

H_IL4_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_IL4_TREATED_MACROPHAGE_DN

GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_UP, GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_UP
GSE25123_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP, GSE25123_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_UP
GSE2405_S_AUREUS_VS_A_PHAGOCYTOPHILUM_NEUTROPHIL_DN, GSE2405_S_AUREUS_VS_A_PHAGOCYTOPHILUM_NEUTROPHIL_DN
GSE32901_TH17_EMRICHD_VS_TH17_NEG_CD4_TCELL_DN, GSE32901_TH17_EMRICHD_VS_TH17_NEG_CD4_TCELL_DN
GSE16385_ROSIGLITAZONE_VS_UNTREATED_IFNG_TNF_STIM_MACROPHAGE_DN, GSE16385_ROSIGLITAZONE_VS_UNTREATED_IFNG_TNF_STIM_MACROPHAGE_DN
GSE3039_CD4_TCELL_VS_ALPHAALPHA_CD8_TCELL_UP, GSE3039_CD4_TCELL_VS_ALPHAALPHA_CD8_TCELL_UP
GSE40274_SATB1_VS_FOXP3_AND_SATB1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_SATB1_VS_FOXP3_AND_SATB1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
WP_INSULIN_SIGNALING, WP_INSULIN_SIGNALING
GSE21380_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_DN, GSE21380_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_DN
GSE17301_CTRL_VS_48H_IFNA2_STIM_CD8_TCELL_DN, GSE17301_CTRL_VS_48H_IFNA2_STIM_CD8_TCELL_DN
GSE29164_DAY3_VS_DAY7_CD8_TCELL_TREATED_MELANOMA_UP, GSE29164_DAY3_VS_DAY7_CD8_TCELL_TREATED_MELANOMA_UP
GSE23114_WT_VS_SLE2C1_MOUSE_SPLEEN_B1A_BCELL_DN, GSE23114_WT_VS_SLE2C1_MOUSE_SPLEEN_B1A_BCELL_DN
GSE12505_WT_VS_E2_2_HET_PDC_DN, GSE12505_WT_VS_E2_2_HET_PDC_DN
SENGUPTA_EBNA1_ANTICORRELATED, SENGUPTA_EBNA1_ANTICORRELATED
GSE37532_VISCERAL_ADIPOSE_TISSUE_VS_LN_DERIVED_PPARG_KO_TREG_CD4_TCELL_DN, GSE37532_VISCERAL_ADIPOSE_TISSUE_VS_LN_DERIVED_PPARG_KO_TREG_CD4_TCELL_DN
MIR6852_5P, MIR6852_5P
GSE10240_IL22_VS_IL22_AND_IL17_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN, GSE10240_IL22_VS_IL22_AND_IL17_STIM_PRIMARY_BRONCHIAL_EPITHELIAL_CELLS_DN
GSE42021_CD24INT_TREG_VS_CD24INT_TCONV_THYMUS_DN, GSE42021_CD24INT_TREG_VS_CD24INT_TCONV_THYMUS_DN
WP_ANGIOPOIETIN_LIKE_PROTEIN_8_REGULATORY_PATHWAY, WP_ANGIOPOIETIN_LIKE_PROTEIN_8_REGULATORY_PATHWAY
GSE28449_WT_VS_LRF_KO_GERMINAL_CENTER_BCELL_DN, GSE28449_WT_VS_LRF_KO_GERMINAL_CENTER_BCELL_DN
GSE22025_UNTREATED_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_UP, GSE22025_UNTREATED_VS_TGFB1_AND_PROGESTERONE_TREATED_CD4_TCELL_UP
GSE22935_UNSTIM_VS_24H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_DN, GSE22935_UNSTIM_VS_24H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_DN
MIR4436B_3P, MIR4436B_3P
YTCCCRNNAGGY_UNKNOWN, YTCCCRNNAGGY_UNKNOWN
GSE16385_UNTREATED_VS_12H_IFNG_TNF_TREATED_MACROPHAGE_UP, GSE16385_UNTREATED_VS_12H_IFNG_TNF_TREATED_MACROPHAGE_UP
MIR7843_5P, MIR7843_5P
MIR6846_5P, MIR6846_5P
GOBP_PLASMA_MEMBRANE_ORGANIZATION, GOBP_PLASMA_MEMBRANE_ORGANIZATION
GSE11884_WT_VS_FURIN_KO_NAIVE_CD4_TCELL_UP, GSE11884_WT_VS_FURIN_KO_NAIVE_CD4_TCELL_UP
MIR6763_5P, MIR6763_5P
MIR3150A_3P, MIR3150A_3P
REACTOME_AKT_PHOSPHORYLATES_TARGETS_IN_THE_CYTOSOL, REACTOME_AKT_PHOSPHORYLATES_TARGETS_IN_THE_CYTOSOL
KEGG_ABC_TRANSPORTERS, KEGG_ABC_TRANSPORTERS
GOCC_MYELIN_SHEATH, GOCC_MYELIN_SHEATH
GSE8515_CTRL_VS_IL6_4H_STIM_MAC_DN, GSE8515_CTRL_VS_IL6_4H_STIM_MAC_DN
REACTOME_SIGNALING_BY_MODERATE_KINASE_ACTIVITY_BRAF_MUTANTS, REACTOME_SIGNALING_BY_MODERATE_KINASE_ACTIVITY_BRAF_MUTANTS
GSE12839_CTRL_VS_IL12_TREATED_PBMCDN, GSE12839_CTRL_VS_IL12_TREATED_PBMCDN
GOBP_POSITIVE_CHEMOTAXIS, GOBP_POSITIVE_CHEMOTAXIS
ZHONG_PFC_C2_THY1_POS_OPC, ZHONG_PFC_C2_THY1_POS_OPC
GOBP_PLASMA_MEMBRANE_TUBULATION, GOBP_PLASMA_MEMBRANE_TUBULATION
YAP1_DN, YAP1_DN
GOBP_POSITIVE_REGULATION_OF_CHOLESTEROL_EFFLUX, GOBP_POSITIVE_REGULATION_OF_CHOLESTEROL_EFFLUX
GOCC_MAIN_AXON, GOCC_MAIN_AXON
KAAB_FAILED_HEART_ATRIUM_UP, KAAB_FAILED_HEART_ATRIUM_UP
GOBP_THYROID_GLAND_DEVELOPMENT, GOBP_THYROID_GLAND_DEVELOPMENT
GOBP_REGULATION_OF_SENSORY_PERCEPTION, GOBP_REGULATION_OF_SENSORY_PERCEPTION
GOBP_CEREBRAL_CORTEX_GABAERGIC_INTERNEURON_DIFFERENTIATION, GOBP_CEREBRAL_CORTEX_GABAERGIC_INTERNEURON_DIFFERENTIATION
GOBP_REGULATION_OF_CELL_MATURATION, GOBP_REGULATION_OF_CELL_MATURATION
GOBP_GABAERGIC_NEURON_DIFFERENTIATION, GOBP_GABAERGIC_NEURON_DIFFERENTIATION
GOBP_CEREBRAL_CORTEX_GABAERGIC_INTERNEURON_DEVELOPMENT, GOBP_CEREBRAL_CORTEX_GABAERGIC_INTERNEURON_DEVELOPMENT
GOBP_CEREBRAL_CORTEX_GABAERGIC_INTERNEURON_MIGRATION, GOBP_CEREBRAL_CORTEX_GABAERGIC_INTERNEURON_MIGRATION
THEODOROU_MAMMARY_TUMORIGENESIS, THEODOROU_MAMMARY_TUMORIGENESIS
chr9p13, chr9p13
GOBP_SUBSTRATE_INDEPENDENT_TELENCEPHALIC_TANGENTIAL_MIGRATION, GOBP_SUBSTRATE_INDEPENDENT_TELENCEPHALIC_TANGENTIAL_MIGRATION
GOBP_INTERNEURON_MIGRATION, GOBP_INTERNEURON_MIGRATION
GOBP_FOREBRAIN_DORSAL_VENTRAL_PATTERN_FORMATION, GOBP_FOREBRAIN_DORSAL_VENTRAL_PATTERN_FORMATION
HP_SCLEROTIC_VERTEBRAL_ENDPLATES, HP_SCLEROTIC_VERTEBRAL_ENDPLATES
GOMF_TYPE_2_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_BINDING, GOMF_TYPE_2_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_BINDING