

## Supplementary data

# Action of multiple rice $\beta$ -glucosidases on abscisic acid glucose ester

Manatchanok Kongdin<sup>1</sup>, Bancha Mahong<sup>2</sup>, Sang- Kyu Lee<sup>2</sup>, Su- Hyeon Shim<sup>2</sup>, Jong-Seong Jeon<sup>2,\*</sup>, and James R. Cairns<sup>1,\*</sup>

<sup>1</sup> School of Chemistry, Institute of Science, Center for Biomolecular Structure, Function and Application, Suranaree University of Technology, Nakhon Ratchasima 30000, Thailand

<sup>2</sup> Graduate School of Biotechnology and Crop Biotech Institute, Kyung Hee University, Yongin 17104, Korea

\* Correspondences: JJ, jjeon@khu.ac.kr, Tel.: +82 31 2012025 and JKC, cairns@sut.ac.th, Tel: +66 44 224304, Fax: +66 44 224185

**Supplementary Table S1.** PCR primers used in this study.

**Supplementary Figure S1** Subcellular localization of Os4BGlu10-GFP and Os4BGlu13-GFP in coleoptile cells of transgenic rice plants

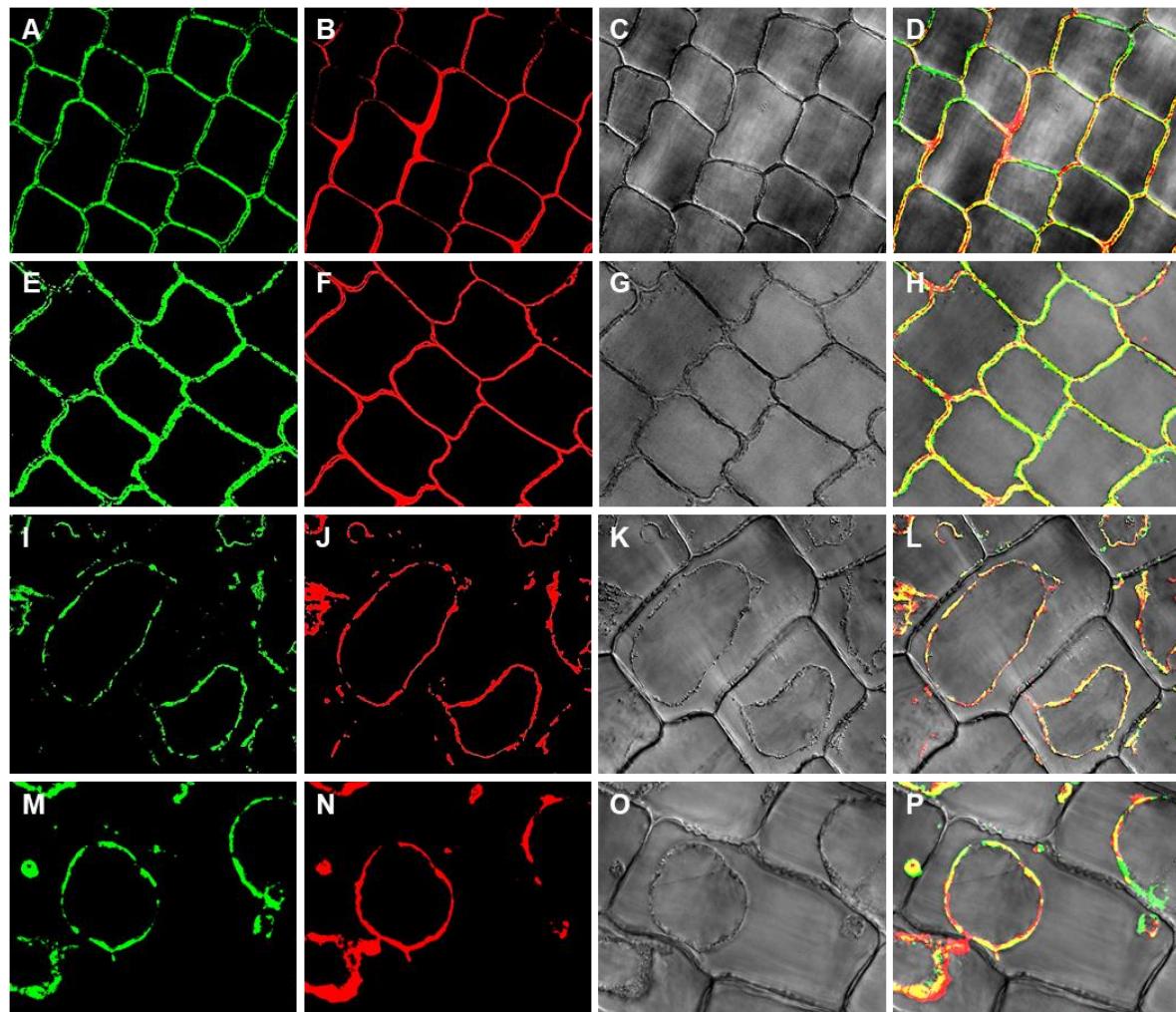
**Supplementary Figure S2** Root and shoot lengths of seedlings of wild type Arabidopsis and transgenic lines expressing rice  $\beta$ -glucosidases under control conditions.

**Supplementary Figure S3.** Sodium dodecyl sulfate polyacrylamide gel electrophoresis analysis of *E. coli*-expressed Os4BGlu12 and Os4BGlu13 purified by IMAC and cut with enterokinase.

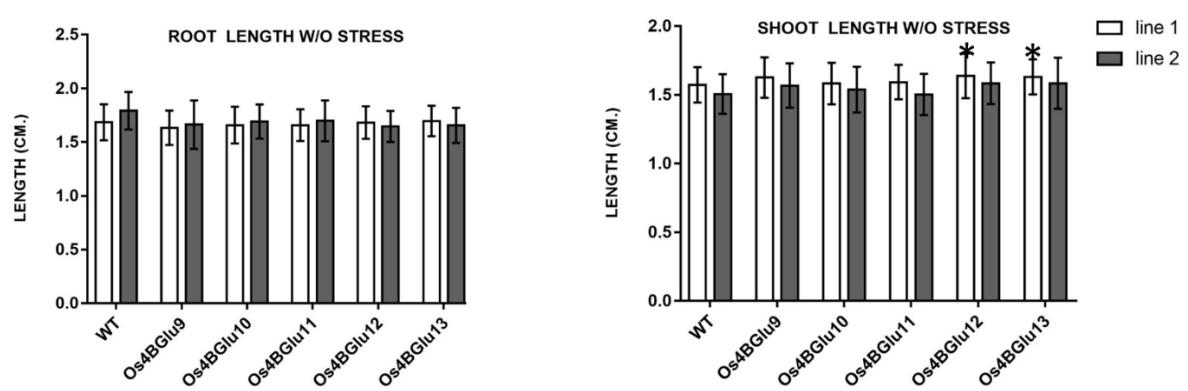
**Supplementary Table S1.** PCR primers used in this study.

Primer name	Sequence (5'→3')
Os4BGlu9 cloning F	5'-CACCGAATTGCAGGTCAAGCTTCGT-3'
Os4BGlu9 cloning R	5'-ACACATTGGGATTAAGTGCCAA-3'
Os4BGlu10 cloning F	5'-CACCCACTCTTCTGTCTATGTAGG -3'
Os4BGlu10 cloning R	5'-CATGCAAGAACAGAGGAGAGTGAG-3'
Os4BGlu11 cloning F	5'-TCTTCTGTCTATGTAGGTGATG-3'
Os4BGlu11cloning R	5'-CTTATATACTATTCGATGTTG-3'
Os4BGlu12 cloning F	5'-GTTGATGGCACCAAACATTG-3'
Os4BGlu12 cloning R	5'-AAGCAATGAGTGCAATGGTGC-3'
Os4BGlu13 cloning F	5'-CACCGCGGGTCACACACACATACT3'
Os4BGlu13 cloning R	5'-GGGAAGAGCAATGTGTGCAACT-3'
GFPBGl9F	5'-CACCATGGCGGTTGCCGGGGCAGTGGCGA-3'
GFPBGl9R	5'-ATGTCTGTACTGGCAAACCTCT-3'
GFPBGl10F	5'-CACCATGGCGGTTGCAGGTGCAATGGTGAT-3'
GFPBGl10R	5'-TTTCCGGAGGAACCTCTTGAACCA-3'
GFPBGl11F	5'-CACCATGGCGGTTGCAGGGCAATGGT-3'
GFPBglu11R	5'-ATTAGAAAAGGCATTGTATGCA-3'
GFPBGl12F	5'-CACCATGGCAGATGGAAGTCTGAGGGTGG-3'
GFPBGl12R	5'-TTTCAGGAGGAACCTCTTGAACCAAT-3'
GFPBGl13F	5'-CACCATGGCAGCTGCAGGGAAAGTGGTGA-3'
GFPBGl13R	5'-TTTCTGGAGGAACTCCTTGAACC-3'
RTBGl9F	5'-GCACTGGAAGATAAAATACAACGGATT-3'
RTBGl9R	5'-TGATATTCTCTGTACAACCGAA-3'
RTBGl10F	5'-TGAAATCAATAATAAGACCATGCGAC-3'
RTBGl10R	5'-TCATTCCGGAGGAACCTCTTGAACCA-3'
RTBGl11F	5'-CACCTCTCTGTCTATGTAGGTGATG -3'
RTBGl11R	5'-TCAATTAGAAAAGGCATTGTATGCAAC-3'
RTBGl12F	5'-CACCGTTGATGGCACCAAACATTG-3'
RTBGl12R	5'-TCATTCAGGAGGAACCTCTTGAACCAAT-3'
RTBGl13F	5'-TGAATTCAACAATAAGACCTTACCACT-3'
RTBGl13R	5'-TCATTCTGGAGGAACTCCTTGAACC-3'

**Supplementary Figure S1** Subcellular localization of Os4BGl10-GFP and Os4BGl13-GFP in coleoptile cells of transgenic rice plants. (A-H) Coleoptile cells of Os4BGl10-GFP (A-D) and Os4BGl13-GFP (E-H). A and E, GFP fluorescence. B and F, FM4-64 red signal. C and G, Bright field. D and H, Merged. (I-P) Plasmolysed coleoptile cells of Os4BGl10-GFP (I-L) and Os4BGl13-GFP (M-P). I and M, GFP fluorescence. J and N, FM4-64 red signal. K and O, Bright field. L and P, Merged.



**Supplementary Figure S2** Root and shoot lengths of seedlings of control Arabidopsis and transgenic lines expressing rice  $\beta$ -glucosidases under control conditions. Control and transgenic Arabidopsis expressing rice phytohormone  $\beta$ -glucosidases Os4BGl9-13 were grown on  $\frac{1}{2}$  MS plates for 7 d and transplanted to  $\frac{1}{2}$  MS for 5 d. To quantify root and shoot growth inhibition, root and shoot lengths were measured in three independent experiments with 20 plants each



**Supplementary Figure S3.** Sodium dodecyl sulfate polyacrylamide gel electrophoresis analysis of *E. coli*-expressed Os4BGl12 and Os4BGl13 purified by IMAC and cut with enterokinase. Lane 1: Low Molecular Weight Protein Marker (GE Healthcare); Lane 2: Os4BGl12; and Lane 3: Os4BGl13.

