

E_VS_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_EFFECTOR_CD8_TCELL_UP

GSE41867_DAY8_VS_DAY15_LCMV_CLONE13_EFFECTOR_CD8_TCELL_DN, GSE41867_DAY8_VS_DAY15_LCMV_CLONE13_EFFECTOR_CD8_TCELL_DN
GSE411_WT_VS_SOCS3_KO_MACROPHAGE_UP, GSE411_WT_VS_SOCS3_KO_MACROPHAGE_UP
GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_T425A_HEXON_INF_UP, GSE36078_WT_VS_IL1R_KO_LUNG_DC_AFTER_AD5_T425A_HEXON_INF_UP
GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_DN, GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_DN
GSE7218_IGM_VS_IGG_SIGNAL_THGOUGH_ANTIGEN_BCELL_UP, GSE7218_IGM_VS_IGG_SIGNAL_THGOUGH_ANTIGEN_BCELL_UP
GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_48H_UP, GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_48H_UP
GSE33424_CD161_INT_VS_NEG_CD8_TCELL_UP, GSE33424_CD161_INT_VS_NEG_CD8_TCELL_UP
GSE36009_WT_VS_NLRP10_KO_DC_UP, GSE36009_WT_VS_NLRP10_KO_DC_UP
GSE12198_CTRL_VS_LOW_IL2_STIM_NK_CELL_DN, GSE12198_CTRL_VS_LOW_IL2_STIM_NK_CELL_DN
GSE3203_HEALTHY_VS_INFLUENZA_INFECTED_LN_BCELL_UP, GSE3203_HEALTHY_VS_INFLUENZA_INFECTED_LN_BCELL_UP
GSE25846_IL10_POS_VS_NEG_CD8_TCELL_DAY7_POST_CORONAVIRUS_BRAIN_DN, GSE25846_IL10_POS_VS_NEG_CD8_TCELL_DAY7_POST_CORONAVIRUS_BRAIN_DN
GSE14308_TH1_VS_NATURAL_TREG_UP, GSE14308_TH1_VS_NATURAL_TREG_UP
MARTORIATI_MDM4_TARGETS_NEUROEPITHELIUM_UP, MARTORIATI_MDM4_TARGETS_NEUROEPITHELIUM_UP
GSE34156_UNTREATED_VS_6H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_DN, GSE34156_UNTREATED_VS_6H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_DN
GSE6092_UNSTIM_VS_IFNG_STIM_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_UP, GSE6092_UNSTIM_VS_IFNG_STIM_AND_B_BURGDORFERI_INF_ENDOTHELIAL_CELL_UP
GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD4_TCELL_UP, GSE26669_CTRL_VS_COSTIM_BLOCK_MLR_CD4_TCELL_UP
GO_MICROBODY, GO_MICROBODY
KEGG_PEROXISOME, KEGG_PEROXISOME
HALLMARK_CHOLESTEROL_HOMEOSTASIS, HALLMARK_CHOLESTEROL_HOMEOSTASIS
GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_6H_UP, GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_6H_UP
LUCAS_HNF4A_TARGETS_UP, LUCAS_HNF4A_TARGETS_UP
GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_2H_ACT_CD4_TCELL_UP, GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_2H_ACT_CD4_TCELL_UP
ZHAN_MULTIPLE_MYELOMA_MS_DN, ZHAN_MULTIPLE_MYELOMA_MS_DN
GO_FATTY_ACID_CATABOLIC_PROCESS, GO_FATTY_ACID_CATABOLIC_PROCESS
GO_NEGATIVE_REGULATION_OF_EPITHELIAL_CELL_PROLIFERATION, GO_NEGATIVE_REGULATION_OF_EPITHELIAL_CELL_PROLIFERATION
GO_MICROBODY_MEMBRANE, GO_MICROBODY_MEMBRANE
GSE16385_MONOCYTE_VS_MACROPHAGE_UP, GSE16385_MONOCYTE_VS_MACROPHAGE_UP
SANA_RESPONSE_TO_IFNG_UP, SANA_RESPONSE_TO_IFNG_UP
GO_ANAPHASE_PROMOTING_COMPLEX, GO_ANAPHASE_PROMOTING_COMPLEX
MILI_PSEUDOPODIA, MILI_PSEUDOPODIA
GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_DN, GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_DN
GO_PEROXISOME_ORGANIZATION, GO_PEROXISOME_ORGANIZATION
SPIRA_SMOKERS_LUNG_CANCER_DN, SPIRA_SMOKERS_LUNG_CANCER_DN
GO_ALCOHOL_BIOSYNTHETIC_PROCESS, GO_ALCOHOL_BIOSYNTHETIC_PROCESS
KEGG_BLADDER_CANCER, KEGG_BLADDER_CANCER
GO_POSITIVE_REGULATION_OF_MITOTIC_SISTER_CHROMATID_SEPARATION, GO_POSITIVE_REGULATION_OF_MITOTIC_SISTER_CHROMATID_SEPARATION
GO_MONOCARBOXYLIC_ACID_CATABOLIC_PROCESS, GO_MONOCARBOXYLIC_ACID_CATABOLIC_PROCESS
GO_POSITIVE_REGULATION_OF_TOR_SIGNALING, GO_POSITIVE_REGULATION_OF_TOR_SIGNALING
chr11q14, chr11q14
GO_DEOXYRIBOSE_PHOSPHATE_CATABOLIC_PROCESS, GO_DEOXYRIBOSE_PHOSPHATE_CATABOLIC_PROCESS
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_DN, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_DN
GO_CELLULAR_RESPONSE_TO_DSRNA, GO_CELLULAR_RESPONSE_TO_DSRNA
GSE13306_LAMINA_PROPRIA_VS_SPLEEN_TREG_DN, GSE13306_LAMINA_PROPRIA_VS_SPLEEN_TREG_DN
BIOCARTA_ARF_PATHWAY, BIOCARTA_ARF_PATHWAY
GSE32901_TH17_EMRICHED_VS_TH17_NEG_CD4_TCELL_UP, GSE32901_TH17_EMRICHED_VS_TH17_NEG_CD4_TCELL_UP
MODULE_432, MODULE_432
GO_INTRINSIC_COMPONENT_OF_PEROXISOMAL_MEMBRANE, GO_INTRINSIC_COMPONENT_OF_PEROXISOMAL_MEMBRANE
GO_VERY_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS, GO_VERY_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS