

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
GSE37416_0H_VS_24H_F_TULARENSIS_1VS_NEUTROPHIL_DN,GSE37416_0H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_DN
GSE3982_BASOPHIL_VS_CENT_MEMORY_CD4_TCELL_UP,GSE3982_BASOPHIL_VS_CENT_MEMORY_CD4_TCELL_UP
GSE32034_1Y6C_HIGH_VS_LOW_MONOCYTE_DN,GSE32034_1Y6C_HIGH_VS_LOW_MONOCYTE_DN
GSE17721_PAM3CSK4_VS_GADIQUIMOD_2H_BMDC_DN,GSE17721_PAM3CSK4_VS_GADIQUIMOD_2H_BMDC_DN
GSE36527_CD69_NEG_VS_POS_TREG_CD62L_LOS_KLRG1_NEG_DN,GSE36527_CD69_NEG_VS_POS_TREG_CD62L_LOS_KLRG1_NEG_DN
GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_DN,GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_DN
GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP,GSE14308_NAIVE_CD4_TCELL_VS_INDUCED_TREG_UP
GSE17721_0.5H_VS_24H_CPG_BMDC_UP,GSE17721_0.5H_VS_24H_CPG_BMDC_UP
GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN,GSE33425_CD161_INT_VS_NEG_CD8_TCELL_DN
GSE15330_WT_VS_IKAROS_KO_LYMPHOID_MULTIPOTENT_PROGENITOR_UP,GSE15330_WT_VS_IKAROS_KO_LYMPHOID_MULTIPOTENT_PROGENITOR_UP
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
HAHTOLA_MYCOSIS_FUNGOIDES_SKIN_UP,HAHTOLA_MYCOSIS_FUNGOIDES_SKIN_UP
GSE27786_LSK_VS_BCELL_DN,GSE27786_LSK_VS_BCELL_DN
GSE36009_WT_VS_NLRP10_KO_DC_LPS_STIM_DN,GSE36009_WT_VS_NLRP10_KO_DC_LPS_STIM_DN
GSE17721_CTRL_VS_LPS_1H_BMDC_UP,GSE17721_CTRL_VS_LPS_1H_BMDC_UP
GSE32164_RESTING_DIFFERENTIATED_VS_ALTERNATIVELY_ACT_M2_MACROPHAGE_DN,GSE32164_RESTING_DIFFERENTIATED_VS_ALTERNATIVELY_ACT_M2_MACROPHAGE_DN
GSE40274_EOS_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP,GSE40274_EOS_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
GSE1448_CTRL_VS_ANTLVBETA5_DP_THYMOCYTE_UP,GSE1448_CTRL_VS_ANTLVBETA5_DP_THYMOCYTE_UP
GSE27786_LSK_VS_CD8_TCELL_DN,GSE27786_LSK_VS_CD8_TCELL_DN
GSE1925_CTRL_VS_24H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN,GSE1925_CTRL_VS_24H_IFNG_STIM_IFNG_PRIMED_MACROPHAGE_DN
GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_UP,GSE36888_STAT5_AB_KNOCKIN_VS_WT_TCELL_IL2_TREATED_6H_UP
LEE_DIFFERENTIATING_T_LYMPHOCYTE,LEE_DIFFERENTIATING_T_LYMPHOCYTE
GSE19401_UNSTIM_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN,GSE19401_UNSTIM_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN
GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_DN,GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_DN
GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN,GSE16385_MONOCYTE_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_DN
GO_NCRNA_TRANSCRIPTION,GO_NCRNA_TRANSCRIPTION
GSE11973_MIR223_KOVS_WT_BONE_MARROW_NEUTROPHIL_UP,GSE11973_MIR223_KOVS_WT_BONE_MARROW_NEUTROPHIL_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
GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_UP,GSE339_CD4POS_VS_CD8POS_DC_IN_CULTURE_UP
GSE1448_ANTL_VALPHA2_VS_VBETA5_DP_THYMOCYTE_UP,GSE1448_ANTL_VALPHA2_VS_VBETA5_DP_THYMOCYTE_UP
GSE17721_0.5H_VS_4H_GARDIQUIMOD_BMDC_DN,GSE17721_0.5H_VS_4H_GARDIQUIMOD_BMDC_DN
KEGG_NEUROTROPHIN_SIGNALING_PATHWAY,KEGG_NEUROTROPHIN_SIGNALING_PATHWAY
GSE39152_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_UP,GSE39152_CD103_NEG_VS_POS_MEMORY_CD8_TCELL_UP
GSE17721_POLYIC_VS_GARDIQUIMOD_24H_BMDC_UP,GSE17721_POLYIC_VS_GARDIQUIMOD_24H_BMDC_UP
GSE27786_LSK_VS_CD4_TCELL_DN,GSE27786_LSK_VS_CD4_TCELL_DN
GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_UP,GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_UP
GSE24142_ADULT_VS_FETAL_EARLY_THYMIC_PROGENITOR_UP,GSE24142_ADULT_VS_FETAL_EARLY_THYMIC_PROGENITOR_UP
GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY3_POST_IMMUNIZATION_UP,GSE39110_UNTREATED_VS_IL2_TREATED_CD8_TCELL_DAY3_POST_IMMUNIZATION_UP
GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_DN,GSE37301_LYMPHOID_PRIMED_MPP_VS_COMMON_LYMPHOID_PROGENITOR_DN
GSE42021_TREG_PLN_VS_CD24HL_TREG_THYMUS_UP,GSE42021_TREG_PLN_VS_CD24HL_TREG_THYMUS_UP
GENTILE_RESPONSE_CLUSTER_D3,GENTILE_RESPONSE_CLUSTER_D3
ETF_Q6,ETF_Q6
GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_4H_UP,GSE30971_CTRL_VS_LPS_STIM_MACROPHAGE_WBP7_HET_4H_UP
GO_LATE_ENDOSOME_MEMBRANE,GO_LATE_ENDOSOME_MEMBRANE
GSE28237_EARLY_VS_LATE_CC_BCELL_DN,GSE28237_EARLY_VS_LATE_CC_BCELL_DN
GSE17721_0.5H_VS_12H_POLYIC_BMDC_DN,GSE17721_0.5H_VS_12H_POLYIC_BMDC_DN
GGCACTT_MIR519E,GGCACTT_MIR519E
GSE41087_WT_VS_FOXP3_MUT_ANTL_CD3_CD28_STIM_CD4_TCELL_UP,GSE41087_WT_VS_FOXP3_MUT_ANTL_CD3_CD28_STIM_CD4_TCELL_UP
AAGGGAT_MIR188,AAGGGAT_MIR188
GSE17721_0.5H_VS_24H_LPS_BMDC_DN,GSE17721_0.5H_VS_24H_LPS_BMDC_DN
GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_DN,GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_52H_DN
GO_ENDOSOME_ORGANIZATION,GO_ENDOSOME_ORGANIZATION
GSE15330_WT_VS_IKAROS_KO_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN,GSE15330_WT_VS_IKAROS_KO_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_DN
GSE17721_CTRL_VS_CPG_12H_BMDC_DN,GSE17721_CTRL_VS_CPG_12H_BMDC_DN
KEGG_NON_SMALL_CELL_LUNG_CANCER,KEGG_NON_SMALL_CELL_LUNG_CANCER
GO_AUTOPHAGOSOME,GO_AUTOPHAGOSOME
GSE17721_POLYIC_VS_GARDIQUIMOD_1H_BMDC_DN,GSE17721_POLYIC_VS_GARDIQUIMOD_1H_BMDC_DN
GSE20715_0H_VS_6H_OZONE_TLR4_KO_LUNG_DN,GSE20715_0H_VS_6H_OZONE_TLR4_KO_LUNG_DN
GSE16385_ROSIGLITAZONE_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP,GSE16385_ROSIGLITAZONE_IFNG_TNF_VS_IL4_STIM_MACROPHAGE_UP
GSE1460_NAIVE_CD4_TCELL_ADULT_BLOOD_VS_THYMIC_STROMAL_CELL_UP,GSE1460_NAIVE_CD4_TCELL_ADULT_BLOOD_VS_THYMIC_STROMAL_CELL_UP
GSE42021_TCONV_PLN_VS_CD24INT_TCONV_THYMUS_UP,GSE42021_TCONV_PLN_VS_CD24INT_TCONV_THYMUS_UP
GSE360_CTRL_VS_T_GONDIL_DC_DN,GSE360_CTRL_VS_T_GONDIL_DC_DN
PID_KIT_PATHWAY,PID_KIT_PATHWAY
GSE17721_CPG_VS_GARDIQUIMOD_4H_BMDC_UP,GSE17721_CPG_VS_GARDIQUIMOD_4H_BMDC_UP
GSE17721_PAM3CSK4_VS_GADIQUIMOD_16H_BMDC_UP,GSE17721_PAM3CSK4_VS_GADIQUIMOD_16H_BMDC_UP
GSE17721_LPS_VS_GARDIQUIMOD_0.5H_BMDC_DN,GSE17721_LPS_VS_GARDIQUIMOD_0.5H_BMDC_DN
KEGG_ENDOMETRIAL_CANCER,KEGG_ENDOMETRIAL_CANCER
GO_POSITIVE_REGULATION_OF_MULTI_ORGANISM_PROCESS,GO_POSITIVE_REGULATION_OF_MULTI_ORGANISM_PROCESS
GSE21670_TGFB_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_DN,GSE21670_TGFB_VS_IL6_TREATED_STAT3_KO_CD4_TCELL_DN
GTATGAT_MIR154_MIR487,GTATGAT_MIR154_MIR487
MARIADASON_REGULATED_BY_HISTONE_ACETYLATION_UP,MARIADASON_REGULATED_BY_HISTONE_ACETYLATION_UP
GNF2_RAP1B,GNF2_RAP1B
PID_P3K_PLC_TRK_PATHWAY,PID_P3K_PLC_TRK_PATHWAY
GSE17721_CTRL_VS_GARDIQUIMOD_0.5H_BMDC_DN,GSE17721_CTRL_VS_GARDIQUIMOD_0.5H_BMDC_DN
GO_INTERCELLULAR_BRIDGE,GO_INTERCELLULAR_BRIDGE
SIG_IL4RECEPTOR_IN_B_LYPHOCYTES,SIG_IL4RECEPTOR_IN_B_LYPHOCYTES
PID_TAP63_PATHWAY,PID_TAP63_PATHWAY
MORL_MATURE_B_LYMPHOCYTE_UP,MORL_MATURE_B_LYMPHOCYTE_UP
AACTGAC_MIR223,AACTGAC_MIR223
GSE37532_WT_VS_PPARG_KO_LN_TCONV_UP,GSE37532_WT_VS_PPARG_KO_LN_TCONV_UP
PID_MYC_PATHWAY,PID_MYC_PATHWAY
PID_TGFB_R_PATHWAY,PID_TGFB_R_PATHWAY
GSE41176_WT_VS_TAK1_KO_UNSTIM_BCELL_UP,GSE41176_WT_VS_TAK1_KO_UNSTIM_BCELL_UP
GO_EARLY_ENDOSOME_MEMBRANE,GO_EARLY_ENDOSOME_MEMBRANE
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_DN,GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_42H_DN
RIZ_ERYTHROID_DIFFERENTIATION_CCNE1,RIZ_ERYTHROID_DIFFERENTIATION_CCNE1
GO_RECYCLING_ENDOSOME,GO_RECYCLING_ENDOSOME
GSE2826_WT_VS_XID_BCELL_DN,GSE2826_WT_VS_XID_BCELL_DN
GO_VESICLE_MEDIATED_TRANSPORT_BETWEEN_ENDOSOMAL_COMPARTMENTS,GO_VESICLE_MEDIATED_TRANSPORT_BETWEEN_ENDOSOMAL_COMPARTMENTS
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_DN,GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_60H_DN
WARTERS_IR_RESPONSE_SGY,WARTERS_IR_RESPONSE_SGY
GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP,GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP
BONCI_TARGETS_OF_MIR15A_AND_MIR16_1,BONCI_TARGETS_OF_MIR15A_AND_MIR16_1
GSE37301_RAG2_KO_VS_RAG2_AND_ETSI1_KO_NK_CELL_UP,GSE37301_RAG2_KO_VS_RAG2_AND_ETSI1_KO_NK_CELL_UP
APRELIKOVA_BRCA1_TARGETS,APRELIKOVA_BRCA1_TARGETS
GSE3982_MAC_VS_NEUTROPHIL_DN,GSE3982_MAC_VS_NEUTROPHIL_DN
GSE17721_CTRL_VS_POLYIC_6H_BMDC_DN,GSE17721_CTRL_VS_POLYIC_6H_BMDC_DN
GO_REGULATION_OF_ERBB_SIGNALING_PATHWAY,GO_REGULATION_OF_ERBB_SIGNALING_PATHWAY
GO_EXTRINSIC_COMPONENT_OF_PLASMA_MEMBRANE,GO_EXTRINSIC_COMPONENT_OF_PLASMA_MEMBRANE
RASHI_RESPONSE_TO_IONIZING_RADIATION_6,RASHI_RESPONSE_TO_IONIZING_RADIATION_6
GSE20500_RETINOIC_ACID_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP,GSE20500_RETINOIC_ACID_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP
GCTCTTG_MIR335,GCTCTTG_MIR335
GO_NEGATIVE_REGULATION_OF_SEQUENCE_SPECIFIC_DNA_BINDING_TRANSCRIPTION_FACTOR_ACTIVITY,GO_NEGATIVE_REGULATION_OF_SEQUENCE_SPECIFIC_DNA_BINDING_TRANSCRIPTION_FACTOR_ACTIVITY
GSE3203_UNTREATED_VS_IFNB_TREATED_LN_BCELL_UP,GSE3203_UNTREATED_VS_IFNB_TREATED_LN_BCELL_UP
GSE15330_LYMPHOID_MULTIPOTENT_VS_PRO_BCELL_UP,GSE15330_LYMPHOID_MULTIPOTENT_VS_PRO_BCELL_UP
GO_PROTEIN_LOCALIZATION_TO_VACUOLE,GO_PROTEIN_LOCALIZATION_TO_VACUOLE
CROONQUIST_NRAS_VS_STROMAL_STIMULATION_UP,CROONQUIST_NRAS_VS_STROMAL_STIMULATION_UP
GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY,GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
GO_POSITIVE_REGULATION_OF_IMMUNE_EFFECTOR_PROCESS,GO_POSITIVE_REGULATION_OF_IMMUNE_EFFECTOR_PROCESS
GSE3982_NEUTROPHIL_VS_BCELL_UP,GSE3982_NEUTROPHIL_VS_BCELL_UP
GO_RESPONSE_TO_EPIDERMAL_GROWTH_FACTOR,GO_RESPONSE_TO_EPIDERMAL_GROWTH_FACTOR
GSE17721_LPS_VS_POLYIC_1H_BMDC_UP,GSE17721_LPS_VS_POLYIC_1H_BMDC_UP
LIN_MELANOMA_COPY_NUMBER_DN,LIN_MELANOMA_COPY_NUMBER_DN
GO_CELLULAR_RESPONSE_TO_EPIDERMAL_GROWTH_FACTOR_STIMULUS,GO_CELLULAR_RESPONSE_TO_EPIDERMAL_GROWTH_FACTOR_STIMULUS
PID_IL8_CXCR2_PATHWAY,PID_IL8_CXCR2_PATHWAY
KEEN_RESPONSE_TO_ROSIGLITAZONE_DN,KEEN_RESPONSE_TO_ROSIGLITAZONE_DN
ST_PHOSPHOINOSITIDE_3_KINASE_PATHWAY,ST_PHOSPHOINOSITIDE_3_KINASE_PATHWAY
ZHAN_LATE_DIFFERENTIATION_GENES_UP,ZHAN_LATE_DIFFERENTIATION_GENES_UP
GSE3920_IFNA_VS_IFNB_TREATED_ENDOTHELIAL_CELL_DN,GSE3920_IFNA_VS_IFNB_TREATED_ENDOTHELIAL_CELL_DN
GO_POSITIVE_REGULATION_OF_VACUOLE_ORGANIZATION,GO_POSITIVE_REGULATION_OF_VACUOLE_ORGANIZATION
GO_POSITIVE_REGULATION_OF_NEURON_DEATH,GO_POSITIVE_REGULATION_OF_NEURON_DEATH
GO_PHOSPHATIDYLINOSITOL_3_PHOSPHATE_BINDING,GO_PHOSPHATIDYLINOSITOL_3_PHOSPHATE_BINDING
GO_BETA_CATENIN_BINDING,GO_BETA_CATENIN_BINDING
KEGG_MTOR_SIGNALING_PATHWAY,KEGG_MTOR_SIGNALING_PATHWAY
KEGG_CHEMOKINE_SIGNALING_PATHWAY,KEGG_CHEMOKINE_SIGNALING_PATHWAY
REACTOME_G_BETA_GAMMA_SIGNALLING_THROUGH_P13KGAMMA,REACTOME_G_BETA_GAMMA_SIGNALLING_THROUGH_P13KGAMMA
KEGG_DORSO_VENTRAL_AXIS_FORMATION,KEGG_DORSO_VENTRAL_AXIS_FORMATION
GO_REGULATION_OF_CELLULAR_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA_STIMULUS,GO_REGULATION_OF_CELLULAR_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA_STIMULUS
WANG_RESPONSE_TO_ANDROGEN_UP,WANG_RESPONSE_TO_ANDROGEN_UP
BIOCARTA_AKT_PATHWAY,BIOCARTA_AKT_PATHWAY
BIOCARTA_KERATINOCYTE_PATHWAY,BIOCARTA_KERATINOCYTE_PATHWAY
TATCTGG_MIR488,TATCTGG_MIR488
GO_PRE_AUTOPHAGOSOMAL_STRUCTURE_MEMBRANE,GO_PRE_AUTOPHAGOSOMAL_STRUCTURE_MEMBRANE
BIOCARTA_PTEN_PATHWAY,BIOCARTA_PTEN_PATHWAY
KANG_DOXORUBICIN_RESISTANCE_DN,KANG_DOXORUBICIN_RESISTANCE_DN
GO_CELLULAR_RESPONSE_TO_AMINO_ACID_STIMULUS,GO_CELLULAR_RESPONSE_TO_AMINO_ACID_STIMULUS
GO_REGULATION_OF_CYTOKINESIS,GO_REGULATION_OF_CYTOKINESIS
GO_POSITIVE_REGULATION_OF_BONE_REMODELING,GO_POSITIVE_REGULATION_OF_BONE_REMODELING
GO_RESPONSE_TO_AMINO_ACID,GO_RESPONSE_TO_AMINO_ACID
GO_REGULATION_OF_NOTCH_SIGNALING_PATHWAY,GO_REGULATION_OF_NOTCH_SIGNALING_PATHWAY
GO_RESPONSE_TO_MONOAMINE,GO_RESPONSE_TO_MONOAMINE
SIG_PIP3_SIGNALING_IN_B_LYMPHOCYTES,SIG_PIP3_SIGNALING_IN_B_LYMPHOCYTES
GO_NEGATIVE_REGULATION_OF_NOTCH_SIGNALING_PATHWAY,GO_NEGATIVE_REGULATION_OF_NOTCH_SIGNALING_PATHWAY
NOUSHMEHR_GBM_GERMLINE_MUTATED,NOUSHMEHR_GBM_GERMLINE_MUTATED
GSE9946_LISTERIA_INF_MATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP,GSE9946_LISTERIA_INF_MATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_UP
GO_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_BINDING,GO_RNA_POLYMERASE_II_TRANSCRIPTION_FACTOR_BINDING
GO_CANONICAL_WNT_SIGNALING_PATHWAY,GO_CANONICAL_WNT_SIGNALING_PATHWAY
chr13q22,chr13q22
GO_POSITIVE_REGULATION_OF_ORGANELLE_ASSEMBLY,GO_POSITIVE_REGULATION_OF_ORGANELLE_ASSEMBLY
GO_POSITIVE_REGULATION_OF_CELLULAR_EXTRAVASATION,GO_POSITIVE_REGULATION_OF_CELLULAR_EXTRAVASATION
GO_REGULATION_OF_POLYSACCHARIDE_METABOLIC_PROCESS,GO_REGULATION_OF_POLYSACCHARIDE_METABOLIC_PROCESS
KYNG_DNA_DAMAGE_BY_4NQO_OR_UV,KYNG_DNA_DAMAGE_BY_4NQO_OR_UV
SCHEIDERREIT_IKK_TARGETS,SCHEIDERREIT_IKK_TARGETS
GO_TRANSCRIPTION_COFACTOR_BINDING,GO_TRANSCRIPTION_COFACTOR_BINDING
GO_BONE_REMODELING,GO_BONE_REMODELING
PID_NOTCH_PATHWAY,PID_NOTCH_PATHWAY
GO_TISSUE_REMODELING,GO_TISSUE_REMODELING
ARGGGTTAA_UNKNOWN,ARGGGTTAA_UNKNOWN
GSE4142_PLASMA_CELL_VS_GC_BCELL_UP,GSE4142_PLASMA_CELL_VS_GC_BCELL_UP
GO_POSITIVE_REGULATION_OF_NEURON_APOPTOTIC_PROCESS,GO_POSITIVE_REGULATION_OF_NEURON_APOPTOTIC_PROCESS
BIOCARTA_ACH_PATHWAY,BIOCARTA_ACH_PATHWAY
GO_PROTEIN_LOCALIZATION_TO_LYSOSOME,GO_PROTEIN_LOCALIZATION_TO_LYSOSOME
REACTOME_SIGNALING_BY_ROBO_RECEPTOR,REACTOME_SIGNALING_BY_ROBO_RECEPTOR
GO_REGULATION_OF_NEURAL_PRECURSOR_CELL_PROLIFERATION,GO_REGULATION_OF_NEURAL_PRECURSOR_CELL_PROLIFERATION
GO_EXTRINSIC_COMPONENT_OF_CYTOPLASMIC_SIDE_OF_PLASMA_MEMBRANE,GO_EXTRINSIC_COMPONENT_OF_CYTOPLASMIC_SIDE_OF_PLASMA_MEMBRANE
GAZIN_EPIGENETIC_SILENCING_BY_KRAS,GAZIN_EPIGENETIC_SILENCING_BY_KRAS
GO_POSITIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA_STIMULUS,GO_POSITIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA_STIMULUS
HUMMEL_BURKITT5_LYMPHOMA_DN,HUMMEL_BURKITT5_LYMPHOMA_DN
TURJANSKI_MAPK1_AND_MAPK2_TARGETS,TURJANSKI_MAPK1_AND_MAPK2_TARGETS
VALK_AML_CLUSTER_1,VALK_AML_CLUSTER_1
CROONQUIST_STROMAL_STIMULATION_DN,CROONQUIST_STROMAL_STIMULATION_DN
GO_MANGANESE_ION_BINDING,GO_MANGANESE_ION_BINDING
GO_PRIMARY_LYSOSOME,GO_PRIMARY_LYSOSOME
GO_PROTEIN_TARGETING_TO_LYSOSOME,GO_PROTEIN_TARGETING_TO_LYSOSOME
BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_24HR_UP,BANDRES_RESPONSE_TO_CARMUSTIN_WITHOUT_MGMT_24HR_UP
BIOCARTA_STRESS_PATHWAY,BIOCARTA_STRESS_PATHWAY
GO_PHOSPHATIDYLINOSITOL_BISPHOSPHATE_BINDING,GO_PHOSPHATIDYLINOSITOL_BISPHOSPHATE_BINDING
GO_NEGATIVE_REGULATION_OF_MUSCLE_CELL_APOPTOTIC_PROCESS,GO_NEGATIVE_REGULATION_OF_MUSCLE_CELL_APOPTOTIC_PROCESS
GSE26023_PHD3_KO_VS_WT_NEUTROPHIL_HYPOXIA_UP,GSE26023_PHD3_KO_VS_WT_NEUTROPHIL_HYPOXIA_UP
GO_REGULATION_OF_LEUKOCYTE_DEGRANULATION,GO_REGULATION_OF_LEUKOCYTE_DEGRANULATION
ST_GAQ_PATHWAY,ST_GAQ_PATHWAY
GO_GASTRULATION,GO_GASTRULATION
GO_PRC1_COMPLEX,GO_PRC1_COMPLEX
GO_NEGATIVE_REGULATION_OF_AXON_EXTENSION,GO_NEGATIVE_REGULATION_OF_AXON_EXTENSION