GSE41867_NAIVE_VS_EFFECTOR_CD8_TCELL_DN, GSE41867_NAIVE_VS_EFFECTOR_CD8_TCELL_DN
GSE21360_SECONDARY_VS_TERTIARY_MEMORY_CD8_TCELL_UP, GSE21360_SECONDARY_VS_TERTIARY_MEMORY_CD8_TCELL_UP
GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_DN, GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_UNTREATED_DN
GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_UP, GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_UP
GSE6674_ANTI_IGM_VS_PL2_3_STIM_BCELL_DN, GSE6674_ANTI_IGM_VS_PL2_3_STIM_BCELL_DN
GSE43863_NAIVE_VS_MEMORY_LY6C_INT_CXCR5POS_CD4_TCELL_D150_LCMV_DN, GSE43863_NAIVE_VS_MEMORY_LY6C_INT_CXCR5POS_CD4_TCELL_D150_LCMV_DN
GO_TRANSLATIONAL_TERMINATION, GO_TRANSLATIONAL_TERMINATION
GSE24671_BAKIMULC_VS_SENDAI_VIRUS_INFECTED_MOUSE_SPLENOCYTES_DN, GSE24671_BAKIMULC_VS_SENDAI_VIRUS_INFECTED_MOUSE_SPLENOCYTES_DN
GO_TRANSLATIONAL_ELONGATION, GO_TRANSLATIONAL_ELONGATION
GSE41867_NAIVE_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP
GSE17721_CTRL_VS_POLYIC_4H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_4H_BMDC_UP
GO_CELLULAR_PROTEIN_COMPLEX_DISASSEMBLY, GO_CELLULAR_PROTEIN_COMPLEX_DISASSEMBLY
GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN, GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN
GSE37301_LYMPHOID_PRIMED_MPP_VS_RAG2_KO_NK_CELL_UP, GSE37301_LYMPHOID_PRIMED_MPP_VS_RAG2_KO_NK_CELL_UP
GO_NUCLEOTIDE_EXCISION_REPAIR, GO_NUCLEOTIDE_EXCISION_REPAIR
GSE16385_MONOCYTE_VS_12H_IFNG_TNF_TREATED_MACROPHAGE_UP, GSE16385_MONOCYTE_VS_12H_IFNG_TNF_TREATED_MACROPHAGE_UP
GSE15624_CTRL_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN, GSE15624_CTRL_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN
GSE17721_PAM3CSK4_VS_GADIQUIMOD_8H_BMDC_UP, GSE17721_PAM3CSK4_VS_GADIQUIMOD_8H_BMDC_UP
GSE17721_LPS_VS_POLYIC_24H_BMDC_UP, GSE17721_LPS_VS_POLYIC_24H_BMDC_UP
GSE17721_POLYIC_VS_PAM3CSK4_12H_BMDC_DN, GSE17721_POLYIC_VS_PAM3CSK4_12H_BMDC_DN
GSE1925_CTRL_VS_3H_IFNG_STIM_MACROPHAGE_UP, GSE1925_CTRL_VS_3H_IFNG_STIM_MACROPHAGE_UP
GSE3982_MAC_VS_NEUTROPHIL_UP, GSE3982_MAC_VS_NEUTROPHIL_UP
GO_TRANSCRIPTION_COUPLED_NUCLEOTIDE_EXCISION_REPAIR, GO_TRANSCRIPTION_COUPLED_NUCLEOTIDE_EXCISION_REPAIR
GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_DN, GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_DN
GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_UP, GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_UP
GSE6674_ANTI_IGM_VS_CPG_STIM_BCELL_DN, GSE6674_ANTI_IGM_VS_CPG_STIM_BCELL_DN
GSE7460_CTRL_VS_TGFB_TREATED_ACT_TREG_UP, GSE7460_CTRL_VS_TGFB_TREATED_ACT_TREG_UP
GSE19888_NO_PRETREAT_VS_ADENOSINE_A3R_INHIBITOR_PRETREATED_MAST_CELL_TCELL_MEMBRANES_ACT_DN, GSE19888_NO_PRETREAT_VS_ADENOSINE_A3R_INHIBITOR_PRETREATED_MAST_CEL
GO_RIBOSOMAL_LARGE_SUBUNIT_BIOGENESIS, GO_RIBOSOMAL_LARGE_SUBUNIT_BIOGENESIS
GSE26030_TH1_VS_TH17_DAY5_POST_POLARIZATION_DN, GSE26030_TH1_VS_TH17_DAY5_POST_POLARIZATION_DN
GSE39820_TGFBETA1_IL6_VS_TGFBETA1_IL6_IL23A_TREATED_CD4_TCELL_DN, GSE39820_TGFBETA1_IL6_VS_TGFBETA1_IL6_IL23A_TREATED_CD4_TCELL_DN
GO_ORGANELLAR_RIBOSOME, GO_ORGANELLAR_RIBOSOME
GSE41867_DAY6_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GSE41867_DAY6_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP
GSE27241_WT_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_TH17_POLARIZING_CONDITIONS_DN, GSE27241_WT_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_TH17_POLARIZING
GSE17721_0.5H_VS_12H_POLYIC_BMDC_UP, GSE17721_0.5H_VS_12H_POLYIC_BMDC_UP
GO_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT, GO_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT
GO_NUCLEOLAR_PART, GO_NUCLEOLAR_PART
GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_DN, GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_DN
GSE14415_ACT_VS_CTRL_NATURAL_TREG_DN, GSE14415_ACT_VS_CTRL_NATURAL_TREG_DN
GSE14308_TH2_VS_NATURAL_TREG_UP, GSE14308_TH2_VS_NATURAL_TREG_UP
GSE15767_MED_VS_SCS_MAC_LN_DN, GSE15767_MED_VS_SCS_MAC_LN_DN
GSE30083_SP1_VS_SP3_THYMOCYTE_UP, GSE30083_SP1_VS_SP3_THYMOCYTE_UP
GSE17721_LPS_VS_CPG_0.5H_BMDC_UP, GSE17721_LPS_VS_CPG_0.5H_BMDC_UP
SHEPARD_CRUSH_AND_BURN_MUTANT_UP, SHEPARD_CRUSH_AND_BURN_MUTANT_UP
GO_COFACTOR_BIOSYNTHETIC_PROCESS, GO_COFACTOR_BIOSYNTHETIC_PROCESS
GSE40277_EOS_AND_LEF1_TRANSDUCED_VS_GATA1_AND_SATB1_TRANSDUCED_CD4_TCELL_DN, GSE40277_EOS_AND_LEF1_TRANSDUCED_VS_GATA1_AND_SATB1_TRANSDUCED_CD4_TCELL_DN
GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_UNTREATED_VS_TGFB_AND_IL4_TREATED_ACT_CD4_TCELL_2H_UP
GSE17721_PAM3CSK4_VS_CPG_8H_BMDC_UP, GSE17721_PAM3CSK4_VS_CPG_8H_BMDC_UP
GSE11924_TFH_VS_TH1_CD4_TCELL_DN, GSE11924_TFH_VS_TH1_CD4_TCELL_DN
GSE30971_2H_VS_4H_LPS_STIM_MACROPHAGE_WBP7_HET_UP, GSE30971_2H_VS_4H_LPS_STIM_MACROPHAGE_WBP7_HET_UP
GSE40274_CTRL_VS_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_CTRL_VS_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE21774_CD56_BRIGHT_VS_DIM_CD62L_POSITIVE_NK_CELL_DN, GSE21774_CD56_BRIGHT_VS_DIM_CD62L_POSITIVE_NK_CELL_DN
GSE3982_EOSINOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN, GSE3982_EOSINOPHIL_VS_EFF_MEMORY_CD4_TCELL_DN
GO_PROTEIN_TRANSMEMBRANE_TRANSPORT, GO_PROTEIN_TRANSMEMBRANE_TRANSPORT
GSE43863_NAIVE_VS_TFH_CD4_EFF_TCELL_D6_LCMV_UP, GSE43863_NAIVE_VS_TFH_CD4_EFF_TCELL_D6_LCMV_UP
GSE13547_WT_VS_ZFX_KO_BCELL_ANTI_IGM_STIM_2H_DN, GSE13547_WT_VS_ZFX_KO_BCELL_ANTI_IGM_STIM_2H_DN
GSE17721_CTRL_VS_GARDIQUIMOD_2H_BMDC_UP, GSE17721_CTRL_VS_GARDIQUIMOD_2H_BMDC_UP
GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP, GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP
GSE22611_NOD2_VS_CTRL_TRANSDUCED_HEK293T_CELL_UP, GSE22611_NOD2_VS_CTRL_TRANSDUCED_HEK293T_CELL_UP
GO_ORGANELLAR_LARGE_RIBOSOMAL_SUBUNIT, GO_ORGANELLAR_LARGE_RIBOSOMAL_SUBUNIT
GO_SULFUR_COMPOUND_BIOSYNTHETIC_PROCESS, GO_SULFUR_COMPOUND_BIOSYNTHETIC_PROCESS
GSE23321_CD8_STEM_CELL_MEMORY_VS_EFFECTOR_MEMORY_CD8_TCELL_UP, GSE23321_CD8_STEM_CELL_MEMORY_VS_EFFECTOR_MEMORY_CD8_TCELL_UP
GSE6674_UNSTIM_VS_CPG_STIM_BCELL_DN, GSE6674_UNSTIM_VS_CPG_STIM_BCELL_DN
GO_RNA_CAPPING, GO_RNA_CAPPING
GO_PIGMENT_BIOSYNTHETIC_PROCESS, GO_PIGMENT_BIOSYNTHETIC_PROCESS
GSE17721_CTRL_VS_CPG_12H_BMDC_DN, GSE17721_CTRL_VS_CPG_12H_BMDC_DN
GO_PRERIBOSOME_LARGE_SUBUNIT_PRECURSOR, GO_PRERIBOSOME_LARGE_SUBUNIT_PRECURSOR
GSE17721_0.5H_VS_4H_POLYIC_BMDC_UP, GSE17721_0.5H_VS_4H_POLYIC_BMDC_UP
GO_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_REMOVAL, GO_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_REMOVAL
REACTOME_ELONGATION_ARREST_AND_RECOVERY, REACTOME_ELONGATION_ARREST_AND_RECOVERY
GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_DAMAGE_RECOGNITION, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_DAMAGE_RECOGNITION
REACTOME_NEP_NS2_INTERACTS_WITH_THE_CELLULAR_EXPORT_MACHINERY, REACTOME_NEP_NS2_INTERACTS_WITH_THE_CELLULAR_EXPORT_MACHINERY
KEGG_PROTEIN_EXPORT, KEGG_PROTEIN_EXPORT
REACTOME_INTERACTIONS_OF_VPR_WITH_HOST_CELLULAR_PROTEINS, REACTOME_INTERACTIONS_OF_VPR_WITH_HOST_CELLULAR_PROTEINS
GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_6H_UP, GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_6H_UP
GO_PORPHYRIN_CONTAINING_COMPOUND_METABOLIC_PROCESS, GO_PORPHYRIN_CONTAINING_COMPOUND_METABOLIC_PROCESS
GSE43863_TFH_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN, GSE43863_TFH_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN
REACTOME_TRANSPORT_OF_RIBONUCLEOPROTEINS_INTO_THE_HOST_NUCLEUS, REACTOME_TRANSPORT_OF_RIBONUCLEOPROTEINS_INTO_THE_HOST_NUCLEUS
GSE40274_LEF1_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_LEF1_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GO_TETRAPYRROLE_BIOSYNTHETIC_PROCESS, GO_TETRAPYRROLE_BIOSYNTHETIC_PROCESS
GO_ALPHA_AMINO_ACID_BIOSYNTHETIC_PROCESS, GO_ALPHA_AMINO_ACID_BIOSYNTHETIC_PROCESS
GO_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_EXONUCLEOLYTIC, GO_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_EXONUCLEOLYTIC
GSE41867_DAY6_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN, GSE41867_DAY6_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN
GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMC_4MONTH_POST_DX_DN, GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMC_4MONTH_POST_DX_DN
GO_HEME_BIOSYNTHETIC_PROCESS, GO_HEME_BIOSYNTHETIC_PROCESS
GSE34515_CD16_NEG_VS_POS_MONOCYTE_UP, GSE34515_CD16_NEG_VS_POS_MONOCYTE_UP
GSE23114_PERITONEAL_CAVITY_B1A_BCELL_VS_SPLEEN_BCELL_IN_SLE2C1_MOUSE_DN, GSE23114_PERITONEAL_CAVITY_B1A_BCELL_VS_SPLEEN_BCELL_IN_SLE2C1_MOUSE_DN
GSE25846_IL10_POS_VS_NEG_CD8_TCELL_DAY7_POST_CORONAVIRUS_BRAIN_UP, GSE25846_IL10_POS_VS_NEG_CD8_TCELL_DAY7_POST_CORONAVIRUS_BRAIN_UP
PURBEY_TARGETS_OF_CTBP1_AND_SATB1_DN, PURBEY_TARGETS_OF_CTBP1_AND_SATB1_DN
REACTOME_REGULATION_OF_GLUCOKINASE_BY_GLUCOKINASE_REGULATORY_PROTEIN, REACTOME_REGULATION_OF_GLUCOKINASE_BY_GLUCOKINASE_REGULATORY_PROTEIN
GSE14308_NAIVE_CD4_TCELL_VS_NATURAL_TREG_UP, GSE14308_NAIVE_CD4_TCELL_VS_NATURAL_TREG_UP
GO_RNA_POLYMERASE_ACTIVITY, GO_RNA_POLYMERASE_ACTIVITY
GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_TREATED_MACROPHAGE_UP, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_TREATED_MACROPHAGE_UP
GSE22443_NAIVE_VS_ACT_AND_IL12_TREATED_CD8_TCELL_DN, GSE22443_NAIVE_VS_ACT_AND_IL12_TREATED_CD8_TCELL_DN
GSE15324_ELF4_KO_VS_WT_ACTIVATED_CD8_TCELL_DN, GSE15324_ELF4_KO_VS_WT_ACTIVATED_CD8_TCELL_DN
MODULE_318, MODULE_318
GO_HEME_METABOLIC_PROCESS, GO_HEME_METABOLIC_PROCESS
GO_PIGMENT_METABOLIC_PROCESS, GO_PIGMENT_METABOLIC_PROCESS
GSE360_T_GONDII_VS_M_TUBERCULOSIS_DC_DN, GSE360_T_GONDII_VS_M_TUBERCULOSIS_DC_DN
REACTOME_VIRAL_MESSENGER_RNA_SYNTHESIS, REACTOME_VIRAL_MESSENGER_RNA_SYNTHESIS
GSE18893_TCONV_VS_TREG_2H_TNF_STIM_UP, GSE18893_TCONV_VS_TREG_2H_TNF_STIM_UP
BIOCARTA_CHREBP2_PATHWAY, BIOCARTA_CHREBP2_PATHWAY
GO_COP9_SIGNALOSOME, GO_COP9_SIGNALOSOME
GO_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GO_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
GSE17721_LPS_VS_POLYIC_0.5H_BMDC_DN, GSE17721_LPS_VS_POLYIC_0.5H_BMDC_DN
PID_P38_MK2_PATHWAY, PID_P38_MK2_PATHWAY
GO_NUCLEAR_EXOSOME_RNASE_COMPLEX_, GO_NUCLEAR_EXOSOME_RNASE_COMPLEX_
MODULE_233, MODULE_233
GSE12707_AT16L1_HYPOMORPH_VS_WT_THYMUS_UP, GSE12707_AT16L1_HYPOMORPH_VS_WT_THYMUS_UP
GSE22282_HYPOXIA_VS_NORMOXIA_MYELOID_DC_UP, GSE22282_HYPOXIA_VS_NORMOXIA_MYELOID_DC_UP
GO_TETRAPYRROLE_METABOLIC_PROCESS, GO_TETRAPYRROLE_METABOLIC_PROCESS
EIF4E_UP, EIF4E_UP
TBK1.DN.48HRS_UP, TBK1.DN.48HRS_UP
GO_SULFUR_AMINO_ACID_METABOLIC_PROCESS, GO_SULFUR_AMINO_ACID_METABOLIC_PROCESS
GO_SULFUR_COMPOUND_CATABOLIC_PROCESS, GO_SULFUR_COMPOUND_CATABOLIC_PROCESS
REACTOME_GLUCOSE_TRANSPORT, REACTOME_GLUCOSE_TRANSPORT
GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION, GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_12, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_12
GO_NCRNA_CATABOLIC_PROCESS, GO_NCRNA_CATABOLIC_PROCESS
GO_MYELOID_DENDRITIC_CELL_DIFFERENTIATION, GO_MYELOID_DENDRITIC_CELL_DIFFERENTIATION
GO_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER
GO_CELLULAR_MODIFIED_AMINO_ACID_CATABOLIC_PROCESS, GO_CELLULAR_MODIFIED_AMINO_ACID_CATABOLIC_PROCESS
GSE26488_WT_VS_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN, GSE26488_WT_VS_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN
GSE42021_TCONV_PLN_VS_CD24LO_TCONV_THYMUS_UP, GSE42021_TCONV_PLN_VS_CD24LO_TCONV_THYMUS_UP
REACTOME_DESTABILIZATION_OF_MRNA_BY_TRISTETRAPROLIN_TTP, REACTOME_DESTABILIZATION_OF_MRNA_BY_TRISTETRAPROLIN_TTP
GO_SERINE_FAMILY_AMINO_ACID_METABOLIC_PROCESS, GO_SERINE_FAMILY_AMINO_ACID_METABOLIC_PROCESS
GO_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GO_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE
GO_ENDORIBONUCLEASE_ACTIVITY, GO_ENDORIBONUCLEASE_ACTIVITY
GAJATE_RESPONSE_TO_TRABECTEDIN_DN, GAJATE_RESPONSE_TO_TRABECTEDIN_DN
GO_ENDORIBONUCLEASE_COMPLEX, GO_ENDORIBONUCLEASE_COMPLEX
GO_ANTEROGRADE_AXONAL_TRANSPORT, GO_ANTEROGRADE_AXONAL_TRANSPORT
JAZAG_TGFB1_SIGNALING_UP, JAZAG_TGFB1_SIGNALING_UP
GO_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER
GO_POSITIVE_REGULATION_OF_RNA_SPLICING, GO_POSITIVE_REGULATION_OF_RNA_SPLICING
GO_TUMOR_NECROSIS_FACTOR_RECEPTOR_SUPERFAMILY_BINDING, GO_TUMOR_NECROSIS_FACTOR_RECEPTOR_SUPERFAMILY_BINDING
GO_FANCONI_ANAEMIA_NUCLEAR_COMPLEX, GO_FANCONI_ANAEMIA_NUCLEAR_COMPLEX
GO_METHIONINE_METABOLIC_PROCESS, GO_METHIONINE_METABOLIC_PROCESS
GSE11884_WT_VS_FURIN_KO_NAIVE_CD4_TCELL_DN, GSE11884_WT_VS_FURIN_KO_NAIVE_CD4_TCELL_DN
RIZKI_TUMOR_INVASIVENESS_2D_UP, RIZKI_TUMOR_INVASIVENESS_2D_UP
GO_FATTY_ACID_BIOSYNTHETIC_PROCESS, GO_FATTY_ACID_BIOSYNTHETIC_PROCESS
REACTOME_DESTABILIZATION_OF_MRNA_BY_BRF1, REACTOME_DESTABILIZATION_OF_MRNA_BY_BRF1
GO_GLUTATHIONE_TRANSFERASE_ACTIVITY, GO_GLUTATHIONE_TRANSFERASE_ACTIVITY
GO_S_ADENOSYLMETHIONINE_METABOLIC_PROCESS, GO_S_ADENOSYLMETHIONINE_METABOLIC_PROCESS
GO_SYNAPTONEMAL_COMPLEX, GO_SYNAPTONEMAL_COMPLEX
GO_FATTY_ACYL_COA_METABOLIC_PROCESS, GO_FATTY_ACYL_COA_METABOLIC_PROCESS
REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION, REACTOME_TRANSCRIPTIONAL_REGULATION_OF_WHITE_ADIPOCYTE_DIFFERENTIATION
GO_TUMOR_NECROSIS_FACTOR_RECEPTOR_BINDING, GO_TUMOR_NECROSIS_FACTOR_RECEPTOR_BINDING
GO_MYELOID_DENDRITIC_CELL_ACTIVATION, GO_MYELOID_DENDRITIC_CELL_ACTIVATION
GO_POSITIVE_REGULATION_OF_OSTEOCLAST_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_OSTEOCLAST_DIFFERENTIATION
PAL_PRMT5_TARGETS_DN, PAL_PRMT5_TARGETS_DN
GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_DN
GO_GTPASE_INHIBITOR_ACTIVITY, GO_GTPASE_INHIBITOR_ACTIVITY
MODULE_132, MODULE_132
GO_BLASTOCYST_DEVELOPMENT, GO_BLASTOCYST_DEVELOPMENT
GO_POSITIVE_REGULATION_OF_PROTEIN_IMPORT, GO_POSITIVE_REGULATION_OF_PROTEIN_IMPORT
GO_TRANSLATION_ELONGATION_FACTOR_ACTIVITY, GO_TRANSLATION_ELONGATION_FACTOR_ACTIVITY
GO_CELLULAR_ALDEHYDE_METABOLIC_PROCESS, GO_CELLULAR_ALDEHYDE_METABOLIC_PROCESS
GO_MEIOSIS_I, GO_MEIOSIS_I
GO_PTERIDINE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GO_PTERIDINE_CONTAINING_COMPOUND_METABOLIC_PROCESS
GSE43863_NAIVE_VS_LY6C_INT_CXCR5POS_CD4_EFF_TCELL_D6_LCMV_DN, GSE43863_NAIVE_VS_LY6C_INT_CXCR5POS_CD4_EFF_TCELL_D6_LCMV_DN
GO_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION, GO_POSITIVE_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_INITIATION
GO_CELLULAR_MODIFIED_AMINO_ACID_BIOSYNTHETIC_PROCESS, GO_CELLULAR_MODIFIED_AMINO_ACID_BIOSYNTHETIC_PROCESS
GO_MONOCARBOXYLIC_ACID_BIOSYNTHETIC_PROCESS, GO_MONOCARBOXYLIC_ACID_BIOSYNTHETIC_PROCESS
GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_I
chr12q12, chr12q12
GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_6H_DN, GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_6H_DN