

KO_VS_STAT4_KO_CD8_TCELL_UP, GSE40666_STAT1_KO_VS_STAT4_KO_CD8_TCELL_UP

- GSE40666_UNTREATED_VS_IFNA_STIM_CD8_TCELL_90MIN_DN, GSE40666_UNTREATED_VS_IFNA_STIM_CD8_TCELL_90MIN_DN
- GSE37416_CTRL_VS_6H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_CTRL_VS_6H_F_TULARENSIS_LVS_NEUTROPHIL_UP
- GSE40666_WT_VS_STAT4_KO_CD8_TCELL_DN, GSE40666_WT_VS_STAT4_KO_CD8_TCELL_DN
- GSE40666_STAT1_KO_VS_STAT4_KO_CD8_TCELL_WITH_IFNA_STIM_90MIN_UP, GSE40666_STAT1_KO_VS_STAT4_KO_CD8_TCELL_WITH_IFNA_STIM_90MIN_UP
- GSE25677_MPL_VS_R848_STIM_BCELL_DN, GSE25677_MPL_VS_R848_STIM_BCELL_DN
- GSE26488_CTRL_VS_PEPTIDE_INJECTION_OT2_THYMOCYTE_DN, GSE26488_CTRL_VS_PEPTIDE_INJECTION_OT2_THYMOCYTE_DN
- GSE411_100MIN_VS_400MIN_IL6_STIM_MACROPHAGE_UP, GSE411_100MIN_VS_400MIN_IL6_STIM_MACROPHAGE_UP
- GSE7568_CTRL_VS_24H_TGFB_TREATED_MACROPHAGES_WITH_IL4_AND_DEXAMETHASONE_UP, GSE7568_CTRL_VS_24H_TGFB_TREATED_MACROPHAGES_WITH_IL4_AND_DEXAMETHASONE_UP
- GSE26928_NAIVE_VS_CENT_MEMORY_CD4_TCELL_UP, GSE26928_NAIVE_VS_CENT_MEMORY_CD4_TCELL_UP
- GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN, GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN
- GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_IRF8_KO_UP, GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_IRF8_KO_UP
- GSE15659_NAIVE_CD4_TCELL_VS_RESTING_TREG_UP, GSE15659_NAIVE_CD4_TCELL_VS_RESTING_TREG_UP
- MIR370_3P, MIR370_3P
- GSE41867_NAIVE_VS_DAY8_LCMV_CLONE13_EFFECTOR_CD8_TCELL_DN, GSE41867_NAIVE_VS_DAY8_LCMV_CLONE13_EFFECTOR_CD8_TCELL_DN
- MIR4451, MIR4451
- MIR633, MIR633
- GOCC_VESICLE_TETHERING_COMPLEX, GOCC_VESICLE_TETHERING_COMPLEX
- MIR640, MIR640
- HP_BROAD_BASED_GAIT, HP_BROAD_BASED_GAIT
- ZBTB1_TARGET_GENES, ZBTB1_TARGET_GENES
- GSE21379_TFH_VS_NON_TFH_CD4_TCELL_UP, GSE21379_TFH_VS_NON_TFH_CD4_TCELL_UP
- GOMF_RNA_POLYMERASE_ACTIVITY, GOMF_RNA_POLYMERASE_ACTIVITY
- GOBP_CENTRIOLE_ASSEMBLY, GOBP_CENTRIOLE_ASSEMBLY
- MIR5583_3P, MIR5583_3P
- BILD_SRC_ONCOGENIC_SIGNATURE, BILD_SRC_ONCOGENIC_SIGNATURE
- KEGG_INOSITOL_PHOSPHATE_METABOLISM, KEGG_INOSITOL_PHOSPHATE_METABOLISM
- GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_4H_DN, GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_4H_DN
- MIR4254, MIR4254
- MIR6165, MIR6165
- GOBP_LAMELLIPODIUM_ASSEMBLY, GOBP_LAMELLIPODIUM_ASSEMBLY
- MIR5089_3P, MIR5089_3P
- WP_IL1_SIGNALING_PATHWAY, WP_IL1_SIGNALING_PATHWAY
- HP_MEDIAL_FLARING_OF_THE_EYEBROW, HP_MEDIAL_FLARING_OF_THE_EYEBROW
- MIR6774_5P, MIR6774_5P
- MIR1293, MIR1293
- GOMF_PHOSPHATIDYLINOSITOL_BISPHOSPHATE_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_BISPHOSPHATE_PHOSPHATASE_ACTIVITY
- MIR4301, MIR4301
- WP_HYPOTHESIZED_PATHWAYS_IN_PATHOGENESIS_OF_CARDIOVASCULAR_DISEASE, WP_HYPOTHESIZED_PATHWAYS_IN_PATHOGENESIS_OF_CARDIOVASCULAR_DISEASE
- GOBP_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION, GOBP_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION
- REACTOME_INOSITOL_PHOSPHATE_METABOLISM, REACTOME_INOSITOL_PHOSPHATE_METABOLISM
- HP_OVERFRIENDLINESS, HP_OVERFRIENDLINESS
- GOBP_UBIQUINONE_METABOLIC_PROCESS, GOBP_UBIQUINONE_METABOLIC_PROCESS
- MIR92A_1_5P, MIR92A_1_5P
- GOBP_MACROPHAGE_DIFFERENTIATION, GOBP_MACROPHAGE_DIFFERENTIATION
- chr7p15, chr7p15
- MIR659_5P, MIR659_5P
- WP_CANNABINOID_RECEPTOR_SIGNALING, WP_CANNABINOID_RECEPTOR_SIGNALING
- GOBP_NEGATIVE_REGULATION_OF_HIPPO_SIGNALING, GOBP_NEGATIVE_REGULATION_OF_HIPPO_SIGNALING
- LIU_TOPBP1_TARGETS, LIU_TOPBP1_TARGETS
- HP_IRREGULAR_VERTEBRAL_ENDPLATES, HP_IRREGULAR_VERTEBRAL_ENDPLATES
- HP_VITILIGO, HP_VITILIGO
- REACTOME_SYNTHESIS_OF_IP2_IP_AND_INS_IN_THE_CYTOSOL, REACTOME_SYNTHESIS_OF_IP2_IP_AND_INS_IN_THE_CYTOSOL
- GOBP_DETERMINATION_OF_DORSAL_VENTRAL_ASYMMETRY, GOBP_DETERMINATION_OF_DORSAL_VENTRAL_ASYMMETRY