

Supplementary Material to “Transcriptome analyses of *Ditylenchus destructor* in responses to cold and desiccation stress”

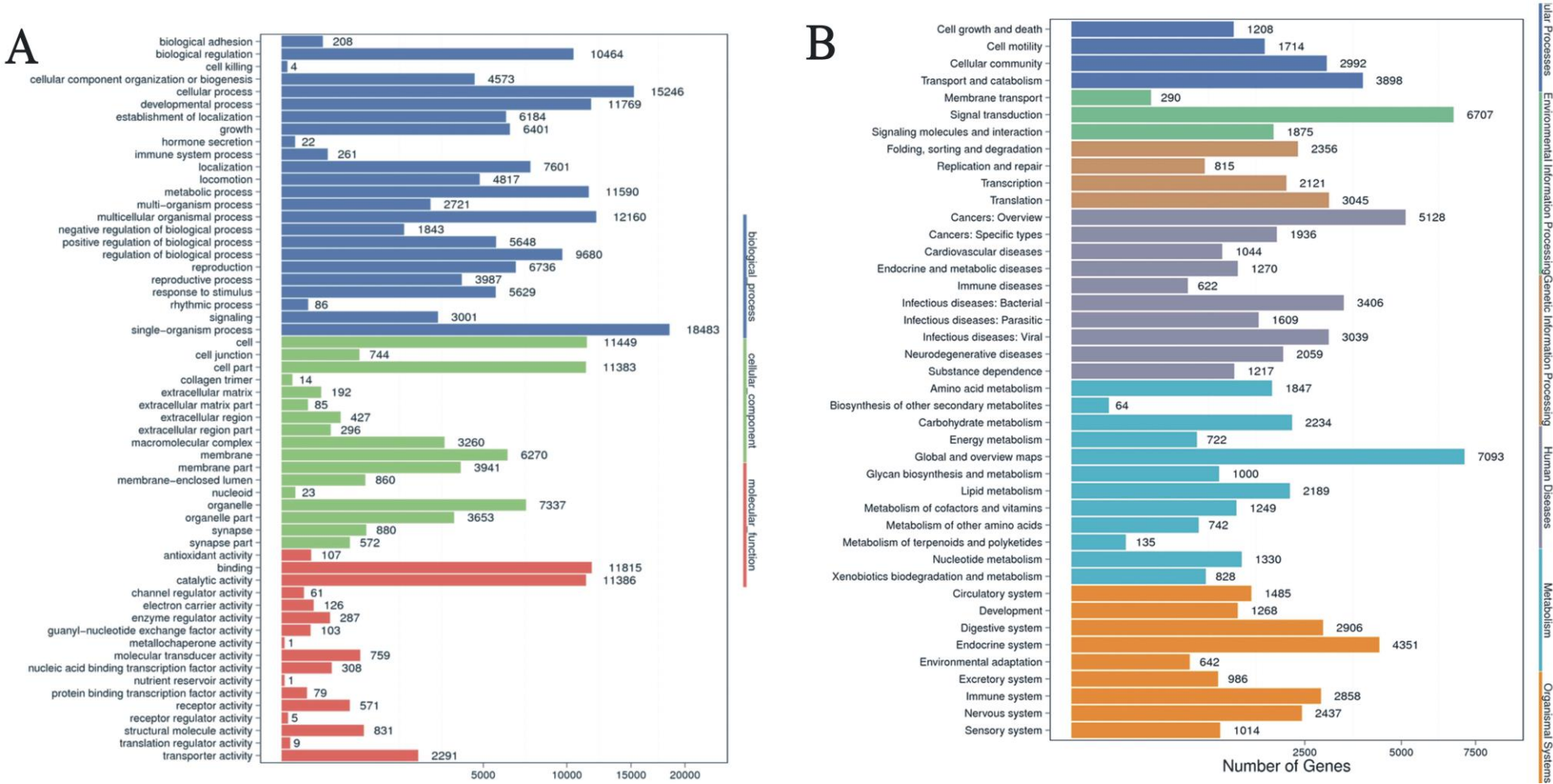


Figure S1 - Functional distribution of GO annotation (A) and KEGG annotation (B). X axis represents the number of Unigenes. Y axis represents the GO functional category/KEGG functional category.