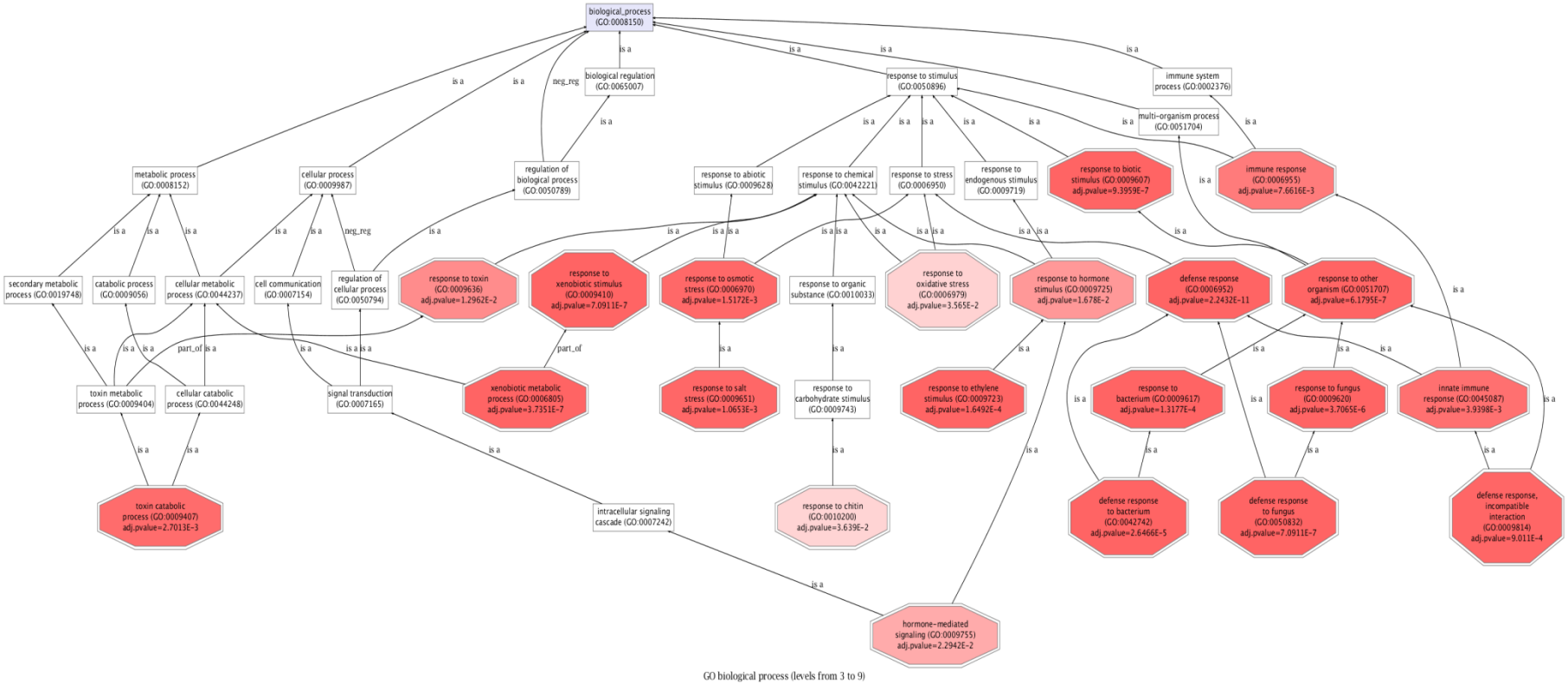Figure S1. Directed acyclic hierarchical graph (DAG) of gene ontology terms significantly over-represented in the genes upregulated in *35S:ERF5* or *35S:ERF6* plants compared to the rest of the Arabidopsis genome. The DAG was generated using FatiGO (http://babelomics.bioinfo.cipf.es). GO terms in red are significantly over-represented in the dataset