



Figure 2. Comparison of enriched ontology clusters among top 150-ranked proteins upregulated 2 h after OFC between OFC-positive and OFC-negative group

We performed enrichment analysis using the top 150-ranked proteins upregulated 2 h after OFC. We selected the term with the lowest p-value within each cluster as the representative term in the dendrogram. The heatmap cells are colored according to their p-values, and grey cells indicate a lack of enrichment for that term in the corresponding gene list.