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Molecular Cloning and Characterization of A Novel Multiple Stressinducible Peroxidase Promoter from Sweetpotato

Sun-Hwa Ryu, Soo-Young Park, Yun-Hee Kim, Suk-Yoon Kwon, Sang-Soo Kwak*

Laboratory of Environmental Biotechnology, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Daejeon 305-806, Korea

Objectives

We have cloned more than 20 peroxidase (POD) cDNAs from suspension cultures of sweetpotato (*Ipomoea batatas* (L.) Lam.). Among them, *swpa4* was most highly induced by both abiotic and biotic stress, suggesting that this gene was regulated by a multiple stress-inducible promoter (Park et al. 2003). In this study, a stress-inducible *SWPA4* promoter was isolated by chromosome walking and its 5'-deletion mutants were characterized in transgenic tobacco plants and transient assay using tobacco BY-2 protoplasts.

Materials and Methods

- Plant Materials: Sweetpotato (*Ipomoea batata* (L.) Lam. cv. White Star), Tobacco BY-2 suspension cells, Tobacco (*Nicotiana tabaccum* cv. Xanthi)
- 2. Methods: Transient assay by PEG transfection
 Agrobacterium tumefaciens-mediated transformation

Results and Discussion

A promoter of genomic clone (referred to SWPA4) with 2,433 bp of 5'-upstream sequence from the translation start site was obtained from sweetpotato by chromosome walking. Sequence analysis reveals that SWPA4 promoter contains putative binding sites for several transcription factors including ELRE, MYBGAHV, ASF1, ABRE, GCN4, W-box and HSF. Employing a transient expression assay in tobacco BY-2 protoplasts, 2.4 kb promoter and five progressive deletion mutants, -1934, -1467, -1199, -818 and -433, were fused to the GUS gene and their GUS activity was analysed. The GUS activity of SWPA4 was enhanced by hydrogen peroxide treatment. Each deletion mutants showed diverse GUS activity. The -1934 bp deletion mutant showed about 8 times higher GUS expression than the CaMV 35S promoter. The further characterization of SWPA4 and deletion promoters is under study in tobacco transgenic plants.

Reference

Park SY et al. (2003) Differential expression of six novel peroxidase cDNAs from cell cultures of sweetpotato in response to stress. Molecular General and Genetics 269: 542-552

^{*} Corresponding author: TEL: 042-860-4432, E-mail: sskwak@kribb.re.kr