



Figure S14. Correlation analysis approaches for comparing developmental stages within *D. melanogaster* and *C. elegans*. Pearson and Spearman correlations are calculated based on all protein-coding gene expression levels in *D. melanogaster* or *C. elegans*. Larger correlations are shown as darker color, corresponding to the scale next to each panel. (A) Pearson correlation for every pair of *D. melanogaster* stages. (B) Pearson correlation for every pair of *C. elegans* stages. (C) Spearman correlation for every pair of *D. melanogaster* stages. (D) Spearman correlation for every pair of *C. elegans* stages. For detailed information on the stage labels, please refer to Table S2.