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论著

## 白芍总苷对系统性红斑狼疮CD<sup>4+</sup> T细胞ITGAL基因表达和启动子甲基化修饰的影响

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摘要:

目的: 探讨白芍总苷(total glucosides of peony, TGP)对系统性红斑狼疮(SLE)患者外周血CD<sup>4+</sup> T细胞甲基化敏感基因ITGAL(CD11a)表达及启动子甲基化修饰的影响。方法: 利用磁珠分选获得外周血CD<sup>4+</sup> T细胞, 用不同浓度TGP(0, 62.5, 312.5和1562.5 mg/L)处理SLE患者CD<sup>4+</sup> T细胞, 48 h后收集细胞。MTT法检测处理后CD<sup>4+</sup> T细胞的活力。定量PCR检测ITGAL基因mRNA表达水平。流式细胞术检测CD<sup>4+</sup> T细胞表面CD11a分子表达水平。亚硫酸盐测序分析ITGAL基因启动子甲基化水平。结果: TGP处理48 h后, 各浓度组CD<sup>4+</sup> T细胞活力无明显差异。与对照组相比, 高浓度组(1562.5 mg/L)ITGAL基因mRNA表达水平显著降低( $P<0.01$ ), CD<sup>4+</sup> T细胞表面CD11a表达水平亦显著降低( $P<0.01$ )。进一步DNA甲基化水平检测显示高浓度TGP处理组ITGAL基因启动子甲基化水平较对照组显著升高( $P<0.01$ )。结论: TGP可通过升高ITGAL基因启动子甲基化水平降低SLE患者外周血CD<sup>4+</sup> T细胞中CD11a表达水平, 初步揭示了TGP抑制SLE自身免疫反应的分子机制。

关键词: 白芍总苷 系统性红斑狼疮 CD<sup>4+</sup> T细胞 ITGAL DNA甲基化

## Effect of total glucosides of peony on expression and DNA methylation status of ITGAL gene in CD<sup>4+</sup> T cells of systemic lupus erythematosus

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Abstract:

Objective: To investigate the effect of total glucosides of peony (TGP) on expression and DNA methylation status of ITGAL gene (CD11a) in CD<sup>4+</sup> T cells from patients with systemic lupus erythematosus (SLE). Methods: CD<sup>4+</sup> T cells were isolated by positive selection using CD4 beads. CD<sup>4+</sup> T cells were treated by TGP at 0, 62.5, 312.5 and 1562.5 mg/L for 48 h. The MTT method was used to assess cell viability; mRNA expression level was measured by realtime-PCR; protein level of CD11a was measured by flow cytometric analysis; DNA methylation status was assayed by bisulfite sequencing. Results: No significant change in cell viability was found in CD<sup>4+</sup> T cells among the different concentration groups ( $P>0.05$ ). Compared with control, the mRNA and protein levels of ITGAL were down-regulated significantly in SLE CD<sup>4+</sup> T cells treated with TGP (1562.5 mg/L) ( $P<0.01$ ). Furthermore, the extent of DNA methylation of ITGAL promoter was increased in TGP (1562.5 mg/L) treated CD<sup>4+</sup> T cells compared with control group ( $P<0.01$ ). Conclusion: TGP can repress CD11a gene expression through enhancing DNA methylation of ITGAL promoter in CD<sup>4+</sup> T cells from patients with SLE. This observation represents a preliminary step in understanding the mechanism of TGP in SLE therapy.

Keywords: total glucosides of peony systemic lupus erythematosus CD<sup>4+</sup> T cells ITGAL DNA methylation

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