

VS_IKAROS_KO_HSC_DN, GSE15330_WT_VS_IKAROS_KO_HSC_DN

GSE26290_WT_VS_PDK1_KO_ANTI_CD3_AND_IL2_STIM_CD8_TCELL_DN, GSE26290_WT_VS_PDK1_KO_ANTI_CD3_AND_IL2_STIM_CD8_TCELL_DN

GSE15330_MEGAKARYOCYTE_ERYTHROID_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_DN, GSE15330_MEGAKARYOCYTE_ERYTHROID_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_DN

GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_DN, GSE15330_LYMPHOID_MULTIPOTENT_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_DN

GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_AND_ROSIGLITAZONE_STIM_UP, GSE25123_WT_VS_PPARG_KO_MACROPHAGE_IL4_AND_ROSIGLITAZONE_STIM_UP

GSE7460_CTRL_VS_TGFB_TREATED_ACT_TREG_DN, GSE7460_CTRL_VS_TGFB_TREATED_ACT_TREG_DN

GO_REGULATION_OF_EXTENT_OF_CELL_GROWTH, GO_REGULATION_OF_EXTENT_OF_CELL_GROWTH

GSE21380_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_UP, GSE21380_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_UP

RYCACNNRNNRNCAG_UNKNOWN, RYCACNNRNNRNCAG_UNKNOWN

PID_LIS1_PATHWAY, PID_LIS1_PATHWAY

GO_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY, GO_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY

GATA_Q6, GATA_Q6

GO_CYTOPLASMIC_MRNA_PROCESSING_BODY_ASSEMBLY, GO_CYTOPLASMIC_MRNA_PROCESSING_BODY_ASSEMBLY

GO_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION, GO_REGULATION_OF_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION

GO_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY, GO_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY