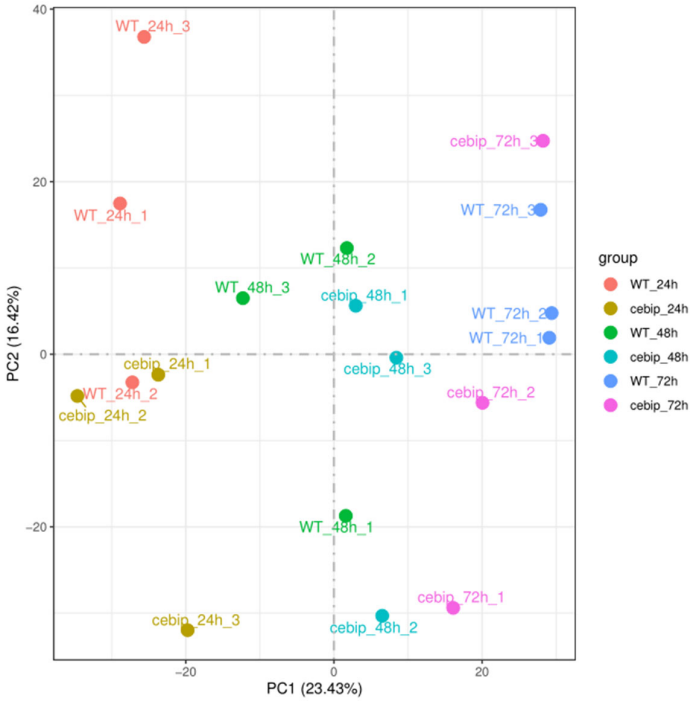
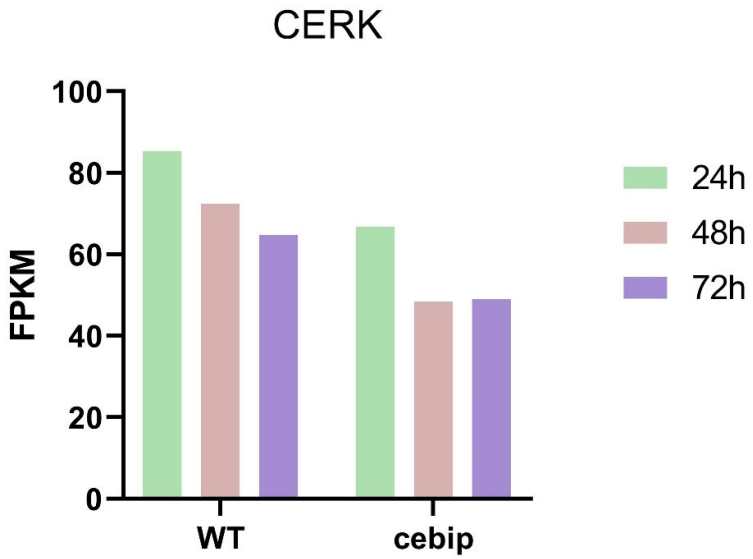
Supplementary Materials



Supplementary Figure S1. The principal component analysis (PCA) of gene expression in rice leaves. The major PC1 was 23.43% and was linked to time points, and the second PC2 was 16.42% and was associated with plant genotype. Each treatment contains three biological replicates.



Supplementary Figure S2. The transcript level of the CERK gene in the WT and cebip lines at 24, 48, and 72 hpi. The bar represents the average level of three biological replicates.