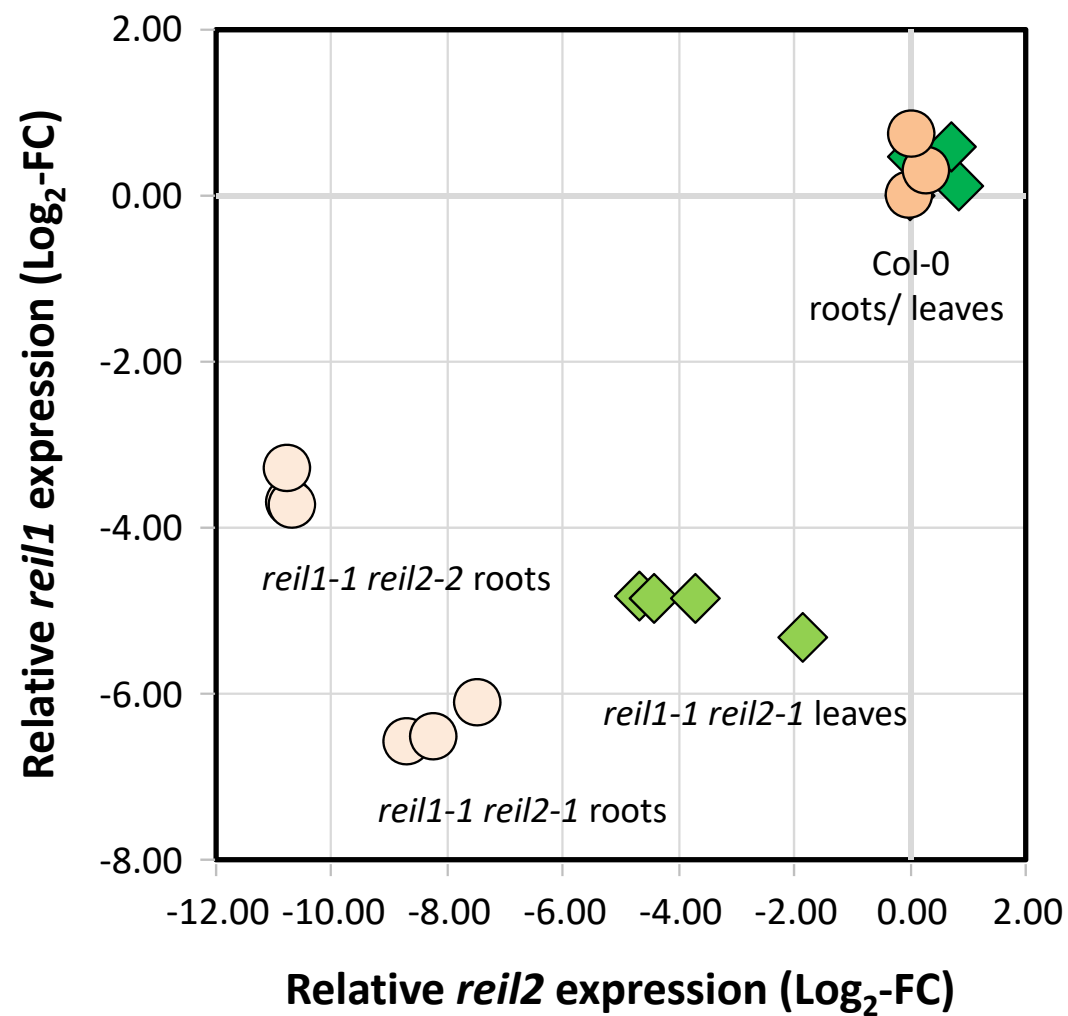




Supplemental Figure S5. Reduced transcript levels of the *reil 1* and *reil2* genes in the *reil1-1 reil2-1* and *reil1-1 reil2-2* double mutants.

Root transcript data are of this study (**Supplemental Table S2**). Leaf transcript data are from soil-grown rosette plants of *reil1-1 reil2-1* from a previous study (Supplemental Table S2 of Beine-Golovchuk et al., 2018). Log₂-transformed transcript data are normalized to the non-acclimated Col-0 wild type, i.e. log₂-fold changes (FC), of each transcriptome experiment, respectively. Roots (circles), leaves (diamonds).