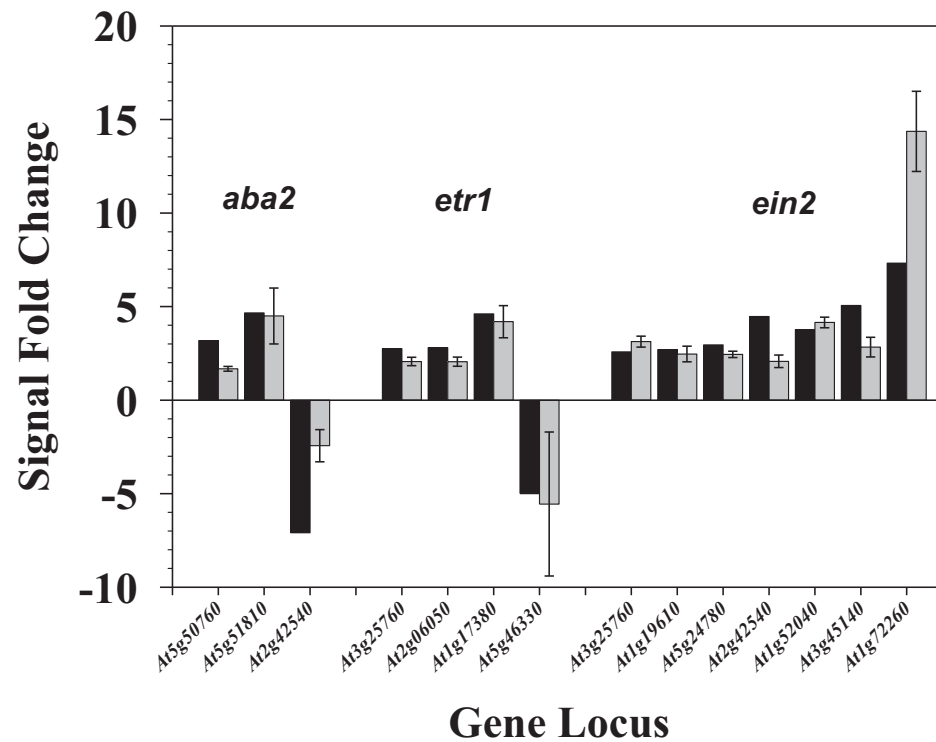


Supplemental material

Figure S1. Comparison of the ABA sensitivity levels in the *etr1-1*, *ctr1*, and *eto1-4* mutants



Supplemental material

Figure S2. Validation of GeneChip signal by semi-quantitative RT-PCR in the *aba2*, *etr1*, and *ein2* mutants

Supplemental data

Figure S1. Comparison of the ABA sensitivity levels in the *etr1-1*, *ctr1*, and *eto1-4* mutants

Seeds were grown on 1% sucrose for five days. Radicle emergence over 1 mm was regarded as germination. Germination rates were measured as germinated seeds/total germinated seeds at day 10. The results shown are the means \pm SD of three independent experiments using different seed batches (each with 60 seeds) and giving consistent results.

Figure S2. Validation of GeneChip signal by semi-quantitative RT-PCR in the *aba2*, *etr1*, and *ein2* mutants

Black bars represent the means of the normalized GeneChip signal of two independent experiments. Grey bars represent the means \pm SD of two independent experiments of semi-quantitative RT-PCR. Each experiment was duplicated. The genes tested here were chosen from Table S1.