

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
GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_2H_DN
GSE7852_LN_VS_THYMUS_TREG_UP, GSE7852_LN_VS_THYMUS_TREG_UP
GSE17721_LPS_VS_CPG_4H_BMDC_UP, GSE17721_LPS_VS_CPG_4H_BMDC_UP
GSE21033_1H_VS_24H_POLYIC_STIM_DC_UP, GSE21033_1H_VS_24H_POLYIC_STIM_DC_UP
MIR3688_5P, MIR3688_5P
GSE360_L_MAJOR_VS_M_TUBERCULOSIS_DC_UP, GSE360_L_MAJOR_VS_M_TUBERCULOSIS_DC_UP
HP_PROMINENT_FOREHEAD, HP_PROMINENT_FOREHEAD
THRA_TARGET_GENES, THRA_TARGET_GENES
YGCANTGCR_UNKNOWN, YGCANTGCR_UNKNOWN
HEN1_02, HEN1_02
MIR6809_5P, MIR6809_5P
GSE20366_CD103_KLRG1_DP_VS_DN_TREG_UP, GSE20366_CD103_KLRG1_DP_VS_DN_TREG_UP
MIR324_3P, MIR324_3P
ZNF777_TARGET_GENES, ZNF777_TARGET_GENES
PR_02, PR_02
MIR938, MIR938
GAO_ESOPHAGUS_25W_C3_FGFR1LOW_EPITHELIAL_CELLS, GAO_ESOPHAGUS_25W_C3_FGFR1LOW_EPITHELIAL_CELLS
GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG2_AND_FOXP3_TRASDUCED_UP, GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_PPARG2_A
GSE22025_UNTREATED_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_DN, GSE22025_UNTREATED_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_DN
GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_DN, GSE46242_TH1_VS_ANERGIC_TH1_CD4_TCELL_DN
MODULE_366, MODULE_366
MIR6892_5P, MIR6892_5P
PR_01, PR_01
KIM_MYCN_AMPLIFICATION_TARGETS_UP, KIM_MYCN_AMPLIFICATION_TARGETS_UP
MIR4289, MIR4289
MIR5684, MIR5684
FOXD3_TARGET_GENES, FOXD3_TARGET_GENES
GOBP_POSITIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_MEDIATED_SIGNA
CLASPER_LYMPHATIC_VESSELS_DURING_METASTASIS_UP, CLASPER_LYMPHATIC_VESSELS_DURING_METASTASIS_UP
VALK_AML_CLUSTER_10, VALK_AML_CLUSTER_10
DESCARTES_FETAL_EYE_AMACRINE_CELLS, DESCARTES_FETAL_EYE_AMACRINE_CELLS