

论文

正常细胞与镉转化细胞抑制消减cDNA文库构建

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摘要:

目的 构建正常人支气管上皮细胞(16HBE)与氯化镉诱导转化16HBE细胞间差异表达基因的消减cDNA文库。方法以16HBE为驱动子,氯化镉诱导转化16HBE细胞为检测子,应用抑制性消减杂交(SSH)方法构建cDNA文库,经2次消减杂交和2次PCR后,将巢式PCR产物插入载体,随机挑选克隆进行鉴定。结果 得到纯度高及完整性好的总RNA和mRNA,并扩增出良好的双链cDNA,cDNA与接头的连接效率>25%,最终使差异表达基因得到富集;经蓝白菌落筛选,获得1 200余个白色阳性克隆;随机挑选50个白色克隆进行PCR扩增,显示96%克隆均有100~600 bp的插入片段,这些片段可能是差异表达基因cDNA片段,提示用SSH法及T/A克隆技术有效构建了两细胞株间差异表达基因的消减cDNA文库。结论 成功创建正常16HBE细胞与镉转化16HBE细胞差异表达基因消减cDNA文库。

关键词: 氯化镉 人支气管上皮细胞(16HBE) 抑制性消减杂交(SSH) cDNA文库

Construction of suppression subtractive cDNA libraries of normal and cadmium-transformed cells

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Abstract:

Objective To construct a subtractive cDNA library of differentially expressed genes in normal and cadmium-transformed cells. **Methods** The human bronchial epithelial cells(16HBE cells) transformed by cadmium chloride was used as a tester, and the normal 16HBE cells as a driver, to construct a cDNA library using suppression subtractive hybridization(SSH). The products were inserted into TA vector after two subtractive hybridization and nested PCR. The clones were picked up randomly and analyzed with PCR. **Results** The total RNA and mRNA of purity and good integrity were extracted, and the double-stranded cDNA was well produced. The link efficiency of cDNA and connector was greater than 25%. And finally the differentially expressed genes were enriched with subtractive hybridization and nested PCR. After blue-white screening, the amplified library contained more than 1 200 white positive clones, and fifty of them were selected randomly and analyzed with PCR. Totally 96% of the clones had the inserted 100-600 bp segment and the segments might be the gene cDNA segments with differential gene expressions. **Conclusion** The subtractive cDNA library of differential genes in transformed 16HBE cells induced by cadmium was successfully established.

Keywords: cadmium chloride human bronchial epithelial cell(16HBE cells) suppression subtractive hybridization cDNA library

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