

Figure S10

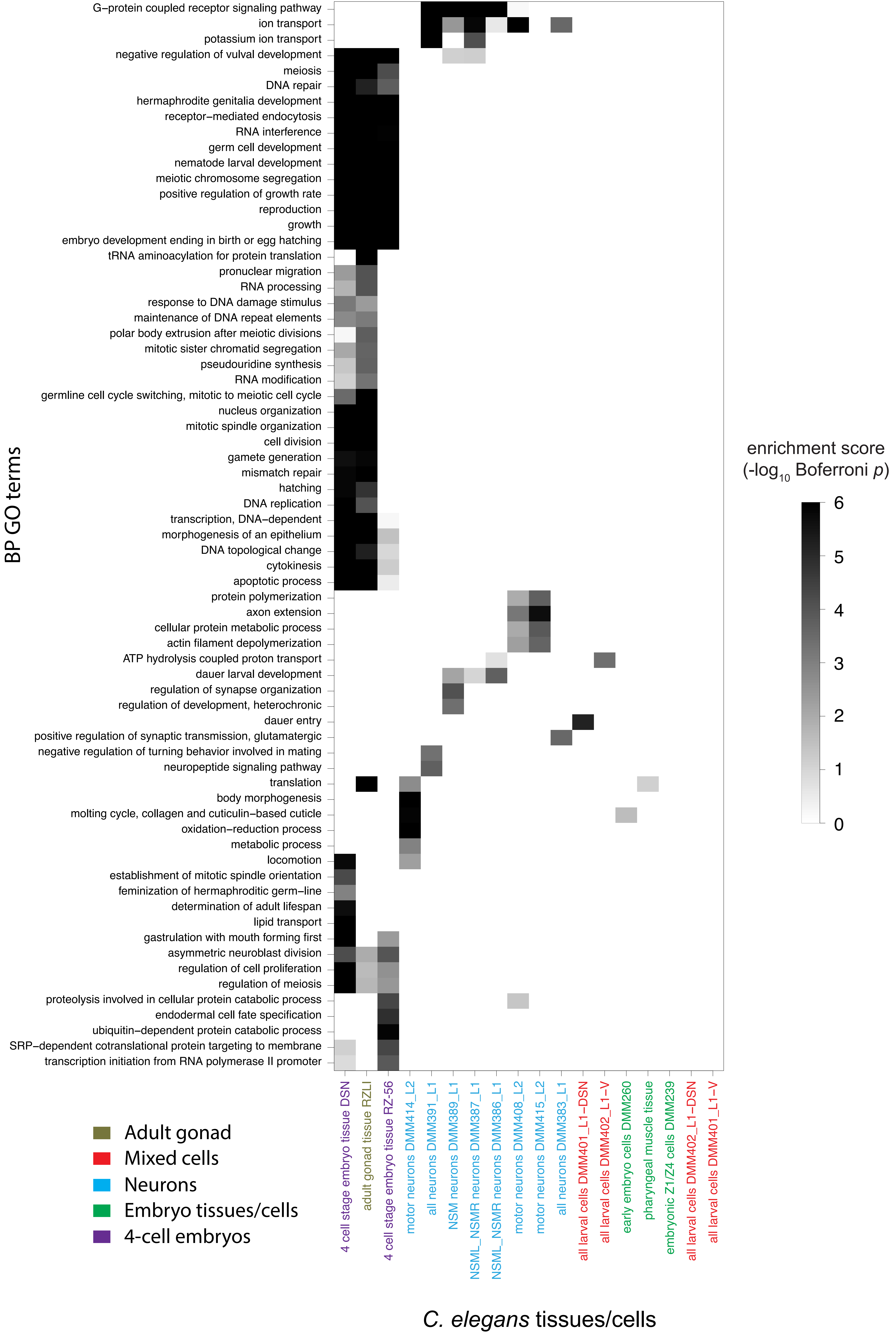


Figure S10. Enrichment patterns of top enriched biological process gene ontology (BP GO) terms across *C. elegans* tissues and dissected cells. The enrichment scores are  $-\log_{10}$  transformed Bonferroni corrected  $p$ -values from hypergeometric tests (see Methods). Hierarchical clustering was applied to order the tissues/cells. Tissues and dissected cells from similar organs are marked with colors. For detailed information on the stage/tissue/cell labels, please refer to Table S2.