LITAZONE_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_TREATED_MACROPHAGE_DN

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GSE23925 DARK ZONE VS NAIVE BCELL UP, GSE23925 DARK ZONE VS NAIVE BCELL UP
GSE16385_MONOCYTE_VS_12H_IL4_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_IL4_TREATED_MACROPHAGE_DN
GSE17721_POLYIC_VS_PAM3CSK4_1H_BMDC_UP, GSE17721_POLYIC_VS_PAM3CSK4_1H_BMDC_UP
GSE6674_UNSTIM_VS_ANTI_IGM_AND_CPG_STIM_BCELL_UP, GSE6674_UNSTIM_VS_ANTI_IGM_AND_CPG_STIM_BCELL_UP
GSE23925_LIGHT_ZONE_VS_NAIVE_BCELL_DN, GSE23925_LIGHT_ZONE_VS_NAIVE_BCELL_DN
GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_DN, GSE42088_UNINF_VS_LEISHMANIA_INF_DC_4H_DN
GSE18893_TCONV_VS_TREG_24H_TNF_STIM_DN, GSE18893_TCONV_VS_TREG_24H_TNF_STIM_DN
GSE17721_PAM3CSK4_VS_CPG_24H_BMDC_DN, GSE17721_PAM3CSK4_VS_CPG_24H_BMDC_DN
GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP, GSE42021_CD24LO_TREG_VS_CD24LO_TCONV_THYMUS_UP
GSE15733_BM_VS_SPLEEN_MEMORY_CD4_TCELL_DN, GSE15733_BM_VS_SPLEEN_MEMORY_CD4_TCELL_DN
GSE15324_ELF4_KO_VS_WT_ACTIVATED_CD8_TCELL_UP, GSE15324_ELF4_KO_VS_WT_ACTIVATED_CD8_TCELL_UP
GSE15330_HSC_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_DN, GSE15330_HSC_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_DN
GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4_KO_BCELL_DN, GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4_KO_BCELL_DN
 GSE41867_NAIVE_VS_DAY15_LCMV_CONE13_EFFECTOR_CD8_TCELL_DN, GSE41867_NAIVE_VS_DAY15_LCMV_CONE13_EFFECTOR_CD8_TCELL_DN
 GSE36527_CD62L_HIGH_VS_CD62L_LOW_TREG_CD69_NEG_KLRG1_NEG_UP, GSE36527_CD62L_HIGH_VS_CD62L_LOW_TREG_CD69_NEG_KLRG1_NEG_UP
 GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_DN, GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_3H_DN
 GSE7460_FOXP3_MUT_VS_WT_ACT_WITH_TGFB_TCONV_UP, GSE7460_FOXP3_MUT_VS_WT_ACT_WITH_TGFB_TCONV_UP
GSE7568_CTRL_VS_24H_TGFB_TREATED_MACROPHAGES_WITH_IL4_AND_DEXAMETHASONE_UP, GSE7568_UP, GSE7668_UP, GSE7668_UP, GSE7668_UP, GSE7666_UP, GSE7666_UP, GSE7660_UP, GSE766_UP, GSE766_UP, GSE766_UP, GSE766_UP, GS
FIGUEROA_AML_METHYLATION_CLUSTER_6_UP, FIGUEROA_AML_METHYLATION_CLUSTER_6_UP
FIGUEROA_AML_METHYLATION_CLUSTER_3_UP, FIGUEROA_AML_METHYLATION_CLUSTER_3_UP
GSE30153_LUPUS_VS_HEALTHY_DONOR_BCELL_UP, GSE30153_LUPUS_VS_HEALTHY_DONOR_BCELL_UP
GSE26669_CD4_VS_CD8_TCELL_IN_MLR_DN, GSE26669_CD4_VS_CD8_TCELL_IN_MLR_DN
GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_UP, GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_UP
GSE17721_PAM3CSK4_VS_CPG_16H_BMDC_UP, GSE17721_PAM3CSK4_VS_CPG_16H_BMDC_UP
 GSE16450_CTRL_VS_IFNA_6H_STIM_IMMATURE_NEURON_CELL_LINE_UP, GSE16450_CTRL_VS_IFNA_6H_STIM_IMMATURE_NEURON_CELL_LINE_UP
 FIGUEROA_AML_METHYLATION_CLUSTER_7_UP, FIGUEROA_AML_METHYLATION_CLUSTER_7_UP
FIGUEROA_AML_METHYLATION_CLUSTER_4_UP, FIGUEROA_AML_METHYLATION_CLUSTER_4_UP
chr15q15, chr15q15
SARTIPY_BLUNTED_BY_INSULIN_RESISTANCE_UP, SARTIPY_BLUNTED_BY_INSULIN_RESISTANCE_UP
GO_REGULATION_OF_PROTEIN_ACETYLATION, GO_REGULATION_OF_PROTEIN_ACETYLATION
GSE12963_UNINF_VS_ENV_AND_NEF_DEFICIENT_HIV1_INF_CD4_TCELL_UP, GSE12963_UNINF_VS_ENV_AND_NEF_DEFICIENT_HIV1_INF_CD4_TCELL_UP
chr21q21, chr21q21
 GSE37301_HEMATOPOIETIC_STEM_CELL_VS_PRO_BCELL_UP, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_PRO_BCELL_UP
 AHRARNT_01, AHRARNT_01
GO_MAINTENANCE_OF_CELL_NUMBER, GO_MAINTENANCE_OF_CELL_NUMBER
 ELLWOOD_MYC_TARGETS_DN, ELLWOOD_MYC_TARGETS_DN
GO_REGULATION_OF_TRANSFORMING_GROWTH_FACTOR_BETA_PRODUCTION, GO_REGULATION_OF_TRANSFORMING_GROWTH_FACTOR_BETA_PRODUCTION
SHI_SPARC_TARGETS_DN, SHI_SPARC_TARGETS_DN
GO_MIDDLE_EAR_MORPHOGENESIS, GO_MIDDLE_EAR_MORPHOGENESIS
GO_CIS_TRANS_ISOMERASE_ACTIVITY, GO_CIS_TRANS_ISOMERASE_ACTIVITY
GO_THYMOCYTE_AGGREGATION, GO_THYMOCYTE_AGGREGATION
RIZ_ERYTHROID_DIFFERENTIATION_APOBEC2, RIZ_ERYTHROID_DIFFERENTIATION_APOBEC2
CCCACAT_MIR2993P, CCCACAT_MIR2993P
GO_HYDROLASE_ACTIVITY_ACTING_ON_ETHER_BONDS, GO_HYDROLASE_ACTIVITY_ACTING_ON_ETHER_BONDS
GO_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_INVOLVED_IN_ENDOCARDIAL_CUSHION_FORMATION, GO_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_INVOLVED_IN_ENDOCAF
GSE19923_WT_VS_HEB_KO_DP_THYMOCYTE_UP, GSE19923_WT_VS_HEB_KO_DP_THYMOCYTE_UP
GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4MID_SORTED_BCELL_DN, GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4MID_SORTED_BCELL_DN
GO_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY_TRANSPOSING_S_S_BONDS, GO_INTRAMOLECULAR_OXIDOREDUCTASE_ACTIVITY_TRANSPOSING_S_S_BONDS
GO_ENDOTHELIUM_DEVELOPMENT, GO_ENDOTHELIUM_DEVELOPMENT
GO_ORGANELLE_MEMBRANE_FUSION, GO_ORGANELLE_MEMBRANE_FUSION
PARK_APL_PATHOGENESIS_UP, PARK_APL_PATHOGENESIS_UP
BROWNE_HCMV_INFECTION_10HR_UP, BROWNE_HCMV_INFECTION_10HR_UP
GO_FOUR_WAY_JUNCTION_DNA_BINDING, GO_FOUR_WAY_JUNCTION_DNA_BINDING
WILLIAMS_ESR1_TARGETS_DN, WILLIAMS_ESR1_TARGETS_DN
GO_EMBRYONIC_DIGIT_MORPHOGENESIS, GO_EMBRYONIC_DIGIT_MORPHOGENESIS
PID_S1P_META_PATHWAY, PID_S1P_META_PATHWAY
GO_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA, GO_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PRO
GO_REGULATION_OF_CELL_PROLIFERATION_INVOLVED_IN_HEART_MORPHOGENESIS, GO_REGULATION_OF_CELL_PROLIFERATION_INVOLVED_IN_HEART_MORPHOGENESIS
GO_REGULATION_OF_HEART_MORPHOGENESIS, GO_REGULATION_OF_HEART_MORPHOGENESIS
GO_PHOSPHOTRANSFERASE_ACTIVITY_FOR_OTHER_SUBSTITUTED_PHOSPHATE_GROUPS, GO_PHOSPHOTRANSFERASE_ACTIVITY_FOR_OTHER_SUBSTITUTED_PHOSPHATE_GROUPS
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GSE14769_UNSTIM_VS_80MIN_LPS_BMDM_UP, GSE14769_UNSTIM_VS_80MIN_LPS_BMDM_UP