

E\_VS\_LN\_DERIVED\_PPARG\_KO\_TCONV\_CD4\_TCELL\_UP, GSE37532\_VISCERAL\_ADIPOSE\_TISSUE\_VS\_LN\_DERIVED\_PPARG\_KO\_TCONV\_CD4\_TCELL\_UP

GSE22886\_NAIVE\_CD4\_TCELL\_VS\_NEUTROPHIL\_UP, GSE22886\_NAIVE\_CD4\_TCELL\_VS\_NEUTROPHIL\_UP  
GSE37532\_VISCERAL\_ADIPOSE\_TISSUE\_VS\_LN\_DERIVED\_PPARG\_KO\_TREG\_CD4\_TCELL\_UP, GSE37532\_VISCERAL\_ADIPOSE\_TISSUE\_VS\_LN\_DERIVED\_PPARG\_KO\_TREG\_CD4\_TCELL\_UP  
GSE12366\_GC\_VS\_MEMORY\_BCELL\_UP, GSE12366\_GC\_VS\_MEMORY\_BCELL\_UP  
GO\_HELICASE\_ACTIVITY, GO\_HELICASE\_ACTIVITY  
GSE3982\_MAST\_CELL\_VS\_MAC\_DN, GSE3982\_MAST\_CELL\_VS\_MAC\_DN  
E2F\_Q2, E2F\_Q2  
GO\_PURINE\_NTP\_DEPENDENT\_HELICASE\_ACTIVITY, GO\_PURINE\_NTP\_DEPENDENT\_HELICASE\_ACTIVITY  
GSE26030\_UNSTIM\_VS\_RESTIM\_TH17\_DAY15\_POST\_POLARIZATION\_UP, GSE26030\_UNSTIM\_VS\_RESTIM\_TH17\_DAY15\_POST\_POLARIZATION\_UP  
GO\_AMINO\_ACID\_ACTIVATION, GO\_AMINO\_ACID\_ACTIVATION  
GO\_LIGASE\_ACTIVITY\_FORMING\_CARBON\_OXYGEN\_BONDS, GO\_LIGASE\_ACTIVITY\_FORMING\_CARBON\_OXYGEN\_BONDS  
KEGG\_PROGESTERONE\_MEDIATED\_OOCYTE\_MATURATION, KEGG\_PROGESTERONE\_MEDIATED\_OOCYTE\_MATURATION  
GO\_CYTOPLASMIC\_MRNA\_PROCESSING\_BODY, GO\_CYTOPLASMIC\_MRNA\_PROCESSING\_BODY  
KOKKINAKIS\_METHIONINE\_DEPRIVATION\_48HR\_DN, KOKKINAKIS\_METHIONINE\_DEPRIVATION\_48HR\_DN  
REACTOME\_NUCLEOTIDE\_BINDING\_DOMAIN\_LEUCINE\_RICH\_REPEAT\_CONTAINING\_RECEPTOR\_NLR\_SIGNALING\_PATHWAYS, REACTOME\_NUCLEOTIDE\_BINDING\_DOMAIN\_LEUCINE\_RICH\_REPEAT\_CONTA  
JOHNSTONE\_PARVB\_TARGETS\_1\_DN, JOHNSTONE\_PARVB\_TARGETS\_1\_DN  
GSE29164\_UNTREATED\_VS\_CD8\_TCELL\_AND\_IL12\_TREATED\_MELANOMA\_DAY7\_DN, GSE29164\_UNTREATED\_VS\_CD8\_TCELL\_AND\_IL12\_TREATED\_MELANOMA\_DAY7\_DN  
GO\_NEGATIVE\_REGULATION\_OF\_MITOTIC\_NUCLEAR\_DIVISION, GO\_NEGATIVE\_REGULATION\_OF\_MITOTIC\_NUCLEAR\_DIVISION  
GO\_REGULATION\_OF\_CELL\_PROJECTION\_ASSEMBLY, GO\_REGULATION\_OF\_CELL\_PROJECTION\_ASSEMBLY  
GO\_ATP\_DEPENDENT\_DNA\_HELICASE\_ACTIVITY, GO\_ATP\_DEPENDENT\_DNA\_HELICASE\_ACTIVITY  
REACTOME\_OPIOID\_SIGNALLING, REACTOME\_OPIOID\_SIGNALLING  
GO\_CELL\_CYCLE\_DNA\_REPLICATION, GO\_CELL\_CYCLE\_DNA\_REPLICATION  
GO\_POSITIVE\_REGULATION\_OF\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSITION, GO\_POSITIVE\_REGULATION\_OF\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSITION  
REACTOME\_GABA\_B\_RECEPTOR\_ACTIVATION, REACTOME\_GABA\_B\_RECEPTOR\_ACTIVATION  
GO\_RESPONSE\_TO\_CADMIUM\_ION, GO\_RESPONSE\_TO\_CADMIUM\_ION  
GSE2770\_IL12\_VS\_TGFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_6H\_UP, GSE2770\_IL12\_VS\_TGFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_6H\_UP  
GO\_CHLORIDE\_CHANNEL\_REGULATOR\_ACTIVITY, GO\_CHLORIDE\_CHANNEL\_REGULATOR\_ACTIVITY  
BIOCARTA\_RANKL\_PATHWAY, BIOCARTA\_RANKL\_PATHWAY