



Figure S15. Whole-genome scan of 214,051 SNPs for association with (A) the relative gene expression of *RKS1-L+S* (i.e. total mRNA) or (B) the relative gene expression of *RKS1-L* ($n = 88$). The y -axis indicates the $-\log^{10} p$ -values using the Wilcoxon model or the EMMAX method. MARF = 0.05.