

**4\_VS\_IL4\_TREATED\_ACT\_CD4\_TCELL\_6H\_UP, GSE2770\_TGFB\_AND\_IL4\_VS\_IL4\_TREATED\_ACT\_CD4\_TCELL\_6H\_UP**

GSE2770\_TGFB\_AND\_IL4\_VS\_TGFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_48H\_UP, GSE2770\_TGFB\_AND\_IL4\_VS\_TGFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_48H\_UP  
MODULE\_257, MODULE\_257  
GSE2770\_TGFB\_AND\_IL4\_ACT\_VS\_ACT\_CD4\_TCELL\_6H\_UP, GSE2770\_TGFB\_AND\_IL4\_ACT\_VS\_ACT\_CD4\_TCELL\_6H\_UP  
GSE45365\_HEALTHY\_VS\_MCMV\_INFECTION\_CD11B\_DC\_IFNAR\_KO\_UP, GSE45365\_HEALTHY\_VS\_MCMV\_INFECTION\_CD11B\_DC\_IFNAR\_KO\_UP  
GSE27786\_LIN\_NEG\_VS\_MONO\_MAC\_DN, GSE27786\_LIN\_NEG\_VS\_MONO\_MAC\_DN  
RB\_P107\_DN.V1\_DN, RB\_P107\_DN.V1\_DN  
GSE3982\_BASOPHIL\_VS\_CENT\_MEMORY\_CD4\_TCELL\_DN, GSE3982\_BASOPHIL\_VS\_CENT\_MEMORY\_CD4\_TCELL\_DN  
GSE2128\_CTRL\_VS\_MIMETOPE\_NEGATIVE\_SELECTION\_DP\_THYMOCYTE\_NOD\_UP, GSE2128\_CTRL\_VS\_MIMETOPE\_NEGATIVE\_SELECTION\_DP\_THYMOCYTE\_NOD\_UP  
GSE22886\_CD8\_VS\_CD4\_NAIVE\_TCELL\_DN, GSE22886\_CD8\_VS\_CD4\_NAIVE\_TCELL\_DN  
KEGG\_ACUTE\_MYELOID\_LEUKEMIA, KEGG\_ACUTE\_MYELOID\_LEUKEMIA  
GO\_TRANSCRIPTIONAL\_REPRESSOR\_ACTIVITY\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_FACTOR\_BINDING, GO\_TRANSCRIPTIONAL\_REPRESSOR\_ACTIVITY\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_FACTOR\_BINDING  
GSE17301\_IFNA2\_VS\_IFNA2\_AND\_ACD3\_ACD28\_STIM\_CD8\_TCELL\_DN, GSE17301\_IFNA2\_VS\_IFNA2\_AND\_ACD3\_ACD28\_STIM\_CD8\_TCELL\_DN  
ESC\_J1\_UP\_EARLY.V1\_UP, ESC\_J1\_UP\_EARLY.V1\_UP  
GSE24574\_BCL6\_LOW\_TFH\_VS\_TCONV\_CD4\_TCELL\_UP, GSE24574\_BCL6\_LOW\_TFH\_VS\_TCONV\_CD4\_TCELL\_UP  
GO\_NEGATIVE\_REGULATION\_OF\_AUTOPHAGY, GO\_NEGATIVE\_REGULATION\_OF\_AUTOPHAGY  
GO\_PYRIMIDINE\_NUCLEOBASE\_METABOLIC\_PROCESS, GO\_PYRIMIDINE\_NUCLEOBASE\_METABOLIC\_PROCESS  
GSE16385\_IFNG\_TNF\_VS\_IL4\_STIM\_MACROPHAGE\_ROSIGLITAZONE\_TREATED\_UP, GSE16385\_IFNG\_TNF\_VS\_IL4\_STIM\_MACROPHAGE\_ROSIGLITAZONE\_TREATED\_UP  
GO\_PROTEIN\_SECRETION, GO\_PROTEIN\_SECRETION  
BMI1\_DN\_MEL18\_DN.V1\_DN, BMI1\_DN\_MEL18\_DN.V1\_DN  
GO\_PHOSPHOTRANSFERASE\_ACTIVITY\_PHOSPHATE\_GROUP\_AS\_ACCEPTOR, GO\_PHOSPHOTRANSFERASE\_ACTIVITY\_PHOSPHATE\_GROUP\_AS\_ACCEPTOR  
GO\_DIPEPTIDASE\_ACTIVITY, GO\_DIPEPTIDASE\_ACTIVITY  
MODULE\_199, MODULE\_199  
GO\_PROTEIN\_O\_LINKED\_GLYCOSYLATION, GO\_PROTEIN\_O\_LINKED\_GLYCOSYLATION  
GO\_REGULATION\_OF\_TRANSCRIPTION\_FROM\_RNA\_POLYMERASE\_III\_PROMOTER, GO\_REGULATION\_OF\_TRANSCRIPTION\_FROM\_RNA\_POLYMERASE\_III\_PROMOTER  
DACOSTA\_ERCC3\_ALLELE\_XPCS\_VS\_TTD\_UP, DACOSTA\_ERCC3\_ALLELE\_XPCS\_VS\_TTD\_UP  
SASAKI\_TARGETS\_OF\_TP73\_AND\_TP63, SASAKI\_TARGETS\_OF\_TP73\_AND\_TP63  
GO\_BICELLULAR\_TIGHT\_JUNCTION\_ASSEMBLY, GO\_BICELLULAR\_TIGHT\_JUNCTION\_ASSEMBLY