



**S4 Fig. Kinetics of *YUC9* expression in *sob3-6*.**

Seedlings were grown under LD conditions for indicated time period (days). Transcript accumulation was analyzed by RT-qPCR. The *eIF4a* gene was used as an internal control. Biological triplicates were averaged and statistically analyzed by two-tailed Student's *t*-test assuming unequal variance ( $*P < 0.05$ ). Bars indicate standard error of the mean. DAG, days after germination.