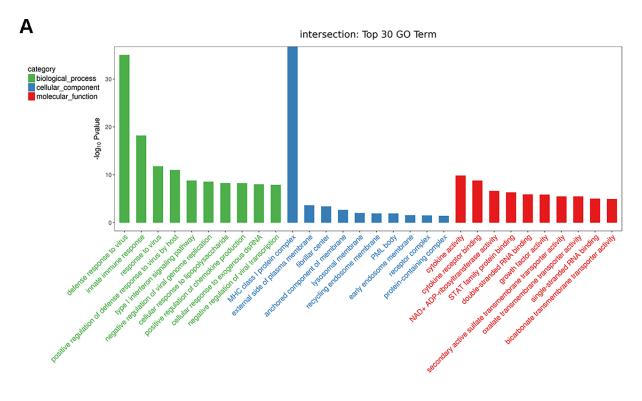
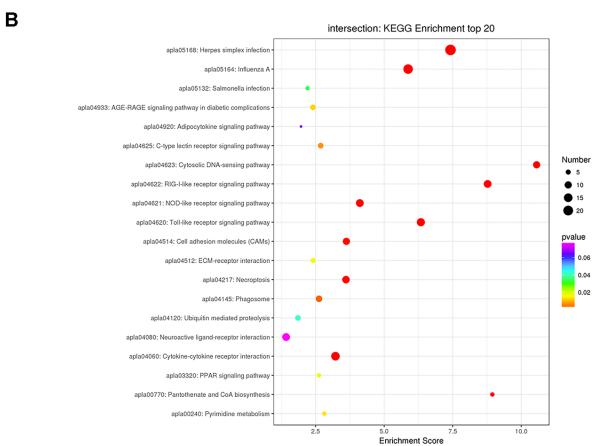
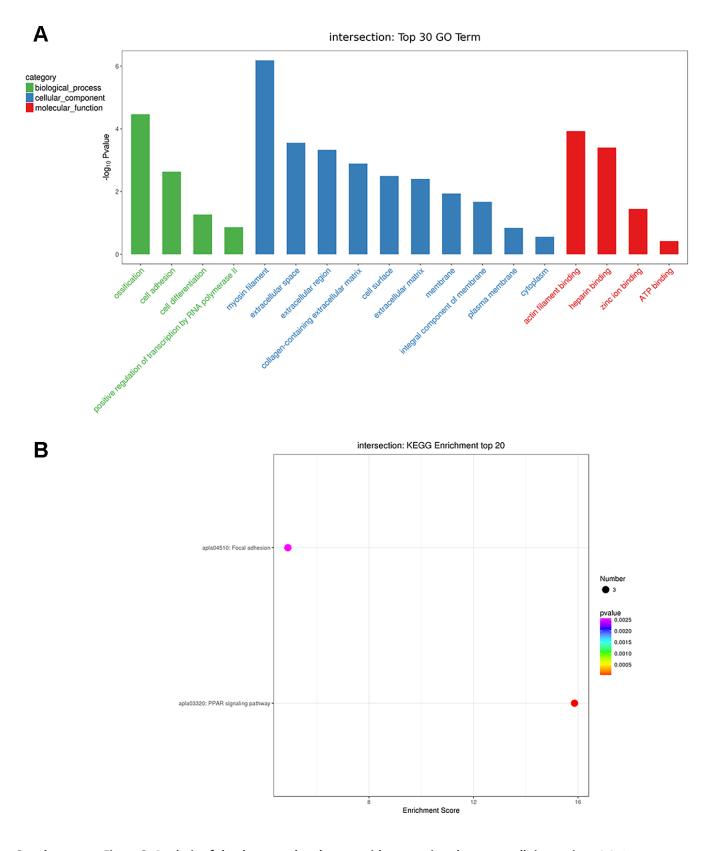


Supplementary Figure 1. Principal component of samples at 12, 24, 36, 48 and 60 hpi. Note: PC1 shows the differences among duck Tembusu virus (DTMUV)-infected samples; PC2 indicates differences between mock- and DTMUV-infected samples. 12h\_Con, mock-infected DEFs at 12 hpi; 24h\_Con, mock-infected DEFs at 24 hpi; 36h\_Con, mock-infected DEFs at 36 hpi; 48h\_Con, mock-infected DEFs at 48 hpi; 60h\_Con, mock-infected DEFs at 36 hpi; 12h\_DTMUV, DTMUV-infected DEFs at 12 hpi, 24h\_DTMUV, DTMUV-infected DEFs at 24 hpi, 36h\_DTMUV, DTMUV-infected DEFs at 36 hpi, 48h\_DTMUV, DTMUV-infected DEFs at 48 hpi, 60h\_DTMUV, DTMUV-infected DEFs at 60 hpi.





Supplementary Figure 2. Analysis of the upregulated genes with expression changes at all time points. (A) The top 30 Gene Ontology (GO) enrichment of differentially expressed genes. (B) The top 20 Kyoto Encyclopedia of Genes and Genomes (KEGG) enrichment analysis of differentially expressed genes.



**Supplementary Figure 3.** Analysis of the downregulated genes with expression changes at all time points. (A) The top 30 Gene Ontology (GO) enrichment of differentially expressed genes. (B) The top 20 Kyoto Encyclopedia of Genes and Genomes (KEGG) enrichment analysis of differentially expressed genes.