

1_KO_8H_ANTI_IGM_STIM_BCELL_DN, GSE21063_WT_VS_NFATC1_KO_8H_ANTI_IGM_STIM_BCELL_DN

GSE15330_GRANULOCYTE_MONOCYTE_PROGENITOR_VS_PRO_BCELL_DN, GSE15330_GRANULOCYTE_MONOCYTE_PROGENITOR_VS_PRO_BCELL_DN
GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_DN, GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_DN
GSE13522_WT_VS_IFNAR_KO_SKING_T_CRUZI_Y_STRAIN_INF_UP, GSE13522_WT_VS_IFNAR_KO_SKING_T_CRUZI_Y_STRAIN_INF_UP
GSE17721_LPS_VS_CPG_4H_BMDC_UP, GSE17721_LPS_VS_CPG_4H_BMDC_UP
GSE7460_FOXP3_MUT_VS_WT_ACT_TCONV_DN, GSE7460_FOXP3_MUT_VS_WT_ACT_TCONV_DN
GSE20366_CD103_KLRG1_DP_VS_DN_TREG_UP, GSE20366_CD103_KLRG1_DP_VS_DN_TREG_UP
GSE17721_0.5H_VS_4H_LPS_BMDC_DN, GSE17721_0.5H_VS_4H_LPS_BMDC_DN
MIR5589_3P, MIR5589_3P
THRA_TARGET_GENES, THRA_TARGET_GENES
MIR3194_5P, MIR3194_5P
GAO_ESOPHAGUS_25W_C3_FGFR1LOW_EPITHELIAL_CELLS, GAO_ESOPHAGUS_25W_C3_FGFR1LOW_EPITHELIAL_CELLS
MIR3613_5P, MIR3613_5P
HP_ABNORMALITY_OF_THE_SCROTUM, HP_ABNORMALITY_OF_THE_SCROTUM
GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_DN, GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_DN
MIR4289, MIR4289
HP_INCOMITANT_STRABISMUS, HP_INCOMITANT_STRABISMUS
HP_APLASIA_HYPOPLASIA_OF_THE_HALLUX, HP_APLASIA_HYPOPLASIA_OF_THE_HALLUX
GOCC_TRANSCRIPTIONALLY_ACTIVE_CHROMATIN, GOCC_TRANSCRIPTIONALLY_ACTIVE_CHROMATIN
HP_ABNORMAL_MUSCLE_GLYCOGEN_CONTENT, HP_ABNORMAL_MUSCLE_GLYCOGEN_CONTENT
GOBP_REGULATION_OF_KETONE_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_KETONE_BIOSYNTHETIC_PROCESS
REACTOME_GLYCOGEN_BREAKDOWN_GLYCOGENOLYSIS, REACTOME_GLYCOGEN_BREAKDOWN_GLYCOGENOLYSIS
GOCC_TROPONIN_COMPLEX, GOCC_TROPONIN_COMPLEX
HP_ROUND_EAR, HP_ROUND_EAR