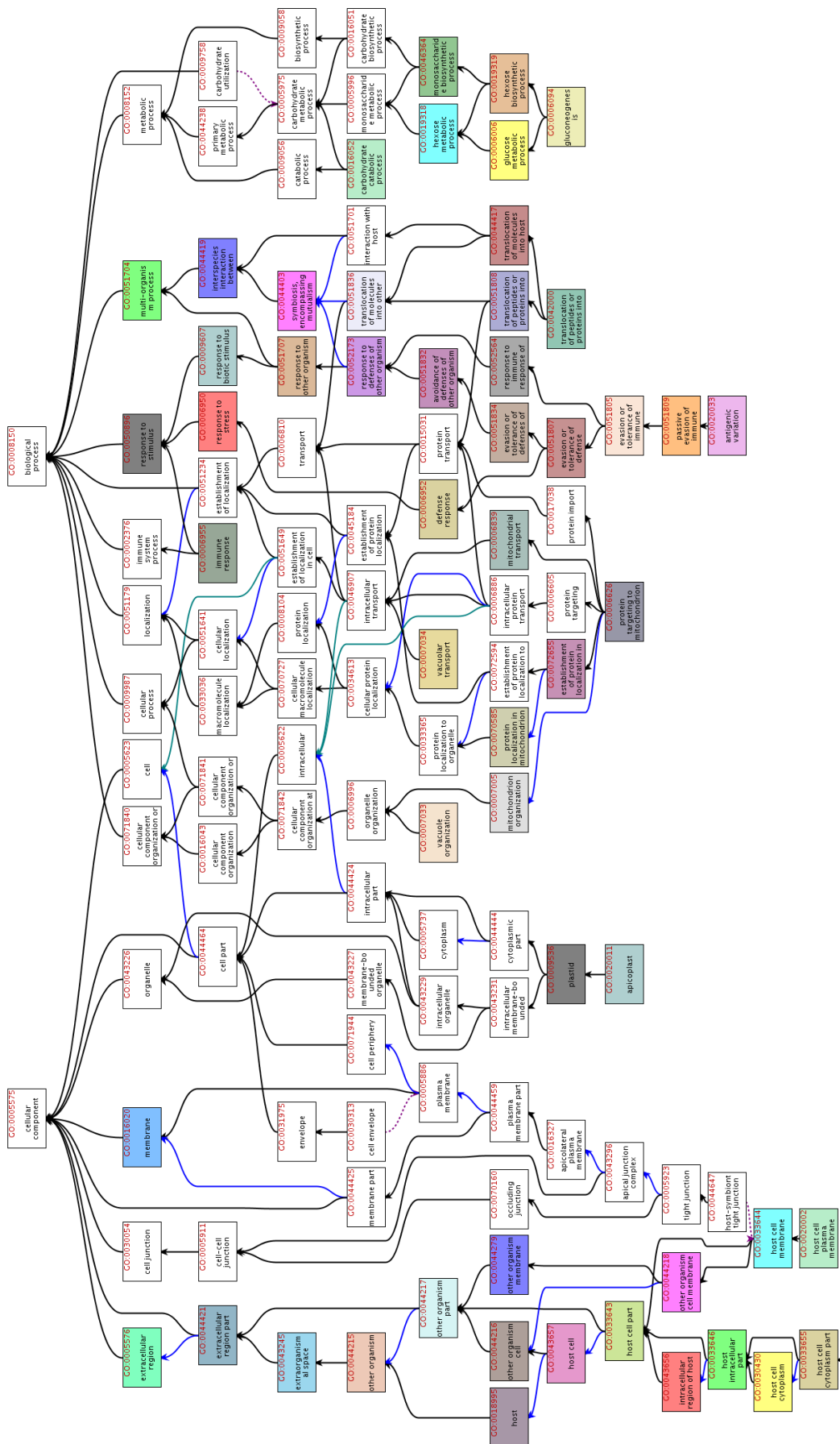
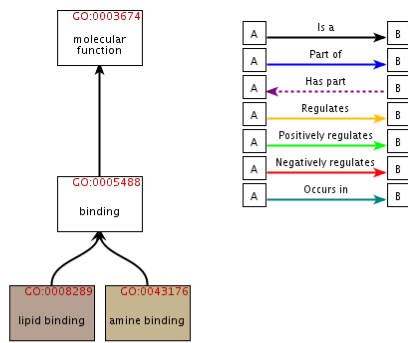


Additional file 3: Hierarchical classification of Gene Ontology terms enriched during the transcriptomic experiment.





QuickGO was used to produce this Ancestral Chart representing significantly overrepresented GOs. This graph is divided into three groups: biological process, cellular component and molecular function.