OD_CD21HIGH_TRANSITIONAL_BCELL_UP, GSE17186_BLOOD_VS_CORD_BLOOD_CD21HIGH_TRANSITIONAL_BCELL_UP

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GSE36888 STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_DN, GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_DN
GSE22601_CD4_SINGLE_POSITIVE_VS_CD8_SINGLE_POSITIVE_THYMOCYTE_UP, GSE22601_CD4_SINGLE_POSITIVE_VS_CD8_SINGLE_POSITIVE_THYMOCYTE_UP
GSE17186_BLOOD_VS_CORD_BLOOD_CD21LOW_TRANSITIONAL_BCELL_DN, GSE17186_BLOOD_VS_CORD_BLOOD_CD21LOW_TRANSITIONAL_BCELL_DN
GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_20MIN_DN, GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_20MIN_DN
HOLLMANN_APOPTOSIS_VIA_CD40_UP, HOLLMANN_APOPTOSIS_VIA_CD40_UP
GSE12392_WT_VS_IFNB_KO_CD8A_NEG_SPLEEN_DC_UP, GSE12392_WT_VS_IFNB_KO_CD8A_NEG_SPLEEN_DC_UP
GSE4748_CTRL_VS_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_DN, GSE4748_CTRL_VS_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_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
GSE42724_NAIVE_VS_MEMORY_BCELL_DN, GSE42724_NAIVE_VS_MEMORY_BCELL_DN
GO_VACUOLE_ORGANIZATION, GO_VACUOLE_ORGANIZATION
GSE40666_WT_VS_STAT1_KO_CD8_TCELL_DN, GSE40666_WT_VS_STAT1_KO_CD8_TCELL_DN
GSE27859_MACROPHAGE_VS_CD11C_INT_F480_HI_MACROPHAGE_UP, GSE27859_MACROPHAGE_VS_CD11C_INT_F480_HI_MACROPHAGE_UP
GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_DN, GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_DN
GSE32986_CURDLAN_HIGHDOSE_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_DN, GSE32986_CURDLAN_HIGHDOSE_VS_GMCSF_AND_CURDLAN_HIGHDOSE_STIM_DC_DN
GSE5589_LPS_AND_IL10_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_DN, GSE5589_LPS_AND_IL10_VS_LPS_AND_IL6_STIM_IL10_KO_MACROPHAGE_45MIN_DN
GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_DN, GSE25088_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_STAT6_KO_MACROPHAGE_DAY10_DN
GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_DN, GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_DN
GSE37301_MULTIPOTENT_PROGENITOR_VS_GRAN_MONO_PROGENITOR_DN, GSE37301_MULTIPOTENT_PROGENITOR_VS_GRAN_MONO_PROGENITOR_DN
GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP, GSE16385_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP
PID_P53_REGULATION_PATHWAY, PID_P53_REGULATION_PATHWAY
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
LIU_VMYB_TARGETS_UP, LIU_VMYB_TARGETS_UP
GSE17721_CTRL_VS_POLYIC_8H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_8H_BMDC_DN
GSE11961_FOLLICULAR_BCELL_VS_MEMORY_BCELL_DAY40_UP, GSE11961_FOLLICULAR_BCELL_VS_MEMORY_BCELL_DAY40_UP
PID_P75_NTR_PATHWAY, PID_P75_NTR_PATHWAY
TCGA_GLIOBLASTOMA_COPY_NUMBER_UP, TCGA_GLIOBLASTOMA_COPY_NUMBER_UP
GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GFP_TCONV_UP, GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GFP_TCONV_UP
PID_ERBB2_ERBB3_PATHWAY, PID_ERBB2_ERBB3_PATHWAY
GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR
GO_POSITIVE_REGULATION_OF_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE, GO_POSITIVE_REGULATION_OF_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALIN
GO_AMINOGLYCAN_BIOSYNTHETIC_PROCESS, GO_AMINOGLYCAN_BIOSYNTHETIC_PROCESS
GSE22611_NOD2_TRANSD_VS_CTRL_TRANSD_HEK293_MDP_STIM_6H_DN, GSE22611_NOD2_TRANSD_VS_CTRL_TRANSD_HEK293_MDP_STIM_6H_DN
 AMIT_SERUM_RESPONSE_240_MCF10A, AMIT_SERUM_RESPONSE_240_MCF10A
WEIGEL_OXIDATIVE_STRESS_RESPONSE, WEIGEL_OXIDATIVE_STRESS_RESPONSE
ST_P38_MAPK_PATHWAY, ST_P38_MAPK_PATHWAY
GO_NEGATIVE_REGULATION_OF_AXONOGENESIS, GO_NEGATIVE_REGULATION_OF_AXONOGENESIS
ZHAN_MULTIPLE_MYELOMA_LB_DN, ZHAN_MULTIPLE_MYELOMA_LB_DN
ST_GA13_PATHWAY, ST_GA13_PATHWAY
GO_CELLULAR_RESPONSE_TO_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_STIMULUS, GO_CELLULAR_RESPONSE_TO_VASCULAR_ENDOTHELIAL_GROWTH_FACTOR_STIMULU
GO_CELLULAR_CARBOHYDRATE_BIOSYNTHETIC_PROCESS, GO_CELLULAR_CARBOHYDRATE_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_FAT_CELL_DIFFERENTIATION, GO_REGULATION_OF_FAT_CELL_DIFFERENTIATION
BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_DN, BREDEMEYER_RAG_SIGNALING_NOT_VIA_ATM_DN
 MEISSNER_NPC_ICP_WITH_H3K4ME3, MEISSNER_NPC_ICP_WITH_H3K4ME3
GO_LIPOPOLYSACCHARIDE_MEDIATED_SIGNALING_PATHWAY, GO_LIPOPOLYSACCHARIDE_MEDIATED_SIGNALING_PATHWAY
GSE40274 SATB1_VS_FOXP3_AND_SATB1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_SATB1_VS_FOXP3_AND_SATB1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GO OXIDOREDUCTASE ACTIVITY ACTING ON THE CH NH2 GROUP OF DONORS, GO OXIDOREDUCTASE ACTIVITY ACTING ON THE CH NH2 GROUP OF DONORS
GO_NEGATIVE_REGULATION_OF_STEM_CELL_PROLIFERATION, GO_NEGATIVE_REGULATION_OF_STEM_CELL_PROLIFERATION
GO ENDOPLASMIC RETICULUM LUMEN, GO ENDOPLASMIC RETICULUM LUMEN
GO OXIDOREDUCTASE ACTIVITY ACTING ON THE CH NH2 GROUP OF DONORS OXYGEN AS ACCEPTOR, GO OXIDOREDUCTASE ACTIVITY ACTING ON THE CH NH2 GROU
REACTOME HEPARAN SULFATE HEPARIN HS GAG METABOLISM, REACTOME HEPARAN SULFATE HEPARIN HS GAG METABOLISM
GO PHOTORECEPTOR CONNECTING CILIUM, GO PHOTORECEPTOR CONNECTING CILIUM
MEISSNER_ES_ICP_WITH_H3K4ME3, MEISSNER_ES_ICP_WITH_H3K4ME3
GO SEMAPHORIN RECEPTOR BINDING, GO SEMAPHORIN RECEPTOR BINDING
GENTLES_LEUKEMIC_STEM_CELL_DN, GENTLES_LEUKEMIC_STEM_CELL_DN
GO_POST_TRANSLATIONAL_PROTEIN_MODIFICATION, GO_POST_TRANSLATIONAL_PROTEIN_MODIFICATION
RADAEVA_RESPONSE_TO_IFNA1_UP, RADAEVA_RESPONSE_TO_IFNA1_UP
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