论著

肿瘤坏死因子受体 II 196位点基因多态性与SLE相关性及功能研究

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摘要 目的:调查中国汉族人群肿瘤坏死因子受体II(TNFRII)196位基因多态性与SLE的关系并构建野生和 突变的逆转录病毒载体以研究其功能差异。 方法: 利用聚合酶链反应-限制性片段长度多态性(PCR-RFLP)方法检测了106例SLE和119例健康人TNFRII 196位的基因型。扩增TNFRII 196M cDNA 克隆到PMD18-T载体上,定点突变为TNFRII 196R,然后亚克隆到逆转录病毒载体PLXSN上(PLXSN-TNFRII 196M和PLXSN-TNFR II 196R)并分别转染大鼠系膜细胞,以 ELISA法观察对系膜细胞产生IL-6的影响。 结果: (1)SLE组TNFR II 196R等位基因型明显高于正常组(35.2% vs 14.3%,P<0.05); (2)成功构建野生型和突变型重组逆转录病毒载体; (3)rhTNFa刺激后196R型转染系膜细胞产生IL-6明显高于196M型(P<0.05)。 结论: TNFRII 196R与中国汉族人群SLE相关,其可以高效转导TNFa的信号,可能参与SLE发病。

关键词 红斑狼疮,系统性; 肿瘤坏死因子受体Ⅱ; 白细胞介素6

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Association and function study of tumor necrosis factor receptor II position 196 polymorphism with systemic lupus erythematosus

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Abstract

AIM: To investigate the association of gene polymorphism at position 196 of tumor necrosis factor receptor II (TNFRII) with systemic lupus erythematosus (SLE) in Chinese, and establish recombinant retroviral vector to analyze the function of the TNFR II 196M/R. METHODS: The genotype at position 196 of TNFR II was determined by PCR-RFLP in 106 SLE patients and 119 healthy controls in china. Human TNFR II 196M cDNA were amplified by PCR and cloned into PMD18-T vector. Then, PMD18-TNFR II 196R was induced by site-directed mutagenesis. The recombinant T vector, PMD18-TNFR II 196M and PMD18-TNFR II 196R, were subcloned into retroviral vector PLXSN. Both normal and variant were transfected into rat mesangial cell. The effects of TNFa on production of sTNFR II and IL-6 were study by ELISA. RESULTS: (1) The frequency of TNFR II 196R allele was significantly higher than those in controls (35.2% vs 14.3%, P<0.05); (2) The recombinant retroviral vector (PLXSN-TNFR 196M and PLXSN-TNFR 196R) was constructed successfully; (3) rhTNFa caused a significant increase in IL-6 production by rat mesangial cells transfected with PLXSN-TNFR II 196R than that with TNFR II 196M. CONCLUSION: These data indicate that TNFR II 196R allele is associated with SLE in the Chinese. TNFR II 196R transduces the signals of TNFa more effectively than TNFR II 196M, which may be involved in the pathogenesis of SLE.

Key words Lupus erythematosus systemic Tumor necrosis factor receptor II Interleukin-6

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