







Figure S1. (A) LOD scores for recorded eQTL at 30 hai (left) and 50 hai (right).
(B) Heritabilities (R2) of eQTL at 30 hai (left) and 50 hai (right). Percentages indicate the relative number of eQTL. F.i. 14% of the eQTL at 30 hai explain between 15% and 20% of the expression variation observed for the genes under their control.