

## Original Article

# Relationship between RECK/matrix metalloproteinase and recurrence/metastasis of giant cell tumor of spine

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**Abstract:** Giant cell tumor of spine (GCTS) is one common benign tumor with high recurrence rate after surgery. The finding of novel index for predicting recurrence is thus of critical importance. Matrix metalloproteinase (MMP) can work as prognostic indexes for various tumors, and is shown to be negatively related with RECK gene expression. This study thus examined the levels of MMP-2, MMP-9 and RECK in GCTS patients, in order to elucidate the correlation between gene expression and recurrent tumors. Tumor tissues were collected to examine the expressional profiles of MMP-2, MMP-9 and RECK genes, along with post-operative follow-ups on all patients. The correlation between gene expression and clinical features was analyzed by SPSS 13.0 software. There was a significant correlation between Enneking stage or Campanacci grade with the expression of MMP-2, MMP-9 and RECK genes ( $P < 0.05$ ). Those genes were unrelated with patients' age or sex ( $P > 0.05$ ). The recurrence of tumor is correlated with expression of RECK, MMP-2 and MMP-9 in tumor tissues ( $P < 0.05$ ). Using Logistic regression, enhanced RECK expression can decrease the risk of GCTS recurrence while MMP-9 up-regulation increased the risk of recurrence ( $P < 0.05$ ). No significant correlation existed between MMP-2 and recurrence. RECK/MMP-9 expressions are closely correlated with the recurrence of GCTS, and can work as important indexes predicting tumor prognosis.

**Keywords:** Giant cell tumor of bone, Matrix metalloproteinase-2, matrix metalloproteinase-9, RECK

## Introduction

Giant cell tumor of bone (GCTB), also named as osteoclastoma, is one common primary benign tumor in bone tissues, occupying 4%~5% of all primary bone tumors. Due to its high recurrence [1] and invasive nature [2], and certain pulmonary metastasis, GCTB is believed to be one potential malignant tumor. GCTB is mostly common in females between 20~45 years old, with higher incidence in Asian people compared to Western population [3]. GCTB is mostly occurred in epiphysis of upper and lower tibia, and frequently leads to pathological bone fracture due to lytic bone lesion. Giant cell tumor of spine (GCTS) occupies about 3%~6% of total cases of GCTB [4]. Due to its special location of occurrence, GCTS may cause dysfunctions to certain extents following compression of spinal cord or nerves, or even causing paralysis. Surgical resection is the major

approach in treating GCTS, but having a post-operative rate as high as 30%~50% [3]. Therefore the prediction of recurrence after GCTS surgery is of critical importance.

Matrix metalloproteinases (MMPs) is a family of zinc ion-dependent endonuclease family with highly conserved sequence. They are able to degrade most proteins within basal membrane and extracellular matrix (ECM). MMPs, including gelatinase, collagenase and matrilysin, are produced in the form of inactive zymogen, and need to be activated for exerting proteinase activity. Certain tumor-inhibitory gene fragments exert their functions by interacting with MMPs [5]. MMPs thus play a crucial role in the invasion and migration of tumors. As one important member of MMPs family, MMP-2 is also called gelatinase A, and is secreted by various cells to degrade multiple ECM. In a cascade reaction initiated by MMPs, MMP-2 is key

enzyme in the activation. Tumor cells can potentiate their invasion and metastatic function via over-expressing MMP-2, which, therefore, can work as prognostic factor for multiple tumors including breast cancer [6, 7], endometrium carcinoma [8] and oral squamous cell carcinoma [9]. Previous study has indicated the expression of MMP-2 in basal cells of GCTB [10]. Microarray analysis also revealed elevated MMP-2 expression in GCTB patients [11]. On the other hand, gelatinase B (MMP-9) is also related with multiple tumors including colorectal cancer [12] and gastric carcinoma [13]. Other studies have also suggested potent expression of MMP-9 in GCTB tissues and the relation with tumor invasion/metastasis [14, 15]. Therefore, MMP-2 and MMP-9 may be both related with recurrence and/or metastasis of GCTB.

Reversion-inducing cysteine-rich protein with Kazal motifs (RECK) were firstly identified in 1998 for inhibiting MMPs expression and activity. RECK expression level is negatively correlated with that in MMPs, and can inhibit tumor's invasion and metastasis [5, 16, 17]. We thus hypothesized that RECK gene expression was correlated with occurrence/progression of RECK. We thus investigated the expression profiles of RECK, MMP-2 and MMP9 on GCTs patients, in order to illustrate its correlation with GCTS recurrence, and to guide the clinical treatment by establishing more sensitive indexes, as well as lowering GCTS recurrent rate.

### Materials and methods

#### *Patients*

A cohort of 42 GCTS patients (19 males and 23 females, aging between 13 and 66 years old) from the department of orthopedics from January 2004 to December 2009 were recruited in The 309th Hospital of Chinese PLA. Inclusive criteria: (1) With confirmed GCTS diagnosis; (2) Primary tumor patients; (3) No metastasis at the primary diagnosis; (4) Having undergone surgical resection; (5) Full medical history; and (6) Available post-operative samples for immunohistochemical (IHC) examinations.

In an Enneking staging system, there were 10 stage I, 13 of stage II and 19 of stage III patients. By Campanacci grading under X-ray,

there were 11 of grade I, 14 patients at grade II, and 17 of grade III patients. This study has been pre-approved by the ethical committee of The 309th Hospital of Chinese PLA and has obtained consents from all participants. With permission, tumor tissues were collected during the surgery.

#### *Post-operative follow-ups*

Follow-ups persisted for 12~60 months until the endpoint or patient death. All cases have successfully finished the follow-up. There were 17 cases of recurrence, 2 patients having pulmonary metastasis, and 23 patients without recurrence.

#### *IHC staining*

All tumor samples were fixed in 10% formalin and were embedded in paraffin. After sectioning into consecutive slices with 4  $\mu$ m thickness, IHC staining was performed using SP method following manual instruction of test kits including MMP-2, MMP-9 and SP (Maixin, Fuzhou, China). Parallel negative controls were also performed using PBS instead. Positive staining was determined as brown-yellow granules in the cytoplasm. Five high-magnification fields were randomly selected to count 100 cells. The staining grade was measured by both percentage score and intensity score as previously documented [18]. Percentage score was scaled as 1, 2 or 3 for those fields with less than 10%, 10%~50% and more than 50% of positive cells, respectively. Staining intensity score was given from 0 to 3 including negative staining, light yellow, moderate, and dark-brown color. The total score (= percentage score + staining intensity score) was divided as negative (equal or less than 4) or positive (larger than 4).

#### *Statistical analysis*

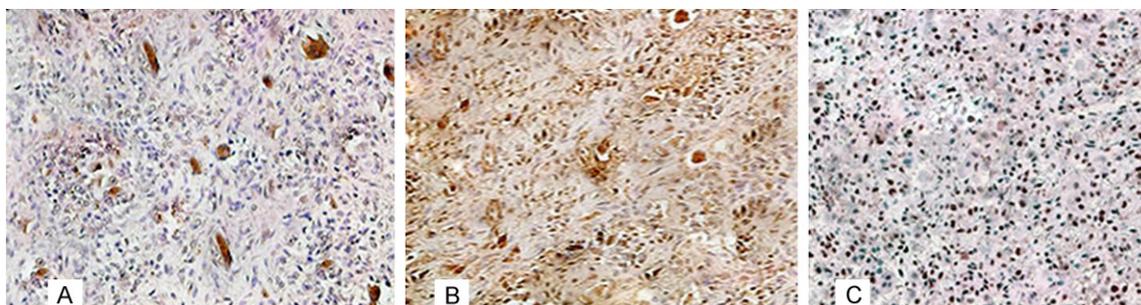
SPSS 13.0 software was used to process all collected data. Enumeration data were compared by chi-square test. Logistic regression analysis was used for multi-factor. The significance level was determined as 0.05.

### Results

#### *Gene expression and clinical features*

IHC staining images of RECK, MMP-2 and MMP-9 genes were shown in **Figure 1**. Among

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**Figure 1.** IHC staining image of MMP-2 (left), MMP-9 (middle) and RECK (right).

**Table 1.** Clinical features and expression levels of RECK, MMP-2 and MMP-9 gene

Clinical index	N	RECK Positive	P value	MMP-2 Positive	P value	MMP-9 Positive	P value
<b>Sex</b>							
Male	19	10	0.55	10	0.55	13	0.11
Female	23	10		10		10	
<b>Age</b>							
<30 years	19	7	0.37	10	0.55	11	0.71
≥30 years	23	13		10		12	
<b>Enneking stage</b>							
Stage I & II	23	15	0.01*	7	0.01*	8	<0.01*
Stage III	19	5		13		15	
<b>Campanacci grade</b>							
Grade I & II	25	16	<0.01*	8	0.01*	10	0.02*
Grade III	17	4		12		13	

Note: \*, P<0.05.

**Table 2.** Clinical features and GCTS recurrence/metastasis

Clinical index	N	No recurrent group	Recurrence-me-tastasis group	$\chi^2$ value	P value
<b>Sex</b>					
Male	19	9	10	0.77	0.38
Female	23	14	9		
<b>Age</b>					
<30 years	19	10	9	0.06	0.80
≥30 years	23	13	10		
<b>Enneking stage</b>					
Stage I & II	23	17	6	7.53	<0.01*
Stage III	19	6	13		
<b>Campanacci grade</b>					
Grade I & II	25	17	8	4.37	0.04*
Grade III	17	6	11		

Note: \*, P<0.05.

all 42 patients, there were 20 positive cases for RECK (47.62%), 20 MMP-2 positive cases (47.62%) and 23 patients with MMP-9 positive

expression (54.76%). Chi-square revealed no significant relation between patients' age and sex with gene expression level. Enneking stage and Campanacci grade, however, is significantly correlated with gene expression levels. (P<0.05, **Table 1**).

### *GCTS recurrence and clinical features*

In all 42 patients undergone follow-ups, there were 17 recurrent cases and 2 cases of pulmonary metastasis, both of which were classified in recurrent group (N=19). Chi-square analysis revealed no significant relation between patients' age and sex with recurrence/metastasis. Enneking

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**Table 3.** Gene expression and GCTS recurrence/metastasis

	N	No recurrent group	Recurrence-me-tastasis group	$\chi^2$ value	P value
RECK					
Positive	20	5	15	6.31	0.01
Negative	22	14	8		
MMP-2					
Positive	20	13	7	6.02	0.01
Negative	22	6	16		
MMP-9					
Positive	23	15	8	8.19	<0.01
Negative	19	4	15		

**Table 4.** Multi-variate analysis of GCTS recurrence

	OR	95% CI	P value
RECK	0.17	0.04~0.83	0.028*
MMP-2	4.05	0.97~16.91	0.055
MMP-9	5.19	1.12~24.04	0.035*

Note: \*, P<0.05.

stage and Campanacci grade, however, is significantly correlated with recurrence/metastasis. (P<0.05, **Table 2**).

### Gene expression and GCTS recurrence/metastasis

Within all 20 patients with RECK-positive expression, there were 3 cases of recurrence or metastasis. Such ratio was 14/20 for MMP-2 and 16/23 for MMP-9. Chi-square analysis revealed the correlation between GCTS recurrence/metastasis with expression of RECK, MMP-2 or MMP-9 gene expression (P<0.05, **Table 3**).

### Multi-factor analysis of GCTS recurrence/metastasis

To rule out possible interfering effects on GCTS recurrence, we employed Logistic regression analysis using sex, age, Enneking stage, Campanacci grade, and expressions of RECK, MMP-2 and MMP-9 as independent variables. Results showed the decreased risk of GCTS recurrence/metastasis by up-regulating RECK, which thus plays as one protective factor (OR=0.17, P=0.028, **Table 4**). MMP-9 overexpression, however, elevated risk of recurrence (OR=5.19, P=0.035, **Table 4**). No significant

correlation existed between MMP-2 and GCTS recurrence (OR=4.05, P=0.055, **Table 4**).

### Discussion

Surgical resection is still the major approach for treating GCTB, whose major postoperative complication is recurrence [1]. GCTB can be classified as grade I to grade III based on Jaffe system according to the

ratio of mononuclear mesenchymal cells against osteoclast cells, and heteromorphism of mononuclear mesenchymal cells [19]. Clinical follow-ups, however, revealed the inability of Jaffe grade system to predict the recurrence of GCTB after surgery. Campanacci further classified GCTB into three grades based on X-ray change [20]. Some studies have higher recurrence rate of GCTB in those patients with advanced Campanacci grades [21, 22]. This study obtained similar results as Campanacci grade III patients had higher recurrent rate compared to grade I and II patients. Some studies, however, rejected the significant correlation between Campanacci grade and GCTB recurrence [23, 24]. This issue thus requires more systematic illustration. Enneking also classified GCTB into three stages based on clinical features, X-ray and pathology [25]. Some scholars suggested elevated risks of pulmonary metastasis in those GCTB patients at stage III [26], a result that occurred also in our study, suggesting the close correlation between Enneking stage and GCTB prognosis.

Current study about MMPs and GCTB mainly focused on the correlation between MMPs expression and GCTB occurrence, but lacked the potency of MMPs as prognostic indicators. This study thus performed IHC staining on GCTS tumor tissues for quantifying MMP-2 and MMP-9 expression, in parallel with follow-ups to elucidate patients' prognosis. Results showed MMP-9 but not MMP-2 as one risk factor for GCTS recurrence. In addition, RECK gene expression was also negatively correlated with MMPs expression. We thus tested the expression of RECK in tumor tissues and found RECK gene as one protective factor preventing GCTS

recurrence or metastasis. Previous report has mentioned higher MMP-9 expression in recurrent GCTB tissues [15], indicating the correlation between MMP-9 overexpression and GCTB recurrence. One recent study has also suggested the close correlation between MMP-9 and GCTB prognosis [27], in addition to the involvement of MMP-2 in GCTB recurrence. This is inconsistent with our results perhaps due to the including of malignancy transformation and different population selected in two studies. No direct study regarding RECK and GCTB has been reported. RECK, although RECK has been suggested to be involved in the progression of other tumors. For example, RECK can inhibit the invasion/metastasis of neuroblastoma and hepatoblastoma via inhibiting MMPs expression [28]. Other study has revealed the modulation on tumor progression by RECK via modulating the expression of vascular endothelial growth factor to regulate tumor angiogenesis, making RECK as one important prognostic indicator for tumors [29].

In summary, this study for the first time revealed the protective function of RECK against GCTS recurrence or metastasis, in addition to the correlation between MMP-9 and GCTS recurrence. Our results provide new insights regarding the evaluation of GCTS prognosis. However, due to the limited sample size, the promotion of RECK and MMP-9 as effective clinical indexes for GCTS needs further validation by multi-centered, large sample study.

#### Disclosure of conflict of interest

None.

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