



Supplementary Figure S3. Confirmation of *ATAF1* expression in WT plants after priming treatment using qRT-PCR and RNA-seq.

(a) Schematic representation of the heat stress (HS) regime applied to assess *ATAF1* expression in response to priming. (b) Expression data from qRT-PCR. Expression is shown as log₂-fold change between HS and control. (c) Expression data from RNA-seq analyses.