## Enhanced Production of Protopanaxdiol (ginsenoside) in Roots of *Panax ginseng* Overexpressing Squalene Sythase Gene

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## **Objectives**

Panax ginseng is an important medicinal plants and pharmaco-logically active component has been accepted as triterpenoid (ginsenoside). Squanene synthase gene represents a putative branch point capable of diverting carbon flow to the biosynthesis of non-saponifiable sterols and saponifiable triterpens. The squanene synthase gene of Panax ginseng could play an important role in triterpene (ginsenoside) production. Here we reported the alteration of ginsenoside contents of Panax ginseng by introducing squanene synthase gene.

## **Materials and Methods**

- 1. Plant materials: Panax ginseng
- 2. Cloning of the squalene synthase gene from *Panax ginseng*: Screned from EST cDNA library.
- 3. Genetic transformation: *Agrobacterium tumefacins* harboring ginseng squalene synthase gene
- Analysis of transgenic plantlets: PCR, Southern, RT-PCR and Nothern analysis.
- 5. Ginsenoside and squalene analysis: HPLC and GC.

## **Results and Discussion**

- 1. Cloning of the squalene synthase gene: A full cDNA of SQ gene was isolated by EST. The nucleotide of SQ gene was 1476 bp and the putative ORF encodes a polypeptide of 415 amino acid residues. A deduced amino acid sequence showed similarity with *Arabidopsis thalina* (75.78%) and *Nicotiana tabacum* (81.45%). A southern analysis indicates that at least 4 copies of squalene synthase genes exist in genome of *Panax ginseng*.
- 2. Genetic transformation: Genetic transformation protocol was followed by the our previous report (Choi et al. 2003).
- Analysis of transgenic plantlets: Transgenic plants was confirmed by PCR, and GFP reaction, and Southern analysis. RT-PCR revealed that expression of SQ gene was enhanced in transgenic ginseng.
- 4. Bioassay for ginsenoside: Adventitious roots were obtained from transgenic and non-transgenic ginseng plants. Ginsenoside analysis of in vitro induced roots revealed that about 25% of ginsenoside of total one increased in transgenic ginseng roots, particularly notable increase only in protopanaxadiol (about double in ginsenoside Rb1). The preferential increase of protopanaxadiol group in transgenic Panax giseng overexpressing SQ gene was similar to the result of Methyl jasmonic acid treatment.

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