

VS\_IFNA\_STIM\_CD8\_TCELL\_90MIN\_DN, GSE40666\_UNTREATED\_VS\_IFNA\_STIM\_CD8\_TCELL\_90MIN\_DN

GSE40666\_STAT1\_KO\_VS\_STAT4\_KO\_CD8\_TCELL\_UP, GSE40666\_STAT1\_KO\_VS\_STAT4\_KO\_CD8\_TCELL\_UP  
GSE37416\_0H\_VS\_24H\_F\_TULARENSIS\_LVS\_NEUTROPHIL\_UP, GSE37416\_0H\_VS\_24H\_F\_TULARENSIS\_LVS\_NEUTROPHIL\_UP  
GSE40666\_WT\_VS\_STAT4\_KO\_CD8\_TCELL\_DN, GSE40666\_WT\_VS\_STAT4\_KO\_CD8\_TCELL\_DN  
GSE3039\_NKT\_CELL\_VS\_B1\_BCELL\_DN, GSE3039\_NKT\_CELL\_VS\_B1\_BCELL\_DN  
GSE40666\_STAT1\_KO\_VS\_STAT4\_KO\_CD8\_TCELL\_WITH\_IFNA\_STIM\_90MIN\_UP, GSE40666\_STAT1\_KO\_VS\_STAT4\_KO\_CD8\_TCELL\_WITH\_IFNA\_STIM\_90MIN\_UP  
GSE2770\_TGFB\_AND\_IL4\_VS\_TGFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_48H\_DN, GSE2770\_TGFB\_AND\_IL4\_VS\_TGFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_48H\_DN  
GSE20715\_WT\_VS\_TLR4\_KO\_48H\_OZONE\_LUNG\_DN, GSE20715\_WT\_VS\_TLR4\_KO\_48H\_OZONE\_LUNG\_DN  
GSE3982\_BCELL\_VS\_BASOPHIL\_UP, GSE3982\_BCELL\_VS\_BASOPHIL\_UP  
GSE1925\_CTRL\_VS\_24H\_IFNG\_STIM\_MACROPHAGE\_DN, GSE1925\_CTRL\_VS\_24H\_IFNG\_STIM\_MACROPHAGE\_DN  
GSE37532\_VISCERAL\_ADIPOSE\_TISSUE\_VS\_LN\_DERIVED\_TREG\_CD4\_TCELL\_DN, GSE37532\_VISCERAL\_ADIPOSE\_TISSUE\_VS\_LN\_DERIVED\_TREG\_CD4\_TCELL\_DN  
HEIDENBLAD\_AMPLICON\_8Q24\_UP, HEIDENBLAD\_AMPLICON\_8Q24\_UP  
PID\_AVB3\_INTEGRIN\_PATHWAY, PID\_AVB3\_INTEGRIN\_PATHWAY  
GO\_TISSUE\_MIGRATION, GO\_TISSUE\_MIGRATION  
GO\_G\_PROTEIN\_COUPLED\_PURINERGIC\_NUCLEOTIDE\_RECEPTOR\_SIGNALING\_PATHWAY, GO\_G\_PROTEIN\_COUPLED\_PURINERGIC\_NUCLEOTIDE\_RECEPTOR\_SIGNALING\_PATHWAY  
GO\_POSITIVE\_REGULATION\_OF\_BLOOD\_CIRCULATION, GO\_POSITIVE\_REGULATION\_OF\_BLOOD\_CIRCULATION  
GO\_INOSITOL\_LIPID\_MEDIATED\_SIGNALING, GO\_INOSITOL\_LIPID\_MEDIATED\_SIGNALING  
GO\_REGULATION\_OF\_ACTIN\_FILAMENT\_BUNDLE\_ASSEMBLY, GO\_REGULATION\_OF\_ACTIN\_FILAMENT\_BUNDLE\_ASSEMBLY  
GO\_REGULATION\_OF\_CHROMATIN\_BINDING, GO\_REGULATION\_OF\_CHROMATIN\_BINDING  
GO\_POSITIVE\_REGULATION\_OF\_MITOTIC\_SISTER\_CHROMATID\_SEPARATION, GO\_POSITIVE\_REGULATION\_OF\_MITOTIC\_SISTER\_CHROMATID\_SEPARATION  
GO\_POSITIVE\_REGULATION\_OF\_ACTIN\_FILAMENT\_BUNDLE\_ASSEMBLY, GO\_POSITIVE\_REGULATION\_OF\_ACTIN\_FILAMENT\_BUNDLE\_ASSEMBLY  
GO\_HISTONE\_H3\_K9\_MODIFICATION, GO\_HISTONE\_H3\_K9\_MODIFICATION  
REACTOME\_GLUCURONIDATION, REACTOME\_GLUCURONIDATION  
GO\_POSITIVE\_REGULATION\_OF\_VASOCONSTRICTION, GO\_POSITIVE\_REGULATION\_OF\_VASOCONSTRICTION  
PID\_ECADHERIN\_KERATINOCYTE\_PATHWAY, PID\_ECADHERIN\_KERATINOCYTE\_PATHWAY  
REACTOME\_GENERATION\_OF\_SECOND\_MESSENGER\_MOLECULES, REACTOME\_GENERATION\_OF\_SECOND\_MESSENGER\_MOLECULES  
REACTOME\_NEF\_MEDIATES\_DOWN\_MODULATION\_OF\_CELL\_SURFACE\_RECEPTORS\_BY\_RECRUITING\_THEM\_TO\_CLATHRIN\_ADAPTERS, REACTOME\_NEF\_MEDIATES\_DOWN\_MODULATION\_OF\_CELL\_SURFACE\_RECEPTORS\_BY\_RECRUITING\_THEM\_TO\_CLATHRIN\_ADAPTERS