

红掌根腐病原鉴定及其PCR 检测方法

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Identification and PCR Detection of the Pathogen Causing Root Rot of *Anthurium andraeanum*

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摘要 利用真菌通用引物ITS1 和ITS4 扩增红掌根腐病菌转录间隔区并进行序列测定, 通过序列比较, 设计了1 对红掌根腐病菌的特异引物SF1/SR2, 对30 个红掌根腐病原菌、8 种其它真菌和2 种细菌基因组DNA 进行PCR 扩增。结果表明, 只有红掌根腐病菌获得572 bp 的特异带。使用引物SF1/SR2 对华丽腐霉进行PCR 扩增, 其检测灵敏度在DNA 水平上可达1 pg。运用设计的引物从红掌根腐病菌基因组DNA 以及人工接种和自然发病的红掌植株中扩增到572 bp 的特异片段, 实现了对红掌根腐病菌的快速可靠的检测。

关键词: 红掌 根腐病菌 ITS 分析 PCR 检测

Abstract: Abstract: Based on the difference in internal transcribed spacer (ITS) sequences of *Pythium splendens* and other *Pythium* spp., a specific pair of primers SF1/SR2 was designed. Among 30 *P. splendens* isolates causing root rot of *Anthurium andraeanum*, and other eight fungi and two bacteria species, the primer pair amplified a single 572 bp product from all isolates of *P. splendens*, but not from any other isolates tested. The sensitivity of detection of the pathogen *P. splendens* with primers SF1/SR2 was 1 pg genomic DNA. It could amplified a specific single product from natural infected *A. andraeanum* plants, that was not amplified from healthy tissue. The results showed that the PCR protocol provides a rapid, sensitive and reliable tool routine detection and identification of *P. splendens*. In addition, this study is beneficial to control root rot disease of *A. andraeanum*.

Keywords: *Anthurium andraeanum*, *Pythium splendens*, ITS analysis, PCR detection

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