

3_KO_4H_LPS_STIM_MACROPHAGE_UP, GSE33162_HDAC3_KO_VS_HDAC3_KO_4H_LPS_STIM_MACROPHAGE_UP

GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_DN, GSE34006_A2AR_KO_VS_A2AR_AGONIST_TREATED_TREG_DN
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
GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP, GSE7348_LPS_VS_TOLERIZED_AND_LPS_STIM_MACROPHAGE_UP
GSE10094_LCMV_VS_LISTERIA_IND_EFF_CD4_TCELL_UP, GSE10094_LCMV_VS_LISTERIA_IND_EFF_CD4_TCELL_UP
GSE3982_BASOPHIL_VS_CENT_MEMORY_CD4_TCELL_UP, GSE3982_BASOPHIL_VS_CENT_MEMORY_CD4_TCELL_UP
GSE32034_LY6C_HIGH_VS_LOW_MONOCYTE_DN, GSE32034_LY6C_HIGH_VS_LOW_MONOCYTE_DN
GSE17721_0.5H_VS_24H_CPG_BMDC_UP, GSE17721_0.5H_VS_24H_CPG_BMDC_UP
GSE1925_CTRL_VS_24H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN, GSE1925_CTRL_VS_24H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN
GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP, GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP
GSE17721_PAM3CSK4_VS_GADIQUIMOD_2H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_2H_BMDC_DN
GSE15330_WT_VS_IKAROS_KO_LYMPHOID_MULTIPOTENT_PROGENITOR_UP, GSE15330_WT_VS_IKAROS_KO_LYMPHOID_MULTIPOTENT_PROGENITOR_UP
GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_DN, GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_DN
GSE27786_LIN_NEG_VS_NKTCELL_DN, GSE27786_LIN_NEG_VS_NKTCELL_DN
GSE1448_CTRL_VS_ANTI_VBETA5_DP_THYMOCYTE_UP, GSE1448_CTRL_VS_ANTI_VBETA5_DP_THYMOCYTE_UP
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
GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_DN, GSE32164_RESTING_DIFFERENTIATED_VS_CMYC_INHIBITED_MACROPHAGE_DN
GSE17721_CTRL_VS_LPS_1H_BMDC_UP, GSE17721_CTRL_VS_LPS_1H_BMDC_UP
GSE11973_MIR223_KOV5_WT_BONE_MARROW_NEUTROPHIL_UP, GSE11973_MIR223_KOV5_WT_BONE_MARROW_NEUTROPHIL_UP
GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN, GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN
GSE15330_WT_VS_IKAROS_KO_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN, GSE15330_WT_VS_IKAROS_KO_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN
GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_DN, GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_DN
GSE40274_EOS_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_EOS_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
GSE17721_CPG_VS_GARDIQUIMOD_8H_BMDC_DN, GSE17721_CPG_VS_GARDIQUIMOD_8H_BMDC_DN
GSE28237_EARLY_VS_LATE_GC_BCELL_DN, GSE28237_EARLY_VS_LATE_GC_BCELL_DN
GSE42021_TCONV_PLN_VS_CD24INT_TCONV_THYMUS_UP, GSE42021_TCONV_PLN_VS_CD24INT_TCONV_THYMUS_UP
GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_UP, GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_UP
GSE36009_WT_VS_NLRP10_KO_DC_LPS_STIM_DN, GSE36009_WT_VS_NLRP10_KO_DC_LPS_STIM_DN
GSE4748_LPS_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP, GSE4748_LPS_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP
GSE17721_0.5H_VS_4H_GARDIQUIMOD_BMDC_DN, GSE17721_0.5H_VS_4H_GARDIQUIMOD_BMDC_DN
GSE17721_LPS_VS_CPG_12H_BMDC_UP, GSE17721_LPS_VS_CPG_12H_BMDC_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
GCM_ERBB2IP, GCM_ERBB2IP
GO_LATE_ENDOSOME_MEMBRANE, GO_LATE_ENDOSOME_MEMBRANE
GSE27786_LSK_VS_CD8_TCELL_DN, GSE27786_LSK_VS_CD8_TCELL_DN
GSE1460_NAIVE_CD4_TCELL_ADULT_BLOOD_VS_THYMIC_STROMAL_CELL_UP, GSE1460_NAIVE_CD4_TCELL_ADULT_BLOOD_VS_THYMIC_STROMAL_CELL_UP
GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_UP, GSE14415_INDUCED_TREG_VS_FOXP3_KO_INDUCED_TREG_IL2_CULTURE_UP
GSE22886_DAY0_VS_DAY1_MONOCYTE_IN_CULTURE_DN, GSE22886_DAY0_VS_DAY1_MONOCYTE_IN_CULTURE_DN
GSE41087_WT_VS_FOXP3_MUT_ANTI_CD3_CD28_STIM_CD4_TCELL_UP, GSE41087_WT_VS_FOXP3_MUT_ANTI_CD3_CD28_STIM_CD4_TCELL_UP
GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_DN, GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_DN
GSE19401_UNSTIM_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN, GSE19401_UNSTIM_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN
GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_4H_UP, GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_4H_UP
GSE37301_HEMATOPOIETIC_STEM_CELL_VS_MULTIPOTENT_PROGENITOR_DN, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_MULTIPOTENT_PROGENITOR_DN
GSE24142_ADULT_VS_FETAL_EARLY_THYMIC_PROGENITOR_UP, GSE24142_ADULT_VS_FETAL_EARLY_THYMIC_PROGENITOR_UP
GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_UP, GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_UP
GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_UP, GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_UP
GSE17721_CTRL_VS_POLYIC_24H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_24H_BMDC_DN
KEGG_NEUROTROPHIN_SIGNALING_PATHWAY, KEGG_NEUROTROPHIN_SIGNALING_PATHWAY
GSE21670_TGFB_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_DN, GSE21670_TGFB_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_DN
GO_AUTOPHAGOSOME, GO_AUTOPHAGOSOME
GSE27786_LSK_VS_BCELL_DN, GSE27786_LSK_VS_BCELL_DN
GSE7852_LN_VS_FAT_TCONV_DN, GSE7852_LN_VS_FAT_TCONV_DN
GSE17721_POLYIC_VS_GARDIQUIMOD_24H_BMDC_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_24H_BMDC_UP
GGCACTT_MIR519E, GGCACTT_MIR519E
GSE17721_PAM3CSK4_VS_CPG_0.5H_BMDC_UP, GSE17721_PAM3CSK4_VS_CPG_0.5H_BMDC_UP
GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY3_POST_IMMUNIZATION_UP, GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY3_POST_IMMUNIZATION_UP
GSE15624_3H_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN, GSE15624_3H_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN
GSE360_CTRL_VS_T_GONDII_DC_DN, GSE360_CTRL_VS_T_GONDII_DC_DN
GSE22935_UNSTIM_VS_48H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_UP, GSE22935_UNSTIM_VS_48H_MBOVIS_BCG_STIM_MYD88_KO_MACROPHAGE_UP
GSE17721_CPG_VS_GARDIQUIMOD_4H_BMDC_UP, GSE17721_CPG_VS_GARDIQUIMOD_4H_BMDC_UP
MORLMATURE_B_LYMPHOCYTE_UP, MORLMATURE_B_LYMPHOCYTE_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
GSE37532_WT_VS_PPARG_KO_LN_TCONV_UP, GSE37532_WT_VS_PPARG_KO_LN_TCONV_UP
GSE5503_MLN_DC_VS_PLN_DC_ACTIVATED_ALLOGENIC_TCELL_DN, GSE5503_MLN_DC_VS_PLN_DC_ACTIVATED_ALLOGENIC_TCELL_DN
GSE4748_CTRL_VS_LPS_STIM_DC_3H_UP, GSE4748_CTRL_VS_LPS_STIM_DC_3H_UP
GSE17721_CTRL_VS_GARDIQUIMOD_0.5H_BMDC_DN, GSE17721_CTRL_VS_GARDIQUIMOD_0.5H_BMDC_DN
GSE39152_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_UP, GSE39152_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_UP
GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_6H_UP
GSE17721_CTRL_VS_CPG_12H_BMDC_DN, GSE17721_CTRL_VS_CPG_12H_BMDC_DN
GO_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI, GO_RETROGRADE_TRANSPORT_ENDOSOME_TO_GOLGI
KEGG_MTOR_SIGNALING_PATHWAY, KEGG_MTOR_SIGNALING_PATHWAY
GSE37301_RAG2_KO_VS_RAG2_AND_ETSI_KO_NK_CELL_UP, GSE37301_RAG2_KO_VS_RAG2_AND_ETSI_KO_NK_CELL_UP
GO_P53_BINDING, GO_P53_BINDING
YRTCANNRCCG_UNKNOWN, YRTCANNRCCG_UNKNOWN
GO_RECYCLING_ENDOSOME, GO_RECYCLING_ENDOSOME
GO_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION, GO_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION
GO_ENDOSOME_ORGANIZATION, GO_ENDOSOME_ORGANIZATION
GO_POSITIVE_REGULATION_OF_MULTIORGANISM_PROCESS, GO_POSITIVE_REGULATION_OF_MULTIORGANISM_PROCESS
GSE17721_LPS_VS_GARDIQUIMOD_0.5H_BMDC_DN, GSE17721_LPS_VS_GARDIQUIMOD_0.5H_BMDC_DN
ETF_Q6, ETF_Q6
GSE42021_CD24HL_TREG_VS_CD24HL_TCONV_THYMUS_DN, GSE42021_CD24HL_TREG_VS_CD24HL_TCONV_THYMUS_DN
GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
GSE27241_CTRL_VS_DIGOXIN_TREATED_CD4_TCELL_IN_THI7_POLARIZING_CONDITIONS_UP, GSE27241_CTRL_VS_DIGOXIN_TREATED_CD4_TCELL_IN_THI7_POLARIZING_CONDITIONS_UP
REACTOME_PKB_MEDIATED_EVENTS, REACTOME_PKB_MEDIATED_EVENTS
GSE4142_PLASMA_CELL_VS_GC_BCELL_UP, GSE4142_PLASMA_CELL_VS_GC_BCELL_UP
KEGG_ENDOMETRIAL_CANCER, KEGG_ENDOMETRIAL_CANCER
GSE22611_UNSTIM_VS_6H_MDP_STIM_NOD2_TRANSDUCED_HEK293T_CELL_UP, GSE22611_UNSTIM_VS_6H_MDP_STIM_NOD2_TRANSDUCED_HEK293T_CELL_UP
GO_LYSOSOMAL_TRANSPORT, GO_LYSOSOMAL_TRANSPORT
GTATGAT_MIR154_MIR487, GTATGAT_MIR154_MIR487
KEGG_PROSTATE_CANCER, KEGG_PROSTATE_CANCER
KEEN_RESPONSE_TO_ROSIGLITAZONE_DN, KEEN_RESPONSE_TO_ROSIGLITAZONE_DN
GSE43955_TH0_VS_TGFB_IL6_THI7_ACT_CD4_TCELL_60H_DN, GSE43955_TH0_VS_TGFB_IL6_THI7_ACT_CD4_TCELL_60H_DN
GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP, GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP
KEGG_NON_SMALL_CELL_LUNG_CANCER, KEGG_NON_SMALL_CELL_LUNG_CANCER
SIG_BCR_SIGNALING_PATHWAY, SIG_BCR_SIGNALING_PATHWAY
PID_P13K_PLC_TRK_PATHWAY, PID_P13K_PLC_TRK_PATHWAY
GO_REGULATION_OF_VACUOLE_ORGANIZATION, GO_REGULATION_OF_VACUOLE_ORGANIZATION
GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_DN, GSE9037_CTRL_VS_LPS_1H_STIM_BMDM_DN
GNF2_RAP1B, GNF2_RAP1B
GSE23114_PERITONEAL_CAVITY_B1A_BCELL_VS_SPLEEN_BCELL_IN_SLE2C1_MOUSE_DN, GSE23114_PERITONEAL_CAVITY_B1A_BCELL_VS_SPLEEN_BCELL_IN_SLE2C1_MOUSE_DN
GSE16385_ROSIGLITAZONE_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP, GSE16385_ROSIGLITAZONE_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP
GO_EXTRINSIC_COMPONENT_OF_PLASMA_MEMBRANE, GO_EXTRINSIC_COMPONENT_OF_PLASMA_MEMBRANE
GO_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY, GO_REGULATION_OF_AUTOPHAGOSOME_ASSEMBLY
GO_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_BINDING, GO_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_BINDING
PID_TAP63_PATHWAY, PID_TAP63_PATHWAY
GO_ENDOSOME_TO_LYSOSOME_TRANSPORT, GO_ENDOSOME_TO_LYSOSOME_TRANSPORT
PID_FAS_PATHWAY, PID_FAS_PATHWAY
GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_UP, GSE22935_24H_VS_48H_MBOVIS_BCG_STIM_MACROPHAGE_UP
REACTOME_P13K_CASCADE, REACTOME_P13K_CASCADE
GSE20500_RETINOIC_ACID_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP, GSE20500_RETINOIC_ACID_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP
GSE2826_WT_VS_XID_BCELL_DN, GSE2826_WT_VS_XID_BCELL_DN
CROONQUIST_NRAS_VS_STROMAL_STIMULATION_UP, CROONQUIST_NRAS_VS_STROMAL_STIMULATION_UP
RIZ_ERYTHROID_DIFFERENTIATION_CCNE1, RIZ_ERYTHROID_DIFFERENTIATION_CCNE1
GO_NEGATIVE_REGULATION_OF_SEQUENCE_SPECIFIC_DNA_BINDING_TRANSCRIPTION_FACTOR_ACTIVITY, GO_NEGATIVE_REGULATION_OF_SEQUENCE_SPECIFIC_DNA_BINDING_TRANSCRIPTION_FACTOR_ACTIVITY
GO_POSITIVE_REGULATION_OF_IMMUNE_EFFECTOR_PROCESS, GO_POSITIVE_REGULATION_OF_IMMUNE_EFFECTOR_PROCESS
PID_KIT_PATHWAY, PID_KIT_PATHWAY
BONCI_TARGETS_OF_MIR15A_AND_MIR16_1, BONCI_TARGETS_OF_MIR15A_AND_MIR16_1
GSE4142_NAIVE_VS_GC_BCELL_UP, GSE4142_NAIVE_VS_GC_BCELL_UP
GO_POSITIVE_REGULATION_OF_NEURON_DEATH, GO_POSITIVE_REGULATION_OF_NEURON_DEATH
GO_REGULATION_OF_LEUKOCYTE_DEGRANULATION, GO_REGULATION_OF_LEUKOCYTE_DEGRANULATION
LIN_MELANOMA_COPY_NUMBER_DN, LIN_MELANOMA_COPY_NUMBER_DN
GO_HISTONE_DEACETYLASE_COMPLEX, GO_HISTONE_DEACETYLASE_COMPLEX
BIOCARTA_AKT_PATHWAY, BIOCARTA_AKT_PATHWAY
GO_RETROMER_COMPLEX, GO_RETROMER_COMPLEX
GO_REGULATION_OF_MAST_CELL_ACTIVATION, GO_REGULATION_OF_MAST_CELL_ACTIVATION
ST_PHOSPHOINOSITIDE_3_KINASE_PATHWAY, ST_PHOSPHOINOSITIDE_3_KINASE_PATHWAY
GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_MAC_UP, GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_MAC_UP
GO_REGULATION_OF_CARBOHYDRATE_BIOSYNTHETIC_PROCESS, GO_REGULATION_OF_CARBOHYDRATE_BIOSYNTHETIC_PROCESS
GNF2_LYN, GNF2_LYN
ST_GAQ_PATHWAY, ST_GAQ_PATHWAY
GSE9946_LISTERIA_INF_MATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP, GSE9946_LISTERIA_INF_MATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP
GO_REGULATION_OF_CELLULAR_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA_STIMULUS, GO_REGULATION_OF_CELLULAR_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA_STIMULUS
REACTOME_PIP3_ACTIVATES_AKT_SIGNALING, REACTOME_PIP3_ACTIVATES_AKT_SIGNALING
GSE43955_10H_VS_30H_ACT_CD4_TCELL_DN, GSE43955_10H_VS_30H_ACT_CD4_TCELL_DN
GSE13485_PRE_VS_POST_YF17D_VACCINATION_PBMIC_UP, GSE13485_PRE_VS_POST_YF17D_VACCINATION_PBMIC_UP
REACTOME_REGULATION_OF_AMPK_ACTIVITY_VIA_LKB1, REACTOME_REGULATION_OF_AMPK_ACTIVITY_VIA_LKB1
VALK_AML_CLUSTER_1, VALK_AML_CLUSTER_1
GSE1925_CTRL_VS_3H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN, GSE1925_CTRL_VS_3H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN
GO_CELLULAR_RESPONSE_TO_ACID_CHEMICAL, GO_CELLULAR_RESPONSE_TO_ACID_CHEMICAL
GO_REGULATION_OF_LIPID_STORAGE, GO_REGULATION_OF_LIPID_STORAGE
GO_RESPONSE_TO_AMINO_ACID, GO_RESPONSE_TO_AMINO_ACID
KANG_DOXORUBICIN_RESISTANCE_DN, KANG_DOXORUBICIN_RESISTANCE_DN
GO_PROTEIN_LOCALIZATION_TO_LYSOSOME, GO_PROTEIN_LOCALIZATION_TO_LYSOSOME
WANG_RESPONSE_TO_ANDROGEN_UP, WANG_RESPONSE_TO_ANDROGEN_UP
KIM_GASTRIC_CANCER_CHEMOSENSITIVITY, KIM_GASTRIC_CANCER_CHEMOSENSITIVITY
RASHI_RESPONSE_TO_IONIZING_RADIATION_6, RASHI_RESPONSE_TO_IONIZING_RADIATION_6
GO_NEGATIVE_REGULATION_OF_TOR_SIGNALING, GO_NEGATIVE_REGULATION_OF_TOR_SIGNALING
GO_REGULATION_OF_CELLULAR_EXTRAVASATION, GO_REGULATION_OF_CELLULAR_EXTRAVASATION
TURJANSKI_MAPK1_AND_MAPK2_TARGETS, TURJANSKI_MAPK1_AND_MAPK2_TARGETS
GO_PRIMARY_LYSOSOME, GO_PRIMARY_LYSOSOME
PID_DELTA_NP63_PATHWAY, PID_DELTA_NP63_PATHWAY
PID_TGFB_R, PID_TGFB_R
GO_CELLULAR_RESPONSE_TO_AMINO_ACID_STIMULUS, GO_CELLULAR_RESPONSE_TO_AMINO_ACID_STIMULUS
GSE34156_NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_DN, GSE34156_NOD2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_DN
GO_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
PID_CD40_PATHWAY, PID_CD40_PATHWAY
GO_GASTRULATION, GO_GASTRULATION
BIOCARTA_PTEN_PATHWAY, BIOCARTA_PTEN_PATHWAY
IWANAGA_CARCIINOGENESIS_BY_KRAS_PTEN_UP, IWANAGA_CARCIINOGENESIS_BY_KRAS_PTEN_UP
GO_POSITIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA_STIMULUS, GO_POSITIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA_STIMULUS
KYNING_DNA_DAMAGE_BY_4NQO_OR_UV_KYNING_DNA_DAMAGE_BY_4NQO_OR_UV
SCHLINGEMANN_SKIN_CARCIINOGENESIS_TPA_DN, SCHLINGEMANN_SKIN_CARCIINOGENESIS_TPA_DN
KEGG_T_CELL_RECEPTOR_SIGNALING_PATHWAY, KEGG_T_CELL_RECEPTOR_SIGNALING_PATHWAY
KEGG_CHEMOKINE_SIGNALING_PATHWAY, KEGG_CHEMOKINE_SIGNALING_PATHWAY
MARSON_FOXP3_CORE_DIRECT_TARGETS, MARSON_FOXP3_CORE_DIRECT_TARGETS
REACTOME_TRAF6_MEDIATED_NFKB_ACTIVATION, REACTOME_TRAF6_MEDIATED_NFKB_ACTIVATION
GO_EARLY_ENDOSOME_MEMBRANE, GO_EARLY_ENDOSOME_MEMBRANE
HUMMEL_BURKITTS_LYMPHOMA_DN, HUMMEL_BURKITTS_LYMPHOMA_DN