

枳抗逆基因*PtrZPT2-2*的克隆与表达分析

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Cloning and Expression Analysis of *PtrZPT2-2* from Trifoliate Orange (*Poncirus trifoliata*)

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摘要 通过电子克隆和RT-PCR 法从枳[Poncirus trifoliata (L.) Raf.]中克隆到1 个C2H2 型锌指蛋白家族基因, 命名为*PtrZPT2-2*。该基因编码159 个氨基酸残基, 预测蛋白分子量为17.28 kD, 等电点为9.51。*PtrZPT2-2* 包含两个C2H2 型锌指结构域, 在其C -末端含有一个转录抑制结构域EAR/DLN-box (DLNL)。*PtrZPT2-2* 与其他植物参与各种非生物胁迫反应的C2H2 型锌指蛋白有较高的一致性。系统发育树分析表明*PtrZPT2-2* 与杨树PtaZFP2, 矮牵牛ZPT2-12, ZPT2-13, 拟南芥ZAT12、ZAT7 亲缘关系较近。半定量PCR 分析表明, 低温、高盐、干旱胁迫和ABA 处理下枳叶、茎、根中*PtrZPT2-2* 均被诱导上调。*PtrZPT2-2* 可能在枳适应低温、高盐、干旱等非生物胁迫和ABA 响应中起着重要作用。

关键词: 枳 C2H2 型锌指蛋白 *PtrZPT2-2*

Abstract: In the present study, a novel C2H2-type zinc finger protein gene, *PtrZPT2-2*, was cloned from trifoliate orange by *in silico* and RT-PCR approaches. The *PtrZPT2-2* encodes a protein of 159 amino acid residues with a predicted molecular mass of 17.28 kD and an isoelectric point of 9.51. The *PtrZPT2-2* protein contains two C2H2-type zinc finger domains, and a putative transcription repression domain (EAR/DLN-box) at the C-terminus. Phylogenetic analysis showed that the *PtrZPT2-2* clustered with PtaZFP2, ZPT2-12, ZPT2-13, ZAT12 and ZAT7. *PtrZPT2-2* shared high identity with other C2H2-type zinc finger proteins involved in abiotic stress responses. Semi-quantitative PCR analysis showed that expression of *PtrZPT2-2* was upregulated by cold, salt, drought and ABA treatments in leaves, stems and roots of trifoliate orange seedlings. Our data suggested that *PtrZPT2-2* is a new member of the C2H2-type zinc finger protein genes and may play an important role in response to cold, salt, drought and ABA stresses in citrus.

Keywords: [trifoliate orange](#), [C2H2-type zinc finger](#), [PtrZPT2-2](#)

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