
ONE_ACTIVITY, GO_HORMONE_ACTIVITY
MODULE_92, MODULE_92
CAHOY_NEURONAL, CAHOY_NEURONAL
REACTOME_PEPTIDE_LIGAND_BINDING_RECEPTORS, REACTOME_PEPTIDE_LIGAND_BINDING_RECEPTORS
GO_NEUROPEPTIDE_HORMONE_ACTIVITY, GO_NEUROPEPTIDE_HORMONE_ACTIVITY
GO_POSITIVE_REGULATION_OF_SYNAPTIC_TRANSMISSION, GO_POSITIVE_REGULATION_OF_SYNAPTIC_TRANSMISSION
GO_FEEDING_BEHAVIOR, GO_FEEDING_BEHAVIOR
GO_POSITIVE_REGULATION_OF_TRANSMEMBRANE_TRANSPORT, GO_POSITIVE_REGULATION_OF_TRANSMEMBRANE_TRANSPORT
KEGG_AUTOIMMUNE_THYROID_DISEASE, KEGG_AUTOIMMUNE_THYROID_DISEASE
GO_DIGESTION, GO_DIGESTION
GO_CELL_SURFACE_RECEPTOR_SIGNALING_PATHWAY_INVOLVED_IN_CELL_CELL_SIGNALING, GO_CELL_SURFACE_RECEPTOR_SIGNALING_PATHWAY_INVOLVED_IN_CELL_CELL_SIGNALING
PTEN_DN.V1_UP, PTEN_DN.V1_UP
REACTOME_REGULATION_OF_BETA_CELL_DEVELOPMENT, REACTOME_REGULATION_OF_BETA_CELL_DEVELOPMENT
KEGG_MATURITY_ONSET_DIABETES_OF_THE_YOUNG, KEGG_MATURITY_ONSET_DIABETES_OF_THE_YOUNG
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_UP, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_UP
NRSF_01, NRSF_01
GO_SIGNAL_RELEASE, GO_SIGNAL_RELEASE
GO_NEUROPEPTIDE_SIGNALING_PATHWAY, GO_NEUROPEPTIDE_SIGNALING_PATHWAY
MODULE_382, MODULE_382
ALK_DN.V1_UP, ALK_DN.V1_UP
GO_REGULATION_OF_RESPONSE_TO_FOOD, GO_REGULATION_OF_RESPONSE_TO_FOOD
GO_REGULATION_OF_BLOOD_PRESSURE, GO_REGULATION_OF_BLOOD_PRESSURE
REACTOME_G_ALPHA_S_SIGNALLING_EVENTS, REACTOME_G_ALPHA_S_SIGNALLING_EVENTS
MIKKELSEN_MCV6_ICP_WITH_H3K27ME3, MIKKELSEN_MCV6_ICP_WITH_H3K27ME3
GO_REGULATION_OF_POSTSYNAPTIC_MEMBRANE_POTENTIAL, GO_REGULATION_OF_POSTSYNAPTIC_MEMBRANE_POTENTIAL
GO_POSITIVE_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORT, GO_POSITIVE_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORT
GNF2_MMP11, GNF2_MMP11
GSE13411_NAIVE_VS_SWITCHED_MEMORY_BCELL_UP, GSE13411_NAIVE_VS_SWITCHED_MEMORY_BCELL_UP
GNF2_KISS1, GNF2_KISS1
GO_REGULATION_OF_CARDIAC_CONDUCTION, GO_REGULATION_OF_CARDIAC_CONDUCTION
GO_REGULATION_OF_APPETITE, GO_REGULATION_OF_APPETITE
GNF2_IGFBP1, GNF2_IGFBP1
GO_EATING_BEHAVIOR, GO_EATING_BEHAVIOR
REACTOME_CLASS_B_2_SECRETIN_FAMILY_RECEPTORS, REACTOME_CLASS_B_2_SECRETIN_FAMILY_RECEPTORS
GO_POSITIVE_REGULATION_OF_VASODILATION, GO_POSITIVE_REGULATION_OF_VASODILATION
GSE18281_CORTICAL_THYMOCYTE_VS_WHOLE_CORTEX_THYMUS_UP, GSE18281_CORTICAL_THYMOCYTE_VS_WHOLE_CORTEX_THYMUS_UP
SMID_BREAST_CANCER_LUMINAL_A_UP, SMID_BREAST_CANCER_LUMINAL_A_UP
GNF2_TIMP2, GNF2_TIMP2
GATA_Q6, GATA_Q6
GO_PERIKARYON, GO_PERIKARYON
GO_STARTLE_RESPONSE, GO_STARTLE_RESPONSE
GNF2_EGFR, GNF2_EGFR
GSE41867_DAY6_VS_DAY15_LCMV_CLONE13_EFFECTOR_CD8_TCELL_DN, GSE41867_DAY6_VS_DAY15_LCMV_CLONE13_EFFECTOR_CD8_TCELL_DN
MODULE_274, MODULE_274
REACTOME_AMINE_DERIVED_HORMONES, REACTOME_AMINE_DERIVED_HORMONES
KIM_BIPOLAR_DISORDER_OLIGODENDROCYTE_DENSITY_CORR_DN, KIM_BIPOLAR_DISORDER_OLIGODENDROCYTE_DENSITY_CORR_DN
GO_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY_COUPLED_TO_CYCLIC_NUCLEOTIDE_SECOND_MESSENGER, GO_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY_COUPLED_TO_CYCLIC_NUCLEOTID
GO_GOLGI_LUMEN, GO_GOLGI_LUMEN
MODULE_234, MODULE_234
GO_ACID_SECRETION, GO_ACID_SECRETION
GO_REGULATION_OF_EXCRETION, GO_REGULATION_OF_EXCRETION
GO_BONE_GROWTH, GO_BONE_GROWTH
GO_POSITIVE_REGULATION_OF_POTASSIUM_ION_TRANSPORT, GO_POSITIVE_REGULATION_OF_POTASSIUM_ION_TRANSPORT
GO_CYCLIC_NUCLEOTIDE_BIOSYNTHETIC_PROCESS, GO_CYCLIC_NUCLEOTIDE_BIOSYNTHETIC_PROCESS
GO_CELLULAR_RESPONSE_TO_VITAMIN, GO_CELLULAR_RESPONSE_TO_VITAMIN
GO_GROWTH_PLATE_CARTILAGE_DEVELOPMENT, GO_GROWTH_PLATE_CARTILAGE_DEVELOPMENT
GO_CGMP_BIOSYNTHETIC_PROCESS, GO_CGMP_BIOSYNTHETIC_PROCESS
GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_FOOD, GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_FOOD
MILICIC_FAMILIAL_ADENOMATOUS_POLYPOSIS_DN, MILICIC_FAMILIAL_ADENOMATOUS_POLYPOSIS_DN
GO_REGULATION_OF_SYNAPTIC_TRANSMISSION_GLUTAMATERGIC, GO_REGULATION_OF_SYNAPTIC_TRANSMISSION_GLUTAMATERGIC
GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_BCELL_UP, GSE29618_LAIV_VS_TIV_FLU_VACCINE_DAY7_BCELL_UP
GSE29618_PRE_VS_DAY7_FLU_VACCINE_MDC_UP, GSE29618_PRE_VS_DAY7_FLU_VACCINE_MDC_UP
GO_REGULATION_OF_POTASSIUM_ION_TRANSPORT, GO_REGULATION_OF_POTASSIUM_ION_TRANSPORT
KANG_GLIS3_TARGETS, KANG_GLIS3_TARGETS
GO_CGMP_METABOLIC_PROCESS, GO_CGMP_METABOLIC_PROCESS
REACTOME_PEPTIDE_HORMONE_BIOSYNTHESIS, REACTOME_PEPTIDE_HORMONE_BIOSYNTHESIS
CAGNWMCNNNGAC_UNKNOWN, CAGNWMCNNNGAC_UNKNOWN
MIKKELSEN_MCV6_LCP_WITH_H3K27ME3, MIKKELSEN_MCV6_LCP_WITH_H3K27ME3
GO_REGULATION_OF_TRANSMISSION_OF_NERVE_IMPULSE, GO_REGULATION_OF_TRANSMISSION_OF_NERVE_IMPULSE
BCAT.100_UP.V1_DN, BCAT.100_UP.V1_DN
YANG_MUC2_TARGETS_DUODENUM_6MO_UP, YANG_MUC2_TARGETS_DUODENUM_6MO_UP
MCGARVEY_SILENCED_BY_METHYLATION_IN_COLON_CANCER, MCGARVEY_SILENCED_BY_METHYLATION_IN_COLON_CANCER
GO_REGULATION_OF_CAMP_METABOLIC_PROCESS, GO_REGULATION_OF_CAMP_METABOLIC_PROCESS
REACTOME_GLYCOPROTEIN_HORMONES, REACTOME_GLYCOPROTEIN_HORMONES
GO_CARTILAGE_DEVELOPMENT_INVOLVED_IN_ENDOCHONDRAL_BONE_MORPHOGENESIS, GO_CARTILAGE_DEVELOPMENT_INVOLVED_IN_ENDOCHONDRAL_BONE_MORPHOGENESIS
PYEON_HPV_POSITIVE_TUMORS_DN, PYEON_HPV_POSITIVE_TUMORS_DN
MODULE_445, MODULE_445
GO_REGULATION_OF_FEMALE_GONAD_DEVELOPMENT, GO_REGULATION_OF_FEMALE_GONAD_DEVELOPMENT
GO_OOCYTE_DIFFERENTIATION, GO_OOCYTE_DIFFERENTIATION
GO_POSITIVE_REGULATION_OF_HEART_RATE, GO_POSITIVE_REGULATION_OF_HEART_RATE
GO_POSITIVE_REGULATION_OF_CYCLIC_NUCLEOTIDE_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_CYCLIC_NUCLEOTIDE_METABOLIC_PROCESS
GO_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORT, GO_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORT
GO_REGULATION_OF_LIPOPROTEIN_METABOLIC_PROCESS, GO_REGULATION_OF_LIPOPROTEIN_METABOLIC_PROCESS
GO_SENSORY_PERCEPTION_OF_PAIN, GO_SENSORY_PERCEPTION_OF_PAIN