

杜梨类钙调磷酸酶B亚基蛋白基因*PbCBL2*的克隆和功能初探李慧^{1,2,*}, 李刚波^{1,3,*}, 丛郁⁴, 常有宏^{1,2,**}, 蔺经¹, 盛宝龙¹

(1江苏省农业科学院园艺研究所, 南京 210014; 2国家农业科技华东(江苏)创新中心——高效园艺作物遗传改良实验室, 南京 210014; 3南京农业大学园艺学院, 南京 210095; 4中国科学院南京土壤研究所土壤与农业可持续发展国家重点实验室, 南京 210008)

Isolation of a Calcineurin B-like Protein Gene *PbCBL2* from *Pyrus betulaefolia* and Preliminary Study of Gene FunctionLI Hui^{1,2,*}, LI Gang-bo^{1,3,*}, CONG Yu⁴, CHANG You-hong^{1,2,**}, LIN Jing¹, and SHENG Bao-long¹

(1Institute of Horticulture, Jiangsu Academy of Agricultural Sciences, Nanjing 210014, China; 2National Agricultural Science and Technology Jiangsu Innovative Center - Efficient Horticulture Crop Genetic Improvement Laboratory, Nanjing 210014, China; 3College of Horticulture, Nanjing Agricultural University, Nanjing 210095, China; 4State Key Laboratory of Soil and Sustainable Agriculture, Institute of Soil Science, Chinese Academy of Sciences, Nanjing 210008, China)

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摘要 类钙调磷酸酶B亚基蛋白(Calcineurin B-like protein, CBLs)是植物中一类重要的钙离子传感器, 参与调控植物生长发育及逆境胁迫响应过程。为了探明杜梨CBLs家族成员*PbCBL2*的序列特征和表达模式, 以杜梨(*Pyrus betulaefolia* Bunge)幼苗为试材, 运用EST搜索结合RACE技术、染色体步移法对*PbCBL2*的cDNA、DNA和启动子进行克隆, 采用半定量RT-PCR和原核表达研究该基因在非生物胁迫下的表达模式。结果表明, *PbCBL2*基因cDNA序列长681 bp, 编码一个含有226个氨基酸残基的蛋白。基因组DNA序列长1 927 bp, 包括8个外显子和7个内含子, 启动子序列包含光反应元件、低氧诱导必需顺式作用元件、赤霉素反应元件和水杨酸响应顺式作用元件。*PbCBL2*编码的多肽具有植物类钙调磷酸酶B亚基蛋白结合Ca²⁺所必需的4个EF手型结构和1个典型的植物钙调磷酸酶A亚基结合位点。未经处理的杜梨幼苗(对照)根和叶中未检测到*PbCBL2*的表达,*PbCBL2*的表达受NaCl、PEG6000、甘露醇和ABA诱导上调。*PbCBL2*转入大肠杆菌BL21 (DE3)后, 能够明显减轻NaCl、甘露醇和PEG6000对该菌株的生长抑制。*PbCBL2*基因具备植物CBLs基因家族的固有特征, 对盐碱、干旱、渗透胁迫和ABA处理均存在转录响应, 大肠杆菌转入该基因后能够提高对盐胁迫和渗透胁迫的耐受能力。

关键词: 杜梨 类钙调磷酸酶B亚基蛋白 基因克隆 基因表达特点 原核表达 逆境胁迫

Abstract: Calcineurin B-like protein (CBLs), as a plant calcium sensor, plays critical role in the regulation of plant growth and stress response process. However, *CBL2* gene sequence feature, expression characteristic and physiological function in birch-leaf pear (*Pyrus betulaefolia* Bunge) are largely unknown. In this study, we isolated the cDNA, genomic DNA and its responding promoter sequences of *PbCBL2* gene from birch-leaf pear seedlings by EST database mining, rapid amplification of cDNA ends (RACE) and genome walking approaches. The results have showed that *PbCBL2* cDNA sequence contained a 681 bp open reading frame which encoded 226 amino acid residues. The length of genomic DNA sequence was 1 927 bp which consists of 8 exons and 7 introns. The promoter region of *PbCBL2* harbored some specific regulatory elements or motifs, such as light responsive element, cis-acting regulatory element essential for the anaerobic induction, gibberellin-responsive element and cis-acting element involved in salicylic acid responsiveness. The deduced *PbCBL2* polypeptide had four EF-hand structure domains (58–71, 95–106, 132–143 and 176–187 amino acids) which was necessary for calcium-binding and one calcineurin A subunit binding sites (156–171 amino acids). Semi-quantitative RT-PCR and prokaryotic expression analyses validated that the mRNA abundance of *PbCBL2* is responsive to different abiotic stresses. However, *PbCBL2* expression was barely detected in roots and leaves of birch-leaf pear seedling without abiotic stresses treatment. The inhibition effects on BL21 (DE3) growth causing by NaCl, mannitol or PEG6000 were significantly alleviated after *PbCBL2* gene transformation. Our studies have suggested that *PbCBL2* gene has the inherent characteristics of the CBLs gene family in plants, which transcription level is respond to salt, drought, osmotic stresses and ABA treatment. *E. coli* BL21 (DE3) tolerance to salt stress and osmotic stress was enhanced by transferred *PbCBL2*.

Keywords: *Pyrus betulaefolia* Bunge, calcineurin B-like protein, gene cloning, gene expression characteristics, prokaryotic expression, environment stress

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- [1] 王凌云^{1,2}, 孙进华¹, 刘保华¹, 王家保^{1,*}.荔枝水孔蛋白基因*LcPIP*的克隆与组织特异性表达研究[J].园艺学报, 2013,40(8): 1456-1464
- [2] 杨德翠, 张玉喜, 郑国生*.牡丹病程相关蛋白1基因的克隆及表达分析[J].园艺学报, 2013,40(8): 1583-1590
- [3] 杨国峰, 沈文涛, 言普, 黎小瑛, 周鹏.原核表达的PRSV HC-Pro 基因同源dsRNA 诱导番木瓜抗性的研究[J].园艺学报, 2013,40(7): 1269-1277
- [4] 梁云, 袁素霞, 冯慧颖, 徐雷锋, 袁迎迎, 刘春, 明军.百合肌动蛋白基因*lilyActin* 的克隆与表达分析[J].园艺学报, 2013,40(7): 1318-1326
- [5] 金雪花^{1,2}, 洪艳¹, 黄河¹, 戴思兰^{1,*}, 朱婧¹.瓜叶菊谷胱甘肽转移酶基因*GST*的分离及表达分析[J].园艺学报, 2013,40(6): 1129-
- [6] 郭勤卫, 李季, 崔利, 张停林, Kere George Mbira, 陈劲枫*.黄瓜生长素响应因子*CsARF10*亚家族3个基因的克隆与表达分析[J].园艺学报, 2013,40(6): 1071-
- [7] 邢爱佳¹, 马小军^{2,3,*}, 莫长明^{1,3}, 潘丽梅^{3,4}, 韦鹏霄¹, 唐春风^{3,4}, 唐其^{3,4,*}.罗汉果葡萄糖基转移酶基因的克隆及原核表达[J].园艺学报, 2013,40(6): 1195-
- [8] 周晨阳, 金基强, 马春雷, 姚明哲, 陈亮.茶树*TIDH*核苷酸多样性及与咖啡碱含量的关联分析[J].园艺学报, 2013,40(5): 981-
- [9] 黄春红, 高燕会, 朱玉球, 童再康, 姜小凤.石蒜黄烷酮3-羟化酶基因*LrF3H*的克隆及表达分析[J].园艺学报, 2013,40(5): 960-
- [10] 施艳, 王振跃, 袁媛, 刘珊珊, 孙虎, 古勤生.瓜类褪绿黄化病毒*P22*基因在大肠杆菌中的表达及抗血清的制备[J].园艺学报, 2013,40(4): 762-
- [11] 曹庆芹, 邓杰, 朱丽静, 白隽帆, 赵天, 朱旭文, 姜奕晨.‘红颜’草莓菌根磷转运蛋白基因的克隆及荧光定量表达分析[J].园艺学报, 2013,40(4): 641-
- [12] 叶阳阳, 陈典, 王勇.洋葱开花相关基因*AcLFY*的克隆与表达分析[J].园艺学报, 2013,40(2): 283-291
- [13] 黄敏玲, 樊荣辉.鹤望兰八氯番茄红素脱氢酶基因*SrPDS*的克隆及表达分析[J].园艺学报, 2013,40(2): 373-379
- [14] 陈新, 梁丽松, 马庆华, 赵天田, 刘庆忠, 王贵禧.平榛脱水素基因的克隆与表达分析[J].园艺学报, 2013,40(1): 32-40
- [15] 黄洁, 刘晓华, 管洁, 吕英民.百合分子育种研究进展[J].园艺学报, 2012,39(9): 1793-1808