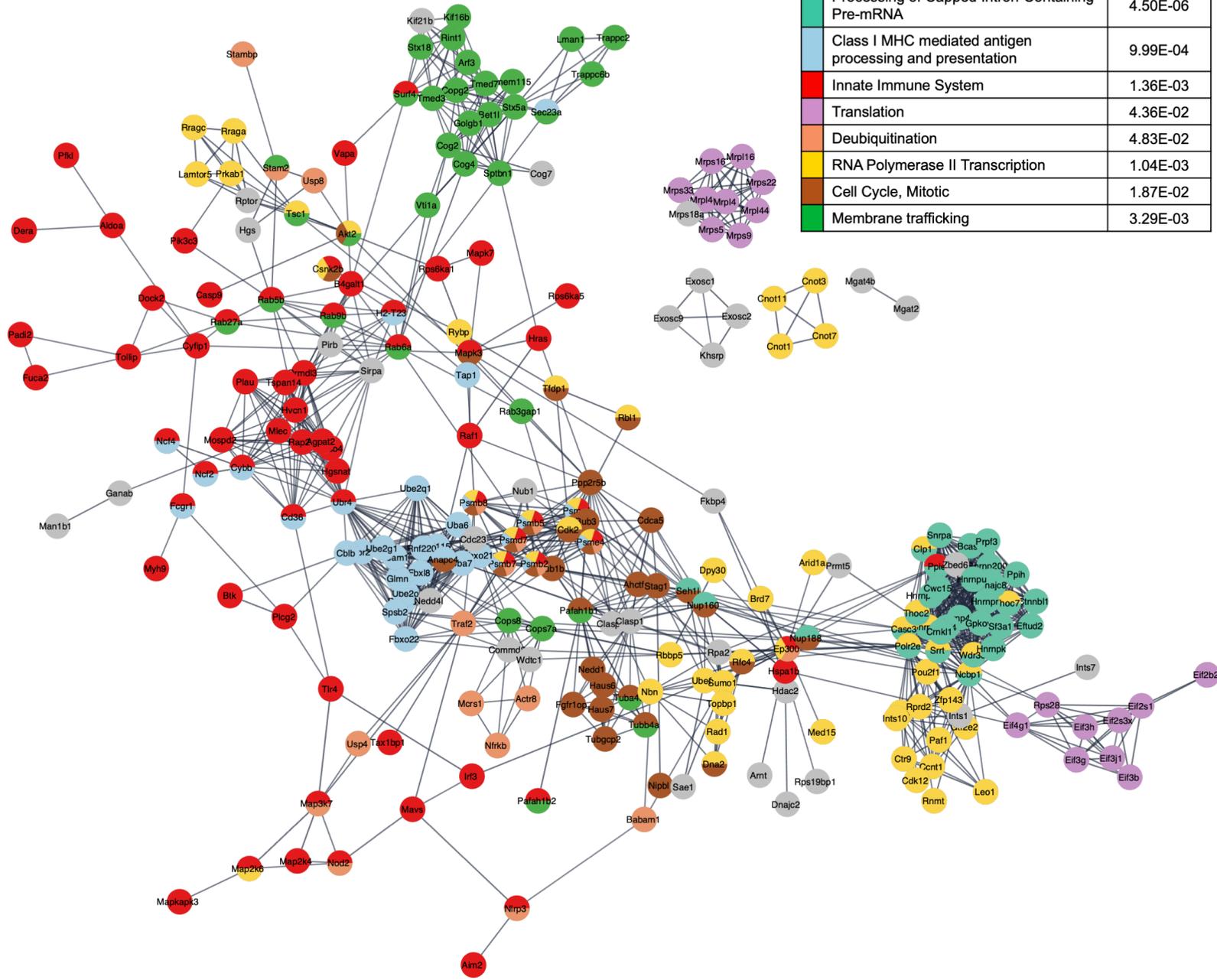


Supplemental Data S1. Protein-Protein Interaction Network of the Late wave (“Supplemental_Data_S1.pdf”). A high-resolution overview of the late active wave (CT15-3) protein-protein interaction network with individual gene symbols displayed on each node. Strings shown are high confidence (>.9) interactions and node color corresponds to a gene’s enriched reactome category. Grey coloring represents genes from enriched reactome categories that did not represent a large proportion of the protein-protein interactome or belonged to redundant categories.



Reactome pathway name	Enrichment
Processing of Capped Intron-Containing Pre-mRNA	4.50E-06
Class I MHC mediated antigen processing and presentation	9.99E-04
Innate Immune System	1.36E-03
Translation	4.36E-02
Deubiquitination	4.83E-02
RNA Polymerase II Transcription	1.04E-03
Cell Cycle, Mitotic	1.87E-02
Membrane trafficking	3.29E-03