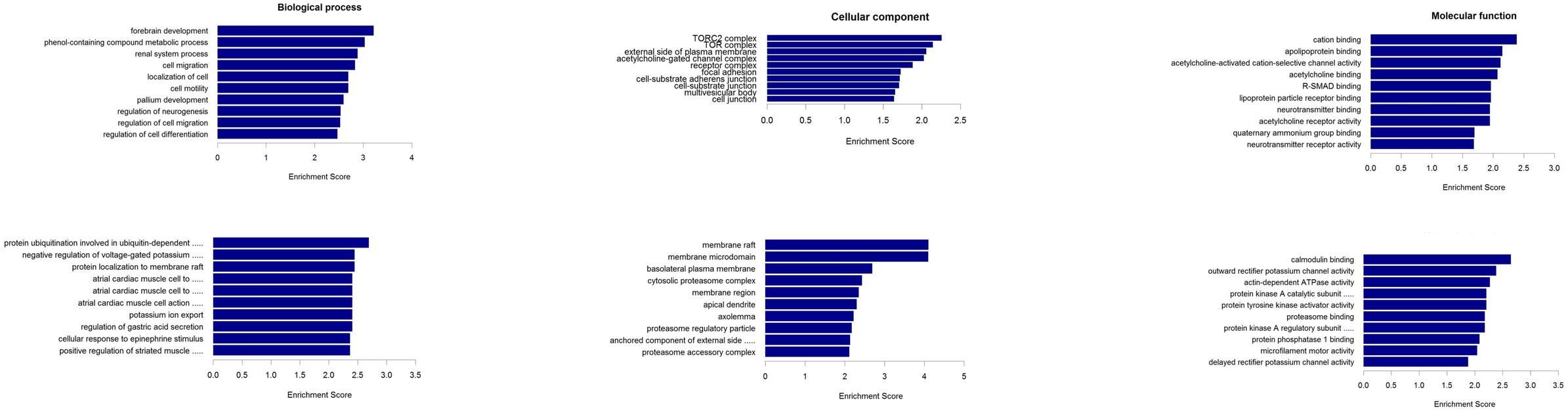

Figure S GO analyses of the differentially expressed lncRNAs

The ontology covers three domains: biological process, cellular component, and molecular function. Target genes of upregulated lncRNAs by GO enrichment analysis (top row). Target genes of downregulated lncRNAs by GO enrichment analysis (bottom row).