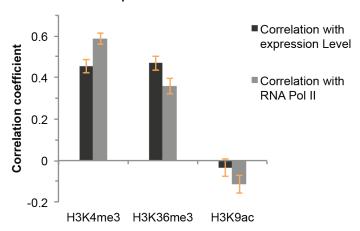
Camponotus floridanus



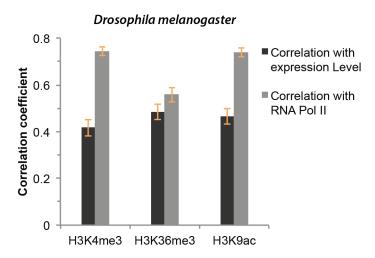


Figure S3. Correlations between transcriptional activity and the histone modifications **H3K4me3**, **H3K36me3**, and **H3K9ac**. Pearson's correlations with 95% confidence intervals are shown for data from *C. floridanus* and *D. melanogaster* (n = 2102 common ortholog groups).