

7_POS_VS_NEG_NK_CELL_UP, GSE23695_CD57_POS_VS_NEG_NK_CELL_UP

GSE22103_UNSTIM_VS_GMCSF_AND_IFNG_STIM_NEUTROPHIL_UP, GSE22103_UNSTIM_VS_GMCSF_AND_IFNG_STIM_NEUTROPHIL_UP
GSE20754_WT_VS_TCF1_KO_MEMORY_CD8_TCELL_DN, GSE20754_WT_VS_TCF1_KO_MEMORY_CD8_TCELL_DN
GSE22935_UNSTIM_VS_12H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_DN, GSE22935_UNSTIM_VS_12H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_DN
GSE34392_ST2_KO_VS_WT_DAY8_LCMV_EFFECTOR_CD8_TCELL_UP, GSE34392_ST2_KO_VS_WT_DAY8_LCMV_EFFECTOR_CD8_TCELL_UP
GSE17721_0.5H_VS_12H_LPS_BMDC_UP, GSE17721_0.5H_VS_12H_LPS_BMDC_UP
GSE8921_3H_VS_24H_TLR1_2_STIM_MONOCYTE_UP, GSE8921_3H_VS_24H_TLR1_2_STIM_MONOCYTE_UP
GSE22589_HEALTHY_VS_SIV_INFECTED_DC_UP, GSE22589_HEALTHY_VS_SIV_INFECTED_DC_UP
GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_MATURE_NEURON_CELL_LINE_UP, GSE16451_CTRL_VS_WEST_EQUINE_ENC_VIRUS_MATURE_NEURON_CELL_LINE_UP
GSE7852_TREG_VS_TCONV_FAT_DN, GSE7852_TREG_VS_TCONV_FAT_DN
GSE13493_DP_VS_CD8POS_THYMOCYTE_DN, GSE13493_DP_VS_CD8POS_THYMOCYTE_DN
GSE40274_FOXP3_VS_FOXP3_AND_XBP1_TRANSDUCECD_ACTIVATED_CD4_TCELL_DN, GSE40274_FOXP3_VS_FOXP3_AND_XBP1_TRANSDUCECD_ACTIVATED_CD4_TCELL_DN
GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN, GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN
GO_REPRESSING_TRANSCRIPTION_FACTOR_BINDING, GO_REPRESSING_TRANSCRIPTION_FACTOR_BINDING
HUTTMANN_B_CLL_POOR_SURVIVAL_DN, HUTTMANN_B_CLL_POOR_SURVIVAL_DN
KEGG_LYSINE_DEGRADATION, KEGG_LYSINE_DEGRADATION
GO_CLATHRIN_ADAPTOR_COMPLEX, GO_CLATHRIN_ADAPTOR_COMPLEX
GO_RNA_POLYMERASE_II_TRANSCRIPTION_COREPRESSOR_ACTIVITY, GO_RNA_POLYMERASE_II_TRANSCRIPTION_COREPRESSOR_ACTIVITY
GSE17721_LPS_VS_PAM3CSK4_2H_BMDC_DN, GSE17721_LPS_VS_PAM3CSK4_2H_BMDC_DN
DUNNE_TARGETS_OF_AML1_MTG8_FUSION_DN, DUNNE_TARGETS_OF_AML1_MTG8_FUSION_DN
IWANAGA_CARCINOGENESIS_BY_KRAS_UP, IWANAGA_CARCINOGENESIS_BY_KRAS_UP
BILANGES_RAPAMYCIN_SENSITIVE_GENES, BILANGES_RAPAMYCIN_SENSITIVE_GENES
GO_NEGATIVE_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GO_NEGATIVE_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY
MODULE_322, MODULE_322
CCANNAGRKGGC_UNKNOWN, CCANNAGRKGGC_UNKNOWN
chr10p14, chr10p14
TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_SUSTAINED_IN_MONOCYTE_UP, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_SUSTAINED_IN_MONOCYTE_UP
CRX_NRL_DN.V1_DN, CRX_NRL_DN.V1_DN
REACTOME_BASIGIN_INTERACTIONS, REACTOME_BASIGIN_INTERACTIONS
GO_AORTA_DEVELOPMENT, GO_AORTA_DEVELOPMENT
GO_MATURATION_OF_LSU_RRNA, GO_MATURATION_OF_LSU_RRNA
GO_REGULATION_OF_CELL_FATE_COMMITMENT, GO_REGULATION_OF_CELL_FATE_COMMITMENT
GO_CARBOHYDRATE_PHOSPHORYLATION, GO_CARBOHYDRATE_PHOSPHORYLATION
chr7q34, chr7q34
GO_CARBOHYDRATE_KINASE_ACTIVITY, GO_CARBOHYDRATE_KINASE_ACTIVITY