

Study on the expression of matrix metalloproteinase-2 mRNA in human gastric cancer

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INTRODUCTION

During tumor invasion and metastasis, malignant cells in primary site acquire the ability to degrade extracellular matrix (ECM) and penetrate tissue barriers. Among the proteolytic enzymes which degrade ECM, matrix metalloproteinase (MMP) is one of the important ones. MMP₂ (72kDa type IV collagenase) is a member of the MMPs gene family which degrades the macromolecules of connective tissue and ECM, such as collagen, proteoglycans, laminin and fibronectin. Thus MMP₂ is believed to play an important role in tumor invasion and metastasis. Several immunohistochemical studies have shown that MMP₂ mRNA is overexpressed in gastric cancer and related to the clinical stage of cancer^[1,2]. However, samples were not enough and lack completeness in these studies. Reverse transcriptase-polymerase chain reaction (RT-PCR) was used in our study to examine the expression of MMP₂ mRNA in tumor and normal tissues adjacent to human gastric cancer.

MATERIALS AND METHODS

Samples

The samples, including tumour tissue and tumour-adjacent normal tissue (2 cm and 5 cm or over 5 cm distance from the tumour) came from twenty gastric cancer patients at the surgery department of our hospital and People's Hospital of Yun-He County, Zhejiang. Gastric tissues from five benign ulcer patients after partial gastrectomy were used as controls. The histological diagnosis was made by the

pathologists of these two hospitals. Patients' clinical features are shown in Table 1. Among these cancer patients, five and fifteen were cases of early and advanced cancer respectively. All samples were quenched in liquid nitrogen immediately after operation and were then stored at -70°C until used for the study.

Reagents and primer synthesis

The dNTP, RNasin and MMLV reverse transcriptase and Taq DNA were provided by Stratagene, La Jolla, CA, U.S.A. MMP₂ primer pair was synthesized by Shanghai Cell Research Institute, Chinese Academy of Sciences^[2] and its sense and anti-sense were 5'-ACAAAGAGTGGCAGTGCAA-3' and 5'-CACGAGCAAAGGCATCATCC-3' respectively. The expected size of MMP-2 product was 302bp.

RT-PCR analysis

Total RNA was extracted from frozen tissues by cesium chloride purifying method. A total amount of 20 µL reaction solution contained 5 µg RNA sample tissue, 1 mmol/L dNTP, 10 U RNasin, 100 mmol/L Tris-HCl pH 8.4, 50 mmol/L KCl, 2.5 mmol/L MgCl₂, 100 mg/mL BSA, 100 pmol random six-polyoligo-nucleotide and 100 U MMLV reverse transcriptase. The reverse transcription condition was 37 °C for 1 h, and 95 °C for 5 min. Twenty µL cDNA reverse transcriptase product was put in PCR reaction solution containing 100 mmol/L Tris-HCl, pH 8.4, 50 mmol/L KCl, 2.5 mmol/L MgCl₂, 100 mg/mL BSA, 30 pmol sense and anti-sense primers, and then 2 U Taq DNA polymerase was added in the solution. The PCR amplification condition was: denatured at 95 °C for 1 min, annealed at 65 °C for 1 min and extended at 72 °C for 1 min. The number of cycles was 35. 10 µL DNA, amplification product was subjected to electrophoresis in 4% agarose gel, stained with ethidium bromide and observed under ultraviolet light. The photographs of PCR results were used to measure the level of optical density (OD) of MMP-2 cDNA bands with densitometry (Backman CD 2000).

Statistical analysis

The significance of differences in expression rates and OD levels among groups was determined by χ^2 test and Student's *t* test respectively.

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RESULTS

Expression of MMP-2 mRNA in tumor and tumor-adjacent tissues (Figure 1)

In 20 cases of gastric cancer, MMP₂ mRNA was expressed in 13 tumor tissues, 11 in 2 cm and 6 in ≥ 5 cm adjacent tissues respectively (Table 1). The positive rate of MMP₂ mRNA expression in tumor tissues was significantly higher than that in ≥ 5 cm adjacent tissues ($P < 0.05$). There was no positive expression of MMP₂ mRNA in the 5 samples of the control group.

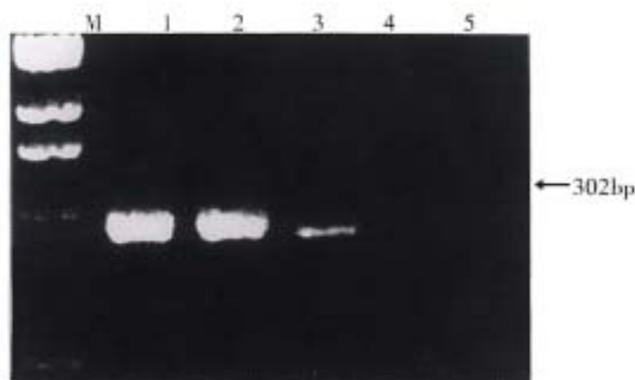


Figure 1 Expression of MMP-2 mRNA in case No.3
Lane M: Marker; Lane 1, 2: tumor tissues; Lane 3-5: 2cm, ≥ 5 cm adjacent tissues and normal gastric tissue of one control.

Table 1 Clinical features of cases and expression of MMP-2 mRNA

No.	Age/sex	Location	Histology	Clinical stage ^b	MMP2 mRNA expression		
					T	2cm	≥ 5 cm
1 ^a	64/M	Pylorus	Poor	T1N0M0(I)	-	-	-
2	48/M	Body	Poor	T4N0M0(IV)	+	+	-
3	72/F	Lesser curve	Poor	T4N2M1(IV)	+	+	-
4	75/F	Lesser curve	Poor	T3N0M0(II)	+	+	-
5	71/M	Body	Poor	T2N1M0(IVA)	+	+	-
6	63/M	Lesser curve	Well	T4N2M0(IV)	-	-	+
7 ^a	73/F	Lesser curve	Poor	T1N1M0(IIIA)	+	-	-
8 ^a	46/M	Lesser curve	Poor	T1N0M0(I)	-	-	-
9	60/F	Body	Poor	T3N0M0(II)	+	-	-
10	57/M	Pylorus	Poor	T3N1M1(IV)	+	+	+
11	53/M	Body	Poor	T4N1M0(IV)	+	+	-
12 ^a	72/F	Antrum	Poor	T1N0M0(I)	-	-	-
13	59/M	Body	Poor	T3N2M0(IIIB)	+	-	-
14 ^a	38/M	Antrum	Well	T1N0M0(I)	-	-	-
15	66/F	Cardia	Poor	T3N1M0(IIIA)	+	+	+
16	55/M	Antrum	Well	T2N0M0(II)	-	-	-
17	74/M	Body	Poor	T4N2M1(IV)	+	+	+
18	67/M	Cardia	Well	T3N2M0(IVB)	-	-	-
19	67/F	Body	Poor	T4N2M1(IV)	+	+	+
20	56/M	Body	Poor	T4N2M1(IV)	+	+	+

T: Tumor tissues; 2cm: 2cm adjacent tissue from tumor; ≥ 5 cm: 5cm or over 5cm adjacent tissue from tumor. ^a: early gastric cancer. Poor: poorly differentiated adenocarcinoma; Well: well differentiated adenocarcinoma. ^b: According to the American Joint Commission Staging of Gastric Cancer.

The OD levels of MMP-2 mRNA in tumor and tumor-adjacent tissues (Figure 2)

The OD of MMP-2 detected cDNA signals ranged

from 1.10 to 19.23 (mean 5.38 ± 0.98) in tumor tissues, 0.86 to 4.17 (mean 2.41 ± 0.30) in 2 cm and 0.78 to 3.80 (mean 1.88 ± 0.22) in ≥ 5 cm adjacent tissues respectively. There was significant difference in OD levels between tumor tissues group and 2 cm or ≥ 5 cm tumoradjacent tissues one ($P < 0.01$), and no significant difference in OD levels between the two adjacent tissues groups.

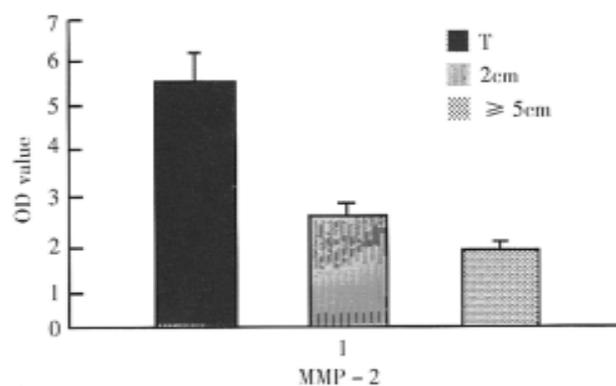


Figure 2 OD of MMP-2 cDNA signals in tumor tissues and tumor-adjacent tissues (2cm and ≥ 5 cm). As compared with tumor tissues: 2cm: $P < 0.01$; ≥ 5 cm: $P < 0.01$.

The OD levels of MMP-2 mRNA in early and advanced cancer (Table 2)

The MMP-2 cDNA signals in tumour and 2cm adjacent tissues of 15 advanced cancer were significantly higher than those in the two corresponding tissues of 5 early cancer respectively ($P < 0.05$). In the ≥ 5 cm adjacent tissues, there was no-significant difference in the signals between the advanced and early cancer.

Table 2 OD of MMP-2 cDNA signals in early and advanced cancer ($\bar{x} \pm s_x$)

	T	2cm	≥ 5 cm
Early cancer (n=5)	1.63 ± 0.42	1.07 ± 0.29	1.50 ± 0.22
Advanced cancer (n=15)	6.63 ± 1.12^a	2.80 ± 0.34^a	1.94 ± 0.29

^a $P < 0.05$, vs early cancer.

DISCUSSION

In 20 gastric cancer cases, 13, 11 and 6 cases positively expressed MMP-2mRNA in tumor, 2 cm adjacent and ≥ 5 cm adjacent tissues respectively. The cases with MMP-2 mRNA expression in tumor tissues and their corresponding tumor adjacent normal tissues (2 cm and/or ≥ 5 cm) were almost poorly differentiated adenocarcinoma and in higher clinical stage. MMP-2 mRNA was seldom expressed in tumor and tumor adjacent tissues of well differentiated or early carcinoma. These results

showed that proliferative and invasive gastric cancer cells had higher MMP-2 secretion. In ultrastructural study, MMP-2 mRNA was expressed markedly in cancer cells with rich false feet and rapid movement in culture, but insignificantly or with few false feet in cancer cells from unmetastatic and uninvase gastric cancerous tissues, indicating that MMP-2 secretion was correlated with the invasion and metastasis of gastric cancer^[3]. In addition, some immunohistochemistry studies have shown that the positive rate of staining cells for MMP-2 protein was consistently higher in poorly differentiated and diffuse gastric carcinoma than that in well differentiated and early gastric carcinoma^[1,4]. The results of these studies were very similar to those of MMP-2 mRNA expression in our study, indicating that the increased MMP-2 positive staining was related to the overexpression of MMP-2 mRNA. This is probably due to increase in MMP-2 transcriptional activity and MMP-2 protein products in the cell proliferative process of gastric cancer which is often accompanied with the acceleration of cancer invasion and metastasis.

In the current study, although the levels of MMP-2 cDNA signals in tumor-adjacent tissues (2 cm and/or ≥ 5 cm) were lower than those in tumor tissues, MMP-2 mRNA was overexpressed in the tumor adjacent tissues in certain extent, suggesting that both gastric cancer cells and adjacent mesenchymal cells, including fibrocyte, endothelium cell, macrophage and lymphocyte have the ability to secrete MMP-2. There may be information exchange between the cancer cells and these mesenchymal cells through the dissolvable intercellular substance and membrane cement factor, and such information exchange may regulate the production of MMP-2. This may be very

important in elucidating the mechanism of invasion and metastasis of cancer cells^[5,6]. In our case No. 6, MMP-2 mRNA was detected only in tumor-adjacent tissues, but not in tumor tissue. The reason for this is unclear. The discrepancy may be due to the necrotic tumor tissue. Overexpression of MMP-2 mRNA only in tumour adjacent tissues also indicates the malignant degree of cancer is rather high.

The prognosis of early gastric cancer is better than that of advanced cancer. Our study showed that the levels of MMP-2 cDNA signal in advanced cancer tissues (tumor and 2 cm tumor-adjacent tissues) were significantly higher. This suggests MMP-2 may play a role in gastric cancer invasion and metastatic progression, and the overexpression may be associated with poor prognosis. Further study on relationship between the expression of MMP-2 mRNA and the survival rate of gastric cancer is being carried out.

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