### **RESEARCH PAPER**



# *PgLOX6* encoding a lipoxygenase contributes to jasmonic acid biosynthesis and ginsenoside production in *Panax ginseng*

#### Shadi Rahimi<sup>1,2</sup>, Yu-Jin Kim<sup>3,4,\*</sup>, Johan Sukweenadhi<sup>1,3</sup>, Dabing Zhang<sup>4,5</sup> and Deok-Chun Yang<sup>1,3,\*</sup>

<sup>1</sup> Graduate School of Biotechnology and Ginseng Bank, College of Life Sciences, Kyung Hee University, Yongin, 446–701, Republic of Korea
<sup>2</sup> Department of Crop Science, Chungbuk National University, Cheongju 361–763, Korea

<sup>3</sup> Department of Oriental Medicinal Biotechnology, College of Life Sciences, Kyung Hee University, Yongin, 446–701, Republic of Korea <sup>4</sup> State Key Laboratory of Hybrid Rice, Shanghai Jiao Tong University–University of Adelaide Joint Centre for Agriculture and Health,

School of Life Sciences and Biotechnology, Shanghai Jiao Tong University, Shanghai 200240, China

<sup>5</sup> School of Agriculture, Food and Wine, University of Adelaide, Waite Campus, Urrbrae, South Australia 5064, Australia

\* Correspondence: yujinkim@khu.ac.kr; dcyang@khu.ac.kr

Received 28 May 2016; Accepted 22 September 2016

Editor: Qiao Zhao, Tsinghua University

## Abstract

Ginsenosides, the valuable pharmaceutical compounds in Panax ginseng, are triterpene saponins that occur mainly in ginseng plants. It was shown that in vitro treatment with the phytohormone jasmonic acid (JA) is able to increase ginsenoside production in ginseng plants. To understand the molecular link between JA biosynthesis and ginsenoside biosynthesis, we identified a JA biosynthetic 13-lipoxygenase gene (PgLOX6) in P. ginseng that promotes ginsenoside production. The expression of PgLOX6 was high in vascular bundles, which corresponds with expression of ginsenoside biosynthetic genes. Consistent with the role of PgLOX6 in synthesizing JA and promoting ginsenoside synthesis, transgenic plants overexpressing PgLOX6 in Arabidopsis had increased amounts of JA and methyl jasmonate (MJ), increased expression of triterpene biosynthetic genes such as squalene synthase (AtSS1) and squalene epoxidase (AtSE1), and increased squalene content. Moreover, transgenic ginseng roots overexpressing PgLOX6 had around 1.4-fold increased ginsenoside content and upregulation of ginsenoside biosynthesis-related genes including PgSS1, PgSE1, and dammarenediol synthase (PgDDS), which is similar to that of treatment with MJ. However, MJ treatment of transgenic ginseng significantly enhanced JA and MJ, associated with a 2.8-fold increase of ginsenoside content compared with the non-treated, non-transgenic control plant, which was 1.4 times higher than the MJ treatment effect on non-transgenic plants. These results demonstrate that PgLOX6 is responsible for the biosynthesis of JA and promotion of the production of triterpenoid saponin through up-regulating the expression of ginsenoside biosynthetic genes. This work provides insight into the role of JA in biosynthesizing secondary metabolites and provides a molecular tool for increasing ginsenoside production.

Key words: Ginsenoside, jasmonic acid, lipoxygenase, Panax ginseng, squalene, secondary metabolite, triterpene, vascular bundles.

## Introduction

*Panax* belongs to the family Araliaceae and contains at least 17 species (Kim *et al.*, 2015). Since ancient time, *Panax ginseng* (known as Korean or Asian ginseng) has been considered

as a healing drug and health tonic in China, Japan, and other Asian countries (Radad *et al.*, 2006). The pharmacological effects of ginseng are correlated with ginsenosides, bioactive

This is an Open Access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/3.0/), which permits unrestricted reuse, distribution, and reproduction in any medium, provided the original work is properly cited.

<sup>©</sup> The Author 2016. Published by Oxford University Press on behalf of the Society for Experimental Biology.