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生物技术一研究报告

烟草CMS相关基因orf25的生物信息学分析

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摘要:

线粒体基因orf25异常可能导致烟草细胞质雄性不育(CMS)。为探讨orf25基因导致CMS的可能机理,本文采用生物信息学方法,分析烟草雄性不育系'MS革新3号'及其保持系中的orf25基因在其编码蛋白各结构层次上存在的差异。结果显示,不育系中orf25基因因碱基改变提前出现终止子从而导致其编码蛋白缺失结构域,这极可能成为干扰线粒体F(0)F(1)-ATP合成酶功能的关键因素,从而成为导致烟草'革新3号'CMS的根本原因之一。

关键词: 不育机理

The Bioinformatics Analysis of orf25 Gene Related to Tobacco Cytoplasmic Male Sterility

Abstract:

It was reported that aberrant mitochondrial gene or f25 might cause to bacco cytoplasmic male sterility (CMS). To understand the mechanism of CMS, the differences of or f25 gene between CMS line 'Gexin 3' and its maintainer line were analyzed by bioinformatics at different structure levels of coding protein. The results showed that the terminal codon appeared early because of the base change in the or f25 gene, which led to the lack of the structural domain in coding protein. It might be the key factor that disturbed the function of f(0)f(1)-ATP synthetase in mitochondrion, and become one of the fundamental causes which led to CMS of to bacco.

Keywords: male sterile mechanism

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