

ARMSTRONG_EFFECTOR_CD8_TCELL_DN, GSE41867_DAY6_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN

GSE41867_NAIVE_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP
GSE24671_BAKIMULC_VS_SENDAL_VIRUS_INFECTED_MOUSE_SPLENOCYTES_DN, GSE24671_BAKIMULC_VS_SENDAL_VIRUS_INFECTED_MOUSE_SPLENOCYTES_DN
GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN, GSE41867_DAY8_EFFECTOR_VS_DAY30_MEMORY_CD8_TCELL_LCMV_ARMSTRONG_DN
GSE17721_CTRL_VS_POLYIC_4H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_4H_BMDC_UP
GSE17721_LPS_VS_POLYIC_24H_BMDC_UP, GSE17721_LPS_VS_POLYIC_24H_BMDC_UP
GSE27786_BCELL_VS_NEUTROPHIL_UP, GSE27786_BCELL_VS_NEUTROPHIL_UP
GSE41867_NAIVE_VS_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_EFFECTOR_CD8_TCELL_UP
REACTOME_NUCLEOTIDE_EXCISION_REPAIR, REACTOME_NUCLEOTIDE_EXCISION_REPAIR
GSE15624_CTRL_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN, GSE15624_CTRL_VS_6H_HALOFUGINONE_TREATED_CD4_TCELL_DN
GSE22103_UNSTIM_VS_LPS_STIM_NEUTROPHIL_UP, GSE22103_UNSTIM_VS_LPS_STIM_NEUTROPHIL_UP
GSE3982_MAC_VS_NEUTROPHIL_UP, GSE3982_MAC_VS_NEUTROPHIL_UP
GSE7460_CTRL_VS_TGFB_TREATED_ACT_TREG_UP, GSE7460_CTRL_VS_TGFB_TREATED_ACT_TREG_UP
GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_DN, GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_DN
GOBP_NUCLEOTIDE_EXCISION_REPAIR, GOBP_NUCLEOTIDE_EXCISION_REPAIR
GSE27241_WT_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_TH17_POLARIZING_CONDITIONS_DN, GSE27241_WT_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_TH17_POLARIZING_CONDITIONS_DN
GSE17721_0.5H_VS_12H_POLYIC_BMDC_UP, GSE17721_0.5H_VS_12H_POLYIC_BMDC_UP
GSE20198_IL12_VS_IL12_IL18_TREATED_ACT_CD4_TCELL_UP, GSE20198_IL12_VS_IL12_IL18_TREATED_ACT_CD4_TCELL_UP
GSE19888_NO_PRETREAT_VS_ADENOSINE_A3R_INHIBITOR_PRETREATED_MAST_CELL_TCELL_MEMBRANES_ACT_DN, GSE19888_NO_PRETREAT_VS_ADENOSINE_A3R_INHIBITOR_PRETREATED_MAST_CELL_TCELL_MEMBRANES_ACT_DN
GSE40277_EOS_AND_LEF1_TRANSDUCED_VS_GATA1_AND_SATB1_TRANSDUCED_CD4_TCELL_DN, GSE40277_EOS_AND_LEF1_TRANSDUCED_VS_GATA1_AND_SATB1_TRANSDUCED_CD4_TCELL_DN
GSE21774_CD56_BRIGHT_VS_DIM_CD62L_POSITIVE_NK_CELL_DN, GSE21774_CD56_BRIGHT_VS_DIM_CD62L_POSITIVE_NK_CELL_DN
GSE33424_CD161_INT_VS_NEG_CD8_TCELL_UP, GSE33424_CD161_INT_VS_NEG_CD8_TCELL_UP
GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_UP, GSE12001_MIR223_KO_VS_WT_NEUTROPHIL_UP
MTOR_UP.N4.V1_UP, MTOR_UP.N4.V1_UP
GSE35685_CD34POS_CD38NEG_VS_CD34POS_CD10POS_BONE_MARROW_DN, GSE35685_CD34POS_CD38NEG_VS_CD34POS_CD10POS_BONE_MARROW_DN
GOCC_PRERIBOSOME, GOCC_PRERIBOSOME
GSE26030_TH1_VS_TH17_DAY5_POST_POLARIZATION_DN, GSE26030_TH1_VS_TH17_DAY5_POST_POLARIZATION_DN
GSE13306_RA_VS_UNTREATED_TCONV_UP, GSE13306_RA_VS_UNTREATED_TCONV_UP
GSE21546_SAPIA_KO_VS_SAPIA_KO_AND_ELK1_KO_ANTL_CD3_STIM_DP_THYMOCYTES_UP, GSE21546_SAPIA_KO_VS_SAPIA_KO_AND_ELK1_KO_ANTL_CD3_STIM_DP_THYMOCYTES_UP
GSE43863_NAIVE_VS_THF_CD4_EFF_TCELL_D6_LCMV_UP, GSE43863_NAIVE_VS_THF_CD4_EFF_TCELL_D6_LCMV_UP
GSE41867_DAY6_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GSE41867_DAY6_VS_DAY8_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP
GSE14415_ACT_VS_CTRL_NATURAL_TREG_DN, GSE14415_ACT_VS_CTRL_NATURAL_TREG_DN
GSE14308_TH2_VS_NATURAL_TREG_UP, GSE14308_TH2_VS_NATURAL_TREG_UP
GSE5542_UNTREATED_VS_IFNA_TREATED_EPITHELIAL_CELLS_24H_DN, GSE5542_UNTREATED_VS_IFNA_TREATED_EPITHELIAL_CELLS_24H_DN
GSE13547_WT_VS_ZFX_KO_BCELL_ANTL_IGM_STIM_2H_DN, GSE13547_WT_VS_ZFX_KO_BCELL_ANTL_IGM_STIM_2H_DN
GSE3982_EOSINOPHIL_VS_CENT_MEMORY_CD4_TCELL_DN, GSE3982_EOSINOPHIL_VS_CENT_MEMORY_CD4_TCELL_DN
GSE17721_CTRL_VS_GARDIQUIMOD_2H_BMDC_UP, GSE17721_CTRL_VS_GARDIQUIMOD_2H_BMDC_UP
GSE30083_SP1_VS_SP3_THYMOCYTE_UP, GSE30083_SP1_VS_SP3_THYMOCYTE_UP
GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_DN, GSE3920_IFNA_VS_IFNG_TREATED_FIBROBLAST_DN
GSE17721_0.5H_VS_4H_POLYIC_BMDC_UP, GSE17721_0.5H_VS_4H_POLYIC_BMDC_UP
REACTOME_SUMOYLATION_OF_UBIQUITINYLATION_PROTEINS, REACTOME_SUMOYLATION_OF_UBIQUITINYLATION_PROTEINS
GSE23321_CD8_STEM_CELL_MEMORY_VS_EFFECTOR_MEMORY_CD8_TCELL_UP, GSE23321_CD8_STEM_CELL_MEMORY_VS_EFFECTOR_MEMORY_CD8_TCELL_UP
GOBP_RESPONSE_TO_HEAT, GOBP_RESPONSE_TO_HEAT
REACTOME_CELLULAR_RESPONSE_TO_HEAT_STRESS, REACTOME_CELLULAR_RESPONSE_TO_HEAT_STRESS
GSE17721_CTRL_VS_CPG_12H_BMDC_DN, GSE17721_CTRL_VS_CPG_12H_BMDC_DN
GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT, GOBP_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT
GOBP_CELLULAR_RESPONSE_TO_HEAT, GOBP_CELLULAR_RESPONSE_TO_HEAT
REACTOME_DNA_DAMAGE_RECOGNITION_IN_GG_NER, REACTOME_DNA_DAMAGE_RECOGNITION_IN_GG_NER
GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP, GSE5542_UNTREATED_VS_IFNG_TREATED_EPITHELIAL_CELLS_6H_UP
GSE14308_NAIVE_CD4_TCELL_VS_NATURAL_TREG_UP, GSE14308_NAIVE_CD4_TCELL_VS_NATURAL_TREG_UP
GSE22443_NAIVE_VS_ACT_AND_IL12_TREATED_CD8_TCELL_DN, GSE22443_NAIVE_VS_ACT_AND_IL12_TREATED_CD8_TCELL_DN
REACTOME_REGULATION_OF_HSF1_MEDIATED_HEAT_SHOCK_RESPONSE, REACTOME_REGULATION_OF_HSF1_MEDIATED_HEAT_SHOCK_RESPONSE
REACTOME_SUMOYLATION_OF_SUMOYLATION_PROTEINS, REACTOME_SUMOYLATION_OF_SUMOYLATION_PROTEINS
HP_PERIPHERAL_AXONAL_NEUROPATHY, HP_PERIPHERAL_AXONAL_NEUROPATHY
GOBP_CELLULAR_MODIFIED_AMINO_ACID_METABOLIC_PROCESS, GOBP_CELLULAR_MODIFIED_AMINO_ACID_METABOLIC_PROCESS
REACTOME_SULFUR_AMINO_ACID_METABOLISM, REACTOME_SULFUR_AMINO_ACID_METABOLISM
REACTOME_SUMOYLATION_OF_RNA_BINDING_PROTEINS, REACTOME_SUMOYLATION_OF_RNA_BINDING_PROTEINS
RAX2_TARGET_GENES, RAX2_TARGET_GENES
GOBP_A XO_DENDRITIC_TRANSPORT, GOBP_A XO_DENDRITIC_TRANSPORT
MIR7155_3P, MIR7155_3P
REACTOME_NUCLEAR_IMPORT_OF_REV_PROTEIN, REACTOME_NUCLEAR_IMPORT_OF_REV_PROTEIN
GOBP_RIBOSOMAL_LARGE_SUBUNIT_ASSEMBLY, GOBP_RIBOSOMAL_LARGE_SUBUNIT_ASSEMBLY
MIR3136_3P, MIR3136_3P
REACTOME_REGULATION_OF_GLUKOKINASE_BY_GLUKOKINASE_REGULATORY_PROTEIN, REACTOME_REGULATION_OF_GLUKOKINASE_BY_GLUKOKINASE_REGULATORY_PROTEIN
GSE18203_CTRL_VS_INTRATUMORAL_CPG_INJ_MC38_TUMOR_UP, GSE18203_CTRL_VS_INTRATUMORAL_CPG_INJ_MC38_TUMOR_UP
GSE26488_WT_VS_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN, GSE26488_WT_VS_HDAC7_KO_DOUBLE_POSITIVE_THYMOCYTE_DN
ESC_V6.5_UP_LATE.V1_DN, ESC_V6.5_UP_LATE.V1_DN
MODULE_318, MODULE_318
GOBP_SULFUR_AMINO_ACID_METABOLIC_PROCESS, GOBP_SULFUR_AMINO_ACID_METABOLIC_PROCESS
GSE12707_AT16L1_HYPOMORPH_VS_WT_THYMUS_UP, GSE12707_AT16L1_HYPOMORPH_VS_WT_THYMUS_UP
MIR889_5P, MIR889_5P
GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_TRICHOSTATINA_72H_CD8_T_CELL_DN
GOBP_REGULATION_OF_CHOLESTEROL_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_CHOLESTEROL_BIOSYNTHETIC_PROCESS
GSE15324_ELF4_KO_VS_WT_ACTIVATED_CD8_TCELL_DN, GSE15324_ELF4_KO_VS_WT_ACTIVATED_CD8_TCELL_DN
GOCC_NUCLEAR_REPLICATION_FORK, GOCC_NUCLEAR_REPLICATION_FORK
GOCC_MEDIATOR_COMPLEX, GOCC_MEDIATOR_COMPLEX
GOBP_NUCLEOTIDE_PHOSPHORYLATION, GOBP_NUCLEOTIDE_PHOSPHORYLATION
GOBP_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS, GOBP_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS
ZNF677_TARGET_GENES, ZNF677_TARGET_GENES
MIR4802_3P, MIR4802_3P
GOBP_FATTY_ACYL_COA_METABOLIC_PROCESS, GOBP_FATTY_ACYL_COA_METABOLIC_PROCESS
GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_DAMAGE_RECOGNITION, GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_DAMAGE_RECOGNITION
WP_CYSSTEINE_AND_METHIONINE_CATABOLISM, WP_CYSSTEINE_AND_METHIONINE_CATABOLISM
REACTOME_FORMATION_OF_INCISION_COMPLEX_IN_GG_NER, REACTOME_FORMATION_OF_INCISION_COMPLEX_IN_GG_NER
GSE11884_WT_VS_FURIN_KO_NAIVE_CD4_TCELL_DN, GSE11884_WT_VS_FURIN_KO_NAIVE_CD4_TCELL_DN
GSE7460_FOXP3_MUT_VS_HET_ACT_TCONV_DN, GSE7460_FOXP3_MUT_VS_HET_ACT_TCONV_DN
GSE43863_THF_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN, GSE43863_THF_VS_LY6C_INT_CXCR5POS_EFFECTOR_CD4_TCELL_DN
GSE29164_DAY3_VS_DAY7_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DN, GSE29164_DAY3_VS_DAY7_CD8_TCELL_AND_IL12_TREATED_MELANOMA_DN
TARTE_PLASMA_CELL_VS_B_LYMPHOCYTE_UP, TARTE_PLASMA_CELL_VS_B_LYMPHOCYTE_UP
GSE15930_STIM_VS_STIM_AND_IL12_72H_CD8_T_CELL_DN, GSE15930_STIM_VS_STIM_AND_IL12_72H_CD8_T_CELL_DN
ZHONG_PFC_C4_UNKNOWN_INP, ZHONG_PFC_C4_UNKNOWN_INP
GOBP_FATTY_ACID_DERIVATIVE_METABOLIC_PROCESS, GOBP_FATTY_ACID_DERIVATIVE_METABOLIC_PROCESS
HP_ABNORMAL_MEAN_CORPUSCULAR_VOLUME, HP_ABNORMAL_MEAN_CORPUSCULAR_VOLUME
MODULE_256, MODULE_256
HP_CEREBRAL_WHITE_MATTER_ATROPHY, HP_CEREBRAL_WHITE_MATTER_ATROPHY
GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_UP, GSE17721_POLYIC_VS_PAM3CSK4_0.5H_BMDC_UP
GOBP_SULFUR_COMPOUND_CATABOLIC_PROCESS, GOBP_SULFUR_COMPOUND_CATABOLIC_PROCESS
BILANGES_SERUM_SENSITIVE_GENES, BILANGES_SERUM_SENSITIVE_GENES
GOMF_TUMOR_NECROSIS_FACTOR_RECEPTOR_BINDING, GOMF_TUMOR_NECROSIS_FACTOR_RECEPTOR_BINDING
RASHI_RESPONSE_TO_IONIZING_RADIATION_1, RASHI_RESPONSE_TO_IONIZING_RADIATION_1
REACTOME_PROTEIN_METHYLATION, REACTOME_PROTEIN_METHYLATION
MIR4687_3P, MIR4687_3P
GOBP_PURINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS, GOBP_PURINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS
REACTOME_FATTY_ACYL_COA_BIOSYNTHESIS, REACTOME_FATTY_ACYL_COA_BIOSYNTHESIS
GOBP_HOMOCYSTEINE_METABOLIC_PROCESS, GOBP_HOMOCYSTEINE_METABOLIC_PROCESS
GOBP_SULFUR_AMINO_ACID_CATABOLIC_PROCESS, GOBP_SULFUR_AMINO_ACID_CATABOLIC_PROCESS
GOBP_SULFUR_AMINO_ACID_BIOSYNTHETIC_PROCESS, GOBP_SULFUR_AMINO_ACID_BIOSYNTHETIC_PROCESS
GOCC_COP9_SIGNALOSOME, GOCC_COP9_SIGNALOSOME
GSE18148_CBF8_KO_VS_WT_TREG_UP, GSE18148_CBF8_KO_VS_WT_TREG_UP
GOBP_REGULATION_OF_STEROID_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_STEROID_BIOSYNTHETIC_PROCESS
MIR134_3P, MIR134_3P
MANN_RESPONSE_TO_AMIFOSTINE_UP, MANN_RESPONSE_TO_AMIFOSTINE_UP
LEE_AGING_MUSCLE_DN, LEE_AGING_MUSCLE_DN
GOBP_S_ADENOSYLMETHIONINE_METABOLIC_PROCESS, GOBP_S_ADENOSYLMETHIONINE_METABOLIC_PROCESS
HORTON_SREBF_TARGETS, HORTON_SREBF_TARGETS
IRX3_TARGET_GENES, IRX3_TARGET_GENES
WP_TRANSLSULFURATION_PATHWAY, WP_TRANSLSULFURATION_PATHWAY
HP_ABNORMAL_CIRCULATING_SULFUR_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_SULFUR_AMINO_ACID_CONCENTRATION
GOBP_LONG_CHAIN_FATTY_ACYL_COA_METABOLIC_PROCESS, GOBP_LONG_CHAIN_FATTY_ACYL_COA_METABOLIC_PROCESS
WP_METHIONINE_METABOLISM_LEADING_TO_SULPHUR_AMINO_ACIDS_AND_RELATED_DISORDERS, WP_METHIONINE_METABOLISM_LEADING_TO_SULPHUR_AMINO_ACIDS_AND_RELATED_DISORDERS
JEON_SMAD6_TARGETS_DN, JEON_SMAD6_TARGETS_DN
HP_HYPERHOMOCYSTINEMIA, HP_HYPERHOMOCYSTINEMIA
PID_CIRCADIAN_PATHWAY, PID_CIRCADIAN_PATHWAY
BLANCO_MELO_INFLUENZA_A_INFECTION_A594_CELLS_UP, BLANCO_MELO_INFLUENZA_A_INFECTION_A594_CELLS_UP
WP_AUTOSOMAL_RECESSIVE_OSTEOPETROSIS_PATHWAYS, WP_AUTOSOMAL_RECESSIVE_OSTEOPETROSIS_PATHWAYS
HP_NARROW_NASAL_RIDGE, HP_NARROW_NASAL_RIDGE
NIKOLSKY_BREAST_CANCER_20Q11_AMPLICON, NIKOLSKY_BREAST_CANCER_20Q11_AMPLICON
GOMF_THIOLESTER_HYDROLASE_ACTIVITY, GOMF_THIOLESTER_HYDROLASE_ACTIVITY
GOBP_RETROGRADE_AXONAL_TRANSPORT, GOBP_RETROGRADE_AXONAL_TRANSPORT
GSE2585_CD80_HIGH_VS_LOW_MTEC_DN, GSE2585_CD80_HIGH_VS_LOW_MTEC_DN
GOBP_PROTEIN_DENEDDYLATION, GOBP_PROTEIN_DENEDDYLATION
GOBP_FATTY_ACID_DERIVATIVE_CATABOLIC_PROCESS, GOBP_FATTY_ACID_DERIVATIVE_CATABOLIC_PROCESS
GOBP_LONG_CHAIN_FATTY_ACID_BIOSYNTHETIC_PROCESS, GOBP_LONG_CHAIN_FATTY_ACID_BIOSYNTHETIC_PROCESS
GOBP_GNANULOCYTE_DIFFERENTIATION, GOBP_GNANULOCYTE_DIFFERENTIATION
GOBP_GLUTATHIONE_METABOLIC_PROCESS, GOBP_GLUTATHIONE_METABOLIC_PROCESS
GOMF_UBIQUITIN_CONJUGATING_ENZYME_BINDING, GOMF_UBIQUITIN_CONJUGATING_ENZYME_BINDING
GOBP_REGULATION_OF_RRNA_PROCESSING, GOBP_REGULATION_OF_RRNA_PROCESSING
GOCC_SYNAPTONEMAL_STRUCTURE, GOCC_SYNAPTONEMAL_STRUCTURE
GOMF_FATTY_ACID_SYNTHASE_ACTIVITY, GOMF_FATTY_ACID_SYNTHASE_ACTIVITY
GOBP_REGULATION_OF_DNA_DAMAGE_CHECKPOINT, GOBP_REGULATION_OF_DNA_DAMAGE_CHECKPOINT
GOBP_AMINO_ACID_BETAINE_BIOSYNTHETIC_PROCESS, GOBP_AMINO_ACID_BETAINE_BIOSYNTHETIC_PROCESS
GOBP_REGULATION_OF_MITOCHONDRIAL_MRNA_STABILITY, GOBP_REGULATION_OF_MITOCHONDRIAL_MRNA_STABILITY
WP_PHOTODYNAMIC_THERAPYINDUCED_NFKB_SURVIVAL_SIGNALING, WP_PHOTODYNAMIC_THERAPYINDUCED_NFKB_SURVIVAL_SIGNALING
KEGG_PROTEIN_EXPORT, KEGG_PROTEIN_EXPORT
GOBP_METHIONINE_METABOLIC_PROCESS, GOBP_METHIONINE_METABOLIC_PROCESS
GOMF_GLYCINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_GLYCINE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
ZHONG_PFC_C1_ASTROCYTE, ZHONG_PFC_C1_ASTROCYTE
RAMPON_ENRICHED_LEARNING_ENVIRONMENT_EARLY_UP, RAMPON_ENRICHED_LEARNING_ENVIRONMENT_EARLY_UP
GOBP_HYDROGEN_SULFIDE_METABOLIC_PROCESS, GOBP_HYDROGEN_SULFIDE_METABOLIC_PROCESS
HP_HYPOCHOLESTEROLEMIA, HP_HYPOCHOLESTEROLEMIA
HP_ABNORMAL_CIRCULATING_ASPARTATE_FAMILY_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_ASPARTATE_FAMILY_AMINO_ACID_CONCENTRATION
BIOCARTA_RANKL_PATHWAY, BIOCARTA_RANKL_PATHWAY
GROSS_ELK3_TARGETS_DN, GROSS_ELK3_TARGETS_DN
RYAAAKNNNNNNTTGW_UNKNOWN, RYAAAKNNNNNNTTGW_UNKNOWN
HP_GENETIC_ANTICIPATION, HP_GENETIC_ANTICIPATION
GOBP_COENZYME_A_BIOSYNTHETIC_PROCESS, GOBP_COENZYME_A_BIOSYNTHETIC_PROCESS
GOBP_ATTACHMENT_OF_MITOTIC_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GOBP_ATTACHMENT_OF_MITOTIC_SPINDLE_MICROTUBULES_TO_KINETOCHORE
REACTOME_SUMOYLATION_OF_IMMUNE_RESPONSE_PROTEINS, REACTOME_SUMOYLATION_OF_IMMUNE_RESPONSE_PROTEINS
REACTOME_METHYLATION, REACTOME_METHYLATION
REACTOME_LIPOPHAGY, REACTOME_LIPOPHAGY
GOBP_POSITIVE_REGULATION_OF_LIPOPOLYSACCHARIDE_MEDIATED_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_LIPOPOLYSACCHARIDE_MEDIATED_SIGNALING_PATHWAY
WP_OSTEOCLAST_SIGNALING, WP_OSTEOCLAST_SIGNALING
GSE25123_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_DN, GSE25123_ROSIGLITAZONE_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_DN
BOYLAN_MULTIPLE_MYELOMA_D_CLUSTER_UP, BOYLAN_MULTIPLE_MYELOMA_D_CLUSTER_UP
HP_PENOSCROTAL_HYPOSPADIAS, HP_PENOSCROTAL_HYPOSPADIAS
HP_CARPAL_BONE_HYPOPLASIA, HP_CARPAL_BONE_HYPOPLASIA
GOBP_POSTTRANSLATIONAL_PROTEIN_TARGETING_TO_ENDOPLASMIC_RETICULUM_MEMBRANE, GOBP_POSTTRANSLATIONAL_PROTEIN_TARGETING_TO_ENDOPLASMIC_RETICULUM_MEMBRANE