GSE37416_0H_VS_12H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_0H_VS_12H_F_TULARENSIS_LVS_NEUTROPHIL_UP GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_6H_DN, GSE2770_IL4_ACT_VS_ACT_CD4_TCELL_6H_DN GSE3982_BASOPHIL_VS_EFF_MEMORY_CD4_TCELL_UP, GSE3982_BASOPHIL_VS_EFF_MEMORY_CD4_TCELL_UP GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_UP, GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_UP GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_ADULT_UP, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_ADULT_UP GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_LPS_STIM_DN, GSE3720_VD1_VS_VD2_GAMMADELTA_TCELL_WITH_LPS_STIM_DN GSE45365_WT_VS_IFNAR_KO_CD8A_DC_MCMV_INFECTION_UP, GSE45365_WT_VS_IFNAR_KO_CD8A_DC_MCMV_INFECTION_UP GSE22935_WT_VS_MYD88_KO_MACROPHAGE_UP, GSE22935_WT_VS_MYD88_KO_MACROPHAGE_UP GSE2706_LPS_VS_R848_AND_LPS_8H_STIM_DC_UP, GSE2706_LPS_VS_R848_AND_LPS_8H_STIM_DC_UP GSE8621_LPS_STIM_VS_LPS_PRIMED_AND_LPS_STIM_MACROPHAGE_DN, GSE8621_LPS_STIM_VS_LPS_PRIMED_AND_LPS_STIM_MACROPHAGE_DN GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_UP, GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_UP GSE21380_NON_TFH_VS_TFH_CD4_TCELL_DN, GSE21380_NON_TFH_VS_TFH_CD4_TCELL_DN GSE369_PRE_VS_POST_IL6_INJECTION_IFNG_KO_LIVER_UP, GSE369_PRE_VS_POST_IL6_INJECTION_IFNG_KO_LIVER_UP GSE40274_CTRL_VS_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_CTRL_VS_XBP1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TCONV_UP, GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TCONV_UP GSE3982_DC_VS_NEUTROPHIL_DN, GSE3982_DC_VS_NEUTROPHIL_DN KOKKINAKIS_METHIONINE_DEPRIVATION_48HR_UP, KOKKINAKIS_METHIONINE_DEPRIVATION_48HR_UP GSE13547 2H VS 12 H ANTI IGM STIM ZFX KO BCELL UP, GSE13547 2H VS 12 H ANTI IGM STIM ZFX KO BCELL UP GTGTTGA_MIR505, GTGTTGA_MIR505 KYNG DNA DAMAGE DN, KYNG DNA DAMAGE DN LUCAS_HNF4A_TARGETS_UP, LUCAS_HNF4A_TARGETS_UP GSE22935_UNSTIM_VS_24H_MBOVIS_BCG_STIM_MACROPHAGE_DN, GSE22935_UNSTIM_VS_24H_MBOVIS_BCG_STIM_MACROPHAGE_DN BIOCARTA_FAS_PATHWAY, BIOCARTA_FAS_PATHWAY KOKKINAKIS_METHIONINE_DEPRIVATION_96HR_UP, KOKKINAKIS_METHIONINE_DEPRIVATION_96HR_UP GO PROTEIN LOCALIZATION TO VACUOLE, GO PROTEIN LOCALIZATION TO VACUOLE GSE17721_PAM3CSK4_VS_CPG_12H_BMDC_DN, GSE17721_PAM3CSK4_VS_CPG_12H_BMDC_DN PID_RHOA_PATHWAY, PID_RHOA_PATHWAY GSE22886_NAIVE_CD4_TCELL_VS_NEUTROPHIL_DN, GSE22886_NAIVE_CD4_TCELL_VS_NEUTROPHIL_DN _F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_0H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_UP GO_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_STRESS, GO_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_STRESS PID_VEGFR1_2_PATHWAY, PID_VEGFR1_2_PATHWAY ST_ERK1_ERK2_MAPK_PATHWAY, ST_ERK1_ERK2_MAPK_PATHWAY GSE3565_DUSP1_VS_WT_SPLENOCYTES_POST_LPS_INJECTION_DN, GSE3565_DUSP1_VS_WT_SPLENOCYTES_POST_LPS_INJECTION_DN GO_PROTEIN_POLYMERIZATION, GO_PROTEIN_POLYMERIZATION REACTOME_SEMAPHORIN_INTERACTIONS, REACTOME_SEMAPHORIN_INTERACTIONS WU_HBX_TARGETS_1_DN, WU_HBX_TARGETS_1_DN SASSON_RESPONSE_TO_GONADOTROPHINS_DN, SASSON_RESPONSE_TO_GONADOTROPHINS_DN GO_REGULATION_OF_CHROMATIN_BINDING, GO_REGULATION_OF_CHROMATIN_BINDING SA_B_CELL_RECEPTOR_COMPLEXES, SA_B_CELL_RECEPTOR_COMPLEXES PID_TRAIL_PATHWAY, PID_TRAIL_PATHWAY GO_NEGATIVE_REGULATION_OF_MAP_KINASE_ACTIVITY, GO_NEGATIVE_REGULATION_OF_MAP_KINASE_ACTIVITY GO_MULTICELLULAR_ORGANISM_GROWTH, GO_MULTICELLULAR_ORGANISM_GROWTH GO_ACTIN_FILAMENT_ORGANIZATION, GO_ACTIN_FILAMENT_ORGANIZATION MORI_EMU_MYC_LYMPHOMA_BY_ONSET_TIME_DN, MORI_EMU_MYC_LYMPHOMA_BY_ONSET_TIME_DN SCHURINGA_STAT5A_TARGETS_DN, SCHURINGA_STAT5A_TARGETS_DN GO_PROTEIN_SECRETION, GO_PROTEIN_SECRETION BIOCARTA_IL12_PATHWAY, BIOCARTA_IL12_PATHWAY ST_P38_MAPK_PATHWAY, ST_P38_MAPK_PATHWAY GSE3982_MAST_CELL_VS_NKCELL_DN, GSE3982_MAST_CELL_VS_NKCELL_DN GO_HEPATICOBILIARY_SYSTEM_DEVELOPMENT, GO_HEPATICOBILIARY_SYSTEM_DEVELOPMENT BASSO_CD40_SIGNALING_DN, BASSO_CD40_SIGNALING_DN GO_AMINE_METABOLIC_PROCESS, GO_AMINE_METABOLIC_PROCESS REACTOME_NOD1_2_SIGNALING_PATHWAY, REACTOME_NOD1_2_SIGNALING_PATHWAY PID_IL6_7_PATHWAY, PID_IL6_7_PATHWAY GSE32034_LY6C_HIGH_VS_LOW_MONOCYTE_UP, GSE32034_LY6C_HIGH_VS_LOW_MONOCYTE_UP VERNOCHET_ADIPOGENESIS, VERNOCHET_ADIPOGENESIS MODULE_139, MODULE_139

GSE37416_0H_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_0H_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_UP GSE37416_12H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_12H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_UP GSE37416_0H_VS_6H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_0H_VS_6H_F_TULARENSIS_LVS_NEUTROPHIL_UP REACTOME_TRAF6_MEDIATED_INDUCTION_OF_NFKB_AND_MAP_KINASES_UPON_TLR7_8_OR_9_ACTIVATION, REACTOME_TRAF6_MEDIATED_INDUCTION_OF_NFKB_AND_MAP_KINASES_UI GO_NEGATIVE_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GO_NEGATIVE_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY REACTOME_MAPK_TARGETS_NUCLEAR_EVENTS_MEDIATED_BY_MAP_KINASES, REACTOME_MAPK_TARGETS_NUCLEAR_EVENTS_MEDIATED_BY_MAP_KINASES REACTOME_CASPASE_MEDIATED_CLEAVAGE_OF_CYTOSKELETAL_PROTEINS, REACTOME_CASPASE_MEDIATED_CLEAVAGE_OF_CYTOSKELETAL_PROTEINS GSE28783_ANTI_MIR33_VS_UNTREATED_ATHEROSCLEROSIS_MACROPHAGE_DN, GSE28783_ANTI_MIR33_VS_UNTREATED_ATHEROSCLEROSIS_MACROPHAGE_DN