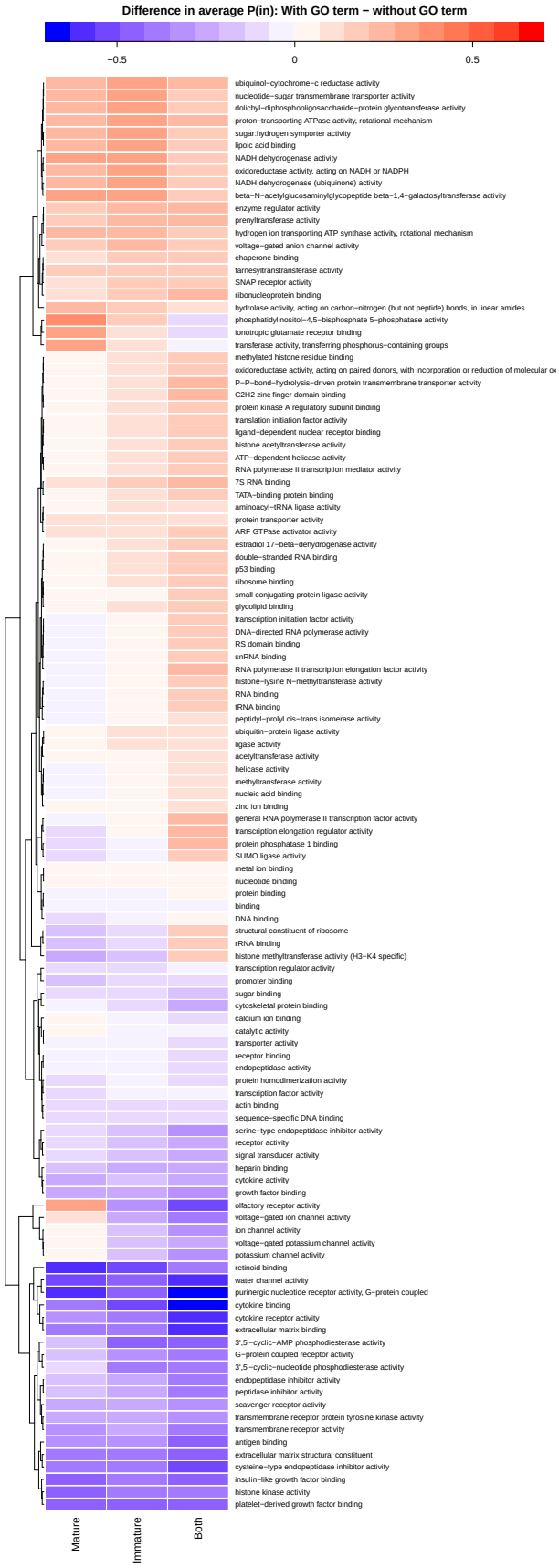

Supplemental Figure 4: Heatmap of average differences in estimated P(in) probabilities between genes annotated with the specified Gene Ontology (GO) molecular function term and genes lacking the term. Probability differences are separated by cell type category. Only terms with significant ($q < 0.05$) differences are listed.