WT_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_UP, GSE46606_IRF4HIGH_VS_WT_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_UP

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GSE21670_STAT3_KO_VS_WT_CD4_TCELL_TGFB_TREATED_UP, GSE21670_STAT3_KO_VS_WT_CD4_TCELL_TGFB_TREATED_UP
  GSE42724_MEMORY_BCELL_VS_PLASMABLAST_DN, GSE42724_MEMORY_BCELL_VS_PLASMABLAST_DN
  GSE42724_NAIVE_BCELL_VS_PLASMABLAST_DN, GSE42724_NAIVE_BCELL_VS_PLASMABLAST_DN
  GSE10240_IL17_VS_IL17_AND_IL22_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_UP, GSE10240_IL17_VS_IL17_AND_IL22_STIM_PRIMARY_BRONCHIAL_EPIT
  GSE33424_CD161_HIGH_VS_NEG_CD8_TCELL_UP, GSE33424_CD161_HIGH_VS_NEG_CD8_TCELL_UP
  GSE11864_CSF1_VS_CSF1_IFNG_PAM3CYS_IN_MAC_UP, GSE11864_CSF1_VS_CSF1_IFNG_PAM3CYS_IN_MAC_UP
  GSE21360_NAIVE_VS_PRIMARY_MEMORY_CD8_TCELL_DN, GSE21360_NAIVE_VS_PRIMARY_MEMORY_CD8_TCELL_DN
  GSE39152_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_UP, GSE39152_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_UP
  GSE9239_CTRL_VS_TNF_INHIBITOR_TREATED_DC_UP, GSE9239_CTRL_VS_TNF_INHIBITOR_TREATED_DC_UP
  GOBP_NEGATIVE_REGULATION_OF_DNA_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_DNA_METABOLIC_PROCESS
  MIR1260A_MIR1260B, MIR1260A_MIR1260B
  GOMF_TRANSFERASE_ACTIVITY_TRANSFERRING_HEXOSYL_GROUPS, GOMF_TRANSFERASE_ACTIVITY_TRANSFERRING_HEXOSYL_GROUPS
  GSE32164 RESTING DIFFERENTIATED VS CMYC INHIBITED MACROPHAGE UP, GSE32164 RESTING DIFFERENTIATED VS CMYC INHIBITED MACROPHAGE
  MIR10526_3P, MIR10526_3P
  MIR3920, MIR3920
  MIR2115_5P, MIR2115_5P
  MIR6509_3P, MIR6509_3P
  MIR3190_3P, MIR3190_3P
  MIR1915_3P, MIR1915_3P
  GOBP_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION, GOBP_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION
  BOYAULT_LIVER_CANCER_SUBCLASS_G6_UP, BOYAULT_LIVER_CANCER_SUBCLASS_G6_UP
  MIR2276_3P, MIR2276_3P
 HP_ABNORMALITY_OF_THE_AUDITORY_CANAL, HP_ABNORMALITY_OF_THE_AUDITORY_CANAL
  GOMF_PROTEIN_SERINE_PHOSPHATASE_ACTIVITY, GOMF_PROTEIN_SERINE_PHOSPHATASE_ACTIVITY
  GOBP_NEGATIVE_REGULATION_OF_DNA_REPAIR, GOBP_NEGATIVE_REGULATION_OF_DNA_REPAIR
  MIR3120_5P, MIR3120_5P
  GOBP_PROTEIN_O_LINKED_GLYCOSYLATION, GOBP_PROTEIN_O_LINKED_GLYCOSYLATION
  DR3_Q4, DR3_Q4
  CHENG_RESPONSE_TO_NICKEL_ACETATE, CHENG_RESPONSE_TO_NICKEL_ACETATE
  MIR1298_3P, MIR1298_3P
  GOMF_UDP_GLYCOSYLTRANSFERASE_ACTIVITY, GOMF_UDP_GLYCOSYLTRANSFERASE_ACTIVITY
  GSE45365_NK_CELL_VS_CD11B_DC_UP, GSE45365_NK_CELL_VS_CD11B_DC_UP
  HP_ABNORMALITY_OF_THE_COLUMELLA, HP_ABNORMALITY_OF_THE_COLUMELLA
  GOBP_GLYCOSPHINGOLIPID_METABOLIC_PROCESS, GOBP_GLYCOSPHINGOLIPID_METABOLIC_PROCESS
  RUVBL2_TARGET_GENES, RUVBL2_TARGET_GENES
  CHIANG_LIVER_CANCER_SUBCLASS_CTNNB1_UP, CHIANG_LIVER_CANCER_SUBCLASS_CTNNB1_UP
   MIR379_5P, MIR379_5P
  GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED VS TGFB AND IL12 TREATED ACT CD4 TCELL 2H UP, GSE2770 UNTREATED ACT CD4 TC
   MODULE_285, MODULE_285
   JAZAERI_BREAST_CANCER_BRCA1_VS_BRCA2_DN, JAZAERI_BREAST_CANCER_BRCA1_VS_BRCA2_DN
  GSE13762_CTRL_VS_125_VITAMIND_DAY5_DC_DN, GSE13762_CTRL_VS_125_VITAMIND_DAY5_DC_DN
   GOBP_REGULATION_OF_CHROMATIN_BINDING, GOBP_REGULATION_OF_CHROMATIN_BINDING
  HP_MENINGITIS, HP_MENINGITIS
 HP_AUTOIMMUNE_HEMOLYTIC_ANEMIA, HP_AUTOIMMUNE_HEMOLYTIC_ANEMIA
  GSE29618_PRE_VS_DAY7_FLU_VACCINE_PDC_UP, GSE29618_PRE_VS_DAY7_FLU_VACCINE_PDC_UP
  GOBP_REGULATION_OF_NUCLEAR_CELL_CYCLE_DNA_REPLICATION, GOBP_REGULATION_OF_NUCLEAR_CELL_CYCLE_DNA_REPLICATION
  GOBP_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY, GOBP_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY
  HP_SLURRED_SPEECH, HP_SLURRED_SPEECH
  LI_CYTIDINE_ANALOGS_CYCTOTOXICITY, LI_CYTIDINE_ANALOGS_CYCTOTOXICITY
  GOBP_FUCOSYLATION, GOBP_FUCOSYLATION
  MIR7109_3P, MIR7109_3P
  HP_BIRTH_LENGTH_LESS_THAN_3RD_PERCENTILE, HP_BIRTH_LENGTH_LESS_THAN_3RD_PERCENTILE
  GOBP_POSITIVE_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION, GOBP_POSITIVE_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION
  GOBP_DERMATAN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS, GOBP_DERMATAN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS
HP_RETINAL_COLOBOMA, HP_RETINAL_COLOBOMA
  HP_SHORT_COLUMELLA, HP_SHORT_COLUMELLA
  GOBP_LYMPHATIC_ENDOTHELIAL_CELL_DIFFERENTIATION, GOBP_LYMPHATIC_ENDOTHELIAL_CELL_DIFFERENTIATION
  HP_CLITORAL_HYPOPLASIA, HP_CLITORAL_HYPOPLASIA
  REACTOME_REPRESSION_OF_WNT_TARGET_GENES, REACTOME_REPRESSION_OF_WNT_TARGET_GENES
  NIKOLSKY_BREAST_CANCER_7P22_AMPLICON, NIKOLSKY_BREAST_CANCER_7P22_AMPLICON
  GOMF_GDP_DISSOCIATION_INHIBITOR_ACTIVITY, GOMF_GDP_DISSOCIATION_INHIBITOR_ACTIVITY
  GOBP_LYMPH_VESSEL_MORPHOGENESIS, GOBP_LYMPH_VESSEL_MORPHOGENESIS
  GOBP_POSITIVE_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLATION
  GOBP_SPERM_AXONEME_ASSEMBLY, GOBP_SPERM_AXONEME_ASSEMBLY
  HP_MICROTIA_THIRD_DEGREE, HP_MICROTIA_THIRD_DEGREE
  GOBP_LYMPHANGIOGENESIS, GOBP_LYMPHANGIOGENESIS
  GOBP_NEGATIVE_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION, GOBP_NEGATIVE_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGANIZATION_ORGA
  HOEK_MONOCYTE_2011_2012_TIV_ADULT_1DY_DN, HOEK_MONOCYTE_2011_2012_TIV_ADULT_1DY_DN
   GOMF_LYSOPHOSPHOLIPID_ACYLTRANSFERASE_ACTIVITY, GOMF_LYSOPHOSPHOLIPID_ACYLTRANSFERASE_ACTIVITY
  GOBP_GANGLIOSIDE_METABOLIC_PROCESS, GOBP_GANGLIOSIDE_METABOLIC_PROCESS
  HP_FIBULAR_APLASIA, HP_FIBULAR_APLASIA
  REACTOME_SIGNALING_BY_BMP, REACTOME_SIGNALING_BY_BMP
  HP_APLASIA_INVOLVING_BONES_OF_THE_LOWER_LIMBS, HP_APLASIA_INVOLVING_BONES_OF_THE_LOWER_LIMBS
  HP_BILATERAL_COXA_VALGA, HP_BILATERAL_COXA_VALGA
   KEGG_GLYCOSPHINGOLIPID_BIOSYNTHESIS_GLOBO_SERIES, KEGG_GLYCOSPHINGOLIPID_BIOSYNTHESIS_GLOBO_SERIES
  chr15q23, chr15q23
  GOMF_HEXOSAMINIDASE_ACTIVITY, GOMF_HEXOSAMINIDASE_ACTIVITY
  REACTOME_SYNTHESIS_OF_PIPS_AT_THE_GOLGI_MEMBRANE, REACTOME_SYNTHESIS_OF_PIPS_AT_THE_GOLGI_MEMBRANE
  GOBP_POSITIVE_REGULATION_OF_NEURON_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_NEURON_DIFFERENTIATION
  GOMF_NARROW_PORE_CHANNEL_ACTIVITY, GOMF_NARROW_PORE_CHANNEL_ACTIVITY
  GOBP_REGULATION_OF_PEPTIDYL_CYSTEINE_S_NITROSYLATION, GOBP_REGULATION_OF_PEPTIDYL_CYSTEINE_S_NITROSYLATION
  HP_APLASIA_HYPOPLASIA_OF_THE_TIBIA, HP_APLASIA_HYPOPLASIA_OF_THE_TIBIA
  GOBP_GLYCOSYLCERAMIDE_METABOLIC_PROCESS, GOBP_GLYCOSYLCERAMIDE_METABOLIC_PROCESS
  HP_PROGRESSIVE_SPASTICITY, HP_PROGRESSIVE_SPASTICITY
  GOBP_CERAMIDE_CATABOLIC_PROCESS, GOBP_CERAMIDE_CATABOLIC_PROCESS
  GOMF TRANSMEMBRANE RECEPTOR PROTEIN SERINE THREONINE KINASE ACTIVITY, GOMF TRANSMEMBRANE RECEPTOR PROTEIN SERINE THREON
  GOBP_GLYCOLIPID_CATABOLIC_PROCESS, GOBP_GLYCOLIPID_CATABOLIC_PROCESS
  GOBP CELLULAR RESPONSE TO ETHANOL, GOBP CELLULAR RESPONSE TO ETHANOL
  GOBP_VENOUS_BLOOD_VESSEL_DEVELOPMENT, GOBP_VENOUS_BLOOD_VESSEL_DEVELOPMENT
  GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_CYSTEINE_S_NITROSYLATION, GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_CYSTEINE_S_NITROSYLATION
  MIR11401, MIR11401
  GOBP_ACID_SECRETION, GOBP_ACID_SECRETION
  HP_PULMONARY_INTERSTITIAL_HIGH_RESOLUTION_COMPUTED_TOMOGRAPHY_ABNORMALITY, HP_PULMONARY_INTERSTITIAL_HIGH_PULMONARY_INTERSTITIAL_HIGH_PULMONARY_INTERSTITIAL_HIGH_PULMONARY_INTERSTITIAL_HIGH_PULMONARY_INTERSTITIAL_HIGH_PULMONARY_INTERSTITIAL_HIGH_PULMONARY_INTERSTITIAL_HIGH_PULMONARY_INTERSTITIAL_HIGH_PULMONARY_INTERSTITIAL_HIGH_PULMONARY_INTERSTITIAL_HIGH_PULMONARY_INTERSTITIAL_HIGH_PULMONARY_
  GOBP RETINA VASCULATURE DEVELOPMENT IN CAMERA TYPE EYE, GOBP RETINA VASCULATURE DEVELOPMENT IN CAMERA TYPE EYE
  GOBP HISTONE DEPHOSPHORYLATION, GOBP HISTONE DEPHOSPHORYLATION
  REACTOME_MULTIFUNCTIONAL_ANION_EXCHANGERS, REACTOME_MULTIFUNCTIONAL_ANION_EXCHANGERS
  GOBP_NEGATIVE_REGULATION_OF_COLLATERAL_SPROUTING, GOBP_NEGATIVE_REGULATION_OF_COLLATERAL_SPROUTING
  GOBP_STORE_OPERATED_CALCIUM_ENTRY, GOBP_STORE_OPERATED_CALCIUM_ENTRY
 HP_SLOW_PUPILLARY_LIGHT_RESPONSE, HP_SLOW_PUPILLARY_LIGHT_RESPONSE
  GOMF_SECONDARY_ACTIVE_SULFATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_SECONDARY_ACTIVE_SULFATE_TRANSMEMBRANE_TRANSPO
  REACTOME ELECTRIC TRANSMISSION ACROSS GAP JUNCTIONS, REACTOME ELECTRIC TRANSMISSION ACROSS GAP JUNCTIONS
  GOMF_STORE_OPERATED_CALCIUM_CHANNEL_ACTIVITY, GOMF_STORE_OPERATED_CALCIUM_CHANNEL_ACTIVITY
  GOBP_LONG_TERM_MEMORY, GOBP_LONG_TERM_MEMORY
  GOBP_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_OXIDATIVE_STRESS, GOBP_NEGATIVE_REGULAR_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATIVE_TO_OXIDATI
  GOCC ATP BINDING CASSETTE ABC TRANSPORTER COMPLEX, GOCC ATP BINDING CASSETTE ABC TRANSPORTER COMPLEX
  ZNF426_TARGET_GENES, ZNF426_TARGET_GENES
  GOBP_COLLATERAL_SPROUTING_IN_ABSENCE_OF_INJURY, GOBP_COLLATERAL_SPROUTING_IN_ABSENCE_OF_INJURY
  GOBP_OXALATE_TRANSPORT, GOBP_OXALATE_TRANSPORT
  GOMF_OXALATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_OXALATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
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GOCC_CUTICULAR_PLATE, GOCC_CUTICULAR_PLATE

GSE24726_WT_VS_E2_2_KO_PDC_DAY6_POST_DELETION_UP, GSE24726_WT_VS_E2_2_KO_PDC_DAY6_POST_DELETION_UP

GSE46606_IRF4MID_VS_WT_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_DN, GSE46606_IRF4MID_VS_WT_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_DN