

GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_16H_UP
GSE14000_UNSTIM_VS_4H_LPS_DC_UP, GSE14000_UNSTIM_VS_4H_LPS_DC_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_8H_UP
GSE14000_TRANSLATED_RNA_VS_MRNA_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_DC_UP
GSE14000_UNSTIM_VS_16H_LPS_DC_TRANSLATED_RNA_UP, GSE14000_UNSTIM_VS_16H_LPS_DC_TRANSLATED_RNA_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_6H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_6H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_10H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_10H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_4H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_4H_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_UP, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_UP
GSE41176_UNSTIM_VS_ANTLJGM_STIM_BCELL_3H_DN, GSE41176_UNSTIM_VS_ANTLJGM_STIM_BCELL_3H_DN
GSE14000_4H_VS_16H_LPS_DC_TRANSLATED_RNA_DN, GSE14000_4H_VS_16H_LPS_DC_TRANSLATED_RNA_DN
GSE2706_UNSTIM_VS_2H_R848_DC_UP, GSE2706_UNSTIM_VS_2H_R848_DC_UP
GSE22611_NOD2_VS_CTRL_TRANSDUCED_HEK293T_CELL_DN, GSE22611_NOD2_VS_CTRL_TRANSDUCED_HEK293T_CELL_DN
GSE6259_BCELL_VS_CD8_TCELL_DN, GSE6259_BCELL_VS_CD8_TCELL_DN
GSE4021_CD24HL_VS_CD24LOW_TREG_THYMUS_UP, GSE4021_CD24HL_VS_CD24LOW_TREG_THYMUS_UP
GSE37534_UNTREATED_VS_ROSIGLITAZONE_TREATED_CD4_TCELL_PPARG1_AND_FOXP3 TRASDUCED_UP, GSE37534_UNTREATED_VS_ROSIGLITAZONE_TREATED_CD4_TCELL_PPARG1_AND_FOXP3 TRASDUCED_UP
GSE3982_CTRL_VS_LPS_48H_DC_UP, GSE3982_CTRL_VS_LPS_48H_DC_UP
GSE19401_PLN_VS_Peyers_PATCH_FOLLICULAR_DC_UP, GSE19401_PLN_VS_Peyers_PATCH_FOLLICULAR_DC_UP
GOBP_MATURATION_OF_5_8S_RRNA, GOBP_MATURATION_OF_5_8S_RRNA
MIR6501_3P, MIR6501_3P
CSE36891_POLYIC_TLR3_VS_PAM_TLR2_STIM_PERITONEAL_MACROPHAGE_DN, GSE36891_POLYIC_TLR3_VS_PAM_TLR2_STIM_PERITONEAL_MACROPHAGE_DN
DESEKT_PERIVENOUS_HEPATOCELLULAR_CARCINOMA_SUBCLASS_UP, DESEKT_PERIVENOUS_HEPATOCELLULAR_CARCINOMA_SUBCLASS_UP
GOMF_3_5_EXONUCLEASE_ACTIVITY, GOMF_3_5_EXONUCLEASE_ACTIVITY
MIR6716_5P, MIR6716_5P
MIR2168_3P, MIR2168_3P
GOBP_MATURATION_OF_5_8S_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_S5U_RRNA_5_8S_RRNA_L5U_RRNA, GOBP_MATURATION_OF_5_8S_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_S5U_RRNA_5_8S_RRNA_L5U_RRNA
MIR3925_5P, MIR3925_5P
GSE21546_WT_VS_SAPIA_KO_DP_THYMOCYTES_DN, GSE21546_WT_VS_SAPIA_KO_DP_THYMOCYTES_DN
GSE40274_FOXP3_VS_FOXP3_AND_GATA1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_FOXP3_VS_FOXP3_AND_GATA1_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
GOBP_MATURATION_OF_S5U_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_S5U_RRNA_5_8S_RRNA_L5U_RRNA, GOBP_MATURATION_OF_S5U_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_S5U_RRNA_5_8S_RRNA_L5U_RRNA
GSE19401_PAM2CSK4_VS_RETINOIC_ACID_STIM_FOLLICULAR_DC_DN, GSE19401_PAM2CSK4_VS_RETINOIC_ACID_STIM_FOLLICULAR_DC_DN
HP_NEOPLASM_OF_THE_PANCREAS, HP_NEOPLASM_OF_THE_PANCREAS
HP_MALIGNANT_NEOPLASM_OF_THE_CENTRAL_NERVOUS_SYSTEM, HP_MALIGNANT_NEOPLASM_OF_THE_CENTRAL_NERVOUS_SYSTEM
GOCC_PRERIBOSOME_LARGE_SUBUNIT_PRECURSOR, GOCC_PRERIBOSOME_LARGE_SUBUNIT_PRECURSOR
MIR6849_3P, MIR6849_3P
MIR6513_5P, MIR6513_5P
GSE22611_MUTANT_NOD2_TRANSDUCED_VS_CTRL_HEK293T_STIMULATED_WITH_MDP_6H_UP, GSE22611_MUTANT_NOD2_TRANSDUCED_VS_CTRL_HEK293T_STIMULATED_WITH_MDP_6H_UP
MIR1304_5P, MIR1304_5P
BHAT_ESR1_TARGETS_NOT_VIA_AKT1_DN, BHAT_ESR1_TARGETS_NOT_VIA_AKT1_DN
HP_NEOPLASM_OF_THE_PITUITARY_GLAND, HP_NEOPLASM_OF_THE_PITUITARY_GLAND
HP_GLIOMA, HP_GLIOMA
GOBP_PEROXISOMAL_MEMBRANE_TRANSPORT, GOBP_PEROXISOMAL_MEMBRANE_TRANSPORT
HP_GONADAL_NEOPLASM, HP_GONADAL_NEOPLASM
MIR3922_3P, MIR3922_3P
HP_PITUITARY_ADENOMA, HP_PITUITARY_ADENOMA
SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5, SHEDDEN_LUNG_CANCER_GOOD_SURVIVAL_A5
GENTILE_UV_RESPONSE_CLUSTER_D9, GENTILE_UV_RESPONSE_CLUSTER_D9
GOCC_NUCLEAR_EXOSOME_RNASE_COMPLEX, GOCC_NUCLEAR_EXOSOME_RNASE_COMPLEX
MIR3117_5P, MIR3117_5P
MIR567, MIR567
ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_6HR_DN, ANDERSON_BLOOD_CN54GP140_ADJUVANTED_WITH_GLA_AF_AGE_18_45YO_6HR_DN
GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_RRNA, GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_RRNA
GOBP_MITOTIC_RECOMBINATION, GOBP_MITOTIC_RECOMBINATION
MIR550A_3P, MIR550A_3P
CSE9946_IMMATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_DN, CSE9946_IMMATURE_VS_PROSTAGLANDINE2_TREATED_MATURE_DC_DN
CSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_DN, GSE25088_CTRL_VS_ROSIGLITAZONE_STIM_MACROPHAGE_DN
GOBP_RRNA_METHYLATION, GOBP_RRNA_METHYLATION
MIR3188, MIR3188
HP_PROGRESSIVE_MUSCLE_WEAKNESS, HP_PROGRESSIVE_MUSCLE_WEAKNESS
GOCC_PRERIBOSOME_SMALL_SUBUNIT_PRECURSOR, GOCC_PRERIBOSOME_SMALL_SUBUNIT_PRECURSOR
KEGG_NON_HOMOLOGOUS_END_JOINING, KEGG_NON_HOMOLOGOUS_END_JOINING
YRCCAknNGNCGC, UNKNOWN, YRCCAknNGNCGC, UNKNOWN
DACOSTA_UV_RESPONSE_VIA_ERCC3_XPCS_UP, DACOSTA_UV_RESPONSE_VIA_ERCC3_XPCS_UP
HP_SPASTIC_DYSARTHRIA, HP_SPASTIC_DYSARTHRIA
MIR5581_5P, MIR5581_5P
HP_INCREASED_CIRCULATING_PROLACTIN_CONCENTRATION, HP_INCREASED_CIRCULATING_PROLACTIN_CONCENTRATION
HP_ABNORMAL_CIRCULATING_FERRITIN_CONCENTRATION, HP_ABNORMAL_CIRCULATING_FERRITIN_CONCENTRATION
DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_GLOBOSE_BASAL_CELLS, DURANTE_ADULT_OLFACTORY_NEUROEPITHELIUM_GLOBOSE_BASAL_CELLS
WP_DISRUPTION_OF_POSTSYNAPTIC_SIGNALLING_BY_CNV, WP_DISRUPTION_OF_POSTSYNAPTIC_SIGNALLING_BY_CNV
GOMF_HYDROLASE_ACTIVITY_HYDROLYZING_N_GLYCOSYL_COMPOUNDS, GOMF_HYDROLASE_ACTIVITY_HYDROLYZING_N_GLYCOSYL_COMPOUNDS
HP_DYSGRAPHIA, HP_DYSGRAPHIA
HP_AMAUROSIS_FUGAX, HP_AMAUROSIS_FUGAX
HP_ESOPHAGITIS, HP_ESOPHAGITIS
GOBP_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS_VIA_SOMATIC_MUTATION, GOBP_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS_VIA_SOMATIC_MUTATION
HP_NEOPLASM_OF_THE_RESPIRATORY_SYSTEM, HP_NEOPLASM_OF_THE_RESPIRATORY_SYSTEM
GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_DN, GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_DN
GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_TRANSLATION, GOBP_POSITIVE_REGULATION_OF_MITOCHONDRIAL_TRANSLATION
GOBP_PROTEIN_IMPORT_INTO_PEROXISOME_MATRIX, GOBP_PROTEIN_IMPORT_INTO_PEROXISOME_MATRIX
GOBP_MITOCHONDRIAL_TRNA_PROCESSING, GOBP_MITOCHONDRIAL_TRNA_PROCESSING
GOBP_MITOCHONDRIAL_RNA_MODIFICATION, GOBP_MITOCHONDRIAL_RNA_MODIFICATION
MIR4781_3P, MIR4781_3P
GOBP_HISTONE_H2A_ACETYLTATION, GOBP_HISTONE_H2A_ACETYLTATION
HP_CHOLESTATIC_LIVER_DISEASE, HP_CHOLESTATIC_LIVER_DISEASE
RYAAKANNNNNTTGW, UNKNOWN, RYAAKANNNNNTTGW, UNKNOWN
GOBP_POLYADENYLATION_DEPENDENT_SNORNA_3_END_PROCESSING, GOBP_POLYADENYLATION_DEPENDENT_SNORNA_3_END_PROCESSING
GOBP_MULTICELLULAR_ORGANISM_AGING, GOBP_MULTICELLULAR_ORGANISM_AGING
GOBP_TELOMERE_MAINTENANCE_IN_RESPONSE_TO_DNA_DAMAGE, GOBP_TELOMERE_MAINTENANCE_IN_RESPONSE_TO_DNA_DAMAGE
HP_GLIOBLASTOMA_MULTIFORME, HP_GLIOBLASTOMA_MULTIFORME
GOBP_TRNA_WOBBLE_BASE_MODIFICATION, GOBP_TRNA_WOBBLE_BASE_MODIFICATION
KEGG_LONG_TERM_POTENTIATION, KEGG_LONG_TERM_POTENTIATION
GOBP_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLTATION, GOBP_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLTATION
GOBP_DEOXYRIBOSE_PHOSPHATE_CATABOLIC_PROCESS, GOBP_DEOXYRIBOSE_PHOSPHATE_CATABOLIC_PROCESS
REACTOME_SYNTHESIS_OF_PIP3_AT_THE_EARLY_ENDOSOME_MEMBRANE, REACTOME_SYNTHESIS_OF_PIP3_AT_THE_EARLY_ENDOSOME_MEMBRANE
GOBP_NEGATIVE_REGULATION_OF_ATPASE_ACTIVITY, GOBP_NEGATIVE_REGULATION_OF_ATPASE_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_SISTER_CHROMATID_COHESION, GOBP_POSITIVE_REGULATION_OF_SISTER_CHROMATID_COHESION
GOBP_REGULATION_OF_CYCLIC_NUCLEOTIDE_PHOSPHODIESTERASE_ACTIVITY, GOBP_REGULATION_OF_CYCLIC_NUCLEOTIDE_PHOSPHODIESTERASE_ACTIVITY
GOCC_MISMATCH_REPAIR_COMPLEX, GOCC_MISMATCH_REPAIR_COMPLEX
GOBP_PYRIMIDINE_NUCLEOTIDE_CATABOLIC_PROCESS, GOBP_PYRIMIDINE_NUCLEOTIDE_CATABOLIC_PROCESS
GOBP_PHOTORECEPTOR_CELL_MAINTENANCE, GOBP_PHOTORECEPTOR_CELL_MAINTENANCE
GOBP_TELOMERE_MAINTENANCE_VIA_RECOMBINATION, GOBP_TELOMERE_MAINTENANCE_VIA_RECOMBINATION
GOBP_DETERMINATION_OF_ADULT_LIFESPAN, GOBP_DETERMINATION_OF_ADULT_LIFESPAN
MIR625_3P, MIR625_3P
GOBP_U4_SNRNA_3_END_PROCESSING, GOBP_U4_SNRNA_3_END_PROCESSING
MIR5684, MIR5684
MIR596, MIR596
MIR3684, MIR3684
GOMF_OXIDIZED_DNA_BINDING, GOMF_OXIDIZED_DNA_BINDING
MODULE_360, MODULE_360
HP_TALIPES_CALCANEVALGUS, HP_TALIPES_CALCANEVALGUS
GOBP_CONSTITUTIVE_SECRETORY_PATHWAY, GOBP_CONSTITUTIVE_SECRETORY_PATHWAY
GOBP_POSITIVE_REGULATION_OF_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY
GOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_4_PHOSPHATASE_ACTIVITY, GOMF_PHOSPHATIDYLINOSITOL_PHOSPHATE_4_PHOSPHATASE_ACTIVITY
GSE21546_UNSTIM_VS_ANTI_CD3_STIM_SAPIA_KO_AND_ELK1_KO_DP_THYMOCYTES_DN, GSE21546_UNSTIM_VS_ANTI_CD3_STIM_SAPIA_KO_AND_ELK1_KO_DP_THYMOCYTES_DN
GOBP_POSITIVE_REGULATION_OF_P38MAPK_CASCADE, GOBP_POSITIVE_REGULATION_OF_P38MAPK_CASCADE
GOBP_ACTIN_POLYMERIZATION_DEPENDENT_CELL_MOTILITY, GOBP_ACTIN_POLYMERIZATION_DEPENDENT_CELL_MOTILITY
HP_DEHYDRATION, HP_DEHYDRATION
HP_DECREASED_LACRIMATION, HP_DECREASED_LACRIMATION
GOMF_MISMATCH_REPAIR_COMPLEX_BINDING, GOMF_MISMATCH_REPAIR_COMPLEX_BINDING
WANG_RECURRENT_LIVER_CANCER_UP, WANG_RECURRENT_LIVER_CANCER_UP
GOBP_NUCLEAR_RETENTION_OF_PRE_MRNA_AT_THE_SITE_OF_TRANSCRIPTION, GOBP_NUCLEAR_RETENTION_OF_PRE_MRNA_AT_THE_SITE_OF_TRANSCRIPTION
PHESSSE_TARGETS_OF_APC_AND_MBD2_DN, PHESSSE_TARGETS_OF_APC_AND_MBD2_DN
LIU_TARGETS_OF_VMYB_VS_CMYB_DN, LIU_TARGETS_OF_VMYB_VS_CMYB_DN
HP_CARDIAC_DIVERTICULUM, HP_CARDIAC_DIVERTICULUM
MIR6798_3P, MIR6798_3P
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_RED_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_RED_DN
MYOGNF1_01, MYOGNF1_01
GOBP_CHONDROCYTE_DEVELOPMENT_INVOLVED_IN_ENDOCHONDRAL_BONE_MORPHOGENESIS, GOBP_CHONDROCYTE_DEVELOPMENT_INVOLVED_IN_ENDOCHONDRAL_BONE_MORPHOGENESIS
GOBP_IMMATURE_B_CELL_DIFFERENTIATION, GOBP_IMMATURE_B_CELL_DIFFERENTIATION
GOMF_ACTIVIN_ACTIVATED_RECEPTOR_ACTIVITY, GOMF_ACTIVIN_ACTIVATED_RECEPTOR_ACTIVITY
NIKOLSKY_BREAST_CANCER_12Q24_AMPLICON, NIKOLSKY_BREAST_CANCER_12Q24_AMPLICON
VALK_AML_CLUSTER_13, VALK_AML_CLUSTER_13
GOBP_NEGATIVE_REGULATION_OF_NEUROTRANSMITTER_SECRETION, GOBP_NEGATIVE_REGULATION_OF_NEUROTRANSMITTER_SECRETION
ZHENG_IL22_SIGNALING_DN, ZHENG_IL22_SIGNALING_DN
GOBP_REGULATION_OF_INNER_EAR_AUDITORY_RECEPTOR_CELL_DIFFERENTIATION, GOBP_REGULATION_OF_INNER_EAR_AUDITORY_RECEPTOR_CELL_DIFFERENTIATION
MIKKELSEN_PLURIPOTENT_STATE_DN, MIKKELSEN_PLURIPOTENT_STATE_DN
HP_NEONATAL ASPHYXIA, HP_NEONATAL ASPHYXIA
GOBP_NEGATIVE_REGULATION_OF_TELOMERE_CAPPING, GOBP_NEGATIVE_REGULATION_OF_TELOMERE_CAPPING
REACTOME_COMPETING_ENDOGENOUS_RNAS_CERNAS_REGULATE_PTEEN_TRANSLATION, REACTOME_COMPETING_ENDOGENOUS_RNAS_CERNAS_REGULATE_PTEEN_TRANSLATION
HP_ABNORMAL_SALIVARY_GLAND_MORPHOLOGY, HP_ABNORMAL_SALIVARY_GLAND_MORPHOLOGY
GOMF_GLYCOSYLTRANSFERASE_ACTIVITY, GOMF_GLYCOSYLTRANSFERASE_ACTIVITY
HP_ABNORMAL_BLOOD_MONOVALENT_INORGANIC_CATION_CONCENTRATION, HP_ABNORMAL_BLOOD_MONOVALENT_INORGANIC_CATION_CONCENTRATION
GOBP_PIGMENT_GRANULE_MATURATION, GOBP_PIGMENT_GRANULE_MATURATION
GOBP_ANTEROGRADE_AXONAL_PROTEIN_TRANSPORT, GOBP_ANTEROGRADE_AXONAL_PROTEIN_TRANSPORT
HP_CLINODACTYL_OF_THE_5TH_TOE, HP_CLINODACTYL_OF_THE_5TH_TOE
HP_ALACRIMA, HP_ALACRIMA
MIR5605, MIR5605
REACTOME_PEXOPHAGY, REACTOME_PEXOPHAGY
GOBP_ACTOMYOSIN_CONTRACTILE_RING_ORGANIZATION, GOBP_ACTOMYOSIN_CONTRACTILE_RING_ORGANIZATION
MORE_JAG1, MORE_JAG1
GOBP_AXO_DENDRITIC_PROTEIN_TRANSPORT, GOBP_AXO_DENDRITIC_PROTEIN_TRANSPORT
GOCC_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_COMPLEX, GOCC_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_COMPLEX
REACTOME_REGULATION_OF_PTEEN_MRNA_TRANSLATION, REACTOME_REGULATION_OF_PTEEN_MRNA_TRANSLATION
MIR181A_3P, MIR181A_3P
GOBP_REGULATION_OF_T_CIRCLE_FORMATION, GOBP_REGULATION_OF_T_CIRCLE_FORMATION
BUSSLINGER_DUODENAL_K_CELLS, BUSSLINGER_DUODENAL_K_CELLS
HP_ERYTHROCYTE_INCLUSION_BODIES, HP_ERYTHROCYTE_INCLUSION_BODIES
HP_EXERCISE_INDUCED_MUSCLE_FATIGUE, HP_EXERCISE_INDUCED_MUSCLE_FATIGUE
GOBP_REGULATION_OF_PROTEIN_COMPLEX_STABILITY, GOBP_REGULATION_OF_PROTEIN_COMPLEX_STABILITY
HP_LACK_OF_SKIN_ELASTICITY, HP_LACK_OF_SKIN_ELASTICITY
MIR1277_3P, MIR1277_3P
GOMF_VOLUME_SENSITIVE_ANION_CHANNEL_ACTIVITY, GOMF_VOLUME_SENSITIVE_ANION_CHANNEL_ACTIVITY
GOMF_7S_RNA_BINDING, GOMF_7S_RNA_BINDING
GOCC_PIP_BODY, GOCC_PIP_BODY
GOBP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS, GOBP_TRANSCRIPTION_FACTOR_CATABOLIC_PROCESS
GOMF_ATPASE_INHIBITOR_ACTIVITY, GOMF_ATPASE_INHIBITOR_ACTIVITY
HP_ARTERIAL_DISSECTION, HP_ARTERIAL_DISSECTION
GOBP_POSITIVE_REGULATION_OF_T_CELL_TOLERANCE_INDUCION, GOBP_POSITIVE_REGULATION_OF_T_CELL_TOLERANCE_INDUCION
HOLLEMAN_ASPARAGINASE_RESISTANCE_B_ALL_DN, HOLLEMAN_ASPARAGINASE_RESISTANCE_B_ALL_DN
HP_STRIAE_DISTENSAE, HP_STRIAE_DISTENSAE
WP_RENIN_ANGIOTENSIN_ALDOSTERONE_SYSTEM_RAAS, WP_RENIN_ANGIOTENSIN_ALDOSTERONE_SYSTEM_RAAS
HP_RIGHT_VENTRICULAR_DILATATION, HP_RIGHT_VENTRICULAR_DILATATION
GOMF_PEROXISOME_TARGETING_SEQUENCE_BINDING, GOMF_PEROXISOME_TARGETING_SEQUENCE_BINDING
HP_OPTIC_NERVE_DYSPLASIA, HP_OPTIC_NERVE_DYSPLASIA
GOBP_TAURINE_TRANSPORT, GOBP_TAURINE_TRANSPORT
GOBP_CHONDROCYTE_DIFFERENTIATION_INVOLVED_IN_ENDOCHONDRAL_BONE_MORPHOGENESIS, GOBP_CHONDROCYTE_DIFFERENTIATION_INVOLVED_IN_ENDOCHONDRAL_BONE_MORPHOGENESIS
GOBP_SULFUR_COMPOUND_TRANSPORT, GOBP_SULFUR_COMPOUND_TRANSPORT
GOBP_CARDIAC_LEFT_VENTRICLE_MORPHOGENESIS, GOBP_CARDIAC_LEFT_VENTRICLE_MORPHOGENESIS
GOBP_PROTEIN_SIDE_CHAIN DEGLUTAMYLATION, GOBP_PROTEIN_SIDE_CHAIN DEGLUTAMYLATION
GOMF_RRNA_CYTOSINE_METHYLTRANSFERASE_ACTIVITY, GOMF_RRNA_CYTOSINE_METHYLTRANSFERASE_ACTIVITY
WP_MITOCHONDRIAL_FATTY_ACID_SYNTHESIS_PATHWAY, WP_MITOCHONDRIAL_FATTY_ACID_SYNTHESIS_PATHWAY
GOBP_ASSEMBLY_OF_ACTOMYOSIN_APPARATUS_INVOLVED_IN_CYTOKINESIS, GOBP_ASSEMBLY_OF_ACTOMYOSIN_APPARATUS_INVOLVED_IN_CYTOKINESIS
GOBP_POSITIVE_REGULATION_OF_FC_GAMMA_RECEPTOR_SIGNALING_PATHWAY_INVOLVED_IN_PHA, GOBP_POSITIVE_REGULATION_OF_FC_GAMMA_RECEPTOR_SIGNALING_PATHWAY_INVOLVED_IN_PHA
GOCC_AP_4_ADAPTOR_COMPLEX, GOCC_AP_4_ADAPTOR_COMPLEX
GOMF_AP_1_ADAPTOR_COMPLEX_BINDING, GOMF_AP_1_ADAPTOR_COMPLEX_BINDING
GOBP_PROTEIN DEGLUTAMYLATION, GOBP_PROTEIN DEGLUTAMYLATION
REACTOME_DEFECTS_IN_BIOTIN_BTEN_METABOLISM, REACTOME_DEFECTS_IN_BIOTIN_BTEN_METABOLISM
HP_TORSION_DYSTONIA, HP_TORSION_DYSTONIA
GOBP_NERVE_DEVELOPMENT, GOBP_NERVE_DEVELOPMENT
REACTOME_GRB2_SOS_PROVIDES_LINKAGE_TO_MAPK_SIGNALING_FOR_INTEGRINS, REACTOME_GRB2_SOS_PROVIDES_LINKAGE_TO_MAPK_SIGNALING_FOR_INTEGRINS
GOBP_PROTEIN_LOCALIZATION_TO_PRESYNAPSE, GOBP_PROTEIN_LOCALIZATION_TO_PRESYNAPSE
GOBP_REGULATION_OF_NITRIC_OXIDE_SYNTHASE_ACTIVITY, GOBP_REGULATION_OF_NITRIC_OXIDE_SYNTHASE_ACTIVITY
ZNF136_TARGET_GENES, ZNF136_TARGET_GENES
HP_REPETITIVE_COMPULSIVE_BEHAVIOR, HP_REPETITIVE_COMPULSIVE_BEHAVIOR
HP_MESOAXIAL_POLYDACTYL, HP_MESOAXIAL_POLYDACTYL
GOBP_HEART_VALVE_DEVELOPMENT, GOBP_HEART_VALVE_DEVELOPMENT
HP_NEPHROGENIC_DIABETES_INSIPIDUS, HP_NEPHROGENIC_DIABETES_INSIPIDUS
GOBP_POSITIVE_REGULATION_OF_STRIATED_MUSCLE_CELL_APOPTOTIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_STRIATED_MUSCLE_CELL_APOPTOTIC_PROCESS
GOBP_PIGMENT_ACCUMULATION, GOBP_PIGMENT_ACCUMULATION
HP_DENTAL_ENAMEL_PITS, HP_DENTAL_ENAMEL_PITS
GOBP_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_INVOLVED_IN_ENDOCARDIAL_CUSHION_FORMATION, GOBP_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION_INVOLVED_IN_ENDOCARDIAL_CUSHION_FORMATION
GALIE_TUMOR_STEMNESS_GENES, GALIE_TUMOR_STEMNESS_GENES
GOBP_TOLERANCE_INDUCION_TO_SELF_ANTIEN, GOBP_TOLERANCE_INDUCION_TO_SELF_ANTIEN
GOBP_PROTEIN_IMPORT_INTO_PEROXISOME_MEMBRANE, GOBP_PROTEIN_IMPORT_INTO_PEROXISOME_MEMBRANE
GOMF_RNA_POLYMERASE_I_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY, GOMF_RNA_POLYMERASE_I_GENERAL_TRANSCRIPTION_INITIATION_FACTOR_ACTIVITY
GOBP_REGULATION_OF_MONOOXYGENASE_ACTIVITY, GOBP_REGULATION_OF_MONOOXYGENASE_ACTIVITY
GOMF_N_ACETYLLACTOSAMINIDE_BETA_1_3_N_ACETYLLGLUCOSAMINYLTRANSFERASE_ACTIVITY, GOMF_N_ACETYLLACTOSAMINIDE_BETA_1_3_N_ACETYLLGLUCOSAMINYLTRANSFERASE_ACTIVITY
GOBP_RRNA_BASE_METHYLATION, GOBP_RRNA_BASE_METHYLATION
GOBP_C_TERMINAL_PROTEIN_AMINO_ACID_MODIFICATION, GOBP_C_TERMINAL_PROTEIN_AMINO_ACID_MODIFICATION
HP_ABNORMAL_COCCYX_MORPHOLOGY, HP_ABNORMAL_COCCYX_MORPHOLOGY
GOBP_GROWTH_PLATE_CARTILAGE_CHONDROCYTE_DIFFERENTIATION, GOBP_GROWTH_PLATE_CARTILAGE_CHONDROCYTE_DIFFERENTIATION
GOBP_REGULATION_OF_NEUROTRANSMITTER_RECEPTOR_LOCALIZATION_TO_POSTSYNAPTIC_SPECIALIZATION_MEMBRANE, GOBP_REGULATION_OF_NEUROTRANSMITTER_RECEPTOR_LOCALIZATION_TO_POSTSYNAPTIC_SPECIALIZATION_MEMBRANE
HP_THICKENED_EARS, HP_THICKENED_EARS
GOCC_PL_BODY, GOCC_PL_BODY
GOBP_LUNG_LOBE_DEVELOPMENT, GOBP_LUNG_LOBE_DEVELOPMENT
HP_RENAL_HAMARTOMA, HP_RENAL_HAMARTOMA
FARMER_BREAST_CANCER_CLUSTER_8, FARMER_BREAST_CANCER_CLUSTER_8
GOBP_BASE_EXCISION_REPAIR_AP_SITE_FORMATION, GOBP_BASE_EXCISION_REPAIR_AP_SITE_FORMATION