

017721.POLYIC.VS.GARDQUIMOD_6H.BMDC.DN,GSE17721.POLYIC.VS.GARDQUIMOD_6H.BMDC.DN
GSE16385.MONOCYTE.VS.12H_IL4.TREATED.MACROPHAGE.DN,GSE16385.MONOCYTE.VS.12H_IL4.TREATED.MACROPHAGE.DN
GSE411.UNSTIM.VS.40MIN_IL4.STIM.MACROPHAGE.DN,GSE411.UNSTIM.VS.40MIN_IL4.STIM.MACROPHAGE.DN
GSE2925.LIGHT_ZONE.VS.NAIVE.BCELL.DN,GSE2925.LIGHT_ZONE.VS.NAIVE.BCELL.DN
GSE14769.UNSTIM.VS.80MIN_ILPS.BMDM.UP,GSE14769.UNSTIM.VS.80MIN_ILPS.BMDM.UP
GSE18893.TCONV.VS.TREG.24H.TNF.STIM.DN,GSE18893.TCONV.VS.TREG.24H.TNF.STIM.DN
GSE13484.3H.UNSTIM.VS.YF1D7.VACCINE.STIM.PBMC.UP,GSE13484.3H.UNSTIM.VS.YF1D7.VACCINE.STIM.PBMC.UP
GSE2925.DARK_ZONE.VS.NAIVE.BCELL.UP,GSE2925.DARK_ZONE.VS.NAIVE.BCELL.UP
GSE4046.AZAR.KO.VS.AZAR.AGONIST.TREATED.TREG.DN,GSE4046.AZAR.KO.VS.AZAR.AGONIST.TREATED.TREG.DN
GSE16385.MONOCYTE.VS.12H_ROSGILITAZONE.FJNG.TNF.TREATED.MACROPHAGE.DN,GSE16385.MONOCYTE.VS.12H_ROSGILITAZONE.FJNG.TNF.TREATED.MACROPHAGE.DN
GSE17721.CPG.VS.GARDQUIMOD_2H.BMDC.DN,GSE17721.CPG.VS.GARDQUIMOD_2H.BMDC.DN
GSE15324.ELF4.KO.VS.WT.ACTIVATED_CD8.TCELL.UP,GSE15324.ELF4.KO.VS.WT.ACTIVATED_CD8.TCELL.UP
GSE32423.MEMORY.VS.NAIVE_CD8.TCELL.DN,GSE32423.MEMORY.VS.NAIVE_CD8.TCELL.DN
GSE28985.CTRL.VS.ROSIGLITAZONE.STIM.MACROPHAGE.UP,GSE28985.CTRL.VS.ROSIGLITAZONE.STIM.MACROPHAGE.UP
GSE17721.LPS.VS.GARDQUIMOD_1H.BMDC.DN,GSE17721.LPS.VS.GARDQUIMOD_1H.BMDC.DN
GCM_HBP1.GCM_HBP1
GSE17721.LPS.VS.PAM3CSK4.24H.BMDC.UP,GSE17721.LPS.VS.PAM3CSK4.24H.BMDC.UP
GSE43955.THO.VS.TGFB_IL4.TH17.ACT_CD4_TCELL_42H.UP,GSE43955.THO.VS.TGFB_IL4.TH17.ACT_CD4_TCELL_42H.UP
MIR640.MIR640
MIR5148.5P.MIR5148.5P
MIR1569.RESTING.VS.ACTIVATED.TREG.DN,GSE1569.RESTING.VS.ACTIVATED.TREG.DN
MIR13C.3P.MIR13C.3P
MIR675A.3P.MIR675A.3P
GSE21063.CTRL.VS.ANTL_ICM.STIM.BCELL_3H.DN,GSE21063.CTRL.VS.ANTL_ICM.STIM.BCELL_3H.DN
GSE3225.UNSTIM.VS.4H_ILPS.STIM.DC.DN,GSE3225.UNSTIM.VS.4H_ILPS.STIM.DC.DN
GSE42021.CD24LO.TREG.VS.CD24LO_TCONV_THYMUS.UP,GSE42021.CD24LO.TREG.VS.CD24LO_TCONV_THYMUS.UP
GSE16385.UNTREATED.VS.12H_ROSGILITAZONE.FJNG.TNF.TREATED.MACROPHAGE.DN,GSE16385.UNTREATED.VS.12H_ROSGILITAZONE.FJNG.TNF.TREATED.MACROPHAGE.DN
GSE17721.LPS.VS.12H.PAM3CSK4.BMDC.UP,GSE17721.LPS.VS.12H.PAM3CSK4.BMDC.UP
GSE9946.MATURE.STIMULATORY.VS.LISTERIA_INF.MATURE_DC.UP,GSE9946.MATURE.STIMULATORY.VS.LISTERIA_INF.MATURE_DC.UP
GSE42088.UNINF.VS.LISHMANIA_INF_DC.24H.DN,GSE42088.UNINF.VS.LISHMANIA_INF_DC.24H.DN
GSE26669.CD4.VS.CD8.TCELL_IN_MLR.DN,GSE26669.CD4.VS.CD8.TCELL_IN_MLR.DN
FIGUEROA_AML_METHYLATION_CLUSTER3.UP,FIGUEROA_AML_METHYLATION_CLUSTER3.UP
MIR6473.MIR6473
GOBP_PROTEIN_INSERTION_INTO_MEMBRANE,GOBP_PROTEIN_INSERTION_INTO_MEMBRANE
GSE1925.CTRL.VS.JFNG.PRIMED.MACROPHAGE.24H.JFNG.STIM.UP,GSE1925.CTRL.VS.JFNG.PRIMED.MACROPHAGE.24H.JFNG.STIM.UP
GSE9601.UNTREATED.VS.P13K.INHIBITOR.TREATED.HCMV_INF.MONOCYTE.DN,GSE9601.UNTREATED.VS.P13K.INHIBITOR.TREATED.HCMV_INF.MONOCYTE.DN
MIR150.3P.MIR150.3P
ETT_Q8.ETT_Q8
MIR3927.3P.MIR3927.3P
GOBP_CYTOSOLIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY,GOBP_CYTOSOLIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY
MIR6730.5P.MIR6730.5P
GSE16385.UNTREATED.VS.12H_IL4.TREATED.MACROPHAGE.DN,GSE16385.UNTREATED.VS.12H_IL4.TREATED.MACROPHAGE.DN
MIR15.3P.MIR15.3P
MIR9718.MIR9718
AHRARNT_01.AHRARNT_01
MIR647.MIR647B,MIR647B,MIR647B
MIR6758.3P.MIR6758.3P
MIR629.5P.MIR629.5P
MIR34.3P.MIR34.3P
FIGUEROA_AML_METHYLATION_CLUSTER6.UP,FIGUEROA_AML_METHYLATION_CLUSTER6.UP
GOBP_B_CELL_DIFFERENTIATION,GOBP_B_CELL_DIFFERENTIATION
HP_CUTS.MARMORATA,HP_CUTS.MARMORATA
MIR625.5P.MIR625.5P
GSE4606.UNSTIM.VS.CD40L_IL5_1DAY.STIMULATED_IRRF4MID_SORTED_BCELL.DN,GSE4606.UNSTIM.VS.CD40L_IL5_1DAY.STIMULATED_IRRF4MID_SORTED_BCELL.DN
GOBP_REGULATION_OF_PROTEIN_ACETYLATION,GOBP_REGULATION_OF_PROTEIN_ACETYLATION
HP_TRIGONOCEPHALY,HP_TRIGONOCEPHALY
MIR183.MIR183
ACTCAGC_MIR345,ACTCAGC_MIR345
FIGUEROA_AML_METHYLATION_CLUSTER7.UP,FIGUEROA_AML_METHYLATION_CLUSTER7.UP
GCM_BCL2L1.GCM_BCL2L1
HP_DECREASED_CIRCULATING_IGG_LEVEL,HP_DECREASED_CIRCULATING_IGG_LEVEL
MIR631.MIR631
GSE34156.UNTREATED.VS.6H_NOD2_AND_TLR1_TLR2_LIGAND.TREATED.MONOCYTE.UP,GSE34156.UNTREATED.VS.6H_NOD2_AND_TLR1_TLR2_LIGAND.TREATED.MONOCYTE.UP
GENTILE_UV_RESPONSE_CLUSTER_D8,GENTILE_UV_RESPONSE_CLUSTER_D8
HAY_BONE_MARROW_CD34_POS_HSC,HAY_BONE_MARROW_CD34_POS_HSC
GOBP_FOUR_WAY_JUNCTION_DNA_BINDING,GOBP_FOUR_WAY_JUNCTION_DNA_BINDING
ELI_UX30.XD.TARGETS.DN,ELI_UX30.XD.TARGETS.DN
IKEDA_MIR133.TARGETS.UP,IKEDA_MIR133.TARGETS.UP
GSE9946.MATURE.STIMULATORY.VS.PROSTAGLANDIN2.TREATED.MATURE_DC.UP,GSE9946.MATURE.STIMULATORY.VS.PROSTAGLANDIN2.TREATED.MATURE_DC.UP
GSE12963.UNINF.VS.ENV_AND_NEF_DEFICIENT_HIV1_INF_CD4_TCELL.UP,GSE12963.UNINF.VS.ENV_AND_NEF_DEFICIENT_HIV1_INF_CD4_TCELL.UP
BECKER_TAMOXIFEN_RESISTANCE.UP,BECKER_TAMOXIFEN_RESISTANCE.UP
CTCTCAGC_MIR342,CTCTCAGC_MIR342
REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_CELL_DEATH_GENES,REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_CELL_DEATH_GENES
GOMF_ADENYLYLTRANSFERASE_ACTIVITY,GOMF_ADENYLYLTRANSFERASE_ACTIVITY
GOMF_FIBROBLAST_APOPTOTIC_PROCESS,GOBP_FIBROBLAST_APOPTOTIC_PROCESS
GOMF_TRANSCRIPTION_COACTIVATOR_BINDING,GOMF_TRANSCRIPTION_COACTIVATOR_BINDING
MIR645.MIR645
GOBP_HEART_VALVE_DEVELOPMENT,GOBP_HEART_VALVE_DEVELOPMENT
PID_HES_HEY_PATHWAY,PID_HES_HEY_PATHWAY
FLECHNER_PBL_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN,FLECHNER_PBL_KIDNEY_TRANSPLANT_OK_VS_DONOR_DN
CCACAT_MIR2969P,CCACAT_MIR2969P
GOBP_POSITIVE_REGULATION_OF_CELLULAR_EXTRAVASATION,GOBP_POSITIVE_REGULATION_OF_CELLULAR_EXTRAVASATION
MIR619.3P.MIR619.3P
TATCTGG.MIR488,TATCTGG.MIR488
GOBP_V_D_J_RECOMBINATION,GOBP_V_D_J_RECOMBINATION
MIR6462A.3P.MIR6462A.3P
MIR4423.5P.MIR4423.5P
GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RESPONSE_TO_CHEMICAL_STIMULUS,GOBP_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RESPONSE_TO_CHEMICAL_STIMULUS
MIR3161.MIR3161
KEGG_SELENOAMINO_ACID_METABOLISM,KEGG_SELENOAMINO_ACID_METABOLISM
MIR69.3P.MIR69.3P
HP_HYPOPROTEINEMIA,HP_HYPOPROTEINEMIA
GSE41176.WT.VS.TAK1_KO.ANTL_ICM.STIM.BCELL_1H.DN,GSE41176.WT.VS.TAK1_KO.ANTL_ICM.STIM.BCELL_1H.DN
PARK_API_PATHOGENESIS.UP,PARK_API_PATHOGENESIS.UP
SKP2_TARGET_GENES,SKP2_TARGET_GENES
BIOCARTA_PCCL2_PATHWAY,BIOCARTA_PCCL2_PATHWAY
GOBP_ENDOCARDIAL_CUSHION_FORMATION,GOBP_ENDOCARDIAL_CUSHION_FORMATION
HP_PSORIASIFORM_DERMATITIS,HP_PSORIASIFORM_DERMATITIS
HP_NASOLACRIMAL_DUCT_OBSTRUCTION,HP_NASOLACRIMAL_DUCT_OBSTRUCTION
GOBP_REGULATION_OF_SYNCYTUM_FORMATION_BY_PLASMA_MEMBRANE_FUSION,GOBP_REGULATION_OF_SYNCYTUM_FORMATION_BY_PLASMA_MEMBRANE_FUSION
GOBP_PRC1_MENSA_PRESSESSING,GOBP_PRC1_MENSA_PRESSESSING
GOMF_PHOSPHOTRANSFERASE_ACTIVITY_FOR_OTHER_SUBSTITUTED_PHOSPHATE_GROUPS,GOMF_PHOSPHOTRANSFERASE_ACTIVITY_FOR_OTHER_SUBSTITUTED_PHOSPHATE_GROUPS
GOCC_MITOTIC_SPINDLE_MICROTUBULE,GOCC_MITOTIC_SPINDLE_MICROTUBULE
BERENJENO_ROCK_SIGNALING_NOT_VIA_RHOA.UP,BERENJENO_ROCK_SIGNALING_NOT_VIA_RHOA.UP
GOBP_TRANSFORMING_GROWTH_FACTOR_BETA_PRODUCTION,GOBP_TRANSFORMING_GROWTH_FACTOR_BETA_PRODUCTION
GOBP_CACO2_EPITHELIAL_TO_MISENCHYMAL_TRANSITION,GOBP_CACO2_EPITHELIAL_TO_MISENCHYMAL_TRANSITION
MIR20C.3P.MIR20C.3P
GOBP_GLOMERULAR_MESANGIUM_DEVELOPMENT,GOBP_GLOMERULAR_MESANGIUM_DEVELOPMENT
GOBP_EARLY_ENDOSOME_TO_GOLGI_TRANSPORT,GOBP_EARLY_ENDOSOME_TO_GOLGI_TRANSPORT
GOBP_VENTRICULAR_SEPTUM_MORPHOGENESIS,GOBP_VENTRICULAR_SEPTUM_MORPHOGENESIS
GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_EARLY_ENDOSOME,GOBP_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_EARLY_ENDOSOME
REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_CYTOSOLIC_C_RELEASE,REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_GENES_INVOLVED_IN_CYTOSOLIC_C_RELEASE
GOBP_EPITHELIAL_TO_MISENCHYMAL_TRANSITION_INVOLVED_IN_ENDOCARDIAL_CUSHION_FORMATION,GOBP_EPITHELIAL_TO_MISENCHYMAL_TRANSITION_INVOLVED_IN_ENDOCARDIAL_CUSHION_FORMATION
GOBP_AMYLOID_FIBRIL_FORMATION,GOBP_AMYLOID_FIBRIL_FORMATION
GOMF_L_ALANINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY,GOMF_L_ALANINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOBP_MUSCLE_HYPERTROPHY,GOBP_MUSCLE_HYPERTROPHY
GOBP_REGULATION_OF_PROTEIN_LIPIDATION,GOBP_REGULATION_OF_PROTEIN_LIPIDATION
GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CHROMATIN,GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CHROMATIN
GOCC_RNA_POLYMERASE_II_TRANSCRIPTION_REPRESSOR_COMPLEX,GOCC_RNA_POLYMERASE_II_TRANSCRIPTION_REPRESSOR_COMPLEX
GOMF_GUANYLATE_CYCLASE_ACTIVITY,GOMF_GUANYLATE_CYCLASE_ACTIVITY
MIR648.MIR648
HP_ABNORMAL_LACRIMAL_DUCT_MORPHOLOGY,HP_ABNORMAL_LACRIMAL_DUCT_MORPHOLOGY
GOBP_L_ALANINE_TRANSPORT,GOBP_L_ALANINE_TRANSPORT
GOBP_EXTRACELLULAR_VESICLE_BIOGENESIS,GOBP