

MIN_IL6_STIM_MACROPHAGE_DN, GSE411_UNSTIM_VS_400MIN_IL6_STIM_MACROPHAGE_DN

GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_48H_DN, GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_48H_DN
GSE2770_UNTREATED_VS_ACT_CD4_TCELL_2H_DN, GSE2770_UNTREATED_VS_ACT_CD4_TCELL_2H_DN
GSE411_WT_VS_SOCS3_KO_MACROPHAGE_UP, GSE411_WT_VS_SOCS3_KO_MACROPHAGE_UP
GSE3039_NKT_CELL_VS_ALPHABETA_CD8_TCELL_UP, GSE3039_NKT_CELL_VS_ALPHABETA_CD8_TCELL_UP
GSE411_UNSTIM_VS_400MIN_IL6_STIM_SOCS3_KO_MACROPHAGE_DN, GSE411_UNSTIM_VS_400MIN_IL6_STIM_SOCS3_KO_MACROPHAGE_DN
GSE2770_IL12_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_DN, GSE2770_IL12_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_DN
GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_MONOCYTE_UP, GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_MONOCYTE_UP
WP_GENES_RELATED_TO_PRIMARY_CILIUM_DEVELOPMENT_BASED_ON_CRISPR, WP_GENES_RELATED_TO_PRIMARY_CILIUM_DEVELOPMENT_BASED_ON_CRISPR
GSE36527_CD62L_HIGH_VS_CD62L_LOW_TREG_CD69_NEG_KLRG1_NEG_DN, GSE36527_CD62L_HIGH_VS_CD62L_LOW_TREG_CD69_NEG_KLRG1_NEG_DN
CTCTGGA_MIR520A_MIR525, CTCTGGA_MIR520A_MIR525
MIR7849_3P, MIR7849_3P
MIR26B_3P, MIR26B_3P
GOBP_PROTEIN_TRANSPORT_ALONG_MICROTUBULE, GOBP_PROTEIN_TRANSPORT_ALONG_MICROTUBULE
GTTNYYNNGGTNA_UNKNOWN, GTTNYYNNGGTNA_UNKNOWN
GSE13887_RESTING_VS_NO_TREATED_CD4_TCELL_DN, GSE13887_RESTING_VS_NO_TREATED_CD4_TCELL_DN
HP_CONE_SHAPED_EPIPYSIS, HP_CONE_SHAPED_EPIPYSIS
MIR218_2_3P, MIR218_2_3P
MIR3146, MIR3146
HP_ABNORMAL_CIRCULATING_FOLLICLE_STIMULATING_HORMONE_LEVEL, HP_ABNORMAL_CIRCULATING_FOLLICLE_STIMULATING_HORMONE_LEVEL
LASTOWSKA_COAMPLIFIED_WITH_MYCN, LASTOWSKA_COAMPLIFIED_WITH_MYCN
GOMF_LIGAND_ACTIVATED_TRANSCRIPTION_FACTOR_ACTIVITY, GOMF_LIGAND_ACTIVATED_TRANSCRIPTION_FACTOR_ACTIVITY
GOBP_NUCLEOSIDE_MONOPHOSPHATE_PHOSPHORYLATION, GOBP_NUCLEOSIDE_MONOPHOSPHATE_PHOSPHORYLATION
MIKKELSEN_IPS_ICP_WITH_H3K27ME3, MIKKELSEN_IPS_ICP_WITH_H3K27ME3
HP_SHOULDER_FLEXION_CONTRACTURE, HP_SHOULDER_FLEXION_CONTRACTURE