

## Erratum to: TTG1-mediated flavonols biosynthesis alleviates root growth inhibition in response to ABA

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Figure 6c is missing in the original publication. The complete figure is reproduced on the following page.

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**Fig. 6** Growth phenotypes of wild-type, *TTG1-OX*, *PAP1-D* and *ttg1* seedlings grown on  $1/2 \times$  MS medium supplemented with  $10 \mu\text{M}$  ABA or  $300 \text{mM}$  mannitol. **a** Four-day-old seedlings were transferred to test medium and allowed to grow for 10 days before being photographed. The seedlings shown were typical of three separate experiments performed at  $23^\circ\text{C}$  ( $n = 15$  seedlings per experiment). **b** The numbers of LRs in these seedlings were counted under a microscope. The data shown are from three separate experiments performed at  $23^\circ\text{C}$  ( $n = 10$  per experiment). One-way ANOVA Tukey's test ( $P < 0.05$ ) was performed to indicate a significant difference in LR number between wild-type and mutants per treatment. Lower-case letters (*a*, *b*, *c*) were marked above the column to reflect significant differences. Vertical bars indicate standard error ( $n = 30$ ). **c** The transcript levels of the stress-marker genes *Cor15A* and *KIN1* were determined for wild-type and *TTG1-OX* #4 and #5 seedlings on normal  $1/2 \times$  MS medium. Several rounds of *RT-PCR* were performed, and *PCR* results from two experiments using different numbers of cycles are shown. *UBC21* was used as the internal loading control. Ten-day-old seedlings were grown on normal  $1/2 \times$  MS medium before total RNA extraction for *RT-PCR*

