

S_TYPE_2_DIABETES_PPMC_AT_DX_UP,GSE9006_TYPE_1_VS_TYPE_2_DIABETES_PPMC_AT_DX_UP

GSE9006_HEALTHY_VS_TYPE_2_DIABETES_PPMC_AT_DX_UP,GSE9006_HEALTHY_VS_TYPE_2_DIABETES_PPMC_AT_DX_UP
HALLMARK_DNA_REPAIR,HALLMARK_DNA_REPAIR
GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_STAT3_KO_CD4_TCELL_DN,GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_STAT3_KO_CD4_TCELL_DN
GOBP_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR,GOBP_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
GSE45365_WT_VS_JFNAR_KO_CD8A_DC_MCMV_INFECTION_UP,GSE45365_WT_VS_JFNAR_KO_CD8A_DC_MCMV_INFECTION_UP
GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_PDC_DN,GSE29618_PRE_VS_DAY7_POST_TIV_FLU_VACCINE_PDC_DN
GSE22886_NAIVE_TCELL_VS_DC_UP,GSE22886_NAIVE_TCELL_VS_DC_UP
GSE22589_HEALTHY_VS_SIV_INFECTED_DC_UP,GSE22589_HEALTHY_VS_SIV_INFECTED_DC_UP
GSE42021_CD24HL_TREG_VS_CD24HL_TCONV_THYMUS_UP,GSE42021_CD24HL_TREG_VS_CD24HL_TCONV_THYMUS_UP
GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_UP,GSE16385_UNTREATED_VS_12H_ROSIGLITAZONE_IL4_TREATED_MACROPHAGE_UP
GSE22886_CD4_TCELL_VS_BCELL_NAIVE_UP,GSE22886_CD4_TCELL_VS_BCELL_NAIVE_UP
GSE16450_CTRL_VS_JFNA_6H_STIM_MATURE_NEURON_CELL_LINE_DN,GSE16450_CTRL_VS_JFNA_6H_STIM_MATURE_NEURON_CELL_LINE_DN
GSE27786_LSK_VS_NKCELL_DN,GSE27786_LSK_VS_NKCELL_DN
FAN_EMBRYONIC_CTX_BRAIN_NAIVE_LIKE_T_CELL,FAN_EMBRYONIC_CTX_BRAIN_NAIVE_LIKE_T_CELL
GSE9601_UNTREATED_VS_P13K_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_UP,GSE9601_UNTREATED_VS_P13K_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_UP
BROWNE_HCMV_INFECTION_14HR_UP,BROWNE_HCMV_INFECTION_14HR_UP
GSE20500_CTRL_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP,GSE20500_CTRL_VS_RARA_ANTAGONIST_TREATED_CD4_TCELL_UP
GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_24H_DN,GSE8921_UNSTIM_VS_TLR1_2_STIM_MONOCYTE_24H_DN
GSE22886_TCELL_VS_BCELL_NAIVE_UP,GSE22886_TCELL_VS_BCELL_NAIVE_UP
GSE16522_ANTL_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_DN,GSE16522_ANTL_CD3CD28_STIM_VS_UNSTIM_NAIVE_CD8_TCELL_DN
GSE6269_HEALTHY_VS_E_COLL_INF_PBMC_UP,GSE6269_HEALTHY_VS_E_COLL_INF_PBMC_UP
HAHTOLA_MYCOSIS_FUNGOIDES_SKIN_UP,HAHTOLA_MYCOSIS_FUNGOIDES_SKIN_UP
GSE21927_SPLEEN_VS_TUMOR_MONOCYTE_BALBC_DN,GSE21927_SPLEEN_VS_TUMOR_MONOCYTE_BALBC_DN
GSE6259_BCELL_VS_CD4_TCELL_UP,GSE6259_BCELL_VS_CD4_TCELL_UP
GSE21380_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_DN,GSE21380_TFH_VS_GERMINAL_CENTER_TFH_CD4_TCELL_DN
MODULE_257,MODULE_257
GSE6259_DEC205_POS_DC_VS_CD8_TCELL_UP,GSE6259_DEC205_POS_DC_VS_CD8_TCELL_UP
GOCC_RNA_POLYMERASE_II_HOLOENZYME,GOCC_RNA_POLYMERASE_II_HOLOENZYME
REACTOME_FORMATION_OF_TC_NER_PRE_INCISION_COMPLEX,REACTOME_FORMATION_OF_TC_NER_PRE_INCISION_COMPLEX
GSE21546_WT_VS_ELK1_KO_DP_THYMOCYTES_DN,GSE21546_WT_VS_ELK1_KO_DP_THYMOCYTES_DN
GSE22886_NAIVE_TCELL_VS_MONOCYTE_UP,GSE22886_NAIVE_TCELL_VS_MONOCYTE_UP
GSE18804_SPLEEN_MACROPHAGE_VS_COLON_TUMORAL_MACROPHAGE_UP,GSE18804_SPLEEN_MACROPHAGE_VS_COLON_TUMORAL_MACROPHAGE_UP
REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_DNA_REPAIR_GENES,REACTOME_TP53_REGULATES_TRANSCRIPTION_OF_DNA_REPAIR_GENES
GSE17721_CPG_VS_GARDIQUIMOD_16H_BMDC_DN,GSE17721_CPG_VS_GARDIQUIMOD_16H_BMDC_DN
GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_TRICHOSTATINA_CD8_TCELL_UP,GSE15930_NAIVE_VS_72H_IN_VITRO_STIM_TRICHOSTATINA_CD8_TCELL_UP
GSE43863_NAIVE_VS_LY6C_LOW_CXCR5NEG_CD4_EFF_TCELL_D6_1CMV_DN,GSE43863_NAIVE_VS_LY6C_LOW_CXCR5NEG_CD4_EFF_TCELL_D6_1CMV_DN
TRAVAGLINI_LUNG_CD8_NAIVE_T_CELL,TRAVAGLINI_LUNG_CD8_NAIVE_T_CELL
GSE3982_DC_VS_EFF_MEMORY_CD4_TCELL_DN,GSE3982_DC_VS_EFF_MEMORY_CD4_TCELL_DN
GSE27786_LSK_VS_NKTCELL_DN,GSE27786_LSK_VS_NKTCELL_DN
GSE5589_UNSTIM_VS_45MIN_LPS_STIM_MACROPHAGE_DN,GSE5589_UNSTIM_VS_45MIN_LPS_STIM_MACROPHAGE_DN
FAN_EMBRYONIC_CTX_BRAIN_B_CELL,FAN_EMBRYONIC_CTX_BRAIN_B_CELL
PURBEY_TARGETS_OF_CTB1_AND_SATB1_DN,PURBEY_TARGETS_OF_CTB1_AND_SATB1_DN
GSE45739_UNSTIM_VS_ACD3_ACD28_STIM_WT_CD4_TCELL_DN,GSE45739_UNSTIM_VS_ACD3_ACD28_STIM_WT_CD4_TCELL_DN
GOBP_REGULATION_OF_BIOLOGICAL_PROCESS_INVOLVED_IN_SYMBIOTIC_INTERACTION,GOBP_REGULATION_OF_BIOLOGICAL_PROCESS_INVOLVED_IN_SYMBIOTIC_INTERACTION
PID_P53_REGULATION_PATHWAY,PID_P53_REGULATION_PATHWAY
DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_NK_CELLS,DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_NK_CELLS
GSE17721_PAM3CSK4_VS_GADIQUIMOD_4H_BMDC_UP,GSE17721_PAM3CSK4_VS_GADIQUIMOD_4H_BMDC_UP
TRAVAGLINI_LUNG_CLUB_CELL,TRAVAGLINI_LUNG_CLUB_CELL
GNF2_HLA_C,GNF2_HLA_C
GSE17721_CTRL_VS_LPS_0.5H_BMDC_DN,GSE17721_CTRL_VS_LPS_0.5H_BMDC_DN
RAPA_EARLY_UP,V1_DN,RAPA_EARLY_UP,V1_DN
REACTOME_POTENTIAL_THERAPEUTICS_FOR_SARS,REACTOME_POTENTIAL_THERAPEUTICS_FOR_SARS
HP_HORSESHOE_KIDNEY,HP_HORSESHOE_KIDNEY
GSE24634_NAIVE_CD4_TCELL_VS_DAY10_IL4_CONV_TREG_UP,GSE24634_NAIVE_CD4_TCELL_VS_DAY10_IL4_CONV_TREG_UP
GSE21927_EL4_VS_MCA203_TUMOR_MONOCYTES_UP,GSE21927_EL4_VS_MCA203_TUMOR_MONOCYTES_UP
GSE7548_NAIVE_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_DN,GSE7548_NAIVE_VS_DAY28_PCC_IMMUNIZATION_CD4_TCELL_DN
GSE17721_POLYIC_VS_GARDIQUIMOD_0.5H_BMDC_UP,GSE17721_POLYIC_VS_GARDIQUIMOD_0.5H_BMDC_UP
GSE5099_MONOCYTE_VS_CLASSICAL_M1_MACROPHAGE_UP,GSE5099_MONOCYTE_VS_CLASSICAL_M1_MACROPHAGE_UP
GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_UP,GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_UP
ZNF16_TARGET_GENES,ZNF16_TARGET_GENES
MCM2_TARGET_GENES,MCM2_TARGET_GENES
GOBP_AUTOPHAGY_OF_MITOCHONDRION,GOBP_AUTOPHAGY_OF_MITOCHONDRION
GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_CORTEX_THYMUS_DN,GSE18281_SUBCAPSULAR_CORTICAL_REGION_VS_WHOLE_CORTEX_THYMUS_DN
MODULE_350,MODULE_350
NIKOLSKY_BREAST_CANCER_11Q12_Q14_AMPLICON,NIKOLSKY_BREAST_CANCER_11Q12_Q14_AMPLICON
GSE21063_WT_VS_NFATC1_KO_BCELL_UP,GSE21063_WT_VS_NFATC1_KO_BCELL_UP
GOMF_BASAL_TRANSCRIPTION_MACHINERY_BINDING,GOMF_BASAL_TRANSCRIPTION_MACHINERY_BINDING
HP_POOR_SUCK,HP_POOR_SUCK
AGCTCCT_MIR28,AGCTCCT_MIR28
GOCC_TRANSCRIPTION_ELONGATION_FACTOR_COMPLEX,GOCC_TRANSCRIPTION_ELONGATION_FACTOR_COMPLEX
GSE15659_NAIVE_VS_PTPRC_NEG_CD4_TCELL_UP,GSE15659_NAIVE_VS_PTPRC_NEG_CD4_TCELL_UP
PID_TCR_PATHWAY,PID_TCR_PATHWAY
GOBP_SELECTIVE_AUTOPHAGY,GOBP_SELECTIVE_AUTOPHAGY
GSE17721_CTRL_VS_CPG_0.5H_BMDC_DN,GSE17721_CTRL_VS_CPG_0.5H_BMDC_DN
HP_ABNORMAL_CORNEAL_EPITHELIUM_MORPHOLOGY,HP_ABNORMAL_CORNEAL_EPITHELIUM_MORPHOLOGY
GSE15659_RESTING_TREG_VS_NONSUPPRESSIVE_TCELL_UP,GSE15659_RESTING_TREG_VS_NONSUPPRESSIVE_TCELL_UP
PID_ERBB1_RECEPTOR_PROXIMAL_PATHWAY,PID_ERBB1_RECEPTOR_PROXIMAL_PATHWAY
REACTOME_SIGNALING_BY_FGFR2,REACTOME_SIGNALING_BY_FGFR2
GOBP_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT,GOBP_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT
GOBP_CHAPERONE_MEDIATED_PROTEIN_FOLDING,GOBP_CHAPERONE_MEDIATED_PROTEIN_FOLDING
BILANGES_RAPAMYCIN_SENSITIVE_GENES,BILANGES_RAPAMYCIN_SENSITIVE_GENES
GOBP_MITOCHONDRIAL_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT,GOBP_MITOCHONDRIAL_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT
GOBP_REGULATION_OF_RESPONSE_TO_CYTOKINE_STIMULUS,GOBP_REGULATION_OF_RESPONSE_TO_CYTOKINE_STIMULUS
HP_PURE_RED_CELL_APLASIA,HP_PURE_RED_CELL_APLASIA
TRAVAGLINI_LUNG_CD4_MEMORY_EFFECTOR_T_CELL,TRAVAGLINI_LUNG_CD4_MEMORY_EFFECTOR_T_CELL
HP_DERMAL_ATROPHY,HP_DERMAL_ATROPHY
GSE41176_WT_VS_TAK1_KO_ANTLJGM_STIM_BCELL_1H_UP,GSE41176_WT_VS_TAK1_KO_ANTLJGM_STIM_BCELL_1H_UP
GNF2_CBBF,GNF2_CBBF
GSE26928_EFF_MEMORY_VS_CXCR5_POS_CD4_TCELL_UP,GSE26928_EFF_MEMORY_VS_CXCR5_POS_CD4_TCELL_UP
HP_VERTEBRAL_SEGMENTATION_DEFECT,HP_VERTEBRAL_SEGMENTATION_DEFECT
GOMF_PROTEIN_PHOSPHORYLATED_AMINO_ACID_BINDING,GOMF_PROTEIN_PHOSPHORYLATED_AMINO_ACID_BINDING
WEIGEL_OXIDATIVE_STRESS_BY_TBH_AND_H2O2,WEIGEL_OXIDATIVE_STRESS_BY_TBH_AND_H2O2
GNF2_CD7,GNF2_CD7
GCM_VAV1,GCM_VAV1
GNF2_IL2RB,GNF2_IL2RB
WP_G13_SIGNALING_PATHWAY,WP_G13_SIGNALING_PATHWAY
REACTOME_ION_TRANSPORT_BY_P_TYPE_ATPASES,REACTOME_ION_TRANSPORT_BY_P_TYPE_ATPASES
GOMF_PEPTIDE_DISULFIDE_OXIDOREDUCTASE_ACTIVITY,GOMF_PEPTIDE_DISULFIDE_OXIDOREDUCTASE_ACTIVITY
GOBP_REGULATION_OF_PROTEIN_DEACETYLATION,GOBP_REGULATION_OF_PROTEIN_DEACETYLATION
GOMF_C2H2_ZINC_FINGER_DOMAIN_BINDING,GOMF_C2H2_ZINC_FINGER_DOMAIN_BINDING
HP_PROMINENT_OCCIPUT,HP_PROMINENT_OCCIPUT
MODULE_419,MODULE_419
GOCC_IMMUNOLOGICAL_SYNAPSE,GOCC_IMMUNOLOGICAL_SYNAPSE
HP_PALPEBRAL_EDEMA,HP_PALPEBRAL_EDEMA
REACTOME_DDX58_IFIH1_MEDIATED_INDUCTION_OF_INTERFERON_ALPHA_BETA,REACTOME_DDX58_IFIH1_MEDIATED_INDUCTION_OF_INTERFERON_ALPHA_BETA
HP_ABNORMAL_BLOOD_POTASSIUM_CONCENTRATION,HP_ABNORMAL_BLOOD_POTASSIUM_CONCENTRATION
GOMF_STEROID_BINDING,GOMF_STEROID_BINDING
GOBP_INFLAMMATORY_CELL_APOPTOTIC_PROCESS,GOBP_INFLAMMATORY_CELL_APOPTOTIC_PROCESS
GOBP_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS,GOBP_FATTY_ACYL_COA_BIOSYNTHETIC_PROCESS
GOMF_ALCOHOL_BINDING,GOMF_ALCOHOL_BINDING
HP_DISPROPORTIONATE_SHORT_TRUNK_SHORT_STATURE,HP_DISPROPORTIONATE_SHORT_TRUNK_SHORT_STATURE
SRF_01,SRF_01
PID_CD8_TCR_PATHWAY,PID_CD8_TCR_PATHWAY
REACTOME_INTERLEUKIN_37_SIGNALING,REACTOME_INTERLEUKIN_37_SIGNALING
GOBP_REGULATION_OF_ANTIEN_RECEPTOR_MEDIATED_SIGNALING_PATHWAY,GOBP_REGULATION_OF_ANTIEN_RECEPTOR_MEDIATED_SIGNALING_PATHWAY
REACTOME_SIGNALING_BY_EGFR_IN_CANCER,REACTOME_SIGNALING_BY_EGFR_IN_CANCER
MODULE_524,MODULE_524
MODULE_429,MODULE_429
HP_HYPOKALEMIA,HP_HYPOKALEMIA
REACTOME_CONSTITUTIVE_SIGNALING_BY_LIGAND_RESPONSIVE_EGFR_CANCER_VARIANTS,REACTOME_CONSTITUTIVE_SIGNALING_BY_LIGAND_RESPONSIVE_EGFR_CANCER_VARIANTS
GOMF_PHOSPHATIDYLINOSITOL_3_KINASE_BINDING,GOMF_PHOSPHATIDYLINOSITOL_3_KINASE_BINDING
GNF2_PTPN4,GNF2_PTPN4
GOMF_CHOLESTEROL_BINDING,GOMF_CHOLESTEROL_BINDING
BIOCARTA_RHO_PATHWAY,BIOCARTA_RHO_PATHWAY
GOCC_POST_MRNA_RELEASE_SPLICEOSOMAL_COMPLEX,GOCC_POST_MRNA_RELEASE_SPLICEOSOMAL_COMPLEX
MODULE_519,MODULE_519
BIOCARTA_PML_PATHWAY
REACTOME_FORMATION_OF_SENESCENCE_ASSOCIATED_HETEROCHROMATIN_FOCI_SAHF,REACTOME_FORMATION_OF_SENESCENCE_ASSOCIATED_HETEROCHROMATIN_FOCI_SAHF
BUCASAS_PPMC_FLUAXR_FLUVIRIN_CAUCASIAN_MALE_AGE_18_40YO_HIGH_RESPONDERS_1DY_TOP_FUNCTIONAL_NETWORK_POSITIVE,BUCASAS_PPMC_FLUAXR_FLUVIRIN_CAUCASIAN_MALE_AGE_18_40YO_HIGH_RESPONDERS_1DY_TOP_FUNCTIONAL_NETWORK_POSITIVE
GOBP_MRNA_TRANSCRIPTION,GOBP_MRNA_TRANSCRIPTION
HOLLEMAN_DAUORUBICIN_B_ALL_UP,HOLLEMAN_DAUORUBICIN_B_ALL_UP
REACTOME_SIGNALING_BY_ERBB2_ECD_MUTANTS,REACTOME_SIGNALING_BY_ERBB2_ECD_MUTANTS
REACTOME_CONSTITUTIVE_SIGNALING_BY_EGFRVIII,REACTOME_CONSTITUTIVE_SIGNALING_BY_EGFRVIII
REACTOME_RAS_PROCESSING,REACTOME_RAS_PROCESSING
MIR6744_5P,MIR6744_5P
REACTOME_SIGNALING_BY_NTRK3_TRKC,REACTOME_SIGNALING_BY_NTRK3_TRKC
GOBP_ACTIN_FILAMENT_SEVERING,GOBP_ACTIN_FILAMENT_SEVERING
GOBP_POSITIVE_REGULATION_OF_LIPID_KINASE_ACTIVITY,GOBP_POSITIVE_REGULATION_OF_LIPID_KINASE_ACTIVITY
HP_THORACIC_SCOLIOSIS,HP_THORACIC_SCOLIOSIS
SCHLINGEMANN_SKIN_CARCIINOGENESIS_TPA_DN,SCHLINGEMANN_SKIN_CARCIINOGENESIS_TPA_DN
GOBP_RELAXATION_OF_CARDIAC_MUSCLE,GOBP_RELAXATION_OF_CARDIAC_MUSCLE
MIR446_3P,MIR446_3P
HP_ACANTHOSIS_NIGRICANS,HP_ACANTHOSIS_NIGRICANS
HP_ABNORMALITY_OF_NEUTROPHIL_MORPHOLOGY,HP_ABNORMALITY_OF_NEUTROPHIL_MORPHOLOGY
REACTOME_ZBP1_DAI_MEDIATED_INDUCTION_OF_TYPE_1_IFNS,REACTOME_ZBP1_DAI_MEDIATED_INDUCTION_OF_TYPE_1_IFNS
GOBP_POSITIVE_THYMIC_T_CELL_SELECTION,GOBP_POSITIVE_THYMIC_T_CELL_SELECTION
GOBP_REGULATION_OF_AMYLOID_FIBRIL_FORMATION,GOBP_REGULATION_OF_AMYLOID_FIBRIL_FORMATION
GOMF_ACTIN_MONOMER_BINDING,GOMF_ACTIN_MONOMER_BINDING
HP_BEADED_RIBS,HP_BEADED_RIBS
GOBP_NEGATIVE_REGULATION_OF_STEM_CELL_DIFFERENTIATION,GOBP_NEGATIVE_REGULATION_OF_STEM_CELL_DIFFERENTIATION
GOBP_RELAXATION_OF_MUSCLE,GOBP_RELAXATION_OF_MUSCLE
GOMF_CALCIIUM_TRANSMEMBRANE_TRANSPORTER_ACTIVITY,PHOSPHORYLATIVE_MECHANISM,GOMF_CALCIIUM_TRANSMEMBRANE_TRANSPORTER_ACTIVITY,PHOSPHORYLATIVE_MECHANISM
HP_VELOPHARYNGEAL_INSUFFICIENCY,HP_VELOPHARYNGEAL_INSUFFICIENCY
GOBP_CELLULAR_POTASSIUM_ION_HOMEOSTASIS,GOBP_CELLULAR_POTASSIUM_ION_HOMEOSTASIS
NOUSHMEHR_GBM_GERMLINE_MUTATED,NOUSHMEHR_GBM_GERMLINE_MUTATED
GOMF_P_TYPE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY,GOMF_P_TYPE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
REACTOME_SIGNALING_BY_ERBB2_IN_CANCER,REACTOME_SIGNALING_BY_ERBB2_IN_CANCER
BLANCO_MELO_MERS_COV_INFECTION_MCR5_CELLS_DN,BLANCO_MELO_MERS_COV_INFECTION_MCR5_CELLS_DN
GOBP_NEGATIVE_REGULATION_OF_NEURAL_PRECURSOR_CELL_PROLIFERATION,GOBP_NEGATIVE_REGULATION_OF_NEURAL_PRECURSOR_CELL_PROLIFERATION
GOMF_PROTON_EXPORTING_ATPASE_ACTIVITY,PHOSPHORYLATIVE_MECHANISM,GOMF_PROTON_EXPORTING_ATPASE_ACTIVITY,PHOSPHORYLATIVE_MECHANISM
GOBP_SODIUM_ION_EXPORT_ACROSS_PLASMA_MEMBRANE,GOBP_SODIUM_ION_EXPORT_ACROSS_PLASMA_MEMBRANE
GOBP_CYTOPLASMIC_SEQUESTERING_OF_TRANSCRIPTION_FACTOR,GOBP_CYTOPLASMIC_SEQUESTERING_OF_TRANSCRIPTION_FACTOR
GOCC_ANCHORED_COMPONENT_OF_SYNAPTIC_MEMBRANE,GOCC_ANCHORED_COMPONENT_OF_SYNAPTIC_MEMBRANE
SOBOLEV_T_CELL_PANDEMRIX_AGE_18_64YO_7DY_DN,SOBOLEV_T_CELL_PANDEMRIX_AGE_18_64YO_7DY_DN
HP_RENAL_MAGNESIUM_WASTING,HP_RENAL_MAGNESIUM_WASTING
GOBP_REGULATION_OF_ACTIN_NUCLEATION,GOBP_REGULATION_OF_ACTIN_NUCLEATION
REACTOME_LYSINE_CATABOLISM,REACTOME_LYSINE_CATABOLISM
GOBP_THYMIC_T_CELL_SELECTION,GOBP_THYMIC_T_CELL_SELECTION
MEISSNER_BRAIN_ICP_WITH_H3K4ME3,MEISSNER_BRAIN_ICP_WITH_H3K4ME3
REACTOME_IRF3_MEDIATED_ACTIVATION_OF_TYPE_1_IFN,REACTOME_IRF3_MEDIATED_ACTIVATION_OF_TYPE_1_IFN
GOBP_RESPONSE_TO_GLYCOSIDE,GOBP_RESPONSE_TO_GLYCOSIDE
GOCC_SPERMATOPROTEASOME_COMPLEX,GOCC_SPERMATOPROTEASOME_COMPLEX
REACTOME_THE_FATTY_ACID_CYCLING_MODEL,REACTOME_THE_FATTY_ACID_CYCLING_MODEL
REACTOME_LRR_FIL1_INTERACTING_PROTEIN_1_LRRFIP1_ACTIVATES_TYPE_1_IFN_PRODUCTION,REACTOME_LRR_FIL1_INTERACTING_PROTEIN_1_LRRFIP1_ACTIVATES_TYPE_1_IFN_PRODUCTION
GOBP_NEGATIVE_REGULATION_OF_VESICLE_FUSION,GOBP_NEGATIVE_REGULATION_OF_VESICLE_FUSION
GOMF_ALPHA_N_ACETYLGALACTOSAMINIDE_ALPHA_2_6_SIALYLTRANSFERASE_ACTIVITY,GOMF_ALPHA_N_ACETYLGALACTOSAMINIDE_ALPHA_2_6_SIALYLTRANSFERASE_ACTIVITY
HP_RENAL_POTASSIUM_WASTING,HP_RENAL_POTASSIUM_WASTING
HP_BLEPHAROCHALASIS,HP_BLEPHAROCHALASIS
GOMF_P_TYPE_SODIUM_POTASSIUM_EXCHANGING_ATPASE_ACTIVITY,GOMF_P_TYPE_SODIUM_POTASSIUM_EXCHANGING_ATPASE_ACTIVITY
REACTOME_MITOCHONDRIAL_UNCOUPLING,REACTOME_MITOCHONDRIAL_UNCOUPLING
HP_LEFT_TO_RIGHT_SHUNT,HP_LEFT_TO_RIGHT_SHUNT
GOBP_REGULATION_OF_CELL_CELL_ADHESION_MEDIATED_BY_INTEGRIN,GOBP_REGULATION_OF_CELL_CELL_ADHESION_MEDIATED_BY_INTEGRIN
GOCC_ANCHORED_COMPONENT_OF_POSTSYNAPTIC_MEMBRANE,GOCC_ANCHORED_COMPONENT_OF_POSTSYNAPTIC_MEMBRANE
GOBP_NEGATIVE_REGULATION_OF_GLUCCORTICOID_METABOLIC_PROCESS,GOBP_NEGATIVE_REGULATION_OF_GLUCCORTICOID_METABOLIC_PROCESS
GOMF_DELTA_CATENIN_BINDING,GOMF_DELTA_CATENIN_BINDING
ZEILSTRA_CD44_TARGETS_UP,ZEILSTRA_CD44_TARGETS_UP
GOBP_TRIGLYCERIDE_CATABOLIC_PROCESS,GOBP_TRIGLYCERIDE_CATABOLIC_PROCESS