RON_CELL_LINE_WEST_EQUINE_ENC_VIRUS_DN, GSE16451_IMMATURE_VS_MATURE_NEURON_CELL_LINE_WEST_EQUINE_ENC_VIRUS_DN

GSE16450_CTRL_VS_IFNA_12H_STIM_IMMATURE_NEURON_CELL_LINE_UP, GSE16450_CTRL_VS_IFNA_12H_STIM_IMMATURE_NEURON_CELL_LINE_UP
GSE17721_POLYIC_VS_GARDIQUIMOD_16H_BMDC_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_16H_BMDC_DN
GSE17974_1.5H_VS_72H_IL4_AND_ANTI_IL12_ACT_CD4_TCELL_DN, GSE17974_1.5H_VS_72H_IL4_AND_ANTI_IL12_ACT_CD4_TCELL_DN
GSE22886_IGM_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_DN, GSE22886_IGM_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_DN
GSE36888_UNTREATED_VS_IL2_TREATED_STAT5_AB_KNOCKIN_TCELL_6H_DN, GSE36888_UNTREATED_VS_IL2_TREATED_STAT5_AB_KNOCKIN_TCELL_6H_DN
GSE5589_LPS_VS_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN, GSE5589_LPS_VS_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN
GSE40274_XBP1_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_XBP1_VS_FOXP3_AND_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE3039_CD4_TCELL_VS_B2_BCELL_DN, GSE3039_CD4_TCELL_VS_B2_BCELL_DN
GSE22886_IGG_IGA_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_DN, GSE22886_IGG_IGA_MEMORY_BCELL_VS_BLOOD_PLASMA_CELL_DN
GSE21678_WT_VS_FOXO1_FOXO3_KO_TREG_DN, GSE21678_WT_VS_FOXO1_FOXO3_KO_TREG_DN
GSE37301_HEMATOPOIETIC_STEM_CELL_VS_RAG2_KO_NK_CELL_UP, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_RAG2_KO_NK_CELL_UP
GO_OUTER_MEMBRANE, GO_OUTER_MEMBRANE
GSE20715_0H_VS_6H_OZONE_TLR4_KO_LUNG_DN, GSE20715_0H_VS_6H_OZONE_TLR4_KO_LUNG_DN
GSE17721_4H_VS_24H_POLYIC_BMDC_UP, GSE17721_4H_VS_24H_POLYIC_BMDC_UP
GSE18804_SPLEEN_MACROPHAGE_VS_TUMORAL_MACROPHAGE_DN, GSE18804_SPLEEN_MACROPHAGE_VS_TUMORAL_MACROPHAGE_DN
GSE14350_IL2RB_KO_VS_WT_TREG_UP, GSE14350_IL2RB_KO_VS_WT_TREG_UP
GSE1460_INTRATHYMIC_T_PROGENITOR_VS_DP_THYMOCYTE_UP, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_DP_THYMOCYTE_UP
GSE41176_UNSTIM_VS_ANTI_IGM_STIM_TAK1_KO_BCELL_6H_UP, GSE41176_UNSTIM_VS_ANTI_IGM_STIM_TAK1_KO_BCELL_6H_UP
GSE21670_STAT3_KO_VS_WT_CD4_TCELL_UP, GSE21670_STAT3_KO_VS_WT_CD4_TCELL_UP
GSE13522_WT_VS_IFNAR_KO_SKING_T_CRUZI_Y_STRAIN_INF_UP, GSE13522_WT_VS_IFNAR_KO_SKING_T_CRUZI_Y_STRAIN_INF_UP
GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_DN, GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_DN
GSE3982_CENT_MEMORY_CD4_TCELL_VS_NKCELL_UP, GSE3982_CENT_MEMORY_CD4_TCELL_VS_NKCELL_UP
GSE25123_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_DAY10_UP, GSE25123_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_DAY10_UP
GSE3039_NKT_CELL_VS_ALPHAALPHA_CD8_TCELL_DN, GSE3039_NKT_CELL_VS_ALPHAALPHA_CD8_TCELL_DN
GO_ALPHA_AMINO_ACID_BIOSYNTHETIC_PROCESS, GO_ALPHA_AMINO_ACID_BIOSYNTHETIC_PROCESS
GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_DN, GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_DN
GSE7218_IGM_VS_IGG_SIGNAL_THGOUGH_ANTIGEN_BCELL_DN, GSE7218_IGM_VS_IGG_SIGNAL_THGOUGH_ANTIGEN_BCELL_DN
GSE2706_UNSTIM_VS_8H_LPS_AND_R848_DC_DN, GSE2706_UNSTIM_VS_8H_LPS_AND_R848_DC_DN
WEI_MIR34A_TARGETS, WEI_MIR34A_TARGETS
GSE14415_NATURAL_TREG_VS_FOXP3_KO_NATURAL_TREG_UP, GSE14415_NATURAL_TREG_VS_FOXP3_KO_NATURAL_TREG_UP
GO_CELLULAR_AMINO_ACID_BIOSYNTHETIC_PROCESS, GO_CELLULAR_AMINO_ACID_BIOSYNTHETIC_PROCESS
GSE39152_SPLEEN_CD103_NEG_VS_BRAIN_CD103_POS_MEMORY_CD8_TCELL_UP, GSE39152_SPLEEN_CD103_NEG_VS_BRAIN_CD103_POS_MEMORY_CD8_TCELL_UP
REACTOME_AMINO_ACID_SYNTHESIS_AND_INTERCONVERSION_TRANSAMINATION, REACTOME_AMINO_ACID_SYNTHESIS_AND_INTERCONVERSION_TRANSAMINATION
GSE360_T_GONDII_VS_M_TUBERCULOSIS_DC_UP, GSE360_T_GONDII_VS_M_TUBERCULOSIS_DC_UP
GSE42088_2H_VS_24H_LEISHMANIA_INF_DC_DN, GSE42088_2H_VS_24H_LEISHMANIA_INF_DC_DN
GSE17721_CTRL_VS_POLYIC_2H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_2H_BMDC_UP
REACTOME_POST_TRANSLATIONAL_MODIFICATION_SYNTHESIS_OF_GPI_ANCHORED_PROTEINS, REACTOME_POST_TRANSLATIONAL_MODIFICATION_SYNTHESIS_OF_GPI_ANCHORED_PROT
REACTOME_SYNTHESIS_OF_GLYCOSYLPHOSPHATIDYLINOSITOL_GPI, REACTOME_SYNTHESIS_OF_GLYCOSYLPHOSPHATIDYLINOSITOL_GPI
GO_GPI_ANCHOR_METABOLIC_PROCESS, GO_GPI_ANCHOR_METABOLIC_PROCESS
PLASARI_TGFB1_TARGETS_10HR_UP, PLASARI_TGFB1_TARGETS_10HR_UP
KEGG_GLYCOSYLPHOSPHATIDYLINOSITOL_GPI_ANCHOR_BIOSYNTHESIS, KEGG_GLYCOSYLPHOSPHATIDYLINOSITOL_GPI_ANCHOR_BIOSYNTHESIS
GO_PREASSEMBLY_OF_GPI_ANCHOR_IN_ER_MEMBRANE, GO_PREASSEMBLY_OF_GPI_ANCHOR_IN_ER_MEMBRANE
GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_MDC_UP, GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_MDC_UP
CGGAARNGGCNG_UNKNOWN, CGGAARNGGCNG_UNKNOWN
GO_MATURATION_OF_5_8S_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA_, GO_MATURATION_OF_5_8S_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIP
GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_FETAL_DN, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_FETAL_DN
PID_MAPK_TRK_PATHWAY, PID_MAPK_TRK_PATHWAY
AMIT_EGF_RESPONSE_60_MCF10A, AMIT_EGF_RESPONSE_60_MCF10A
GSE43863_DAY6_EFF_VS_DAY150_MEM_TFH_CD4_TCELL_DN, GSE43863_DAY6_EFF_VS_DAY150_MEM_TFH_CD4_TCELL_DN
AMIT_SERUM_RESPONSE_40_MCF10A, AMIT_SERUM_RESPONSE_40_MCF10A
GO_PROTEIN_IMPORT_INTO_MITOCHONDRIAL_MATRIX, GO_PROTEIN_IMPORT_INTO_MITOCHONDRIAL_MATRIX
GO_REGULATION_OF_ORGAN_GROWTH, GO_REGULATION_OF_ORGAN_GROWTH
LIU_TARGETS_OF_VMYB_VS_CMYB_UP, LIU_TARGETS_OF_VMYB_VS_CMYB_UP
WANG_BARRETTS_ESOPHAGUS_AND_ESOPHAGUS_CANCER_UP, WANG_BARRETTS_ESOPHAGUS_AND_ESOPHAGUS_CANCER_UP
GO_PROTEIN_MANNOSYLATION, GO_PROTEIN_MANNOSYLATION
GILMORE_CORE_NFKB_PATHWAY, GILMORE_CORE_NFKB_PATHWAY
GO_PROTEIN_CHANNEL_ACTIVITY, GO_PROTEIN_CHANNEL_ACTIVITY
MODULE_427, MODULE_427
GO_GLYCOLIPID_BIOSYNTHETIC_PROCESS, GO_GLYCOLIPID_BIOSYNTHETIC_PROCESS
GO_MANNOSYLATION, GO_MANNOSYLATION
GO_PTERIDINE_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS, GO_PTERIDINE_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS
GO_POSITIVE_REGULATION_OF_CALCIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GO_POSITIVE_REGULATION_OF_CALCIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_GRANULOCYTE_UP, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_GRANULOCYTE_UP
GO_TRANSFERASE_ACTIVITY_TRANSFERRING_ALKYL_OR_ARYL_OTHER_THAN_METHYL_GROUPS, GO_TRANSFERASE_ACTIVITY_TRANSFERRING_ALKYL_OR_ARYL_OTHER_THAN_METHYL_G
GSE43955_1H_VS_42H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN, GSE43955_1H_VS_42H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN
HUNSBERGER_EXERCISE_REGULATED_GENES, HUNSBERGER_EXERCISE_REGULATED_GENES
GO_CELLULAR_METABOLIC_COMPOUND_SALVAGE, GO_CELLULAR_METABOLIC_COMPOUND_SALVAGE
GO_PROTEIN_O_LINKED_MANNOSYLATION, GO_PROTEIN_O_LINKED_MANNOSYLATION
GO_OUTER_MITOCHONDRIAL_MEMBRANE_PROTEIN_COMPLEX, GO_OUTER_MITOCHONDRIAL_MEMBRANE_PROTEIN_COMPLEX
GO_CELLULAR_MODIFIED_AMINO_ACID_BIOSYNTHETIC_PROCESS, GO_CELLULAR_MODIFIED_AMINO_ACID_BIOSYNTHETIC_PROCESS
GSE5589_WT_VS_IL6_KO_LPS_STIM_MACROPHAGE_180MIN_UP, GSE5589_WT_VS_IL6_KO_LPS_STIM_MACROPHAGE_180MIN_UP
GO_REGULATION_OF_CELLULAR_RESPIRATION, GO_REGULATION_OF_CELLULAR_RESPIRATION
GSE10211_UV_INACT_SENDAI_VS_LIVE_SENDAI_VIRUS_TRACHEAL_EPITHELIAL_CELLS_DN, GSE10211_UV_INACT_SENDAI_VS_LIVE_SENDAI_VIRUS_TRACHEAL_EPITHELIAL_CELLS_DN
SANSOM_WNT_PATHWAY_REQUIRE_MYC, SANSOM_WNT_PATHWAY_REQUIRE_MYC
BONOME_OVARIAN_CANCER_POOR_SURVIVAL_DN, BONOME_OVARIAN_CANCER_POOR_SURVIVAL_DN
GO_HEME_BIOSYNTHETIC_PROCESS, GO_HEME_BIOSYNTHETIC_PROCESS
GO_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_OUTER_MEMBRANE, GO_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_OUTER_MEMBRANE
MODULE_120, MODULE_120
GO_T_CELL_MEDIATED_IMMUNITY, GO_T_CELL_MEDIATED_IMMUNITY
GO_POSITIVE_REGULATION_OF_ORGAN_GROWTH, GO_POSITIVE_REGULATION_OF_ORGAN_GROWTH