

DEXAMETHASONE_TREATED_THYMOCYTE_UP, GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOCYTE_UP

GSE22601_IMMATURE_CD4_SINGLE_POSITIVE_VS_CD4_SINGLE_POSITIVE_THYMOCYTE_UP, GSE22601_IMMATURE_CD4_SINGLE_POSITIVE_VS_CD4_SINGLE_POSITIVE_THYMOCYTE_UP
GSE15271_CXCR4_POS_VS_NEG_GC_BCELL_DN, GSE15271_CXCR4_POS_VS_NEG_GC_BCELL_DN
GSE12003_MIR223_KO_VS_WT_BM_PROGENITOR_4D_CULTURE_UP, GSE12003_MIR223_KO_VS_WT_BM_PROGENITOR_4D_CULTURE_UP
GSE15330_HSC_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_UP, GSE15330_HSC_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_IKAROS_KO_UP
GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_DN, GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_DN
TOOKER_GEMCITABINE_RESISTANCE_DN, TOOKER_GEMCITABINE_RESISTANCE_DN
GSE5455_HEALTHY_VS_TUMOR_BEARING_MOUSE_SPLEEN_MONOCYTE_DN, GSE5455_HEALTHY_VS_TUMOR_BEARING_MOUSE_SPLEEN_MONOCYTE_DN
GSE17721_CTRL_VS_LPS_12H_BMDC_DN, GSE17721_CTRL_VS_LPS_12H_BMDC_DN
GO_REGULATION_OF_CELL_CYCLE_ARREST, GO_REGULATION_OF_CELL_CYCLE_ARREST
GSE20715_0H_VS_48H_OZONE_TLR4_KO_LUNG_UP, GSE20715_0H_VS_48H_OZONE_TLR4_KO_LUNG_UP
GO_POSITIVE_REGULATION_OF_CELL_CYCLE_ARREST, GO_POSITIVE_REGULATION_OF_CELL_CYCLE_ARREST
GSE3565_CTRL_VS_LPS_INJECTED_DUSP1_KO_SPLENOCYTES_UP, GSE3565_CTRL_VS_LPS_INJECTED_DUSP1_KO_SPLENOCYTES_UP
GSE40666_UNTREATED_VS_IFNA_STIM_EFFECTOR_CD8_TCELL_90MIN_UP, GSE40666_UNTREATED_VS_IFNA_STIM_EFFECTOR_CD8_TCELL_90MIN_UP
GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_IL12_CD8_TCELL_UP, GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_IL12_CD8_TCELL_UP
GO_KINASE_REGULATOR_ACTIVITY, GO_KINASE_REGULATOR_ACTIVITY
E2F1_UP.V1_DN, E2F1_UP.V1_DN
HALLMARK_MYOGENESIS, HALLMARK_MYOGENESIS
GO_ENTRAINMENT_OF_CIRCADIAN_CLOCK, GO_ENTRAINMENT_OF_CIRCADIAN_CLOCK
REACTOME_GROWTH_HORMONE_RECEPTOR_SIGNALING, REACTOME_GROWTH_HORMONE_RECEPTOR_SIGNALING
ISSAEVA_MLL2_TARGETS, ISSAEVA_MLL2_TARGETS
HOFFMANN_PRE_BI_TO_LARGE_PRE_BII_LYMPHOCYTE_DN, HOFFMANN_PRE_BI_TO_LARGE_PRE_BII_LYMPHOCYTE_DN
GO_REGULATION_OF_CALCIUM_ION_TRANSMEMBRANE_TRANSPORT, GO_REGULATION_OF_CALCIUM_ION_TRANSMEMBRANE_TRANSPORT
GO_DEVELOPMENTAL_GROWTH_INVOLVED_IN_MORPHOGENESIS, GO_DEVELOPMENTAL_GROWTH_INVOLVED_IN_MORPHOGENESIS
GO_T_TUBULE, GO_T_TUBULE
MORF_MYST2, MORF_MYST2
ALK_DN.V1_DN, ALK_DN.V1_DN
GSE19941_UNSTIM_VS_LPS_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_DN, GSE19941_UNSTIM_VS_LPS_STIM_IL10_KO_NFKBP50_KO_MACROPHAGE_DN
GSE34156_UNTREATED_VS_24H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP, GSE34156_UNTREATED_VS_24H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP
GO_ENDODERM_DEVELOPMENT, GO_ENDODERM_DEVELOPMENT
GSE19772_HCMV_INFL_VS_HCMV_INF_MONOCYTES_AND_PI3K_INHIBITION_UP, GSE19772_HCMV_INFL_VS_HCMV_INF_MONOCYTES_AND_PI3K_INHIBITION_UP
DELASERNA_MYOD_TARGETS_UP, DELASERNA_MYOD_TARGETS_UP
JAZAG_TGFB1_SIGNALING_VIA_SMAD4_DN, JAZAG_TGFB1_SIGNALING_VIA_SMAD4_DN
GO_CYTOKINE_RECEPTOR_ACTIVITY, GO_CYTOKINE_RECEPTOR_ACTIVITY
GO_HIPPO_SIGNALING, GO_HIPPO_SIGNALING
KEGG_INTESTINAL_IMMUNE_NETWORK_FOR_IGA_PRODUCTION, KEGG_INTESTINAL_IMMUNE_NETWORK_FOR_IGA_PRODUCTION
REACTOME_SIGNALING_BY_HIPPO, REACTOME_SIGNALING_BY_HIPPO
GO_MAMMARY_GLAND_DEVELOPMENT, GO_MAMMARY_GLAND_DEVELOPMENT
GO_NEGATIVE_REGULATION_OF_CELL_PROJECTION_ORGANIZATION, GO_NEGATIVE_REGULATION_OF_CELL_PROJECTION_ORGANIZATION
WEINMANN_ADAPTATION_TO_HYPOXIA_DN, WEINMANN_ADAPTATION_TO_HYPOXIA_DN
GO_POSITIVE_REGULATION_OF_EXTRACELLULAR_MATRIX_ORGANIZATION, GO_POSITIVE_REGULATION_OF_EXTRACELLULAR_MATRIX_ORGANIZATION
GO_CARDIAC_MUSCLE_TISSUE_MORPHOGENESIS, GO_CARDIAC_MUSCLE_TISSUE_MORPHOGENESIS