```
GSE40666_UNTREATED_VS_IFNA_STIM_CD8_TCELL_90MIN_DN, GSE40666_UNTREATED_VS_IFNA_STIM_CD8_TCELL_90MIN_DN
GSE23308_WT_VS_MINERALCORTICOID_REC_KO_MACROPHAGE_CORTICOSTERONE_TREATED_UP, GSE23308_WT_VS_MINERALCORTICOID_REC_KO_MACROPHAGE_CORTICOSTERONE_TREATED_UP
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
GSE411_100MIN_VS_400MIN_IL6_STIM_MACROPHAGE_UP, GSE411_100MIN_VS_400MIN_IL6_STIM_MACROPHAGE_UP
GSE17721_POLYIC_VS_CPG_6H_BMDC_UP, GSE17721_POLYIC_VS_CPG_6H_BMDC_UP
GSE26488_CTRL_VS_PEPTIDE_INJECTION_OT2_THYMOCYTE_DN, GSE26488_CTRL_VS_PEPTIDE_INJECTION_OT2_THYMOCYTE_DN
GSE13547_WT_VS_ZFX_KO_BCELL_ANTI_IGM_STIM_12H_DN, GSE13547_WT_VS_ZFX_KO_BCELL_ANTI_IGM_STIM_12H_DN
GOCC REPLICATION FORK, GOCC REPLICATION FORK
GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN, GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN
GSE12963_UNINF_VS_ENV_AND_NEF_AND_VPR_DEFICIENT_HIV1_INF_CD4_TCELL_UP, GSE12963_UNINF_VS_ENV_AND_NEF_AND_VPR_DEFICIENT_HIV1_INF_CD4_TCELL_UP
MIR633, MIR633
GSE37416_CTRL_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_CTRL_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_UP
MIR302F, MIR302F
GSE21927_SPLEEN_VS_4T1_TUMOR_MONOCYTE_BALBC_UP, GSE21927_SPLEEN_VS_4T1_TUMOR_MONOCYTE_BALBC_UP
GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION, GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION
MIR4451, MIR4451
BILD_SRC_ONCOGENIC_SIGNATURE, BILD_SRC_ONCOGENIC_SIGNATURE
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
GOCC_CUL4_RING_E3_UBIQUITIN_LIGASE_COMPLEX, GOCC_CUL4_RING_E3_UBIQUITIN_LIGASE_COMPLEX
GSE43957_UNTREATED_VS_NACL_TREATED_ANTI_CD3_CD28_STIM_CD4_TCELL_UP, GSE43957_UNTREATED_VS_NACL_TREATED_ANTI_CD3_CD28_STIM_CD4_TCELL_UP
MIR4536_5P, MIR4536_5P
BAELDE_DIABETIC_NEPHROPATHY_UP, BAELDE_DIABETIC_NEPHROPATHY_UP
KYNG ENVIRONMENTAL STRESS RESPONSE UP, KYNG ENVIRONMENTAL STRESS RESPONSE UP
MIR4254, MIR4254
MIR617, MIR617
NIKOLSKY_BREAST_CANCER_20Q11_AMPLICON, NIKOLSKY_BREAST_CANCER_20Q11_AMPLICON
HP_MALE_REPRODUCTIVE_SYSTEM_NEOPLASM, HP_MALE_REPRODUCTIVE_SYSTEM_NEOPLASM
GOCC_EXTRINSIC_COMPONENT_OF_ORGANELLE_MEMBRANE, GOCC_EXTRINSIC_COMPONENT_OF_ORGANELLE_MEMBRANE
MIR6774_5P, MIR6774_5P
GOCC_INO80_TYPE_COMPLEX, GOCC_INO80_TYPE_COMPLEX
MIR3529_5P, MIR3529_5P
MARZEC_IL2_SIGNALING_DN, MARZEC_IL2_SIGNALING_DN
GOBP_P38MAPK_CASCADE, GOBP_P38MAPK_CASCADE
GOBP_REGULATION_OF_HIPPO_SIGNALING, GOBP_REGULATION_OF_HIPPO_SIGNALING
WP_HYPOTHESIZED_PATHWAYS_IN_PATHOGENESIS_OF_CARDIOVASCULAR_DISEASE, WP_HYPOTHESIZED_PATHWAYS_IN_PATHOGENESIS_OF_CARDIOVASCULAR_DISEASE
REACTOME_TELOMERE_C_STRAND_SYNTHESIS_INITIATION, REACTOME_TELOMERE_C_STRAND_SYNTHESIS_INITIATION
HP VITILIGO, HP VITILIGO
WALLACE_JAK2_TARGETS_UP, WALLACE_JAK2_TARGETS_UP
GOBP_POLYOL_CATABOLIC_PROCESS, GOBP_POLYOL_CATABOLIC_PROCESS
GOBP_MITOCHONDRIAL_DNA_METABOLIC_PROCESS, GOBP_MITOCHONDRIAL_DNA_METABOLIC_PROCESS
GOBP MEMBRANE BIOGENESIS, GOBP MEMBRANE BIOGENESIS
HP_NEOPLASM_OF_THE_MALE_EXTERNAL_GENITALIA, HP_NEOPLASM_OF_THE_MALE_EXTERNAL_GENITALIA
 GOMF_UBIQUITIN_UBIQUITIN_LIGASE_ACTIVITY, GOMF_UBIQUITIN_UBIQUITIN_LIGASE_ACTIVITY
GOBP_NEGATIVE_REGULATION_OF_HIPPO_SIGNALING, GOBP_NEGATIVE_REGULATION_OF_HIPPO_SIGNALING
ZNF454_TARGET_GENES, ZNF454_TARGET_GENES
GOBP_PHOSPHORYLATED_CARBOHYDRATE_DEPHOSPHORYLATION, GOBP_PHOSPHORYLATED_CARBOHYDRATE_DEPHOSPHORYLATION
HP_RENAL_DUPLICATION, HP_RENAL_DUPLICATION
WP_CANNABINOID_RECEPTOR_SIGNALING, WP_CANNABINOID_RECEPTOR_SIGNALING
MIR6863, MIR6863
WP_HEME_BIOSYNTHESIS, WP_HEME_BIOSYNTHESIS
GOBP_PROTOPORPHYRINOGEN_IX_BIOSYNTHETIC_PROCESS, GOBP_PROTOPORPHYRINOGEN_IX_BIOSYNTHETIC_PROCESS
GOBP_NUCLEOBASE_CONTAINING_SMALL_MOLECULE_CATABOLIC_PROCESS, GOBP_NUCLEOBASE_CONTAINING_SMALL_MOLECULE_CATABOLIC_PROCESS
MIR659_5P, MIR659_5P
GOBP_POSITIVE_REGULATION_OF_RUFFLE_ASSEMBLY, GOBP_POSITIVE_REGULATION_OF_RUFFLE_ASSEMBLY
MIR1293, MIR1293
GOBP_INOSITOL_PHOSPHATE_CATABOLIC_PROCESS, GOBP_INOSITOL_PHOSPHATE_CATABOLIC_PROCESS
HP_PEPTIC_ULCER, HP_PEPTIC_ULCER
GOBP_CELLULAR_RESPONSE_TO_HYDROPEROXIDE, GOBP_CELLULAR_RESPONSE_TO_HYDROPEROXIDE
SRPK2_TARGET_GENES, SRPK2_TARGET_GENES
 chr1q41, chr1q41
GOBP_PHOTORECEPTOR_CELL_MAINTENANCE, GOBP_PHOTORECEPTOR_CELL_MAINTENANCE
GOBP_PROTEIN_AUTO_ADP_RIBOSYLATION, GOBP_PROTEIN_AUTO_ADP_RIBOSYLATION
HP_HYPERPIGMENTATION_IN_SUN_EXPOSED_AREAS, HP_HYPERPIGMENTATION_IN_SUN_EXPOSED_AREAS
ONO_AML1_TARGETS_DN, ONO_AML1_TARGETS_DN
WILLIAMS_ESR1_TARGETS_DN, WILLIAMS_ESR1_TARGETS_DN
GOBP_REGULATION_OF_RUFFLE_ASSEMBLY, GOBP_REGULATION_OF_RUFFLE_ASSEMBLY
TANG_SENESCENCE_TP53_TARGETS_UP, TANG_SENESCENCE_TP53_TARGETS_UP
MATSUMIYA_PBMC_MODIFIED_VACCINIA_ANKARA_VACCINE_AGE_18_55YO_VACCINATED_VS_CONTROL_TREATED_IN_VITRO_WITH_WILD_TYPE_MVA_6HR_UP, MATSUMIYA_PBMC_MODIFIED_VACCINATED_VS_CONTROL_TREATED_IN_VITRO_WITH_WILD_TYPE_MVA_6HR_UP, MATSUMIYA_PBMC_MODIFIED_VACCINATED_VS_CONTROL_TREATED_IN_VITRO_WITH_WILD_TYPE_MVA_6HR_UP, MATSUMIYA_PBMC_MODIFIED_VACCINATED_VS_CONTROL_TREATED_IN_VITRO_WITH_WILD_TYPE_MVA_6HR_UP, MATSUMIYA_PBMC_MODIFIED_VACCINATED_VS_CONTROL_TREATED_IN_VITRO_WITH_WILD_TYPE_MVA_6HR_UP, MATSUMIYA_PBMC_MODIFIED_VACCINATED_VS_CONTROL_TREATED_IN_VITRO_WITH_WILD_TYPE_MVA_6HR_UP, MATSUMIYA_PBMC_WITH_WILD_TYPE_MVA_6HR_UP, MATSUMIYA_PBMC_WITH_WILD_TYPE_MVA_6HR_UP, MATSUMIYA_PBMC_WITH_WILD_TYPE_MVA_6HR_UP, MATSUMIYA_PBMC_WITH_WILD_TYPE_MVA_6HR_UP, MATSUMIYA_PBMC_WITH_WILD_TYPE_WITH_WILD_TYPE_WITH_WILD_TYPE_WITH_WILD_TYPE_WITH_WILD_TYPE_WITH_WILD_TYPE_WITH_WILD_TYPE_WITH_WILD_TYPE_WITH_WILD_TYPE_WITH_WILD_TYPE_WIT
 GOBP_MICROTUBULE_SEVERING, GOBP_MICROTUBULE_SEVERING
GOBP_DETERMINATION_OF_DORSAL_VENTRAL_ASYMMETRY, GOBP_DETERMINATION_OF_DORSAL_VENTRAL_ASYMMETRY
ZNF501_TARGET_GENES, ZNF501_TARGET_GENES
HP_HYPOCALCIURIA, HP_HYPOCALCIURIA
GOBP_AUTOPHAGIC_CELL_DEATH, GOBP_AUTOPHAGIC_CELL_DEATH
GOBP_REGULATION_OF_DNA_METHYLATION, GOBP_REGULATION_OF_DNA_METHYLATION
GARGALOVIC RESPONSE TO OXIDIZED PHOSPHOLIPIDS LIGHTGREEN DN, GARGALOVIC RESPONSE TO OXIDIZED PHOSPHOLIPIDS LIGHTGREEN DN
GOBP_POSITIVE_REGULATION_OF_RECEPTOR_BINDING, GOBP_POSITIVE_REGULATION_OF_RECEPTOR_BINDING
GOCC_GATOR2_COMPLEX, GOCC_GATOR2_COMPLEX
GOBP_PROTEIN_OXIDATION, GOBP_PROTEIN_OXIDATION
GOBP_RESPONSE_TO_ZINC_ION, GOBP_RESPONSE_TO_ZINC_ION
GOMF_RNA_POLYMERASE_II_CTD_HEPTAPEPTIDE_REPEAT_PHOSPHATASE_ACTIVITY, GOMF_RNA_POLYMERASE_II_CTD_HEPTAPEPTIDE_REPEAT_PHOSPHATASE_ACTIVITY
HP_SILVER_GRAY_HAIR, HP_SILVER_GRAY_HAIR
GOBP_BLOOD_VESSEL_MATURATION, GOBP_BLOOD_VESSEL_MATURATION
GOBP_PROTEIN_POLY_ADP_RIBOSYLATION, GOBP_PROTEIN_POLY_ADP_RIBOSYLATION
GOMF_CARD_DOMAIN_BINDING, GOMF_CARD_DOMAIN_BINDING
 STARK BRAIN 22Q11 DELETION, STARK BRAIN 22Q11 DELETION
GOBP_REGULATION_OF_SMAD_PROTEIN_COMPLEX_ASSEMBLY, GOBP_REGULATION_OF_SMAD_PROTEIN_COMPLEX_ASSEMBLY
HP_INCREASED_CUP_TO_DISC_RATIO, HP_INCREASED_CUP_TO_DISC_RATIO
GOBP_PEPTIDYL_LYSINE_OXIDATION, GOBP_PEPTIDYL_LYSINE_OXIDATION
GOMF_PROTEIN_LYSINE_6_OXIDASE_ACTIVITY, GOMF_PROTEIN_LYSINE_6_OXIDASE_ACTIVITY
GOBP PEPTIDE CROSS LINKING, GOBP PEPTIDE CROSS LINKING
GOBP_MUCUS_SECRETION, GOBP_MUCUS_SECRETION
GOBP_REGULATION_OF_T_HELPER_17_CELL_LINEAGE_COMMITMENT, GOBP_REGULATION_OF_T_HELPER_17_CELL_LINEAGE_COMMITMENT
GOBP_NEGATIVE_REGULATION_OF_JUN_KINASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_JUN_KINASE_ACTIVITY
HP_HIPPOCAMPAL_ATROPHY, HP_HIPPOCAMPAL_ATROPHY
GOBP_B_CELL_PROLIFERATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_B_CELL_PROLIFERATION_INVOLVED_IN_IMMUNE_RESPONSE
IIZUKA LIVER CANCER PROGRESSION G1 G2 UP, IIZUKA LIVER CANCER PROGRESSION G1 G2 UP
RAMJAUN_APOPTOSIS_BY_TGFB1_VIA_MAPK1_UP, RAMJAUN_APOPTOSIS_BY_TGFB1_VIA_MAPK1_UP
GOMF_FLOPPASE_ACTIVITY, GOMF_FLOPPASE_ACTIVITY
LIU_TOPBP1_TARGETS, LIU_TOPBP1_TARGETS
GOBP_POSITIVE_REGULATION_OF_MAST_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GOBP_POSITIVE_REGULATION_OF_MAST_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE
WANG METASTASIS OF BREAST CANCER, WANG METASTASIS OF BREAST CANCER
GOBP_G_PROTEIN_COUPLED_GLUTAMATE_RECEPTOR_SIGNALING_PATHWAY, GOBP_G_PROTEIN_COUPLED_GLUTAMATE_RECEPTOR_SIGNALING_PATHWAY
HP_RETROCEREBELLAR_CYST, HP_RETROCEREBELLAR_CYST
TERAMOTO_OPN_TARGETS_CLUSTER_5, TERAMOTO_OPN_TARGETS_CLUSTER_5
GOBP_HISTAMINE_PRODUCTION_INVOLVED_IN_INFLAMMATORY_RESPONSE, GOBP_HISTAMINE_PRODUCTION_INVOLVED_IN_INFLAMMATORY_RESPONSE
GOBP_MYOFIBROBLAST_DIFFERENTIATION, GOBP_MYOFIBROBLAST_DIFFERENTIATION
GOBP POSITIVE REGULATION OF AMYLOID BETA CLEARANCE, GOBP POSITIVE REGULATION OF AMYLOID BETA CLEARANCE
GOBP_POLYPHOSPHATE_CATABOLIC_PROCESS, GOBP_POLYPHOSPHATE_CATABOLIC_PROCESS
GOBP_RESPONSE_TO_GLYCOSIDE, GOBP_RESPONSE_TO_GLYCOSIDE
GOBP_RESPONSE_TO_ALDOSTERONE, GOBP_RESPONSE_TO_ALDOSTERONE
GOBP_NEGATIVE_REGULATION_OF_ATP_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_ATP_BIOSYNTHETIC_PROCESS
GOMF_EXOPOLYPHOSPHATASE_ACTIVITY, GOMF_EXOPOLYPHOSPHATASE_ACTIVITY
GOMF_PHOSPHATIDYLCHOLINE_FLOPPASE_ACTIVITY, GOMF_PHOSPHATIDYLCHOLINE_FLOPPASE_ACTIVITY
GOBP_PROTEIN_C_LINKED_GLYCOSYLATION, GOBP_PROTEIN_C_LINKED_GLYCOSYLATION
JONES_TCOF1_TARGETS, JONES_TCOF1_TARGETS
GOBP_NEGATIVE_REGULATION_OF_NUCLEOTIDE_BIOSYNTHETIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_NUCLEOTIDE_BIOSYNTHETIC_PROCESS
GOMF_INTERLEUKIN_17_RECEPTOR_ACTIVITY, GOMF_INTERLEUKIN_17_RECEPTOR_ACTIVITY
GOBP_NEGATIVE_REGULATION_OF_B_CELL_RECEPTOR_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_B_CELL_RECEPTOR_SIGNALING_PATHWAY
HP_BROAD_CHIN, HP_BROAD_CHIN
GOBP_REGULATION_OF_B_CELL_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_B_CELL_RECEPTOR_SIGNALING_PATHWAY
GOBP_CANNABINOID_SIGNALING_PATHWAY, GOBP_CANNABINOID_SIGNALING_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_CELL_FATE_COMMITMENT, GOBP_NEGATIVE_REGULATION_OF_CELL_FATE_COMMITMENT
ZIRN_TRETINOIN_RESPONSE_WT1_DN, ZIRN_TRETINOIN_RESPONSE_WT1_DN
HP_CENTRAL_PRIMITIVE_NEUROECTODERMAL_TUMOR, HP_CENTRAL_PRIMITIVE_NEUROECTODERMAL_TUMOR
GOCC_PROXIMAL_DENDRITE, GOCC_PROXIMAL_DENDRITE
GOMF_1_PHOSPHATIDYLINOSITOL_4_KINASE_ACTIVITY, GOMF_1_PHOSPHATIDYLINOSITOL_4_KINASE_ACTIVITY
FAN_EMBRYONIC_CTX_IN_6_INTERNEURON, FAN_EMBRYONIC_CTX_IN_6_INTERNEURON
GOBP_NEGATIVE_REGULATION_OF_T_HELPER_CELL_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_T_HELPER_CELL_DIFFERENTIATION
HP_VITREORETINOPATHY, HP_VITREORETINOPATHY
HOLLERN_ADENOMYOEPITHELIAL_BREAST_TUMOR, HOLLERN_ADENOMYOEPITHELIAL_BREAST_TUMOR
GOBP_REGULATION_OF_APOPTOTIC_CELL_CLEARANCE, GOBP_REGULATION_OF_APOPTOTIC_CELL_CLEARANCE
chr7p15, chr7p15
GOBP_APOLIPOPROTEIN_A_I_MEDIATED_SIGNALING_PATHWAY, GOBP_APOLIPOPROTEIN_A_I_MEDIATED_SIGNALING_PATHWAY
HP SMALL FOREHEAD, HP SMALL FOREHEAD
GOBP_NEGATIVE_REGULATION_OF_NECROTIC_CELL_DEATH, GOBP_NEGATIVE_REGULATION_OF_NECROTIC_CELL_DEATH
MURAKAMI_UV_RESPONSE_24HR, MURAKAMI_UV_RESPONSE_24HR
HERNANDEZ_ABERRANT_MITOSIS_BY_DOCETACEL_4NM_UP, HERNANDEZ_ABERRANT_MITOSIS_BY_DOCETACEL_4NM_UP
CALVET_IRINOTECAN_SENSITIVE_VS_RESISTANT_UP, CALVET_IRINOTECAN_SENSITIVE_VS_RESISTANT_UP
HP_CHOROID_PLEXUS_CARCINOMA, HP_CHOROID_PLEXUS_CARCINOMA
GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_23_PRODUCTION, GOBP_POSITIVE_REGULATION_OF_INTERLEUKIN_23_PRODUCTION
GOBP_CHEMOKINE_C_X_C_MOTIF_LIGAND_1_PRODUCTION, GOBP_CHEMOKINE_C_X_C_MOTIF_LIGAND_1_PRODUCTION
GOCC_HOST_CELL_PART, GOCC_HOST_CELL_PART
REACTOME_ARACHIDONATE_PRODUCTION_FROM_DAG, REACTOME_ARACHIDONATE_PRODUCTION_FROM_DAG
HP_PERIPHERAL_HYPERMYELINATION, HP_PERIPHERAL_HYPERMYELINATION
GOBP_PHOSPHOLIPID_EFFLUX, GOBP_PHOSPHOLIPID_EFFLUX
chr4q12, chr4q12
HP_FOCAL_HEMICLONIC_SEIZURE, HP_FOCAL_HEMICLONIC_SEIZURE
GOBP_ENDOCANNABINOID_SIGNALING_PATHWAY, GOBP_ENDOCANNABINOID_SIGNALING_PATHWAY
MEISSNER_NPC_ICP_WITH_H3_UNMETHYLATED, MEISSNER_NPC_ICP_WITH_H3_UNMETHYLATED
GOBP_REGULATION_OF_NECROTIC_CELL_DEATH, GOBP_REGULATION_OF_NECROTIC_CELL_DEATH
MURAKAMI UV RESPONSE 6HR DN, MURAKAMI UV RESPONSE 6HR DN
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KO VS STAT4 KO CD8 TCELL UP, GSE40666 STAT1 KO VS STAT4 KO CD8 TCELL UP