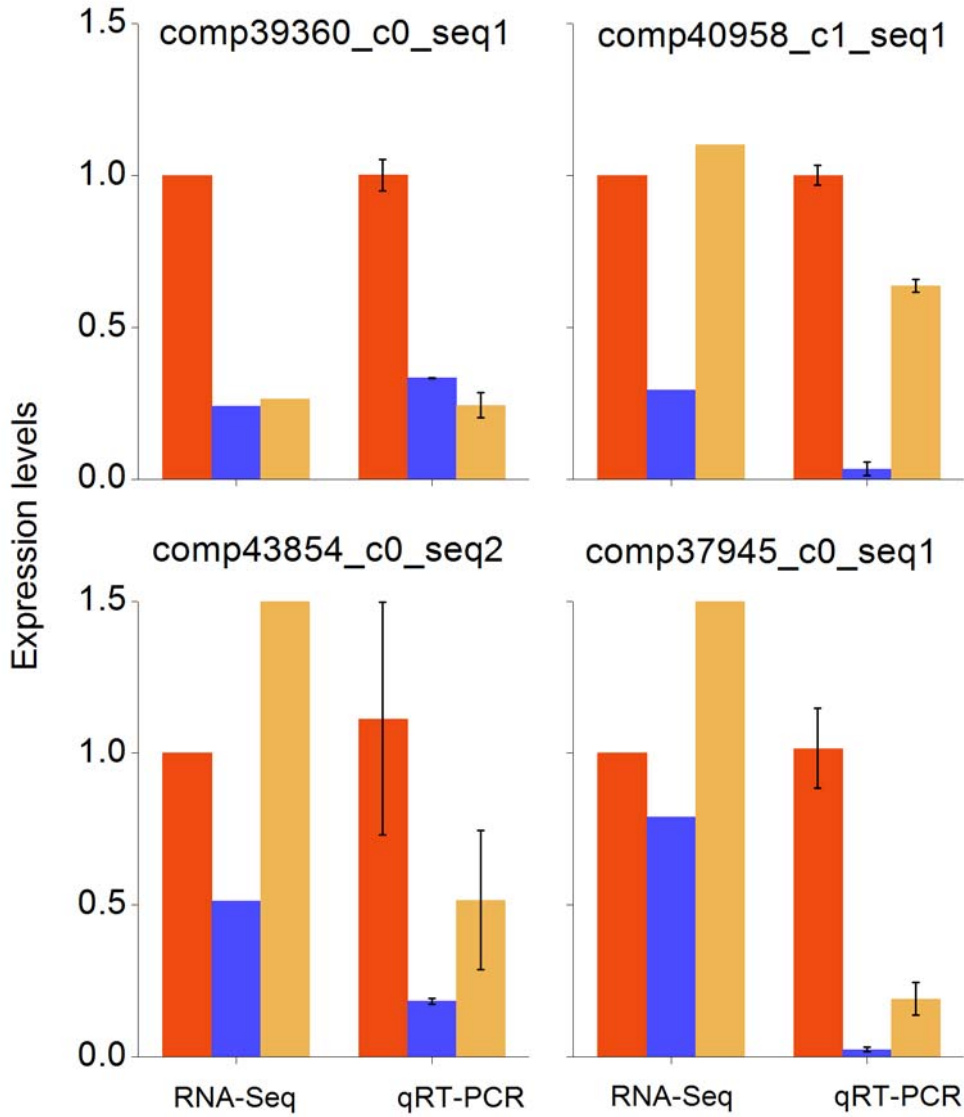


1 **Supplemental Figure S6**



2

3 **Fig. S6 qRT-PCR for GA and ABA associated genes.**

4 Values of expression levels (RPKM) by RNA-seq were standardized, with the value for primary
5 dormant seeds set to 1. Expression levels of genes detected by qRT-PCR were calculated using
6 the $2^{-\Delta\Delta C_t}$ method. PD, primary dormant seeds; ND, non-dormant seeds; SD, secondary dormant
7 seeds.

8