KO\_VS\_STAT4\_KO\_CD8\_TCELL\_UP, GSE40666\_STAT1\_KO\_VS\_STAT4\_KO\_CD8\_TCELL\_UP

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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 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
GSE46606_IRF4_KO_VS_WT_UNSTIM_BCELL_UP, GSE46606_IRF4_KO_VS_WT_UNSTIM_BCELL_UP
GSE25677_MPL_VS_R848_STIM_BCELL_DN, GSE25677_MPL_VS_R848_STIM_BCELL_DN
GSE23308_WT_VS_MINERALCORTICOID_REC_KO_MACROPHAGE_CORTICOSTERONE_TREATED_UP, GSE23308_WT_VS_MINERALCORTICOID_REC_KO_MACROPHAGE_CORTICOSTERONE_TREATED_UP, GSE23308_WT_VS_MINERALCORTICOID_REC_KO_MACROPHAGE_CORTICOSTERONE
GSE40666_WT_VS_STAT4_KO_CD8_TCELL_DN, GSE40666_WT_VS_STAT4_KO_CD8_TCELL_DN
GSE35825_UNTREATED_VS_IFNA_STIM_MACROPHAGE_UP, GSE35825_UNTREATED_VS_IFNA_STIM_MACROPHAGE_UP
GSE17721_CTRL_VS_LPS_6H_BMDC_UP, GSE17721_CTRL_VS_LPS_6H_BMDC_UP
GSE26488_CTRL_VS_PEPTIDE_INJECTION_OT2_THYMOCYTE_DN, GSE26488_CTRL_VS_PEPTIDE_INJECTION_OT2_THYMOCYTE_DN
GSE45365_HEALTHY_VS_MCMV_INFECTION_CD8A_DC_IFNAR_KO_UP, GSE45365_HEALTHY_VS_MCMV_INFECTION_CD8A_DC_IFNAR_KO_UP
 GSE411_100MIN_VS_400MIN_IL6_STIM_MACROPHAGE_UP, GSE411_100MIN_VS_400MIN_IL6_STIM_MACROPHAGE_UP
GSE17721_POLYIC_VS_GARDIQUIMOD_4H_BMDC_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_4H_BMDC_UP
GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN, GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN
GOCC_REPLICATION_FORK, GOCC_REPLICATION_FORK
GSE37416_CTRL_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_CTRL_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_UP
GCM_ERBB2IP, GCM_ERBB2IP
MIR4451, MIR4451
GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP
MIR302F, MIR302F
GSE41867 NAIVE VS DAY8 LCMV CLONE13 EFFECTOR CD8 TCELL DN, GSE41867 NAIVE VS DAY8 LCMV CLONE13 EFFECTOR CD8 TCELL DN
GSE7460_TCONV_VS_TREG_LN_UP, GSE7460_TCONV_VS_TREG_LN_UP
MIR370_3P, MIR370_3P
GSE15659_NAIVE_CD4_TCELL_VS_RESTING_TREG_UP, GSE15659_NAIVE_CD4_TCELL_VS_RESTING_TREG_UP
MIR640, MIR640
TAYLOR_METHYLATED_IN_ACUTE_LYMPHOBLASTIC_LEUKEMIA, TAYLOR_METHYLATED_IN_ACUTE_LYMPHOBLASTIC_LEUKEMIA
GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_4H_DN, GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_4H_DN
MIR5583_3P, MIR5583_3P
GOBP_CENTRIOLE_ASSEMBLY, GOBP_CENTRIOLE_ASSEMBLY
HP_MEDIAL_FLARING_OF_THE_EYEBROW, HP_MEDIAL_FLARING_OF_THE_EYEBROW
MIR5089_3P, MIR5089_3P
WARTERS_IR_RESPONSE_5GY, WARTERS_IR_RESPONSE_5GY
MODULE_333, MODULE_333
MIR1265, MIR1265
GSE43957_UNTREATED_VS_NACL_TREATED_ANTI_CD3_CD28_STIM_CD4_TCELL_UP, GSE43957_UNTREATED_VS_NACL_TREATED_ANTI_CD3_CD28_STIM_CD4_TCELL_UP
WP_IL1_SIGNALING_PATHWAY, WP_IL1_SIGNALING_PATHWAY
HP_INCREASED_LACTATE_DEHYDROGENASE_LEVEL, HP_INCREASED_LACTATE_DEHYDROGENASE_LEVEL
MIR6774_5P, MIR6774_5P
ROSS_AML_OF_FAB_M7_TYPE, ROSS_AML_OF_FAB_M7_TYPE
FONTAINE FOLLICULAR THYROID ADENOMA UP, FONTAINE FOLLICULAR THYROID ADENOMA UP
GOBP_P38MAPK_CASCADE, GOBP_P38MAPK_CASCADE
GOBP_POSITIVE_REGULATION_OF_PROTEIN_BINDING, GOBP_POSITIVE_REGULATION_OF_PROTEIN_BINDING
MIR3138, MIR3138
GOBP_MITOCHONDRIAL_DNA_METABOLIC_PROCESS, GOBP_MITOCHONDRIAL_DNA_METABOLIC_PROCESS
GSE12707_AT16L1_HYPOMORPH_VS_WT_THYMUS_DN, GSE12707_AT16L1_HYPOMORPH_VS_WT_THYMUS_DN
GOBP INOSITOL PHOSPHATE METABOLIC PROCESS, GOBP INOSITOL PHOSPHATE METABOLIC PROCESS
GOBP_REGULATION_OF_HIPPO_SIGNALING, GOBP_REGULATION_OF_HIPPO_SIGNALING
GOBP_QUINONE_METABOLIC_PROCESS, GOBP_QUINONE_METABOLIC_PROCESS
PLASARI_TGFB1_SIGNALING_VIA_NFIC_10HR_DN, PLASARI_TGFB1_SIGNALING_VIA_NFIC_10HR_DN
GOBP_POSITIVE_REGULATION_OF_P38MAPK_CASCADE, GOBP_POSITIVE_REGULATION_OF_P38MAPK_CASCADE
REACTOME_INHIBITION_OF_REPLICATION_INITIATION_OF_DAMAGED_DNA_BY_RB1_E2F1, REACTOME_INHIBITION_OF_REPLICATION_INITIATION_OF_DAMAGED_DNA
GOBP_POLYOL_CATABOLIC_PROCESS, GOBP_POLYOL_CATABOLIC_PROCESS
SRPK2_TARGET_GENES, SRPK2_TARGET_GENES
GOBP_CELLULAR_RESPONSE_TO_ZINC_ION, GOBP_CELLULAR_RESPONSE_TO_ZINC_ION
GOBP_UBIQUINONE_METABOLIC_PROCESS, GOBP_UBIQUINONE_METABOLIC_PROCESS
GSE3720 LPS VS PMA STIM VD2 GAMMADELTA TCELL DN, GSE3720 LPS VS PMA STIM VD2 GAMMADELTA TCELL DN
ROVERSI_GLIOMA_COPY_NUMBER_DN, ROVERSI_GLIOMA_COPY_NUMBER_DN
GOBP NUCLEOTIDE EXCISION REPAIR PREINCISION COMPLEX STABILIZATION, GOBP NUCLEOTIDE EXCISION REPAIR PREINCISION COMPLEX STABILIZATION
GOBP_FRUCTOSE_2_6_BISPHOSPHATE_METABOLIC_PROCESS, GOBP_FRUCTOSE_2_6_BISPHOSPHATE_METABOLIC_PROCESS
GOMF_FRUCTOSE_2_6_BISPHOSPHATE_2_PHOSPHATASE_ACTIVITY, GOMF_FRUCTOSE_2_6_BISPHOSPHATE_2_PHOSPHATASE_ACTIVITY
REACTOME_TELOMERE_C_STRAND_SYNTHESIS_INITIATION, REACTOME_TELOMERE_C_STRAND_SYNTHESIS_INITIATION
LIU_TOPBP1_TARGETS, LIU_TOPBP1_TARGETS
GOMF_PROTEIN_GLUTAMIC_ACID_LIGASE_ACTIVITY, GOMF_PROTEIN_GLUTAMIC_ACID_LIGASE_ACTIVITY
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