

Available online at www.sciencedirect.com





Biochemical and Biophysical Research Communications 304 (2003) 215-216

www.elsevier.com/locate/ybbrc

Erratum

Erratum to "Molecular cloning and mRNA expression analysis of a novel rice (Oryza sativa L.) MAPK kinase kinase, OsEDR1, an ortholog of Arabidopsis AtEDR1, reveal its role in defense/stress signalling pathways and development" [Biochem. Biophys. Res. Commun. 300 (2003) 868–876]**

Jung-A. Kim,^a Ganesh K. Agrawal,^b Randeep Rakwal,^c Keon-Seon Han,^a Kyung-Nam Kim,^a Choong-Hyo Yun,^d Sunggi Heu,^d Sook-Young Park,^e Yong-Hwan Lee,^e and Nam-Soo Jwa^{a,*}

^a Department of Molecular Biology, College of Natural Science, Sejong University, Seoul 143-747, Republic of Korea ^b Research Laboratory for Agricultural Biotechnology and Biochemistry (RLABB), GPO Box 8207, Kathmandu, Nepal ^c Molecular and Microbial Ecology Research Group, Institute for Biological Resources and Functions, National Institute of Advanced Industrial Science and Technology (AIST),

Central 6, 1-1-1 Higashi, Tsukuba, Ibaraki 305-8566, Japan ^d Plant Pathology Division, National Institute of Agricultural Science and Technology, Rural Development Administration, Suwon 441-707, Republic of Korea

In Fig. 3A and Figs. 5A and D, the Northern images are missing. For the reader's convenience, the correct figures are reproduced here with their legends.

E-mail address: nsjwa@sejong.ac.kr (N.-S. Jwa).

^e School of Agricultural Biotechnology, Seoul National University, Suwon 441-744, Republic of Korea

[★] DOI of original article: 10.1016/S0006-291X(02)02944-3.

^{*}Corresponding author. Fax: +82-2-3408-3661.

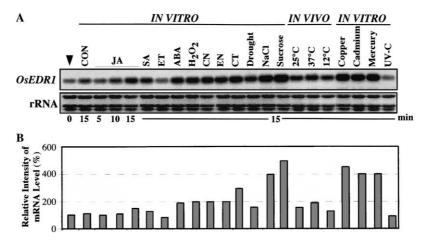


Fig. 3. Activation of the OsEDR1 transcript within 5 and 15 min in response to multiple stresses in rice seedling leaves. (A) Total RNA was extracted from leaf segments treated with $100 \,\mu\text{M}$ each of JA, SA, ABA, CN, EN, and copper, cadmium, and mercury; 0.1% CT; $1 \,\text{mM}$ ET; $10 \,\text{mM}$ H₂O₂; drought; $150 \,\text{mM}$ each of NaCl and sucrose; and UV-C irradiation (in vitro). CON refers to the wounding by cut control. Intact seedlings (in vivo) were placed at various temperatures ($25/37/12 \,^{\circ}\text{C}$). Arrowhead refers to sampling at the start of the experiments. Treatments were done under continuous light ($150 \,\mu\text{mol m}^{-2}\,\text{s}^{-1}$). The stresses and sampling times are indicated above and below each lane, respectively. The blots were hybridized to a [α - 32 P]dCTP-labeled OsEDR1 cDNA probe and single hybridizing band of ca. $3.4 \,\text{kbp}$ is shown. Equal loading ($20 \,\mu\text{g}$) was confirmed by staining of membranes with methylene blue and a part of rRNA is shown. Northern analysis was carried out as described in Materials and methods. (B) The histograms show relative intensity of mRNA level in percentage considering the constitutively expressed $OsEDR1 \,\text{mRNA}$ level at $0 \,\text{min}$ as 100%.

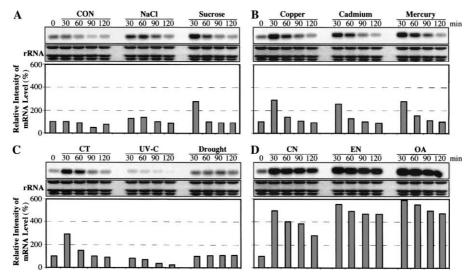


Fig. 5. Differential and transient regulation of the OsEDR1 mRNA expression in leaves upon treatment with other environmental stressors (A–D). Leaf segments were irradiated with UV-C, treated with NaCl, sucrose, heavy metals copper, cadmium, and mercury, CT, UV-C irradiation, drought, CN, EN, and OA. Except for OA (1 μ M), concentrations are as given in Fig. 3 and equal loading, hybridization, and histograms are as described in Fig. 3.