

M\_RAB10, GCM\_RAB10

GSE24574\_NAIVE\_VS\_TCONV\_CD4\_TCELL\_DN, GSE24574\_NAIVE\_VS\_TCONV\_CD4\_TCELL\_DN  
GCM\_RAN, GCM\_RAN  
GCM\_GSPT1, GCM\_GSPT1  
GSE39556\_CD8A\_DC\_VS\_NK\_CELL\_MOUSE\_3H\_POST\_POLYIC\_INJ\_DN, GSE39556\_CD8A\_DC\_VS\_NK\_CELL\_MOUSE\_3H\_POST\_POLYIC\_INJ\_DN  
GCM\_MAP4K4, GCM\_MAP4K4  
GSE17721\_LPS\_VS\_GARDIQUIMOD\_4H\_BMDC\_DN, GSE17721\_LPS\_VS\_GARDIQUIMOD\_4H\_BMDC\_DN  
GSE13229\_MATURE\_VS\_INTMATURE\_NKCELL\_DN, GSE13229\_MATURE\_VS\_INTMATURE\_NKCELL\_DN  
GSE17721\_LPS\_VS\_GARDIQUIMOD\_16H\_BMDC\_DN, GSE17721\_LPS\_VS\_GARDIQUIMOD\_16H\_BMDC\_DN  
GSE21546\_ELK1\_KO\_VS\_SAP1A\_KO\_AND\_ELK1\_KO\_ANTI\_CD3\_STIM\_DP\_THYMOCYTES\_UP, GSE21546\_ELK1\_KO\_VS\_SAP1A\_KO\_AND\_ELK1\_KO\_ANTI\_CD3\_STIM\_DP\_THYMOCYTES\_UP  
GCM\_NCAM1, GCM\_NCAM1  
GCM\_BMPR2, GCM\_BMPR2  
GCM\_RBM8A, GCM\_RBM8A  
GCM\_PTK2, GCM\_PTK2  
GO\_ERAD\_PATHWAY, GO\_ERAD\_PATHWAY  
GCM\_CALM1, GCM\_CALM1  
MODULE\_323, MODULE\_323  
GSE33292\_WT\_VS\_TCF1\_KO\_DN3\_THYMOCYTE\_UP, GSE33292\_WT\_VS\_TCF1\_KO\_DN3\_THYMOCYTE\_UP  
GO\_ER\_ASSOCIATED\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GO\_ER\_ASSOCIATED\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS  
GSE4748\_CTRL\_VS\_LPS\_STIM\_DC\_3H\_UP, GSE4748\_CTRL\_VS\_LPS\_STIM\_DC\_3H\_UP  
GCM\_DLG1, GCM\_DLG1  
GCM\_CRKL, GCM\_CRKL  
GGGCATT\_MIR365, GGGCATT\_MIR365  
GSE6259\_33D1\_POS\_DC\_VS\_CD4\_TCELL\_DN, GSE6259\_33D1\_POS\_DC\_VS\_CD4\_TCELL\_DN  
ATGTTTC\_MIR494, ATGTTTC\_MIR494  
BYSTROEM\_CORRELATED\_WITH\_IL5\_DN, BYSTROEM\_CORRELATED\_WITH\_IL5\_DN  
GO\_POST\_GOLGI\_VESICLE\_MEDIATED\_TRANSPORT, GO\_POST\_GOLGI\_VESICLE\_MEDIATED\_TRANSPORT  
GCM\_GSTA4, GCM\_GSTA4  
GO\_GOLGI\_TO\_VACUOLE\_TRANSPORT, GO\_GOLGI\_TO\_VACUOLE\_TRANSPORT  
GCM\_MAP1B, GCM\_MAP1B  
GCM\_SIRT2, GCM\_SIRT2  
GSE32901\_TH1\_VS\_TH17\_NEG\_CD4\_TCELL\_UP, GSE32901\_TH1\_VS\_TH17\_NEG\_CD4\_TCELL\_UP  
GSE10500\_ARTHRITIC\_SYNOVIAL\_FLUID\_VS\_HEALTHY\_MACROPHAGE\_DN, GSE10500\_ARTHRITIC\_SYNOVIAL\_FLUID\_VS\_HEALTHY\_MACROPHAGE\_DN  
GCM\_AQP4, GCM\_AQP4  
GO\_PROTEIN\_K48\_LINKED\_DEUBIQUITINATION, GO\_PROTEIN\_K48\_LINKED\_DEUBIQUITINATION  
GSE17721\_PAM3CSK4\_VS\_GADIQUIMOD\_4H\_BMDC\_UP, GSE17721\_PAM3CSK4\_VS\_GADIQUIMOD\_4H\_BMDC\_UP  
GO\_GOLGI\_TO\_ENDOSOME\_TRANSPORT, GO\_GOLGI\_TO\_ENDOSOME\_TRANSPORT  
CCTGTGA\_MIR513, CCTGTGA\_MIR513  
GO\_PROTEIN\_DEGLYCOSYLATION, GO\_PROTEIN\_DEGLYCOSYLATION  
chr14q22, chr14q22  
GO\_REGULATION\_OF\_EARLY\_ENDOSOME\_TO\_LATE\_ENDOSOME\_TRANSPORT, GO\_REGULATION\_OF\_EARLY\_ENDOSOME\_TO\_LATE\_ENDOSOME\_TRANSPORT  
EIF4E\_DN, EIF4E\_DN  
GO\_ACTIN\_CYTOSKELETON\_REORGANIZATION, GO\_ACTIN\_CYTOSKELETON\_REORGANIZATION  
GSE26023\_PHD3\_KO\_VS\_WT\_NEUTROPHIL\_HYPOXIA\_DN, GSE26023\_PHD3\_KO\_VS\_WT\_NEUTROPHIL\_HYPOXIA\_DN  
GSE41978\_ID2\_KO\_VS\_BIM\_KO\_KLRG1\_LOW\_EFFECTOR\_CD8\_TCELL\_DN, GSE41978\_ID2\_KO\_VS\_BIM\_KO\_KLRG1\_LOW\_EFFECTOR\_CD8\_TCELL\_DN  
VERNELL\_RETINOBLASTOMA\_PATHWAY\_DN, VERNELL\_RETINOBLASTOMA\_PATHWAY\_DN  
GO\_ASYMMETRIC\_PROTEIN\_LOCALIZATION, GO\_ASYMMETRIC\_PROTEIN\_LOCALIZATION  
chr12q14, chr12q14  
GO\_CYTOPLASMIC\_UBIQUITIN\_LIGASE\_COMPLEX, GO\_CYTOPLASMIC\_UBIQUITIN\_LIGASE\_COMPLEX  
GTAAAG\_MIR302B, GTAAAG\_MIR302B  
GO\_PHOSPHATIDIC\_ACID\_METABOLIC\_PROCESS, GO\_PHOSPHATIDIC\_ACID\_METABOLIC\_PROCESS  
CAATGCA\_MIR33, CAATGCA\_MIR33  
HEDENFALK\_BREAST\_CANCER\_BRACX\_DN, HEDENFALK\_BREAST\_CANCER\_BRACX\_DN  
GO\_PROTEIN\_DEMANNOSYLATION, GO\_PROTEIN\_DEMANNOSYLATION