Figure S11

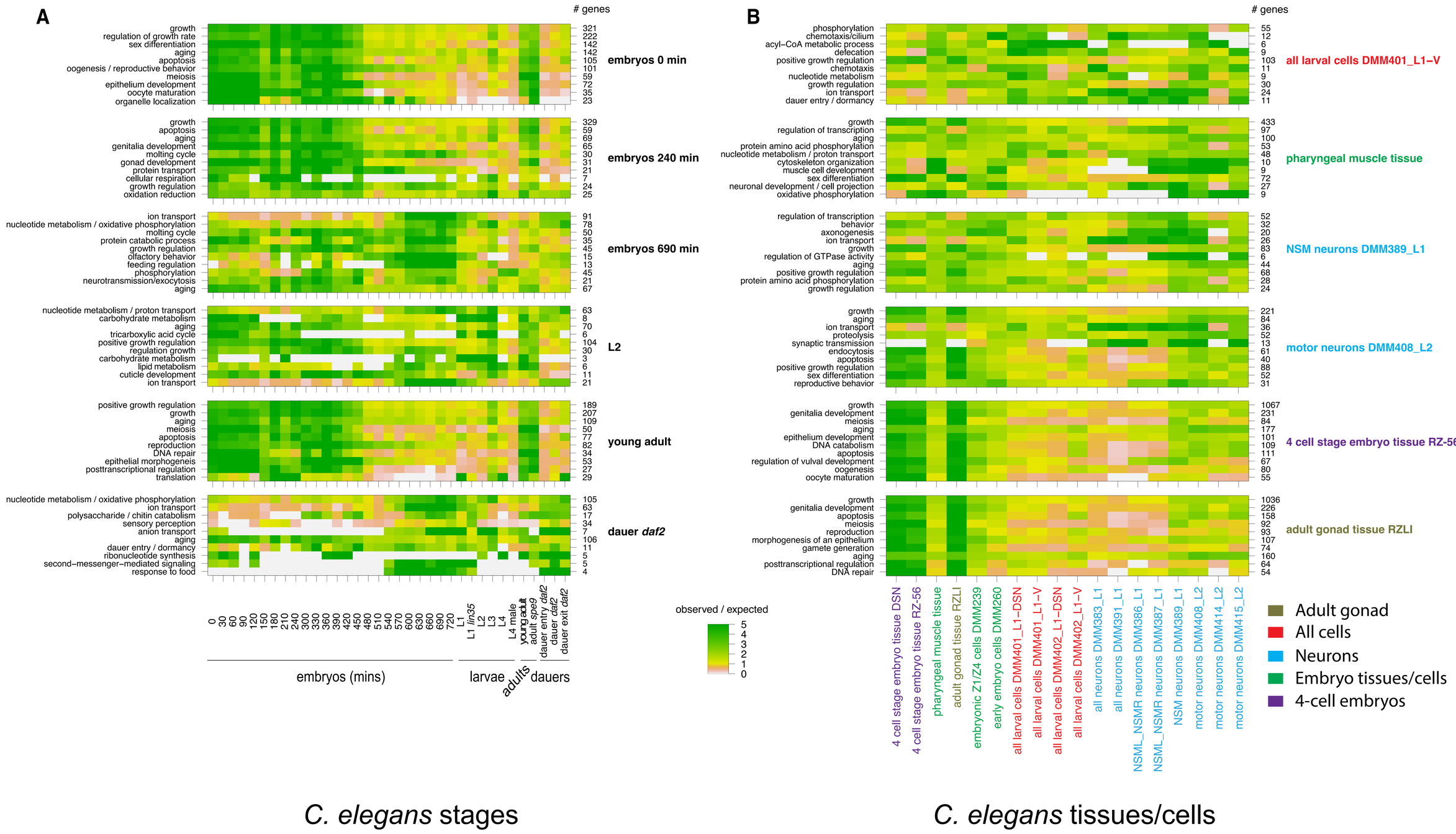


Figure S11. Top 10 enriched GO clusters in (A) six example worm developmental stages and (B) six example worm tissues/dissected cells. The heatmaps show the ratios of observed gene counts / expected gene counts of every GO cluster in every stage or tissue/dissected cell. The observed gene counts of the GO clusters in every example sample (stage or tissue/dissected cell) are also listed.