

3_KO_4H_LPS_STIM_MACROPHAGE_UP, GSE33162_HDAC3_KO_VS_HDAC3_KO_4H_LPS_STIM_MACROPHAGE_UP

GSE41000_TRANSLATED_RNA_VS_MRNA_4H_LPS_DC_UP, GSE41000_TRANSLATED_RNA_VS_MRNA_4H_LPS_DC_UP
GSE37416_CTRL_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GSE37416_CTRL_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_DN, GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_DN
GSE10094_LCMV_VS_LISTERIA_IND_EFF_CD4_TCELL_UP, GSE10094_LCMV_VS_LISTERIA_IND_EFF_CD4_TCELL_UP
GSE17721_PAM3CSK4_VS_GADIQUIMOD_2H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_2H_BMDC_DN
GSE21063_WT_VS_NFATC1_KO_8H_ANTLI_IJM_STIM_BCELL_DN, GSE21063_WT_VS_NFATC1_KO_8H_ANTLI_IJM_STIM_BCELL_DN
GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP, GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP
GSE15330_WT_VS_IKAROS_KO_LYMPHOID_MULTIPOTENT_PROGENITOR_UP, GSE15330_WT_VS_IKAROS_KO_LYMPHOID_MULTIPOTENT_PROGENITOR_UP
GSE36527_CD69_NEG_VS_POS_TREG_CD62L_LOS_KLRG1_NEG_DN, GSE36527_CD69_NEG_VS_POS_TREG_CD62L_LOS_KLRG1_NEG_DN
GSE17721_LPS_VS_GARDIQUIMOD_1H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_1H_BMDC_DN
GSE32034_LY6C_HIGH_VS_LOW_MONOCYTE_DN, GSE32034_LY6C_HIGH_VS_LOW_MONOCYTE_DN
GSE42724_MEMORY_BCELL_VS_PLASMABLAST_DN, GSE42724_MEMORY_BCELL_VS_PLASMABLAST_DN
GSE32164_RESTING_DIFFERENTIATED_VS_ALTERNATIVELY_ACT_M2_MACROPHAGE_DN, GSE32164_RESTING_DIFFERENTIATED_VS_ALTERNATIVELY_ACT_M2_MACROPHAGE_DN
GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_UP, GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_UP
GSE3982_BASOPHIL_VS_CENT_MEMORY_CD4_TCELL_UP, GSE3982_BASOPHIL_VS_CENT_MEMORY_CD4_TCELL_UP
GSE39820_CTRL_VS_IL1B_IL6_IL23A_CD4_TCELL_UP, GSE39820_CTRL_VS_IL1B_IL6_IL23A_CD4_TCELL_UP
GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP, GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP
GSE4748_LPS_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP, GSE4748_LPS_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP
GSE36009_WT_VS_NLRP10_KO_DC_LPS_STIM_DN, GSE36009_WT_VS_NLRP10_KO_DC_LPS_STIM_DN
GSE3920_IFNA_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP, GSE3920_IFNA_VS_IFNG_TREATED_ENDOTHELIAL_CELL_UP
GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_DN, GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_DN
GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_DN, GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_DN
GSE27786_LSK_VS_BCELL_DN, GSE27786_LSK_VS_BCELL_DN
HAHTOLA_MYCOSIS_FUNGOIDES_SKIN_UP, HAHTOLA_MYCOSIS_FUNGOIDES_SKIN_UP
GSE11973_MIR223_KOVS_WT_BONE_MARROW_NEUTROPHIL_UP, GSE11973_MIR223_KOVS_WT_BONE_MARROW_NEUTROPHIL_UP
GSE1925_CTRL_VS_24H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN, GSE1925_CTRL_VS_24H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN
GSE27786_LIN_NEG_VS_NKTCCELL_DN, GSE27786_LIN_NEG_VS_NKTCCELL_DN
GSE40274_EOS_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_EOS_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_DN, GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_DN
GSE17721_0.5H_VS_4H_GARDIQUIMOD_BMDC_DN, GSE17721_0.5H_VS_4H_GARDIQUIMOD_BMDC_DN
ATACTGT_MIR144, ATACTGT_MIR144
GSE39152_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_UP, GSE39152_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_UP
AAGGGAT_MIR188, AAGGGAT_MIR188
GSE38681_WT_VS_LYL1_KO_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_DN, GSE38681_WT_VS_LYL1_KO_LYMPHOID_PRIMED_MULTIPOTENT_PROGENITOR_DN
GSE17721_POLYIC_VS_GARDIQUIMOD_2H_BMDC_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_2H_BMDC_UP
GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_DN, GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_DN
GSE17721_0.5H_VS_12H_POLYIC_BMDC_DN, GSE17721_0.5H_VS_12H_POLYIC_BMDC_DN
GSE27434_WT_VS_DNMT1_KO_TREG_DN, GSE27434_WT_VS_DNMT1_KO_TREG_DN
GSE17721_POLYIC_VS_GARDIQUIMOD_1H_BMDC_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_1H_BMDC_DN
GSE33424_CD161_HIGH_VS_NEG_CD8_TCELL_UP, GSE33424_CD161_HIGH_VS_NEG_CD8_TCELL_UP
GSE15330_WT_VS_IKAROS_KO_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN, GSE15330_WT_VS_IKAROS_KO_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN
GSE24142_ADULT_VS_FETAL_EARLY_THYMIC_PROGENITOR_UP, GSE24142_ADULT_VS_FETAL_EARLY_THYMIC_PROGENITOR_UP
GSE17721_PAM3CSK4_VS_CPG_0.5H_BMDC_UP, GSE17721_PAM3CSK4_VS_CPG_0.5H_BMDC_UP
GSE21670_TGFB_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_DN, GSE21670_TGFB_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_DN
GSE17721_LPS_VS_CPG_8H_BMDC_UP, GSE17721_LPS_VS_CPG_8H_BMDC_UP
GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_UP, GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_UP
GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN
GSE29618_PRE_VS_DAY7_POST_LAIY_FLU_VACCINE_MDC_DN, GSE29618_PRE_VS_DAY7_POST_LAIY_FLU_VACCINE_MDC_DN
GO_NCRNA_TRANSCRIPTION, GO_NCRNA_TRANSCRIPTION
GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_UP, GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_UP
GSE43955_TGFB_IL6_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_60H_UP, GSE43955_TGFB_IL6_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_60H_UP
GO_REGULATION_OF_MRNA_METABOLIC_PROCESS, GO_REGULATION_OF_MRNA_METABOLIC_PROCESS
GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_4H_UP, GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_4H_UP
GSE27786_LSK_VS_CD4_TCELL_DN, GSE27786_LSK_VS_CD4_TCELL_DN
GSE41087_WT_VS_FOXP3_MUT_ANTLI_CD3_CD28_STIM_CD4_TCELL_UP, GSE41087_WT_VS_FOXP3_MUT_ANTLI_CD3_CD28_STIM_CD4_TCELL_UP
GSE369_SOCS3_KO_VS_WT_LIVER_POST_IL6_INJECTION_DN, GSE369_SOCS3_KO_VS_WT_LIVER_POST_IL6_INJECTION_DN
GSE22935_UNSTIM_VS_48H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_UP, GSE22935_UNSTIM_VS_48H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_UP
GCM_ERBB2IP, GCM_ERBB2IP
GSE42021_TREG_PLN_VS_CD24HL_TREG_THYMUS_UP, GSE42021_TREG_PLN_VS_CD24HL_TREG_THYMUS_UP
KEEN_RESPONSE_TO_ROSIGLITAZONE_DN, KEEN_RESPONSE_TO_ROSIGLITAZONE_DN
GSE42021_TCONV_PLN_VS_CD24INT_TCONV_THYMUS_UP, GSE42021_TCONV_PLN_VS_CD24INT_TCONV_THYMUS_UP
GO_REGULATION_OF_RNA_SPLICING, GO_REGULATION_OF_RNA_SPLICING
YAGL_AML_SURVIVAL, YAGL_AML_SURVIVAL
GSE17721_LPS_VS_PAM3CSK4_12H_BMDC_UP, GSE17721_LPS_VS_PAM3CSK4_12H_BMDC_UP
GSE7852_THYMUS_VS_FAT_TCONV_DN, GSE7852_THYMUS_VS_FAT_TCONV_DN
GCM_CSNK1A1, GCM_CSNK1A1
GO_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION, GO_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION
GSE1448_ANTLI_VALPHA2_VS_YBETA5_DP_THYMOCYTE_UP, GSE1448_ANTLI_VALPHA2_VS_YBETA5_DP_THYMOCYTE_UP
KEGG_NEUROTROPHIN_SIGNALING_PATHWAY, KEGG_NEUROTROPHIN_SIGNALING_PATHWAY
GOLDRATH_NAIVE_VS_MEMORY_CD8_TCELL_DN, GOLDRATH_NAIVE_VS_MEMORY_CD8_TCELL_DN
GSE7768_OVA_ALONE_VS_OVA_WITH_LPS_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_DN, GSE7768_OVA_ALONE_VS_OVA_WITH_LPS_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_DN
GGCACTT_MIR519E, GGCACTT_MIR519E
GSE17721_LPS_VS_GARDIQUIMOD_0.5H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_0.5H_BMDC_DN
GSE17721_CTRL_VS_CPG_12H_BMDC_DN, GSE17721_CTRL_VS_CPG_12H_BMDC_DN
GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP
GSE8621_UNSTIM_VS_LPS_PRIMED_AND_LPS_STIM_MACROPHAGE_DN, GSE8621_UNSTIM_VS_LPS_PRIMED_AND_LPS_STIM_MACROPHAGE_DN
GSE4142_PLASMA_CELL_VS_GC_BCELL_UP, GSE4142_PLASMA_CELL_VS_GC_BCELL_UP
GSE1460_INTRATHYMIC_T_PROGENITOR_VS_THYMIC_STROMAL_CELL_UP, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_THYMIC_STROMAL_CELL_UP
GAVIN_FOXP3_TARGETS_CLUSTER_T4, GAVIN_FOXP3_TARGETS_CLUSTER_T4
MODULE_331, MODULE_331
GSE40274_LEF1_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_LEF1_VS_FOXP3_AND_LEF1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE17721_POLYIC_VS_GARDIQUIMOD_24H_BMDC_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_24H_BMDC_UP
GCGSCMNTTT_UNKNOWN, GCGSCMNTTT_UNKNOWN
GSE19923_E2A_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP, GSE19923_E2A_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP
GENTILE_RESPONSE_CLUSTER_D3, GENTILE_RESPONSE_CLUSTER_D3
GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
GO_REGULATION_OF_ORGANELLE_ASSEMBLY, GO_REGULATION_OF_ORGANELLE_ASSEMBLY
GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_6H_DN, GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_6H_DN
GSE7852_LN_VS_FAT_TCONV_DN, GSE7852_LN_VS_FAT_TCONV_DN
GSE37532_WT_VS_PPARG_KO_LN_TCONV_UP, GSE37532_WT_VS_PPARG_KO_LN_TCONV_UP
GSE7852_TREG_VS_TCONV_THYMUS_UP, GSE7852_TREG_VS_TCONV_THYMUS_UP
GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY3_POST_IMMUNIZATION_UP, GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY3_POST_IMMUNIZATION_UP
GSE7460_TCONV_VS_TREG_THYMUS_DN, GSE7460_TCONV_VS_TREG_THYMUS_DN
GO_LYSOSOMAL_TRANSPORT, GO_LYSOSOMAL_TRANSPORT
GO_TRANSCRIPTIONAL_REPRESSOR_COMPLEX, GO_TRANSCRIPTIONAL_REPRESSOR_COMPLEX
AACTGAC_MIR223, AACTGAC_MIR223
MORI_MATURE_B_LYMPHOCYTE_UP, MORI_MATURE_B_LYMPHOCYTE_UP
SIG_PIP3_SIGNALING_IN_CARDIAC_MYOCYTES, SIG_PIP3_SIGNALING_IN_CARDIAC_MYOCYTES
GTATGAT_MIR154_MIR487, GTATGAT_MIR154_MIR487
GO_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY, GO_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY
GSE360_CTRL_VS_T_GONDIL_DC_DN, GSE360_CTRL_VS_T_GONDIL_DC_DN
MARIADASON_REGULATED_BY_HISTONE_ACETYLATION_UP, MARIADASON_REGULATED_BY_HISTONE_ACETYLATION_UP
GO_POSITIVE_REGULATION_OF_MULTIORGANISM_PROCESS, GO_POSITIVE_REGULATION_OF_MULTIORGANISM_PROCESS
GO_REGULATION_OF_DNA_REPLICATION, GO_REGULATION_OF_DNA_REPLICATION
GSE3982_MAC_VS_NEUTROPHIL_DN, GSE3982_MAC_VS_NEUTROPHIL_DN
GNF2_RAP1B, GNF2_RAP1B
GSE43955_10H_VS_30H_ACT_CD4_TCELL_DN, GSE43955_10H_VS_30H_ACT_CD4_TCELL_DN
PID_TGFB_R_PATHWAY, PID_TGFB_R_PATHWAY
GO_REGULATION_OF_VACUOLE_ORGANIZATION, GO_REGULATION_OF_VACUOLE_ORGANIZATION
GO_RECYCLING_ENDOSOME, GO_RECYCLING_ENDOSOME
ST_PHOSPHOINOSITIDE_3_KINASE_PATHWAY, ST_PHOSPHOINOSITIDE_3_KINASE_PATHWAY
GCTCTTG_MIR335, GCTCTTG_MIR335
ATTACAT_MIR3803P, ATTACAT_MIR3803P
GO_LATE_ENDOSOME_MEMBRANE, GO_LATE_ENDOSOME_MEMBRANE
GSE7568_IL4_VS_IL4_AND_DEXAMETHASONE_TREATED_MACROPHAGE_DN, GSE7568_IL4_VS_IL4_AND_DEXAMETHASONE_TREATED_MACROPHAGE_DN
GSE15330_LYMPHOID_MULTIPOTENT_VS_PRO_BCELL_UP, GSE15330_LYMPHOID_MULTIPOTENT_VS_PRO_BCELL_UP
CROONQUIST_NRAS_VS_STROMAL_STIMULATION_UP, CROONQUIST_NRAS_VS_STROMAL_STIMULATION_UP
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_DN, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_DN
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_DN, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_DN
GSE9946_LISTERIA_INF_MATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP, GSE9946_LISTERIA_INF_MATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP
GO_ENDOSOME_ORGANIZATION, GO_ENDOSOME_ORGANIZATION
PID_SMAD2_3NUCLEAR_PATHWAY, PID_SMAD2_3NUCLEAR_PATHWAY
GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_MAC_UP, GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_MAC_UP
SIG_BCR_SIGNALING_PATHWAY, SIG_BCR_SIGNALING_PATHWAY
GSE43955_10H_VS_60H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN, GSE43955_10H_VS_60H_ACT_CD4_TCELL_WITH_TGFB_IL6_DN
GO_INTERCELLULAR_BRIDGE, GO_INTERCELLULAR_BRIDGE
KEGG_T_CELL_RECEPTOR_SIGNALING_PATHWAY, KEGG_T_CELL_RECEPTOR_SIGNALING_PATHWAY
BURTON_ADIPOGENESIS_I1, BURTON_ADIPOGENESIS_I1
GO_PROTEIN_TARGETING_TO_LYSOSOME, GO_PROTEIN_TARGETING_TO_LYSOSOME
GSE17721_LPS_VS_PAM3CSK4_0.5H_BMDC_UP, GSE17721_LPS_VS_PAM3CSK4_0.5H_BMDC_UP
GO_EXTRINSIC_COMPONENT_OF_CYTOPLASMIC_SIDE_OF_PLASMA_MEMBRANE, GO_EXTRINSIC_COMPONENT_OF_CYTOPLASMIC_SIDE_OF_PLASMA_MEMBRANE
GSE19888_ADENOSINE_A3R_ACT_VS_A3R_ACT_WITH_A3R_INH_PRETREATMENT_IN_M, GSE19888_ADENOSINE_A3R_ACT_VS_A3R_ACT_WITH_A3R_INH_PRETREATMENT_IN_M
GO_BETA_CATENIN_BINDING, GO_BETA_CATENIN_BINDING
GSE16385_ROSIGLITAZONE_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP, GSE16385_ROSIGLITAZONE_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP
GO_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI, GO_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI
BIOCARTA_STRESS_PATHWAY, BIOCARTA_STRESS_PATHWAY
GSE3203_UNTREATED_VS_IFNB_TREATED_LN_BCELL_UP, GSE3203_UNTREATED_VS_IFNB_TREATED_LN_BCELL_UP
YRTCANNRCCG_UNKNOWN, YRTCANNRCCG_UNKNOWN
GO_TISSUE_REMODELING, GO_TISSUE_REMODELING
KEGG_CHEMOKINE_SIGNALING_PATHWAY, KEGG_CHEMOKINE_SIGNALING_PATHWAY
KEGG_ENDOMETRIAL_CANCER, KEGG_ENDOMETRIAL_CANCER
GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_DN, GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_DN
KEGG_PROSTATE_CANCER, KEGG_PROSTATE_CANCER
GO_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_BINDING, GO_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_BINDING
ZHAN_MULTIPLE_MYELOMA_CD1_AND_CD2_DN, ZHAN_MULTIPLE_MYELOMA_CD1_AND_CD2_DN
GO_POSITIVE_REGULATION_OF_RNA_SPLICING, GO_POSITIVE_REGULATION_OF_RNA_SPLICING
ST_T_CELL_SIGNAL_TRANSDUCTION, ST_T_CELL_SIGNAL_TRANSDUCTION
GO_REGULATION_OF_NEURON_APOPTOTIC_PROCESS, GO_REGULATION_OF_NEURON_APOPTOTIC_PROCESS
GO_SIGNAL_TRANSDUCTION_IN_ABSENCE_OF_LIGAND, GO_SIGNAL_TRANSDUCTION_IN_ABSENCE_OF_LIGAND
BIOCARTA_JGFIMTOR_PATHWAY, BIOCARTA_JGFIMTOR_PATHWAY
GO_SNRNA_BINDING, GO_SNRNA_BINDING
GSE2826_WT_VS_XID_BCELL_DN, GSE2826_WT_VS_XID_BCELL_DN
PID_TAP63_PATHWAY, PID_TAP63_PATHWAY
GO_POSITIVE_REGULATION_OF_IMMUNE_EFFECTOR_PROCESS, GO_POSITIVE_REGULATION_OF_IMMUNE_EFFECTOR_PROCESS
REACTOME_SIGNALING_BY_NOTCH1, REACTOME_SIGNALING_BY_NOTCH1
GO_REGULATION_OF_ERBB_SIGNALING_PATHWAY, GO_REGULATION_OF_ERBB_SIGNALING_PATHWAY
REACTOME_SMAD2_SMAD3_SMAD4_HETEROTRIMER_REGULATES_TRANSCRIPTION, REACTOME_SMAD2_SMAD3_SMAD4_HETEROTRIMER_REGULATES_TRANSCRIPTION
GO_POSITIVE_REGULATION_OF_NEURON_DEATH, GO_POSITIVE_REGULATION_OF_NEURON_DEATH
GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE
GO_NEGATIVE_REGULATION_OF_SEQUENCE_SPECIFIC_DNA_BINDING_TRAN, GO_NEGATIVE_REGULATION_OF_SEQUENCE_SPECIFIC_DNA_BINDING_TRAN
GO_GASTRULATION_WITH_MOUTH_FORMING_SECOND, GO_GASTRULATION_WITH_MOUTH_FORMING_SECOND
GO_PROTEIN_LOCALIZATION_TO_VACUOLE, GO_PROTEIN_LOCALIZATION_TO_VACUOLE
GO_T_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE, GO_T_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE
GO_CELLULAR_RESPONSE_TO_EPIDERMAL_GROWTH_FACTOR_STIMULUS, GO_CELLULAR_RESPONSE_TO_EPIDERMAL_GROWTH_FACTOR_STIMULUS
GO_P53_BINDING, GO_P53_BINDING
GO_PHOSPHATIDYLINOSITOL_PHOSPHATE_BINDING, GO_PHOSPHATIDYLINOSITOL_PHOSPHATE_BINDING
GO_RESPONSE_TO_MONOAMINE, GO_RESPONSE_TO_MONOAMINE
SCHEIDERIT_JKK_TARGETS, SCHEIDERIT_JKK_TARGETS
SIG_IL4RECEPTOR_IN_B_LYPHOCYTES, SIG_IL4RECEPTOR_IN_B_LYPHOCYTES
PID_DELTA_NP63_PATHWAY, PID_DELTA_NP63_PATHWAY
BIOCARTA_GSK3_PATHWAY, BIOCARTA_GSK3_PATHWAY
BAUS_TFF2_TARGETS_UP, BAUS_TFF2_TARGETS_UP
GO_BLASTOCYST_GROWTH, GO_BLASTOCYST_GROWTH
KANG_DOXORUBICIN_RESISTANCE_DN, KANG_DOXORUBICIN_RESISTANCE_DN
GSE22045_TREG_VS_TCONV_UP, GSE22045_TREG_VS_TCONV_UP
GO_POSITIVE_REGULATION_OF_NEURON_APOPTOTIC_PROCESS, GO_POSITIVE_REGULATION_OF_NEURON_APOPTOTIC_PROCESS
GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP, GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP
VALK_AML_CLUSTER_1, VALK_AML_CLUSTER_1
ST_GAQ_PATHWAY, ST_GAQ_PATHWAY
KEGG_MTOR_SIGNALING_PATHWAY, KEGG_MTOR_SIGNALING_PATHWAY
KYNQ_DNA_DAMAGE_BY_4NQO_OR_UV, KYNQ_DNA_DAMAGE_BY_4NQO_OR_UV
HUMMEL_BURKITTs_LYMPHOMA_DN, HUMMEL_BURKITTs_LYMPHOMA_DN
GO_BLASTOCYST_DEVELOPMENT, GO_BLASTOCYST_DEVELOPMENT
GO_NEGATIVE_REGULATION_OF_BINDING, GO_NEGATIVE_REGULATION_OF_BINDING
ARGGGTTAA_UNKNOWN, ARGGGTTAA_UNKNOWN
GO_MANGANESE_ION_BINDING, GO_MANGANESE_ION_BINDING
GNF2_LYN, GNF2_LYN
RASHI_NFKB1_TARGETS, RASHI_NFKB1_TARGETS
GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_UP, GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_UP
WARTERS_IR_RESPONSE_SGY, WARTERS_IR_RESPONSE_SGY
GSE20500_RETINOIC_ACID_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP, GSE20500_RETINOIC_ACID_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP
GO_PROTEIN_LOCALIZATION_TO_LYSOSOME, GO_PROTEIN_LOCALIZATION_TO_LYSOSOME
MULLIGAN_NTF3_SIGNALING_VIA_INSR_AND_IGFIR_UP, MULLIGAN_NTF3_SIGNALING_VIA_INSR_AND_IGFIR_UP
KIM_GASTRIC_CANCER_CHEMOSENSITIVITY, KIM_GASTRIC_CANCER_CHEMOSENSITIVITY
BIOCARTA_KERATINOCYTE_PATHWAY, BIOCARTA_KERATINOCYTE_PATHWAY
GO_INNER_CELL_MASS_CELL_PROLIFERATION, GO_INNER_CELL_MASS_CELL_PROLIFERATION
GO_NEGATIVE_REGULATION_OF_NEURAL_PRECURSOR_CELL_PROLIFERATION, GO_NEGATIVE_REGULATION_OF_NEURAL_PRECURSOR_CELL_PROLIFERATION
BIOCARTA_PTEN_PATHWAY, BIOCARTA_PTEN_PATHWAY
REACTOME_PKB_CASCADE, REACTOME_PKB_CASCADE
GO_POSITIVE_REGULATION_OF_ORGANELLE_ASSEMBLY, GO_POSITIVE_REGULATION_OF_ORGANELLE_ASSEMBLY
PID_FAS_PATHWAY, PID_FAS_PATHWAY
GO_CANONICAL_WNT_SIGNALING_PATHWAY, GO_CANONICAL_WNT_SIGNALING_PATHWAY
GO_PROTEIN_K63_LINKED_DEUBIQUITINATION, GO_PROTEIN_K63_LINKED_DEUBIQUITINATION
GO_RNA_POLYMERASE_II_REPRESSING_TRANSCRIPTION_FACTOR_BINDING, GO_RNA_POLYMERASE_II_REPRESSING_TRANSCRIPTION_FACTOR_BINDING
PID_CD40_PATHWAY, PID_CD40_PATHWAY
GO_REGULATION_OF_EPIDERMAL_GROWTH_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY, GO_REGULATION_OF_EPIDERMAL_GROWTH_FACTOR_ACTIVATED_RECEPTOR_ACTIVITY
BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_24HR_UP, BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_24HR_UP
GSE17721_CTRL_VS_POLYIC_6H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_6H_BMDC_DN
KEGG_NON_SMALL_CELL_LUNG_CANCER, KEGG_NON_SMALL_CELL_LUNG_CANCER
GO_POSITIVE_REGULATION_OF_PEPTIDYL_THREONINE_PHOSPHORYLATION, GO_POSITIVE_REGULATION_OF_PEPTIDYL_THREONINE_PHOSPHORYLATION
KEGG_RIG_I_LIKE_RECEPTOR_SIGNALING_PATHWAY, KEGG_RIG_I_LIKE_RECEPTOR_SIGNALING_PATHWAY
TATCTGG_MIR488, TATCTGG_MIR488
PID_MYC_PATHWAY, PID_MYC_PATHWAY
GO_RRNA_TRANSCRIPTION, GO_RRNA_TRANSCRIPTION
WILLIAMS_ESR2_TARGETS_DN, WILLIAMS_ESR2_TARGETS_DN