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GSE41867_DAY6_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN, GSE41867_DAY6_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN
GSE23321_EFFECTOR_MEMORY_VS_NAIVE_CD8_TCELL_DN, GSE23321_EFFECTOR_MEMORY_VS_NAIVE_CD8_TCELL_DN
GSE17721_LPS_VS_POLYIC_24H_BMDC_UP, GSE17721_LPS_VS_POLYIC_24H_BMDC_UP
GSE41867_NAIVE_VS_DAY15_LCMV_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY15_LCMV_EFFECTOR_CD8_TCELL_UP
GSE39820_CTRL_VS_TGFBETA3_IL6_CD4_TCELL_UP, GSE39820_CTRL_VS_TGFBETA3_IL6_CD4_TCELL_UP
GSE3039_NKT_CELL_VS_ALPHABETA_CD8_TCELL_UP, GSE3039_NKT_CELL_VS_ALPHABETA_CD8_TCELL_UP
GSE3982_MAC_VS_NEUTROPHIL_UP, GSE3982_MAC_VS_NEUTROPHIL_UP
GSE27786_LSK_VS_CD4_TCELL_UP, GSE27786_LSK_VS_CD4_TCELL_UP
GO_PROTEIN_IMPORT, GO_PROTEIN_IMPORT
GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_UP, GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_UP
GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_24H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_24H_CD8_T_CELL_DN
GSE17721_PAM3CSK4_VS_GADIQUIMOD_24H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_24H_BMDC_DN
GSE8621_UNSTIM_VS_LPS_PRIMED_UNSTIM_MACROPHAGE_UP, GSE8621_UNSTIM_VS_LPS_PRIMED_UNSTIM_MACROPHAGE_UP
GSE17721_LPS_VS_POLYIC_8H_BMDC_UP, GSE17721_LPS_VS_POLYIC_8H_BMDC_UP
GSE31082_DP_VS_CD8_SP_THYMOCYTE_DN, GSE31082_DP_VS_CD8_SP_THYMOCYTE_DN
GSE14308_TH1_VS_NATURAL_TREG_UP, GSE14308_TH1_VS_NATURAL_TREG_UP
GO_PROTEIN_TRANSPORTER_ACTIVITY, GO_PROTEIN_TRANSPORTER_ACTIVITY
GSE7509_DC_VS_MONOCYTE_DN, GSE7509_DC_VS_MONOCYTE_DN
GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_UP, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_UP
GSE19923_E2A_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP, GSE19923_E2A_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP
GSE2128_CTRL_VS_MIMETOPE_NEGATIVE_SELECTION_DP_THYMOCYTE_C57BL6_DN, GSE2128_CTRL_VS_MIMETOPE_NEGATIVE_SELECTION_DP_THYMOCYTE_C57BL6_DN
GSE11924_TH1_VS_TH2_CD4_TCELL_UP, GSE11924_TH1_VS_TH2_CD4_TCELL_UP
GO_SMALL_SUBUNIT_PROCESSOME, GO_SMALL_SUBUNIT_PROCESSOME
GSE37301_HEMATOPOIETIC_STEM_CELL_VS_RAG2_KO_NK_CELL_UP, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_RAG2_KO_NK_CELL_UP
GO_MRNA_SPLICE_SITE_SELECTION, GO_MRNA_SPLICE_SITE_SELECTION
HALLMARK_REACTIVE_OXIGEN_SPECIES_PATHWAY, HALLMARK_REACTIVE_OXIGEN_SPECIES_PATHWAY
GO_SIGNAL_SEQUENCE_BINDING, GO_SIGNAL_SEQUENCE_BINDING
GSE4984_GALECTIN1_VS_LPS_STIM_DC_DN, GSE4984_GALECTIN1_VS_LPS_STIM_DC_DN
LAU_APOPTOSIS_CDKN2A_UP, LAU_APOPTOSIS_CDKN2A_UP
GO_NUCLEAR_LOCALIZATION_SEQUENCE_BINDING, GO_NUCLEAR_LOCALIZATION_SEQUENCE_BINDING
MARZEC_IL2_SIGNALING_UP, MARZEC_IL2_SIGNALING_UP
GO_MITOCHONDRIAL_RNA_METABOLIC_PROCESS, GO_MITOCHONDRIAL_RNA_METABOLIC_PROCESS
GSE33162_HDAC3_KO_VS_HDAC3_KO_MACROPHAGE_UP, GSE33162_HDAC3_KO_VS_HDAC3_KO_MACROPHAGE_UP
GO_AXON_CYTOPLASM, GO_AXON_CYTOPLASM
GO_AXO_DENDRITIC_TRANSPORT, GO_AXO_DENDRITIC_TRANSPORT
GO_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_EXONUCLEOLYTIC, GO_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_EXONUCLEOLYTIC
GSE22282_HYPOXIA_VS_NORMOXIA_MYELOID_DC_UP, GSE22282_HYPOXIA_VS_NORMOXIA_MYELOID_DC_UP
GSE33374_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_UP, GSE33374_CD8_ALPHAALPHA_VS_ALPHABETA_CD161_HIGH_TCELL_UP
GSE31622_WT_VS_KLF3_KO_BCELL_DN, GSE31622_WT_VS_KLF3_KO_BCELL_DN
chr1q42, chr1q42
GO_RESPONSE_TO_OXYGEN_RADICAL, GO_RESPONSE_TO_OXYGEN_RADICAL
REACTOME_MRNA_DECAY_BY_3_TO_5_EXORIBONUCLEASE, REACTOME_MRNA_DECAY_BY_3_TO_5_EXORIBONUCLEASE
TURASHVILI_BREAST_NORMAL_DUCTAL_VS_LOBULAR_UP, TURASHVILI_BREAST_NORMAL_DUCTAL_VS_LOBULAR_UP
GO PROTEIN HOMOTETRAMERIZATION, GO PROTEIN HOMOTETRAMERIZATION
PRAMOONJAGO SOX4 TARGETS DN, PRAMOONJAGO SOX4 TARGETS DN
HOUSTIS ROS, HOUSTIS ROS
GO_MLL1_2_COMPLEX, GO_MLL1_2_COMPLEX
GO_TRANSCRIPTIONAL_REPRESSOR_COMPLEX, GO_TRANSCRIPTIONAL_REPRESSOR_COMPLEX
GO RESPONSE TO ANTIBIOTIC, GO RESPONSE TO ANTIBIOTIC
REACTOME_ACTIVATION_OF_BH3_ONLY_PROTEINS, REACTOME_ACTIVATION_OF_BH3_ONLY_PROTEINS
REACTOME_DESTABILIZATION_OF_MRNA_BY_TRISTETRAPROLIN_TTP, REACTOME_DESTABILIZATION_OF_MRNA_BY_TRISTETRAPROLIN_TTP
STARK_HYPPOCAMPUS_22Q11_DELETION_DN, STARK_HYPPOCAMPUS_22Q11_DELETION_DN
GO_CELL_PROJECTION_CYTOPLASM, GO_CELL_PROJECTION_CYTOPLASM
REACTOME_DESTABILIZATION_OF_MRNA_BY_BRF1, REACTOME_DESTABILIZATION_OF_MRNA_BY_BRF1
BERTUCCI INVASIVE CARCINOMA DUCTAL VS LOBULAR UP, BERTUCCI INVASIVE CARCINOMA DUCTAL VS LOBULAR UP
RB_DN.V1_DN, RB_DN.V1_DN
GSE17721_LPS_VS_PAM3CSK4_2H_BMDC_DN, GSE17721_LPS_VS_PAM3CSK4_2H_BMDC_DN
GO_TUMOR_NECROSIS_FACTOR_RECEPTOR_BINDING, GO_TUMOR_NECROSIS_FACTOR_RECEPTOR_BINDING
MODULE_427, MODULE_427
HINATA_NFKB_TARGETS_KERATINOCYTE_UP, HINATA_NFKB_TARGETS_KERATINOCYTE_UP
GO_GLIAL_CELL_DEVELOPMENT, GO_GLIAL_CELL_DEVELOPMENT
chr3q21, chr3q21
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREEN_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREEN_DN
GO_CELLULAR_RESPONSE_TO_OXYGEN_RADICAL, GO_CELLULAR_RESPONSE_TO_OXYGEN_RADICAL
CHANGOLKAR_H2AFY_TARGETS_DN, CHANGOLKAR_H2AFY_TARGETS_DN
BROWNE_HCMV_INFECTION_8HR_UP, BROWNE_HCMV_INFECTION_8HR_UP
WEIGEL_OXIDATIVE_STRESS_RESPONSE, WEIGEL_OXIDATIVE_STRESS_RESPONSE
GO_SUPEROXIDE_METABOLIC_PROCESS, GO_SUPEROXIDE_METABOLIC_PROCESS
chr3p13, chr3p13
GO_POSITIVE_REGULATION_OF_FATTY_ACID_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_FATTY_ACID_METABOLIC_PROCESS
GO_OLIGODENDROCYTE_DEVELOPMENT, GO_OLIGODENDROCYTE_DEVELOPMENT
BIOCARTA_LONGEVITY_PATHWAY, BIOCARTA_LONGEVITY_PATHWAY
GSE29618_PRE_VS_DAY7_POST_LAIV_FLU_VACCINE_MDC_UP, GSE29618_PRE_VS_DAY7_POST_LAIV_FLU_VACCINE_MDC_UP
MODULE_480, MODULE_480
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GO_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GO_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_

GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_UP, GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_UP

_VS_EFFECTOR_CD8_TCELL_DN, GSE41867_NAIVE_VS_EFFECTOR_CD8_TCELL_DN