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[1]韩利涛,杨守萍,喻德跃,等.大豆细胞质雄性不育系与保持系atp6基因的RNA编辑比较研究[J].大豆科学,2010,29(03):361-365.  
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## 大豆细胞质雄性不育系与保持系atp6基因的RNA编辑比较研究

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摘要: 对大豆细胞质雄性不育系NJCMS1A与其保持系NJCMS1B的atp6基因的RNA编辑进行比较研究。结果在不不育系NJCMS1A与保持系NJCMS1B的atp6-3基因保守区中均发现2个编辑位点, 但互不相同, 并且导致了氨基酸的不同变化; mtDNA 序列分析显示, atp6-3基因转录本保守区在不不育系NJCMS1A与保持系NJCMS1B间存在1个碱基的差异; 另外还发现atp6-1、atp6-2和atp6-3的表达在不不育系NJCMS1A与保持系NJCMS1B间存在明显差异。

Abstract: The atp6 gene is an important mitochondrial functional gene. The studies on some plant species showed that RNA editing of atp6 gene was related to the cytoplasmic male sterility. In this paper, the comparative studies of RNA editing of atp6 gene between the cytoplasmic male sterile line NJCMS1A and its maintainer line NJCMS1B in soybean (Glycine max L. Merr.) were conducted. The results showed that two editing sites were found on the conservative region of the atp6 gene of NJCMS1A and NJCMS1B, but the two editing sites of the atp6 gene of NJCMS1A were different from those of NJCMS1B, and led to the different changes of amino acid. The results of the mtDNA sequence comparison showed that one base difference was found in the conservative region of atp6-3 gene between NJCMS1A and NJCMS1B. In addition, the obvious difference was found on the expression of atp6-1, atp6-2 and atp6-3 between NJCMS1A and NJCMS1B.

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