoplastic tumorigenesis, the stroma facilitates tumor progression and metastasis. Caveolin-1, an essential structural protein of endocytic caveolar plasma membrane invaginations, is present in most stromal cells.

Peer review

In the manuscript "Loss of stromal caveolin-1 expression in colorectal cancer predicts poor survival", the authors found loss of stromal CAV-1 was closely associated with histological type, pathologic TNM stage, pathologic N stage and recurrence in colorectal cancer. The higher stromal CAV-1 expression, the better patients' survival. It indicates that CAV-1 could be a prognostic factor for patients with CRC.

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