

Article Addendum

NaCl-induced expression of *ASCORBATE PEROXIDASE 8* in roots of rice (*Oryza sativa* L.) seedlings is not associated with osmotic component

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Abbreviations: ABA, abscisic acid; APx, ascorbate peroxidase; ELISA, enzyme-linked immunosorbent assay; Man, mannitol; *OsRab 16A*, the abscisic acid responsive gene; ROS, reactive oxygen species

Key words: abscisic acid, ascorbate peroxidase, *Oryza sativa*, osmotic effect, salt stress

Ascorbate peroxidase (APx; EC 1.11.1.11) plays an important role in scavenging the toxic effects of H₂O₂ in higher plants. Eight types of APx have been described for *Oryza sativa*: two cytosolic (*OsAPx1* and *OsAPx2*), two putative peroxisomal (*OsAPx3* and *OsAPx4*), and four chloroplastic isoforms (*OsAPx5*, *OsAPx6*, *OsAPx7* and *OsAPx8*). We have recently demonstrated that Na⁺ but not Cl⁻ is required for the NaCl-induced expression of *OsAPx8* in rice roots. Evidence is also provided to show that Na⁺-induced expression of *OsAPx8* is mediated through an accumulation of ABA. In addition to its known component of ion toxicity, there is an osmotic effect resulting from salt concentration in the soil. Here we show that ABA level but not *OsAPx8* expression was enhanced at a concentration of mannitol iso-osmotic with 150 mM NaCl suggests that NaCl-enhanced *OsAPx8* expression is not associated with osmotic component.

Soil salinity is one of the major abiotic stresses affecting plant growth and productivity globally. A consequence of salt stress in plants is the excessive generation of ROS.^{1,2} Plants have evolved both enzymatic and non-enzymatic mechanisms for ROS scavenging.³ APx (EC 1.11.1.11) plays a crucial role in the detoxification of cellular H₂O₂, the toxic product of superoxide dismutase.

Expression of APx has been reported to be enhanced in plants by NaCl.^{2,4-7} In contrast, the expression of APx in sweet potato leaves has been shown to be reduced by NaCl.⁸ Moreover, it has been demonstrated that expression of APx was not affected by NaCl.⁹⁻¹¹

The plant hormone ABA is a sesquiterpenoid derived from xanthophyll and appears to influence several physiological and developmental events.¹² The level of ABA in plants increases upon

their exposure to NaCl stress.¹³⁻¹⁵ It has been shown that ABA treatment increased the expression of APx in pea, rice and sweet potato leaves^{8,16,17} but had no effect on APx gene expression in *Brassica napus*¹⁸ and BY-2 cells.¹⁹

When growing in saline soil, roots have to cope with two types of stress. The first of these is ionic stress induced by changes in the concentrations of Na⁺, Cl⁻ or both in the root-growing medium and within root tissues. The second is an osmotic stress resulting from salt concentration in the soil that results in lowered water potential and a consequent loss of cell turgor in roots.

Recently, we demonstrated that NaCl-enhanced expression of *OsAPx8* in rice roots is mediated through an accumulation of ABA.¹⁵ Evidence is also provided to show that Na⁺ but not Cl⁻ is required for enhancing *OsAPx8* expression, APx activity and ABA accumulation in rice roots treated with NaCl.¹⁵ To understand the role of osmotic effect in NaCl-induced expression of *OsAPx8*, rice roots were treated with 276 mM mannitol (Man), which is a concentration iso-osmotic with 150 mM NaCl. A key observation of this study is that ABA accumulation but not *OsAPx8* expression was observed in rice roots treated with Man (Fig. 1). In this study, the level of ABA was judged by the transcript level of *OsRab 16A*, an ABA responsive gene^{20,21} (Fig. 1). Using ELISA method, Man treatment also resulted in an accumulation of ABA (data not shown) in rice roots. ABA is known to enhance the expression of *OsAPx8* in rice roots.¹⁹ *OsAPx8* is a putative thylakoid isoform.⁷ It seems that ABA levels in subcellular compartments of rice roots treated with NaCl differ from those treated with Man. The results obtained so far seem to conclude that NaCl-inducing *OsAPx8* expression in rice roots is associated with ionic but not osmotic component.

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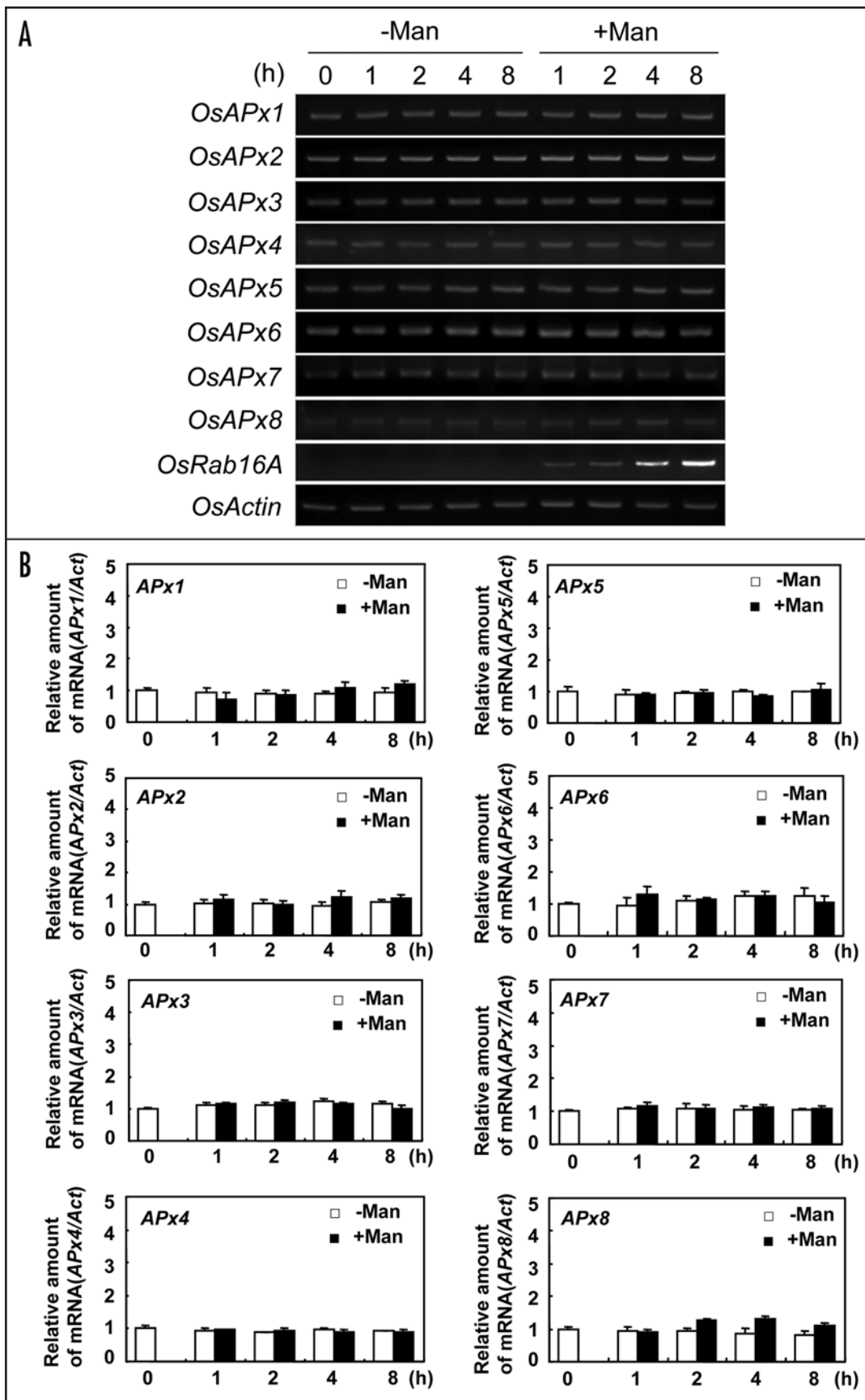


Figure 1. Changes in mRNA levels for *OsAPx* genes and *OsRab 16A* in rice roots in the presence or absence of mannitol (Man). Two-day-old rice seedlings were treated with or without Man (276 mM). Semi-quantitative RT-PCR for *OsAPx* genes and *Rab 16A* was performed. The values of mRNA for the *OsAPx* genes were adjusted by the corresponding amount of *OsActin* mRNA for equality of loading. After the adjustment by *OsActin*, the reaction with the roots without Man was treated as the normalized reference, with a value of one, for determination of the relative amount of mRNA of *OsAPx* genes. Bars show means \pm SE ($n = 4-6$). Statistical differences between measurements on different times were analysed using the LSD test. No significant differences were observed between -Man and +Man at $p > 0.05$.

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