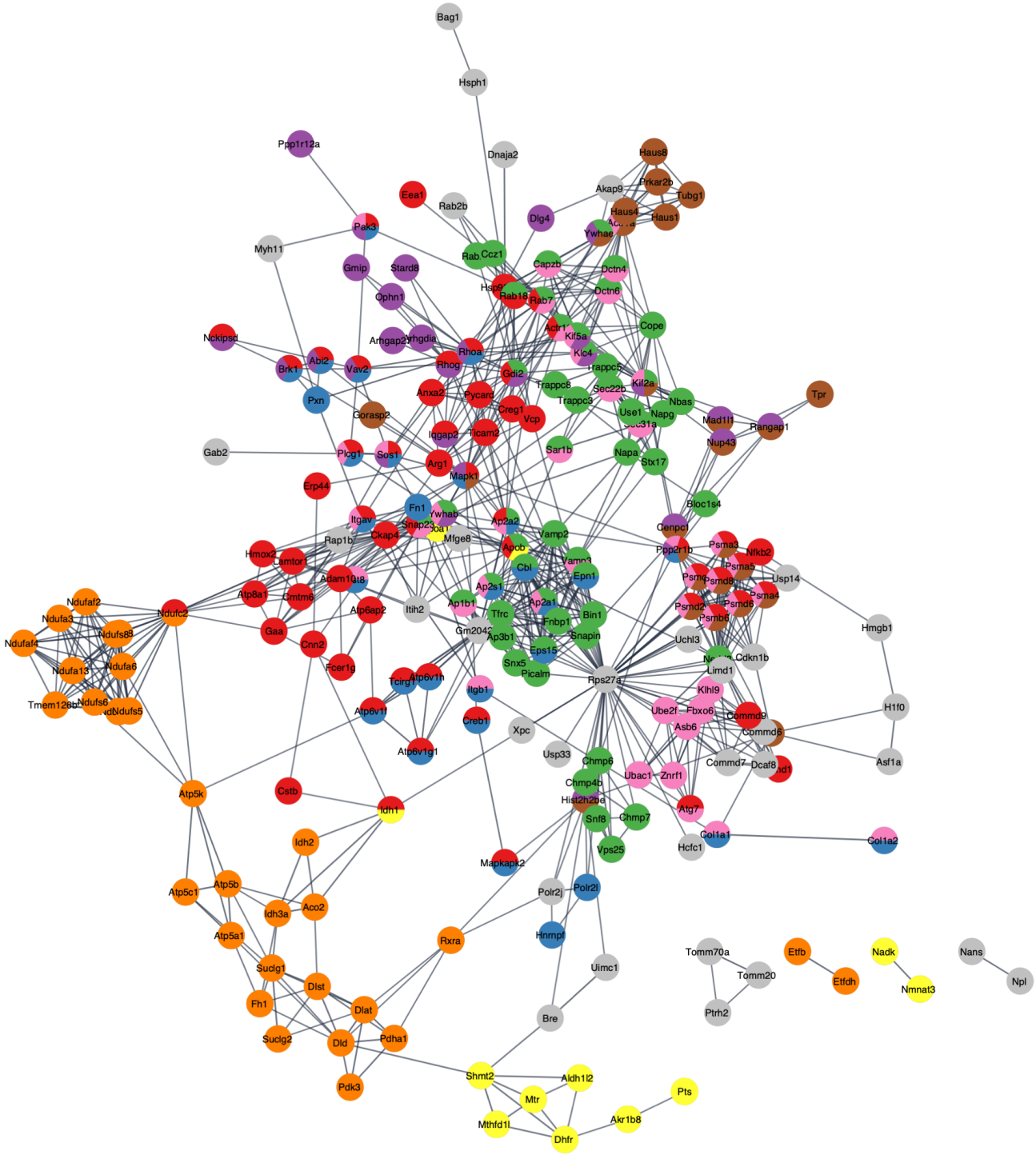
Supplemental Data S2. Protein-protein interaction network of the Early wave ("Supplemental_Data_S2.pdf"). A high-resolution overview of the late inactive wave (CT3-15) protein-protein interaction network with individual gene symbols displayed on each node. Strings shown are high confidence (>.9) interactions 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
	Signaling by Rho GTPases	1.79E-02
	M phase	4.82E-03
	Innate Immune System	6.41E-05
	Adaptive Immune System	1.74E-03
	The citric acid cycle and respiratory electron transport	7.00E-03
	Metabolism of vitamins and cofactors	9.50E-04
	Signaling by receptor tyrosine kinases	7.90E-03
	Membrane trafficking	4.02E-09