Thermotolerant Transgenic Ginseng (*Panax ginseng* C.A. Meyer) by Introducing Isoprene Synthase Gene through Agrobacterium tumefaciens-mediated Transformation

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ABSTRACT: The cost of conventional cultivation of ginseng (Panax ginseng C.A. Meyer) is very expensive, because shadow condition should be maintained during cultivation periods owing to inherently weak plant for high-temperature. Therefore, application of plant biotechnology may be possible to overcome these difficulties caused by conventional breeding of ginseng. Transgenic plants were produced via Agrobacterium tumefaciens GV3101, both carrying the binary plasmid pBI121mLPISO with nptII and Iso (isoprene synthase) gene. Integration of the transgenes into the P. ginseng nuclear genome was confirmed by PCR analysis using nptII primers and Iso primers. RT-PCR result also demonstrated the foreign isoprene synthase gene in three transgenic plant lines (T1, T3, and T5) which was expressed at the transcriptional level. When whole plants of transgenic ginseng were exposed to high temperature at 46 °C for 1 h, a non-transformed plant was wilted from heat shock, whereas a transgenic plant appeared to remain healthy. We suggest that the introduction of exogenous isoprene synthase is considered as alternative methods for generating thermotolerance ginseng.

Key words: Agrobacterium tumefaciens, Panax ginseng, isoprene synthase, thermotolerance

INTRODUCTION

Panax ginseng C.A. Meyer, commonly known as ginseng, one of the most medicinally important genera in the Orient. This plant have been applied as tonic, anti-cancer, vaccine for common cold and/or influenza, prophylacetic, anti-stress, and anti-aging agents (See et al., 1997; Scaglione et al., 1996; Takahashi, 1992). The cultivation of ginseng is troublesome and a period of 4-6 years is required to harvest the roots. The cost of conventional cultivation of ginseng is very expensive, because shadow condition should be maintained during cultivation periods owing to inherently weak plant for high-temperature. Therefore, application of plant biotechnology may be possible to overcome these difficulties caused by conventional breeding of ginseng.

Until now, the best characterized aspect of acquired thermotolerance has been production of heat shock protein (HSP) (Vierling, 1991; Howarth & Skot, 1994; Burke, 2001). Majority of HSPs are functionally linked to cellular chaperon activities and maintain homeostasis of the cell. Recently, it has been reported that isoprene emission against environmental stress

contributes to thermotolerance in plants. Isoprene, made from dimethylallyl diphosphate by isoprene synthase, is an important biological process because it plays a large role in atmospheric chemistry (Thompson, 1992). Several studies have independently confirmed the hypothesis (Singsaas & Shakey, 2000; Shakey et al., 2001; Velikova & Loreto, 2005), plants make isoprene for thermotolerance, especially protection from damage caused by rapid temperature fluctuation (Shakey & Singsaas, 1995). The study of characterization of *Populous alba* isoprene synthase (*PalspS*) showed the gene expression was strongly induced by heat stress (40 °C) (Sasaki et al., 2005). We suggest that the introduction of exogenous isoprene synthase is considered as alternative methods for generating thermotolerance ginseng.

A few research studies have been conducted to improve its tolerance to environmental stress such as high temperature using molecular breeding technology. But the generation of transgenic ginseng plants shown tolerance for high-temperature stress has not been reported. One of several reasons still has been remain to understand the mechanism of environmental stress, which is intricate. The present study was aimed at

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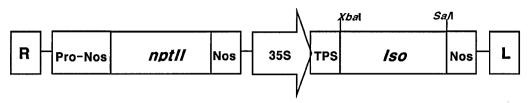


Fig. 1. A vector construction with pBI121mLPISO containing neomycin phosphotransferase (nptII) and Populous alba × Populous tremula isoprene synthase genes (ISO, Accession number: AJ294819). For P. ginseng transformation, Locations are indicated for the nopaline synthase (NOS term), transit peptide sequence (TPS), CaMV 35S promoters (35S), left (L) and right (R) borders.

exploiting transgenic plants for thermotolerance. This work firstly describes the possibility that the introduction of isoprene synthase in ginseng plants gene plays important role on thermotolerance.

MATERIALS AND METHODS

Plant material

Stratified *P. ginseng* seeds were immersed in 70% ethanol for 1 min, then in 3% sodium hypochlorite solution for 10 min, followed by three washes in sterile distilled water. Carefully dissection of the zygotic embryos from seeds was performed as described by Choi *et al.*, (1998). The cotyledons as an explant were prepared for *Agrobacterium* transformation.

Conditions of Co-cultivation

A single clone of the disarmed *A. tumefaciens* strain GV3101 containing the binary vector pBI121mLPISO (Fig. 1) consisting of neomycin phosphotransferase (*nptII*) gene as a selectable marker and *Populous alba* × *Populous tremula* isoprene synthase genes (*ISO*, Accession number: AJ294819) controlled under CaMV 35S promoter was selected. *A. tumefaciens* cultures were grown at 28°C on a gyratory shaker at 220 rpm in liquid YEP medium containing 50 mg/ ℓ kanamycin. The bacterial cells were collected by centrifugation for 5 min at 3000 rpm and resuspended in liquid inoculation medium (MS medium, Murashige & Skoog, 1962).

Production of transgenic plants

Excised cotyledons were dipped into the A. tumefaciens culture in liquid inoculation medium 20 min, blotted dry on steril filter paper, and incubated in the dark at 20 °C on callus induction medium (MS medium containing 3% sucrose, 1 mg/ ℓ 2,4-D, 0.8% agar). After 3-days of co-cultivation with A. tumefaciens, the cotyledons were transferred to fresh same medium except added 400 mg/ ℓ cefotaxim to eliminate bacteria. After 4 month of inoculation, embryogenic calli were subcultured on the MS medium supplemented with 100 mg/ ℓ kanamycin without plant growth regulators for selecting putative trans-

genic somatic embryos. Induced somatic embryos for germination were cultured on the MS medium containing 5 mg/ ℓ GA₃, 3% sucrose, 100 mg/ ℓ kanamycin, and 0.8% agar. Regenerated putative transgenic plantlets were grown in the growth chamber at 25 °C under 16 h photoperiod. Plantlets were then transferred to plastic pots containing soil, sand, and peat 4:4:3 (v/v) in a greenhouse. In vitro culture of transformed P ginseng plantlet (over 10 cm in height) was used for investigating high temperature stress. The plantlet was cultured in a bottle $(7 \times 12 \text{ cm})$ containing 70 m ℓ MS medium at 46 °C for 1 h under continuous light conditions. The treated plants were transferred to normal conditions (25 °C) for recovery from the stress. After 3 weeks of transfer to normal conditions, changes of morphological characterization were observed.

DNA analysis in transgenic plants

Genomic DNAs of the non-transformed leaves and transformed plants were isolated with DNeasy Plant Mini Kit (Qiagen). PCR was performed to detect the nptII and Iso genes in both the T-DNA of the Agrobacterium plasmid and genomic DNAs of the non-transformed leaves and hairy roots. The plasmid DNA of A. tumefaciens GV3101 strain containing pBI121mLPISO was used as a positive control. Primers for PCR were designed according to the DNA sequence of the nptII gene. The primer pair used for nptII gene amplification of an 700-bp fragment was: 5'-GAGGCTATTCGGCTAT-AGCTG-3' and 5'-ATCGGGAGCGGCGATACCGTA-3'; for the Iso gene amplification of an 810-bp fragment was: 5'-TCAATACAGTGATTGCCGTAATTCA-3' and 5'-TTATCTC TCAAAGGGTAGAATAGG-3'. The PCR mixture consisted of DNA, 0.4 mM of each primer, 0.2 mM of each dNTP, 2.5U Taq DNA polymerase (5U/ $\mu\ell$, Takara, Shiga, Japan), and 10 $\mu\ell$ 10× buffer in a final volume of 100 $\mu\ell$. The PCR was run under the following conditions: an initial denaturation at 94 °C for 5 min; denaturation at 94 °C for 1 min; primer annealing at 58°C for 1 min; elongation at 72°C for 1 min, 30 cycles; a final extension at 72 °C for 10 min. The amplification PCR products were examined by electrophoresis on a 1.5% (w/v) agarose gel.

RT-PCR analysis

First strand cDNA was synthesized using AMV reverse transcriptase (Promega) from 2 microgram total RNA plants that were isolated from the leaves of transgenic ginseng. For each reaction, 1 μℓ of the RT reaction was used as template for PCR. For normalization, an actin fragment amplified by the primer designed from *P. ginseng* was employed as an internal standard. The forward and reverse sequences used for PCR amplification of actin were 5'-GATGACATGGAAAAGAT TTGGCATC-3' and 5'-AAGGATGGCATGAGGGAGGGCGTAA-3', respectively. PCR conditions were as follows: 95 °C for 5 min followed by 95 °C, 1 min; 58 °C, 1 min; 72 °C, 1 min; 30 cycles. The amplification PCR products were examined by electrophoresis on a 1.5% (w/v) agarose gel.

RESULTS AND DISCUSSION

Excised cotyledon explants were used to facilitate infection with *Agrobacterium tumefaciens* and the induction of transformed primary callus. The formation of primary callus was induced on MS medium containing 1 mg/ ℓ 2,4-D, 3% sucrose, 0.8% agar, and 400 mg/ ℓ cefotaxim. To select transgenic lines, callus induced from explants was cultured on the medium containing 100 mg/ ℓ kanamycin. At high concentration of kanamycin, the growth of callus was not inhibited (Fig. 2A). After 2 month growth in the selection medium, callus was developed to somatic embryogenic cells (Fig. 2B). Putative transgenic somatic embryos were transferred to the regeneration medium containing 5 mg/ ℓ GA₃ to germinate them. After this treatment, a root and shoot were formed and most of them grew to plantlet (Fig. 2C, D). To avoid infection by fungus or

other microorganism, we used autoclaved soil including sand, and peat as described by Choi *et al.* (1998). Choi *et al.* (2003) reported that not all of the plantlets survived in soil that had not been autoclaved because of fungal infection. As shown in Fig. 2E, a transgenic ginseng plant was successfully trans-

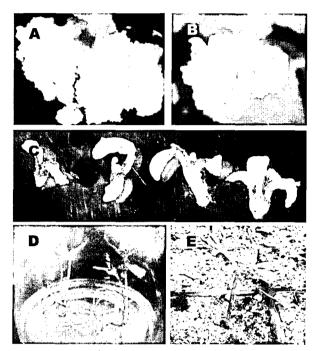


Fig. 2. Transgenic plants of *P. ginseng*. A, Embryogenic callus on MS medium supplemented with 3% sucrose, 1% agar, 400 mg/l cefotaxim, and 100 mg/l kanamycin; B, Surviving embryos on the medium containing high concentration of kanamycin (120 mg/l); C, Transgenic plantlets on the medium supplemented with 150 mg/l kanmycin.; D, Successful transfer to the soil.



Fig. 3. PCR analysis of the *nptll* and *lso* genes in *P. ginseng* transgenic plants. Lanes: P, plasmid DNA isolated from *A. tumefaciens* GV3101 strain as a positive control; T1, T2, T3, T4, and T5, DNA isolated from leaves of transgenic plants; N, DNA isolated from non-transformed leaves as a negative control.

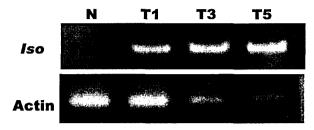


Fig. 4. RT-PCR analysis of *Iso* gene in leaves of wild-type and transgenic *P. ginseng*. Total RNA extracted from non-transformed leaves (N) and transgenic plants (T1, T3, and T5) were subjected to RT-PCR analysis. An actin fragment was amplified as internal loading control.



Fig. 5. Effects of high temperature (46°C) on transgenic ginseng plant. After high temperature treatment for 1 h, plants were transferred to normal conditions (25°C) for recovery from the stress. Visible differential damages in the leaves of a non-transformed plant (N) and a transformed plant (T).

ferred to the soil.

The PCR analysis of ginseng transgenic plants showed that they contained the *nptII* and *Iso* genes (Fig. 3). Amplification with the *nptII* primers showed a 740-bp band for the transgenic plants (lane T1-T5) and for the pBI121mLP*ISO* plasmid (lane P), while no band was observed for the untransformed plants (lane N). When the *Iso* primers were used for amplification, a 800-bp band was visualized only for transgenic plant clones (lane T1-T5). This result demonstrated that transformation of ginseng was confirmed by polymerase chain reaction analysis.

To investigate whether *Iso gene* transcription is tissue specific, total RNA was isolated from leaves of transformed ginseng plantlets. For reverse transcriptase-polymease chain raction (RT-PCR) analysis, we designed two primers to amplify a 800-bp fragment corresponding to *Populous alba* ×

Populous tremula isoprene synthase cDNA. As shown in Fig. 4, only *Iso* gene transcripts were observed in leaves of transgenic plants, but were not detected in a non-transgenic plant. RT-PCR result demonstrated the foreign isoprene synthase gene in three transgenic plant lines (T1, T3, and T5) which were expressed at the transcriptional level.

When whole plants of transgenic ginseng were exposed to high temperature at 46 °C for 1 h, a non-transformed plant was wilted from heat shock, whereas a transgenic plant appeared to remain healthy (Fig. 5). This result revealed that transgenic plants exhibited a tolerance to high temperature. Several genes for environmental stress of high temperature have been employed to generate transgenic plants. Tang et al. (2006) demonstrated enhanced tolerance to high temperature by introducing the genes, which consist of Cu/Zn superoxide dismutase and ascorbate peroxidase expressed in chloroplasts under the control of an oxidative stress-inducible SWPA2 promotor. in potato. Although our first attempt could show the possibility that isoprene synthase gene plays important role on thermotolerance in ginseng plants, further experiments should be performed to exactly demonstrate thermotolerance of transgenic ginseng.

ACKNOWLEDGEMENT

This study was supported by the Post Doctoral Course Program of the National Institute of Crop Sciences, Rural Development Administration, Republic of Korea.

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