

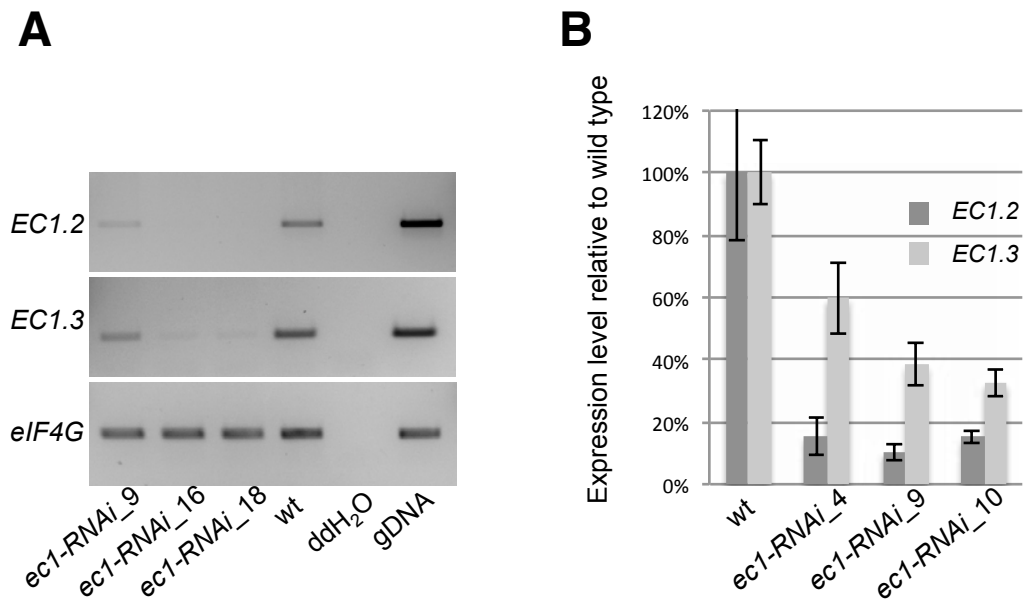


Figure S6.6 | *EC1.2* and *EC1.3* transcript levels are reduced in *ec1-RNAi* lines.

Emasculated mature pistils of wild type and of three independent heterozygous *ec1-RNAi* lines were used to analyze the relative transcript amounts of *EC1.2* and *EC1.3* by either semi-quantitative RT-PCR **(A)** or by real-time-quantitative-PCR **(B)**. Mean expression levels relative to wild type $\pm$ SD of three technical replicates are indicated. *EUKARYOTIC TRANSLATION INITIATION FACTOR 4 GAMMA* (*eIF4G*) was used as control.