VS_IL3_IL33_TREATED_MAST_CELL_UP, GSE39382_IL3_VS_IL3_IL33_TREATED_MAST_CELL_UP

```
GSE11864 CSF1 VS CSF1 PAM3CYS IN MAC UP, GSE11864 CSF1 VS CSF1 PAM3CYS IN MAC UP
GSE411_UNSTIM_VS_400MIN_IL6_STIM_MACROPHAGE_UP, GSE411_UNSTIM_VS_400MIN_IL6_STIM_MACROPHAGE_UP
GSE5589 LPS VS LPS AND IL6 STIM IL6 KO MACROPHAGE 45MIN DN, GSE5589 LPS VS LPS AND IL6 STIM IL6 KO MACROPHAGE 45MIN DN
GSE36826 NORMAL VS STAPH AUREUS INF IL1R KO SKIN UP, GSE36826 NORMAL VS STAPH AUREUS INF IL1R KO SKIN UP
GSE36888 UNTREATED_VS_IL2_TREATED_TCELL_17H_UP, GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_17H_UP
GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG1_AND_FOXP3_TRASDUCED_DN, GSE37533_UNTREATED_TCELL_PPARG1_AND_FOXP3_TCELL_PPARG1_AND_FOXP3_TCELL_PPARG1_AND_FOXP3_TCELL_PPARG1_AND_FOXP3_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_PPARG1_TCELL_TCELL_PPARG1_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCELL_TCE
GSE19198_6H_VS_24H_IL21_TREATED_TCELL_DN, GSE19198_6H_VS_24H_IL21_TREATED_TCELL_DN
GSE11864_CSF1_VS_CSF1_IFNG_IN_MAC_UP, GSE11864_CSF1_VS_CSF1_IFNG_IN_MAC_UP
GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_UP, GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_UP
GSE35435_RESTING_VS_IL4_TREATED_MACROPHAGE_DN, GSE35435_RESTING_VS_IL4_TREATED_MACROPHAGE_DN
GO FC GAMMA RECEPTOR SIGNALING PATHWAY, GO FC GAMMA RECEPTOR SIGNALING PATHWAY
KEGG_INOSITOL_PHOSPHATE_METABOLISM, KEGG_INOSITOL_PHOSPHATE_METABOLISM
GO POSITIVE REGULATION OF LIPID BIOSYNTHETIC PROCESS, GO POSITIVE REGULATION OF LIPID BIOSYNTHETIC PROCESS
LE_EGR2_TARGETS_DN, LE_EGR2_TARGETS_DN
LEIN_ASTROCYTE_MARKERS, LEIN_ASTROCYTE_MARKERS
 YAO_HOXA10_TARGETS_VIA_PROGESTERONE_UP, YAO_HOXA10_TARGETS_VIA_PROGESTERONE_UP
chr9q33, chr9q33
REACTOME IL RECEPTOR SHC SIGNALING, REACTOME IL RECEPTOR SHC SIGNALING
GO MONOSACCHARIDE TRANSMEMBRANE TRANSPORTER ACTIVITY, GO MONOSACCHARIDE TRANSMEMBRANE TRANSPORTER ACTIVITY
GO CARBON SULFUR LYASE ACTIVITY, GO CARBON SULFUR LYASE ACTIVITY
GO_REGULATION_OF_WNT_SIGNALING_PATHWAY_PLANAR_CELL_POLARITY_PATHWAY, GO_REGULATION_OF_WNT_SIGNALING_PATHWAY_PLANAR_CELL_POLARITY_PATHWAY
GEISS_RESPONSE_TO_DSRNA_DN, GEISS_RESPONSE_TO_DSRNA_DN
HAHTOLA MYCOSIS FUNGOIDES DN. HAHTOLA MYCOSIS FUNGOIDES DN
```