

Supplementary Information

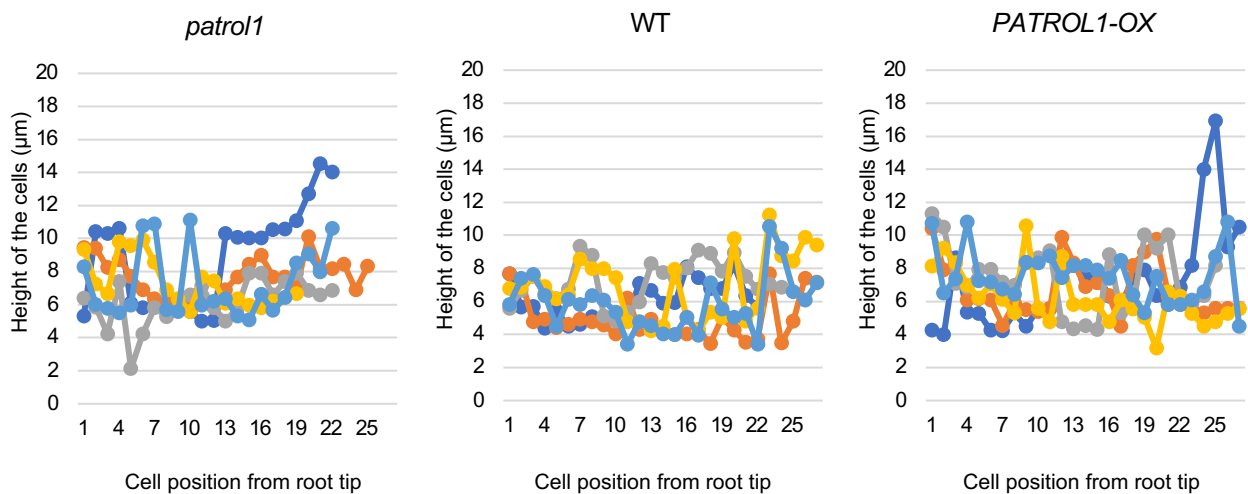
The PATROL1 function in roots contributes to the increase in shoot biomass

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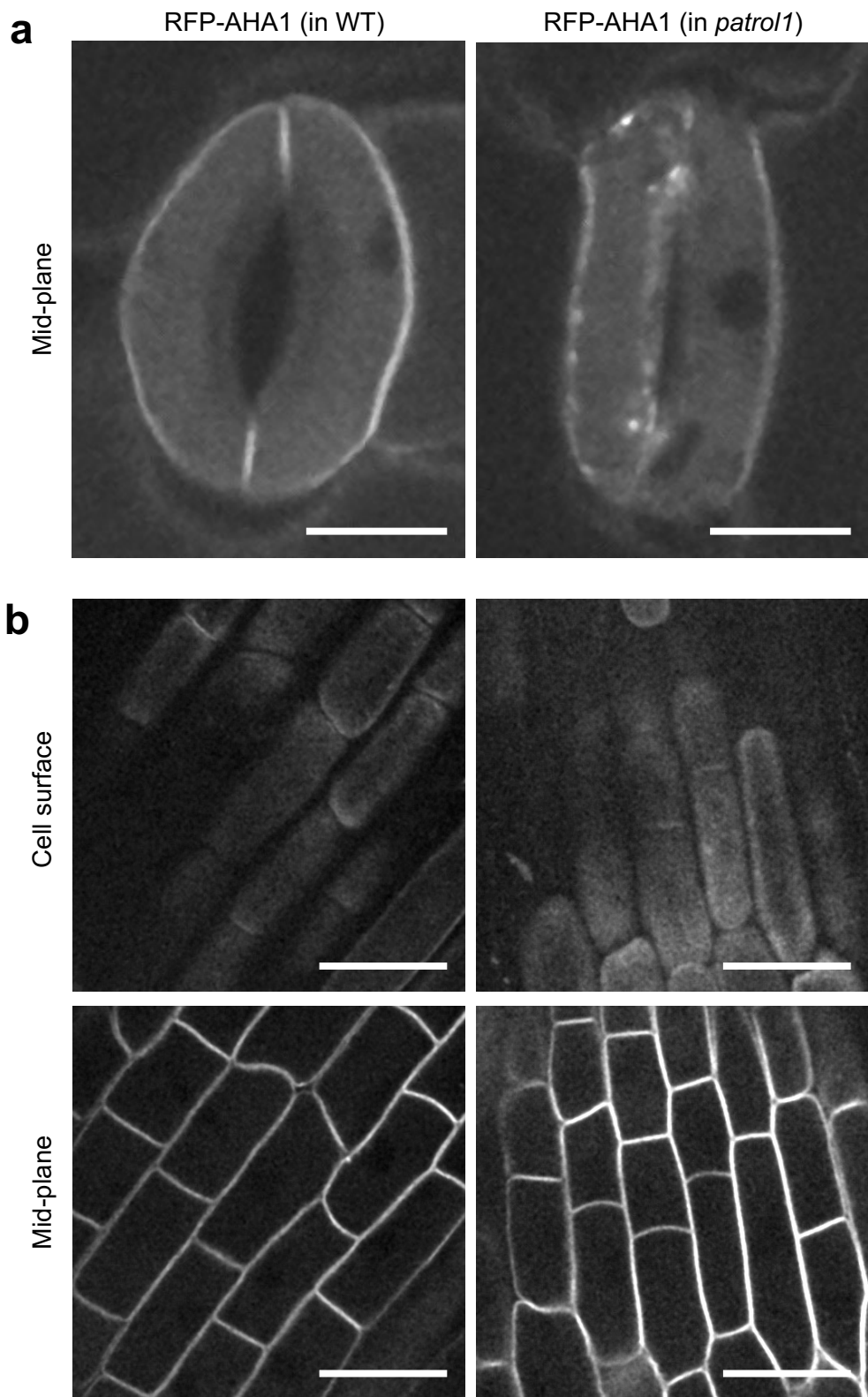
- **Supplementary Figs. S1–S3.**

- **Supplementary Movie S1.**

Dynamic motion of GFP-PATROL1 dots in the root cells. Time-sequential images were acquired at 500 ms intervals for 30 s (60 frames).



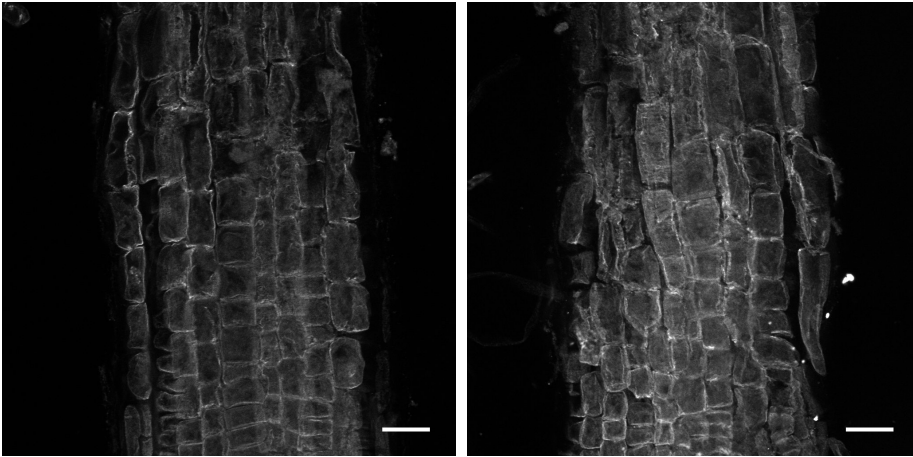
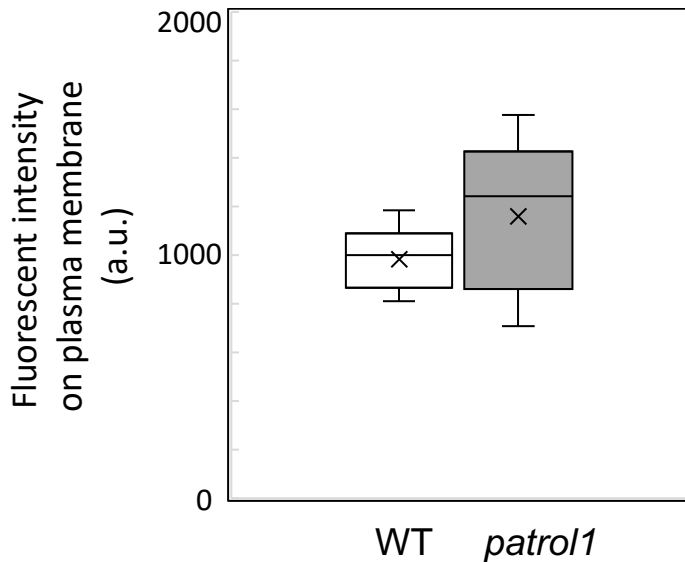
Supplementary Fig. S1. The height of the cells in meristematic zone of the root of *patrol1* mutant, wild-type (WT), and *PATROL1-OX*. The vertical length of cortical cells secondary from outer layer were measured for five independent plants.



Supplementary Fig. S2. PATROL1 is involved in AHA1 localization at the plasma membrane of stomatal guard cells but not in roots. **a** Representative confocal images of RFP-AHA1 in guard cells of cotyledons in wild-type (WT) background (left) and *patrol1* mutant (right). **b** Confocal images of the surface and mid-plane of the root cortex in wild-type (WT) background (left) and *patrol1* mutant (right). Scale bars are 5 μm in **(a)** and 10 μm in **(b)**.

a

Anti-AHA2 antibody (in WT)

Anti-AHA2 antibody (in *patrol1*)**b**

Supplementary Fig. S3. Immunohistochemical estimation of plasma membrane H⁺-ATPase in 5-day-old Arabidopsis roots. Plasma membrane H⁺-ATPase proteins were detected using unti-Thr881 of AHA2 antibody. **a** Representative confocal images of the surface of the root in wild-type (WT, left) and *patrol1* mutant (right). Scale bars are 10 μ m. **b** Box plot of the fluorescence intensity of the plasma membrane of the roots in wild-type (WT) and *patrol1* mutant. Crosses inside boxes show average values, horizontal lines within boxes mark the median value. Each boxplot consists of average values of roots from 11 plants. Each root average value derives from 8 to 27 cells. Significance between wild-type and *patrol1* mutant was calculated using two-paired Student's t-test: $P = 0.088$.