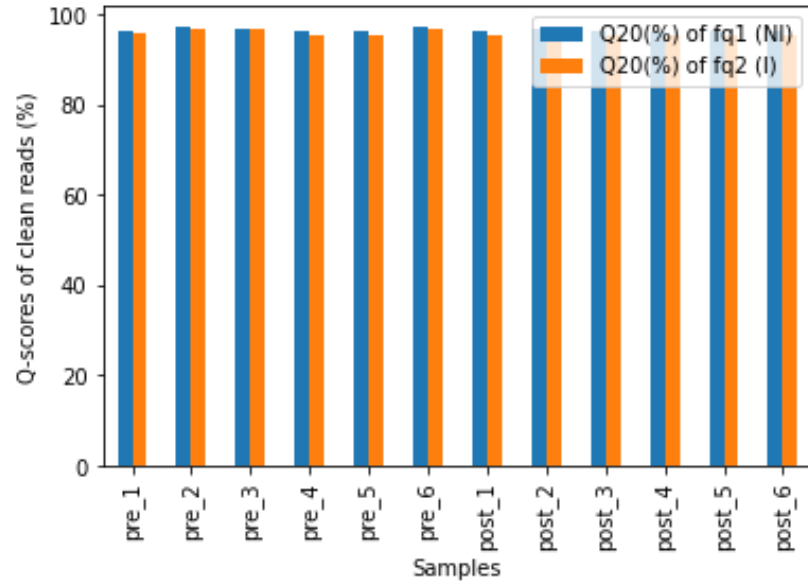
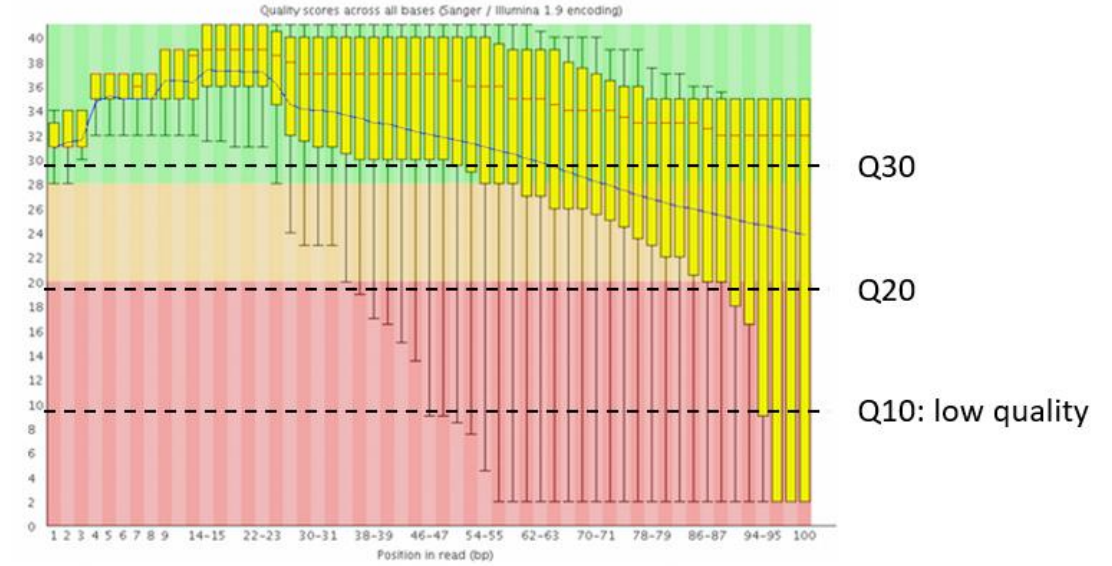


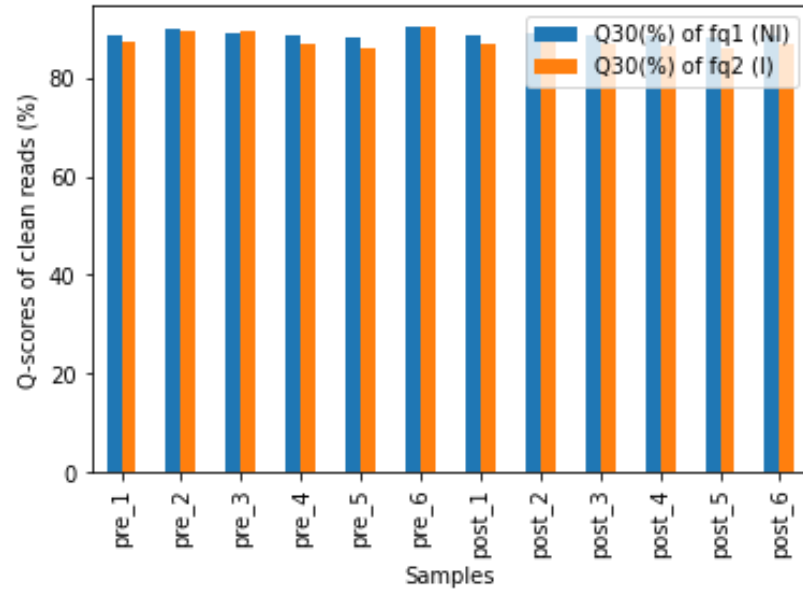
Pre and post FMDV infection Q-scores across all bases for non-infected and infected porcine



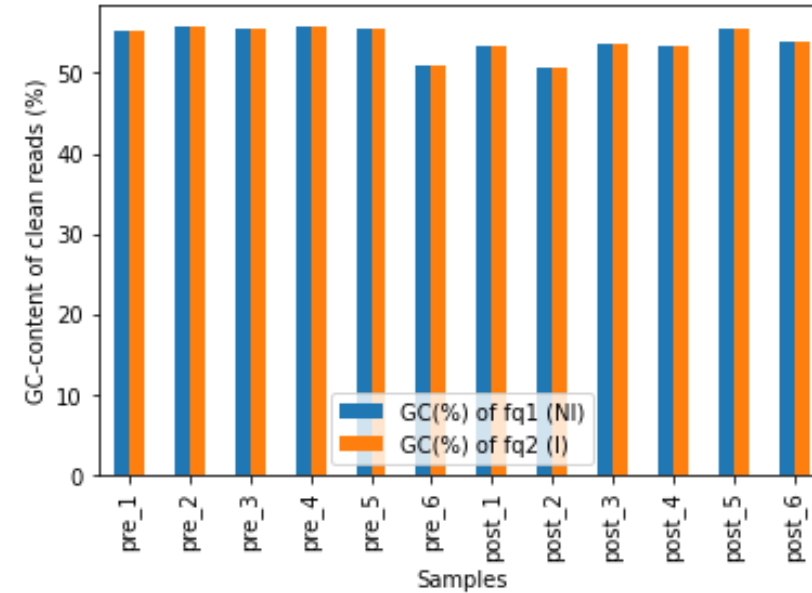
Per base sequence quality trimming



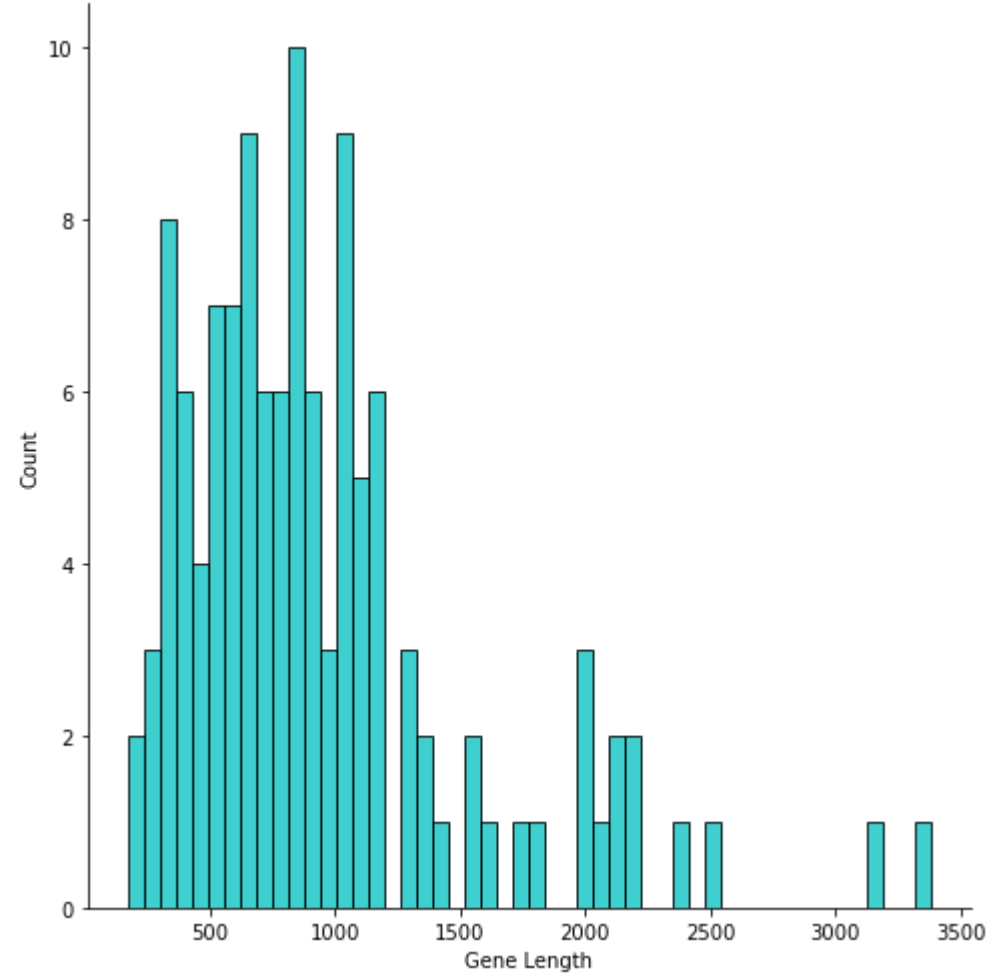
Pre and post FMDV infection Q-scores across all bases for non-infected and infected porcine



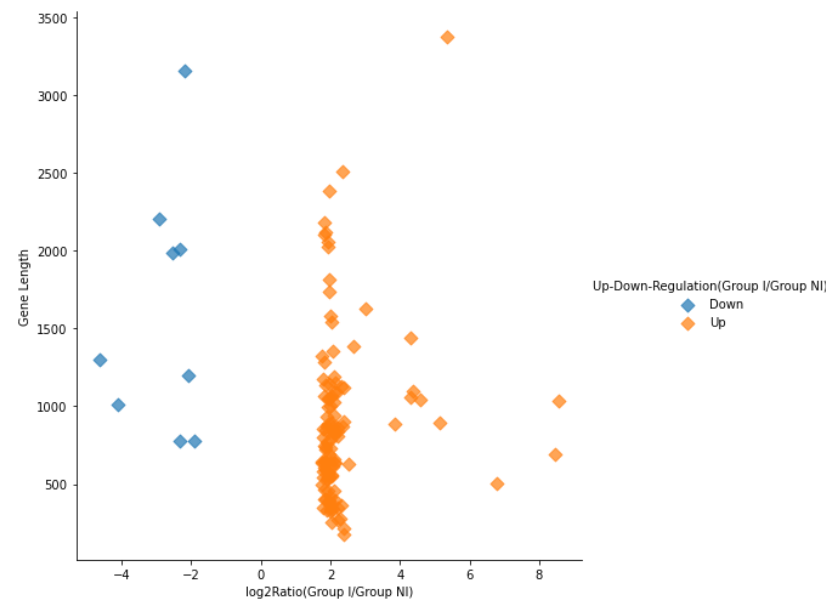
GC-content of samples pre and post FMDV infection



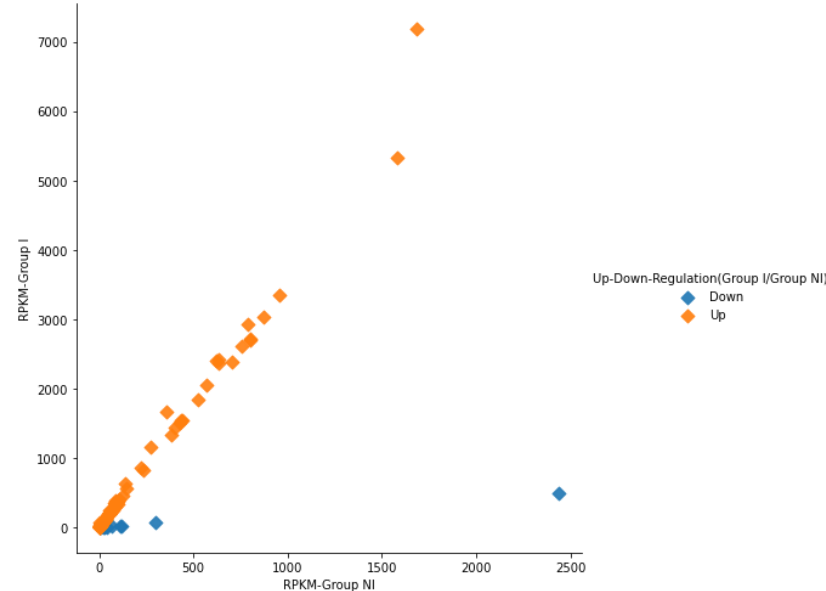
The distribution of gene length



Log2Ration vs Gene Length

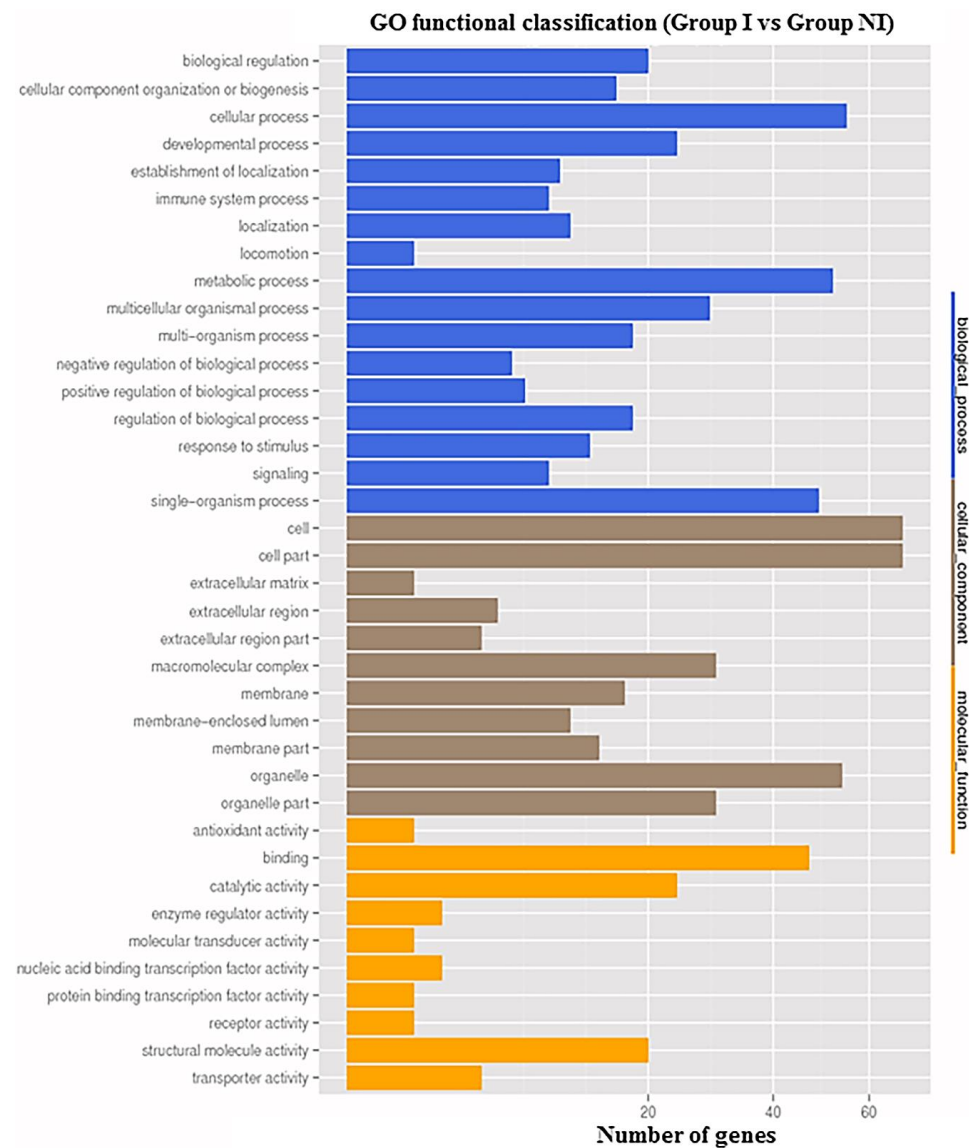


RPKM(infected) vs RPKM(non-infected)



Gene Ontology functional classification

Original Fig.2

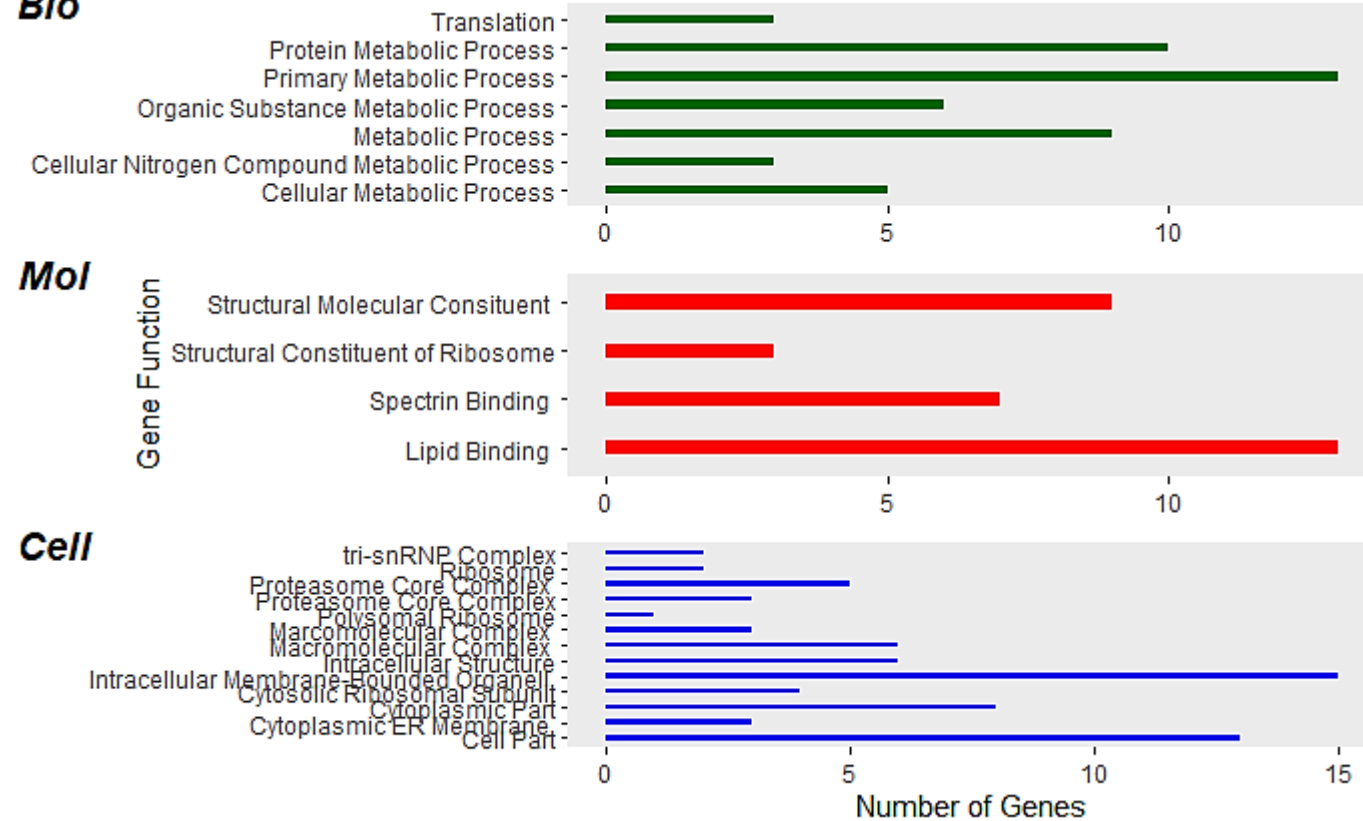


Reproduced Fig.2

Bio

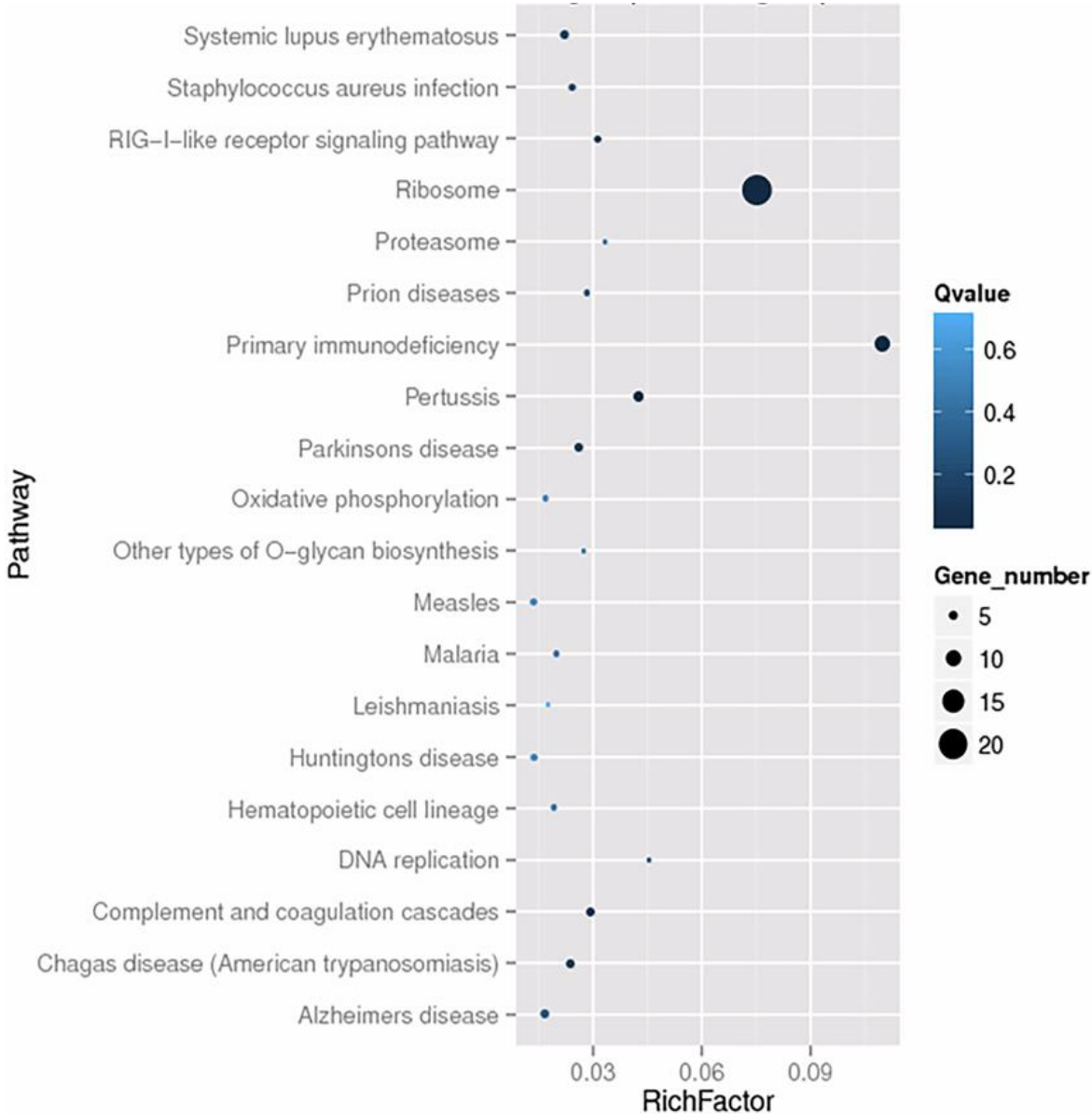
Mol

Cell

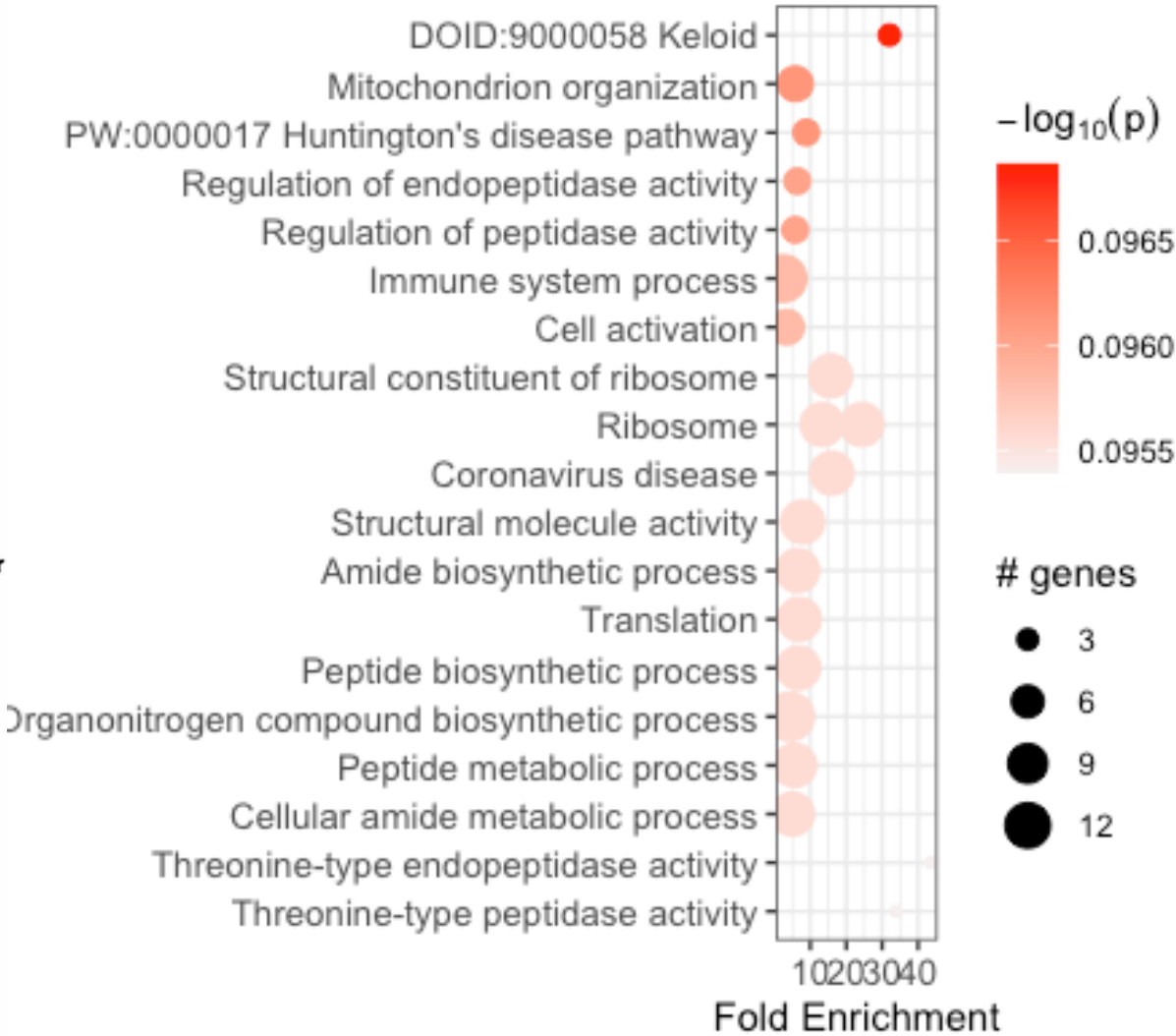


KEGG Enrichment Analysis

Original Fig.3

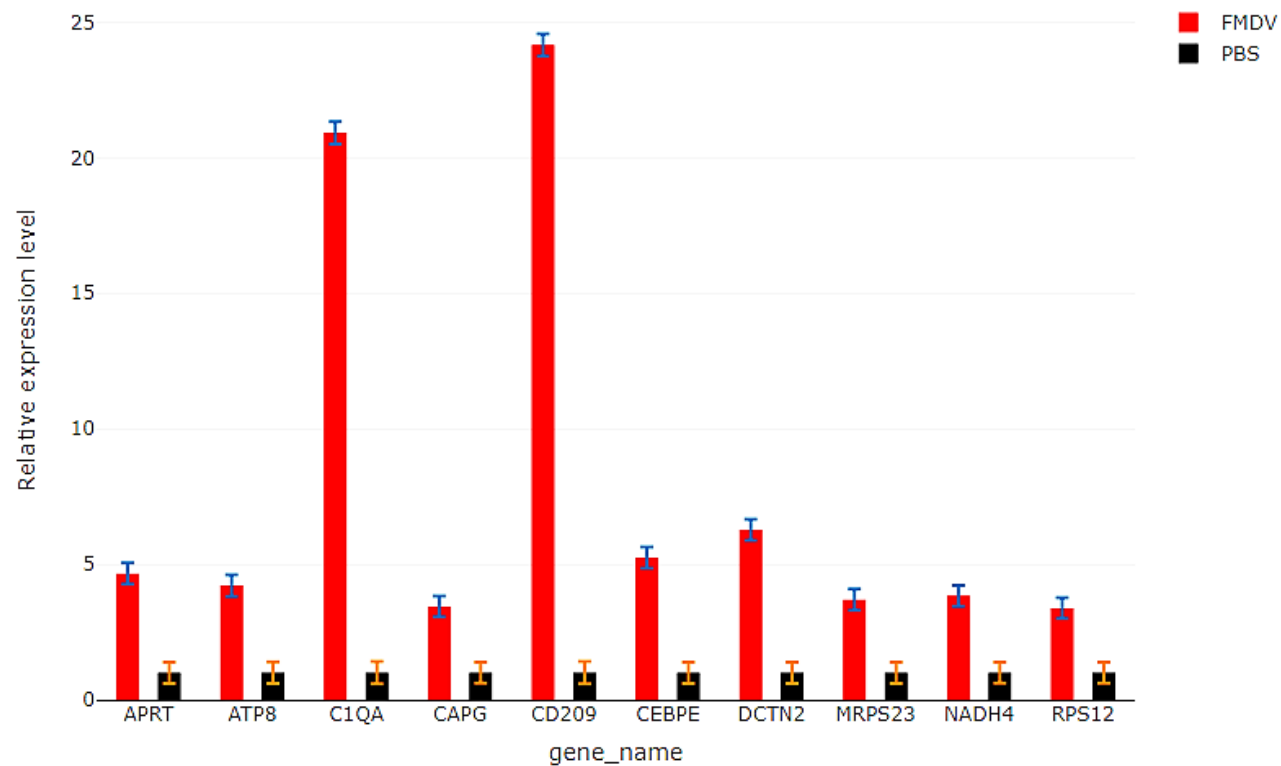


Reproduced Fig.3



Validation of the of DEGs by qRT-PCR

Reproduced Fig.4



Original Fig.4

