

## 论著

### 高血流性肺动脉高压大鼠肺组织中TGF- $\beta$ 1和CTGF的动态变化及意义

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#### 摘要:

目的:观察转化生长因子- $\beta$ 1 (TGF- $\beta$ 1) 与结缔组织生长因子(CTGF) 在高血流性肺动脉高压大鼠肺组织的动态表达变化, 探讨TGF- $\beta$ 1 和CTGF 在高血流性肺动脉高压过程中的作用。方法:50 只成年雄性SD 大鼠随机分成假手术组(S 组)、右肺切除1 周组(PE1 组)、2 周组(PE2 组)、4 周组(PE4 组) 和6 周组(PE6 组), 每组10 只。测量各组大鼠平均肺动脉压(mPAP)、右室肥大指数(RVHI) 和血管形态学指标;免疫组织化学染色和Western 印迹检测肺组织TGF- $\beta$ 1 和CTGF 蛋白表达;RT-PCR 检测TGF- $\beta$ 1 和CTGF mRNA 的表达。结果:与S 组比较, 右肺切除各时间组大鼠mPAP 和RVHI 均显著升高( $P<0.05$ ), PE1 和PE 2 组肌型动脉(MA) 和部分肌型动脉(PMA) 占肺中小血管总数的百分比[(MA+PMA)%]、MA 的相对中膜厚度(RMT) 和相对中膜面积(RMA) 与S 组比较无明显差异, PE4 和PE6 组(MA+PMA)%、RMT 及RMA 明显增高( $P<0.05$ );免疫组织化学染色显示, 右肺切除各时间组大鼠TGF- $\beta$ 1 和CTGF 较S 组分布广泛, 表达增强。Western 印迹结果表明, 与S 组比较, 右肺切除各时间组TGF- $\beta$ 1 蛋白均显著上升( $P<0.01$ ), 峰值出现在PE2 组;CTGF 蛋白在PE1 组和PE2 组较S 组升高, 但差异无统计学意义, PE4 组和PE6 组较S 组明显增高( $P<0.05$ )。与S 组比较, 右肺切除各时间组TGF- $\beta$ 1mRNA 表达均显著升高( $P<0.01$ ), PE2 组达峰, PE4 组和PE6 组维持在高水平。PE1 组CTGF mRNA 增高不明显, PE2 组、PE4 组和PE6 组明显上升( $P<0.01$ )。相关分析表明, CTGF 蛋白和mRNA 表达与RMT 和RMA 呈正相关( $P<0.05$ );TGF- $\beta$ 1 蛋白和mRNA 表达与RMT 和RMA无相关性;TGF- $\beta$ 1 mRNA 表达与CTGF mRNA 表达之间无明显相关性。结论:TGF- $\beta$ 1 和CTGF 参与了大鼠高血流性肺动脉高压的病理生理过程。

关键词: 转化生长因子- $\beta$ 1 结缔组织生长因子 高肺血流 肺动脉高压

### Changes of TGF- $\beta$ 1 and CTGF in rats with increased blood flow-induced pulmonary artery hypertension

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#### Abstract:

Objective: To evaluate the role of transforming growth factor- $\beta$ 1 (TGF- $\beta$ 1) and connective tissue growth factor (CTGF) in the pathogenesis of pulmonary artery hypertension, we observed the dynamic expression of TGF- $\beta$ 1 and CTGF in rats with high blood flow.

Methods: Fifty adult male SD rats were randomly divided into 5 groups: a sham group (group S) and groups with right pneumonectomy for 1, 2, 4 and 6 weeks (PE1, PE2, PE4 and PE6 group), 10 rats per group. The mean pulmonary arterial pressure (mPAP), vessel morphometry and right ventricle hypertrophy index (RVHI) were measured. TGF- $\beta$ 1 and CTGF protein expression in the lung tissues were determined with immunohistochemistry and Western blot. The expression of TGF- $\beta$ 1 mRNA and CTGF mRNA in the lung tissues was evaluated by RT-PCR.

Results: Compared with group S, the mPAP and RVHI in the rats were significantly increased in group PE1, PE2, PE4, and PE6 ( $P<0.05$ ); the indicators of vascular remodeling [(MA+PMA)%, RMT, and RMA] were markedly elevated in group PE4 and PE6 ( $P<0.05$ ), but not in group PE1 and PE2.

Immunohistochemical staining of TGF- $\beta$ 1 and CTGF were more prominent in all of the right pneumonectomy groups than in the sham group. Western blot showed that the level of TGF- $\beta$ 1 protein was significantly increased in all of the right pneumonectomy groups ( $P<0.01$ ), and the peak was observed in group PE2, whereas the level of CTGF protein was markedly elevated in group PE4 and PE6 ( $P<0.05$ ), but no change was noticed in group PE1 and PE2. Compare to group S, the mRNA level of TGF- $\beta$ 1 was dramatically increased in all right pneumonectomy groups ( $P<0.01$ ), peaked at group PE2, and remained high in group PE4 and PE6. In contrast, the elevation of mRNA level of CTGF was not significant in group PE1, but group PE2, PE4 and PE6 demonstrated significant mRNA level of CTGF ( $P<0.01$ ). Correlation analysis showed that the protein and mRNA levels of CTGF were positively correlated with RMT and RMA ( $r=0.743$ ,  $r=0.906$ ;  $P<0.05$ ), while no correlation between the protein and

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mRNA level of TGF- $\beta$ 1 with RMT or RMA. There was no correlation between the mRNA level of TGF- $\beta$ 1 and CTGF.

Conclusion: TGF- $\beta$ 1 and CTGF play a role in the pathogenesis of increased pulmonary bloodflow-induced pulmonary hypertension.

Keywords: transforming growth factor- $\beta$ 1 connective tissue growth factor high pulmonary blood flow pulmonary artery hypertension

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