

甜瓜 *CmACO I* 启动子组织特异性表达研究

毛 娟¹, 陆 璐², 陈佰鸿¹, 褚明宇¹, 赵长增^{1,*}

(¹甘肃农业大学农学院, 兰州 730070; ²西北师范大学生命科学学院, 兰州 730070)

Studies on Tissue Specific Regulation of *CmACO I* Promoter in Melon

MAO Juan¹, LU Lu², CHEN Bai-hong¹, CHU Ming-yul, and ZHAO Chang-zeng^{1,*}

(¹College of Agronomy, Gansu Agricultural University, Lanzhou 730070, China; ²College of Life Science, Northwest Normal University, Lanzhou 730070, China)

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摘要 为进一步明确甜瓜果实软化的分子机理, 构建了ACC氧化酶I基因(*CmACO I*)启动子与GUS基因融合的植物表达载体, 采用根癌农杆菌介导法转化甜瓜‘甘甜一号’。通过卡那霉素抗性和PCR检测筛选呈阳性的转化植株, 取不同组织进行X-Gluc染色。结果表明, 转化植株的根、茎、叶、花、果实等器官组织经X-Gluc染色后, 只在花药组织和成熟果果皮中出现蓝色斑点, 其余组织均未检出, 表明甜瓜*CmACO I*启动子能够驱动GUS基因在转基因甜瓜花药和成熟果果皮中特异表达。

关键词: 甜瓜 *CmACO I* 启动子 GUS基因 组织特异性

Abstract: To provide an important reference for further clarifying the molecular mechanism of fruitsoftening, GUS histochemical staining method was employed to detect tissue specific regulation of the *CmACO I* promoter in melon (*Cucumis melo* L.). The plant expression vector with a fusion gene, *CmACO I* promoter and *GUS* gene was established and melon ‘GT-1’ was transformed by using *Agrobacterium*-mediated transformation system. Then the roots, stems, leaves, flowers, fruits and other organs in transformed plants during different growth stages were dyed with X-Gluc. The transformed plants with Kanamycin resistance and positive detection by PCR were obtained. It showed that some blue spots in the anther tissue and peel organ of the after-ripening fruit in the plants with a fusion gene (*CmACO I* promoter and *GUS* gene) were detected by using X-Gluc dye, while the other organs did not appear blue spots. It showed that *GUS* gene was driven by melon *CmACO I* promoter and expressed specifically in anther tissue and peel organ of the after-ripening fruit.

Keywords: melon, *CmACO I* promoter, *GUS* gene, tissue specificity

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