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梨两个休眠相关MADS-box 基因特征及在其休眠过程中的表达分析

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Characteristics of Two Dormancy-associated MADS-box Genes in Pear and Their Expression Analysis During the Dormancy

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摘要 *PpMADS1* 和*PpMADS2* 是从'酥梨'(*Pyrus pyrifolia* white pear group'Suli')休眠芽转录组文库筛选的两个与休眠相关的MADS-box 基因序列。为了解序列的特征,对其进行了相关生物信息学分析,并以'翠冠'和'圆黄'梨为试材,用实时定量PCR 技术分析其花芽休眠不同阶段的表达变化。结果表明:两个基因都具有MADS-box 家族的特征序列MIKC-基序,*PpMADS1* 和*PpMADS2* 分别与日本梨(*P. pyrifolia* 'Kosui')休眠相关的两个MADS-box 基因*PpMADS13-1* 和*PpMADS13-2* 聚在一起,且单独聚为一支,与李属植物休眠相关的MADS-box 基因关系最近。在两个品种的休眠过程中,*PpMADS1* 和*PpMADS1* 和*PpMADS2* 的表达呈现相似的变化趋势,都有一个表达高峰,但'翠冠'梨*PpMADS1* 和*PpMADS2* 的表达高峰均出现在11 月15 日,而'圆黄'梨*PpMADS1* 的表达高峰延后到12 月30 日,PpMADS2 的表达高峰出现在12 月15 日。两个品种*PpMADS1* 和*PpMADS2* 的表达高峰出现在12 月15 日。两个品种*PpMADS1* 和*PpMADS2* 的表达高峰都出现在内休眠解除之前,随内休眠解除其表达量下调,在生态休眠阶段表达量维持在较低的水平。据此推测*PpMADS1* 和*PpMADS2* 的表达对梨芽内休眠的解除具有调控作用

关键词: 梨 芽 休眠 MADS-box 基因 表达分析

Abstract: Two dormancy-associated MADS-box genes (PpMADS1 and PpMADS2) were obtained from transcriptome library constructed Pyrus pyrifolia white pear group 'Suli'. To understand sequence characteristics of two genes, bioinformatics analysis was carried out. Furthermore the gene expression patterns in lateral flower buds of both pear cultivars 'Cuiguan' and 'Wonhwang' were analyzed during the period of dormancy by quantitative real-time PCR. The results showed that PpMADS1 and PpMADS2 contained well-conserved MIKC-motifs of MADS-box. A phylogenetic analysis revealed that PpMADS1 was closely related to PpMADS13-1 of Japanese pear (P. pyrifolia 'Kosui'), whereas PpMADS2 was similar to PpMADS13-2 of Japanese pear. The pear DAM genes formed an independent subclade and were closely related to those of Prunus spp. The expression patterns of PpMADS1 and PpMADS2 were similar with one peak in two cultivars during the period of dormancy. However, the timing of gene expression peak differed between two cultivars. The PpMADS1 and PpMADS2 expression peak occured on the November 15th in 'Cuiquan' pear, while the PpMADS1 expression peak on the December 30th, and the PpMADS2 expression peak on the December 15th in 'Wonhwang' pear. The PpMADS1 and PpMADS2 expression peak appeared prior to the release of endodormancy, and their expression levels were gradually down-regulated after the turn over of endodormancy and kept at a lower level during the period of ecodormancy. These results indicated that PpMADS1 and PpMADS2 might play an important role in the regulation of bud endodormancy-release in pear.

Keywords: pear, bud, dormancy, MADS-box gene, expression analysis

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