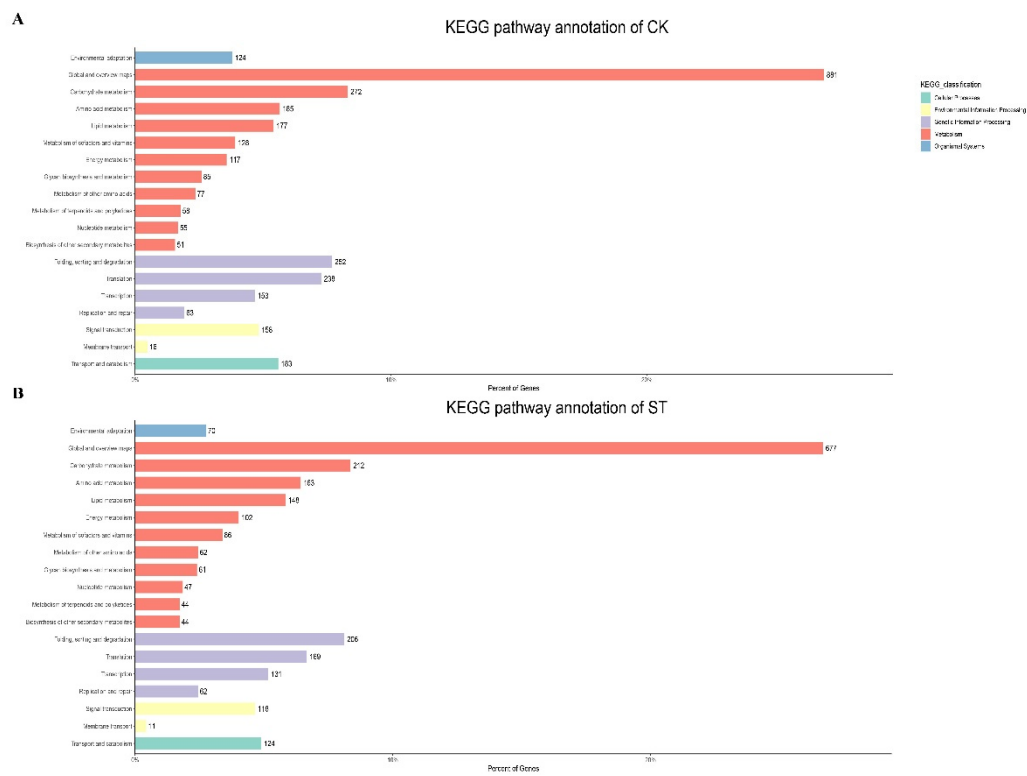
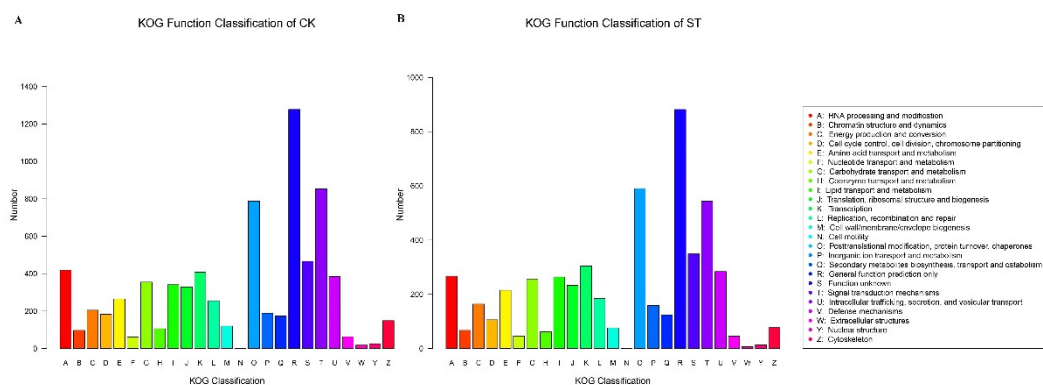


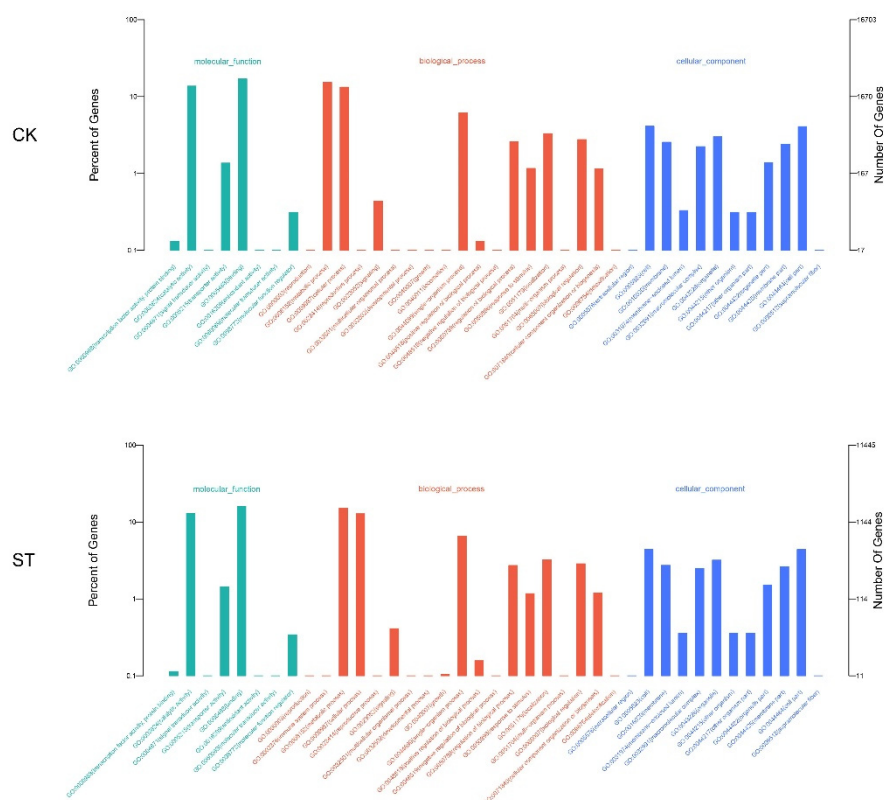
Supplementary Figure S1: Species taxonomy plot of NR annotation



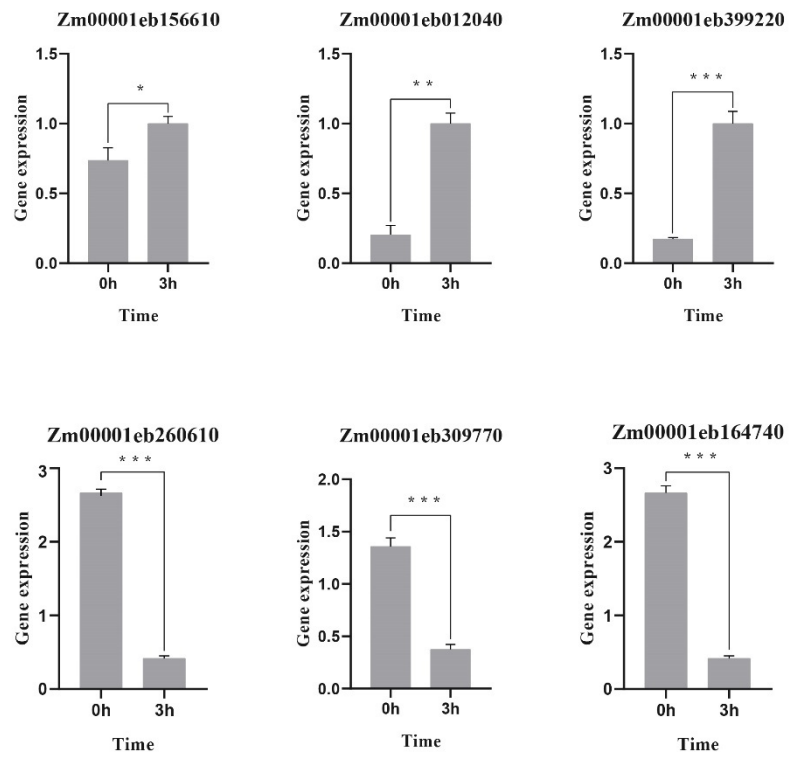
Supplementary Figure S2: KEGG annotation statistics for CK and ST transcripts



Supplementary Figure S3: KOG annotation statistics for CK and ST transcripts



Supplementary Figure S4: GO annotation statistics for CK and ST transcripts



Supplementary Figure S5: The qRT-PCR results figure. “ * ” indicates $P < 0.05$, “ ** ” indicates $P < 0.01$, and “ *** ” indicates $P < 0.001$.