ARTICLE ADDENDUM

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Drought stress promotes xylem differentiation by modulating the interaction between cytokinin and jasmonic acid

Geupil Jang and Yang Do Choi 🝺

Department of Agricultural Biotechnology and Research Institute of Agriculture and Life Sciences, Seoul National University, Seoul, Korea

ABSTRACT

Drought stress provokes jasmonic acid (JA) signaling, which mediates plant stress responses; moreover, growing numbers of studies suggest that JA is involved in the modulation of root development under drought stress. Recently, we showed that JA promotes differentiation of xylem from procambial cells in Arabidopsis roots. Further molecular and genetic approaches revealed that the effect of JA on xylem development is caused by suppression of cytokinin responses, suggesting that JA antagonistically interacts with cytokinin to modulate xylem development. Here, we showed that, similar to JA, drought stress promotes xylem development. This suggests that the antagonistic interaction between JA and cytokinin is involved in drought-mediated xylem development, a hypothesis supported by the observation that drought stress promotes xylem development, and the antagonistic interaction between JA and cytokinin is deeply involved in this process.

Plants dynamically coordinate their development and defenses in response to environmental stresses such as drought, and modulation of growth may help plants survive environmental stresses.¹⁻⁴ Drought stress affects root development and growth and our recent study revealed that drought stress promotes the differentiation of xylem from meristematic procambial cells in root vascular tissues.⁵ (Fig. 1) In Arabidopsis roots, xylem cells develop in a single axis.⁶ Wild-type plants grown in polyethylene glycol (PEG)-containing medium designed to simulate drought stress conditions formed extra xylem adjacent to the xylem axis, in contrast to wild-type plants grown in normal conditions. Quantification of the number of xylem cells showed that roots of wild-type plants grown in normal conditions developed approximately 4 xylem cells but wild-type plants grown on PEG-containing medium developed approximately 5 xylem cells, suggesting that the PEG-mediated stress promotes xylem development.

The interaction between hormones, especially between the hormones that govern stress responses and development, regulates developmental flexibility under stress conditions.^{3,4,7-11} Cytokinin is an essential hormone controlling cell proliferation and root xylem development.^{6,12,13,14} For example, *wooden leg* (*wol*) mutants with severe defects in cytokinin signaling display all-xylem phenotypes in their vascular tissues, and the mutant plants that lack expression of Type-B *ARABIDOPSIS RESPONSE REGULATORs* (*ARRs*) such as *ARR1* and *ARR12*

produce more xylem.^{6,13,14} These observations indicate that cytokinin is a key negative regulator inhibiting xylem development in root vascular tissues. The finding that exogenous cytokinin treatment suppresses xylem formation supports this hypothesis (Fig. 2A). Together with previous findings that expression of the genes responsible for cytokinin responses tended to be downregulated by environmental stresses,¹⁵⁻¹⁷ these findings suggest that modulation of the cytokinin response is deeply involved in xylem development under PEG-mediated drought stress conditions.

In contrast to cytokinin, JA and its related stresses such as drought and oxidative stress promote xylem differentiation,^{5,18} in agreement with our finding that JA promotes formation of extra xylem (Fig. 2A). This indicates that cytokinin and JA have opposite functions in xylem development and suggests that cytokinin and JA antagonistically interact in root xylem development. When the number of xylem cells was quantified, the plants co-treated with cytokinin and JA had more xylem cells compared with the plants treated with cytokinin alone, but fewer xylem cells compared with the plants treated with JA alone (Fig. 2B). These results support the idea that cytokinin and JA antagonistically interact in root xylem development. The results also suggest that PEG-mediated drought stress promotes xylem development by modulating the JA and cytokinin responses. To explore this, we analyzed JA and cytokinin responses in roots grown on PEG-containing medium by quantifying expression levels

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CONTACT Yang Do Choi Schoiyngd@snu.ac.kr Department of Agricultural Biotechnology and Research Institute of Agriculture and Life Sciences, Seoul National University, Seoul 151-921, Korea.

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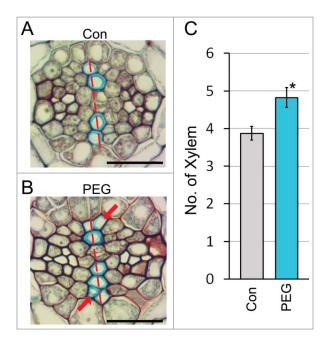


Figure 1. PEG-mediated drought stress promotes xylem development in roots. Xylem development of wild-type (Col-0) roots grown in PEG-untreated (A) and -treated (B) conditions. Four-day-old seedlings germinated on normal Murashige and Skoog (MS) media were transferred to the control media (1/2 MS medium, water potential = -0.25 MPa) or PEG-containing media (water potential = -0.70 MPa), respectively. After 8 days, the roots collected from these plants were sectioned transversly and stained with toluidine blue. Root sections were taken from the root maturation zone 2 mm above the transition zone, which is located between the meristem and basal differentiation region. The red arrows and dotted lines indicate extra xylem cells and the xylem axis. (C) Quantification of the number of xylem cells formed in these roots (n > 15). Error bars represent S.E. and asterisks indicate statistically significant differences between the corresponding samples and their control (p < 0.01, Student's t-test). Scale bar = $20 \,\mu$ m.

of JA-induced genes such as LOX2 and MYC2 and cytokinininduced genes such as ARR15 (Fig. 2C). Transcript levels of JA-induced genes such as LOX2 and MYC2 increased while transcript levels of cytokinin-induced genes such as ARR15 decreased in response to the stress. Unlike ARR15, expression of AHP6 whose expression is downregulated by cytokinin⁶ was upregulated by the stress. These findings indicated that the PEG-mediated drought stress activates JA responses and suppresses cytokinin responses, suggesting that the PEG-mediated drought stress affects xylem development by modulating the JA and cytokinin responses. The previous result showing that JA-responsive transcription factor MYC2 promotes xylem differentiation by activating expression of AHP6, a negative regulator of cytokinin response⁵ partially supports this. Collectively, our findings suggest that the antagonistic interaction between cytokinin and JA mediates xylem differentiation under drought stress.

Growing numbers of studies have proposed that JA antagonistically interacts with cytokinin in various aspects of plant physiology and development. For example, JA strongly inhibits cytokinin-induced callus growth¹⁹ and suppresses the effect of cytokinin on chloroplast development and the plant defense system.^{20,21} A recent study of circadian stress responses supports the antagonistic interaction between JA and cytokinin.²² These suggest that the JA–cytokinin interaction is largely involved in the modulation of plant physiology and development in response to environmental stresses such as drought.

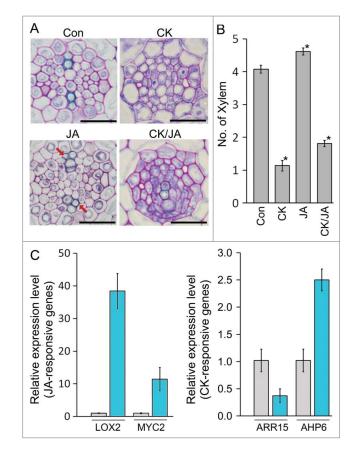


Figure 2. JA and cytokinin responses in PEG-treated roots. (A) Xylem development of Col-0 roots grown in the indicated conditions for 7 days (cytokinin [CK], 50 nM BAP; JA, 10 μ M MeJA; CK/JA, 10 μ M MeJA and 50 nM BAP). The red arrows indicate extra xylem. (B) Quantification of the number of xylem cells formed in these roots (n>21). Error bars represent S.E. and asterisks indicate statistically significant differences between the corresponding samples and the control (p < 0.01, Student's t-test). (C) Expression of JA-responsive and cytokinin-responsive genes in the roots grown in the PEG-untreated (grey) and -treated (blue) conditions. Total RNAs were extracted from 8-day-old Col-0 roots grown in PEG-untreated (water potential = -0.25 MPa) and -treated (water potential = -0.70 MPa) conditions. Error bars represent S.D. Scale bar = 20 μ m.

However, the molecular mechanisms controlling the JA-cytokinin interaction are largely unknown. Further studies will expand our understanding of the molecular mechanisms underlying this process.

Disclosure of potential conflicts of interest

No potential conflicts of interest were disclosed.

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ORCID

Yang Do Choi (D) http://orcid.org/0000-0003-4706-5154

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