

Supplementary Material

Supplementary Figures

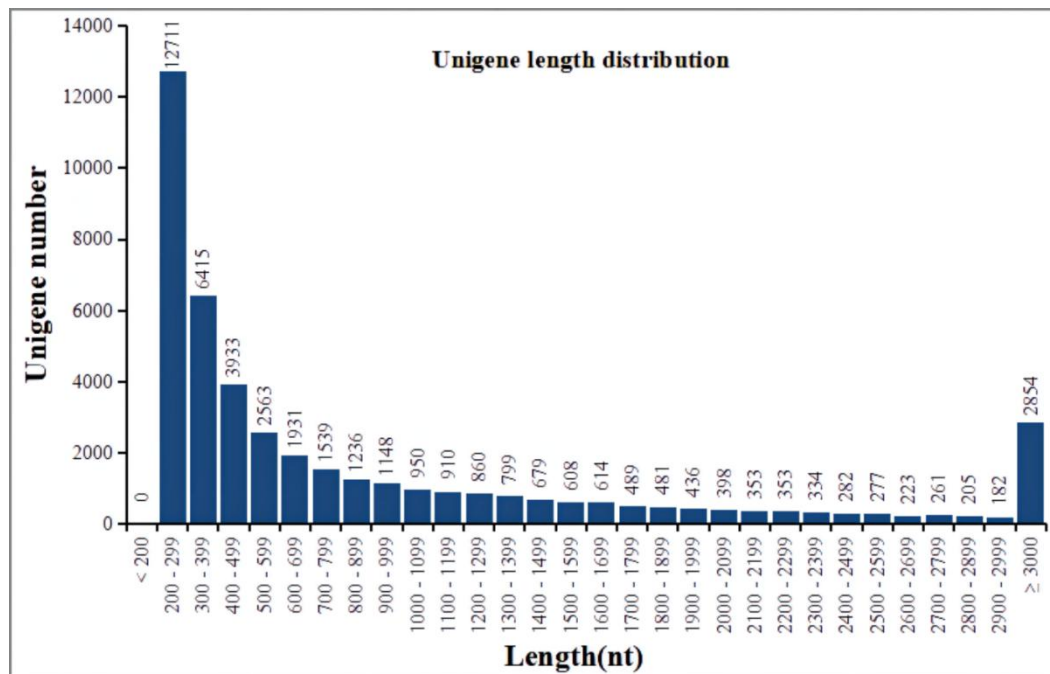


Figure S1. Length distribution of unigenes in *C. dieckmanni*. The histogram bars represent the numbers of unigenes in each length category.

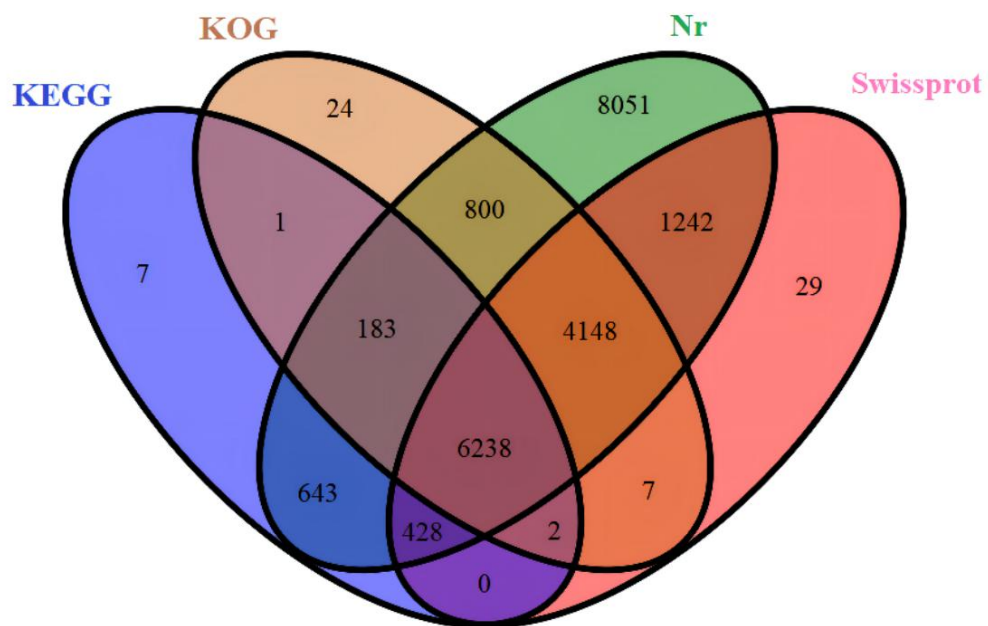


Figure S2. Venn diagrams of genes that were annotated in NR, Swiss-Port, KEGG and COG/KOG databases.

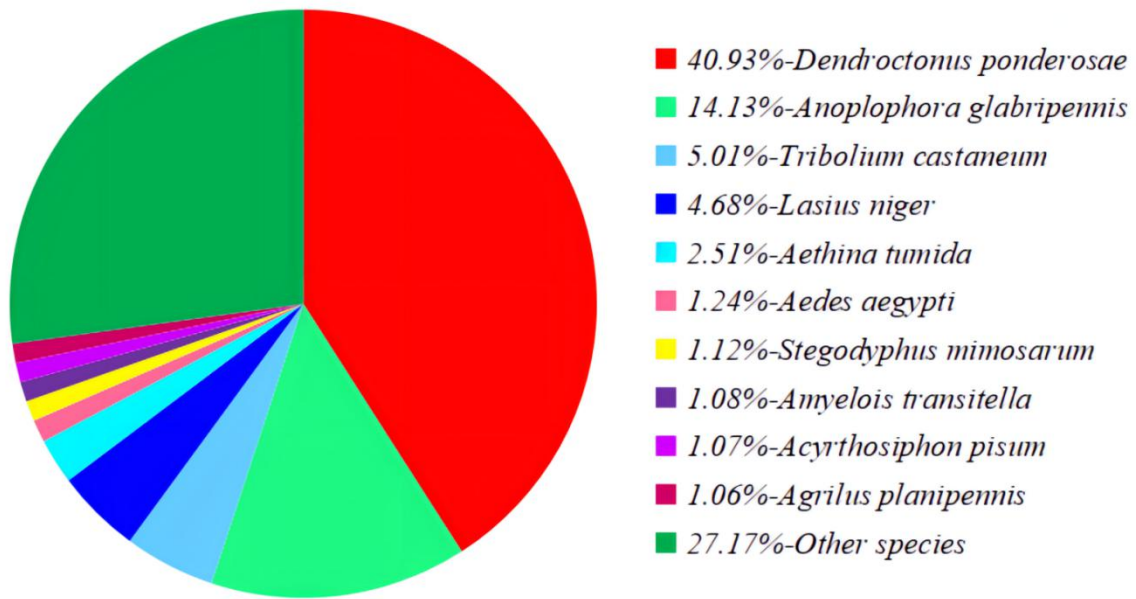


Figure S3. Species distribution of annotated genes in the Nr database.

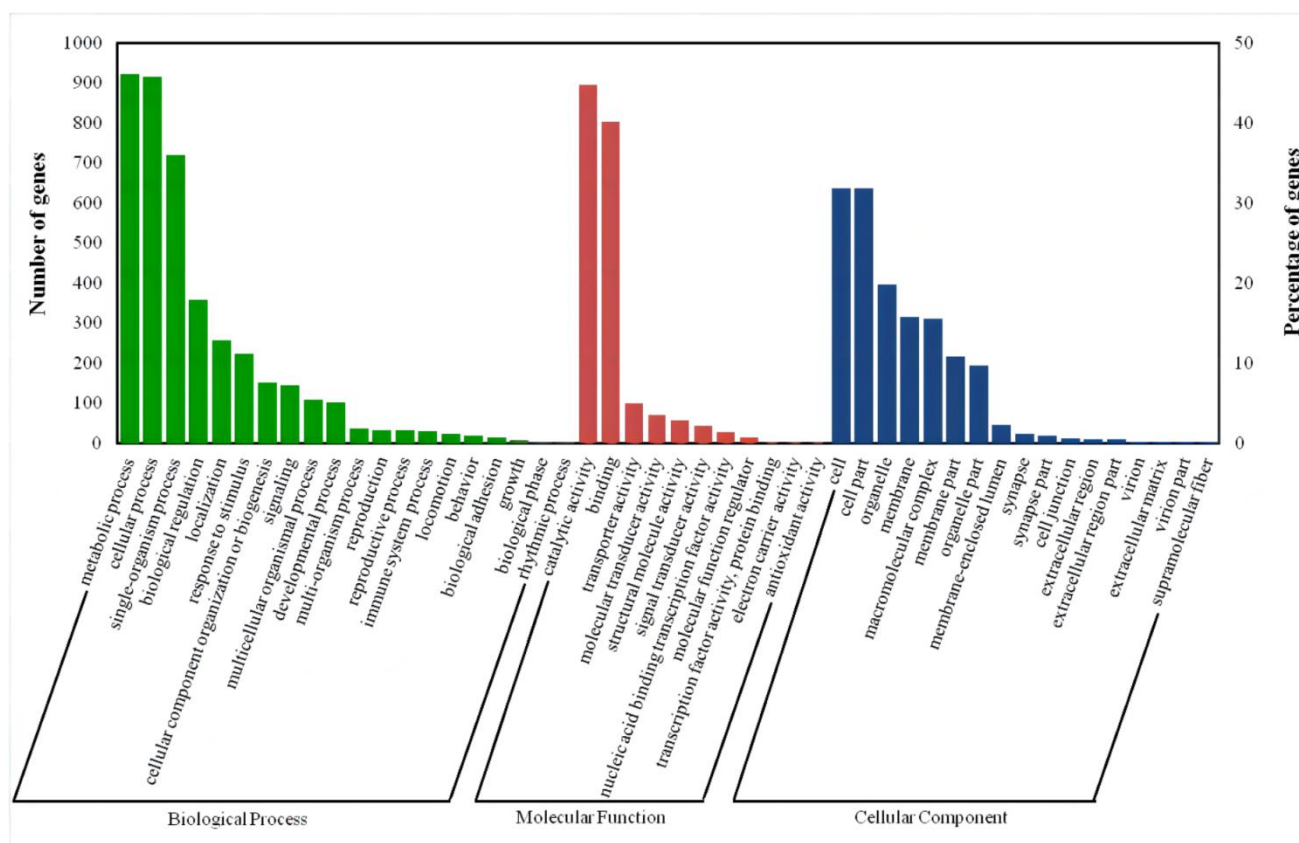


Figure S4. Gene Ontology (GO) classifications of *C. dieckmanni* unigenes.

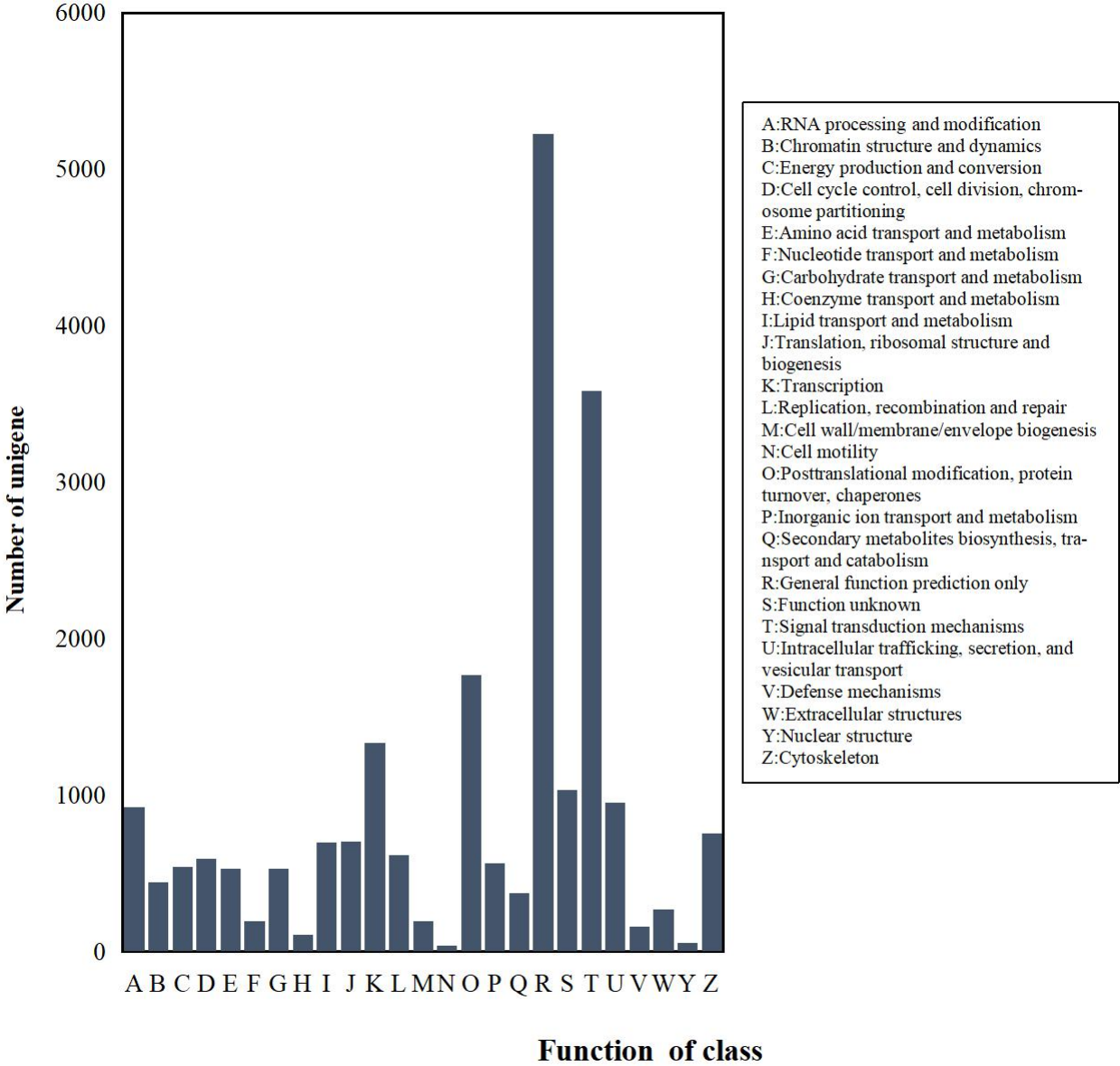


Figure S5. Karyotic Orthologous Groups classification of *C. dieckmanni* unigenes.