

_VS_IL12_TREATED_ACT_CD4_TCELL_UP, GSE20198_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_UP

GSE20198_IL12_VS_IFNA_TREATED_ACT_CD4_TCELL_UP, GSE20198_IL12_VS_IFNA_TREATED_ACT_CD4_TCELL_UP
GSE25890_CTRL_VS_IL33_IL7_TREATED_NUOCYTES_DN, GSE25890_CTRL_VS_IL33_IL7_TREATED_NUOCYTES_DN
GSE19888_CTRL_VS_T_CELL_MEMBRANES_ACT_MAST_CELL_UP, GSE19888_CTRL_VS_T_CELL_MEMBRANES_ACT_MAST_CELL_UP
GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_DN, GSE20198_UNTREATED_VS_IFNA_TREATED_ACT_CD4_TCELL_DN
GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_6H_DN, GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_6H_DN
GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_DN, GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_DN
GSE27786_BCELL_VS_CD4_TCELL_UP, GSE27786_BCELL_VS_CD4_TCELL_UP
GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_UP
GSE14308_INDUCED_VS_NATURAL_TREG_UP, GSE14308_INDUCED_VS_NATURAL_TREG_UP
GSE11961_MEMORY_BCELL_DAY7_VS_MEMORY_BCELL_DAY40_DN, GSE11961_MEMORY_BCELL_DAY7_VS_MEMORY_BCELL_DAY40_DN
GOBP_REGULATION_OF_CELL_PROJECTION_ASSEMBLY, GOBP_REGULATION_OF_CELL_PROJECTION_ASSEMBLY
MIR3168, MIR3168
MIR5691_MIR6805_3P, MIR5691_MIR6805_3P
MIR939_5P, MIR939_5P
GOBP_REGULATION_OF_CILIUM_ASSEMBLY, GOBP_REGULATION_OF_CILIUM_ASSEMBLY
HEY2_TARGET_GENES, HEY2_TARGET_GENES
WP_WNTBETACATENIN_SIGNALING_PATHWAY_IN_LEUKEMIA, WP_WNTBETACATENIN_SIGNALING_PATHWAY_IN_LEUKEMIA
REACTOME_ARACHIDONIC_ACID_METABOLISM, REACTOME_ARACHIDONIC_ACID_METABOLISM
GOBP_REGULATION_OF_TUBE_SIZE, GOBP_REGULATION_OF_TUBE_SIZE
GOMF_INTERLEUKIN_1_RECEPTOR_BINDING, GOMF_INTERLEUKIN_1_RECEPTOR_BINDING
GOBP_ENDOCARDIAL_CUSHION_DEVELOPMENT, GOBP_ENDOCARDIAL_CUSHION_DEVELOPMENT
VERNELL_RETINOBLASTOMA_PATHWAY_DN, VERNELL_RETINOBLASTOMA_PATHWAY_DN
GOMF_DYNEIN_LIGHT_INTERMEDIATE_CHAIN_BINDING, GOMF_DYNEIN_LIGHT_INTERMEDIATE_CHAIN_BINDING
VANDESLUIS_COMMD1_TARGETS_GROUP_4_UP, VANDESLUIS_COMMD1_TARGETS_GROUP_4_UP
KEGG_ARACHIDONIC_ACID_METABOLISM, KEGG_ARACHIDONIC_ACID_METABOLISM
GOBP_AMPA_GLUTAMATE_RECEPTOR_CLUSTERING, GOBP_AMPA_GLUTAMATE_RECEPTOR_CLUSTERING
WP_EICOSANOID_METABOLISM_VIA_LIPO_OXYGENASES_LOX, WP_EICOSANOID_METABOLISM_VIA_LIPO_OXYGENASES_LOX
GOBP_VASOCONSTRICTION, GOBP_VASOCONSTRICTION
HP_ABNORMAL_BLOOD_GAS_LEVEL, HP_ABNORMAL_BLOOD_GAS_LEVEL