

论著

survivin基因启动子区-31C/G多态性与散发性结直肠癌遗传易感性的关系

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摘要 目的: 探讨生存素survivin基因启动子区-31C/G单核苷酸多态性与中国华南地区散发性结直肠癌(CRC)易感性的关系。方法: 采用聚合酶链反应-限制性片段长度多态性法(PCR-RFLP)检测华南地区711例健康人和702例CRC的survivin基因-31C/G位点单核苷酸多态性。结果: 结直肠癌患者CC基因型的频率明显高于对照人群(36.5% vs 26.2%, $\chi^2=17.89$, $P<0.01$),与CC基因型相比,CG、GG基因型和等位基因G携带者的CRC发病风险分别显著下降至0.61倍(95%CI=0.46-0.80, $P<0.01$)、0.52倍(95%CI=0.38-0.71, $P<0.01$)和0.58倍(95%CI=0.45-0.74, $P<0.01$)。结论: survivin基因-31C/G多态与CRC发病风险有关, -31G变异基因型是中国南方人群散发性结直肠癌独立保护因素。

关键词 [结直肠肿瘤](#); [基因,survivin](#); [多态性,单核苷酸](#)

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Association between survivin promoter -31C/G polymorphism and genetic susceptibility to sporadic colorectal cancer

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Abstract

AIM: To investigate the association between -31C/G polymorphism in the promoter of survivin gene and the susceptibility to sporadic colorectal cancer (CRC) in southern Chinese population. METHODS: survivin -31C/G genotypes were determined by PCR-RFLP in 711 healthy controls and 702 CRC cases. RESULTS: The number of CRC patients carrying with CC genotype was much higher than that of controls (36.5 % vs 26.2%, $\chi^2=17.89$, $P<0.01$). Compared to CC genotypes, CG, GG genotypes and G allele carriers had a significantly decreased risk of CRC, with the decrease being 0.61-fold (95% confidence interval=0.46-0.80, $P<0.01$), 0.52-fold (95% confidence interval=0.38-0.71, $P<0.01$) and 0.58-fold (95% confidence interval=0.45-0.74, $P<0.01$), respectively. CONCLUSION: survivin gene -31C/G polymorphism is associated with sporadic CRC risk, the G variant genotype is the independent protective factors against sporadic CRC in southern Chinese population.

Key words [Colorectal neoplasms](#) [Genes](#) [survivin](#) [Polymorphism](#) [single nucleotide](#)

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