

GSE32986_UNSTIM_VS_CURDLAN_HIGHDOSE_STIM_DC_UP, GSE32986_UNSTIM_VS_CURDLAN_LOWDOSE_STIM_DC_UP, GSE10325_BCELL_VS_LUPUS_BCELL_DN, GSE10325_BCELL_VS_LUPUS_BCELL_DN, GSE13738_RESTING_VS_TCR_ACTIVATED_CD4_TCELL_DN, GSE13738_RESTING_VS_TCR_ACTIVATED_CD4_TCELL_DN, GSE40666_UNTREATED_VS_IFNA_STIM_STAT4_KO_EFFECTOR_CD8_TCELL_90MIN_UP, GSE40666_UNTREATED_VS_IFNA_STIM_STAT4_KO_EFFECTOR_CD8_TCELL_90MIN_UP, GSE36826_NORMAL_VS_STAPH_AUREUS_INF_SKIN_DN, GSE36826_NORMAL_VS_STAPH_AUREUS_INF_SKIN_DN, GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_6H_DN, GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_6H_DN, GSE11057_NAIVE_VS_CENT_MEMORY_CD4_TCELL_DN, GSE11057_NAIVE_VS_CENT_MEMORY_CD4_TCELL_DN, GSE11057_NAIVE_VS_MEMORY_CD4_TCELL_DN, GSE11057_NAIVE_VS_MEMORY_CD4_TCELL_DN, GSE22886_NAIVE_TCELL_VS_NKCELL_DN, GSE22886_NAIVE_TCELL_VS_NKCELL_DN, GSE22140_GERMFREE_VS_SPF_ARTHRITIC_MOUSE_CD4_TCELL_DN, GSE22140_GERMFREE_VS_SPF_ARTHRITIC_MOUSE_CD4_TCELL_DN, GSE2770_TCFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_TCFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_UP, GSE55085_FETAL_BM_VS_ADULT_BM_SP4_THYMIC_IMPLANT_DN, GSE55085_FETAL_BM_VS_ADULT_BM_SP4_THYMIC_IMPLANT_DN, GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_UP, GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_UP, GSE5542_UNTREATED_VS_IFNA_TREATED_EPITHELIAL_CELLS_24H_UP, GSE5542_UNTREATED_VS_IFNA_TREATED_EPITHELIAL_CELLS_24H_UP, GSE4984_LPS_VS_VEHICLE_CTRL_TREATED_DC_DN, GSE4984_LPS_VS_VEHICLE_CTRL_TREATED_DC_DN, GSE22886_NAIVE_BCELL_VS_NEUTROPHIL_DN, GSE22886_NAIVE_BCELL_VS_NEUTROPHIL_DN, GSE24634_TEFF_VS_TCONV_DAY3_IN_CULTURE_DN, GSE24634_TEFF_VS_TCONV_DAY3_IN_CULTURE_DN, GSE7218_UNSTIM_VS_ANTIGEN_STIM_THROUGH_IGG_BCELL_UP, GSE7218_UNSTIM_VS_ANTIGEN_STIM_THROUGH_IGG_BCELL_UP, GSE17721_POLYIC_VS_PAM3CSK4_12H_BMDC_UP, GSE17721_POLYIC_VS_PAM3CSK4_12H_BMDC_UP, GSE29164_UNTREATED_VS_CD8_TCELL_TREATED_MELANOMA_DAY7_UP, GSE29164_UNTREATED_VS_CD8_TCELL_TREATED_MELANOMA_DAY7_UP, GSE37416_0H_VS_12H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_0H_VS_12H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE26495_NAIVE_VS_PDILOW_CD8_TCELL_DN, GSE26495_NAIVE_VS_PDILOW_CD8_TCELL_DN, GSE26495_NAIVE_VS_PDIIHIGH_CD8_TCELL_DN, GSE26495_NAIVE_VS_PDIIHIGH_CD8_TCELL_DN, GSE16385_UNTREATED_VS_12H_IFNG_TNF_TREATED_MACROPHAGE_DN, GSE16385_UNTREATED_VS_12H_IFNG_TNF_TREATED_MACROPHAGE_DN, GSE16755_CTRL_VS_IFNA_TREATED_MAC_DN, GSE16755_CTRL_VS_IFNA_TREATED_MAC_DN, GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_UP, GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_UP, SASAKI_ADULT_T_CELL_LEUKEMIA, SASAKI_ADULT_T_CELL_LEUKEMIA, GSE21927_C26GM_VS_4T1_TUMOR_MONOCYTE_BALBC_UP, GSE21927_C26GM_VS_4T1_TUMOR_MONOCYTE_BALBC_UP, GSE37301_PRO_BCELL_VS GRANULOCYTE_MONOCYTE_PROGENITOR_DN, GSE37301_PRO_BCELL_VS GRANULOCYTE_MONOCYTE_PROGENITOR_DN, GSE44732_UNSTIM_VS_IL27_STIM_IMATURE_DC_DN, GSE44732_UNSTIM_VS_IL27_STIM_IMATURE_DC_DN, GSE22935_WT_VS_MYD88_KO_MACROPHAGE_48H_MBOVIS_BCG_STIM_DN, GSE22935_WT_VS_MYD88_KO_MACROPHAGE_48H_MBOVIS_BCG_STIM_DN, GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_DN, GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_DN, GSE4590_LARGE_PRE_BCELL_VS_VPREB_POS_LARGE_PRE_BCELL_DN, GSE4590_LARGE_PRE_BCELL_VS_VPREB_POS_LARGE_PRE_BCELL_DN, GSE22886_NAIVE_CD8_TCELL_VS_NKCELL_DN, GSE22886_NAIVE_CD8_TCELL_VS_NKCELL_DN, GSE32986_UNSTIM_VS_CURDLAN_HIGHDOSE_STIM_DC_DN, GSE32986_UNSTIM_VS_CURDLAN_HIGHDOSE_STIM_DC_DN, GSE40666_UNTREATED_VS_IFNA_STIM_EFFECTOR_CD8_TCELL_90MIN_UP, GSE40666_UNTREATED_VS_IFNA_STIM_EFFECTOR_CD8_TCELL_90MIN_UP, GSE37301_HEMATOPOIETIC STEM_CELL_VS GRAN_MONO_PROGENITOR_DN, GSE37301_HEMATOPOIETIC STEM_CELL_VS GRAN_MONO_PROGENITOR_DN, GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_24H_UP, GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_24H_UP, YAGI_AML_FAB_MARKERS, YAGI_AML_FAB_MARKERS, GSE17301_CTRL_VS_48H_ACD3_ACD28_STIM_CD8_TCELL_DN, GSE17301_CTRL_VS_48H_ACD3_ACD28_STIM_CD8_TCELL_DN, GSE12366_NAIVE_VS_MEMORY_BCELL_DN, GSE12366_NAIVE_VS_MEMORY_BCELL_DN, GSE46606_IRF4_KO_VS_WT_CD40L_IL2_IL5_3DAY_STIMULATED_BCELL_UP, GSE46606_IRF4_KO_VS_WT_CD40L_IL2_IL5_3DAY_STIMULATED_BCELL_UP, GSE3982_DC_VS_BCELL_UP, GSE3982_DC_VS_BCELL_UP, GSE5589_LPS_VS_LPS_AND_IL6_STIM_IL6_KO_MACROPHAGE_45MIN_UP, GSE5589_LPS_VS_LPS_AND_IL6_STIM_IL6_KO_MACROPHAGE_45MIN_UP, GSE10325_CD4_TCELL_VS_LUPUS_CD4_TCELL_DN, GSE10325_CD4_TCELL_VS_LUPUS_CD4_TCELL_DN, GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_DN, GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_DN, GSE6259_DEC205_POS_DC_VS_CD4_TCELL_UP, GSE6259_DEC205_POS_DC_VS_CD4_TCELL_UP, GSE3982_FOSINOPHIL_VS_EFF_MEMORY_CD4_TCELL_UP, GSE3982_FOSINOPHIL_VS_EFF_MEMORY_CD4_TCELL_UP, GSE6259_33D1_POS_VS_DEC205_POS_SPLENIC_DC_DN, GSE6259_33D1_POS_VS_DEC205_POS_SPLENIC_DC_DN, GSE17721_0.5H_VS_4H_CPG_BMDC_DN, GSE17721_0.5H_VS_4H_CPG_BMDC_DN, GSE6259_CD4_TCELL_VS_CD8_TCELL_DN, GSE6259_CD4_TCELL_VS_CD8_TCELL_DN, GSE7460_CTRL_VS_TGFB_TREATED_ACT_FOXP3_MUT_TCONV_UP, GSE7460_CTRL_VS_TGFB_TREATED_ACT_FOXP3_MUT_TCONV_UP, GO_LAMELLIPODIUM, GO_LAMELLIPODIUM, GSE6259_BCELL_VS_CD8_TCELL_UP, GSE6259_BCELL_VS_CD8_TCELL_UP, GSE17974_IL4_AND_ANTILIL12_VS_UNTREATED_72H_ACT_CD4_TCELL_DN, GSE17974_IL4_AND_ANTILIL12_VS_UNTREATED_72H_ACT_CD4_TCELL_DN, GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_DN, GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_DN, GO_MYELOID_CELL_DIFFERENTIATION, GO_MYELOID_CELL_DIFFERENTIATION, GSE11864_CSF1_VS_CSF1_PAM3CVS_IN_MAC_DN, GSE11864_CSF1_VS_CSF1_PAM3CVS_IN_MAC_DN, GSE40274_FOXP3_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_FOXP3_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GO_HISTONE_BINDING, GO_HISTONE_BINDING, GSE21546_SAPIA_KO_VS_SAPIA_KO_AND_ELK1_KO_ANTILCD3_STIM_DP_THYMOCYTES_UP, GSE21546_SAPIA_KO_VS_SAPIA_KO_AND_ELK1_KO_ANTILCD3_STIM_DP_T, GSE32255_WT_UNSTIM_VS_JMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_DN, GSE32255_WT_UNSTIM_VS_JMJD2D_KNOCKDOWN_4H_LPS_STIM_DC_DN, GSE11057_EFF_MEM_VS_CENT_MEM_CD4_TCELL_UP, GSE11057_EFF_MEM_VS_CENT_MEM_CD4_TCELL_UP, GSE20366_CD103_KLRG1_DP_VS_DN_TREG_DN, GSE20366_CD103_KLRG1_DP_VS_DN_TREG_DN, GSE13522_CTRL_VS_T_CRUZLY_STRAIN_INF_SKIN_IFNAR_KO_DN, GSE13522_CTRL_VS_T_CRUZLY_STRAIN_INF_SKIN_IFNAR_KO_DN, GSE24142_ADULT_VS_FETAL_DN3_THYMOCYTE_DN, GSE24142_ADULT_VS_FETAL_DN3_THYMOCYTE_DN, LANDIS_ERBB2_BREAST_TUMORS_324_UP, LANDIS_ERBB2_BREAST_TUMORS_324_UP, GSE3039_CD4_TCELL_VS_ALPHAALPHA_CD8_TCELL_UP, GSE3039_CD4_TCELL_VS_ALPHAALPHA_CD8_TCELL_UP, GSE3982_DC_VS_BASOPHIL_UP, GSE3982_DC_VS_BASOPHIL_UP, GSE17721_POLYIC_VS_CPG_6H_BMDC_UP, GSE17721_POLYIC_VS_CPG_6H_BMDC_UP, GSE3982_DC_VS_NKCELL_UP, GSE3982_DC_VS_NKCELL_UP, GSE14350_IL2RB_KO_VS_WT_TREG_DN, GSE14350_IL2RB_KO_VS_WT_TREG_DN, GSE25087_TREG_VS_TCONV_ADULT_UP, GSE25087_TREG_VS_TCONV_ADULT_UP, GSE18893_CTRL_VS_TNF_TREATED_TREG_24H_DN, GSE18893_CTRL_VS_TNF_TREATED_TREG_24H_DN, GSE2935_UV_INACTIVATED_VS_LIVE_SENDAI_VIRUS_INF_MACROPHAGE_DN, GSE2935_UV_INACTIVATED_VS_LIVE_SENDAI_VIRUS_INF_MACROPHAGE_DN, GSE17301_CTRL_VS_48H_ACD3_ACD28_IFNA2_STIM_CD8_TCELL_DN, GSE17301_CTRL_VS_48H_ACD3_ACD28_IFNA2_STIM_CD8_TCELL_DN, GSE7460_TCONV_VS_TREG_LN_DN, GSE7460_TCONV_VS_TREG_LN_DN, GSE24081_CONTROLLER_VS_PROGRESSOR_HIV_SPECIFIC_CD8_TCELL_DN, GSE24081_CONTROLLER_VS_PROGRESSOR_HIV_SPECIFIC_CD8_TCELL_DN, GSE6092_UNSTIM_VS_IFNG_STIM_ENDOTHELIAL_CELL_DN, GSE6092_UNSTIM_VS_IFNG_STIM_ENDOTHELIAL_CELL_DN, MODULE_261, MODULE_261, GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_2H_DN, GSE15735_CTRL_VS_HDAC_INHIBITOR_TREATED_CD4_TCELL_2H_DN, GSE17721_4_VS_24H_GARDIQUIMOD_BMDC_UP, GSE17721_4_VS_24H_GARDIQUIMOD_BMDC_UP, WOOD_EBV_EBNA1_TARGETS_UP, WOOD_EBV_EBNA1_TARGETS_UP, GSE4590_LARGE_PRE_BCELL_VS_VPREB_POS_LARGE_PRE_BCELL_UP, GSE4590_LARGE_PRE_BCELL_VS_VPREB_POS_LARGE_PRE_BCELL_UP, STEIN_ESRRA_TARGETS_DN, STEIN_ESRRA_TARGETS_DN, GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_MATURE_NEURON_CELL_LINE_DN, GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_MATURE_NEURON_CELL_LINE_DN, GSE8685_IL2_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP, GSE8685_IL2_ACT_IL2_STARVED_VS_IL21_ACT_IL2_STARVED_CD4_TCELL_UP, GSE41867_NAIVE_VS_DAY30_LCMV_CLONEI3_EXHAUSTED_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY30_LCMV_CLONEI3_EXHAUSTED_CD8_TCELL_UP, GSE8868_SPLEEN_VS_INTESTINE_CD11B_POS_CD11C_NEG_DC_DN, GSE8868_SPLEEN_VS_INTESTINE_CD11B_POS_CD11C_NEG_DC_DN, GSE25088_WT_VS_STAT6_KO_MACROPHAGE_ROSIGLITAZONE_AND_IL4_STIM_DN, GSE25088_WT_VS_STAT6_KO_MACROPHAGE_ROSIGLITAZONE_AND_IL4_STIM_DN, GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_DN, GSE32986_GMCSF_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_DN, MODULE_358, MODULE_358, GSE5679_RARA_AGONIST_AM580_VS_AM580_AND_ROSIGLITAZONE_TREATED_DC_UP, GSE5679_RARA_AGONIST_AM580_VS_AM580_AND_ROSIGLITAZONE_TREATED_DC, GSE5679_RARA_AGONIST_AM580_VS_AM580_AND_ROSIGLITAZONE_TREATED_DC, GSE5679_RARA_AGONIST_AM580_VS_AM580_AND_ROSIGLITAZONE_TREATED_DC, REACTOME_DEPOSITION_OF_NEW_CENPA_CONTAINING_NUCLEOSOMES_AT_THE_CENTROMERE, REACTOME_DEPOSITION_OF_NEW_CENPA_CONTAINING_NUCLEOSOMES_AT_THE_CENTROMERE, GERY_CEBP_TARGETS, GERY_CEBP_TARGETS, GSE32034_UNTREATED_VS_ROSIGLITAZONE_TREATED_LY6C_HIGH_MONOCYTE_UP, GSE32034_UNTREATED_VS_ROSIGLITAZONE_TREATED_LY6C_HIGH_MONOCYTE_UP, KEGG_ADIPOCYTOKINE_SIGNALING_PATHWAY, KEGG_ADIPOCYTOKINE_SIGNALING_PATHWAY, GSE37605_TREG_VS_TCONV_C57BL6_FOXP3_FUSION_GFP_DN, GSE37605_TREG_VS_TCONV_C57BL6_FOXP3_FUSION_GFP_DN, GSE29949_CD8_NEG_DC_SPLEEN_VS_DC_BRAIN_DN, GSE29949_CD8_NEG_DC_SPLEEN_VS_DC_BRAIN_DN, GSE26928_NAIVE_VS_EFF_MEMORY_CD4_TCELL_DN, GSE26928_NAIVE_VS_EFF_MEMORY_CD4_TCELL_DN, GO_HOMEOSTASIS_OF_NUMBER_OF_CELLS, GO_HOMEOSTASIS_OF_NUMBER_OF_CELLS, GSE6259_FLT3L_INDUCED_33D1_POS_DC_VS_CD8_TCELL_UP, GSE6259_FLT3L_INDUCED_33D1_POS_DC_VS_CD8_TCELL_UP, REACTOME_RNA_POL_I_TRANSCRIPTION, REACTOME_RNA_POL_I_TRANSCRIPTION, GSE21927_SPLEEN_VS_BONE_MARROW_MONOCYTE_BALBC_DN, GSE21927_SPLEEN_VS_BONE_MARROW_MONOCYTE_BALBC_DN, GSE32255_UNSTIM_VS_4H_LPS_STIM_DC_UP, GSE32255_UNSTIM_VS_4H_LPS_STIM_DC_UP, GSE13522_WT_VS_IFNAR_KO_SKIN_DN, GSE13522_WT_VS_IFNAR_KO_SKIN_DN, GSE2886_IIGA_VS_IGM_MEMORY_BCELL_UP, GSE2886_IIGA_VS_IGM_MEMORY_BCELL_UP, GSE13522_WT_VS_IFNAR_KO_SKING_T_CRUZLY_STRAIN_INF_UP, GSE13522_WT_VS_IFNAR_KO_SKING_T_CRUZLY_STRAIN_INF_UP, GSE11961_MARGINAL_ZONE_BCELL_VS_GERMINAL_CENTER_BCELL_DAY7_UP, GSE11961_MARGINAL_ZONE_BCELL_VS_GERMINAL_CENTER_BCELL_DAY7_UP, GSE36095_WT_VS_HDAC9_KO_TREG_UP, GSE36095_WT_VS_HDAC9_KO_TREG_UP, REACTOME_MEIOTIC_RECOMBINATION, REACTOME_MEIOTIC_RECOMBINATION, GSE9037_WT_VS_IRAK4_KO_LPS_1H_STIM_BMDM_DN, GSE9037_WT_VS_IRAK4_KO_LPS_1H_STIM_BMDM_DN, GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY6_POST_IMMUNIZATION_UP, GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY6_POST_IMMUNIZATION_UP, GSE5589_WT_VS_IL6_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN, GSE5589_WT_VS_IL6_KO_LPS_AND_IL6_STIM_MACROPHAGE_45MIN_DN, NIKOLSKY_BREAST_CANCER_8Q12_Q22_AMPLICON, NIKOLSKY_BREAST_CANCER_8Q12_Q22_AMPLICON, GO_RESPONSE_TO_UV, GO_RESPONSE_TO_UV, GO_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS, GO_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS, COULOUARN_TEMPORAL_TGFB1_SIGNATURE_DN, COULOUARN_TEMPORAL_TGFB1_SIGNATURE_DN, REACTOME_RNA_POL_I_PROMOTER_OPENING, REACTOME_RNA_POL_I_PROMOTER_OPENING, GSE32128_INOS_DEPENDENT_VS_INOS_INDEPENDENT_ACTIVATED_TCELL_DN, GSE32128_INOS_DEPENDENT_VS_INOS_INDEPENDENT_ACTIVATED_TCELL_DN, GSE2826_WT_VS_BTK_KO_BCELL_DN, GSE2826_WT_VS_BTK_KO_BCELL_DN, GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_DN, GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_DN, GO_ZYMOGEN_ACTIVATION, GO_ZYMOGEN_ACTIVATION, GSE20366_TREG_VS_NAIVE_CD4_TCELL_HOMEOSTATIC_CONVERSION_DN, GSE20366_TREG_VS_NAIVE_CD4_TCELL_HOMEOSTATIC_CONVERSION_DN, GSE2706_R848_VS_LPS_8H_STIM_DC_DN, GSE2706_R848_VS_LPS_8H_STIM_DC_DN, BROWNE_HCMV_INFECTION_24HR_DN, BROWNE_HCMV_INFECTION_24HR_DN, GO_KINASE_REGULATOR_ACTIVITY, GO_KINASE_REGULATOR_ACTIVITY, GSE23321_CENTRAL_VS_EFFECTOR_MEMORY_CD8_TCELL_UP, GSE23321_CENTRAL_VS_EFFECTOR_MEMORY_CD8_TCELL_UP, AKT_UP.V1_UP, AKT_UP.V1_UP, FOSTER_TOLERANT_MACROPHAGE_UP, FOSTER_TOLERANT_MACROPHAGE_UP, BURTON_ADIPOGENESIS_PEAK_AT_0HR, BURTON_ADIPOGENESIS_PEAK_AT_0HR, GSE2128_CTRL_VS_MIMETOPE_NEGATIVE_SELECTION_DP_THYMOCYTE_NOD_DN, GSE2128_CTRL_VS_MIMETOPE_NEGATIVE_SELECTION_DP_THYMOCYTE_NOD_DN, GSE17974_IL4_AND_ANTILIL12_VS_UNTREATED_48H_ACT_CD4_TCELL_DN, GSE17974_IL4_AND_ANTILIL12_VS_UNTREATED_48H_ACT_CD4_TCELL_DN, GSE360_DC_VS_MAC_M_TUBERCULOSIS_UP, GSE360_DC_VS_MAC_M_TUBERCULOSIS_UP, GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_UP, GSE3691_CONVENTIONAL_VS_PLASMACYTOID_DC_SPLEEN_UP, GSE21063_WT_VS_NFATC1_KO_3H_ANTILIGM_STIM_BCELL_UP, GSE21063_WT_VS_NFATC1_KO_3H_ANTILIGM_STIM_BCELL_UP, KANNAN_TP53_TARGETS_UP, KANNAN_TP53_TARGETS_UP, GO_CELLULAR_RESPONSE_TO_RADIATION, GO_CELLULAR_RESPONSE_TO_RADIATION, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_YELLOW_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_YELLOW_UP, GSE41867_DAY6_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GSE41867_DAY6_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GO_DNA_PACKAGING_COMPLEX, GO_DNA_PACKAGING_COMPLEX, GO_CHROMATIN_SILENCING_AT_RDNA, GO_CHROMATIN_SILENCING_AT_RDNA, GSE17721_0.5H_VS_4H_LPS_BMDC_DN, GSE17721_0.5H_VS_4H_LPS_BMDC_DN, GSE17721_LPS_VS_CPG_6H_BMDC_UP, GSE17721_LPS_VS_CPG_6H_BMDC_UP, GSE34156_UNTREATED_VS_24H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP, GSE34156_UNTREATED_VS_24H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP, KOBAYASHI_EGFR_SIGNALING_24HR_UP, KOBAYASHI_EGFR_SIGNALING_24HR_UP, KRCTCNNNNMANAGC_UNKNOWN, KRCTCNNNNMANAGC_UNKNOWN, GAVIN_FOXP3_TARGETS_CLUSTER_P4, GAVIN_FOXP3_TARGETS_CLUSTER_P4, ZHAN_MULTIPLE_MYELOMA_CD1_DN, ZHAN_MULTIPLE_MYELOMA_CD1_DN, GSE24574_BCL6_HIGH_TFH_VS_TCONV_CD4_TCELL_DN, GSE24574_BCL6_HIGH_TFH_VS_TCONV_CD4_TCELL_DN, GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_1_DN, GSE7460_CTRL_VS_FOXP3_OVEREXPR_TCONV_1_DN, KERLEY_RESPONSE_TO_CISPLATIN_UP, KERLEY_RESPONSE_TO_CISPLATIN_UP, WAMUNYOKOLI_OVARIAN_CANCER_GRADES_1_2_UP, WAMUNYOKOLI_OVARIAN_CANCER_GRADES_1_2_UP, GSE46242_CTRL_VS_EGR2_DELETED_THI_CD4_TCELL_DN, GSE46242_CTRL_VS_EGR2_DELETED_THI_CD4_TCELL_DN, SESTO_RESPONSE_TO_UV_C8, SESTO_RESPONSE_TO_UV_C8, GSE5455_EX_VIVO_VS_POST_24H_INCUBATION_MONOCYTES_FROM_TUMOR_BEARING_MOUSE_DN, GSE5455_EX_VIVO_VS_POST_24H_INCUBATION_MONOCYTES_FROM_TUMOR_BEARING_MOUSE_DN, GSE17721_CTRL_VS_PAM3CSK4_4H_BMDC_DN, GSE17721_CTRL_VS_PAM3CSK4_4H_BMDC_DN, ACOSTA_PROLIFERATION_INDEPENDENT_MYC_TARGETS_DN, ACOSTA_PROLIFERATION_INDEPENDENT_MYC_TARGETS_DN, GSE20366_TREG_VS_NAIVE_CD4_TCELL_DN, GSE20366_TREG_VS_NAIVE_CD4_TCELL_DN, GSE3982_MEMORY_CD4_TCELL_VS_BCELL_UP, GSE3982_MEMORY_CD4_TCELL_VS_BCELL_UP, GO_XENOPHAGY, GO_XENOPHAGY, REACTOME_CD28_DEPENDENT_PI3K_AKT_SIGNALING, REACTOME_CD28_DEPENDENT_PI3K_AKT_SIGNALING, GO_NEURON_APOPTOTIC_PROCESS, GO_NEURON_APOPTOTIC_PROCESS, chr6p22, chr6p22, SAFFORD_T_LYMPHOCYTE_ANERGY, SAFFORD_T_LYMPHOCYTE_ANERGY, GSE32986_GMCSF_AND_CURDLAN_LOWDOSE_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_DN, GSE32986_GMCSF_AND_CURDLAN_LOWDOSE_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_DN, KEGG_SYSTEMIC_LUPUS_ERYTHEMATOSUS, KEGG_SYSTEMIC_LUPUS_ERYTHEMATOSUS, CYCLIN_D1_UP.V1_UP, CYCLIN_D1_UP.V1_UP, REACTOME_PACKAGING_OF_TELOMERE_ENDS, REACTOME_PACKAGING_OF_TELOMERE_ENDS, GSE34156_UNTREATED_VS_24H_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP, GSE34156_UNTREATED_VS_24H_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP, KEGG_SPHINGOLIPID_METABOLISM, KEGG_SPHINGOLIPID_METABOLISM, GOLDRATH_IMMUNE_MEMORY, GOLDRATH_IMMUNE_MEMORY, GO_NUCLEAR_NUCLEOSOME, GO_NUCLEAR_NUCLEOSOME, GO_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_BY_HOST, GO_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_BY_HOST, BROWNE_HCMV_INFECTION_20HR_DN, BROWNE_HCMV_INFECTION_20HR_DN, BOHN_PRIMARY_IMMUNODEFICIENCY_SYNDROM_DN, BOHN_PRIMARY_IMMUNODEFICIENCY_SYNDROM_DN, WENDT_COHESIN_TARGETS_UP, WENDT_COHESIN_TARGETS_UP, GSE3565_DUSP1_VS_WT_SPLENOCYTES_POST_LPS_INJECTION_DN, GSE3565_DUSP1_VS_WT_SPLENOCYTES_POST_LPS_INJECTION_DN, DARWICHE_PAPILLOMA_RISK_HIGH_DN, DARWICHE_PAPILLOMA_RISK_HIGH_DN, MODULE_291, MODULE_291, GO_PROTEASE_BINDING, GO_PROTEASE_BINDING, ZHU_CMV_24_HR_DN, ZHU_CMV_24_HR_DN, GO_RESPONSE_TO ESTRADIOL, GO_RESPONSE_TO ESTRADIOL, GSE360_L_DONOVANI_VS_T_GONDI_DC_UP, GSE360_L_DONOVANI_VS_T_GONDI_DC_UP, STARK_HYPPOCAMPUS_22Q11_DELETION_UP, STARK_HYPPOCAMPUS_22Q11_DELETION_UP, GO_ALDO_KETO_REDUCTASE_NADP_ACTIVITY, GO_ALDO_KETO_REDUCTASE_NADP_ACTIVITY, TENEDINI_MEGAKARYOCYTE_MARKERS, TENEDINI_MEGAKARYOCYTE_MARKERS, GO_ESTABLISHMENT_OF_ENDOTHELIAL_BARRIER, GO_ESTABLISHMENT_OF_ENDOTHELIAL_BARRIER, GO_MEMBRANE_PROTEIN_PROTEOLYSIS, GO_MEMBRANE_PROTEIN_PROTEOLYSIS, GO_FEMALE_SEX_DIFFERENTIATION, GO_FEMALE_SEX_DIFFERENTIATION, MANTOVANI_NFKB_TARGETS_DN, MANTOVANI_NFKB_TARGETS_DN, KEGG_GLYCEROLIPID_METABOLISM, KEGG_GLYCEROLIPID_METABOLISM, LEE_LIVER_CANCER_ACOX1_UP, LEE_LIVER_CANCER_ACOX1_UP, REACTOME_AMYLOIDS, REACTOME_AMYLOIDS, chr1p33, chr1p33, GSE6566_STRONG_VS_WEAK_DC_STIMULATED_CD4_TCELL_UP, GSE6566_STRONG_VS_WEAK_DC_STIMULATED_CD4_TCELL_UP, HOFFMANN_IMMATURE_TO_MATURE_B_LYMPHOCYTE_UP, HOFFMANN_IMMATURE_TO_MATURE_B_LYMPHOCYTE_UP, GO_REGENERATION, GO_REGENERATION, GO_REGULATION_OF_INTERLEUKIN_1_BETA_PRODUCTION, GO_REGULATION_OF_INTERLEUKIN_1_BETA_PRODUCTION, LINDSTEDT_DENDRITIC_CELL_MATURATION_A, LINDSTEDT_DENDRITIC_CELL_MATURATION_A, GO_POSITIVE_REGULATION_OF_TRANSMEMBRANE_TRANSPORT, GO_POSITIVE_REGULATION_OF_TRANSMEMBRANE_TRANSPORT, NIELSEN_LEIOMYOSARCOMA_UP, NIELSEN_LEIOMYOSARCOMA_UP, GO_LYSOSOMAL_LUMEN, GO_LYSOSOMAL_LUMEN, GO_REGULATION_OF_STRIATED_MUSCLE_CELL_APOPTOTIC_PROCESS, GO_REGULATION_OF_STRIATED_MUSCLE_CELL_APOPTOTIC_PROCESS, GO_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING, GO_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING, GO_REGULATION_OF_SODIUM_ION_TRANSPORT, GO_REGULATION_OF_SODIUM_ION_TRANSPORT, GO_REGULATION_OF_CYTOKINE_SECRETION, GO_REGULATION_OF_CYTOKINE_SECRETION, NIKOLSKY_BREAST_CANCER_8P12_P11_AMPLICON, NIKOLSKY_BREAST_CANCER_8P12_P11_AMPLICON, GO_RESPONSE_TO_PH, GO_RESPONSE_TO_PH, MODULE_241, MODULE_241, GO_PHAGOCYTIC_CUP, GO_PHAGOCYTIC_CUP, SMID_BREAST_CANCER_ERBB2_UP, SMID_BREAST_CANCER_ERBB2_UP