GSE17721_LPS_VS_PAM3CSK4_4H_BMDC_DN, GSE17721_LPS_VS_PAM3CSK4_4H_BMDC_DN
GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP
GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_DN, GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_DN
GSE32164_RESTING_DIFFERENTIATED_VS_ALTERNATIVELY_ACT_M2_MACROPHAGE_DN, GSE32164_RESTING_DIFFERENTIATED_VS_ALTERNATIVELY_ACT_M2_MACROPHAGE_DN
GSE17721_PAM3CSK4_VS_GADIQUIMOD_6H_BMDC_UP, GSE17721_PAM3CSK4_VS_GADIQUIMOD_6H_BMDC_UP
GSE17721_CTRL_VS_LPS_1H_BMDC_UP, GSE17721_CTRL_VS_LPS_1H_BMDC_UP
GSE14908_RESTING_VS_HDM_STIM_CD4_TCELL_ATOPIC_PATIENT_DN, GSE14908_RESTING_VS_HDM_STIM_CD4_TCELL_ATOPIC_PATIENT_DN
GSE369_PRE_VS_POST_IL6_INJECTION_SOCS3_KO_LIVER_UP, GSE369_PRE_VS_POST_IL6_INJECTION_SOCS3_KO_LIVER_UP
GSE411_UNSTIM_VS_100MIN_IL6_STIM_MACROPHAGE_UP, GSE411_UNSTIM_VS_100MIN_IL6_STIM_MACROPHAGE_UP
GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_UP, GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_UP
GSE17721_0.5H_VS_24H_PAM3CSK4_BMDC_UP, GSE17721_0.5H_VS_24H_PAM3CSK4_BMDC_UP
GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_T425A_HEXON_INF_UP, GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_T425A_HEXON_INF_UP
GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_UP, GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_UP
GSE38681_WT_VS_LYL1_KO_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_UP, GSE38681_WT_VS_LYL1_KO_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_UP
GSE23568_ID3_TRANSDUCED_VS_ID3_KO_CD8_TCELL_UP, GSE23568_ID3_TRANSDUCED_VS_ID3_KO_CD8_TCELL_UP
GSE15330_WT_VS_IKAROS_KO_GRANULOCYTE_MONOCYTE_PROGENITOR_DN, GSE15330_WT_VS_IKAROS_KO_GRANULOCYTE_MONOCYTE_PROGENITOR_DN
GSE21927_UNTREATED_VS_GMCSF_GCSF_TREATED_BONE_MARROW_DN, GSE21927_UNTREATED_VS_GMCSF_GCSF_TREATED_BONE_MARROW_DN
GSE17721_POLYIC_VS_GARDIQUIMOD_2H_BMDC_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_2H_BMDC_UP
GSE33513_TCF7_KO_VS_HET_EARLY_THYMIC_PROGENITOR_DN, GSE33513_TCF7_KO_VS_HET_EARLY_THYMIC_PROGENITOR_DN
SREBP1_01, SREBP1_01
GSE21360_SECONDARY_VS_QUATERNARY_MEMORY_CD8_TCELL_DN, GSE21360_SECONDARY_VS_QUATERNARY_MEMORY_CD8_TCELL_DN
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_10H_UP, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_10H_UP
GO_NEGATIVE_REGULATION_OF_CYTOPLASMIC_TRANSPORT, GO_NEGATIVE_REGULATION_OF_CYTOPLASMIC_TRANSPORT
GSE11961_MEMORY_BCELL_DAY7_VS_PLASMA_CELL_DAY7_DN, GSE11961_MEMORY_BCELL_DAY7_VS_PLASMA_CELL_DAY7_DN
GO_RECYCLING_ENDOSOME, GO_RECYCLING_ENDOSOME
GSE25147_UNSTIM_VS_HELIOBACTER_PYLORI_LPS_STIM_MKN45_CELL_UP, GSE25147_UNSTIM_VS_HELIOBACTER_PYLORI_LPS_STIM_MKN45_CELL_UP
GSE17721_CTRL_VS_LPS_0.5H_BMDC_DN, GSE17721_CTRL_VS_LPS_0.5H_BMDC_DN
GO_NEGATIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT, GO_NEGATIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT
GSE27786_CD4_TCELL_VS_NEUTROPHIL_DN, GSE27786_CD4_TCELL_VS_NEUTROPHIL_DN
HALLMARK_ANDROGEN_RESPONSE, HALLMARK_ANDROGEN_RESPONSE
GSE43955_10H_VS_30H_ACT_CD4_TCELL_DN, GSE43955_10H_VS_30H_ACT_CD4_TCELL_DN
PID_CXCR4_PATHWAY, PID_CXCR4_PATHWAY
GO_NEGATIVE_REGULATION_OF_INTRACELLULAR_TRANSPORT, GO_NEGATIVE_REGULATION_OF_INTRACELLULAR_TRANSPORT
GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_UP, GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_UP
GSE21670_UNTREATED_VS_IL6_TREATED_CD4_TCELL_UP, GSE21670_UNTREATED_VS_IL6_TREATED_CD4_TCELL_UP
HOSHIDA_LIVER_CANCER_SURVIVAL_DN, HOSHIDA_LIVER_CANCER_SURVIVAL_DN
TONG_INTERACT_WITH_PTTG1, TONG_INTERACT_WITH_PTTG1
GO_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION, GO_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION
GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH_GROUP_OF_DONORS_NAD_OR_NADP_AS_ACCEPTOR, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH_GROUP_OF_DO
GSE17721_CPG_VS_GARDIQUIMOD_2H_BMDC_UP, GSE17721_CPG_VS_GARDIQUIMOD_2H_BMDC_UP
GO_POSITIVE_REGULATION_OF_RESPONSE_TO_EXTRACELLULAR_STIMULUS, GO_POSITIVE_REGULATION_OF_RESPONSE_TO_EXTRACELLULAR_STIMULUS
DORSAM_HOXA9_TARGETS_UP, DORSAM_HOXA9_TARGETS_UP
GO_EPITHELIAL_CELL_APOPTOTIC_PROCESS, GO_EPITHELIAL_CELL_APOPTOTIC_PROCESS
GSE2585_THYMIC_DC_VS_THYMIC_MACROPHAGE_UP, GSE2585_THYMIC_DC_VS_THYMIC_MACROPHAGE_UP
RAY_TARGETS_OF_P210_BCR_ABL_FUSION_DN, RAY_TARGETS_OF_P210_BCR_ABL_FUSION_DN
GO_SNARE_BINDING, GO_SNARE_BINDING
GO_VESICLE_DOCKING_INVOLVED_IN_EXOCYTOSIS, GO_VESICLE_DOCKING_INVOLVED_IN_EXOCYTOSIS
GO_POSITIVE_REGULATION_OF_INSULIN_RECEPTOR_SIGNALING_PATHWAY, GO_POSITIVE_REGULATION_OF_INSULIN_RECEPTOR_SIGNALING_PATHWAY
GO_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_PERMEABILITY_INVOLVED_IN_APOPTOTIC_PROCESS, GO_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_PERMEABILITY_INVOLVED_I
GO_CELLULAR_RESPONSE_TO_LITHIUM_ION, GO_CELLULAR_RESPONSE_TO_LITHIUM_ION
GO_FOLIC_ACID_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS, GO_FOLIC_ACID_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_RAC_PROTEIN_SIGNAL_TRANSDUCTION, GO_REGULATION_OF_RAC_PROTEIN_SIGNAL_TRANSDUCTION
GO_TAU_PROTEIN_KINASE_ACTIVITY, GO_TAU_PROTEIN_KINASE_ACTIVITY
GALLUZZI_PERMEABILIZE_MITOCHONDRIA, GALLUZZI_PERMEABILIZE_MITOCHONDRIA
GO_GLIOGENESIS, GO_GLIOGENESIS
GO_RESPONSE_TO_LITHIUM_ION, GO_RESPONSE_TO_LITHIUM_ION
GO_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION, GO_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION
MODULE_169, MODULE_169
LANDIS_BREAST_CANCER_PROGRESSION_UP, LANDIS_BREAST_CANCER_PROGRESSION_UP
YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_6, YAO_TEMPORAL_RESPONSE_TO_PROGESTERONE_CLUSTER_6
GO_EXTRINSIC_COMPONENT_OF_PLASMA_MEMBRANE, GO_EXTRINSIC_COMPONENT_OF_PLASMA_MEMBRANE
GO_NEGATIVE_REGULATION_OF_GLIAL_CELL_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_GLIAL_CELL_DIFFERENTIATION
GO_POSITIVE_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_PERMEABILITY, GO_POSITIVE_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_PERMEABILITY
GO_REGULATION_OF_STEM_CELL_DIFFERENTIATION, GO_REGULATION_OF_STEM_CELL_DIFFERENTIATION
GSE34205_RSV_VS_FLU_INF_INFANT_PBMC_UP, GSE34205_RSV_VS_FLU_INF_INFANT_PBMC_UP
GO_EXOCYTIC_VESICLE, GO_EXOCYTIC_VESICLE
GO_STEM_CELL_DIFFERENTIATION, GO_STEM_CELL_DIFFERENTIATION
GO_PERINUCLEAR_ENDOPLASMIC_RETICULUM, GO_PERINUCLEAR_ENDOPLASMIC_RETICULUM
REACTOME_G_PROTEIN_ACTIVATION, REACTOME_G_PROTEIN_ACTIVATION
ST_WNT_CA2_CYCLIC_GMP_PATHWAY, ST_WNT_CA2_CYCLIC_GMP_PATHWAY

A2AR_AGONIST_TREATED_TREG_UP, GSE34006_UNTREATED_VS_A2AR_AGONIST_TREATED_TREG_UP