

1_KO_8H_ANTI_IGM_STIM_BCELL_DN, GSE21063_WT_VS_NFATC1_KO_8H_ANTI_IGM_STIM_BCELL_DN

GSE8835_HEALTHY_VS_CLL_CD8_TCELL_UP, GSE8835_HEALTHY_VS_CLL_CD8_TCELL_UP
GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP, GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP
GSE43863_TH1_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN, GSE43863_TH1_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN
GSE23321_CENTRAL_VS_EFFECTOR_MEMORY_CD8_TCELL_UP, GSE23321_CENTRAL_VS_EFFECTOR_MEMORY_CD8_TCELL_UP
GSE17580_TREG_VS_TEFF_UP, GSE17580_TREG_VS_TEFF_UP
GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_DN, GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_DN
GSE7460_FOXP3_MUT_VS_WT_ACT_TCONV_DN, GSE7460_FOXP3_MUT_VS_WT_ACT_TCONV_DN
GSE21033_1H_VS_24H_POLYIC_STIM_DC_UP, GSE21033_1H_VS_24H_POLYIC_STIM_DC_UP
ACCATTT_MIR522, ACCATTT_MIR522
GSE17721_0.5H_VS_4H_LPS_BMDC_DN, GSE17721_0.5H_VS_4H_LPS_BMDC_DN
MIR423_5P, MIR423_5P
MIR3184_5P, MIR3184_5P
GSE20366_CD103_KLRG1_DP_VS_DN_TREG_UP, GSE20366_CD103_KLRG1_DP_VS_DN_TREG_UP
GSE18203_CTRL_VS_INTRATUMORAL_CPG_INJ_MC38_TUMOR_UP, GSE18203_CTRL_VS_INTRATUMORAL_CPG_INJ_MC38_TUMOR_UP
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_DN, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_DN
GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_DN, GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_DN
GSE12198_CTRL_VS_HIGH_IL2_STIM_NK_CELL_UP, GSE12198_CTRL_VS_HIGH_IL2_STIM_NK_CELL_UP
GGGCATT_MIR365, GGCATT_MIR365
GSE17721_PAM3CSK4_VS_GADIQUIMOD_1H_BMDC_UP, GSE17721_PAM3CSK4_VS_GADIQUIMOD_1H_BMDC_UP
THRA_TARGET_GENES, THRA_TARGET_GENES
HP_DISPROPORTIONATE_TALL_STATURE, HP_DISPROPORTIONATE_TALL_STATURE
GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG2_AND_FOXP3_TRASDUCED_UP, GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG2_A
HP_DEVIATION_OF_THE_HALLUX, HP_DEVIATION_OF_THE_HALLUX
HP_ABNORMAL_FINGERTIP_MORPHOLOGY, HP_ABNORMAL_FINGERTIP_MORPHOLOGY
GSE22886_TH1_VS_TH2_48H_ACT_UP, GSE22886_TH1_VS_TH2_48H_ACT_UP
GTTATAT_MIR410, GTTATAT_MIR410
AR_01, AR_01
HMEF2_Q6, HMEF2_Q6
GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_DN, GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_DN
GOBP_T_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_T_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE
GOBP_POSITIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
MCBRYAN_PUBERTAL_TGFB1_TARGETS_DN, MCBRYAN_PUBERTAL_TGFB1_TARGETS_DN
HP_ABNORMAL_HAND_BONE_OSSIFICATION, HP_ABNORMAL_HAND_BONE_OSSIFICATION
CLASPER_LYMPHATIC_VESSELS_DURING_METASTASIS_UP, CLASPER_LYMPHATIC_VESSELS_DURING_METASTASIS_UP
VALK_AML_CLUSTER_10, VALK_AML_CLUSTER_10
GOBP_NEGATIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GOBP_NEGATIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
GOBP_POSITIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNA
MIR6865_3P, MIR6865_3P
GOBP_C21_STEROID_HORMONE_BIOSYNTHETIC_PROCESS, GOBP_C21_STEROID_HORMONE_BIOSYNTHETIC_PROCESS
GSE41867_DAY8_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN, GSE41867_DAY8_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN
MODULE_210, MODULE_210
GOMF_ORGANIC_CATION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_ORGANIC_CATION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
BEGUM_TARGETS_OF_PAX3_FOXO1_FUSION_AND_PAX3, BEGUM_TARGETS_OF_PAX3_FOXO1_FUSION_AND_PAX3
HP_ROUND_EAR, HP_ROUND_EAR
GOBP_MUCOSA_ASSOCIATED_LYMPHOID_TISSUE_DEVELOPMENT, GOBP_MUCOSA_ASSOCIATED_LYMPHOID_TISSUE_DEVELOPMENT
GOBP_PROGESTERONE_BIOSYNTHETIC_PROCESS, GOBP_PROGESTERONE_BIOSYNTHETIC_PROCESS
GOBP_OLEFINIC_COMPOUND_BIOSYNTHETIC_PROCESS, GOBP_OLEFINIC_COMPOUND_BIOSYNTHETIC_PROCESS
ALONSO_METASTASIS_DN, ALONSO_METASTASIS_DN
GOBP_GASTRULATION_WITH_MOUTH_FORMING_SECOND, GOBP_GASTRULATION_WITH_MOUTH_FORMING_SECOND