

CagA⁺幽门螺杆菌胃癌前及癌变中NF-κB的表达增强

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广东省科技厅资助项目, No. 2003C30307
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收稿日期: 2005-06-28 接受日期: 2005-07-15

Increased expression of NF-κB p65 in CagA⁺ *H pylori*-related gastric precancerous lesions and carcinoma

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Supported by Science and Technology Department of Guangdong Province, No.2003C30307

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Received: 2005-06-28 Accepted: 2005-07-15

Abstract

AIM: To determine the expression of nuclear factor kappa B (NF-κB) in human gastric precancerous lesions and carcinoma and its correlation with CagA⁺ *H pylori* infection.

METHODS: The expression of NF-κB p65 was detected by immunohistochemistry (SABC assay), and *H pylori* were examined using ¹⁴C-breath test, rapid urease test and Warthin-Starry staining in patients with chronic superficial gastritis (CSG: n = 34), intestinal metaplasia (IM: n = 31), atypical dysplasia (AD: n = 34) and gastric cancer (GC: n = 55). Serum CagA IgG antibody was detected by dot immunogold filtration assay. The correlations of NF-κB p65 expression with CagA⁺ *H pylori* infection as well as the histological types, clinicopathological stages and lymph node metastasis were analyzed.

RESULTS: The expression of NF-κB p65 in CSG, IM, AD, GC was 15.0%, 41.9%, 64.7%, and 78.2%, respectively, and there were significant differences between them ($\chi^2 = 43.98, P < 0.01$). The rates of *H pylori* infection were 70.0%, 67.7%, 73.5%, and 54.5%, respectively, and there were no significant differences between them ($P > 0.05$). The percentage of CagA⁺ *H pylori* infection were 53.6%, 61.9%, 68.0%, and 73.3%, respectively, and there were no significant differences ($P > 0.05$). In IM, the positive rate of NF-κB p65 expression in *H pylori* or CagA⁺ *H pylori* positive patients were significantly higher than that in patients without *H pylori* infection or with CagA⁻ *H pylori* infection (57.1%, 76.9% vs 10%, 25.0%, $P < 0.05$). In GC patients, the positive expression of NF-κB p65 was correlation with the T stages ($\chi^2 = 5.91, P < 0.05$) and lymph node metastasis ($\chi^2 = 7.47, P < 0.01$), but not with the pathohistological types ($P > 0.05$).

CONCLUSION: NF-κB is constitutively activated in human gastric precancerous lesions and carcinoma tissue and correlates with tumor progression. The early activation may be related to CagA⁺ *H pylori* infection.

Key Words: Nuclear factor kappa B; CagA⁺ *H pylori*; Gastric cancer; Precancerous lesions

Li GP, Wu LF, Pu ZJ, Feng JL, Zheng ZM, Wang BZ. Increased expression of NF-κB p65 in CagA⁺ *H pylori* gastric precancerous lesions and carcinoma. Shijie Huaren Xiaohua Zazhi 2005;13 (17):2064-2068

摘要

目的: 探讨核因子-κB (nuclear factor-Kappa B, NF-κB)在癌前病变及胃癌组织中的表达及其与细胞毒素相关抗原A幽门螺杆菌(CagA⁺ *H pylori*)感染之间的关系。

方法: 慢性浅表性胃炎34例, 肠腺化生31例, 不典型增生34例, 胃癌55例, 应用免疫组化方法(SABC法)检测NF-κB p65的表达, 以¹⁴C-呼气试验、快速尿素酶试验和Warthin-Starry银染色检测*H pylori*, 采用斑点金免疫渗滤法检测患者血清抗*H pylori* CagA IgG抗体, 分析NF-κB p65表达与CagA⁺ *H pylori*感染之间、以及与胃癌组织学分型、临床病理分期、淋巴结转移的关系。

结果: 在慢性浅表性胃炎、肠腺化生、不典型增生和胃癌组中, NF-κB p65阳性表达率分别为15.0%, 41.9%, 64.7%和78.2%, 呈逐渐增高趋势, 各组间有显著性差异($\chi^2 = 43.98, P < 0.01$); *H pylori*感染率分别为70.0%, 67.7%, 73.5%和54.5%, 各组间无显著性差异($P > 0.05$). CagA⁺ *H pylori*构成比分别为53.6%, 61.9%, 68.0%和73.3%, 各组间无显著性差异($P > 0.05$). 在肠化组中, *H pylori* 阳性和CagA⁺ *H pylori*感染的患者NF-κB p65阳性表达率分别为57.1%和76.9%, 显著高于同组无*H pylori*感染的10.0% ($\chi^2 = 6.18, P < 0.05$) 和CagA⁻ *H pylori*感染者的25.0% ($\chi^2 = 5.45, P < 0.05$). 在胃癌组中, NF-κB p65阳性表达与T分期($\chi^2 = 5.91, P < 0.05$)及淋巴结转移有关($\chi^2 = 7.47, P < 0.05$), 但与胃癌组织学分型无关($P > 0.05$).

结论: NF-κB的异常活化在胃癌前病变及癌变过程中起作用, 早期的活化与CagA⁺ *H pylori*感染有关.

关键词: 核因子-κB; CagA⁺ *H pylori*; 胃癌; 癌前病变

李国平, 吴灵飞, 蒲泽锦, 冯家琳, 王炳周, 郑宗茂. CagA⁺ 幽门螺杆菌胃癌前及癌变中NF-κB的表达增强. 世界华人消化杂志 2005;13(17):2064–2068
http://www.wjgnet.com/1009-3079/17/2064.asp

0 引言

幽门螺杆菌 (*H pylori*) 感染是慢性活动性胃炎的主要病因, 是萎缩性胃炎肠上皮化生和不典型增生等癌前病变的促进因素^[1,5]. 流行病学调查资料表明, *H pylori* 感染者发生胃癌的危险性较非感染者6倍^[6], 但确切机制仍不清楚. 核因子-κB (nuclear factor kappa B, NF-κB) 作为一种多向转录调节因子^[7], 广泛参与多种生理和病理过程的基因调控^[8,11], 最近发现它在肿瘤发生发展中亦起重要作用^[12,15], 但有关在胃癌发病中的作用报道尚少. 有文献报道, 表达细胞毒素相关抗原A的*H pylori* (CagA⁺ *H pylori*) 可上调NF-κB的表达^[16]. 我们研究胃癌及癌前病变中NF-κB的表达以及*H pylori*, 尤其是CagA⁺ *H pylori*的感染状态, 旨在探讨二者之间的相互作用关系及其对胃癌发生发展的影响.

1 材料和方法

1.1 材料 2002–5/2004–12我院胃癌患者55例, 男30例, 女25例, 年龄32–77(平均62.4)岁. 高分化癌25例, 中分化癌13例, 低分化癌11例, 黏液腺癌6例. T₁期2例, T₂期11例, T₃期29例, T₄期13例. 淋巴结转移40例, 无淋巴结转移15例. 所有患者术前均未行化疗或放疗. 胃镜取材活检标本中慢性浅表性胃炎40例, 慢性萎缩性胃炎伴肠化31例, 不典型增生34例, 患者

均经病理组织学诊断证实. 胃黏膜活检标本取自胃窦或病变区, 手术标本均取癌灶及癌旁2 cm组织, 采用40 g/L多聚甲醛固定, 常规脱水、石蜡包埋, 备测. 取胃窦部黏膜作快速尿素酶试验、Warthin-starry银染色结合¹⁴C-呼气试验, 其中2项阳性定为*H pylori* 感染. 微量胶囊法¹⁴C-呼气试验所需试剂由深圳养和生物科技公司提供, 按说明书检测样品每分钟衰变数(dmp), ≥200 dmp为(+), ≤150 dmp为(-); Warthin-Starry银染色按常规组织病理学技术进行, 深棕色细长弯曲为*H pylori*染色. CagA⁺ *H pylori* 检测 采用斑点金免疫渗滤法. 试剂盒购自西安联尔生物技术公司, 仅对*H pylori*阳性者检测, 反应完成后有1个红色斑点为阴性, 有2个红色斑点为阳性, 无红色斑点为无效. 阳性者被视为CagA⁺ *H pylori*感染.

1.2 方法 采用链霉亲和素-生物素-过氧化物酶复合物(SABC)法 兔抗人NF-κB p65及SP试剂盒购自北京中山生物公司. 4 μm厚连续切片, 二甲苯脱蜡、酒精水化后, 浸入30 mL/L的双氧水中30 min以阻断内源性过氧化物酶的活性, 微波抗原修复, 山羊血清封闭, 分别依次滴加抗NF-κB p65抗体(工作浓度1:100), 4°C过夜, PBS液洗涤3次. 生物素化二抗, 加辣根过氧化物酶标记的链霉卵白素工作液, 室温、湿盒中作用30 min, PBS液洗涤3次, DAB液显色, 苏木精轻度复染后脱水、透明、封片. 以PBS代替一抗作阴性对照, 用已知胃癌阳性切片作阳性对照. 在组织切片中显示胞质或胞核染为淡黄至棕黄色者为阳性细胞. 先在低倍镜下选择阳性细胞最集中的5个视野, 每个视野观察100个细胞, 最后取5个视野中阳性细胞平均百分数作为每张切片的观察结果. 根据Handel et al^[17]的分级标准采用双盲法记分, 根据阳性细胞所占比例判断结果:<5%为阴性(-), 5–24%为弱阳性(+), 25–50%为中度阳性(++)>50%为强阳性(+++). 根据染色强度将阳性信号分为3类: 染色强度弱(+)—淡黄色或仅个别细胞呈黄至棕黄色染色, 染色强度强(+++)—呈棕黄至棕褐色染色, 中等染色强度(++)—染色强度介于弱阳性与强阳性之间. 染色强度×阳性细胞百分数为每例组织染色的综合记分. 综合记分≥1判为表达阳性, <1则为阴性.

统计学处理 采用 χ^2 检验和Spearman等级相关分析, 应用SPSS 10.0软件包, 检验水准 $\alpha = 0.05$.

2 结果

2.1 胃部疾病中 *H pylori* 及CagA⁺ *H pylori*感染和NF-κB p65表达 慢性浅表性胃炎40例中, *H pylori* 阳性28例(CagA⁺ *H pylori* 15例, 占53.6%), *H pylori* 阴性12例, *H pylori*感染率为70.0%. 慢性萎缩性胃炎

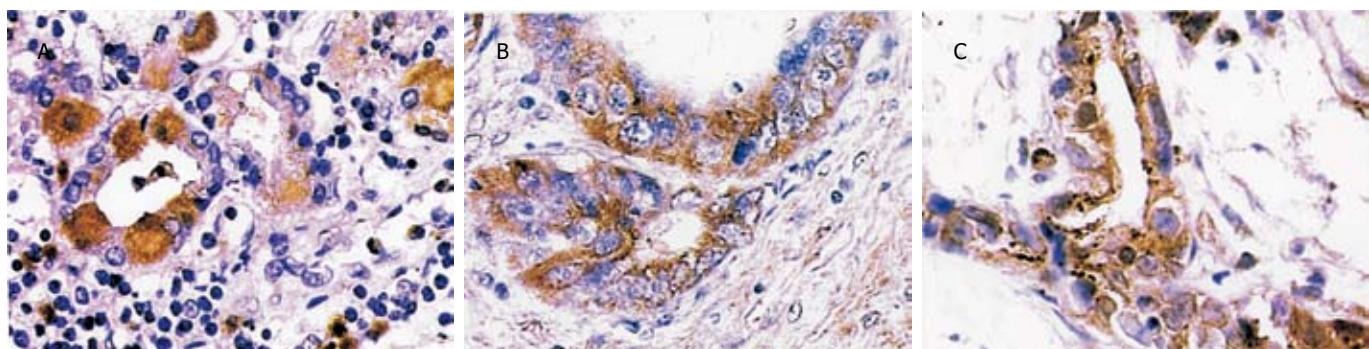


图1 NF-κB p65的表达(SP法, $\times 400$). A: 不典型增生; B: 高分化胃腺癌; C: 中分化胃腺癌.

表1 NF-κB p65表达与Hp感染的关系 n (%)

分组	<i>H pylori</i> (+)	NF-κB p65	<i>H pylori</i> (-)	NF-κB p65
浅表性胃炎	28	5(17.9)	12	1(8.3)
肠化	21	12(57.1)	10	1(10.0) ^a
不典型增生	25	16(64.0)	9	6(66.7)
胃癌	30	24(80.0)	25	19(76.0)

^a $P < 0.05$ vs *H pylori*(+).

伴肠化31例中, *H pylori*阳性21例(*CagA*⁺ *H pylori* 13例, 占61.9%), *H pylori*阴性10例, *H pylori*感染率为67.7%. 不典型增生34例中, *H pylori*阳性25例(*CagA*⁺ *H pylori* 17例, 占68.0%), *H pylori*阴性9例, *H pylori*感染率为73.5%. 55例胃癌组织中, *H pylori*阳性30例(*CagA*⁺ *H pylori* 22例, 占73.3%), *H pylori*阴性25例, *H pylori*感染率为54.5%. 各组 *H pylori*感染率无显著性差异($\chi^2 = 4.32$, $P > 0.05$). *CagA*⁺ *H pylori*感染的比例在胃炎、胃癌前病变至胃癌组中有逐渐增高趋势, 但无统计学意义($\chi^2 = 2.69$, $P > 0.05$). NF-κB p65蛋白表达呈棕黄色颗粒, 定位于细胞质, 部分亦可见于细胞核. 慢性浅表性胃炎组织中胞质、胞核中很少呈现表达. 癌前病变组织中其表达信号逐渐增强, 在不典型增生中炎性细胞浸润明显, 腺体结构破坏, 异型细胞胞质中可见其表达, 为棕黄颗粒(图1A). 胃癌细胞中则常见其表达, 呈粗细不等的棕黄或棕褐色颗粒, 胞质、胞核同时着色, 表达强度为中等或强阳性(图1B-C). 慢性浅表性胃炎、慢性萎缩性胃炎伴肠化、不典型增生和胃癌组织中NF-κB p65的表达率分别为15.0%, 41.9%, 64.7%和78.2%, 呈递增趋势, 经统计学处理, 差异有显著性($\chi^2 = 43.98$, $P < 0.005$).

2.2 *H pylori*与NF-κB p65表达的关系 按照*H pylori*感染情况分组, 浅表性胃炎中*H pylori*(+)组NF-κB p65表达率高于*H pylori*(-)组, 但无显著性差异($P > 0.05$). 肠化患者中*H pylori*(+)组NF-κB p65表达率显著高于*H pylori*(-)组, 差异有统计学意

义($\chi^2 = 6.18$, $P < 0.05$). 不典型增生和胃癌组中*H pylori*(+)与*H pylori*(-)患者NF-κB p65表达阳性率相近, 经统计学处理, 均无显著性差异($P > 0.05$) (表1). 按照*CagA*情况分组, 慢性萎缩性胃炎伴肠化组中*CagA*⁺ *H pylori*患者NF-κB p65阳性率为76.9%, *CagA*⁻ *H pylori*患者的阳性率为25.0%, 二者相比, 差异有统计学意义($\chi^2 = 5.45$, $P < 0.05$). 其余各组组内比较均无显著性差异($P > 0.05$) (表2).

表2 NF-κB p65表达与*CagA*⁺ *H pylori*感染的关系 n (%)

分组	<i>CagA</i> ⁺ <i>H pylori</i>	NF-κB p65	<i>CagA</i> ⁻ <i>H pylori</i>	NF-κB p65
浅表性胃炎	15	4(26.7)	13	1(7.7)
肠化	13	10(76.9)	8	2(25.0) ^a
不典型增生	17	13(76.5)	8	3(37.5)
胃癌	22	18(81.8)	8	6(75.0)

^a $P < 0.05$ vs *CagA*⁺ *H pylori*.

2.3 NF-κB p65与胃癌临床病理特征的关系 NF-κB p65表达阳性率与胃癌病理学类型无关, 与胃癌的临床分期、浸润深度及淋巴结转移有关(表3).

表3 NF-κB p65表达与胃癌临床病理特征的关系 n (%)

临床病理因素	n	NF-κB p65
高分化腺癌	25	22(88.0)
中分化腺癌	13	10(76.9)
未分化/黏液癌	17	11(64.7)
T ₁ +T ₂	13	7(53.8)
T ₃ +T ₄	42	36(85.7) ^a
无淋巴结转移	15	8(53.3)
有淋巴结转移	40	35(87.5) ^b

^a $P < 0.05$ vs T₁+T₂, ^b $P < 0.01$, 无淋巴结转移.

3 讨论

流行病学调查资料表明, 胃癌发生率与当地*H pylori*感染率呈正相关, 然而, *H pylori*如何引起胃癌的机制并不清楚. 一般认为, 胃癌的发生是一个有多因素参与的复杂而漫长的过程. 大多从慢性浅表性胃炎至

萎缩性胃炎, 逐渐出现肠上皮化生和不典型增生等癌前病变, 最终才转化成癌。本结果表明, 在有胃病症状的人群中 *H pylori* 的感染率为 54.5~73.5%, 与文献报道相同^[18, 19]。胃癌组患者的感染率较低(54.5%), 我们认为可能与研究方法有关。通过血清学方法, 我们发现在胃炎、胃癌前病变至胃癌组中 CagA⁺ *H pylori* 感染比例有逐渐增高趋势, 显示检测抗体的方法可能比实时检测细菌更能反映 *H pylori* 对胃部疾病的长期影响, 新近 Storskrubb *et al* 在 *H pylori* 对胃癌前病变的一组流行病学调查报告中亦支持此观点^[20]。慢性萎缩性胃炎伴肠化 *H pylori* 阳性或 CagA⁺ *H pylori* 患者 NF-κB p65 的阳性表达率显著高于 *H pylori* 阴性或 CagA⁻ *H pylori* 患者, 慢性浅表性胃炎组中 *H pylori* 阳性 NF-κB p65 表达虽高于 *H pylori* 阴性患者, 但未显出差异且均在较低的水平, 提示 *H pylori* 感染引起 NF-κB p65 表达上调需要一定的时间。不典型增生及胃癌组 NF-κB p65 的表达均明显增高, 且与 *H pylori* 或 CagA⁺ *H pylori* 感染无关, 说明一旦进入病变严重或不可逆阶段, 由于涉及多种遗传物质的改变, *H pylori* 的作用已不再易于显示出来, NF-κB p65 的激活是多因素作用长期累积的结果^[21, 22]。CagA⁺ *H pylori* 感染的胃炎向胃癌转变过程中 NF-κB p65 可能是重要的分子生物学环节。*H pylori*, 特别是 CagA⁺ *H pylori* 引起 NF-κB p65 活化, 并与其它因素一起启动靶基因 *c-myc*, *Cyclin D* 和 *bcl-2* 的转录^[23, 27], 破坏正常细胞的分化过程, 最终导致胃上皮细胞向恶性方向转化^[28], 可能是其参与胃癌发生的机制之一。最近研究发现, NF-κB 在肿瘤的发生、发展中起重要作用, 可调控细胞的增殖、分化、凋亡及恶性转化, 其活性失控常与哺乳动物肿瘤发生有关^[29, 31]。有证据表明, NF-κB 的持续活化可作为结肠癌、胰腺癌、乳腺癌、前列腺癌等多种实体肿瘤的标志^[7~11, 22, 29]。Wang *et al*^[32] 检测胰腺组织中 NF-κB p65 的表达, 结果胰腺癌的表达阳性率为 67%, 而正常胰腺组织表达罕见。在前列腺癌的研究中, Huang *et al*^[33] 报道 NF-κB 活化后可抑制细胞凋亡, 促进其增殖。Sasaki *et al*^[15] 应用免疫组织化学方法检测 NF-κB p65 在胃癌细胞胞质和胞核中的表达情况, 结果肿瘤组织中 NF-κB p65 的表达明显高于癌旁正常上皮细胞; 电泳迁移率改变分析显示胃癌组织中 NF-κB 的 DNA 结合活性与核转位的活性均增加。本研究中 NF-κB p65 在慢性浅表性胃炎、慢性萎缩性胃炎伴肠化、不典型增生及胃癌组织中的表达阳性率呈递增趋势, 各组之间差异有统计学意义, 且随着组织学病变程度的加重, 阳性细胞数量逐渐增多, 分布范围也逐渐扩大, 与 Sasaki *et al*^[15] 结果一致。NF-κB p65 在癌前病变中有较高的阳性率, 表

明 NF-κB p65 蛋白表达参与了胃癌前病变的进展, 为胃癌癌变的早期事件, 有可能成为检测早期癌变的标志物。本组 55 例胃癌中, NF-κB p65 表达阳性率在高分化腺癌中较高, 但与中分化腺癌、低分化或黏液癌无显著性差异, 提示 NF-κB 的表达与肿瘤发生的组织学类型无关。比较不同肿瘤分期中 NF-κB p65 的表达, T₃, T₄ 期的胃癌此蛋白的表达显著高于 T₁, T₂ 期, 而且伴淋巴结转移者显著高于无淋巴结转移者, 显示 NF-κB p65 表达与胃癌的分期、浸润深度及淋巴结转移等病理学特性关系密切, 越近晚期的癌肿表达阳性率相对也越高, 与史朝晖 *et al*^[34]、王维 *et al*^[35] 的报道相似, 提示 NF-κB p65 的异常活化与胃癌的发病密切相关外, 还可能对胃癌侵袭生长施加影响, 并可增加其转移潜能^[13, 14]。从另一角度分析, 此蛋白表达与胃癌侵润及转移有关, 则有可能作为判断胃癌预后的指标^[36]。

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编辑 潘伯荣 审读 张海宁