

# 胃癌耐药相关抗体 MGr1 筛选文库获得的基因 MGr1 - Ag1 在胃癌组织中表达的研究

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## Expression of cDNA fragment encoding MGr1-Ag1 detected by MDR related antibody MGr1 in gastric cancer

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## Abstract

**AIM:** To analyze the distribution and significance of the possible cDNA fragments encoding MGr1-Ag1 in human gastric cancer.

**METHODS:** We investigated MGr1-Ag1 expression in gastric cancer using *in situ* hybridization (ISH) techniques in 42 cases of gastric carcinomas in cryostatic sections collected from Xijing Hospital.

**RESULTS:** MGr1-Ag1 mRNA was positive in cytoplasm of gastric glandular epithelia with intensive staining in 50.00% (21/42) cases with gastric cancer, and 71.42% (30/42) cases in paracancerous gastric tissues. Eleven of 16 (68.75%) cases were well differentiated, 8/14 (57.14%) cases were moderately differentiated and 1/10 (10.00%) cases were poorly differentiated. The positive rate of MGr1-Ag1 mRNA in gastric cancer was significantly different from that in paracancerous gastric tissues (71.42%). There was no significant difference in positive rate of MGr1-Ag1 mRNA among normal, paracancerous gastric tissues, well and moderately differentiated tissues, but the positive rate in the poorly

differentiated tissues was low, being significantly different from that in well or moderately differentiated tissues.

**CONCLUSION:** MGr1-Ag1 mRNA is mainly localized in epithelial cells, and the level of its expression is correlated with the tumor differentiation.

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## 摘要

**目的:** 为研究肿瘤的多药耐药现象, 应用胃癌耐药相关的单克隆抗体 MGr1 筛选肿瘤耐药细胞表达文库获得的一个稳定阳性克隆的 cDNA 片段 MGr1-Ag1, 检测 MGr1-Ag1 在胃癌组织中的分布及表达水平。

**方法:** 利用原位杂交技术, 对 MGr1-Ag1 mRNA 在胃癌组织中的分布进行了检测, 以了解其在胃癌中的表达情况及意义。

**结果:** MGr1-Ag1 mRNA 定位在胃腺体细胞的胞质中, 为蓝色小颗粒, 呈弥漫性分布。MGr1-Ag1 mRNA 在 SGC7901/VCR 阳性信号明显高于 SGC7901, 胃癌与癌旁组织相比, MGr1-Ag1 mRNA 的阳性表达率均有显著差异 ( $P < 0.05$ )。MGr1-Ag1 mRNA 在胃癌 (50.00%)、癌旁组织 (71.42%)、高分化腺癌 (68.75%) 和中分化腺癌组织 (57.14%) 中的阳性表达率无显著差别, 但高分化组和中分化组都与低分化腺癌组织 (10.00%) 差别显著 ( $P < 0.05$ )。此外, MGr1-Ag1 mRNA 还分布于小肠及大肠腺体内, 因例数偏少, 未做统计。

**结论:** MGr1-Ag1 与胃癌的分化程度有关, 随着组织的分化程度增高, MGr1-Ag1 mRNA 的阳性表达率增加。

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## 0 引言

肿瘤多药耐药现象 (multidrug resistance, MDR) 是肿瘤治疗的热点和难点。但目前的已知机制并不能充分解释肿瘤多药耐药性的发生, 而与耐药有关的分子和机制不断的被提出<sup>[1-9]</sup>。为扩充肿瘤细胞产生 MDR 的机制, 以耐

长春新碱人胃癌细胞 SGC7901/VCR 为免疫原制备了一株与胃癌耐药相关的单克隆抗体 MGr1<sup>[10]</sup>, 利用 MGr1 筛选肿瘤耐药细胞表位文库, 获得了一个稳定阳性克隆 MGr1 - Ag1, 携带长约 262 bp 的 cDNA 片段, 但其中 218 bp 的序列与定位于人染色体 7p22-p21 的部分基因组序列完全相同<sup>[11]</sup>. 为分析 MGr1 - Ag1 片段可能的意义及功能, 利用 mRNA 分子原位杂交等技术, 对 MGr1 抗体筛选的 cDNA 片段 MGr1 - Ag1 在胃癌的表达情况进行了检测.

## 1 材料和方法

1.1 材料 42 例胃癌组织及其癌旁组织取自第四军医大学西京医院于 1999-08/1999-12 间行胃癌切除术的患者, 正常胃黏膜标本 25 例取自胃镜活检. 所有标本均于离体后 30 min 内完成取材并置于液氮中保存. 制备 8  $\mu$ m 冰冻切片, 40 g/L 多聚甲醛固定、干燥, -70  $^{\circ}$ C 冰箱保存. 所有病理标本均经病理科两位以上医师诊断. 胃癌细胞 SGC7901 和胃癌耐长春新碱 SGC7901/VCR 为我所保存的细胞. 胰酶和小牛血清为 Sigma 公司产品. RPMI1640 和 BSA 为 Gibco BRL 产品. 多聚赖氨酸胶购于武汉博士德公司. 限制内切酶、DNA 纯化试剂盒及其他常规试剂购于华美生物工程公司. DIG DNA Labeling and Detection Kit 购于德国 Boehringer Mannheim 公司.

### 1.2 方法

1.2.1 组织切片准备 培养细胞在多聚赖氨酸处理的盖玻片上进行培养, 培养液为 80 ml/L 小牛血清 RPMI1640. 细胞长好后 10 mol/L PBS(pH7.4) 洗 3  $\times$  2 min. 细胞爬片和冰冻切片在 40 g/L 多聚甲醛室温固定 10 min. 10 mol/L PBS(pH7.4) 洗 3  $\times$  2 min. 干燥后冰冻保存.

1.2.2 探针的制备和标记 MGr1 - Ag1 cDNA 片段克隆在 pSPORT1 载体 Sal 和 Not 位点之间, 两侧有 BamH 和 Kpn 的位点. BamH 和 Kpn 消化质粒. 将酶切产物进行 10 g/L 琼脂糖凝胶电泳, 得到约 0.3 kb cDNA 片段. 试剂盒纯化回收 cDNA 片段并定量. 取 1-3  $\mu$ g cDNA 煮沸 10 min, 冰水中冷却. 用地高辛 DNA 标记试剂盒进行标记和纯化.

1.2.3 mRNA 原位杂交 切片于 70  $\times$  2  $\times$  SSC 预热 15 min, PBS 洗切片 2 min. 切片上滴加 0.5 mol/L TBS (pH7.2-7.6) 稀释的 1:1000 Proteinase K, 37  $^{\circ}$ C 消化 2-15 min, 暴露 mRNA 核酸片段. 置 4  $^{\circ}$ C 预冷的 40 g/L 多聚甲醛中平衡 5 min, PBS 洗 2  $\times$  2 min. 2  $\times$  SSC 洗 2  $\times$  2 min. 逐级乙醇脱水. 每张切片上加适量预杂交液, 湿盒 70  $^{\circ}$ C 孵育 8 min, 37  $^{\circ}$ C 孵育 1 h. 将标记的 cDNA 探针放入 95-100  $^{\circ}$ C 水中煮沸 5-10 min 后, 迅速冷却 3 min. 按 0.5-1.0  $\mu$ g/ml 在杂交液加入标记探针. 37  $^{\circ}$ C 湿盒中杂交 16-24 h 后, 室温 2  $\times$  SSC 洗 20 min, 1  $\times$  SSC 洗 20 min, 43  $^{\circ}$ C 0.5  $\times$  SSC 洗 20 min, 室温 0.5  $\times$  SSC 洗 20 min. Buffer 室温振洗 1  $\times$  5 min, 1  $\times$  30 min, Buffer 作用 5 min,

加抗地高辛抗体 -AP 复合物(1:500), 37  $^{\circ}$ C 2h. Buffer 浸洗 2  $\times$  15 min, 振洗 1  $\times$  5 min, Buffer 浸洗 5 min, 加显色液显色 0.5-12 h. 染色理想时终止反应. 上述所用试剂和材料均经去 RNA 酶处理.

1.2.4 结果判定 显微镜下选取染色结果较理想部位, 于高倍镜下, 每视野统计 200 个细胞数. 阳性细胞数比率 < 10 % 或无阳性着色为阴性(-); 阳性细胞数比率 < 50 % 为弱阳性(+); 阳性细胞数比率 > 50 % 为阳性(++); 阳性细胞数比率 > 75 % 为强阳性(+++).

统计学处理 采用 SPSS9.0 统计软件进行  $\chi^2$  检验, 以  $P < 0.05$  确定为有统计学意义.

## 2 结果

2.1 MGr1 - Ag1 cDNA 片段的鉴定 用 BamH 和 Kpn 酶消化 pSPORT1 质粒, 得到一长约 0.3 kb 片段, 携带 262 bp 的 MGr1 - Ag1 cDNA 片段.

2.2 MGr1 - Ag1 mRNA 在胃癌组织及细胞系中的表达 MGr1 - Ag1 mRNA 在胃癌耐药细胞系 SGC7901/VCR 中的表达强度明显高于 SGC7901. MGr1 - Ag1 mRNA 定位在 SGC7901/VCR 和 SGC7901 细胞的胞质中, 为蓝色小颗粒, 呈弥漫性分布(图 1、图 2). 在胃组织中, MGr1 - Ag1 的 mRNA 定位在胃腺体细胞, 在胃癌组织中阳性率为 50.00 %; 在癌旁胃黏膜中的阳性表达率为 71.42 %; 在正常胃黏膜的阳性表达率为 68.00 %. 胃癌与癌旁组织相比, MGr1 - Ag1 mRNA 的阳性表达率有显著差异( $P < 0.05$ )(表 1).

表 1 在胃癌、癌旁组织和正常胃黏膜中的表达

Group	MGr1 - Ag1				T	R(%)
	-	+	++	+++		
GC 42	21	9	10	2	21	50.00
PC 42	12	12	15	3	30	71.42
NT 25	8	5	9	3	17	68.00

NT: normal gastric tissue; GC: gastric cancer; PC: paracancerous gastric mucosa  
Pearson Chi-Square Tests:  $\chi^2 = 4.043$ ;  $\eta^2 = 1$ ;  $P = 0.044 < 0.05$ ; GC vs PC.

表 2 MGr1 - Ag1 mRNA 的表达和胃癌分型

Group	Grade of intensity				T	R(%)
	-	+	++	+++		
WD 16	5	5	5	1	11	68.75
MD 14	6	3	4	1	8	57.14
PD 10	9	1	0	0	1	10.00
MC 2	1	0	1	0	1	50.00

MC: mucous carcinoma including mucous adenocarcinoma and signet ring cell carcinoma

WD: well differentiated adenocarcinomas

MD: moderately differentiated adenocarcinoma

PD: poorly differentiated adenocarcinoma

Fisher's Exact Test:  $P = 0.005 < 0.05$  WD vs PD;  $P = 0.033 < 0.05$  MD vs PD

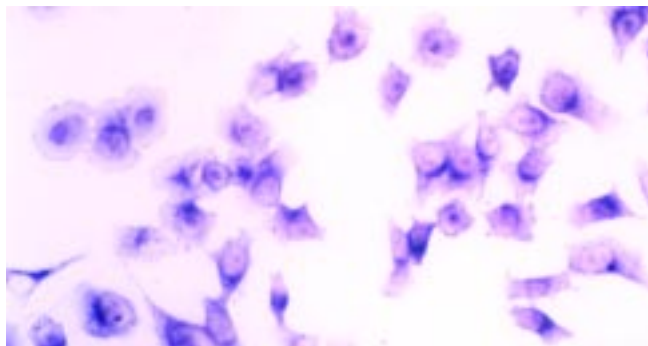


图1 原位杂交显示, MGr1-Ag1 mRNA 在人胃癌耐长春新碱细胞 SGC7901/VCR 胞质中呈强阳性表达 NBT/BCIP 显色  $\times 400$ .

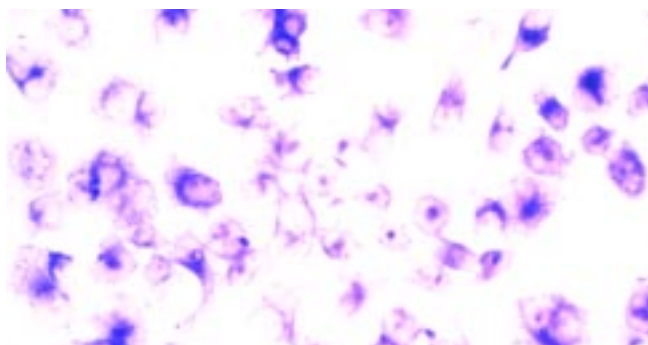


图2 原位杂交显示, MGr1-Ag1 mRNA 在人亲本胃癌细胞 SGC7901 胞质中呈阳性表达, 其阳性信号明显弱于 SGC7901/VCR 细胞  $\times 400$ .

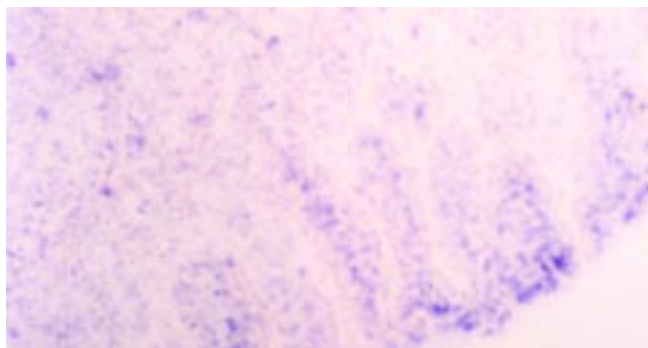


图3 原位杂交显示, 正常胃黏膜中 MGr1-Ag1 mRNA 呈阳性表达, 阳性信号分布于上皮细胞的胞质中  $\times 200$ .

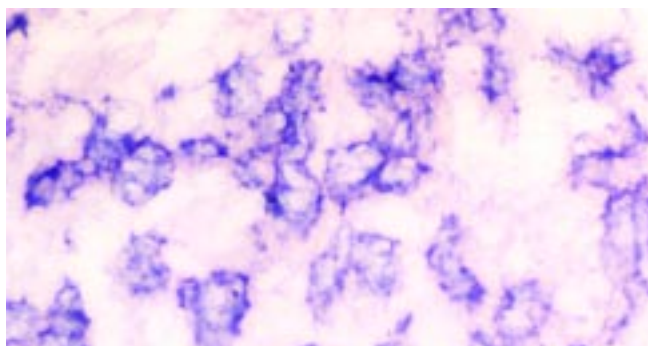


图4 原位杂交显示, 高分化腺癌中 MGr1-Ag1 mRNA 呈阳性表达, 阳性信号分布于腺体细胞的胞质中  $\times 400$ .

### 2.3 MGr1-Ag1 mRNA 的表达与胃癌的关系 MGr1-Ag1

mRNA 的表达水平在癌旁组织、正常胃黏膜组织、高分化腺癌(68.75%)和中分化腺癌(57.14%)的阳性表达率无显著差别(图3、图4),但高分化组和中分化组与低分化腺癌(10.00%)的阳性表达率差别显著( $P < 0.05$ )(表2).此外, MGr1-Ag1 mRNA 还分布于小肠及大肠腺体内, 因例数偏少,未做统计.

### 3 讨论

胃癌对化疗药物的反应性通常很差<sup>[12,13]</sup>,肿瘤细胞可通过各种途径产生对化疗的抵抗性,其机制复杂多样.肿瘤细胞能通过 P-糖蛋白和 MRP 增加药物的外排<sup>[14-18]</sup>;位于胞质的 MRP1 和肺耐药蛋白能改变细胞内药物的分布<sup>[19-23]</sup>;GST 通过催化 GSH 与抗肿瘤药物结合增强细胞解毒功能<sup>[24-26]</sup>;Topo 发生改变,可引起以其为靶点的药物诱导的 DNA 稳定断裂复合物形成减少, DNA 双链断裂减少,细胞产生耐药性<sup>[27,28]</sup>.此外一些代谢酶类及涉及凋亡和信号转导的分子也参与耐药性的形成<sup>[29-33]</sup>.

我所采用大剂量间歇诱导法以长春新碱诱导人胃癌细胞 SGC7901,获得了人胃癌耐药细胞株 SGC7901/VCR,该细胞具有 MDR 表型,耐受 5-氟尿嘧啶、顺铂、柔红霉素等多种结构不同的药物.以 SGC7901/VCR 为免疫原,采用杂交瘤技术制备了胃癌细胞耐药相关性单克隆抗体,命名为 MGr1<sup>[10]</sup>.MGr1 抗原在 SGC7901/VCR 的胞膜上和胞质中表达,且显著高于亲本细胞 SGC7901.Western blot 结果表明 MGr1 抗原分子量大约为 40 KDa.MGr1 抗体能部分逆转 SGC7901/VCR 对阿霉素、长春新碱和 5-氟尿嘧啶的耐药性,显著增加 SGC7901/VCR 细胞内阿霉素的滞留.说明 MGr1 是肿瘤耐药相关性抗体.为了克隆 MGr1 抗原的编码基因,研究 MGr1 抗原的分子结构和功能,利用 MGr1 抗体筛选肿瘤耐药细胞表达文库,获得了一个稳定阳性克隆 MGr1-Ag1.MGr1-Ag1 mRNA 的表达水平在癌旁组织、正常胃黏膜组织、高分化腺癌和中分化腺癌的阳性表达率无显著差别,但与低分化腺癌的阳性表达率差别显著 ( $P < 0.05$ ),提示 MGr1-Ag1 与胃癌的分化程度有关,随着组织的分化程度增高, MGr1-Ag1 mRNA 的阳性表达率也增加,该结果与利用免疫组化和 MGr1 抗体检测 MGr1-Ag 在胃癌组织中的表达分布情况基本一致.

我们将继续用 MGr1 抗体筛选肿瘤 cDNA 文库,采用 RACE 技术克隆 MGr1-Ag1 全长 cDNA,并通过基因转染分析 MGr1-Ag1 在胃癌 MDR 中的作用,从而判断 MGr1-Ag1 是否与肿瘤细胞耐药现象有关,并最终获得 MGr1-Ag 的编码基因,为充实胃癌 MDR 机制和从基因水平寻找新的耐药逆转措施奠定良好的基础.

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