```
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_18H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_18H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_UP
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_10H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_10H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_6H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_6H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_4H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_4H_UP
GSE40666_UNTREATED_VS_IFNA_STIM_CD8_TCELL_90MIN_DN, GSE40666_UNTREATED_VS_IFNA_STIM_CD8_TCELL_90MIN_DN
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_18H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_18H_UP
GSE10239_NAIVE_VS_KLRG1INT_EFF_CD8_TCELL_DN, GSE10239_NAIVE_VS_KLRG1INT_EFF_CD8_TCELL_DN
GSE18804_SPLEEN_MACROPHAGE_VS_TUMORAL_MACROPHAGE_DN, GSE18804_SPLEEN_MACROPHAGE_VS_TUMORAL_MACROPHAGE_DN
GSE46606_IRF4_KO_VS_WT_CD40L_IL2_IL5_1DAY_STIMULATED_BCELL_UP, GSE46606_IRF4_KO_VS_WT_CD40L_IL2_IL5_1DAY_STIMULATED_BCELL_UP
GSE42021_CD24HI_VS_CD24LOW_TREG_THYMUS_UP, GSE42021_CD24HI_VS_CD24LOW_TREG_THYMUS_UP
GSE26669 CTRL VS COSTIM BLOCK MLR CD4 TCELL DN, GSE26669 CTRL VS COSTIM BLOCK MLR CD4 TCELL DN
GSE6259_FLT3L_INDUCED_33D1_POS_DC_VS_CD8_TCELL_UP, GSE6259_FLT3L_INDUCED_33D1_POS_DC_VS_CD8_TCELL_UP
GSE42021_TREG_PLN_VS_TREG_PRECURSORS_THYMUS_UP, GSE42021_TREG_PLN_VS_TREG_PRECURSORS_THYMUS_UP
HP_ABNORMAL_PERIPHERAL_MYELINATION, HP_ABNORMAL_PERIPHERAL_MYELINATION
GSE20152_HTNFA_OVERXPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_DN, GSE20152_HTNFA_OVERXPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_DN
BRUINS_UVC_RESPONSE_MIDDLE, BRUINS_UVC_RESPONSE_MIDDLE
HP_POLYNEUROPATHY, HP_POLYNEUROPATHY
GSE9601_UNTREATED_VS_NFKB_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_UP, GSE9601_UNTREATED_VS_NFKB_INHIBITOR_TREATED_HCMV_INF_MONOCYTE_UP
GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_A_SULFUR_GROUP_OF_DONORS, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_A_SULFUR_GROUP_OF_DONORS
KATSANOU_ELAVL1_TARGETS_DN, KATSANOU_ELAVL1_TARGETS_DN
GNF2_MAP2K3, GNF2_MAP2K3
MIR4425, MIR4425
GNF2_BNIP3L, GNF2_BNIP3L
MIR383_5P, MIR383_5P
HP_SUPRANUCLEAR_GAZE_PALSY, HP_SUPRANUCLEAR_GAZE_PALSY
MIR4455, MIR4455
MIR6892_3P, MIR6892_3P
GNF2_SPTA1, GNF2_SPTA1
HP_HYPOHIDROSIS, HP_HYPOHIDROSIS
GSE34156_UNTREATED_VS_6H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP, GSE34156_UNTREATED_VS_6H_NOD2_AND_TLR1_TLR2_LIGAND_TREATED_MONOCYTE_UP
GOBP_REGULATION_OF_INSULIN_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_INSULIN_RECEPTOR_SIGNALING_PATHWAY
LIN_NPAS4_TARGETS_DN, LIN_NPAS4_TARGETS_DN
HP_LACRIMATION_ABNORMALITY, HP_LACRIMATION_ABNORMALITY
DAVICIONI_PAX_FOXO1_SIGNATURE_IN_ARMS_UP, DAVICIONI_PAX_FOXO1_SIGNATURE_IN_ARMS_UP
GNF2_TAL1, GNF2_TAL1
GNF2_ANK1, GNF2_ANK1
GNF2 SPTB, GNF2 SPTB
HP AMBLYOPIA, HP AMBLYOPIA
GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT, GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT
GOCC SMN COMPLEX, GOCC SMN COMPLEX
LEE LIVER CANCER MYC TGFA DN, LEE LIVER CANCER MYC TGFA DN
MIR6507 3P, MIR6507 3P
GOBP_REGULATION_OF_LIPID_KINASE_ACTIVITY, GOBP_REGULATION_OF_LIPID_KINASE_ACTIVITY
MIR4633_3P_MIR6500_5P, MIR4633_3P_MIR6500_5P
BROWNE_HCMV_INFECTION_30MIN_DN, BROWNE_HCMV_INFECTION_30MIN_DN
HP_ABNORMAL_TISSUE_METABOLITE_CONCENTRATION, HP_ABNORMAL_TISSUE_METABOLITE_CONCENTRATION
HARALAMBIEVA_PBMC_TIV_AGE_50_74YO_CORRELATED_WITH_MEMORY_B_CELL_RESPONSE_3DY_NEGATIVE, HARALAMBIEVA_PBMC_TIV_AGE_50_74YO_CORRELATED_WITH_MEMORY_B_CELL_RESPONSE_3DY_NEGATIVE
GOMF_GLUCOSYLTRANSFERASE_ACTIVITY, GOMF_GLUCOSYLTRANSFERASE_ACTIVITY
GOBP CELLULAR MODIFIED AMINO ACID CATABOLIC PROCESS, GOBP CELLULAR MODIFIED AMINO ACID CATABOLIC PROCESS
GRABARCZYK_BCL11B_TARGETS_DN, GRABARCZYK_BCL11B_TARGETS_DN
MIR3684, MIR3684
PID_EPHA2_FWD_PATHWAY, PID_EPHA2_FWD_PATHWAY
MIR409 5P, MIR409 5P
HASLINGER_B_CLL_WITH_CHROMOSOME_12_TRISOMY, HASLINGER_B_CLL_WITH_CHROMOSOME_12_TRISOMY
HP_DYSDIADOCHOKINESIS, HP_DYSDIADOCHOKINESIS
GOBP_RECEPTOR_RECYCLING, GOBP_RECEPTOR_RECYCLING
GOMF_NUCLEOTIDE_DIPHOSPHATASE_ACTIVITY, GOMF_NUCLEOTIDE_DIPHOSPHATASE_ACTIVITY
HP_ABNORMALITY_OF_THE_DENTAL_PULP, HP_ABNORMALITY_OF_THE_DENTAL_PULP
MIR3934_5P, MIR3934_5P
GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CENTROSOME, GOBP_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CENTROSOME
GOMF_CROSSOVER_JUNCTION_ENDODEOXYRIBONUCLEASE_ACTIVITY, GOMF_CROSSOVER_JUNCTION_ENDODEOXYRIBONUCLEASE_ACTIVITY
MIR7106_3P, MIR7106_3P
chr6q23, chr6q23
GOCC_TRANSCRIPTIONALLY_ACTIVE_CHROMATIN, GOCC_TRANSCRIPTIONALLY_ACTIVE_CHROMATIN
WANG_RESPONSE_TO_BEXAROTENE_UP, WANG_RESPONSE_TO_BEXAROTENE_UP
GOBP_CYCLIC_NUCLEOTIDE_MEDIATED_SIGNALING, GOBP_CYCLIC_NUCLEOTIDE_MEDIATED_SIGNALING
GOBP_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_ACTIVITY, GOBP_REGULATION_OF_PHOSPHATIDYLINOSITOL_3_KINASE_ACTIVITY
HP_HETEROCHROMIA_IRIDIS, HP_HETEROCHROMIA_IRIDIS
REACTOME_METABOLISM_OF_COFACTORS, REACTOME_METABOLISM_OF_COFACTORS
HP_ABNORMAL_MAGNESIUM_CONCENTRATION, HP_ABNORMAL_MAGNESIUM_CONCENTRATION
GOBP_DEVELOPMENTAL_PIGMENTATION, GOBP_DEVELOPMENTAL_PIGMENTATION
HP_INCREASED_SERUM_IRON, HP_INCREASED_SERUM_IRON
NIKOLSKY_BREAST_CANCER_6P24_P22_AMPLICON, NIKOLSKY_BREAST_CANCER_6P24_P22_AMPLICON
GOBP_SNO_S_RNA_3_END_PROCESSING, GOBP_SNO_S_RNA_3_END_PROCESSING
HP_ABNORMAL_PUPILLARY_FUNCTION, HP_ABNORMAL_PUPILLARY_FUNCTION
GOERING_BLOOD_HDL_CHOLESTEROL_QTL_TRANS, GOERING_BLOOD_HDL_CHOLESTEROL_QTL_TRANS
GOCC_AXONAL_GROWTH_CONE, GOCC_AXONAL_GROWTH_CONE
GOBP_GOLGI_DISASSEMBLY, GOBP_GOLGI_DISASSEMBLY
GOMF_TRANSMEMBRANE_RECEPTOR_PROTEIN_TYROSINE_KINASE_ADAPTOR_ACTIVITY, GOMF_TRANSMEMBRANE_RECEPTOR_PROTEIN_TYROSINE_KINASE_ADAPTOR_ACTIVITY
REACTOME_RET_SIGNALING, REACTOME_RET_SIGNALING
GOBP_POSITIVE_REGULATION_OF_LIPID_KINASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_LIPID_KINASE_ACTIVITY
GOMF_TRANSLATION_REPRESSOR_ACTIVITY, GOMF_TRANSLATION_REPRESSOR_ACTIVITY
HP_DRY_HAIR, HP_DRY_HAIR
XPO1_TARGET_GENES, XPO1_TARGET_GENES
GOBP_SIGNAL_COMPLEX_ASSEMBLY, GOBP_SIGNAL_COMPLEX_ASSEMBLY
WELCH GATA1 TARGETS, WELCH GATA1 TARGETS
HP ABNORMAL CIRCULATING ARGININE CONCENTRATION, HP ABNORMAL CIRCULATING ARGININE CONCENTRATION
GOMF_ENDODEOXYRIBONUCLEASE_ACTIVITY_PRODUCING_3_PHOSPHOMONOESTERS, GOMF_ENDODEOXYRIBONUCLEASE_ACTIVITY_PRODUCING_3_PHOSPHOMONOESTERS
MODULE_539, MODULE_539
HP ZONULAR CATARACT, HP ZONULAR CATARACT
GOBP GLYCOSIDE CATABOLIC PROCESS, GOBP GLYCOSIDE CATABOLIC PROCESS
GOCC_HOLLIDAY_JUNCTION_RESOLVASE_COMPLEX, GOCC_HOLLIDAY_JUNCTION_RESOLVASE_COMPLEX
GOCC_BBSOME, GOCC_BBSOME
GOMF MONOAMINE TRANSMEMBRANE TRANSPORTER ACTIVITY, GOMF MONOAMINE TRANSMEMBRANE TRANSPORTER ACTIVITY
WP_ALANINE_AND_ASPARTATE_METABOLISM, WP_ALANINE_AND_ASPARTATE_METABOLISM
HP_UNDETECTABLE_LIGHT_AND_DARK_ADAPTED_ELECTRORETINOGRAM, HP_UNDETECTABLE_LIGHT_AND_DARK_ADAPTED_ELECTRORETINOGRAM
GOCC_ENDODEOXYRIBONUCLEASE_COMPLEX, GOCC_ENDODEOXYRIBONUCLEASE_COMPLEX
GOMF_GLUCURONOSYL_N_ACETYLGALACTOSAMINYL_PROTEOGLYCAN_4_BETA_N_ACETYLGALACTOSAMINYLTRANSFERASE_ACTIVITY, GOMF_GLUCURONOSYL_N_ACETYLGALACTOSAMINYL_PROTEOGLYCAN_4_BETA_N_ACETYLGALACTOSAMINYLTRANSFERASE_ACTIVITY, GOMF_GLUCURONOSYL_N_ACETYLGALACTOSAMINYL_PROTEOGLYCAN_4_BETA_N_ACETYLGAL
MOTAMED_RESPONSE_TO_ANDROGEN_DN, MOTAMED_RESPONSE_TO_ANDROGEN_DN
ZNF20 TARGET GENES, ZNF20 TARGET GENES
GOBP_SPLICEOSOMAL_CONFORMATIONAL_CHANGES_TO_GENERATE_CATALYTIC_CONFORMATION, GOBP_SPLICEOSOMAL_CONFORMATIONAL_CHANGES_TO_GENERATE_CATALYTIC_CONFORMATION
chr16q21, chr16q21
GOBP_UREA_METABOLIC_PROCESS, GOBP_UREA_METABOLIC_PROCESS
HP_ALACRIMA, HP_ALACRIMA
WP_MBDNF_AND_PROBDNF_REGULATION_OF_GABA_NEUROTRANSMISSION, WP_MBDNF_AND_PROBDNF_REGULATION_OF_GABA_NEUROTRANSMISSION
HP_ANKLE_CLONUS, HP_ANKLE_CLONUS
GOBP_N_ACYLPHOSPHATIDYLETHANOLAMINE_METABOLIC_PROCESS, GOBP_N_ACYLPHOSPHATIDYLETHANOLAMINE_METABOLIC_PROCESS
HP_CHRONIC_LYMPHATIC_LEUKEMIA, HP_CHRONIC_LYMPHATIC_LEUKEMIA
GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT_ENDOSOME_TO_PLASMA_MEMBRANE, GOBP_NEUROTRANSMITTER_RECEPTOR_TRANSPORT_ENDOSOME_TO_PLASMA_MEMBRANE
PID_EPHRINB_REV_PATHWAY, PID_EPHRINB_REV_PATHWAY
FOXF1_TARGET_GENES, FOXF1_TARGET_GENES
GOBP_SYNAPTIC_VESICLE_RECYCLING_VIA_ENDOSOME, GOBP_SYNAPTIC_VESICLE_RECYCLING_VIA_ENDOSOME
GOCC DNA RECOMBINASE MEDIATOR COMPLEX, GOCC DNA RECOMBINASE MEDIATOR COMPLEX
GOCC_PHOSPHOLIPID_TRANSLOCATING_ATPASE_COMPLEX, GOCC_PHOSPHOLIPID_TRANSLOCATING_ATPASE_COMPLEX
SABATES_COLORECTAL_ADENOMA_SIZE_DN, SABATES_COLORECTAL_ADENOMA_SIZE_DN
REACTOME_SIGNALING_BY_NTRK2_TRKB, REACTOME_SIGNALING_BY_NTRK2_TRKB
HP_DECREASED_LACRIMATION, HP_DECREASED_LACRIMATION
REACTOME_P130CAS_LINKAGE_TO_MAPK_SIGNALING_FOR_INTEGRINS, REACTOME_P130CAS_LINKAGE_TO_MAPK_SIGNALING_FOR_INTEGRINS
HOEK_NEUTROPHIL_2011_2012_TIV_ADULT_3DY_UP, HOEK_NEUTROPHIL_2011_2012_TIV_ADULT_3DY_UP
GOBP_GLYCOSYLCERAMIDE_METABOLIC_PROCESS, GOBP_GLYCOSYLCERAMIDE_METABOLIC_PROCESS
GOBP_RESPONSE_TO_GROWTH_HORMONE, GOBP_RESPONSE_TO_GROWTH_HORMONE
GOMF_ANION_CATION_SYMPORTER_ACTIVITY, GOMF_ANION_CATION_SYMPORTER_ACTIVITY
GOBP_BEHAVIORAL_RESPONSE_TO_COCAINE, GOBP_BEHAVIORAL_RESPONSE_TO_COCAINE
GOBP_WATER_HOMEOSTASIS, GOBP_WATER_HOMEOSTASIS
GOBP_PROTEIN_OXIDATION, GOBP_PROTEIN_OXIDATION
GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_VIA_MHC_CLASS_IB, GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_VIA_MHC_CLASS_IB
GOBP_AMINOPHOSPHOLIPID_TRANSPORT, GOBP_AMINOPHOSPHOLIPID_TRANSPORT
REACTOME_ROBO_RECEPTORS_BIND_AKAP5, REACTOME_ROBO_RECEPTORS_BIND_AKAP5
GOBP_POLYADENYLATION_DEPENDENT_SNORNA_3_END_PROCESSING, GOBP_POLYADENYLATION_DEPENDENT_SNORNA_3_END_PROCESSING
GOBP_NEGATIVE_REGULATION_OF_OSTEOCLAST_DIFFERENTIATION, GOBP_NEGATIVE_REGULATION_OF_OSTEOCLAST_DIFFERENTIATION
GOBP_HEAT_ACCLIMATION, GOBP_HEAT_ACCLIMATION
GOBP_N_ACYLETHANOLAMINE_METABOLIC_PROCESS, GOBP_N_ACYLETHANOLAMINE_METABOLIC_PROCESS
GOBP_CELLULAR_HYPOTONIC_RESPONSE, GOBP_CELLULAR_HYPOTONIC_RESPONSE
HP_ABNORMAL_CIRCULATING_GLUTAMINE_CONCENTRATION, HP_ABNORMAL_CIRCULATING_GLUTAMINE_CONCENTRATION
GOCC_EXTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_MEMBRANE, GOCC_EXTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_MEMBRANE
REACTOME_UBIQUINOL_BIOSYNTHESIS, REACTOME_UBIQUINOL_BIOSYNTHESIS
GOMF_MONOCARBOXYLIC_ACID_BINDING, GOMF_MONOCARBOXYLIC_ACID_BINDING
GOBP_HYPOTONIC_RESPONSE, GOBP_HYPOTONIC_RESPONSE
REACTOME_ACETYLCHOLINE_REGULATES_INSULIN_SECRETION, REACTOME_ACETYLCHOLINE_REGULATES_INSULIN_SECRETION
GOMF_BILE_ACID_BINDING, GOMF_BILE_ACID_BINDING
GOBP_SEQUESTERING_OF_ZINC_ION, GOBP_SEQUESTERING_OF_ZINC_ION
REACTOME_FATTY_ACIDS_BOUND_TO_GPR40_FFAR1_REGULATE_INSULIN_SECRETION, REACTOME_FATTY_ACIDS_BOUND_TO_GPR40_FFAR1_REGULATE_INSULIN_SECRETION
TCGA_GLIOBLASTOMA_MUTATED, TCGA_GLIOBLASTOMA_MUTATED
WP_HIJACK_OF_UBIQUITINATION_BY_SARSCOV2, WP_HIJACK_OF_UBIQUITINATION_BY_SARSCOV2
GOBP_TRNA_TRANSCRIPTION, GOBP_TRNA_TRANSCRIPTION
GOBP_TROPHECTODERMAL_CELL_DIFFERENTIATION, GOBP_TROPHECTODERMAL_CELL_DIFFERENTIATION
MIKKELSEN_MEF_ICP_WITH_H3K4ME3_AND_H3K27ME3, MIKKELSEN_MEF_ICP_WITH_H3K4ME3_AND_H3K27ME3
HP_ABNORMALITY_OF_THE_DISTAL_PHALANX_OF_THE_5TH_FINGER, HP_ABNORMALITY_OF_THE_DISTAL_PHALANX_OF_THE_5TH_FINGER
HP_PRIMARY_HYPERPARATHYROIDISM, HP_PRIMARY_HYPERPARATHYROIDISM
GOBP_AMMONIUM_TRANSPORT, GOBP_AMMONIUM_TRANSPORT
GOMF_AMMONIUM_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_AMMONIUM_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOCC_TERMINAL_BOUTON, GOCC_TERMINAL_BOUTON
HAY_BONE_MARROW_CD34_POS_ERP, HAY_BONE_MARROW_CD34_POS_ERP
BIOCARTA_PLC_PATHWAY, BIOCARTA_PLC_PATHWAY
GOBP_NEGATIVE_REGULATION_OF_FEEDING_BEHAVIOR, GOBP_NEGATIVE_REGULATION_OF_FEEDING_BEHAVIOR
ZNF136_TARGET_GENES, ZNF136_TARGET_GENES
HP_HYPOPLASTIC_FRONTAL_SINUSES, HP_HYPOPLASTIC_FRONTAL_SINUSES
GOBP_RESPONSE_TO_ISOLATION_STRESS, GOBP_RESPONSE_TO_ISOLATION_STRESS
HP_URINARY_GLYCOSAMINOGLYCAN_EXCRETION, HP_URINARY_GLYCOSAMINOGLYCAN_EXCRETION
HP_YELLOW_BROWN_DISCOLORATION_OF_THE_TEETH, HP_YELLOW_BROWN_DISCOLORATION_OF_THE_TEETH
HP_KNEE_CLONUS, HP_KNEE_CLONUS
HP HYPERMAGNESIURIA, HP HYPERMAGNESIURIA
GOMF_BETA_GLUCOSIDASE_ACTIVITY, GOMF_BETA_GLUCOSIDASE_ACTIVITY
HP STOOPED POSTURE, HP STOOPED POSTURE
GOBP_5S_CLASS_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_III, GOBP_5S_CLASS_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_III
GOBP_REGULATION_OF_FOCAL_ADHESION_DISASSEMBLY, GOBP_REGULATION_OF_FOCAL_ADHESION_DISASSEMBLY
GOCC TRANSCRIPTION FACTOR TFIIIC COMPLEX, GOCC TRANSCRIPTION FACTOR TFIIIC COMPLEX
GOMF_NUCLEOSOMAL_HISTONE_BINDING, GOMF_NUCLEOSOMAL_HISTONE_BINDING
STEINER_ERYTHROCYTE_MEMBRANE_GENES, STEINER_ERYTHROCYTE_MEMBRANE_GENES
REACTOME_THE_ACTIVATION_OF_ARYLSULFATASES, REACTOME_THE_ACTIVATION_OF_ARYLSULFATASES
GOMF_CATION_CHLORIDE_SYMPORTER_ACTIVITY, GOMF_CATION_CHLORIDE_SYMPORTER_ACTIVITY
ILK_TARGET_GENES, ILK_TARGET_GENES
GOBP_MEMBRANE_RAFT_ORGANIZATION, GOBP_MEMBRANE_RAFT_ORGANIZATION
GOBP_SEROTONIN_UPTAKE, GOBP_SEROTONIN_UPTAKE
GOBP_DNA_DEAMINATION, GOBP_DNA_DEAMINATION
GROSS_HIF1A_TARGETS_UP, GROSS_HIF1A_TARGETS_UP
HP_HYPERMAGNESEMIA, HP_HYPERMAGNESEMIA
MIR433_5P, MIR433_5P
GOBP_PARALLEL_ACTIN_FILAMENT_BUNDLE_ASSEMBLY, GOBP_PARALLEL_ACTIN_FILAMENT_BUNDLE_ASSEMBLY
GOBP NEUROTRANSMITTER RECEPTOR TRANSPORT POSTSYNAPTIC ENDOSOME TO LYSOSOME, GOBP NEUROTRANSMITTER RECEPTOR TRANSPORT POSTSYNAPTIC ENDOSOME TO LYSOSOME
GOBP_REGULATION_OF_MACROPHAGE_PROLIFERATION, GOBP_REGULATION_OF_MACROPHAGE_PROLIFERATION
GOBP_HEPARIN_METABOLIC_PROCESS, GOBP_HEPARIN_METABOLIC_PROCESS
GOBP_NEUROTRANSMITTER_LOADING_INTO_SYNAPTIC_VESICLE, GOBP_NEUROTRANSMITTER_LOADING_INTO_SYNAPTIC_VESICLE
GOMF CARBON NITROGEN LYASE ACTIVITY, GOMF CARBON NITROGEN LYASE ACTIVITY
GOBP_NEUROTRANSMITTER_CATABOLIC_PROCESS, GOBP_NEUROTRANSMITTER_CATABOLIC_PROCESS
REACTOME_TETRAHYDROBIOPTERIN_BH4_SYNTHESIS_RECYCLING_SALVAGE_AND_REGULATION, REACTOME_TETRAHYDROBIOPTERIN_BH4_SYNTHESIS_RECYCLING_SALVAGE_AND_REGULATION
GOBP_CENTRAL_NERVOUS_SYSTEM_MYELIN_MAINTENANCE, GOBP_CENTRAL_NERVOUS_SYSTEM_MYELIN_MAINTENANCE
GOBP_MUCOCILIARY_CLEARANCE, GOBP_MUCOCILIARY_CLEARANCE
GOMF TROPONIN I BINDING, GOMF TROPONIN I BINDING
HP_HYPERGLUTAMINEMIA, HP_HYPERGLUTAMINEMIA
GOBP_DIPHOSPHOINOSITOL_POLYPHOSPHATE_METABOLIC_PROCESS, GOBP_DIPHOSPHOINOSITOL_POLYPHOSPHATE_METABOLIC_PROCESS
GOMF_ENDOPOLYPHOSPHATASE_ACTIVITY, GOMF_ENDOPOLYPHOSPHATASE_ACTIVITY
GOMF_TYPE_2A_SEROTONIN_RECEPTOR_BINDING, GOMF_TYPE_2A_SEROTONIN_RECEPTOR_BINDING
GOMF_ERBB_3_CLASS_RECEPTOR_BINDING, GOMF_ERBB_3_CLASS_RECEPTOR_BINDING
HP_JERKY_OCULAR_PURSUIT_MOVEMENTS, HP_JERKY_OCULAR_PURSUIT_MOVEMENTS
GOBP_SCHWANN_CELL_MIGRATION, GOBP_SCHWANN_CELL_MIGRATION
GOBP_ESTABLISHMENT_OF_MEIOTIC_SPINDLE_LOCALIZATION, GOBP_ESTABLISHMENT_OF_MEIOTIC_SPINDLE_LOCALIZATION
GOBP_REGULATION_OF_SEQUESTERING_OF_ZINC_ION, GOBP_REGULATION_OF_SEQUESTERING_OF_ZINC_ION
GOMF_INTERFERON_RECEPTOR_ACTIVITY, GOMF_INTERFERON_RECEPTOR_ACTIVITY
GOBP_PHOSPHOLIPASE_C_ACTIVATING_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY, GOBP_PHOSPHOLIPASE_C_ACTIVATING_DOPAMINE_RECEPTOR_SIGNALING_PATHWAY
GOBP_TETRAHYDROBIOPTERIN_METABOLIC_PROCESS, GOBP_TETRAHYDROBIOPTERIN_METABOLIC_PROCESS
GOBP_REGULATION_OF_REMOVAL_OF_SUPEROXIDE_RADICALS, GOBP_REGULATION_OF_REMOVAL_OF_SUPEROXIDE_RADICALS
GOCC DOPAMINERGIC SYNAPSE, GOCC DOPAMINERGIC SYNAPSE
GOBP_MACROPHAGE_PROLIFERATION, GOBP_MACROPHAGE_PROLIFERATION
MIR4258, MIR4258
HP_CENTRAL_HETEROCHROMIA, HP_CENTRAL_HETEROCHROMIA
HP_FRONTAL_HIRSUTISM, HP_FRONTAL_HIRSUTISM
GOBP_DIADENOSINE_POLYPHOSPHATE_CATABOLIC_PROCESS, GOBP_DIADENOSINE_POLYPHOSPHATE_CATABOLIC_PROCESS
GOBP_PROTEIN_LOCALIZATION_TO_ADHERENS_JUNCTION, GOBP_PROTEIN_LOCALIZATION_TO_ADHERENS_JUNCTION
GOBP_REGULATION_OF_NK_T_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_NK_T_CELL_DIFFERENTIATION
GOMF_PHOSPHATIDYLETHANOLAMINE_FLIPPASE_ACTIVITY, GOMF_PHOSPHATIDYLETHANOLAMINE_FLIPPASE_ACTIVITY
REACTOME_ACTIVATED_NTRK3_SIGNALS_THROUGH_PI3K, REACTOME_ACTIVATED_NTRK3_SIGNALS_THROUGH_PI3K
KEGG FOLATE BIOSYNTHESIS, KEGG FOLATE BIOSYNTHESIS
GOBP NK_T_CELL_DIFFERENTIATION, GOBP_NK_T_CELL_DIFFERENTIATION
REACTOME PI3K AKT ACTIVATION, REACTOME PI3K AKT ACTIVATION
HP_HYPOPLASTIC_IRIS_STROMA, HP_HYPOPLASTIC_IRIS_STROMA
GOBP AMINOPHOSPHOLIPID TRANSLOCATION, GOBP AMINOPHOSPHOLIPID TRANSLOCATION
GOMF AMINOPHOSPHOLIPID FLIPPASE ACTIVITY, GOMF AMINOPHOSPHOLIPID FLIPPASE ACTIVITY
GOMF_AMIDINE_LYASE_ACTIVITY, GOMF_AMIDINE_LYASE_ACTIVITY
FOURNIER_ACINAR_DEVELOPMENT_LATE_UP, FOURNIER_ACINAR_DEVELOPMENT_LATE_UP
HP_ABNORMAL_VERTEBRAL_SEGMENTATION_AND_FUSION, HP_ABNORMAL_VERTEBRAL_SEGMENTATION_AND_FUSION
HP_GEOGRAPHIC_ATROPHY, HP_GEOGRAPHIC_ATROPHY
GOBP_RESPONSE_TO_METHOTREXATE, GOBP_RESPONSE_TO_METHOTREXATE
GOCC_CLATHRIN_SCULPTED_MONOAMINE_TRANSPORT_VESICLE, GOCC_CLATHRIN_SCULPTED_MONOAMINE_TRANSPORT_VESICLE
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GOMF_PHOSPHATIDYLSERINE_FLIPPASE_ACTIVITY, GOMF_PHOSPHATIDYLSERINE_FLIPPASE_ACTIVITY

GOBP_NEGATIVE_REGULATION_OF_MOLECULAR_FUNCTION_IN_OTHER_ORGANISM, GOBP_NEGATIVE_REGULATION_OF_MOLECULAR_FUNCTION_IN_OTHER_ORGANISM

REACTOME_RETINOID_CYCLE_DISEASE_EVENTS, REACTOME_RETINOID_CYCLE_DISEASE_EVENTS

GOBP_NEGATIVE_REGULATION_OF_BEHAVIOR, GOBP_NEGATIVE_REGULATION_OF_BEHAVIOR

GOMF_GLUCURONOSYLTRANSFERASE_ACTIVITY, GOMF_GLUCURONOSYLTRANSFERASE_ACTIVITY

GOBP_AXON_ENSHEATHMENT_IN_CENTRAL_NERVOUS_SYSTEM, GOBP_AXON_ENSHEATHMENT_IN_CENTRAL_NERVOUS_SYSTEM

GOMF_TOXIC_SUBSTANCE_BINDING, GOMF_TOXIC_SUBSTANCE_BINDING

HP_MICROGLOSSIA, HP_MICROGLOSSIA

GOCC_CLATHRIN_SCULPTED_VESICLE, GOCC_CLATHRIN_SCULPTED_VESICLE

GOBP_SEQUESTERING_OF_METAL_ION, GOBP_SEQUESTERING_OF_METAL_ION

_NEWCASTLE_VIRUS_DC_14H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_14H_UP