

VE\_TCELL\_VS\_NKCELL\_DN, GSE22886\_NAIVE\_TCELL\_VS\_NKCELL\_DN

GSE22886\_NAIVE\_CD8\_TCELL\_VS\_NKCELL\_DN, GSE22886\_NAIVE\_CD8\_TCELL\_VS\_NKCELL\_DN  
GSE5542\_UNTREATED\_VS\_IFNG\_TREATED\_EPITHELIAL\_CELLS\_6H\_DN, GSE5542\_UNTREATED\_VS\_IFNG\_TREATED\_EPITHELIAL\_CELLS\_6H\_DN  
GSE4984\_LPS\_VS\_VEHICLE\_CTRL\_TREATED\_DC\_DN, GSE4984\_LPS\_VS\_VEHICLE\_CTRL\_TREATED\_DC\_DN  
GSE5542\_UNTREATED\_VS\_IFNA\_TREATED\_EPITHELIAL\_CELLS\_24H\_UP, GSE5542\_UNTREATED\_VS\_IFNA\_TREATED\_EPITHELIAL\_CELLS\_24H\_UP  
KEGG\_ENDOCYTOSIS, KEGG\_ENDOCYTOSIS  
GSE1460\_DP\_THYMOCYTE\_VS\_THYMIC\_STROMAL\_CELL\_DN, GSE1460\_DP\_THYMOCYTE\_VS\_THYMIC\_STROMAL\_CELL\_DN  
GSE46606\_IRF4\_KO\_VS\_WT\_CD40L\_IL2\_IL5\_3DAY\_STIMULATED\_BCELL\_DN, GSE46606\_IRF4\_KO\_VS\_WT\_CD40L\_IL2\_IL5\_3DAY\_STIMULATED\_BCELL\_DN  
GSE1432\_6H\_VS\_24H\_IFNG\_MICROGLIA\_UP, GSE1432\_6H\_VS\_24H\_IFNG\_MICROGLIA\_UP  
GSE36392\_TYPE\_2\_MYELOID\_VS\_EOSINOPHIL\_IL25\_TREATED\_LUNG\_DN, GSE36392\_TYPE\_2\_MYELOID\_VS\_EOSINOPHIL\_IL25\_TREATED\_LUNG\_DN  
GSE36392\_EOSINOPHIL\_VS\_NEUTROPHIL\_IL25\_TREATED\_LUNG\_UP, GSE36392\_EOSINOPHIL\_VS\_NEUTROPHIL\_IL25\_TREATED\_LUNG\_UP  
GO\_TRANS\_GOLGI\_NETWORK, GO\_TRANS\_GOLGI\_NETWORK  
ACTGTAG\_MIR139, ACTGTAG\_MIR139  
KIM\_WT1\_TARGETS\_12HR\_UP, KIM\_WT1\_TARGETS\_12HR\_UP  
GSE4590\_LARGE\_PRE\_BCELL\_VS\_VPREB\_POS\_LARGE\_PRE\_BCELL\_UP, GSE4590\_LARGE\_PRE\_BCELL\_VS\_VPREB\_POS\_LARGE\_PRE\_BCELL\_UP  
GSE43955\_1H\_VS\_60H\_ACT\_CD4\_TCELL\_UP, GSE43955\_1H\_VS\_60H\_ACT\_CD4\_TCELL\_UP  
GSE17721\_LPS\_VS\_CPG\_8H\_BMDC\_UP, GSE17721\_LPS\_VS\_CPG\_8H\_BMDC\_UP  
ROSS\_LEUKEMIA\_WITH\_MLL\_FUSIONS, ROSS\_LEUKEMIA\_WITH\_MLL\_FUSIONS  
GRAHAM\_CML\_DIVIDING\_VS\_NORMAL QUIESCENT\_DN, GRAHAM\_CML\_DIVIDING\_VS\_NORMAL QUIESCENT\_DN  
WAMUNYOKOLI\_OVARIAN\_CANCER\_GRADES\_1\_2\_UP, WAMUNYOKOLI\_OVARIAN\_CANCER\_GRADES\_1\_2\_UP  
GO\_RAGE\_RECEPTOR\_BINDING, GO\_RAGE\_RECEPTOR\_BINDING  
GO\_REGULATION\_OF\_CELLULAR\_RESPONSE\_TO\_TRANSFORMING\_GROWTH\_FACTOR\_BETA\_STIMULUS, GO\_REGULATION\_OF\_CELLULAR\_RESPONSE\_TO\_TRANSFORMING\_GROWTH\_FACTOR\_BETA\_STIMULUS  
REACTOME\_SEMA4D\_IN\_SEMAPHORIN\_SIGNALING, REACTOME\_SEMA4D\_IN\_SEMAPHORIN\_SIGNALING  
GSE17721\_CTRL\_VS\_POLYIC\_0.5H\_BMDC\_DN, GSE17721\_CTRL\_VS\_POLYIC\_0.5H\_BMDC\_DN  
GO\_REGULATION\_OF\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSITION, GO\_REGULATION\_OF\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSITION  
GO\_DEFENSE\_RESPONSE\_TO\_FUNGUS, GO\_DEFENSE\_RESPONSE\_TO\_FUNGUS  
GO\_REGULATION\_OF\_EMBRYONIC\_DEVELOPMENT, GO\_REGULATION\_OF\_EMBRYONIC\_DEVELOPMENT  
AGCYRWTTT\_UNKNOWN, AGCYRWTTT\_UNKNOWN  
REACTOME\_ION\_TRANSPORT\_BY\_P\_TYPE\_ATPASES, REACTOME\_ION\_TRANSPORT\_BY\_P\_TYPE\_ATPASES  
GSE23321\_CENTRAL\_MEMORY\_VS\_NAIVE\_CD8\_TCELL\_DN, GSE23321\_CENTRAL\_MEMORY\_VS\_NAIVE\_CD8\_TCELL\_DN  
JI\_CARCINOGENESIS\_BY\_KRAS\_AND\_STK11\_DN, JI\_CARCINOGENESIS\_BY\_KRAS\_AND\_STK11\_DN  
GO\_REGULATION\_OF\_HISTONE\_DEACETYLATION, GO\_REGULATION\_OF\_HISTONE\_DEACETYLATION  
GO\_SPECIFIC GRANULE, GO\_SPECIFIC GRANULE  
GO\_LEUKOCYTE\_MIGRATION\_INVOLVED\_IN\_INFLAMMATORY\_RESPONSE, GO\_LEUKOCYTE\_MIGRATION\_INVOLVED\_IN\_INFLAMMATORY\_RESPONSE  
CHEOK\_RESPONSE\_TO\_MERCAPTOPURINE\_AND\_LD\_MTX\_DN, CHEOK\_RESPONSE\_TO\_MERCAPTOPURINE\_AND\_LD\_MTX\_DN  
chr17p12, chr17p12  
GO\_MALE\_SEX\_DIFFERENTIATION, GO\_MALE\_SEX\_DIFFERENTIATION  
GO\_POSITIVE\_REGULATION\_OF\_PROTEIN\_MATURATION, GO\_POSITIVE\_REGULATION\_OF\_PROTEIN\_MATURATION