GSE17721_LPS_VS_PAM3CSK4_24H_BMDC_DN, GSE17721_LPS_VS_PAM3CSK4_24H_BMDC_DN GSE4590_PRE_BCELL_VS_LARGE_PRE_BCELL_DN, GSE4590_PRE_BCELL_VS_LARGE_PRE_BCELL_DN GSE7509_UNSTIM_VS_FCGRIIB_STIM_MONOCYTE_DN, GSE7509_UNSTIM_VS_FCGRIIB_STIM_MONOCYTE_DN GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_UP, GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_UP GO CELL CYCLE ARREST, GO CELL CYCLE ARREST PID_P53_REGULATION_PATHWAY, PID_P53_REGULATION_PATHWAY GSE7764_NKCELL_VS_SPLENOCYTE_UP, GSE7764_NKCELL_VS_SPLENOCYTE_UP GSE3982_DC_VS_MAC_UP, GSE3982_DC_VS_MAC_UP KEGG_CHRONIC_MYELOID_LEUKEMIA, KEGG_CHRONIC_MYELOID_LEUKEMIA GGCAGTG_MIR3243P, GGCAGTG_MIR3243P GSE7509_UNSTIM_VS_FCGRIIB_STIM_DC_UP, GSE7509_UNSTIM_VS_FCGRIIB_STIM_DC_UP GO_REGULATION_OF_PROTEIN_BINDING, GO_REGULATION_OF_PROTEIN_BINDING KEGG_PANCREATIC_CANCER, KEGG_PANCREATIC_CANCER GO_REGULATION_OF_CYTOKINESIS, GO_REGULATION_OF_CYTOKINESIS GO_PROTEIN_DEACETYLASE_ACTIVITY, GO_PROTEIN_DEACETYLASE_ACTIVITY IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP, GSE7768_OVA_ALONE_VS_OVA_WITH_MPL_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP_ ACTWSNACTNY_UNKNOWN, ACTWSNACTNY_UNKNOWN GO_CELLULAR_CARBOHYDRATE_METABOLIC_PROCESS, GO_CELLULAR_CARBOHYDRATE_METABOLIC_PROCESS PHONG_TNF_RESPONSE_VIA_P38_PARTIAL, PHONG_TNF_RESPONSE_VIA_P38_PARTIAL HUTTMANN_B_CLL_POOR_SURVIVAL_DN, HUTTMANN_B_CLL_POOR_SURVIVAL_DN GO_DEACETYLASE_ACTIVITY, GO_DEACETYLASE_ACTIVITY CRX_NRL_DN.V1_UP, CRX_NRL_DN.V1_UP EGR3_01, EGR3_01 GO PERICENTRIC HETEROCHROMATIN, GO PERICENTRIC HETEROCHROMATIN BOYAULT_LIVER_CANCER_SUBCLASS_G5_DN, BOYAULT_LIVER_CANCER_SUBCLASS_G5_DN GO_HOMOLOGOUS_CHROMOSOME_SEGREGATION, GO_HOMOLOGOUS_CHROMOSOME_SEGREGATION TGASTMAGC_NFE2_01, TGASTMAGC_NFE2_01 BIOCARTA_HDAC_PATHWAY, BIOCARTA_HDAC_PATHWAY ZIRN_TRETINOIN_RESPONSE_WT1_UP, ZIRN_TRETINOIN_RESPONSE_WT1_UP HORTON_SREBF_TARGETS, HORTON_SREBF_TARGETS KIM_MYCL1_AMPLIFICATION_TARGETS_DN, KIM_MYCL1_AMPLIFICATION_TARGETS_DN GO_B_CELL_RECEPTOR_SIGNALING_PATHWAY, GO_B_CELL_RECEPTOR_SIGNALING_PATHWAY REACTOME_TRIGLYCERIDE_BIOSYNTHESIS, REACTOME_TRIGLYCERIDE_BIOSYNTHESIS GO_REGULATION_OF_SPROUTING_ANGIOGENESIS, GO_REGULATION_OF_SPROUTING_ANGIOGENESIS GO_CRANIAL_SKELETAL_SYSTEM_DEVELOPMENT, GO_CRANIAL_SKELETAL_SYSTEM_DEVELOPMENT GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREEN_UP, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_GREEN_UP GO_NEGATIVE_REGULATION_OF_I_KAPPAB_KINASE_NF_KAPPAB_SIGNALING, GO_NEGATIVE_REGULATION_OF_I_KAPPAB_KINASE_NF_KAPPAB_SIGNALING GO_MEMBRANE_DEPOLARIZATION, GO_MEMBRANE_DEPOLARIZATION GO_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION, GO_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION GO_ACTIVATION_OF_GTPASE_ACTIVITY, GO_ACTIVATION_OF_GTPASE_ACTIVITY GO_REGULATION_OF_CARTILAGE_DEVELOPMENT, GO_REGULATION_OF_CARTILAGE_DEVELOPMENT GO_EMBRYONIC_CRANIAL_SKELETON_MORPHOGENESIS, GO_EMBRYONIC_CRANIAL_SKELETON_MORPHOGENESIS MATHEW_FANCONI_ANEMIA_GENES, MATHEW_FANCONI_ANEMIA_GENES GO_REGULATION_OF_EXTRACELLULAR_MATRIX_ORGANIZATION, GO_REGULATION_OF_EXTRACELLULAR_MATRIX_ORGANIZATION GO_REGULATION_OF_CELL_MIGRATION_INVOLVED_IN_SPROUTING_ANGIOGENESIS, GO_REGULATION_OF_CELL_MIGRATION_INVOLVED_IN_SPROUTING_A

PIGF UP.V1 DN, PIGF UP.V1 DN

HOWLIN_CITED1_TARGETS_1_UP, HOWLIN_CITED1_TARGETS_1_UP

GO PROTEIN DEPOLYMERIZATION, GO PROTEIN DEPOLYMERIZATION

RUAN_RESPONSE_TO_TNF_TROGLITAZONE_DN, RUAN_RESPONSE_TO_TNF_TROGLITAZONE_DN

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GSE7768_OVA_WITH_LPS_VS_OVA_WITH_MPL_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP, GSE7768_OVA_WITH_LPS_VS_OVA_WITH_MPL_IMMUNIZED_MO
GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_DN, GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_DN
GSE35825_UNTREATED_VS_IFNG_STIM_MACROPHAGE_DN, GSE35825_UNTREATED_VS_IFNG_STIM_MACROPHAGE_DN
GSE4590 PRE BCELL VS VPREB POS LARGE PRE BCELL DN, GSE4590 PRE BCELL VS VPREB POS LARGE PRE BCELL DN
GSE7218_UNSTIM_VS_ANTIGEN_STIM_THROUGH_IGG_BCELL_UP, GSE7218_UNSTIM_VS_ANTIGEN_STIM_THROUGH_IGG_BCELL_UP
GSE7348_UNSTIM_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP, GSE7348_UNSTIM_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP
GSE7218_IGM_VS_IGG_SIGNAL_THGOUGH_ANTIGEN_BCELL_UP, GSE7218_IGM_VS_IGG_SIGNAL_THGOUGH_ANTIGEN_BCELL_UP
GSE6092_IFNG_VS_IFNG_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_UP, GSE6092_IFNG_VS_IFNG_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_UP
GSE32986_GMCSF_AND_CURDLAN_LOWDOSE_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_UP, GSE32986_GMCSF_AND_CURDLAN_LOWDOSE_VS_GMCSI
GSE17186_CD21LOW_VS_CD21HIGH_TRANSITIONAL_BCELL_DN, GSE17186_CD21LOW_VS_CD21HIGH_TRANSITIONAL_BCELL_DN
GSE7768_OVA_ALONE_VS_OVA_WITH_LPS_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP, GSE7768_OVA_ALONE_VS_OVA_WITH_LPS_IMMUNIZED_MOUSE_WI
GOLDRATH_NAIVE_VS_MEMORY_CD8_TCELL_UP, GOLDRATH_NAIVE_VS_MEMORY_CD8_TCELL_UP
GSE21360_TERTIARY_VS_QUATERNARY_MEMORY_CD8_TCELL_UP, GSE21360_TERTIARY_VS_QUATERNARY_MEMORY_CD8_TCELL_UP
GSE46606_IRF4MID_VS_WT_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_UP, GSE46606_IRF4MID_VS_WT_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_UP
GSE3920_IFNA_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP, GSE3920_IFNA_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP
GSE37301_HEMATOPOIETIC_STEM_CELL_VS_LYMPHOID_PRIMED_MPP_UP, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_LYMPHOID_PRIMED_MPP_UP
GSE17301 CTRL VS 48H ACD3 ACD28 IFNA2 STIM CD8 TCELL DN, GSE17301 CTRL VS 48H ACD3 ACD28 IFNA2 STIM CD8 TCELL DN
GSE6259_FLT3L_INDUCED_VS_WT_SPLENIC_DC_33D1_POS_UP, GSE6259_FLT3L_INDUCED_VS_WT_SPLENIC_DC_33D1_POS_UP
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
GSE21360_SECONDARY_VS_TERTIARY_MEMORY_CD8_TCELL_DN, GSE21360_SECONDARY_VS_TERTIARY_MEMORY_CD8_TCELL_DN
 GSE20500_CTRL_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP, GSE20500_CTRL_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP
 GSE22601_IMMATURE_CD4_SINGLE_POSITIVE_VS_CD4_SINGLE_POSITIVE_THYMOCYTE_DN, GSE22601_IMMATURE_CD4_SINGLE_POSITIVE_VS_CD4_SINGLE_POS
 GSE12003_4D_VS_8D_CULTURE_MIR223_KO_BM_PROGENITOR_DN, GSE12003_4D_VS_8D_CULTURE_MIR223_KO_BM_PROGENITOR_DN
 GSE37301_MULTIPOTENT_PROGENITOR_VS_CD4_TCELL_UP, GSE37301_MULTIPOTENT_PROGENITOR_VS_CD4_TCELL_UP
GSE2128_CTRL_VS_MIMETOPE_NEGATIVE_SELECTION_DP_THYMOCYTE_NOD_DN, GSE2128_CTRL_VS_MIMETOPE_NEGATIVE_SELECTION_DP_THYMOCYTE_NO
GSE21063_3H_VS_16H_ANTI_IGM_STIM_NFATC1_KOBCELL_UP, GSE21063_3H_VS_16H_ANTI_IGM_STIM_NFATC1_KOBCELL_UP
GSE29164_DAY3_VS_DAY7_CD8_TCELL_TREATED_MELANOMA_UP, GSE29164_DAY3_VS_DAY7_CD8_TCELL_TREATED_MELANOMA_UP
GO_NUCLEOBASE_CONTAINING_COMPOUND_KINASE_ACTIVITY, GO_NUCLEOBASE_CONTAINING_COMPOUND_KINASE_ACTIVITY
GSE37301_MULTIPOTENT_PROGENITOR_VS_PRO_BCELL_UP, GSE37301_MULTIPOTENT_PROGENITOR_VS_PRO_BCELL_UP
 GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP
 GO_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GO_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY
 GSE41176_UNSTIM_VS_ANTI_IGM_STIM_TAK1_KO_BCELL_24H_DN, GSE41176_UNSTIM_VS_ANTI_IGM_STIM_TAK1_KO_BCELL_24H_DN
GO_FATTY_ACID_BETA_OXIDATION_USING_ACYL_COA_OXIDASE, GO_FATTY_ACID_BETA_OXIDATION_USING_ACYL_COA_OXIDASE
GO_ACTIVATION_OF_ANAPHASE_PROMOTING_COMPLEX_ACTIVITY, GO_ACTIVATION_OF_ANAPHASE_PROMOTING_COMPLEX_ACTIVITY
GO_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVELOPMENT, GO_EPITHELIAL_CELL_DIFFERENTIATION_INVOLVED_IN_KIDNEY_DEVEL
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