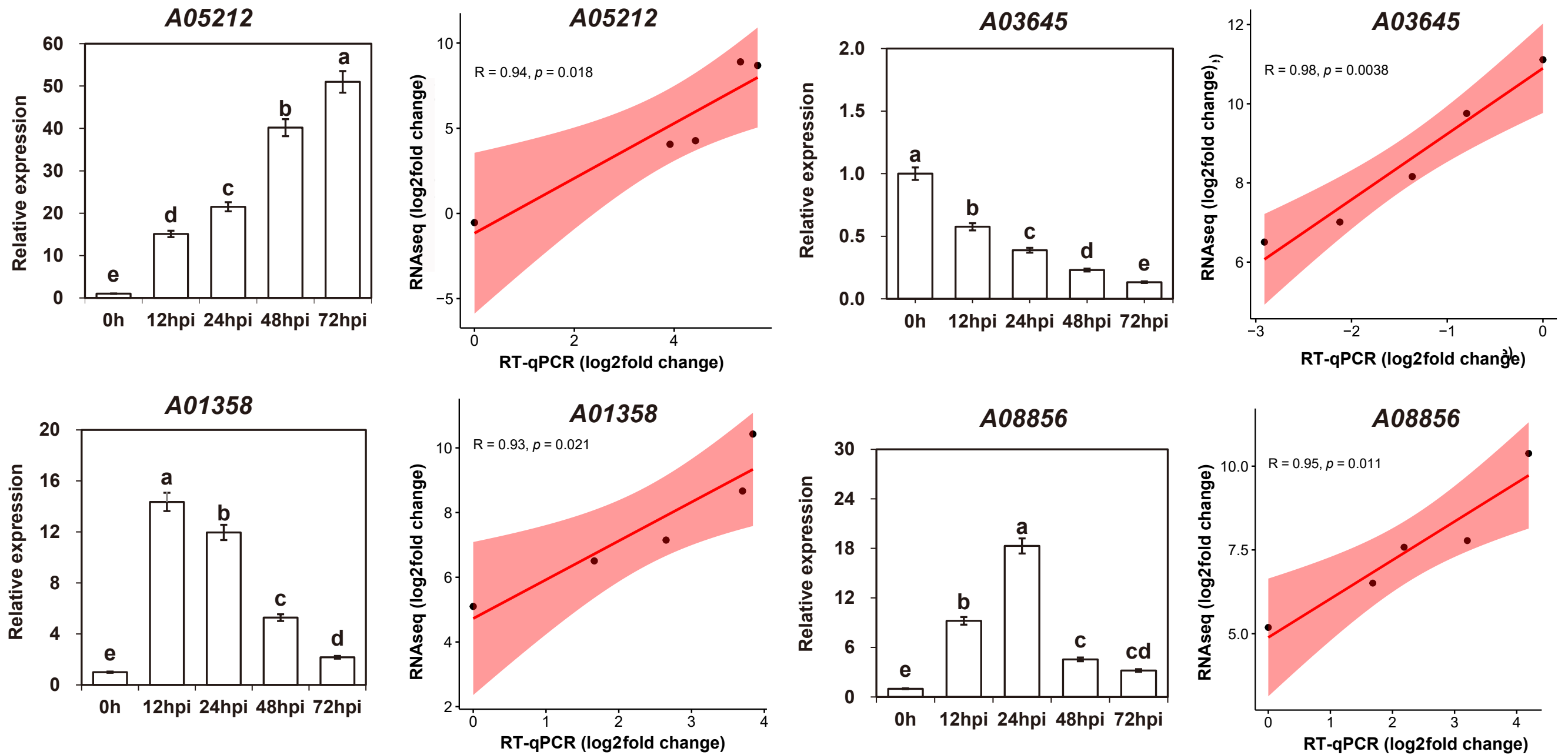




**Fig. S5.** Expression analysis of four randomly selected genes at five time points (0 h, 12, 24, 48, and 72 hpi) using RT-qPCR. Bar plots show relative expression levels, different letters above bars indicate significant differences ( $p < 0.05$ ). Line plots show Pearson correlation between RNA-seq and RT-qPCR log2 fold change values for each gene.