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GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_UP
GSE10239 NAIVE VS MEMORY CD8 TCELL UP, GSE10239 NAIVE VS MEMORY CD8 TCELL UP
GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_48H_DN, GSE2770_UNTREATED_VS_IL4_TREATED_ACT_CD4_TCELL_48H_DN
GSE10239 MEMORY VS_KLRG1INT_EFF_CD8_TCELL_DN, GSE10239 MEMORY_VS_KLRG1INT_EFF_CD8_TCELL_DN
GSE411_WT_VS_SOCS3_KO_MACROPHAGE_UP, GSE411_WT_VS_SOCS3_KO_MACROPHAGE_UP
GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN
GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_DN, GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_DN
GSE7831_UNSTIM_VS_CPG_STIM_PDC_4H_UP, GSE7831_UNSTIM_VS_CPG_STIM_PDC_4H_UP
GSE19772_HCMV_INFL_VS_HCMV_INF_MONOCYTES_AND_PI3K_INHIBITION_DN, GSE19772_HCMV_INFL_VS_HCMV_INF_MONOCYTES_AND_PI3K_INHIBITION_DN
GSE14699_DELETIONAL_TOLERANCE_VS_ACTIVATED_CD8_TCELL_UP, GSE14699_DELETIONAL_TOLERANCE_VS_ACTIVATED_CD8_TCELL_UP
GSE19923 E2A KO VS E2A AND HEB KO DP THYMOCYTE UP, GSE19923 E2A KO VS E2A AND HEB KO DP THYMOCYTE UP
GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_DN, GSE1925_CTRL_VS_IFNG_PRIMED_MACROPHAGE_24H_IFNG_STIM_DN
GSE8835 HEALTHY VS CLL CD8 TCELL UP, GSE8835 HEALTHY VS CLL CD8 TCELL UP
GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_IL12_CD8_TCELL_DN, GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_IL12_CD8_TCELL_DN
GSE3982 BASOPHIL VS TH1_UP, GSE3982 BASOPHIL VS TH1_UP
GSE6259 BCELL VS CD4 TCELL DN, GSE6259 BCELL VS CD4 TCELL DN
GSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_DN, GSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_DN
GSE21379_WT_VS_SAP_KO_CD4_TCELL_DN, GSE21379_WT_VS_SAP_KO_CD4_TCELL_DN
GSE7460_TREG_VS_TCONV_ACT_UP, GSE7460_TREG_VS_TCONV_ACT_UP
GSE27670_CTRL_VS_LMP1_TRANSDUCED_GC_BCELL_UP, GSE27670_CTRL_VS_LMP1_TRANSDUCED_GC_BCELL_UP
GSE21670_STAT3_KO_VS_WT_CD4_TCELL_IL6_TREATED_UP, GSE21670_STAT3_KO_VS_WT_CD4_TCELL_IL6_TREATED_UP
HALLMARK_HEME_METABOLISM, HALLMARK_HEME_METABOLISM
GSE45365_HEALTHY_VS_MCMV_INFECTION_CD8A_DC_DN, GSE45365_HEALTHY_VS_MCMV_INFECTION_CD8A_DC_DN
GO PHOSPHATIDYLINOSITOL BINDING, GO PHOSPHATIDYLINOSITOL BINDING
GSE14386_UNTREATED_VS_IFNA_TREATED_ACT_PBMC_MS_PATIENT_DN, GSE14386_UNTREATED_VS_IFNA_TREATED_ACT_PBMC_MS_PATIENT_DN
GSE20366_CD103_KLRG1_DP_VS_DN_TREG_DN, GSE20366_CD103_KLRG1_DP_VS_DN_TREG_DN
GSE37605_TREG_VS_TCONV_C57BL6_FOXP3_FUSION_GFP_DN, GSE37605_TREG_VS_TCONV_C57BL6_FOXP3_FUSION_GFP_DN
GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TREG_DN, GSE37532_WT_VS_PPARG_KO_VISCERAL_ADIPOSE_TISSUE_TREG_DN
GO_PROTEIN_TETRAMERIZATION, GO_PROTEIN_TETRAMERIZATION
GSE26669_CD4_VS_CD8_TCELL_IN_MLR_COSTIM_BLOCK_UP, GSE26669_CD4_VS_CD8_TCELL_IN_MLR_COSTIM_BLOCK_UP
GSE30153_LUPUS_VS_HEALTHY_DONOR_BCELL_UP, GSE30153_LUPUS_VS_HEALTHY_DONOR_BCELL_UP
GSE7348_UNSTIM_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_DN, GSE7348_UNSTIM_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_DN
GSE16450_CTRL_VS_IFNA_6H_STIM_IMMATURE_NEURON_CELL_LINE_UP, GSE16450_CTRL_VS_IFNA_6H_STIM_IMMATURE_NEURON_CELL_LINE_UP
GSE23114_WT_VS_SLE2C1_MOUSE_PERITONEAL_CAVITY_B1A_BCELL_UP, GSE23114_WT_VS_SLE2C1_MOUSE_PERITONEAL_CAVITY_B1A_BCELL_UP
 GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_ADULT_DN, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_ADULT_DN
LINDGREN_BLADDER_CANCER_CLUSTER_2A_DN, LINDGREN_BLADDER_CANCER_CLUSTER_2A_DN
GSE32034_LY6C_HIGH_VS_LOW_ROSIGLIZATONE_TREATED_MONOCYTE_DN, GSE32034_LY6C_HIGH_VS_LOW_ROSIGLIZATONE_TREATED_MONOCYTE_DN
GO_INTERCELLULAR_BRIDGE, GO_INTERCELLULAR_BRIDGE
SESTO_RESPONSE_TO_UV_C8, SESTO_RESPONSE_TO_UV_C8
GO_REGULATION_OF_TOR_SIGNALING, GO_REGULATION_OF_TOR_SIGNALING
GO ESTABLISHMENT OR MAINTENANCE OF CELL POLARITY, GO ESTABLISHMENT OR MAINTENANCE OF CELL POLARITY
GO GOLGI TO VACUOLE TRANSPORT, GO GOLGI TO VACUOLE TRANSPORT
GSE7831_1H_VS_4H_INFLUENZA_STIM_PDC_DN, GSE7831_1H_VS_4H_INFLUENZA_STIM_PDC_DN
GSE22886_TH1_VS_TH2_12H_ACT_DN, GSE22886_TH1_VS_TH2_12H_ACT_DN
GSE37301_MULTIPOTENT_PROGENITOR_VS_GRAN_MONO_PROGENITOR_UP, GSE37301_MULTIPOTENT_PROGENITOR_VS_GRAN_MONO_PROGENITOR_UP
GO_CELLULAR_MODIFIED_AMINO_ACID_BIOSYNTHETIC_PROCESS, GO_CELLULAR_MODIFIED_AMINO_ACID_BIOSYNTHETIC_PROCESS
GO_TBP_CLASS_PROTEIN_BINDING, GO_TBP_CLASS_PROTEIN_BINDING
GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_IRF8_KO_DN, GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_IRF8_KO_DN
 GSE27786 NKCELL VS MONO MAC DN, GSE27786 NKCELL VS MONO MAC DN
GO_NUCLEOSIDE_SALVAGE, GO_NUCLEOSIDE_SALVAGE
GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN, GSE26488_HDAC7_KO_VS_VP16_TRANSGENIC_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN
 GAZDA DIAMOND BLACKFAN ANEMIA PROGENITOR UP, GAZDA DIAMOND BLACKFAN ANEMIA PROGENITOR UP
GSE14026_TH1_VS_TH17_DN, GSE14026_TH1_VS_TH17_DN
 GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_MONOCYTE_UP, GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_MONOCYTE_UP
GO_ESTABLISHMENT_OF_CELL_POLARITY, GO_ESTABLISHMENT_OF_CELL_POLARITY
chr2q33, chr2q33
GO_POSITIVE_REGULATION_OF_TOR_SIGNALING, GO_POSITIVE_REGULATION_OF_TOR_SIGNALING
BOHN_PRIMARY_IMMUNODEFICIENCY_SYNDROM_DN, BOHN_PRIMARY_IMMUNODEFICIENCY_SYNDROM_DN
GSE21927_SPLENIC_C26GM_TUMOROUS_VS_4T1_TUMOR_MONOCYTES_UP, GSE21927_SPLENIC_C26GM_TUMOROUS_VS_4T1_TUMOR_MONOCYTES_UP
ST_JNK_MAPK_PATHWAY, ST_JNK_MAPK_PATHWAY
PETRETTO_HEART_MASS_QTL_CIS_UP, PETRETTO_HEART_MASS_QTL_CIS_UP
GO_NUCLEAR_INNER_MEMBRANE, GO_NUCLEAR_INNER_MEMBRANE
GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_UP, GSE32901_TH1_VS_TH17_NEG_CD4_TCELL_UP
GO_NUCLEAR_ENVELOPE_REASSEMBLY, GO_NUCLEAR_ENVELOPE_REASSEMBLY
GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_A_SULFUR_GROUP_OF_DONORS, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_A_SULFUR_GROUP_OF_DONORS
GENTILE_UV_RESPONSE_CLUSTER_D2, GENTILE_UV_RESPONSE_CLUSTER_D2
GSE25123_CTRL_VS_IL4_STIM_PPARG_KO_MACROPHAGE_DN, GSE25123_CTRL_VS_IL4_STIM_PPARG_KO_MACROPHAGE_DN
REACTOME_LYSOSOME_VESICLE_BIOGENESIS, REACTOME_LYSOSOME_VESICLE_BIOGENESIS
GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_UP, GSE7460_TREG_VS_TCONV_ACT_WITH_TGFB_UP
GO_CYTOKINETIC_PROCESS, GO_CYTOKINETIC_PROCESS
GAVIN_FOXP3_TARGETS_CLUSTER_P4, GAVIN_FOXP3_TARGETS_CLUSTER_P4
GO_GOLGI_TO_ENDOSOME_TRANSPORT, GO_GOLGI_TO_ENDOSOME_TRANSPORT
BIOCARTA_P38MAPK_PATHWAY, BIOCARTA_P38MAPK_PATHWAY
ACTACCT_MIR196A_MIR196B, ACTACCT_MIR196A_MIR196B
NADLER_HYPERGLYCEMIA_AT_OBESITY, NADLER_HYPERGLYCEMIA_AT_OBESITY
REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATION_MEDIATED_BY_ACTIVATED_HUMAN_TAK1, REACTOME_JNK_C_JUN_KINASES_PHOSPHORYLATION_AND_ACTIVATION_MEDIA
ROME_INSULIN_TARGETS_IN_MUSCLE_DN, ROME_INSULIN_TARGETS_IN_MUSCLE_DN
BRUECKNER_TARGETS_OF_MIRLET7A3_UP, BRUECKNER_TARGETS_OF_MIRLET7A3_UP
chr8q23, chr8q23
chr7p15, chr7p15
GUO HEX TARGETS_DN, GUO_HEX_TARGETS_DN
GO_QUINONE_METABOLIC_PROCESS, GO_QUINONE_METABOLIC_PROCESS
YANG_BREAST_CANCER_ESR1_DN, YANG_BREAST_CANCER_ESR1_DN
chr6q25, chr6q25
RAY TARGETS OF P210 BCR ABL FUSION UP, RAY TARGETS OF P210 BCR ABL FUSION UP
GO REGULATION OF DENDRITE DEVELOPMENT, GO REGULATION OF DENDRITE DEVELOPMENT
GSE40184_HEALTHY_VS_HCV_INFECTED_DONOR_PBMC_UP, GSE40184_HEALTHY_VS_HCV_INFECTED_DONOR_PBMC_UP
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PID_TCR_JNK_PATHWAY, PID_TCR_JNK_PATHWAY

GO_PIGMENT_CELL_DIFFERENTIATION, GO_PIGMENT_CELL_DIFFERENTIATION

KLRG1HIGH_EFF_CD8_TCELL_DN, GSE10239_MEMORY_VS_KLRG1HIGH_EFF_CD8_TCELL_DN