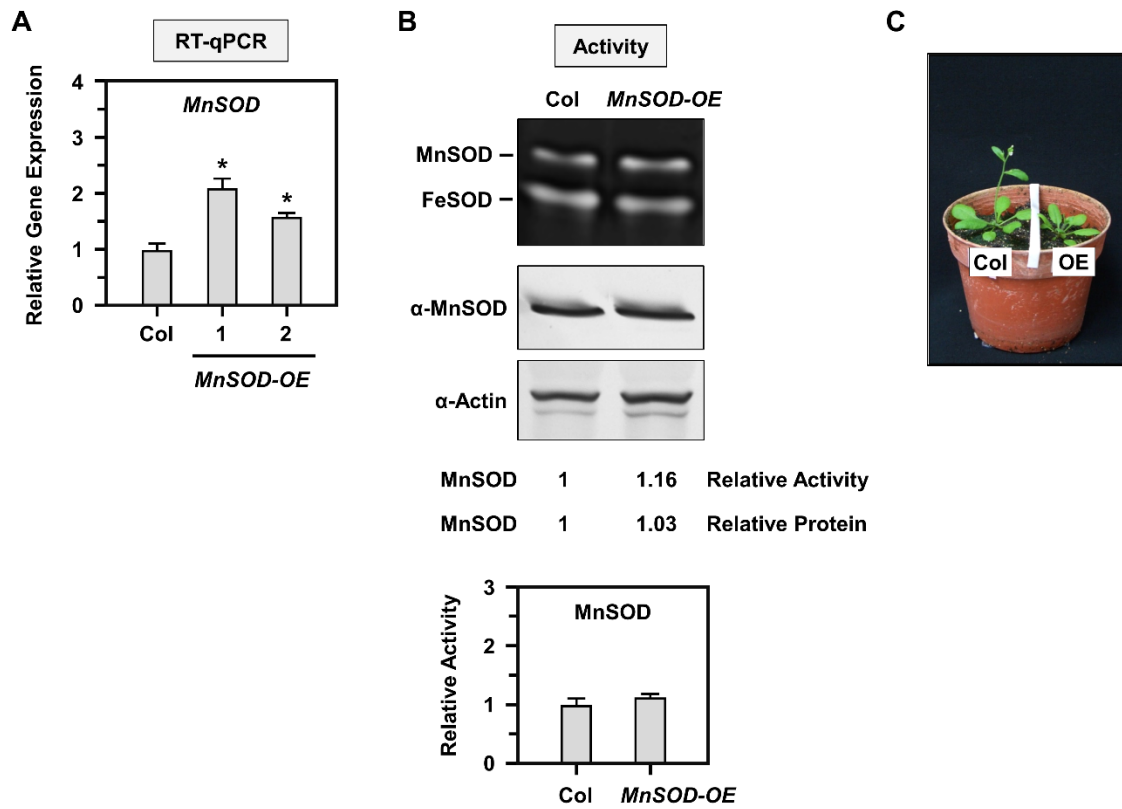


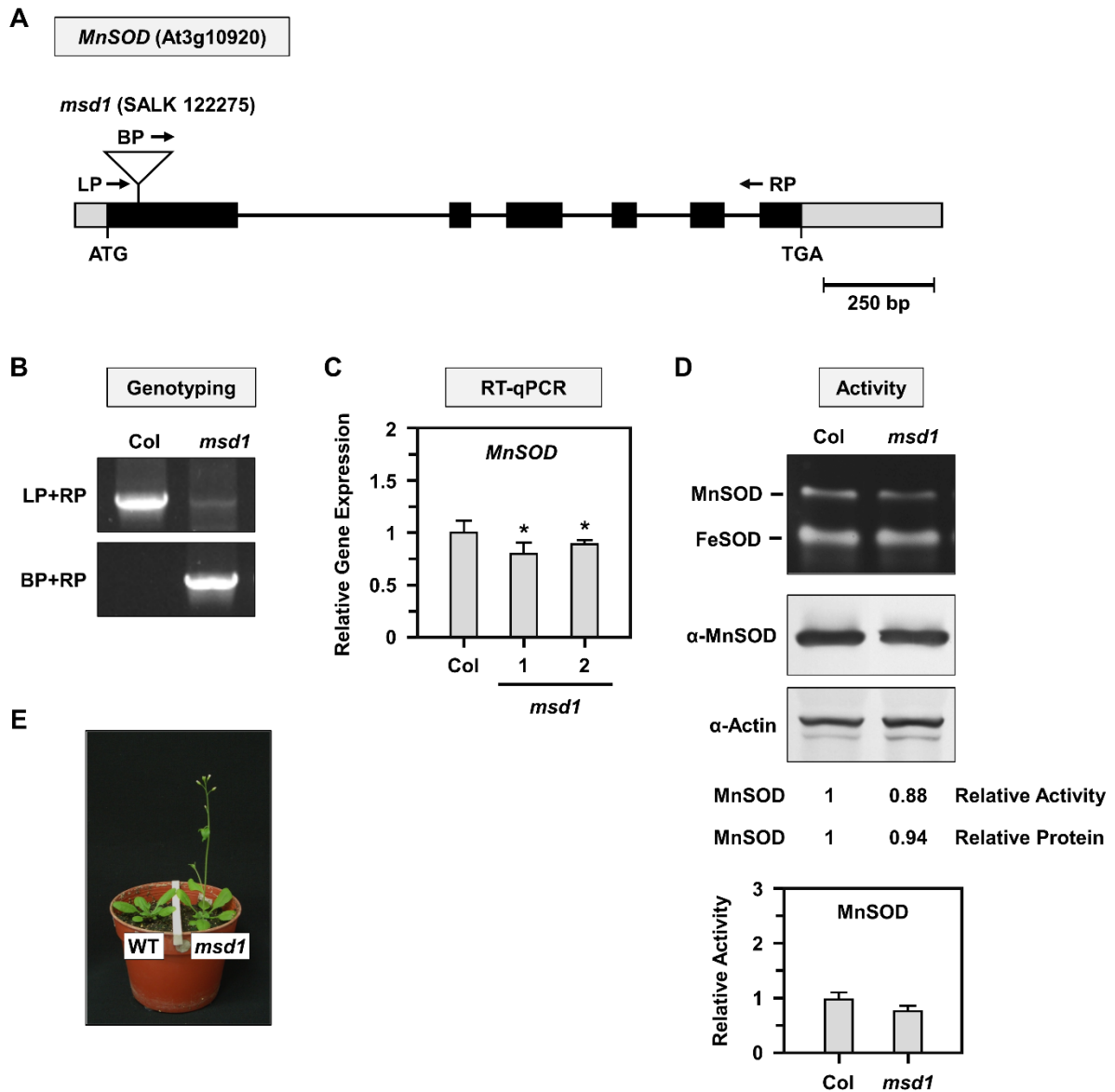
**Supplementary Table S1.** Primers for cloning, genotyping, RT-qPCR, and accession numbers of genes.

Purpose	Sequence (5'→3')	Primer Set
Construction of <i>AtMnSOD-OE</i>	<u>GAGCTC</u> TTTGGCGACCACTAGAAGGAGA	<u>SacI</u> - <i>AtMnSOD-OE-Fw</i>
	TCTAGA <u>AGCTGAGCTGGA</u> ACTGGTTCATCTC	<u>XbaI</u> - <i>AtMnSOD-OE-Rv</i>
Genotyping of <i>msd1</i>	CGATTCGTTGTGTAGCGAG	<i>AtMnSOD-Fw</i> (LP)
	TTCAGTTGTTTTCTTCTCATAAAC	<i>AtMnSOD-Rv</i> (RP)
T-DNA left border	ATTCAGTACATTA AAAACGTCCGCAAT	BP
RT-qPCR for <i>AtMTM1</i>	AGAGGACTGTTTCATGGGAATGG	<i>AtMTM1-RT-qPCR-Fw</i>
	ATACTTGACCACTTCGTAGAAGGAAAC	<i>AtMTM1-RT-qPCR-Rv</i>
RT-qPCR for <i>AtMTM2</i>	GAGCCATGACAATGACCACAAG	<i>AtMTM2-RT-qPCR-Fw</i>
	CTCTAGCCCCTGCTCCACTAAAC	<i>AtMTM2-RT-qPCR-Rv</i>
RT-qPCR for <i>AtMnSOD</i>	TGCCATTGACGCTCACTTTG	<i>AtMnSOD-RT-qPCR-Fw</i>
	TTGTCTAGTCCGAGCCACAC	<i>AtMnSOD-RT-qPCR-Rv</i>
RT-qPCR for <i>AtAOX1A</i>	GAATGTTCTGCTCCGGCTAT	<i>AtAOX1A-RT-qPCR-Fw</i>
	TCAGCACGAACAACCATCACA	<i>AtAOX1A-RT-qPCR-Rv</i>
Internal control of RT-qPCR	CCTGCGGTAATAACTGCATCT	<i>AtPP2A-RT-qPCR-Fw</i>
	CTTCACTTAGCTCCACCAAGCA	<i>AtPP2A-RT-qPCR-Rv</i>

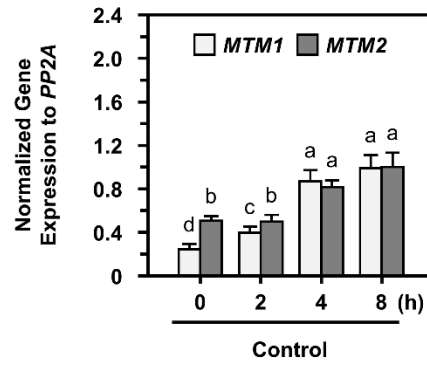
Sequence data in this article can be found in the TAIR database under the following accession numbers: *AtAOX1A* (At3G22370), *AtMnSOD* (*AtMSD1*; At3G10920), *AtMTM1* (At4G27940), *AtMTM2* (At2G46320), *AtPP2A* (At1G13320), and *AtUBQ10* (At4G05320).



**Supplementary Figure S1.** Characterizations of *AtMnSOD*-overexpressing (*MnSOD-OE*) plant. **(A)** *AtMnSOD* gene expression in Col and *MnSOD-OE* seedlings was analyzed by RT-qPCR. *PP2A* was used as internal control. *AtMnSOD* gene expression level in *MnSOD-OE* was measured relative to that in Col. Data are mean  $\pm$  SD of three biological replicates. \* Significant at  $P < 0.05$ . **(B)** In-gel SOD activity assay (top) and immunoblotting with  $\alpha$ -MnSOD and  $\alpha$ -Actin antibodies (bottom). Actin was used as a loading control. MnSOD activity and protein levels in *MnSOD-OE* were measured relative to that in Col. **(D)** Late-flowering phenotype of one-month-old *MnSOD-OE* plant.



**Supplementary Figure S2.** Characterizations of *AtMnSOD* T-DNA knockdown (*msd1*) mutant. **(A)** Schematic map of *AtMnSOD* gene (At3G10920), T-DNA insertion site, and primers (arrows) used for genotyping. **(B)** Genotyping by PCR in *msd1* was performed with DNA flanking sequence primers (LP and RP) and T-DNA border primer (BP). **(C)** The *AtMnSOD* gene expression level in Col and *msd1* mutant was analyzed by RT-qPCR. *PP2A* was used as internal control. *AtMnSOD* gene expression level in *msd1* was measured relative to that in Col. Data are mean  $\pm$  SD of three biological replicates. \* Significant at  $P < 0.05$ . **(D)** In-gel SOD activity assay (top) and immunoblotting with  $\alpha$ -MnSOD and  $\alpha$ -Actin antibodies (middle and bottom). Actin was used as a loading control. MnSOD activity and protein levels in *msd1* were measured relative to that in Col. **(F)** Early-flowering phenotype of one-month-old *msd1* mutant.



**Supplementary Figure S3.** Control experiment of *AtMTM1* and *AtMTM2* gene expression levels in Col. 14-day-old seedlings were incubated in 1/2 MS liquid medium without stressor treatment. *PP2A* was used as internal control. Data are mean  $\pm$  SD of three biological replicates. The statistical significances ( $P < 0.05$ ) are indicated as different letters (Duncan's multiple range test).