定量PCR的非同源对照模板的构建 Generation of Non-homologous Competitor DNA for Competitive Q uantitative PCR

陈燃,金袭,毛裕民 CHEN Ran, JIN Zhe, MAO Yu-min

复旦大学生命科学学院遗传所遗传工程国家重点实验室,上海 200433 State Key Lab of Genetic Engineering, Institute of Genetics, Sc hool of Life Sciences, Fudan University, Shanghai 200433, China 收稿日期 修回日期 网络版发布日期 接受日期

摘要 建立了HCVRNA和hTERTmRNA的竞争定量PCR系统。对照模板的构建方法是:利用计算机辅助优选设计连接弓物,低严紧型扩增大肠杆菌DNA,回收并克隆预期的DNA片段。该片段与靶序列除两端引物序列完全相同外,无同源性,因此可作为非同源对照模板。这种构建非同源对照模板的方法,简便易行,适应面广。

Abstract: It cannot be neglected sometimes that the error caus ed by the "heteroduplex DNA" that occurs accompanying with the homologous comp etitor DNA which is usually used in the competitive quantitative PCR (CQ-PCR). Here a method is developed to generate non-homologous competitor DNA for CQ-PCR detection of the interest DNA, based on low-stringency amplification of cross-species' DNA with a pair of linker-primers which are designed according to partly homologous sites of the interest DNA primers in cross-species' DNA. With the method, the non-homologous competitor DNAs for the HCV RNA and hTERT mRNA are generated from E. coli DNA respectively, then the CQ-PCR systems are established for the 2 species RNAs with the RRTR (Repeated reverse transcription reaction). The method is multi-adaptive and easy to apply.

关键词竞争定量PCR对照模板异源双链 Keywordscompetitive quantitative PCRcompetitor DNAheteroduplex DNA

分类号

扩展功能

本文信息

- ▶ Supporting info
- ▶ <u>PDF</u>(0KB)
- ▶[HTML全文](0KB)
- ▶参考文献

服务与反馈

- ▶把本文推荐给朋友
- ▶加入我的书架
- ▶加入引用管理器
- ▶复制索引
- ▶ Email Alert
- ▶文章反馈
- ▶浏览反馈信息

相关信息

▶ <u>本刊中 包含"竞争定量PCR"的</u> 相关文章

▶本文作者相关文章

- 陈燃
- 金袭
- · 毛裕民CHEN Ran
- JIN Zhe
- · MAO Yu-min

Abstract

Key words

DOI:

通讯作者