



Supplemental Figure 1. The relative *GUS* transcript levels in the corresponding tissues of *SPMS* promoter-*GUS* transgenics and *ACL5* promoter-*GUS* transgenics were quantified by real-time RT-PCR and normalized to a control gene, *CBP20*. (A) The relative *GUS* transcript levels in the corresponding tissues of *SPMS* promoter-*GUS* transgenics. (B) The relative *GUS* transcript levels in the corresponding tissues of *ACL5* promoter-*GUS* transgenics. The inset has magnified scales in ordinate. The *GUS* transcript levels detected in embryos of *ACL5* promoter-*GUS* transgenics were set as 1 and the others were relatively displayed.