

	GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_DN, GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_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 GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_DN, GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_DN GSE36527_CD69_NEG_VS_POS_TREG_CD62L_LOS_KLRG1_NEG_DN, GSE36527_CD69_NEG_VS_POS_TREG_CD62L_LOS_KLRG1_NEG_DN GSE36009_WT_VS_NLRP10_KO_DC_LPS_STIM_DN, GSE36009_WT_VS_NLRP10_KO_DC_LPS_STIM_DN GSE17721_PAM3CSK4_VS_GADIQUIMOD_2H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_2H_BMDC_DN GSE37416_0H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN, GSE37416_0H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN GSE40274_EOS_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_EOS_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP GSE32164_RESTING_DIFFERENTIATED_VS_ALTERNATIVELY_ACT_M2_MACROPHAGE_DN, GSE32164_RESTING_DIFFERENTIATED_VS_ALTERNATIVELY_ACT_M2_MACROPHAGE_DN GSE17721_CPG_VS_GARDIQUIMOD_8H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_8H_BMDC_DN GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_UP, GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_UP GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_DN, GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_DN GSE3982_BASOPHIL_VS_CENT_MEMORY_CD4_TCELL_UP, GSE3982_BASOPHIL_VS_CENT_MEMORY_CD4_TCELL_UP GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN, GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN GO_VACUOLE_ORGANIZATION, GO_VACUOLE_ORGANIZATION GSE1925_CTRL_VS_24H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN, GSE1925_CTRL_VS_24H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN HAHTOLA_MYCOSIS_FUNGOIDES_SKIN_UP, HAHTOLA_MYCOSIS_FUNGOIDES_SKIN_UP GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN GSE1460_NAIVE_CD4_TCELL_ADULT_BLOOD_VS_THYMIC_STROMAL_CELL_UP, GSE1460_NAIVE_CD4_TCELL_ADULT_BLOOD_VS_THYMIC_STROMAL_CELL_UP GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP, GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP GSE1448_CTRL_VS_ANTLVBETA5_DP_THYMOCYTE_UP, GSE1448_CTRL_VS_ANTLVBETA5_DP_THYMOCYTE_UP GSE17721_LPS_VS_CPG_12H_BMDC_UP, GSE17721_LPS_VS_CPG_12H_BMDC_UP GSE27786_LIN_NEG_VS_NKTCELL_DN, GSE27786_LIN_NEG_VS_NKTCELL_DN GSE15330_WT_VS_IKAROS_KO_LYMPHOID_MULTIPOTENT_PROGENITOR_UP, GSE15330_WT_VS_IKAROS_KO_LYMPHOID_MULTIPOTENT_PROGENITOR_UP GSE17721_0.5H_VS_4H_GARDIQUIMOD_BMDC_DN, GSE17721_0.5H_VS_4H_GARDIQUIMOD_BMDC_DN GSE11973_MIR223_KOVS_WT_BONE_MARROW_NEUTROPHIL_UP, GSE11973_MIR223_KOVS_WT_BONE_MARROW_NEUTROPHIL_UP GSE37532_WT_VS_PPARG_KO_LN_TCONV_UP, GSE37532_WT_VS_PPARG_KO_LN_TCONV_UP GSE17721_LPS_VS_CPG_4H_BMDC_DN, GSE17721_LPS_VS_CPG_4H_BMDC_DN GSE17721_0.5H_VS_12H_POLYIC_BMDC_DN, GSE17721_0.5H_VS_12H_POLYIC_BMDC_DN GSE15330_WT_VS_IKAROS_KO_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN, GSE15330_WT_VS_IKAROS_KO_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN GSE27786_LSK_VS_CD8_TCELL_DN, GSE27786_LSK_VS_CD8_TCELL_DN GSE19401_UNSTIM_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN, GSE19401_UNSTIM_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_UP, GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_UP GSE1448_ANTLVALPHA2_VS_VBETA5_DP_THYMOCYTE_UP, GSE1448_ANTLVALPHA2_VS_VBETA5_DP_THYMOCYTE_UP GSE28237_EARLY_VS_LATE_GC_BCELL_DN, GSE28237_EARLY_VS_LATE_GC_BCELL_DN GSE24142_ADULT_VS_FETAL_EARLY_THYMIC_PROGENITOR_UP, GSE24142_ADULT_VS_FETAL_EARLY_THYMIC_PROGENITOR_UP GSE16385_ROSIGLITAZONE_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP, GSE16385_ROSIGLITAZONE_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_UP, GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_UP 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 GSE39152_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_UP, GSE39152_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_UP GSE43955_TH0_VS_TGFB_IL6_IL23_THI7_ACT_CD4_TCELL_52H_DN, GSE43955_TH0_VS_TGFB_IL6_IL23_THI7_ACT_CD4_TCELL_52H_DN GSE37301_HEMATOPOIETIC_STEM_CELL_VS_MULTIPOTENT_PROGENITOR_DN, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_MULTIPOTENT_PROGENITOR_DN GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_DN, GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_DN GSE21670_TGFB_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_DN, GSE21670_TGFB_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_DN GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_UP, GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_UP GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY3_POST_IMMUNIZATION_UP, GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY3_POST_IMMUNIZATION_UP GSE33424_CD161_HIGH_VS_NEG_CD8_TCELL_UP, GSE33424_CD161_HIGH_VS_NEG_CD8_TCELL_UP PID_KIT_PATHWAY, PID_KIT_PATHWAY GSE17721_LPS_VS_GARDIQUIMOD_0.5H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_0.5H_BMDC_DN GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_UP, GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_UP GSE17721_POLYIC_VS_GARDIQUIMOD_24H_BMDC_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_24H_BMDC_UP GSE17721_CTRL_VS_POLYIC_24H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_24H_BMDC_DN YAGL_AML_SURVIVAL, YAGL_AML_SURVIVAL GSE17721_POLYIC_VS_GARDIQUIMOD_1H_BMDC_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_1H_BMDC_DN GO_POSITIVE_REGULATION_OF_MULTL_ORGANISM_PROCESS, GO_POSITIVE_REGULATION_OF_MULTL_ORGANISM_PROCESS GSE20715_0H_VS_6H_OZONE_TLR4_KO_LUNG_DN, GSE20715_0H_VS_6H_OZONE_TLR4_KO_LUNG_DN GO_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI, GO_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_DN, GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_DN GSE41087_WT_VS_FOXP3_MUT_ANTL_CD3_CD28_STIM_CD4_TCELL_UP, GSE41087_WT_VS_FOXP3_MUT_ANTL_CD3_CD28_STIM_CD4_TCELL_UP GGCACTT_MIR519E, GGCACTT_MIR519E GSE42021_TCONV_PLN_VS_CD24INT_TCONV_THYMUS_UP, GSE42021_TCONV_PLN_VS_CD24INT_TCONV_THYMUS_UP GNF2_RAP1B, GNF2_RAP1B GO_LYSOSOMAL_TRANSPORT, GO_LYSOSOMAL_TRANSPORT ETF_Q6, ETF_Q6 AACTGAC_MIR223, AACTGAC_MIR223 GO_LATE_ENDOSOME_MEMBRANE, GO_LATE_ENDOSOME_MEMBRANE GSE17721_CTRL_VS_GARDIQUIMOD_0.5H_BMDC_DN, GSE17721_CTRL_VS_GARDIQUIMOD_0.5H_BMDC_DN GSE360_CTRL_VS_T_GONDIL_DC_DN, GSE360_CTRL_VS_T_GONDIL_DC_DN GO_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION, GO_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION GO_ENDOSOME_TO_LYSOSOME_TRANSPORT, GO_ENDOSOME_TO_LYSOSOME_TRANSPORT GSE3982_EOSINOPHIL_VS_TH1_UP, GSE3982_EOSINOPHIL_VS_TH1_UP GSE19888_NO_PRETREAT_VS_ADENOSINE_A3R_INHIBITOR_PRETREATED_MAST_CELL_TCELL_MEMBRANES_ACT_UP, GSE19888_NO_PRETREAT_VS_ADENOSINE_A3R_INHIBITOR_PRETREATED_MAST_CELL_TCELL_MEMBRANES_ACT_UP GSE17721_PAM3CSK4_VS_GADIQUIMOD_16H_BMDC_UP, GSE17721_PAM3CSK4_VS_GADIQUIMOD_16H_BMDC_UP GSE15330_LYMPHOID_MULTIPOTENT_VS_PRO_BCELL_UP, GSE15330_LYMPHOID_MULTIPOTENT_VS_PRO_BCELL_UP GSE3920_IFNA_VS_IFNB_TREATED_ENDOTHELIAL_CELL_DN, GSE3920_IFNA_VS_IFNB_TREATED_ENDOTHELIAL_CELL_DN GO_REGULATION_OF_VACUOLE_ORGANIZATION, GO_REGULATION_OF_VACUOLE_ORGANIZATION GSE3982_MAC_VS_NEUTROPHIL_DN, GSE3982_MAC_VS_NEUTROPHIL_DN GO_POSITIVE_REGULATION_OF_IMMUNE_EFFECTOR_PROCESS, GO_POSITIVE_REGULATION_OF_IMMUNE_EFFECTOR_PROCESS AAGGGAT_MIR188, AAGGGAT_MIR188 GO_GDP_BINDING, GO_GDP_BINDING GO_PROTEIN_LOCALIZATION_TO_VACUOLE, GO_PROTEIN_LOCALIZATION_TO_VACUOLE GSE41176_WT_VS_TAK1_KO_UNSTIM_BCELL_UP, GSE41176_WT_VS_TAK1_KO_UNSTIM_BCELL_UP GSE1925_CTRL_VS_3H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN, GSE1925_CTRL_VS_3H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP GCTCTTG_MIR335, GCTCTTG_MIR335 WARTERS_IR_RESPONSE_5GY, WARTERS_IR_RESPONSE_5GY GSE43955_10H_VS_30H_ACT_CD4_TCELL_DN, GSE43955_10H_VS_30H_ACT_CD4_TCELL_DN GO_MANGANESE_ION_BINDING, GO_MANGANESE_ION_BINDING PID_FAS_PATHWAY, PID_FAS_PATHWAY REACTOME_PKB_MEDIATED_EVENTS, REACTOME_PKB_MEDIATED_EVENTS RIZ_ERYTHROID_DIFFERENTIATION_CCNE1, RIZ_ERYTHROID_DIFFERENTIATION_CCNE1 GTATGAT_MIR154_MIR487, GTATGAT_MIR154_MIR487 GO_EXTRINSIC_COMPONENT_OF_PLASMA_MEMBRANE, GO_EXTRINSIC_COMPONENT_OF_PLASMA_MEMBRANE GSE13485_PRE_VS_POST_YF17D_VACCINATION_PBMG_UP, GSE13485_PRE_VS_POST_YF17D_VACCINATION_PBMG_UP GO_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY, GO_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY GSE22935_UNSTIM_VS_48H_MBOVIS_BCG_STIM_MYPD88_KO_MACROPHAGE_UP, GSE22935_UNSTIM_VS_48H_MBOVIS_BCG_STIM_MYPD88_KO_MACROPHAGE_UP GO_ENDOSOME_ORGANIZATION, GO_ENDOSOME_ORGANIZATION PID_TGFBR_PATHWAY, PID_TGFBR_PATHWAY GO_TISSUE_REMODELING, GO_TISSUE_REMODELING KEGG_MTOR_SIGNALING_PATHWAY, KEGG_MTOR_SIGNALING_PATHWAY GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_DN, GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_DN BIOCARTA_KERATINOCYTE_PATHWAY, BIOCARTA_KERATINOCYTE_PATHWAY GSE43955_TH0_VS_TGFB_IL6_THI7_ACT_CD4_TCELL_60H_DN, GSE43955_TH0_VS_TGFB_IL6_THI7_ACT_CD4_TCELL_60H_DN GO_NEGATIVE_REGULATION_OF_BINDING, GO_NEGATIVE_REGULATION_OF_BINDING PID_TAP63_PATHWAY, PID_TAP63_PATHWAY GO_POSITIVE_REGULATION_OF_ORGANELLE_ASSEMBLY, GO_POSITIVE_REGULATION_OF_ORGANELLE_ASSEMBLY GO_INTERCELLULAR_BRIDGE, GO_INTERCELLULAR_BRIDGE KANG_DOXORUBICIN_RESISTANCE_DN, KANG_DOXORUBICIN_RESISTANCE_DN MORLMATURE_B_LYMPHOCYTE_UP, MORLMATURE_B_LYMPHOCYTE_UP LIN_MELANOMA_COPY_NUMBER_DN, LIN_MELANOMA_COPY_NUMBER_DN GO_EARLY_ENDOSOME_MEMBRANE, GO_EARLY_ENDOSOME_MEMBRANE REACTOME_SIGNALING_BY_NOTCH1, REACTOME_SIGNALING_BY_NOTCH1 ST_T_CELL_SIGNAL_TRANSDUCTION, ST_T_CELL_SIGNAL_TRANSDUCTION ST_PHOSPHOINOSITIDE_3_KINASE_PATHWAY, ST_PHOSPHOINOSITIDE_3_KINASE_PATHWAY GO_REGULATION_OF_CELLULAR_EXTRAVASATION, GO_REGULATION_OF_CELLULAR_EXTRAVASATION GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY GO_RETROMER_COMPLEX, GO_RETROMER_COMPLEX GO_POSITIVE_REGULATION_OF_NEURON_DEATH, GO_POSITIVE_REGULATION_OF_NEURON_DEATH chr2p22, chr2p22 GO_REGULATION_OF_ERBB_SIGNALING_PATHWAY, GO_REGULATION_OF_ERBB_SIGNALING_PATHWAY IWANAGA_CARCIINOGENESIS_BY_KRAS_PTEIN_UP, IWANAGA_CARCIINOGENESIS_BY_KRAS_PTEIN_UP PID_IL8_CXCR2_PATHWAY, PID_IL8_CXCR2_PATHWAY GO_REGULATION_OF_MAST_CELL_ACTIVATION, GO_REGULATION_OF_MAST_CELL_ACTIVATION KEEN_RESPONSE_TO_ROSIGLITAZONE_DN, KEEN_RESPONSE_TO_ROSIGLITAZONE_DN GO_REGULATION_OF_CHOLESTEROL_EFFLUX, GO_REGULATION_OF_CHOLESTEROL_EFFLUX SIG_BCR_SIGNALING_PATHWAY, SIG_BCR_SIGNALING_PATHWAY GO_EXTRINSIC_COMPONENT_OF_CYTOPLASMIC_SIDE_OF_PLASMA_MEMBRANE, GO_EXTRINSIC_COMPONENT_OF_CYTOPLASMIC_SIDE_OF_PLASMA_MEMBRANE GO_P53_BINDING, GO_P53_BINDING GO_RECEPTOR_CATABOLIC_PROCESS, GO_RECEPTOR_CATABOLIC_PROCESS GSE15659_TREG_VS_TCONV_UP, GSE15659_TREG_VS_TCONV_UP GO_SIGNAL_TRANSDUCTION_IN_ABSENCE_OF_LIGAND, GO_SIGNAL_TRANSDUCTION_IN_ABSENCE_OF_LIGAND GSE45365_NK_CELL_VS_CD8_TCELL_DN, GSE45365_NK_CELL_VS_CD8_TCELL_DN MARIADASON_REGULATED_BY_HISTONE_ACETYLATION_UP, MARIADASON_REGULATED_BY_HISTONE_ACETYLATION_UP GO_VESICLE_MEDIATED_TRANSPORT_BETWEEN_ENDOSOMAL_COMPARTMENTS, GO_VESICLE_MEDIATED_TRANSPORT_BETWEEN_ENDOSOMAL_COMPARTMENTS GGCCATT_MIR365, GGCCATT_MIR365 GO_GASTRULATION, GO_GASTRULATION RASHI_NFKB1_TARGETS, RASHI_NFKB1_TARGETS WILLIAMS_ESR2_TARGETS_DN, WILLIAMS_ESR2_TARGETS_DN GO_REGULATION_OF_LIPID_STORAGE, GO_REGULATION_OF_LIPID_STORAGE ACATATC_MIR190, ACATATC_MIR190 VALK_AML_CLUSTER_1, VALK_AML_CLUSTER_1 MARSON_FOXP3_CORE_DIRECT_TARGETS, MARSON_FOXP3_CORE_DIRECT_TARGETS CROONQUIST_STROMAL_STIMULATION_DN, CROONQUIST_STROMAL_STIMULATION_DN RASHI_RESPONSE_TO_IONIZING_RADIATION_6, RASHI_RESPONSE_TO_IONIZING_RADIATION_6 KEGG_ENDOMETRIAL_CANCER, KEGG_ENDOMETRIAL_CANCER BIOCARTA_GSK3_PATHWAY, BIOCARTA_GSK3_PATHWAY GO_REGULATION_OF_VIRAL_RELEASE_FROM_HOST_CELL, GO_REGULATION_OF_VIRAL_RELEASE_FROM_HOST_CELL POS_RESPONSE_TO_HISTAMINE_UP, POS_RESPONSE_TO_HISTAMINE_UP ST_GAQ_PATHWAY, ST_GAQ_PATHWAY GO_REGULATION_OF_NEURON_APOPTOTIC_PROCESS, GO_REGULATION_OF_NEURON_APOPTOTIC_PROCESS GO_SCF_UBIQUITIN_LIGASE_COMPLEX, GO_SCF_UBIQUITIN_LIGASE_COMPLEX SIG_IL4RECEPTOR_IN_B_LYPHOCYTES, SIG_IL4RECEPTOR_IN_B_LYPHOCYTES GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_MAC_UP, GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_MAC_UP PID_IL8_CXCR1_PATHWAY, PID_IL8_CXCR1_PATHWAY GO_NEGATIVE_REGULATION_OF_SEQUENCE_SPECIFIC_DNA_BINDING_TRANSCRIPTION_FACTOR_ACTIVITY, GO_NEGATIVE_REGULATION_OF_SEQUENCE_SPECIFIC_DNA_BINDING_TRANSCRIPTION_FACTOR_ACTIVITY GNF2_LYN, GNF2_LYN BRUNO_HEMATOPOIESIS, BRUNO_HEMATOPOIESIS GO_RESPONSE_TO_PROSTAGLANDIN_E, GO_RESPONSE_TO_PROSTAGLANDIN_E GO_CANONICAL_WNT_SIGNALING_PATHWAY, GO_CANONICAL_WNT_SIGNALING_PATHWAY GO_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY, GO_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY BIOCARTA_PTEIN_PATHWAY, BIOCARTA_PTEIN_PATHWAY CROONQUIST_NRAS_VS_STROMAL_STIMULATION_UP, CROONQUIST_NRAS_VS_STROMAL_STIMULATION_UP WANG_RESPONSE_TO_ANDROGEN_UP, WANG_RESPONSE_TO_ANDROGEN_UP PID_DELTA_Np63_PATHWAY, PID_DELTA_Np63_PATHWAY KEGG_NON_SMALL_CELL_LUNG_CANCER, KEGG_NON_SMALL_CELL_LUNG_CANCER KEGG_CHEMOKINE_SIGNALING_PATHWAY, KEGG_CHEMOKINE_SIGNALING_PATHWAY GO_REGULATION_OF_MYELINATION, GO_REGULATION_OF_MYELINATION PID_NOTCH_PATHWAY, PID_NOTCH_PATHWAY GO_NEGATIVE_REGULATION_OF_NF_KAPPAB_TRANSCRIPTION_FACTOR_ACTIVITY, GO_NEGATIVE_REGULATION_OF_NF_KAPPAB_TRANSCRIPTION_FACTOR_ACTIVITY GO_NEGATIVE_REGULATION_OF_NOTCH_SIGNALING_PATHWAY, GO_NEGATIVE_REGULATION_OF_NOTCH_SIGNALING_PATHWAY
3_KO_4H_LPS_STIM_MACROPHAGE_UP, GSE33162_HDAC3_KO_VS_HDAC3_KO_4H_LPS_STIM_MACROPHAGE_UP	