

1_VS_8H_LPS_AND_R848_DC_UP, GSE2706_UNSTIM_VS_8H_LPS_AND_R848_DC_UP

GSE16755_CTRL_VS_IFNA_TREATED_MAC_UP, GSE16755_CTRL_VS_IFNA_TREATED_MAC_UP
GSE27859_MACROPHAGE_VS_CD11C_INT_F480_HI_MACROPHAGE_DN, GSE27859_MACROPHAGE_VS_CD11C_INT_F480_HI_MACROPHAGE_DN
GSE14000_UNSTIM_VS_16H_LPS_DC_UP, GSE14000_UNSTIM_VS_16H_LPS_DC_UP
GSE30971_WBP7_HET_VS_KO_MACROPHAGE_4H_LPS_STIM_DN, GSE30971_WBP7_HET_VS_KO_MACROPHAGE_4H_LPS_STIM_DN
GSE35825_UNTREATED_VS_IFNA_STIM_MACROPHAGE_UP, GSE35825_UNTREATED_VS_IFNA_STIM_MACROPHAGE_UP
GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_2H_UP, GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_2H_UP
GSE9037_CTRL_VS_LPS_4H_STIM_BMDM_UP, GSE9037_CTRL_VS_LPS_4H_STIM_BMDM_UP
GSE2706_LPS_VS_R848_AND_LPS_8H_STIM_DC_UP, GSE2706_LPS_VS_R848_AND_LPS_8H_STIM_DC_UP
GSE3039_NKT_CELL_VS_B2_BCELL_UP, GSE3039_NKT_CELL_VS_B2_BCELL_UP
GSE9037_WT_VS_IRAK4_KO_LPS_4H_STIM_BMDM_DN, GSE9037_WT_VS_IRAK4_KO_LPS_4H_STIM_BMDM_DN
GSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_DN, GSE17721_CTRL_VS_PAM3CSK4_24H_BMDC_DN
GSE2706_R848_VS_R848_AND_LPS_8H_STIM_DC_UP, GSE2706_R848_VS_R848_AND_LPS_8H_STIM_DC_UP
HP_RECURRENT_URINARY_TRACT_INFECTIONS, HP_RECURRENT_URINARY_TRACT_INFECTIONS
HP_ABNORMAL_TOENAIL_MORPHOLOGY, HP_ABNORMAL_TOENAIL_MORPHOLOGY
CHAF1A_TARGET_GENES, CHAF1A_TARGET_GENES
HP_ABNORMAL_FINGERNAIL_MORPHOLOGY, HP_ABNORMAL_FINGERNAIL_MORPHOLOGY
GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_180MIN_UP, GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_180MIN_UP
MIR628_3P, MIR628_3P
GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_129_MOUSE_DN, GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_129_MOUSE_DN
HP_RECURRENT_LOWER_RESPIRATORY_TRACT_INFECTIONS, HP_RECURRENT_LOWER_RESPIRATORY_TRACT_INFECTIONS
GSE339_CD4POS_VS_CD4CD8DN_DC_IN_CULTURE_DN, GSE339_CD4POS_VS_CD4CD8DN_DC_IN_CULTURE_DN
GOBP_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GOBP_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION
GSE15659_CD45RA_NEG_CD4_TCELL_VS_RESTING_TREG_UP, GSE15659_CD45RA_NEG_CD4_TCELL_VS_RESTING_TREG_UP
GOBP_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_DIFFERENTIATION
GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_UNTREATED_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_UP
HOXA13_TARGET_GENES, HOXA13_TARGET_GENES
MIR569, MIR569
MIR6804_5P, MIR6804_5P
MIR3934_5P, MIR3934_5P
MIR3650, MIR3650
HP_ABNORMALITY_OF_THE_ACETABULUM, HP_ABNORMALITY_OF_THE_ACETABULUM
BIOCARTA_HSWI_SNF_PATHWAY, BIOCARTA_HSWI_SNF_PATHWAY
REACTOME_GAMMA_CARBOXYLATION_HYPUSINE_FORMATION_AND_ARYLSULFATASE_ACTIVATION, REACTOME_GAMMA_CARBOXYLATION_HYPUSINE_FORMATION_AND_ARYLSULFATASE_ACTIVATION
HP_FLAT_FOREHEAD, HP_FLAT_FOREHEAD
GOBP_NEUTRAL_AMINO_ACID_TRANSPORT, GOBP_NEUTRAL_AMINO_ACID_TRANSPORT
GOBP_POSITIVE_REGULATION_OF_SMAD_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_POSITIVE_REGULATION_OF_SMAD_PROTEIN_SIGNAL_TRANSDUCTION
REACTOME_ACYL_CHAIN_REMODELLING_OF_PC, REACTOME_ACYL_CHAIN_REMODELLING_OF_PC
HP_YOUNG_ADULT_ONSET, HP_YOUNG_ADULT_ONSET
ABDELMOHSEN_ELAVL4_TARGETS, ABDELMOHSEN_ELAVL4_TARGETS
GOBP_BONE_MORPHOGENESIS, GOBP_BONE_MORPHOGENESIS
HOWLIN_CITED1_TARGETS_2_UP, HOWLIN_CITED1_TARGETS_2_UP
REACTOME_ACYL_CHAIN_REMODELLING_OF_PG, REACTOME_ACYL_CHAIN_REMODELLING_OF_PG
GOMF_CARBOHYDRATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_CARBOHYDRATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
REACTOME_ACYL_CHAIN_REMODELLING_OF_PS, REACTOME_ACYL_CHAIN_REMODELLING_OF_PS
REACTOME_REDUCTION_OF_CYTOSOLIC_CA_LEVELS, REACTOME_REDUCTION_OF_CYTOSOLIC_CA_LEVELS
GOBP_PHOSPHATIDYLSERINE_ACYL_CHAIN_REMODELING, GOBP_PHOSPHATIDYLSERINE_ACYL_CHAIN_REMODELING
GOBP_FLUID_TRANSPORT, GOBP_FLUID_TRANSPORT