

IL2RAHIGH_DAY3_EFF_CD8_TCELL_DN, GSE19825_NAIVE_VS_IL2RAHIGH_DAY3_EFF_CD8_TCELL_DN

GSE19825_NAIVE_VS_DAY3_EFF_CD8_TCELL_DN, GSE19825_NAIVE_VS_DAY3_EFF_CD8_TCELL_DN
GSE19825_NAIVE_VS_IL2RALOW_DAY3_EFF_CD8_TCELL_DN, GSE19825_NAIVE_VS_IL2RALOW_DAY3_EFF_CD8_TCELL_DN
GSE37563_WT_VS_CTLA4_KO_CD4_TCELL_D4_POST_IMMUNIZATION_DN, GSE37563_WT_VS_CTLA4_KO_CD4_TCELL_D4_POST_IMMUNIZATION_DN
GSE15624_CTRL_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN, GSE15624_CTRL_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN
GSE45365_WT_VS_IFNAR_KO_CD11B_DC_MCMV_INFECTION_UP, GSE45365_WT_VS_IFNAR_KO_CD11B_DC_MCMV_INFECTION_UP
GSE46242_CTRL_VS_EGR2_DELETED_ANERGIC_TH1_CD4_TCELL_UP, GSE46242_CTRL_VS_EGR2_DELETED_ANERGIC_TH1_CD4_TCELL_UP
GSE8678_IL7R_LOW_VS_HIGH_EFF_CD8_TCELL_UP, GSE8678_IL7R_LOW_VS_HIGH_EFF_CD8_TCELL_UP
GSE7460_CTRL_VS_TGFB_TREATED_ACT_CD8_TCELL_UP, GSE7460_CTRL_VS_TGFB_TREATED_ACT_CD8_TCELL_UP
GSE21670_UNTREATED_VS_IL6_TREATED_CD4_TCELL_UP, GSE21670_UNTREATED_VS_IL6_TREATED_CD4_TCELL_UP
GO_REGULATION_OF_CELL_CYCLE_ARREST, GO_REGULATION_OF_CELL_CYCLE_ARREST
GO_RIBOSOME_BINDING, GO_RIBOSOME_BINDING
GO_HOMEOSTASIS_OF_NUMBER_OF_CELLS, GO_HOMEOSTASIS_OF_NUMBER_OF_CELLS
QUINTENS_EMBRYONIC_BRAIN_RESPONSE_TO_IR, QUINTENS_EMBRYONIC_BRAIN_RESPONSE_TO_IR
REACTOME_FATTY_ACID_TRIACYLGLYCEROL_AND_KETONE_BODY_METABOLISM, REACTOME_FATTY_ACID_TRIACYLGLYCEROL_AND_KETONE_BODY_METABOLISM
GO_REGULATION_OF_FIBROBLAST_PROLIFERATION, GO_REGULATION_OF_FIBROBLAST_PROLIFERATION
GSE18281_PERIMEDULLARY_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_UP, GSE18281_PERIMEDULLARY_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_UP
REACTOME_TRIGLYCERIDE_BIOSYNTHESIS, REACTOME_TRIGLYCERIDE_BIOSYNTHESIS
chr7q22, chr7q22
GO_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION, GO_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION
CHEN_HOXA5_TARGETS_9HR_DN, CHEN_HOXA5_TARGETS_9HR_DN
GO_MYELOID_CELL_DEVELOPMENT, GO_MYELOID_CELL_DEVELOPMENT
GO_POSITIVE_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_MUSCLE_CELL_DIFFERENTIATION
HOFFMANN_SMALL_PRE_BII_TO_IMMATURE_B_LYMPHOCYTE_UP, HOFFMANN_SMALL_PRE_BII_TO_IMMATURE_B_LYMPHOCYTE_UP
GO_TRANS_GOLGI_NETWORK_MEMBRANE, GO_TRANS_GOLGI_NETWORK_MEMBRANE
GO_BONE_CELL_DEVELOPMENT, GO_BONE_CELL_DEVELOPMENT
GO_POSITIVE_REGULATION_OF_STRIATED_MUSCLE_CELL_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_STRIATED_MUSCLE_CELL_DIFFERENTIATION
GO_LYSOPHOSPHOLIPID_ACYLTRANSFERASE_ACTIVITY, GO_LYSOPHOSPHOLIPID_ACYLTRANSFERASE_ACTIVITY
PID_HDAC_CLASSIII_PATHWAY, PID_HDAC_CLASSIII_PATHWAY
GO_ACYLGLYCEROL_O_ACYLTRANSFERASE_ACTIVITY, GO_ACYLGLYCEROL_O_ACYLTRANSFERASE_ACTIVITY
GSE43863_DAY6_EFF_VS_DAY150_MEM_TFH_CD4_TCELL_DN, GSE43863_DAY6_EFF_VS_DAY150_MEM_TFH_CD4_TCELL_DN
GO_REGULATION_OF_STRIATED_MUSCLE_CELL_DIFFERENTIATION, GO_REGULATION_OF_STRIATED_MUSCLE_CELL_DIFFERENTIATION
GO_CYTOPLASMIC_MICROTUBULE, GO_CYTOPLASMIC_MICROTUBULE
GO_MUSCLE_MYOSIN_COMPLEX, GO_MUSCLE_MYOSIN_COMPLEX
GO_MEGAKARYOCYTE_DEVELOPMENT, GO_MEGAKARYOCYTE_DEVELOPMENT
GO_POSITIVE_REGULATION_OF_MYOTUBE_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_MYOTUBE_DIFFERENTIATION
GO_T_CELL_PROLIFERATION, GO_T_CELL_PROLIFERATION
DELASERNA_MYOD_TARGETS_UP, DELASERNA_MYOD_TARGETS_UP
GO_MEGAKARYOCYTE_DIFFERENTIATION, GO_MEGAKARYOCYTE_DIFFERENTIATION
GO_REGULATION_OF_LIPID_KINASE_ACTIVITY, GO_REGULATION_OF_LIPID_KINASE_ACTIVITY
GO_REGULATION_OF_MYOTUBE_DIFFERENTIATION, GO_REGULATION_OF_MYOTUBE_DIFFERENTIATION
PAX3_01, PAX3_01
GO_POSITIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_IMMUNITY, GO_POSITIVE_REGULATION_OF_LEUKOCYTE_MEDIATED_IMMUNITY
GO_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS, GO_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS
GO_N_TERMINAL_PROTEIN_AMINO_ACID_ACETYLTATION, GO_N_TERMINAL_PROTEIN_AMINO_ACID_ACETYLTATION
GSE1566_WT_VS_EZH2_KO_LN_TCELL_DN, GSE1566_WT_VS_EZH2_KO_LN_TCELL_DN
GUTIERREZ_WALDENSTROEMS_MACROGLOBULINEMIA_2, GUTIERREZ_WALDENSTROEMS_MACROGLOBULINEMIA_2
GSE33513_TCF7_KO_VS_HET_EARLY_THYMIC_PROGENITOR_UP, GSE33513_TCF7_KO_VS_HET_EARLY_THYMIC_PROGENITOR_UP
GO_MYOSIN_II_COMPLEX, GO_MYOSIN_II_COMPLEX
GO_GOLGI_RIBBON_FORMATION, GO_GOLGI_RIBBON_FORMATION
GO_N_TERMINAL_PROTEIN_ACETYLTTRANSFERASE_COMPLEX, GO_N_TERMINAL_PROTEIN_ACETYLTTRANSFERASE_COMPLEX