GSE339_EX_VIVO_VS_IN_CULTURE_CD8POS_DC_UP, GSE339_EX_VIVO_VS_IN_CULTURE_CD8POS_DC_UP GSE7852_THYMUS_VS_FAT_TREG_UP, GSE7852_THYMUS_VS_FAT_TREG_UP GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_UNTREATED_VS_IL12_TREATED_ACT_CD4_TCELL_2H_DN GSE339_EX_VIVO_VS_IN_CULTURE_CD4CD8DN_DC_UP, GSE339_EX_VIVO_VS_IN_CULTURE_CD4CD8DN_DC_UP GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN GSE17721_0.5H_VS_4H_LPS_BMDC_UP, GSE17721_0.5H_VS_4H_LPS_BMDC_UP GSE34392_ST2_KO_VS_WT_DAY8_LCMV_EFFECTOR_CD8_TCELL_UP, GSE34392_ST2_KO_VS_WT_DAY8_LCMV_EFFECTOR_CD8_TCELL_UP GSE17721_CTRL_VS_CPG_0.5H_BMDC_UP, GSE17721_CTRL_VS_CPG_0.5H_BMDC_UP GSE17721_CTRL_VS_PAM3CSK4_6H_BMDC_UP, GSE17721_CTRL_VS_PAM3CSK4_6H_BMDC_UP GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_UP, GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL10_STIM_MACROPHAGE_180MIN_UP GSE37416_CTRL_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_CTRL_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_UP GSE9037_CTRL_VS_LPS_4H_STIM_BMDM_UP, GSE9037_CTRL_VS_LPS_4H_STIM_BMDM_UP GSE17721_LPS_VS_PAM3CSK4_12H_BMDC_DN, GSE17721_LPS_VS_PAM3CSK4_12H_BMDC_DN GSE15330_MEGAKARYOCYTE_ERYTHROID_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_DN, GSE15330_MEGAKARYOCYTE_ERYTHROID_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_DN GSE15139_GMCSF_TREATED_VS_UNTREATED_NEUTROPHILS_UP, GSE15139_GMCSF_TREATED_VS_UNTREATED_NEUTROPHILS_UP GSE5679_RARA_AGONIST_AM580_VS_AM580_AND_ROSIGLITAZONE_TREATED_DC_UP, GSE5679_RARA_AGONIST_AM580_VS_AM580_AND_ROSIGLITAZONE_TREATED_DC_UP GSE44649_NAIVE_VS_ACTIVATED_CD8_TCELL_UP, GSE44649_NAIVE_VS_ACTIVATED_CD8_TCELL_UP GSE21033_3H_VS_24H_POLYIC_STIM_DC_DN, GSE21033_3H_VS_24H_POLYIC_STIM_DC_DN GSE16522 MEMORY VS NAIVE ANTI CD3CD28 STIM CD8 TCELL UP, GSE16522 MEMORY VS NAIVE ANTI CD3CD28 STIM CD8 TCELL UP GSE24726_WT_VS_E2-2_KO_PDC_DN, GSE24726_WT_VS_E2-2_KO_PDC_DN GSE12198 CTRL VS HIGH IL2 STIM NK CELL DN, GSE12198 CTRL VS HIGH IL2 STIM NK CELL DN GSE27092_WT_VS_HDAC7_PHOSPHO_DEFICIENT_CD8_TCELL_DN, GSE27092_WT_VS_HDAC7_PHOSPHO_DEFICIENT_CD8_TCELL_DN GSE37301_RAG2_KO_VS_RAG2_AND_ETS1_KO_NK_CELL_DN, GSE37301_RAG2_KO_VS_RAG2_AND_ETS1_KO_NK_CELL_DN GSE19198_CTRL_VS_IL21_TREATED_TCELL_24H_DN, GSE19198_CTRL_VS_IL21_TREATED_TCELL_24H_DN GSE18893 CTRL VS_TNF_TREATED_TCONV_24H_DN, GSE18893_CTRL_VS_TNF_TREATED_TCONV_24H_DN GSE17721_CPG_VS_GARDIQUIMOD_24H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_24H_BMDC_DN ROSS_AML_WITH_AML1_ETO_FUSION, ROSS_AML_WITH_AML1_ETO_FUSION GSE20715_0H_VS_24H_OZONE_TLR4_KO_LUNG_UP, GSE20715_0H_VS_24H_OZONE_TLR4_KO_LUNG_UP GSE32423_CTRL_VS_IL7_IL4_MEMORY_CD8_TCELL_DN, GSE32423_CTRL_VS_IL7_IL4_MEMORY_CD8_TCELL_DN GSE32255_WT_VS_JMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_UP, GSE32255_WT_VS_JMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_UP GO_STEROID_HORMONE_RECEPTOR_BINDING, GO_STEROID_HORMONE_RECEPTOR_BINDING CUI_GLUCOSE_DEPRIVATION, CUI_GLUCOSE_DEPRIVATION GSE16385_MONOCYTE_VS_MACROPHAGE_UP, GSE16385_MONOCYTE_VS_MACROPHAGE_UP GSE32128_INOS_DEPENDENT_VS_INOS_INDEPENDENT_ACTIVATED_TCELL_DN, GSE32128_INOS_DEPENDENT_VS_INOS_INDEPENDENT_ACTIVATED_TCELL_DN GSE35543_IN_VITRO_ITREG_VS_CONVERTED_EX_ITREG_DN, GSE35543_IN_VITRO_ITREG_VS_CONVERTED_EX_ITREG_DN GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_UP, GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_UP GO_PHOSPHOTRANSFERASE_ACTIVITY_PHOSPHATE_GROUP_AS_ACCEPTOR, GO_PHOSPHOTRANSFERASE_ACTIVITY_PHOSPHATE_GROUP_AS_ACCEPTOR GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE TOMLINS PROSTATE CANCER UP, TOMLINS PROSTATE CANCER UP GO_NUCLEOTIDE_KINASE_ACTIVITY, GO_NUCLEOTIDE_KINASE_ACTIVITY GRABARCZYK_BCL11B_TARGETS_DN, GRABARCZYK_BCL11B_TARGETS_DN GO_PERICENTRIC_HETEROCHROMATIN, GO_PERICENTRIC_HETEROCHROMATIN GSE22886 NAIVE BCELL VS MONOCYTE UP, GSE22886 NAIVE BCELL VS MONOCYTE UP GO_PURINE_DEOXYRIBONUCLEOTIDE_METABOLIC_PROCESS, GO_PURINE_DEOXYRIBONUCLEOTIDE_METABOLIC_PROCESS GO_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS_VIA_THE_MULTIVESICULAR_BODY_SORTING_PATHWAY, GO_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS_VIA_THE_MULTIVESICULAR_ REACTOME_TRAFFICKING_OF_GLUR2_CONTAINING_AMPA_RECEPTORS, REACTOME_TRAFFICKING_OF_GLUR2_CONTAINING_AMPA_RECEPTORS GSE26928_EFF_MEM_VS_CENTR_MEM_CD4_TCELL_UP, GSE26928_EFF_MEM_VS_CENTR_MEM_CD4_TCELL_UP GO FATTY ACID BETA OXIDATION USING ACYL COA OXIDASE, GO FATTY ACID BETA OXIDATION USING ACYL COA OXIDASE GO_MULTIVESICULAR_BODY_ORGANIZATION, GO_MULTIVESICULAR_BODY_ORGANIZATION LABBE_TARGETS_OF_TGFB1_AND_WNT3A_DN, LABBE_TARGETS_OF_TGFB1_AND_WNT3A_DN GO_SH3_SH2_ADAPTOR_ACTIVITY, GO_SH3_SH2_ADAPTOR_ACTIVITY GO_GTP_METABOLIC_PROCESS, GO_GTP_METABOLIC_PROCESS PID_P38_ALPHA_BETA_PATHWAY, PID_P38_ALPHA_BETA_PATHWAY GO_EXOPEPTIDASE_ACTIVITY, GO_EXOPEPTIDASE_ACTIVITY GO_CALCIUM_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GO_CALCIUM_DEPENDENT_PROTEIN_KINASE_ACTIVITY BIOCARTA CCR5 PATHWAY, BIOCARTA CCR5 PATHWAY REACTOME_ENDOSOMAL_SORTING_COMPLEX_REQUIRED_FOR_TRANSPORT_ESCRT, REACTOME_ENDOSOMAL_SORTING_COMPLEX_REQUIRED_FOR_TRANSPORT_ESCRT GO_POLYOL_BIOSYNTHETIC_PROCESS, GO_POLYOL_BIOSYNTHETIC_PROCESS PID_ALK1_PATHWAY, PID_ALK1_PATHWAY GO_ALPHA_LINOLENIC_ACID_METABOLIC_PROCESS, GO_ALPHA_LINOLENIC_ACID_METABOLIC_PROCESS MODULE_530, MODULE_530 BIOCARTA_ETS_PATHWAY, BIOCARTA_ETS_PATHWAY BIOCARTA_GATA3_PATHWAY, BIOCARTA_GATA3_PATHWAY BIOCARTA_ARENRF2_PATHWAY, BIOCARTA_ARENRF2_PATHWAY GO_ENDOTHELIAL_TUBE_MORPHOGENESIS, GO_ENDOTHELIAL_TUBE_MORPHOGENESIS BIOCARTA_CXCR4_PATHWAY, BIOCARTA_CXCR4_PATHWAY

GSE31082 DP_VS_CD4_SP_THYMOCYTE_UP, GSE31082 DP_VS_CD4_SP_THYMOCYTE_UP

ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_2H_UP 🕻