

Supplementary Table 1: Primers designed and used in this study

Primers for reverse transcription-quantitative (RT-qPCR)	
Forward	5'- GCAATCAACATGGCCTAGTG -3'
Reverse	5'- AAAGTCACTAGGACCCAAACA -3'
<i>Ubiquitin</i>	
Forward	5'- CACCAAGCCAAAGAAGATCA -3'
Reverse	5'- TCAGCATTAGGGCACTCCTT -3'
PCR condition	98°C 10s, 55°C 30s, 72°C 10s (40 cycles)
Infusion cloning primers for pENTR/D-TOPO vector and SIACB4	
Forward	5'- GCCGCCCCCTTCACCATGGCGGAGACTACTGAAGGG-3'
Reverse	5'- GGCGCGCCACCCCTTTATGCGGTGGTGTGTAGTTGC-3'
Primers for linearization of pENTR/D-TOPO vector	
Forward	5'-AAGGGTGGGCGCGCCGAC-3'
Reverse	5'-GGTGAAGGGGGCGGCCGC-3'

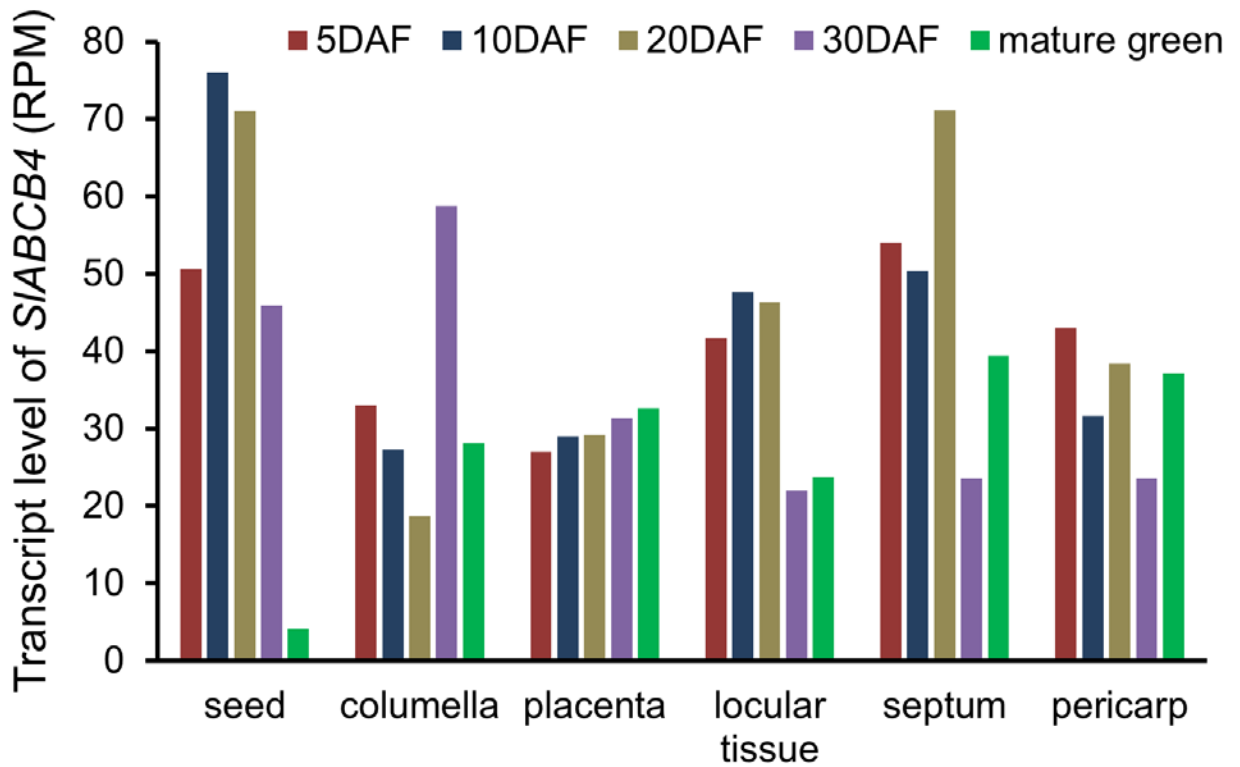


Figure S1. Tissue expression analysis of *SlABCB4* in developing fruit. High-resolution transcript data of *SlABCB4* was obtained from Tomato Expression Atlas (Tea; <http://tea.solgenomics.net/>).