

Figure S6

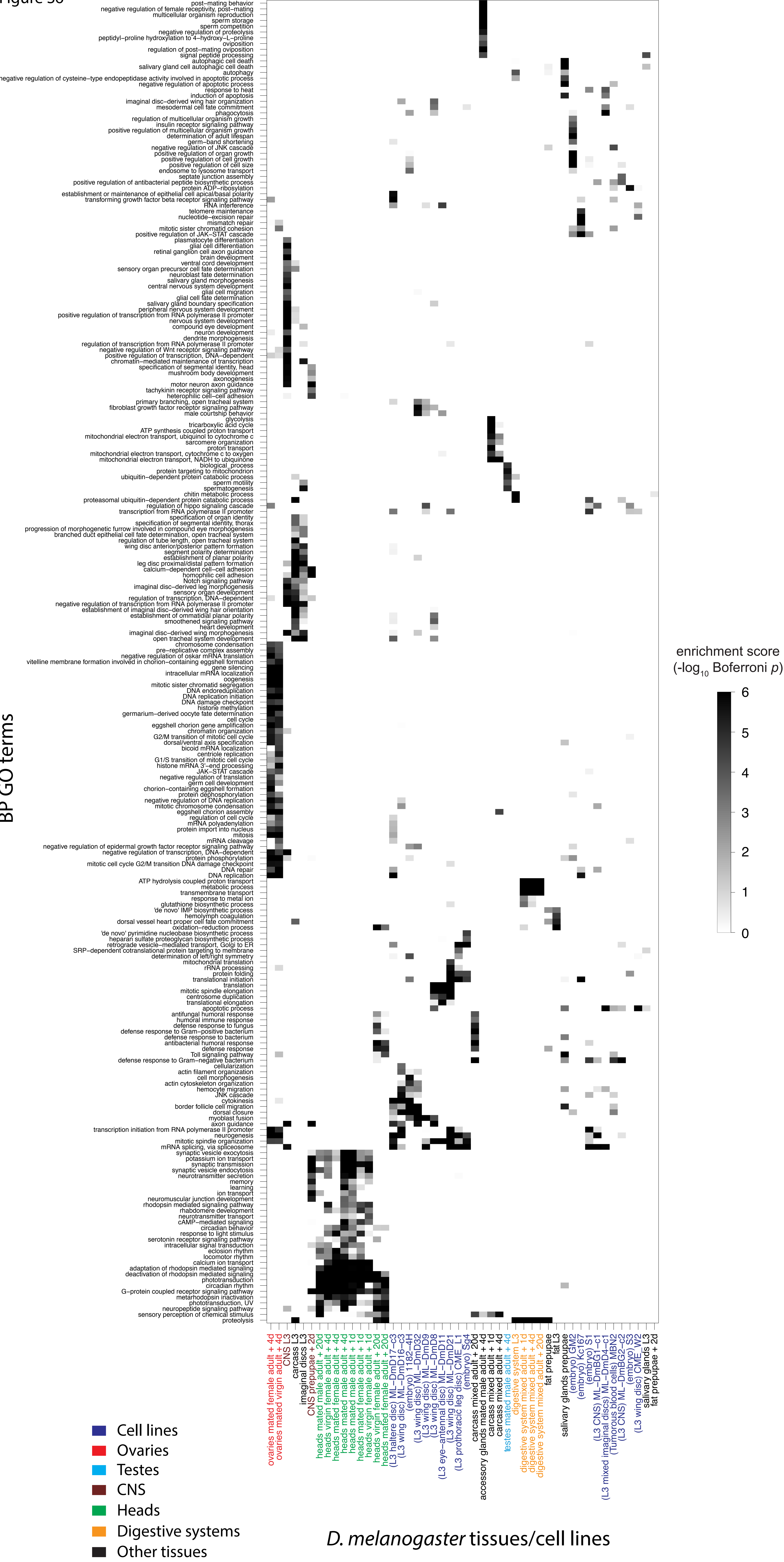


Figure S6. Enrichment patterns of top enriched biological process gene ontology (BP GO) terms across *D. melanogaster* tissues and cell lines. The enrichment scores are $-\log_{10}$ transformed Bonferroni corrected p -values from hypergeometric tests (see Methods). Hierarchical clustering was applied to order the tissues/cell lines. Tissues from similar organs and cell lines are marked with colors. For detailed information on the stage/tissue/cell labels, please refer to Table S2.