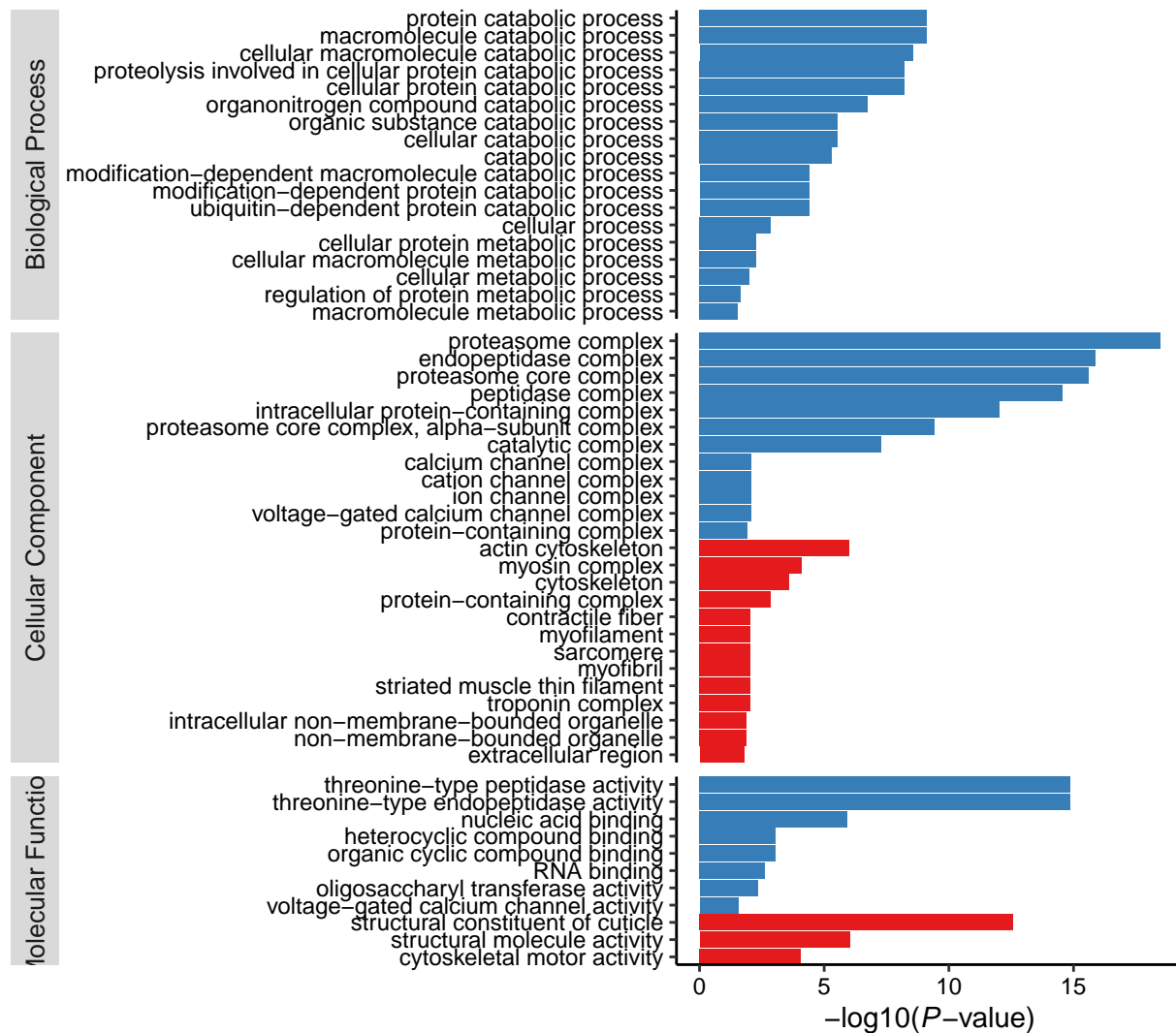


A. Gene ontology (GO) term enrichment analysis



B. Distribution of co-expressed transcripts modules

