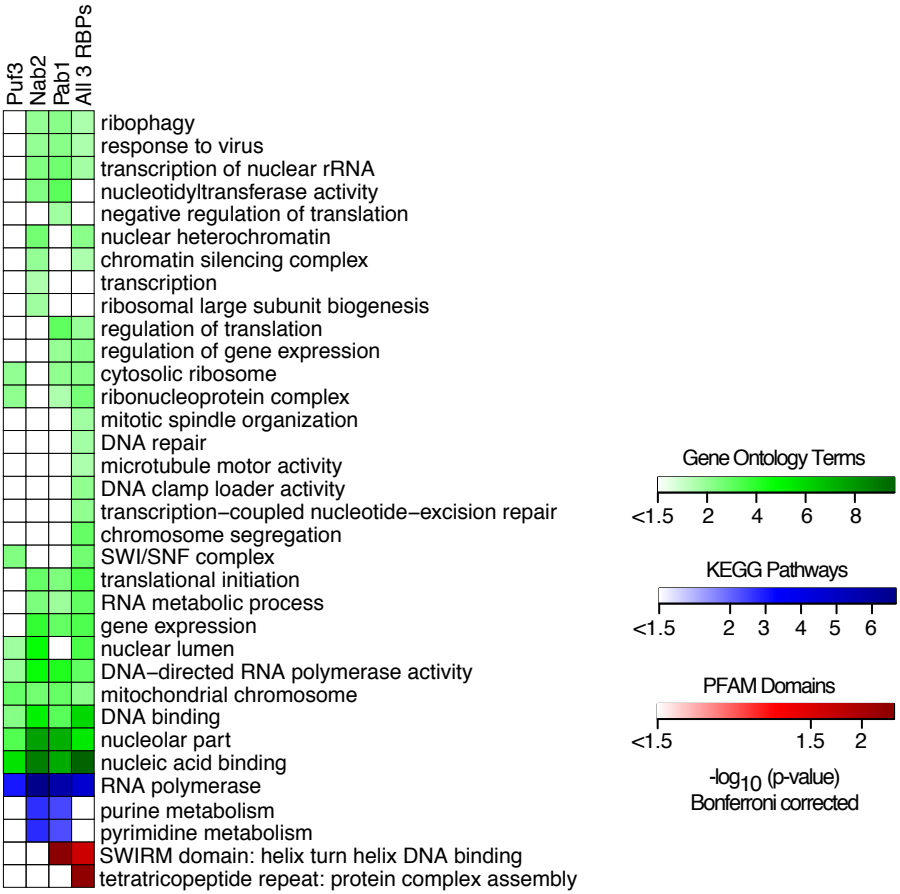
Puf3
Nab2
Pab1
All 3 RBPs
ribophagy
response to virus
transcription of nuclear rRNA
nucleotidyltransferase activity
negative regulation of translation
nuclear heterochromatin
chromatin silencing complex
transcription
ribosomal large subunit biogenesis
regulation of translation
regulation of gene expression
cytosolic ribosome
ribonucleoprotein complex
mitotic spindle organization
DNA repair
microtubule motor activity
DNA clamp loader activity
transcription−coupled nucleotide−excision repair
chromosome segregation
SWI/SNF complex
translational initiation
RNA metabolic process
gene expression
nuclear lumen
DNA−directed RNA polymerase activity
mitochondrial chromosome
DNA binding
nucleolar part
nucleic acid binding
RNA polymerase
purine metabolism
pyrimidine metabolism
SWIRM domain: helix turn helix DNA binding
tetratricopeptide repeat: protein complex assembly
Gene Ontology Terms
<1.5 2 4 6 8
KEGG Pathways
<1.5 2 3 4 5 6
PFAM Domains
<1.5 1.5 2
-log10 (p-value)
Bonferroni corrected