

Supplementary Figures

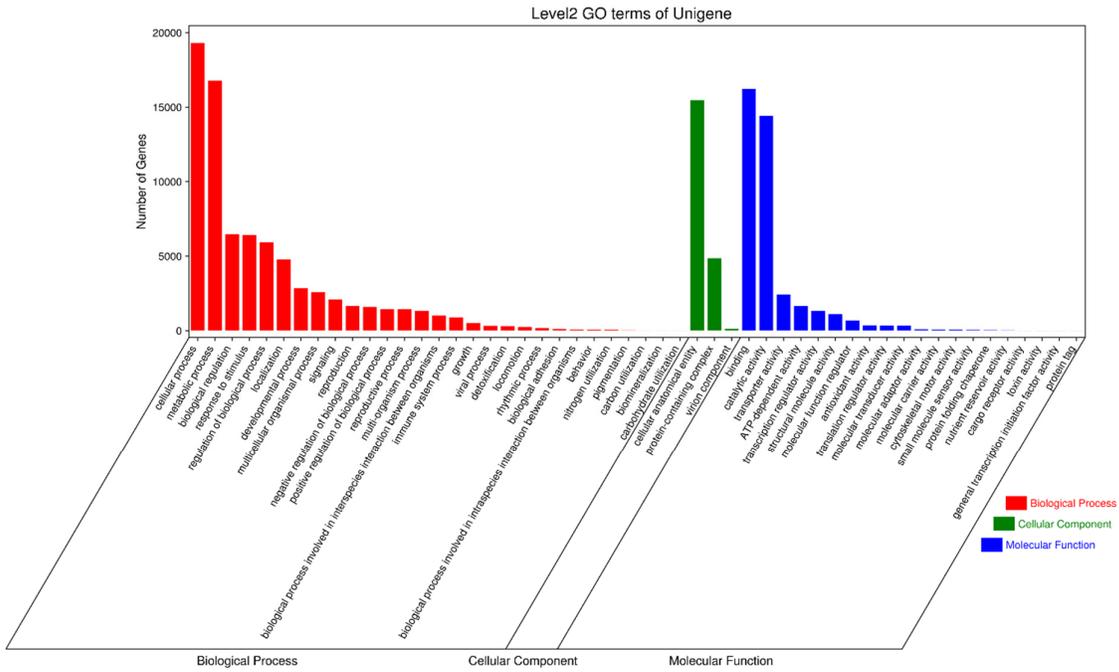


Figure S1. Gene ontology (GO) classification analysis different expression genes (DEGs) during the current-year flower development in *Hydrangea arborescens* ‘Annabelle’

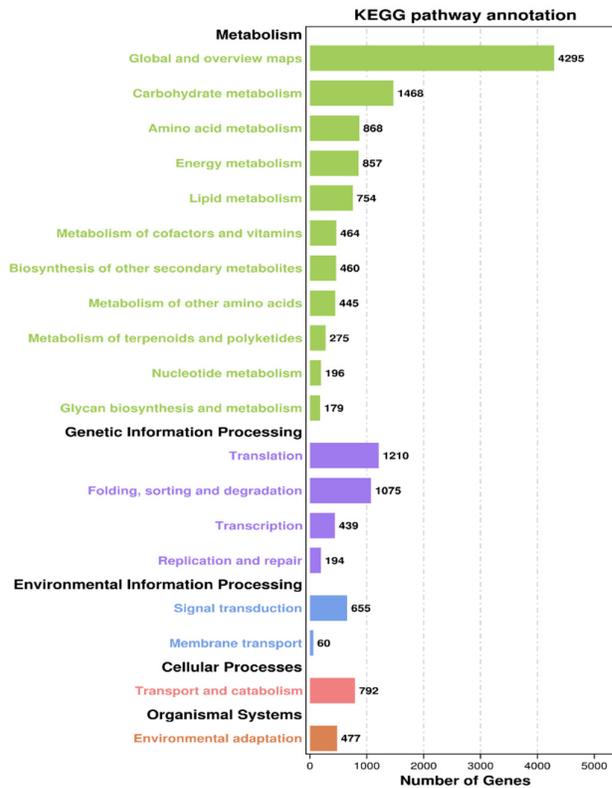
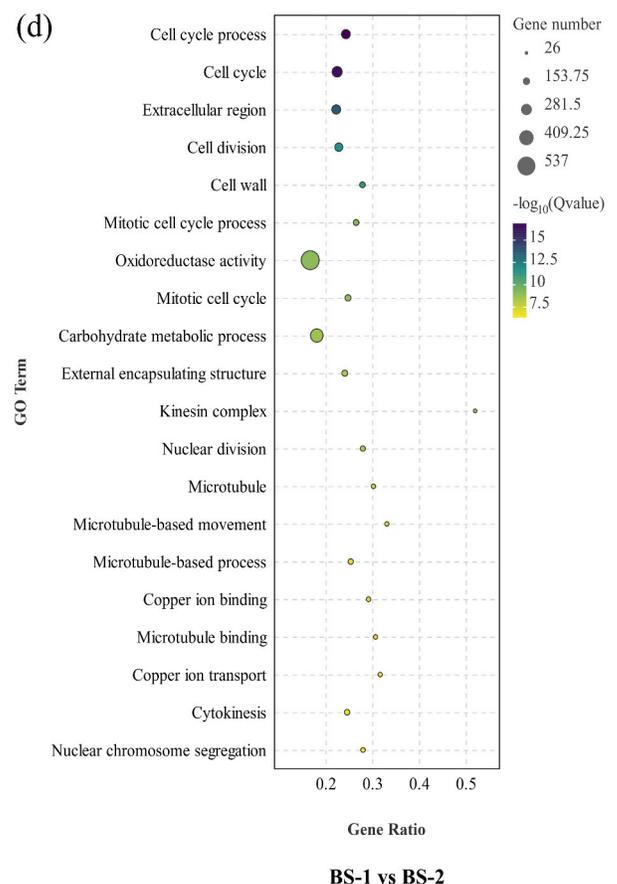
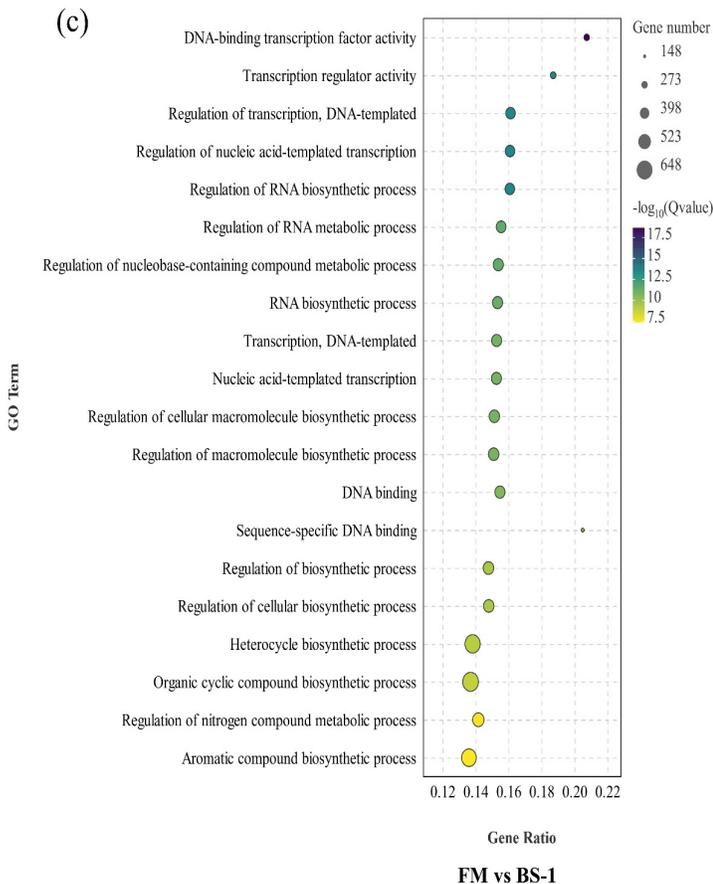
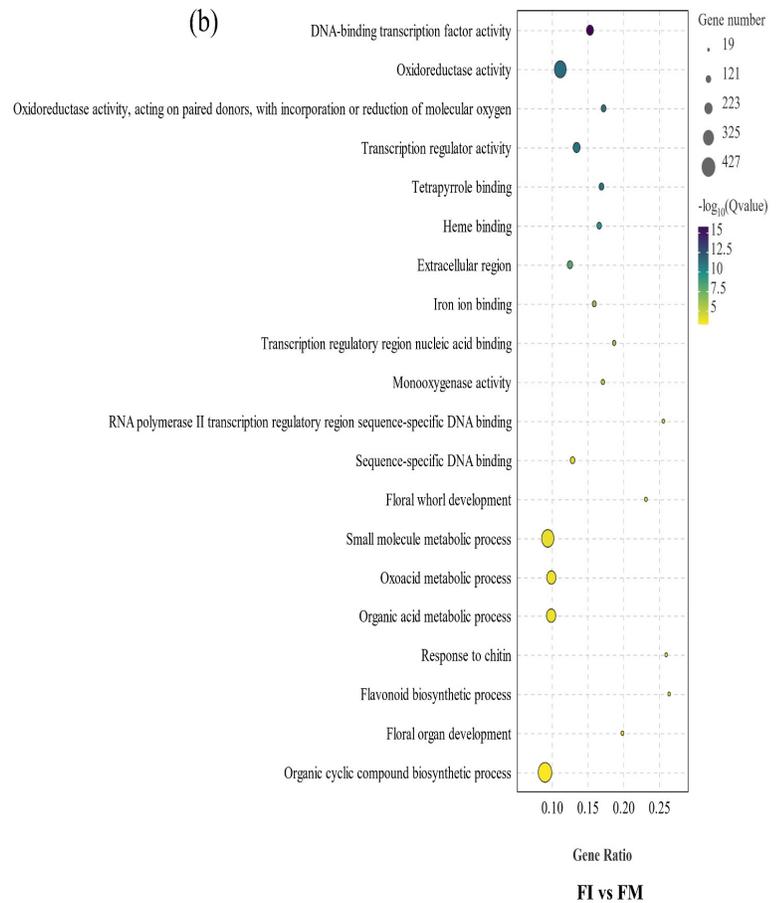
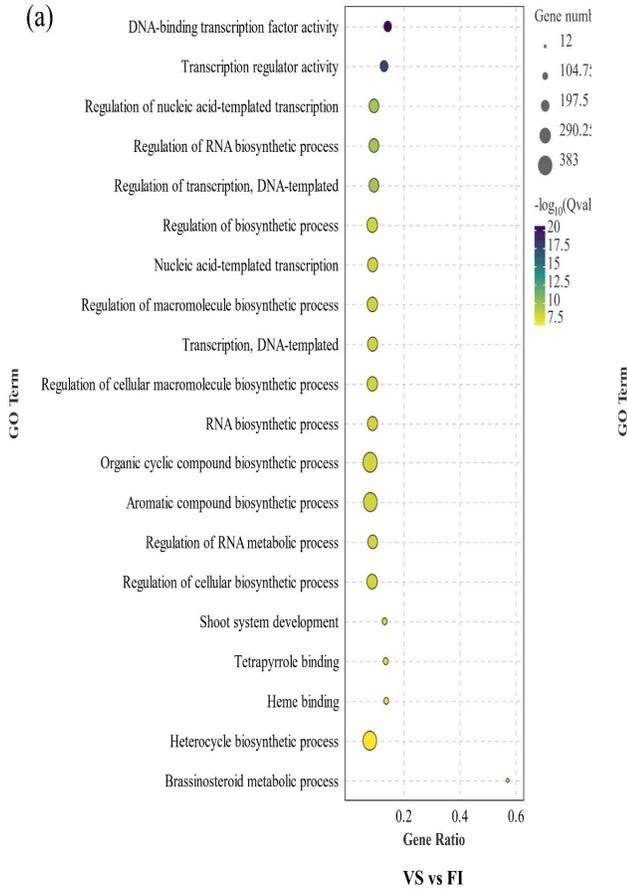


Figure S2. KEGG classification analysis different expression genes (DEGs) during the current-year flower development in *Hydrangea arborescens* ‘Annabelle’



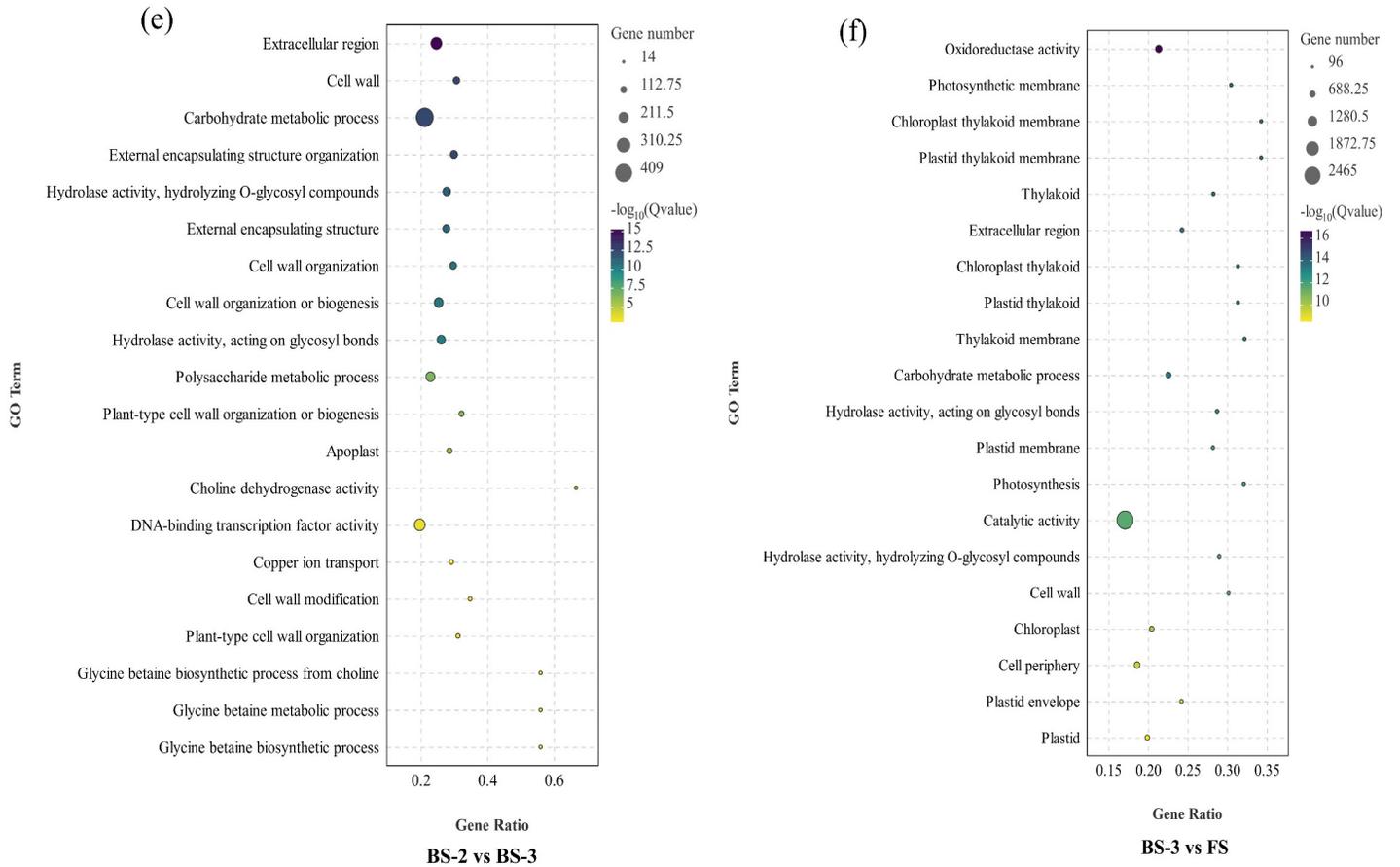


Figure S3. GO term enrichment analysis of DEGs during flower development

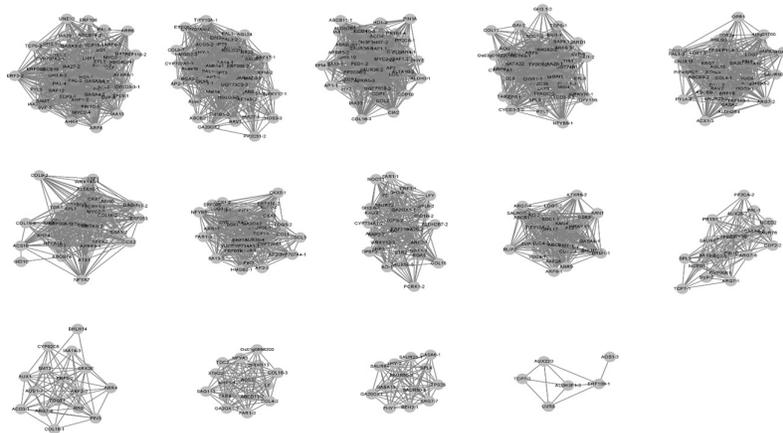


Figure S4. Summary of MCL gene co-expression network (GCN) . The network consisted of 14 biological meaningful clusters.