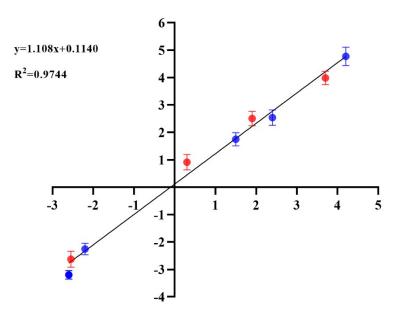$y=1.108x+0.1140$

$R^2=0.9744$

Fig. S1 Correlation analysis of the gene expression levels between qPCR and transcriptome data for nine

selected genes. The nine analyzed genes and primers are listed in Table S1. Each point represents a value

of fold-change of the expression level of acid–stress treatment compared with the control group.