Corrigendum

ROS-mediated vascular homeostatic control of root-to-shoot soil Na delivery in *Arabidopsis*

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Since the publication of this article, the authors have discovered a small error in the labelling of Figure 6B; 'sss1-1/atrbohF-F3' should read 'rss1-1/atrbohF-F3'.

The correct Figure 6 is shown here.

The authors apologize for any inconvenience caused.

Figure 6 *rss1-1* is a novel mutant *AtCesA7* allele. (A) Soil-grown plants (genotypes as indicated) grown under control conditions (Control) and 7 days after salt treatment (Salt). Bar = 1 cm. (B) Chlorophyll content of soil-grown plants. Genotypes as indicated (salt treatments were as described in Figure 1A). (C) The mutation causing *rss1-1* phenotype. The top panel shows the 2-bp deletion (–CA; highlighted by the red arrow) that interrupts *AtCesA7* in *rss1-1* and *atrbohF-F3* are aligned against TAIR9. The bottom panel shows the genomic structure of *AtCesA7*. Blue boxes represent coding sequence, green boxes introns, open blue boxes untranslated regions (UTR). (D) Demonstration that *rss1-1* is a mutant *AtCesA7* allele. *atrbohF-F3/AtCesA7^{rx3-4}* is less sensitive to soil salinity compared to *atrbohF-F3* and shows a salt sensitivity similar to that of *atrbohF-F3/AtCesA7^{rss1-1}*. Bar = 1 cm.