

F\_NPM1, MORF\_NPM1

MORF\_ACTG1, MORF\_ACTG1  
GSE41978\_KLRG1\_HIGH\_VS\_LOW\_EFFECTOR\_CD8\_TCELL\_DN, GSE41978\_KLRG1\_HIGH\_VS\_LOW\_EFFECTOR\_CD8\_TCELL\_DN  
MORF\_TPT1, MORF\_TPT1  
GNF2\_NPM1, GNF2\_NPM1  
REACTOME\_3\_UTR\_MEDIATED\_TRANSLATIONAL\_REGULATION, REACTOME\_3\_UTR\_MEDIATED\_TRANSLATIONAL\_REGULATION  
REACTOME\_INFLUENZA\_VIRAL\_RNA\_TRANSCRIPTION\_AND\_REPLICATION, REACTOME\_INFLUENZA\_VIRAL\_RNA\_TRANSCRIPTION\_AND\_REPLICATION  
REACTOME\_TRANSLATION, REACTOME\_TRANSLATION  
GO\_CYTOSOLIC\_RIBOSOME, GO\_CYTOSOLIC\_RIBOSOME  
MORF\_NME2, MORF\_NME2  
REACTOME\_PEPTIDE\_CHAIN\_ELONGATION, REACTOME\_PEPTIDE\_CHAIN\_ELONGATION  
GO\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_ENDOPLASMIC\_RETICULUM, GO\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_ENDOPLASMIC\_RETICULUM  
GNF2\_EIF3S6, GNF2\_EIF3S6  
KEGG\_RIBOSOME, KEGG\_RIBOSOME  
GO\_TRANSLATIONAL\_INITIATION, GO\_TRANSLATIONAL\_INITIATION  
REACTOME\_SRP\_DEPENDENT\_COTRANSLATIONAL\_PROTEIN\_TARGETING\_TO\_MEMBRANE, REACTOME\_SRP\_DEPENDENT\_COTRANSLATIONAL\_PROTEIN\_TARGETING\_TO\_MEMBRANE  
GNF2\_ST13, GNF2\_ST13  
GO\_CYTOSOLIC\_LARGE\_RIBOSOMAL\_SUBUNIT, GO\_CYTOSOLIC\_LARGE\_RIBOSOMAL\_SUBUNIT  
GSE2405\_0H\_VS\_24H\_A\_PHAGOCYTOPHILUM\_STIM\_NEUTROPHIL\_UP, GSE2405\_0H\_VS\_24H\_A\_PHAGOCYTOPHILUM\_STIM\_NEUTROPHIL\_UP  
GSE14000\_TRANSLATED\_RNA\_VS\_MRNA\_DC\_DN, GSE14000\_TRANSLATED\_RNA\_VS\_MRNA\_DC\_DN  
GO\_RIBOSOMAL\_SUBUNIT, GO\_RIBOSOMAL\_SUBUNIT  
GO\_PROTEIN\_LOCALIZATION\_TO\_ENDOPLASMIC\_RETICULUM, GO\_PROTEIN\_LOCALIZATION\_TO\_ENDOPLASMIC\_RETICULUM  
GCM\_TPT1, GCM\_TPT1  
GO\_MYELIN\_SHEATH, GO\_MYELIN\_SHEATH  
GO\_LARGE\_RIBOSOMAL\_SUBUNIT, GO\_LARGE\_RIBOSOMAL\_SUBUNIT  
MORF\_EIF4A2, MORF\_EIF4A2  
REACTOME\_INFLUENZA\_LIFE\_CYCLE, REACTOME\_INFLUENZA\_LIFE\_CYCLE  
GSE41978\_ID2\_KO\_VS\_ID2\_KO\_AND\_BIM\_KO\_KLRG1\_LOW\_EFFECTOR\_CD8\_TCELL\_DN, GSE41978\_ID2\_KO\_VS\_ID2\_KO\_AND\_BIM\_KO\_KLRG1\_LOW\_EFFECTOR\_CD8\_TCELL\_DN  
GNF2\_DAP3, GNF2\_DAP3  
GO\_PROTEIN\_TARGETING\_TO\_MEMBRANE, GO\_PROTEIN\_TARGETING\_TO\_MEMBRANE  
GSE42088\_UNINF\_VS\_LEISHMANIA\_INF\_DC\_2H\_DN, GSE42088\_UNINF\_VS\_LEISHMANIA\_INF\_DC\_2H\_DN  
KEGG\_PATHOGENIC\_ESCHERICHIA\_COLI\_INFECTION, KEGG\_PATHOGENIC\_ESCHERICHIA\_COLI\_INFECTION  
GNF2\_FBL, GNF2\_FBL  
GSE3039\_CD4\_TCELL\_VS\_B2\_BCELL\_DN, GSE3039\_CD4\_TCELL\_VS\_B2\_BCELL\_DN  
GO\_MULTI\_ORGANISM\_METABOLIC\_PROCESS, GO\_MULTI\_ORGANISM\_METABOLIC\_PROCESS  
MORF\_ANP32B, MORF\_ANP32B  
REACTOME\_NONSENSE\_MEDIATED\_DECAY\_ENHANCED\_BY\_THE\_EXON\_JUNCTION\_COMPLEX, REACTOME\_NONSENSE\_MEDIATED\_DECAY\_ENHANCED\_BY\_THE\_EXON\_JUNCTION\_COMPLEX  
GCM\_NPM1, GCM\_NPM1  
DANG\_MYC\_TARGETS\_UP, DANG\_MYC\_TARGETS\_UP  
GNF2\_TPT1, GNF2\_TPT1  
GSE25123\_WT\_VS\_PPARG\_KO\_MACROPHAGE\_DN, GSE25123\_WT\_VS\_PPARG\_KO\_MACROPHAGE\_DN  
GSE7460\_TREG\_VS\_TCONV\_ACT\_WITH\_TGFB\_DN, GSE7460\_TREG\_VS\_TCONV\_ACT\_WITH\_TGFB\_DN  
MORF\_FBL, MORF\_FBL  
GCM\_CSNK2B, GCM\_CSNK2B  
GSE5099\_MONOCYTE\_VS\_CLASSICAL\_M1\_MACROPHAGE\_DN, GSE5099\_MONOCYTE\_VS\_CLASSICAL\_M1\_MACROPHAGE\_DN  
GSE2405\_0H\_VS\_9H\_A\_PHAGOCYTOPHILUM\_STIM\_NEUTROPHIL\_DN, GSE2405\_0H\_VS\_9H\_A\_PHAGOCYTOPHILUM\_STIM\_NEUTROPHIL\_DN  
GSE42724\_NAIVE\_VS\_B1\_BCELL\_DN, GSE42724\_NAIVE\_VS\_B1\_BCELL\_DN  
GO\_UNFOLDED\_PROTEIN\_BINDING, GO\_UNFOLDED\_PROTEIN\_BINDING  
PECE\_MAMMARY\_STEM\_CELL\_UP, PECE\_MAMMARY\_STEM\_CELL\_UP  
GO\_NUCLEAR\_TRANSCRIBED\_MRNA\_CATABOLIC\_PROCESS\_NONSENSE\_MEDIATED\_DECAY, GO\_NUCLEAR\_TRANSCRIBED\_MRNA\_CATABOLIC\_PROCESS\_NONSENSE\_MEDIATED\_DECAY  
BILANGES\_RAPAMYCIN\_SENSITIVE\_VIA\_TSC1\_AND\_TSC2, BILANGES\_RAPAMYCIN\_SENSITIVE\_VIA\_TSC1\_AND\_TSC2