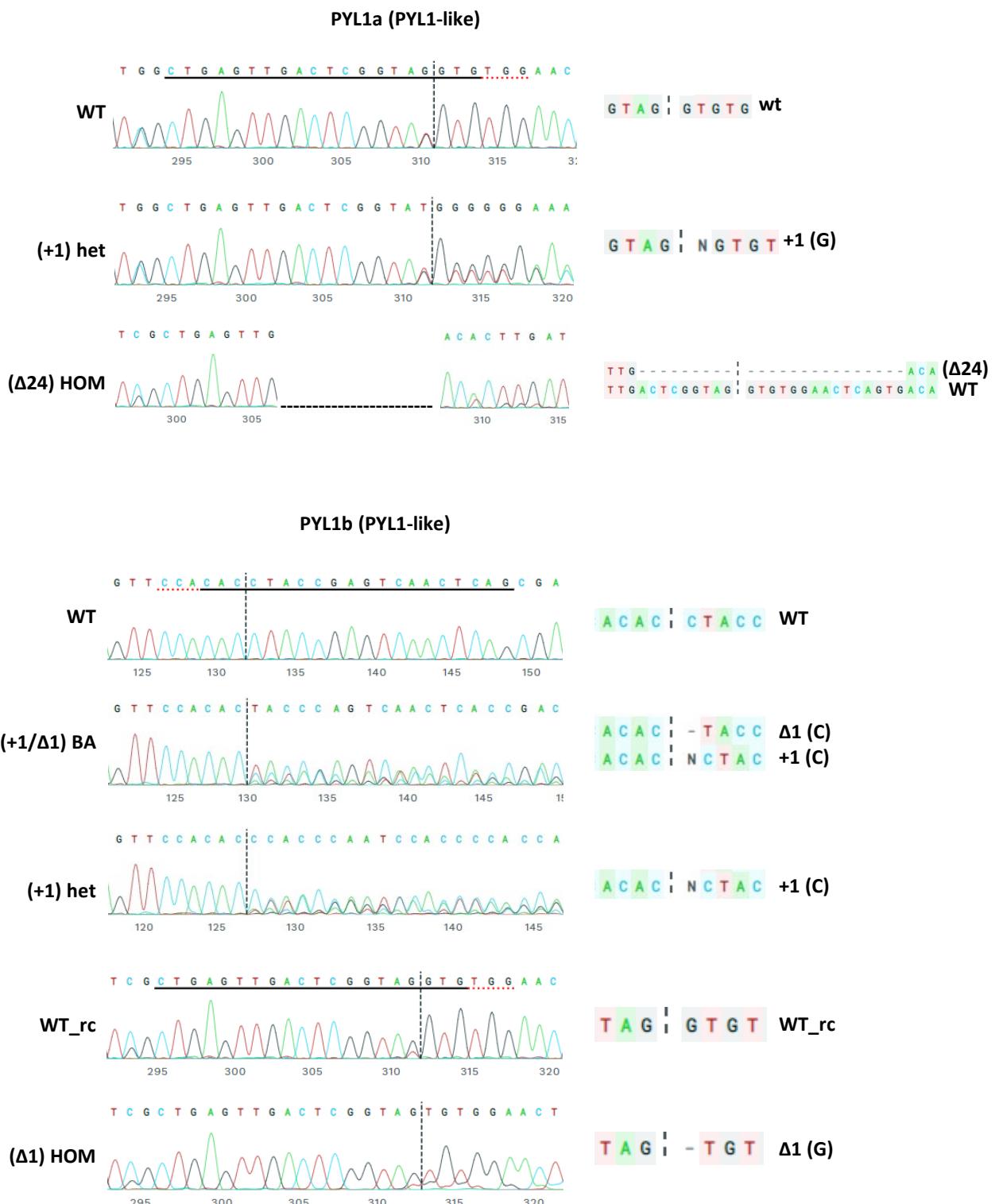
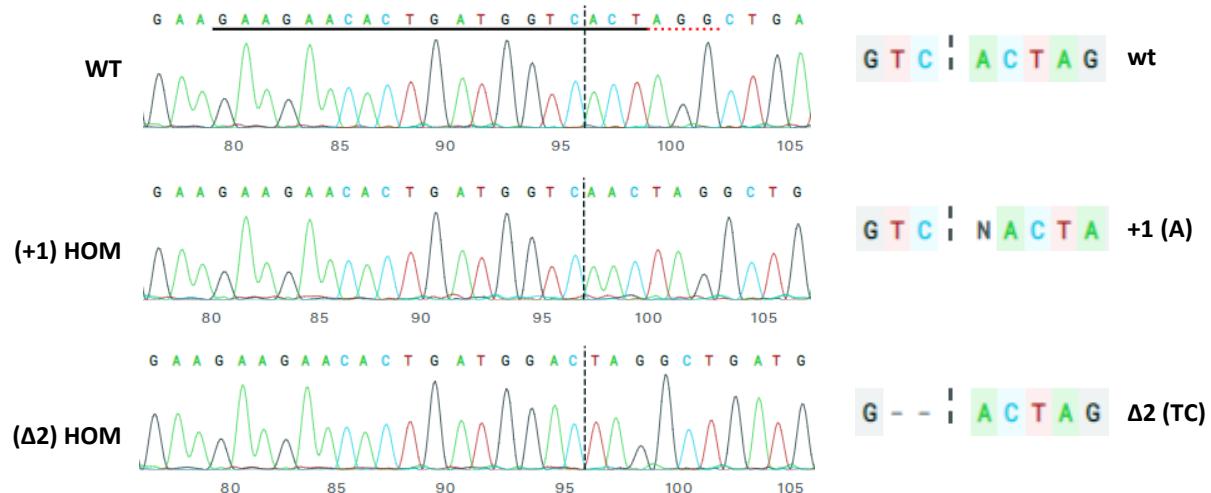


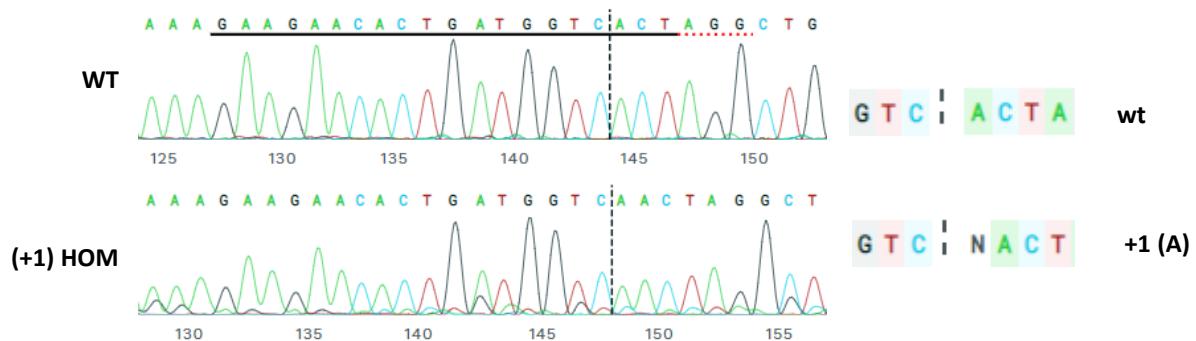
Figure S2. Chromatograms of the indicated gene sequences for wild-type and mutant plants. Sequence analysis reveals the different insertions (+1) or deletions ($\Delta 1$, $\Delta 2$, $\Delta 4$, $\Delta 24$) detected in the CRISPR/Cas9-edited alleles of each ABA receptor



PYL8a (PYL9-like)



PYL8b (PYL9-like)



PYL8c (PYL8-like)

