
GSE31082_DP_VS_CD8_SP_THYMOCYTE_UP, GSE31082_DP_VS_CD8_SP_THYMOCYTE_UP
GSE22886_NAIVE_TCELL_VS_MONOCYTE_DN, GSE22886_NAIVE_TCELL_VS_MONOCYTE_DN
GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_DN, GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_DN
GSE22886_NAIVE_CD4_TCELL_VS_MONOCYTE_DN, GSE22886_NAIVE_CD4_TCELL_VS_MONOCYTE_DN
GSE29164_DAY3_VS_DAY7_UNTREATED_MELANOMA_DN, GSE29164_DAY3_VS_DAY7_UNTREATED_MELANOMA_DN
GSE17186_BLOOD_VS_CORD_BLOOD_NAIVE_BCELL_UP, GSE17186_BLOOD_VS_CORD_BLOOD_NAIVE_BCELL_UP
GSE9509_10MIN_VS_30MIN_LPS_STIM_IL10_KO_MACROPHAGE_UP, GSE9509_10MIN_VS_30MIN_LPS_STIM_IL10_KO_MACROPHAGE_UP
GSE17186_BLOOD_VS_CORD_BLOOD_CD21LOW_TRANSITIONAL_BCELL_DN, GSE17186_BLOOD_VS_CORD_BLOOD_CD21LOW_TRANSITIONAL_BCELL_DN
GSE12392_WT_VS_IFNB_KO_CD8A_POS_SPLEEN_DC_UP, GSE12392_WT_VS_IFNB_KO_CD8A_POS_SPLEEN_DC_UP
GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_20MIN_DN, GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_20MIN_DN
GSE411_WT_VS_SOCS3_KO_MACROPHAGE_UP, GSE411_WT_VS_SOCS3_KO_MACROPHAGE_UP
GSE17186_MEMORY_VS_CD21LOW_TRANSITIONAL_BCELL_UP, GSE17186_MEMORY_VS_CD21LOW_TRANSITIONAL_BCELL_UP
GSE17186_NAIVE_VS_CD21HIGH_TRANSITIONAL_BCELL_DN, GSE17186_NAIVE_VS_CD21HIGH_TRANSITIONAL_BCELL_DN
GSE17186_CD21LOW_VS_CD21HIGH_TRANSITIONAL_BCELL_CORD_BLOOD_UP, GSE17186_CD21LOW_VS_CD21HIGH_TRANSITIONAL_BCELL_CORD_BLOOD_UP
GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNAR_KO_DN, GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_IFNAR_KO_DN
GSE37563_WT_VS_CTLA4_KO_CD4_TCELL_D4_POST_IMMUNIZATION_DN, GSE37563_WT_VS_CTLA4_KO_CD4_TCELL_D4_POST_IMMUNIZATION_DN
GSE557_WT_VS_CIITA_KO_DC_UP, GSE557_WT_VS_CIITA_KO_DC_UP
GSE40666_WT_VS_STAT1_KO_CD8_TCELL_DN, GSE40666_WT_VS_STAT1_KO_CD8_TCELL_DN
GSE8685_IL2_ACT_IL2_STARVED_VS_IL15_ACT_IL2_STARVED_CD4_TCELL_DN, GSE8685_IL2_ACT_IL2_STARVED_VS_IL15_ACT_IL2_STARVED_CD4_TCELL_DN
GSE11057_NAIVE_CD4_VS_PBMC_CD4_TCELL_DN, GSE11057_NAIVE_CD4_VS_PBMC_CD4_TCELL_DN
GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IFNG_TNF_TREATED_MACROPHAGE_DN
GSE26030_UNSTIM_VS_RESTIM_TH1_DAY5_POST_POLARIZATION_DN, GSE26030_UNSTIM_VS_RESTIM_TH1_DAY5_POST_POLARIZATION_DN
GSE21670_TGFB_VS_TGFB_AND_IL6_TREATED_STAT3_KO_CD4_TCELL_DN, GSE21670_TGFB_VS_TGFB_AND_IL6_TREATED_STAT3_KO_CD4_TCELL_DN
GSE12392_WT_VS_IFNB_KO_CD8A_NEG_SPLEEN_DC_UP, GSE12392_WT_VS_IFNB_KO_CD8A_NEG_SPLEEN_DC_UP
GSE14386_UNTREATED_VS_IFNA_TREATED_ACT_PBMC_MS_PATIENT_DN, GSE14386_UNTREATED_VS_IFNA_TREATED_ACT_PBMC_MS_PATIENT_DN
GSE34392_ST2_KO_VS_WT_DAY8_LCMV_EFFECTOR_CD8_TCELL_DN, GSE34392_ST2_KO_VS_WT_DAY8_LCMV_EFFECTOR_CD8_TCELL_DN
GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN, GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN
GNF2_CASP1, GNF2_CASP1
GSE42724_NAIVE_VS_MEMORY_BCELL_DN, GSE42724_NAIVE_VS_MEMORY_BCELL_DN
GSE22935_WT_VS_MYD88_KO_MACROPHAGE_UP, GSE22935_WT_VS_MYD88_KO_MACROPHAGE_UP
GSE46606_IRF4MID_VS_WT_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_UP, GSE46606_IRF4MID_VS_WT_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_UP
GSE15767_MED_VS_SCS_MAC_LN_UP, GSE15767_MED_VS_SCS_MAC_LN_UP
GSE24210_TCONV_VS_TREG_DN, GSE24210_TCONV_VS_TREG_DN
GSE26343_UNSTIM_VS_LPS_STIM_NFAT5_KO_MACROPHAGE_DN, GSE26343_UNSTIM_VS_LPS_STIM_NFAT5_KO_MACROPHAGE_DN
GSE16697_CD4_TCELL_VS_TFH_CD4_TCELL_UP, GSE16697_CD4_TCELL_VS_TFH_CD4_TCELL_UP
GSE7219_UNSTIM_VS_LPS_AND_ANTI_CD40_STIM_DC_DN, GSE7219_UNSTIM_VS_LPS_AND_ANTI_CD40_STIM_DC_DN
GSE17186_MEMORY_VS_NAIVE_BCELL_DN, GSE17186_MEMORY_VS_NAIVE_BCELL_DN
GSE6269_HEALTHY_VS_FLU_INF_PBMC_DN, GSE6269_HEALTHY_VS_FLU_INF_PBMC_DN
GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TREG_DN, GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TREG_DN
GSE8921_3H_VS_24H_TLR1_2_STIM_MONOCYTE_DN, GSE8921_3H_VS_24H_TLR1_2_STIM_MONOCYTE_DN
GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_DN, GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_DN
GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
GSE30083_SP3_VS_SP4_THYMOCYTE_DN, GSE30083_SP3_VS_SP4_THYMOCYTE_DN
HALLMARK_COMPLEMENT, HALLMARK_COMPLEMENT
GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_DN, GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_DN
GNF2_HCK, GNF2_HCK
GSE339_CD8POS_VS_CD4CD8DN_DC_UP, GSE339_CD8POS_VS_CD4CD8DN_DC_UP
GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_UP, GSE29164_UNTREATED_VS_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DAY3_UP
PID_P53_REGULATION_PATHWAY, PID_P53_REGULATION_PATHWAY
GSE7852_TREG_VS_TCONV_THYMUS_UP, GSE7852_TREG_VS_TCONV_THYMUS_UP
GSE339_CD4POS_VS_CD8POS_DC_DN, GSE339_CD4POS_VS_CD8POS_DC_DN
GSE17721_CTRL_VS_POLYIC_8H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_8H_BMDC_DN
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_UP, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_UP
GSE17721_CPG_VS_GARDIQUIMOD_4H_BMDC_UP, GSE17721_CPG_VS_GARDIQUIMOD_4H_BMDC_UP
GSE3982_MAC_VS_TH1_UP, GSE3982_MAC_VS_TH1_UP
GSE25087_FETAL_VS_ADULT_TCONV_DN, GSE25087_FETAL_VS_ADULT_TCONV_DN
GSE7852_TREG_VS_TCONV_UP, GSE7852_TREG_VS_TCONV_UP
GSE45365_NK_CELL_VS_CD8A_DC_UP, GSE45365_NK_CELL_VS_CD8A_DC_UP
GSE3039_NKT_CELL_VS_B2_BCELL_UP, GSE3039_NKT_CELL_VS_B2_BCELL_UP
GSE23568_CTRL_TRANSDUCED_VS_WT_CD8_TCELL_UP, GSE23568_CTRL_TRANSDUCED_VS_WT_CD8_TCELL_UP
GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP, GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP
GSE11961_MARGINAL_ZONE_BCELL_VS_GERMINAL_CENTER_BCELL_DAY40_DN, GSE11961_MARGINAL_ZONE_BCELL_VS_GERMINAL_CENTER_BCELL_DAY40_DN
GSE22886_NAIVE_BCELL_VS_BM_PLASMA_CELL_DN, GSE22886_NAIVE_BCELL_VS_BM_PLASMA_CELL_DN
GSE11961_MARGINAL_ZONE_BCELL_VS_MEMORY_BCELL_DAY40_UP, GSE11961_MARGINAL_ZONE_BCELL_VS_MEMORY_BCELL_DAY40_UP
GSE3982_DC_VS_TH2_UP, GSE3982_DC_VS_TH2_UP
GSE3982_MAST_CELL_VS_MAC_DN, GSE3982_MAST_CELL_VS_MAC_DN
GSE3982_EFF_MEMORY_CD4_TCELL_VS_NKCELL_DN, GSE3982_EFF_MEMORY_CD4_TCELL_VS_NKCELL_DN
LIU_CMYB_TARGETS_UP, LIU_CMYB_TARGETS_UP
GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_UP, GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_UP
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_10H_UP, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_10H_UP
GSE27670_CTRL_VS_LMP1_TRANSDUCED_GC_BCELL_UP, GSE27670_CTRL_VS_LMP1_TRANSDUCED_GC_BCELL_UP
GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_48H_CD8_T_CELL_UP, GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_48H_CD8_T_CELL_UP
GSE37534_GW1929_VS_ROSIGLITAZONE_TREATED_CD4_TCELL_PPARG1_FOXP3_TRANSDUCED_UP, GSE37534_GW1929_VS_ROSIGLITAZONE_TREATED_CD4_TCELL_PPARG1_FOXP3_TRANSDUCED_UP
LANDIS_ERBB2_BREAST_TUMORS_324_UP, LANDIS_ERBB2_BREAST_TUMORS_324_UP
GSE45365_WT_VS_IFNAR_KO_CD8A_DC_DN, GSE45365_WT_VS_IFNAR_KO_CD8A_DC_DN
GSE23925_LIGHT_ZONE_VS_DARK_ZONE_BCELL_DN, GSE23925_LIGHT_ZONE_VS_DARK_ZONE_BCELL_DN
GSE21927_SPLEEN_VS_C26GM_TUMOR_MONOCYTE_BALBC_DN, GSE21927_SPLEEN_VS_C26GM_TUMOR_MONOCYTE_BALBC_DN
GSE5542_IFNG_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_24H_DN, GSE5542_IFNG_VS_IFNA_AND_IFNG_TREATED_EPITHELIAL_CELLS_24H_DN
GO_RESPONSE_TO_FLUID_SHEAR_STRESS, GO_RESPONSE_TO_FLUID_SHEAR_STRESS
GO_AMINOGLYCAN_BIOSYNTHETIC_PROCESS, GO_AMINOGLYCAN_BIOSYNTHETIC_PROCESS
GO_AMINOGLYCAN_METABOLIC_PROCESS, GO_AMINOGLYCAN_METABOLIC_PROCESS
ZHAN_MULTIPLE_MYELOMA_CD1_AND_CD2_DN, ZHAN_MULTIPLE_MYELOMA_CD1_AND_CD2_DN
BASSO_CD40_SIGNALING_UP, BASSO_CD40_SIGNALING_UP
GSE37301_MULTIPOTENT_PROGENITOR_VS_GRAN_MONO_PROGENITOR_DN, GSE37301_MULTIPOTENT_PROGENITOR_VS_GRAN_MONO_PROGENITOR_DN
BOYLAN_MULTIPLE_MYELOMA_D_DN, BOYLAN_MULTIPLE_MYELOMA_D_DN
GO_POSITIVE_REGULATION_OF_HEMOPOIESIS, GO_POSITIVE_REGULATION_OF_HEMOPOIESIS
GSE20715_WT_VS_TLR4_KO_24H_OZONE_LUNG_UP, GSE20715_WT_VS_TLR4_KO_24H_OZONE_LUNG_UP
GO_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS, GO_MUCOPOLYSACCHARIDE_METABOLIC_PROCESS
HASLINGER_B_CLL_WITH_CHROMOSOME_12_TRISOMY, HASLINGER_B_CLL_WITH_CHROMOSOME_12_TRISOMY
GSE37532_VISCERAL_ADIPOSE_TISSUE_VS_LN_DERIVED_TREG_CD4_TCELL_UP, GSE37532_VISCERAL_ADIPOSE_TISSUE_VS_LN_DERIVED_TREG_CD4_TCELL_UP
GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_DN, GSE39820_TGFBETA1_VS_TGFBETA3_IN_IL6_IL23A_TREATED_CD4_TCELL_DN
REACTOME_GLYCOSAMINOGLYCAN_METABOLISM, REACTOME_GLYCOSAMINOGLYCAN_METABOLISM
GO_CELLULAR_RESPONSE_TO_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_STIMULUS, GO_CELLULAR_RESPONSE_TO_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_STIMULUS
GO_PROTEIN_EXPORT_FROM_NUCLEUS, GO_PROTEIN_EXPORT_FROM_NUCLEUS
SIRNA_EIF4GI_DN, SIRNA_EIF4GI_DN
GSE15659_NAIVE_CD4_TCELL_VS_RESTING_TREG_UP, GSE15659_NAIVE_CD4_TCELL_VS_RESTING_TREG_UP
GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GFP_TCONV_UP, GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GFP_TCONV_UP
GSE26023_PHD3_KO_VS_WT_NEUTROPHIL_HYPOXIA_UP, GSE26023_PHD3_KO_VS_WT_NEUTROPHIL_HYPOXIA_UP
GNF2_CD14, GNF2_CD14
GO_PROTEIN_TYROSINE_KINASE_BINDING, GO_PROTEIN_TYROSINE_KINASE_BINDING
BOYLAN_MULTIPLE_MYELOMA_D_CLUSTER_DN, BOYLAN_MULTIPLE_MYELOMA_D_CLUSTER_DN
KYNG_RESPONSE_TO_H2O2, KYNG_RESPONSE_TO_H2O2
BIOCARTA_GLEEVEC_PATHWAY, BIOCARTA_GLEEVEC_PATHWAY
GSE3982_NEUTROPHIL_VS_TH2_UP, GSE3982_NEUTROPHIL_VS_TH2_UP
PID_IL4_2PATHWAY, PID_IL4_2PATHWAY
MMS_MOUSE_LYMPH_HIGH_4HRS_UP, MMS_MOUSE_LYMPH_HIGH_4HRS_UP
MODULE_356, MODULE_356
GO_SITE_OF_POLARIZED_GROWTH, GO_SITE_OF_POLARIZED_GROWTH
ZHAN_MULTIPLE_MYELOMA_LB_DN, ZHAN_MULTIPLE_MYELOMA_LB_DN
BIOCARTA_P38MAPK_PATHWAY, BIOCARTA_P38MAPK_PATHWAY
GO_PROTEIN_DEGLYCOSYLATION, GO_PROTEIN_DEGLYCOSYLATION
BCAT_GDS748_DN, BCAT_GDS748_DN
GO_LIPOPOLYSACCHARIDE_MEDIATED_SIGNALING_PATHWAY, GO_LIPOPOLYSACCHARIDE_MEDIATED_SIGNALING_PATHWAY
GO_LYMPHOCYTE_HOMEOSTASIS, GO_LYMPHOCYTE_HOMEOSTASIS
GSE29164_DAY3_VS_DAY7_CD8_TCELL_TREATED_MELANOMA_DN, GSE29164_DAY3_VS_DAY7_CD8_TCELL_TREATED_MELANOMA_DN
GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_UP, GSE17721_CPG_VS_GARDIQUIMOD_12H_BMDC_UP
MODULE_576, MODULE_576
GO_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE, GO_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE
GO_POSITIVE_REGULATION_OF_LEUKOCYTE_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_LEUKOCYTE_DIFFERENTIATION
MODULE_195, MODULE_195
PDGF_ERK_DN.V1_UP, PDGF_ERK_DN.V1_UP
GO_POSITIVE_REGULATION_OF_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION, GO_POSITIVE_REGULATION_OF_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION
GO_T_CELL_HOMEOSTASIS, GO_T_CELL_HOMEOSTASIS
GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR
KRIEG_HYPOXIA_VIA_KDM3A, KRIEG_HYPOXIA_VIA_KDM3A
GSE21927_SPLEEN_VS_4T1_TUMOR_MONOCYTE_BALBC_UP, GSE21927_SPLEEN_VS_4T1_TUMOR_MONOCYTE_BALBC_UP
GO_NEGATIVE_REGULATION_OF_AXONOGENESIS, GO_NEGATIVE_REGULATION_OF_AXONOGENESIS
CHEN_HOXA5_TARGETS_9HR_DN, CHEN_HOXA5_TARGETS_9HR_DN
WILENSKY_RESPONSE_TO_DARAPLADIB, WILENSKY_RESPONSE_TO_DARAPLADIB
PID_NEPHRIN_NEPH1_PATHWAY, PID_NEPHRIN_NEPH1_PATHWAY
GO_PROTEIN_DEMANNOSYLATION, GO_PROTEIN_DEMANNOSYLATION
GO_POSITIVE_REGULATION_OF_MITOCHONDRIAL_OUTER_MEMBRANE_PERMEABILIZATION_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY, GO_POSITIVE_REGULATION_OF_MITOCHONDRIAL_OUTER_MEMBRANE_PERMEABILIZATION_INVOLV
WOO_LIVER_CANCER_RECURRENCE_UP, WOO_LIVER_CANCER_RECURRENCE_UP
GAVIN_IL2_RESPONSIVE_FOXP3_TARGETS_UP, GAVIN_IL2_RESPONSIVE_FOXP3_TARGETS_UP
GSE29949_DC_BRAIN_VS_MONOCYTE_BONE_MARROW_DN, GSE29949_DC_BRAIN_VS_MONOCYTE_BONE_MARROW_DN
GO_ERBB_SIGNALING_PATHWAY, GO_ERBB_SIGNALING_PATHWAY
GO_GOLGI_CISTERNA_MEMBRANE, GO_GOLGI_CISTERNA_MEMBRANE
TCGA_GLIOBLASTOMA_COPY_NUMBER_UP, TCGA_GLIOBLASTOMA_COPY_NUMBER_UP
GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN, GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN
GO_MYELOID_LEUKOCYTE_ACTIVATION, GO_MYELOID_LEUKOCYTE_ACTIVATION
BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_UP, BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_UP
GSE2585_AIRE_KO_VS_WT_CD80_LOW_MTEC_DN, GSE2585_AIRE_KO_VS_WT_CD80_LOW_MTEC_DN
GO_POSITIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_LIPID_METABOLIC_PROCESS
LEE_METASTASIS_AND_ALTERNATIVE_SPLICING_UP, LEE_METASTASIS_AND_ALTERNATIVE_SPLICING_UP
GSE22611_NOD2_TRANSD_VS_CTRL_TRANSD_HEK293_MDP_STIM_6H_DN, GSE22611_NOD2_TRANSD_VS_CTRL_TRANSD_HEK293_MDP_STIM_6H_DN
GAURNIER_PSMD4_TARGETS, GAURNIER_PSMD4_TARGETS
MYAATNNNNNNNGGC_UNKNOWN, MYAATNNNNNNNGGC_UNKNOWN
HALLMARK_IL6_JAK_STAT3_SIGNALING, HALLMARK_IL6_JAK_STAT3_SIGNALING
GSE43955_1H_VS_60H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN, GSE43955_1H_VS_60H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN
GO_REGULATION_OF_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION, GO_REGULATION_OF_BLOOD_VESSEL_ENDOTHELIAL_CELL_MIGRATION
GO_PROTEOGLYCAN_METABOLIC_PROCESS, GO_PROTEOGLYCAN_METABOLIC_PROCESS
GSE25677_MPL_VS_MPL_AND_R848_STIM_BCELL_DN, GSE25677_MPL_VS_MPL_AND_R848_STIM_BCELL_DN
RAMJAUN_APOPTOSIS_BY_TGFB1_VIA_MAPK1_DN, RAMJAUN_APOPTOSIS_BY_TGFB1_VIA_MAPK1_DN
GO_REGULATION_OF_PHAGOCYTOSIS, GO_REGULATION_OF_PHAGOCYTOSIS
GSE12003_MIR223_KO_VS_WT_BM_PROGENITOR_8D_CULTURE_UP, GSE12003_MIR223_KO_VS_WT_BM_PROGENITOR_8D_CULTURE_UP
RADAEVA_RESPONSE_TO_IFNA1_UP, RADAEVA_RESPONSE_TO_IFNA1_UP
PARK_TRETINOIN_RESPONSE_AND_PML_RARA_FUSION, PARK_TRETINOIN_RESPONSE_AND_PML_RARA_FUSION
GO_NUCLEAR_INNER_MEMBRANE, GO_NUCLEAR_INNER_MEMBRANE
PID_ERBB2_ERBB3_PATHWAY, PID_ERBB2_ERBB3_PATHWAY
GO_POSITIVE_REGULATION_OF_COAGULATION, GO_POSITIVE_REGULATION_OF_COAGULATION
GO_B_CELL_HOMEOSTASIS, GO_B_CELL_HOMEOSTASIS
GO_RESPONSE_TO_LAMINAR_FLUID_SHEAR_STRESS, GO_RESPONSE_TO_LAMINAR_FLUID_SHEAR_STRESS
GTCTACC_MIR379, GTCTACC_MIR379
GSE7831_CPG_VS_INFLUENZA_STIM_PDC_1H_DN, GSE7831_CPG_VS_INFLUENZA_STIM_PDC_1H_DN
BIOCARTA_BCELLSURVIVAL_PATHWAY, BIOCARTA_BCELLSURVIVAL_PATHWAY
LEIN_LOCALIZED_TO_DISTAL_AND_PROXIMAL_DENDRITES, LEIN_LOCALIZED_TO_DISTAL_AND_PROXIMAL_DENDRITES
GO_REGULATION_OF_B_CELL_DIFFERENTIATION, GO_REGULATION_OF_B_CELL_DIFFERENTIATION
PID_P75_NTR_PATHWAY, PID_P75_NTR_PATHWAY
GO_TRANSCRIPTION_FACTOR_ACTIVITY_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_BINDING, GO_TRANSCRIPTION_FACTOR_ACTIVITY_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_BINDING
GO_POSITIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION, GO_POSITIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION
GO_CALCIUM_CHANNEL_REGULATOR_ACTIVITY, GO_CALCIUM_CHANNEL_REGULATOR_ACTIVITY
GO_MONOSACCHARIDE_BINDING, GO_MONOSACCHARIDE_BINDING
REACTOME_ANTIGEN_PRESENTATION_FOLDING_ASSEMBLY_AND_PEPTIDE_LOADING_OF_CLASS_I_MHC, REACTOME_ANTIGEN_PRESENTATION_FOLDING_ASSEMBLY_AND_PEPTIDE_LOADING_OF_CLASS_I_MHC
GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH2_GROUP_OF_DONORS_OXYGEN_AS_ACCEPTOR, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH2_GROUP_OF_DONORS_OXYGEN_AS_ACCEPTOR
GO_POSITIVE_REGULATION_OF_MAST_CELL_ACTIVATION, GO_POSITIVE_REGULATION_OF_MAST_CELL_ACTIVATION
GSE7218_UNSTIM_VS_ANTIGEN_STIM_THROUGH_IGM_BCELL_UP, GSE7218_UNSTIM_VS_ANTIGEN_STIM_THROUGH_IGM_BCELL_UP
REACTOME_DOWNREGULATION_OF_ERBB2_ERBB3_SIGNALING, REACTOME_DOWNREGULATION_OF_ERBB2_ERBB3_SIGNALING
POS_HISTAMINE_RESPONSE_NETWORK, POS_HISTAMINE_RESPONSE_NETWORK
GO_DEATH_RECEPTOR_BINDING, GO_DEATH_RECEPTOR_BINDING
GO_POSITIVE_REGULATION_OF_FIBROBLAST_MIGRATION, GO_POSITIVE_REGULATION_OF_FIBROBLAST_MIGRATION
GO_POST_TRANSLATIONAL_PROTEIN_MODIFICATION, GO_POST_TRANSLATIONAL_PROTEIN_MODIFICATION
GO_CILIARY_TRANSITION_ZONE, GO_CILIARY_TRANSITION_ZONE
LANDIS_BREAST_CANCER_PROGRESSION_UP, LANDIS_BREAST_CANCER_PROGRESSION_UP
GO_NEGATIVE_REGULATION_OF_INTERLEUKIN_6_PRODUCTION, GO_NEGATIVE_REGULATION_OF_INTERLEUKIN_6_PRODUCTION
RYTGCNWTGGNR_UNKNOWN, RYTGCNWTGGNR_UNKNOWN
REACTOME_HS_GAG_BIOSYNTHESIS, REACTOME_HS_GAG_BIOSYNTHESIS
GSE45365_HEALTHY_VS_MCMV_INFECTION_CD8_TCELL_IFNAR_KO_UP, GSE45365_HEALTHY_VS_MCMV_INFECTION_CD8_TCELL_IFNAR_KO_UP
GO_REGULATION_OF_FAT_CELL_DIFFERENTIATION, GO_REGULATION_OF_FAT_CELL_DIFFERENTIATION
ST_WNT_BETA_CATENIN_PATHWAY, ST_WNT_BETA_CATENIN_PATHWAY
GO_CHANNEL_INHIBITOR_ACTIVITY, GO_CHANNEL_INHIBITOR_ACTIVITY
OD_CD21HIGH_TRANSITIONAL_BCELL_UP, GSE17186_BLOOD_VS_CORD_BLOOD_CD21HIGH_TRANSITIONAL_BCELL_UP