

(05-1-32)

## The *Arabidopsis* SUMO E3 ligase negatively regulates the salicylic acid-mediated defense pathway

Jiyoung Lee<sup>1</sup>, Kenji Miura<sup>2</sup>, JingBo Jin<sup>2</sup>, Chan Yul Yoo<sup>2</sup>, Ray A. Bressan<sup>2</sup>,  
Paul M. Hasegawa<sup>2</sup>, Jaesung Nam<sup>3</sup>, and Dae-Jin Yun<sup>1,\*</sup>

<sup>1</sup>Division of Applied Life Science, Graduate School of Gyeongsang National University, Jinju 660-701;

<sup>2</sup>Center for Plant Environmental Stress Physiology, Purdue University, West Lafayette, IN 47906, USA;

<sup>3</sup>Faculty of Plant Biotechnology, Dong-A University, Busan, 604-714, Korea

### Objectives

To find the function of *Arabidopsis* SUMO E3 ligase (*AtSIZ1*) in the plant defense response pathway.

### Materials and Methods

#### 1. Material

Plant-*Arabidopsis thaliana* plants (ecotype C24), and T-DNA insertional mutated alleles of *AtSIZ1*

#### 2. Methods

Pathogen test, Northern and microarray analysis, RT-PCR analysis

### Results and Discussion

Small Ubiquitin-like Modifier (SUMO) family proteins covalently attached to other proteins as post-translational modifications. SUMO modifies many proteins that participate in diverse cellular processes including transcriptional regulation, nuclear transport, maintenance of genome integrity, and signal transduction. Systemic acquired resistance (SAR) is a plant immune response induced after a local infection by necrotizing pathogen. Here, we show that the *Arabidopsis* SUMO E3 ligase (*AtSIZ1*) regulates pathogen defense response. Mutation in the *SIZ1* gene showed resistant to virulence bacteria strain *Pseudomonas syringae* DC3000. Northern and microarray analysis reveals that pathogenesis-inducible genes are constitutively expressed in *siz1* seedlings without pathogen treatment. However, jasmonate induced the expression of PDF1.2 to the same extent in *siz1* and W.T seedlings. High temperature (28°C) suppresses SA-mediated defense response pathway. When *siz1* plants were grown at 28°C, they show similar morphology to that of wild-type and PR gene expression level. Together, these results suggest that *AtSIZ1* regulates SA-mediated defense pathway.

[Supported by EB-NCRC and Biogreen 21 program]