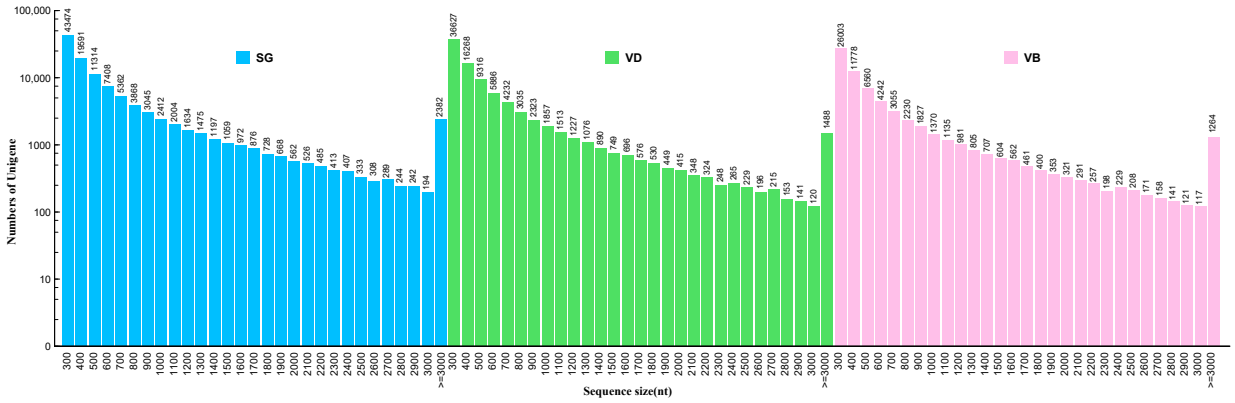
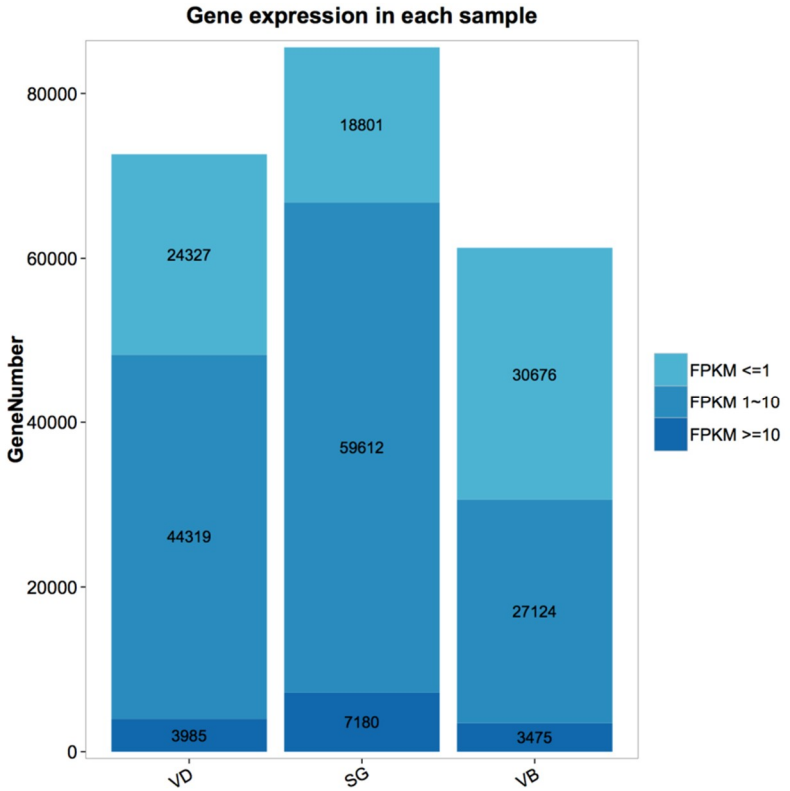
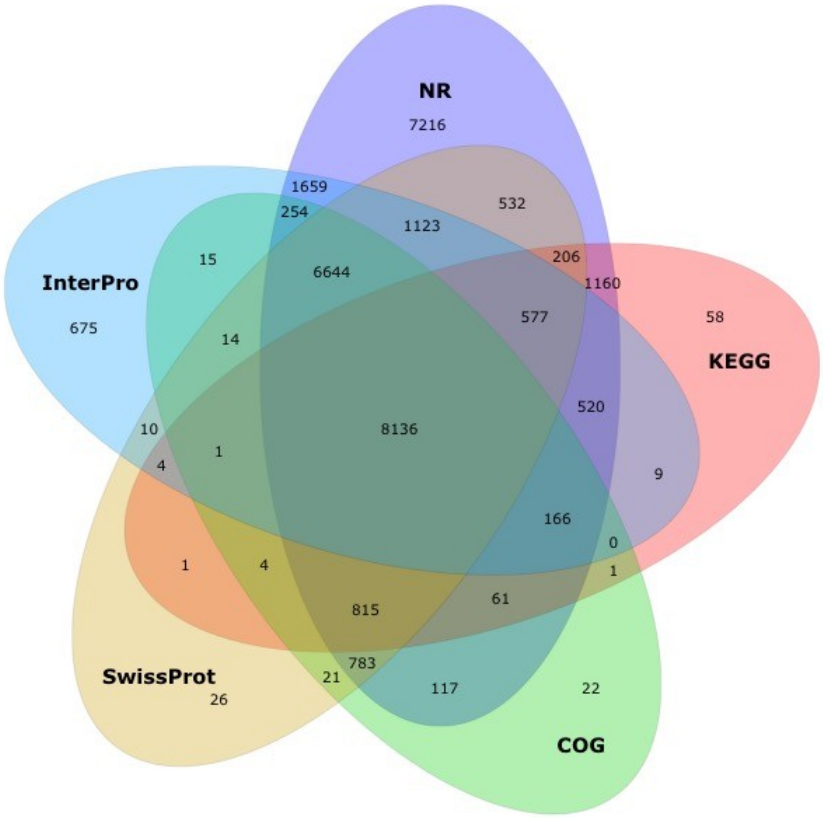


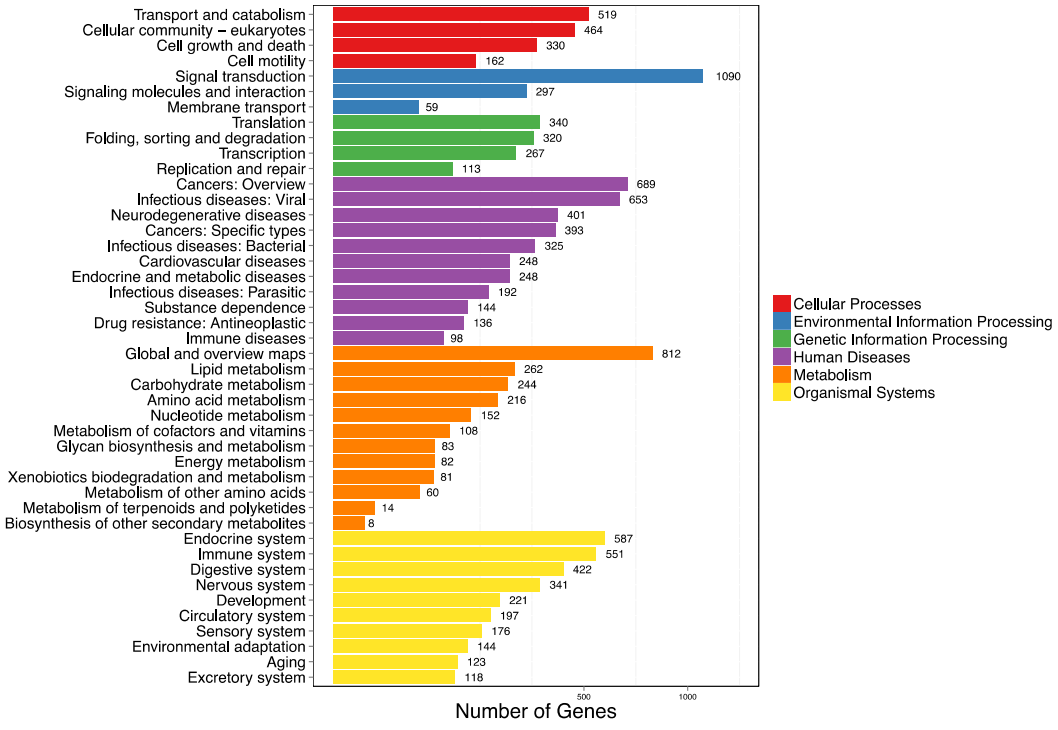
Supplementary Figure S1. Length distribution of unigenes for venom duct (VD), salivary gland (SG) and venom bulb (VB).



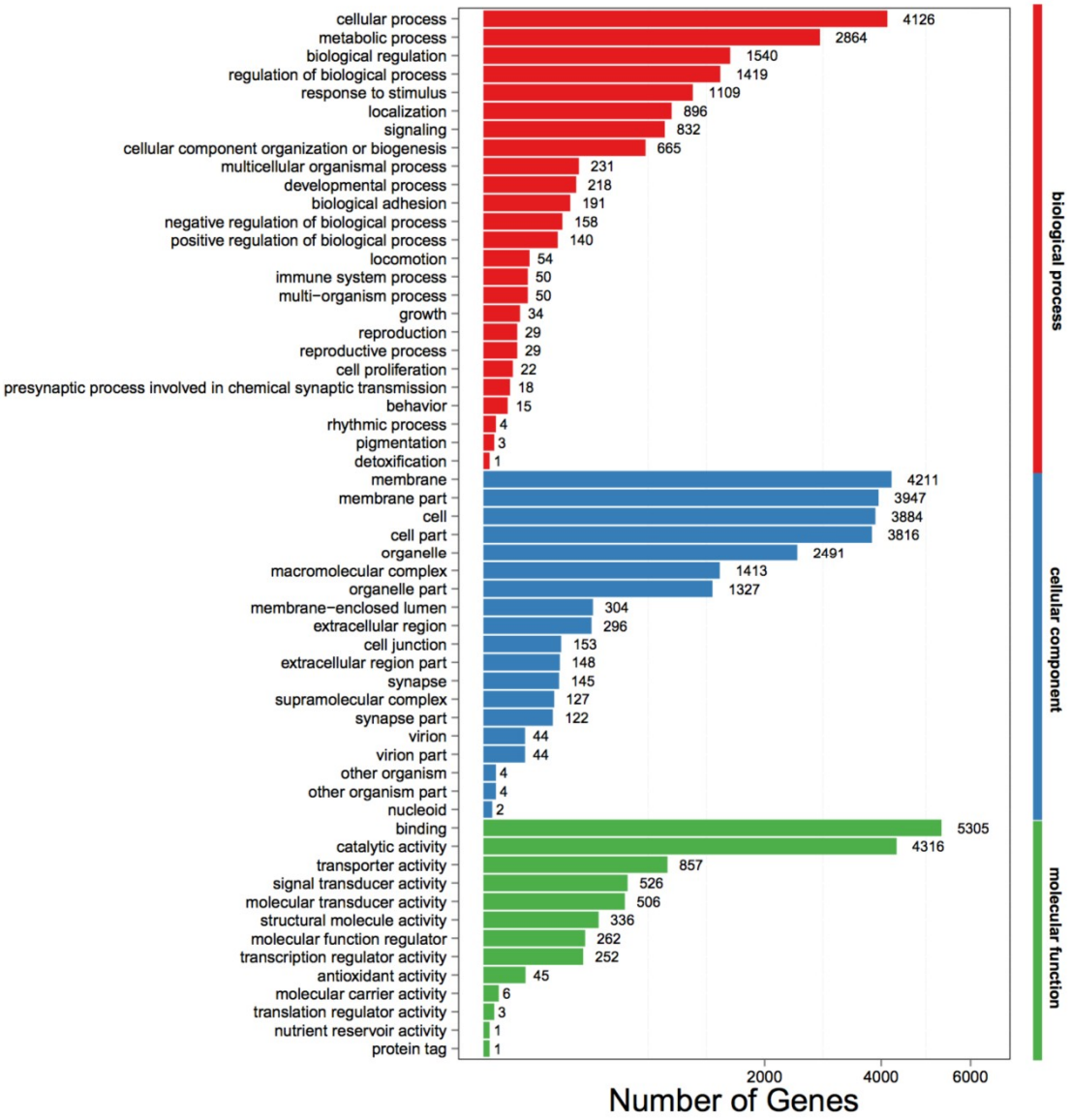
Supplementary Figure S2. Distribution of gene transcription levels in venom duct (VD), salivary gland (SG) and venom bulb (VB).



Supplementary Figure S3. Functional annotation of all assembled genes in venom duct (VD), salivary gland (SG) and venom bulb (VB) based on multiple public databases.



Supplementary Figure S4. Enriched KEGG pathways and number distribution of annotated genes in each pathway.



Supplementary Figure S5. Enriched GO terms and number distribution of annotated genes in each term.