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LAKE ADULT KIDNEY C27 VASCULAR SMOOTH MUSCLE CELLS AND PERICYTES, LAKE ADULT KIDNEY C27 VASCULAR SMOOTH MUSCLE CELLS AND PERICYTES
 HALLMARK_EPITHELIAL_MESENCHYMAL_TRANSITION, HALLMARK_EPITHELIAL_MESENCHYMAL_TRANSITION
 MIR199A_3P_MIR199B_3P, MIR199A_3P_MIR199B_3P
 MIR3129_5P, MIR3129_5P
 WU_CELL_MIGRATION, WU_CELL_MIGRATION
 GSE5589 WT VS IL6 KO LPS AND IL6 STIM MACROPHAGE 45MIN DN, GSE5589 WT VS IL6 KO LPS AND IL6 STIM MACROPHAGE 45MIN DN
 AKT UP.V1 UP, AKT UP.V1 UP
GOBP_CALCIUM_MEDIATED_SIGNALING, GOBP_CALCIUM_MEDIATED_SIGNALING
ZHENG CORD BLOOD C2 PUTATIVE BASOPHIL EOSINOPHIL MAST CELL PROGENITOR, ZHENG CORD BLOOD C2 PUTATIVE BASOPHIL EOSINOPHIL MAST CELL PROGENITOR
 MODULE 122, MODULE 122
VART_KSHV_INFECTION_ANGIOGENIC_MARKERS_UP, VART_KSHV_INFECTION_ANGIOGENIC_MARKERS_UP
GOMF_EXTRACELLULAR_MATRIX_STRUCTURAL_CONSTITUENT, GOMF_EXTRACELLULAR_MATRIX_STRUCTURAL_CONSTITUENT
HALLMARK_MYOGENESIS, HALLMARK_MYOGENESIS
GSE16385_ROSIGLITAZONE_VS_UNTREATED_IFNG_TNF_STIM_MACROPHAGE_UP, GSE16385_ROSIGLITAZONE_VS_UNTREATED_IFNG_TNF_STIM_MACROPHAGE_UP
ESC_V6.5_UP_LATE.V1_UP, ESC_V6.5_UP_LATE.V1_UP
MIR4670_3P, MIR4670_3P
GOBP_ACTIN_FILAMENT_BASED_MOVEMENT, GOBP_ACTIN_FILAMENT_BASED_MOVEMENT
CHIARADONNA_NEOPLASTIC_TRANSFORMATION_CDC25_UP, CHIARADONNA_NEOPLASTIC_TRANSFORMATION_CDC25_UP
GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY, GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY
REACTOME MUSCLE CONTRACTION, REACTOME MUSCLE CONTRACTION
FAN_EMBRYONIC_CTX_EX_4_EXCITATORY_NEURON, FAN_EMBRYONIC_CTX_EX_4_EXCITATORY_NEURON
GOBP_STRIATED_MUSCLE_CONTRACTION, GOBP_STRIATED_MUSCLE_CONTRACTION
GOBP_ACTION_POTENTIAL, GOBP_ACTION_POTENTIAL
FAN_EMBRYONIC_CTX_ASTROCYTE_2, FAN_EMBRYONIC_CTX_ASTROCYTE_2
GOCC_BASEMENT_MEMBRANE, GOCC_BASEMENT_MEMBRANE
GOMF_GROWTH_FACTOR_BINDING, GOMF_GROWTH_FACTOR_BINDING
GSE3982 BCELL VS EFF MEMORY CD4 TCELL DN, GSE3982 BCELL VS EFF MEMORY CD4 TCELL DN
AIZARANI LIVER C33 STELLATE CELLS 2, AIZARANI LIVER C33 STELLATE CELLS 2
KEGG ECM RECEPTOR INTERACTION, KEGG ECM RECEPTOR INTERACTION
GOBP_ADHERENS_JUNCTION_ORGANIZATION, GOBP_ADHERENS_JUNCTION_ORGANIZATION
REACTOME_ECM_PROTEOGLYCANS, REACTOME_ECM_PROTEOGLYCANS
REACTOME INTEGRIN CELL SURFACE INTERACTIONS, REACTOME INTEGRIN CELL SURFACE INTERACTIONS
GOCC_FILOPODIUM, GOCC_FILOPODIUM
REACTOME_CELL_CELL_COMMUNICATION, REACTOME_CELL_CELL_COMMUNICATION
GOBP_SODIUM_ION_TRANSMEMBRANE_TRANSPORT, GOBP_SODIUM_ION_TRANSMEMBRANE_TRANSPORT
GOCC_SARCOPLASM, GOCC_SARCOPLASM
GOBP_CARDIAC_CONDUCTION, GOBP_CARDIAC_CONDUCTION
TRAVAGLINI_LUNG_ARTERY_CELL, TRAVAGLINI_LUNG_ARTERY_CELL
TOMLINS_PROSTATE_CANCER_DN, TOMLINS_PROSTATE_CANCER_DN
HP_WADDLING GAIT, HP_WADDLING_GAIT
GOBP CYTOSOLIC CALCIUM ION TRANSPORT, GOBP CYTOSOLIC CALCIUM ION TRANSPORT
 MENON_FETAL_KIDNEY_3_STROMAL_CELLS, MENON_FETAL_KIDNEY_3_STROMAL_CELLS
 GOCC_SARCOLEMMA, GOCC_SARCOLEMMA
 GOBP CALCIUM ION TRANSPORT INTO CYTOSOL, GOBP CALCIUM ION TRANSPORT INTO CYTOSOL
 GOBP REGULATION OF PROTEIN DEPOLYMERIZATION, GOBP REGULATION OF PROTEIN DEPOLYMERIZATION
 MIR450A_2_3P, MIR450A_2_3P
 GOBP_NEURON_PROJECTION_EXTENSION, GOBP_NEURON_PROJECTION_EXTENSION
GOBP REGULATION OF STEROID BIOSYNTHETIC PROCESS, GOBP REGULATION OF STEROID BIOSYNTHETIC PROCESS
BRUECKNER_TARGETS_OF_MIRLET7A3_DN, BRUECKNER_TARGETS_OF_MIRLET7A3_DN
GOBP_APPENDAGE_DEVELOPMENT, GOBP_APPENDAGE_DEVELOPMENT
GOBP_CELL_CELL_JUNCTION_ASSEMBLY, GOBP_CELL_CELL_JUNCTION_ASSEMBLY
IWANAGA_CARCINOGENESIS_BY_KRAS_UP, IWANAGA_CARCINOGENESIS_BY_KRAS_UP
MIR4714_5P, MIR4714_5P
REACTOME NON INTEGRIN MEMBRANE ECM INTERACTIONS, REACTOME NON INTEGRIN MEMBRANE ECM INTERACTIONS
GOBP_IMPORT_ACROSS_PLASMA_MEMBRANE, GOBP_IMPORT_ACROSS_PLASMA_MEMBRANE
BERENJENO ROCK SIGNALING NOT VIA RHOA DN, BERENJENO ROCK SIGNALING NOT VIA RHOA DN
MIR3925 5P, MIR3925 5P
 GSE9006_1MONTH_VS_4MONTH_POST_TYPE_1_DIABETES_DX_PBMC_DN, GSE9006_1MONTH_VS_4MONTH_POST_TYPE_1_DIABETES_DX_PBMC_DN
 REACTOME_LAMININ_INTERACTIONS, REACTOME_LAMININ_INTERACTIONS
GOBP_AXON_EXTENSION, GOBP_AXON_EXTENSION
GOBP_CARDIAC_VENTRICLE_DEVELOPMENT, GOBP_CARDIAC_VENTRICLE_DEVELOPMENT
GOBP_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL, GOBP_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL
GOMF_CELL_ADHESION_MEDIATOR_ACTIVITY, GOMF_CELL_ADHESION_MEDIATOR_ACTIVITY
GOCC PLASMA MEMBRANE RAFT, GOCC PLASMA MEMBRANE RAFT
GOBP_NEURAL_PRECURSOR_CELL_PROLIFERATION, GOBP_NEURAL_PRECURSOR_CELL_PROLIFERATION
GOMF_SODIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_SODIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOBP_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL_INVOLVED_IN_CONTRACTION, GOBP_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL_INVOLVED_IN_CONTRACTION
PID_INTEGRIN1_PATHWAY, PID_INTEGRIN1_PATHWAY
GOBP_REGULATION_OF_CATION_CHANNEL_ACTIVITY, GOBP_REGULATION_OF_CATION_CHANNEL_ACTIVITY
GSE11961_MARGINAL_ZONE_BCELL_VS_MEMORY_BCELL_DAY7_DN, GSE11961_MARGINAL_ZONE_BCELL_VS_MEMORY_BCELL_DAY7_DN
DESCARTES_FETAL_EYE_SKELETAL_MUSCLE_CELLS, DESCARTES_FETAL_EYE_SKELETAL_MUSCLE_CELLS
GOMF_CELL_CELL_ADHESION_MEDIATOR_ACTIVITY, GOMF_CELL_CELL_ADHESION_MEDIATOR_ACTIVITY
GOBP_BASEMENT_MEMBRANE_ORGANIZATION, GOBP_BASEMENT_MEMBRANE_ORGANIZATION
 GSE40274_SATB1_VS_FOXP3_AND_SATB1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_SATB1_VS_FOXP3_AND_SATB1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
 REACTOME_RND3_GTPASE_CYCLE, REACTOME_RND3_GTPASE_CYCLE
 GOBP_VENTRICULAR_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL, GOBP_VENTRICULAR_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL
 REACTOME_ION_HOMEOSTASIS, REACTOME_ION_HOMEOSTASIS
GOBP_GASTRULATION, GOBP_GASTRULATION
 REACTOME_CARDIAC_CONDUCTION, REACTOME_CARDIAC_CONDUCTION
 HP_ABNORMALITY_OF_LOWER_LIMB_EPIPHYSIS_MORPHOLOGY, HP_ABNORMALITY_OF_LOWER_LIMB_EPIPHYSIS_MORPHOLOGY
 GOBP POSITIVE REGULATION OF CATION TRANSMEMBRANE TRANSPORT, GOBP POSITIVE REGULATION OF CATION TRANSMEMBRANE TRANSPORT
 GOBP_REGULATION_OF_HEART_RATE, GOBP_REGULATION_OF_HEART_RATE
 BEGUM_TARGETS_OF_PAX3_FOXO1_FUSION_DN, BEGUM_TARGETS_OF_PAX3_FOXO1_FUSION_DN
 HP_ATRIAL_ARRHYTHMIA, HP_ATRIAL_ARRHYTHMIA
 KORKOLA_YOLK_SAC_TUMOR, KORKOLA_YOLK_SAC_TUMOR
 HP_EPIPHYSEAL_DYSPLASIA, HP_EPIPHYSEAL_DYSPLASIA
 HP_DENTAL_CROWDING, HP_DENTAL_CROWDING
 NABA_BASEMENT_MEMBRANES, NABA_BASEMENT_MEMBRANES
 REACTOME_ACTIVATION_OF_MATRIX_METALLOPROTEINASES, REACTOME_ACTIVATION_OF_MATRIX_METALLOPROTEINASES
 CLASPER LYMPHATIC VESSELS DURING METASTASIS DN, CLASPER LYMPHATIC VESSELS DURING METASTASIS DN
REACTOME_RND1_GTPASE_CYCLE, REACTOME_RND1_GTPASE_CYCLE
 PLASARI_NFIC_TARGETS_BASAL_UP, PLASARI_NFIC_TARGETS_BASAL_UP
GOBP_VASCULOGENESIS, GOBP_VASCULOGENESIS
GOBP_NEGATIVE_REGULATION_OF_VASCULAR_PERMEABILITY, GOBP_NEGATIVE_REGULATION_OF_VASCULAR_PERMEABILITY
GOCC INTRINSIC COMPONENT OF POSTSYNAPTIC MEMBRANE, GOCC INTRINSIC COMPONENT OF POSTSYNAPTIC MEMBRANE
 GTCAACC_MIR3805P, GTCAACC_MIR3805P
 MODULE_329, MODULE_329
 LEE_TARGETS_OF_PTCH1_AND_SUFU_DN, LEE_TARGETS_OF_PTCH1_AND_SUFU_DN
 GOMF_EXTRACELLULAR_MATRIX_BINDING, GOMF_EXTRACELLULAR_MATRIX_BINDING
 GOBP_REGULATION_OF_CALCIUM_ION_TRANSPORT_INTO_CYTOSOL, GOBP_REGULATION_OF_CALCIUM_ION_TRANSPORT_INTO_CYTOSOL
GOBP POSITIVE REGULATION OF MORPHOGENESIS OF AN EPITHELIUM, GOBP POSITIVE REGULATION OF MORPHOGENESIS OF AN EPITHELIUM
 MODULE_387, MODULE_387
 GOBP_BUNDLE_OF_HIS_CELL_TO_PURKINJE_MYOCYTE_COMMUNICATION, GOBP_BUNDLE_OF_HIS_CELL_TO_PURKINJE_MYOCYTE_COMMUNICATION
 GOBP POSITIVE REGULATION OF BLOOD CIRCULATION, GOBP POSITIVE REGULATION OF BLOOD CIRCULATION
 GOBP REGULATION OF BIOMINERALIZATION, GOBP REGULATION OF BIOMINERALIZATION
GOBP_MEMBRANE_REPOLARIZATION_DURING_ACTION_POTENTIAL, GOBP_MEMBRANE_REPOLARIZATION_DURING_ACTION_POTENTIAL
 GOCC_NEURON_PROJECTION_MEMBRANE, GOCC_NEURON_PROJECTION_MEMBRANE
 GOMF_ACTININ_BINDING, GOMF_ACTININ_BINDING
 GOCC_T_TUBULE, GOCC_T_TUBULE
 GOMF EXTRACELLULAR MATRIX STRUCTURAL CONSTITUENT CONFERRING TENSILE STRENGTH, GOMF EXTRACELLULAR MATRIX STRUCTURAL CONSTITUENT CONFERRING TENSILE STRENGTH
 HP VENTRICULAR FIBRILLATION, HP VENTRICULAR FIBRILLATION
PACHER_TARGETS_OF_IGF1_AND_IGF2_UP, PACHER_TARGETS_OF_IGF1_AND_IGF2_UP
GOBP_CELL_CELL_SIGNALING_INVOLVED_IN_CARDIAC_CONDUCTION, GOBP_CELL_CELL_SIGNALING_INVOLVED_IN_CARDIAC_CONDUCTION
GOBP_REGULATION_OF_CARDIAC_CONDUCTION, GOBP_REGULATION_OF_CARDIAC_CONDUCTION
 GOBP_REGULATION_OF_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE, GOBP_REGULATION_OF_MORPHOGENESIS_OF_A_BRANCHING_STRUCTURE
REACTOME_TRANSPORT_OF_BILE_SALTS_AND_ORGANIC_ACIDS_METAL_IONS_AND_AMINE_COMPOUNDS, REACTOME_TRANSPORT_ORGANIC_ACIDS_METAL_IONS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANICACIDS_AND_ORGANIC
GOBP_ATRIAL_CARDIAC_MUSCLE_CELL_TO_AV_NODE_CELL_SIGNALING, GOBP_ATRIAL_CARDIAC_MUSCLE_CELL_TO_AV_NODE_CELL_SIGNALING
GOMF_STRUCTURAL_CONSTITUENT_OF_MUSCLE, GOMF_STRUCTURAL_CONSTITUENT_OF_MUSCLE
REACTOME_MET_ACTIVATES_PTK2_SIGNALING, REACTOME_MET_ACTIVATES_PTK2_SIGNALING
 MODULE_201, MODULE_201
GOBP_METAL_ION_EXPORT, GOBP_METAL_ION_EXPORT
HP_CENTRAL_OPACIFICATION_OF_THE_CORNEA, HP_CENTRAL_OPACIFICATION_OF_THE_CORNEA
GOBP_POSITIVE_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORT
 TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_ERYTHROCYTE_DN, TONKS_TARGETS_OF_RUNX1_RUNX1T1_FUSION_ERYTHROCYTE_DN
GOCC_COLLAGEN_TRIMER, GOCC_COLLAGEN_TRIMER
GOBP_POSITIVE_REGULATION_OF_POTASSIUM_ION_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_POTASSIUM_ION_TRANSPORT
 GOMF_PROTEIN_BINDING_INVOLVED_IN_HETEROTYPIC_CELL_CELL_ADHESION, GOMF_PROTEIN_BINDING_INVOLVED_IN_HETEROTYPIC_CELL_CELL_ADHESION
NABA_COLLAGENS, NABA_COLLAGENS
 REACTOME_COLLAGEN_CHAIN_TRIMERIZATION, REACTOME_COLLAGEN_CHAIN_TRIMERIZATION
WP_DEVELOPMENT_OF_URETERIC_COLLECTION_SYSTEM, WP_DEVELOPMENT_OF_URETERIC_COLLECTION_SYSTEM
REACTOME_COLLAGEN_BIOSYNTHESIS_AND_MODIFYING_ENZYMES, REACTOME_COLLAGEN_BIOSYNTHESIS_AND_MODIFYING_ENZYMES
GSE9946_MATURE_STIMULATORY_VS_LISTERIA_INF_MATURE_DC_DN, GSE9946_MATURE_STIMULATORY_VS_LISTERIA_INF_MATURE_DC_DN
 GOBP_HETEROPHILIC_CELL_CELL_ADHESION_VIA_PLASMA_MEMBRANE_CELL_ADHESION_MOLECULES, GOBP_HETEROPHILIC_CELL_CELL_ADHESION_VIA_PLASMA_MEMBRANE_CELL_ADHESION_MOLECULES
GOBP_REGULATION_OF_MORPHOGENESIS_OF_AN_EPITHELIUM, GOBP_REGULATION_OF_MORPHOGENESIS_OF_AN_EPITHELIUM
 GOBP_EXCITATORY_SYNAPSE_ASSEMBLY, GOBP_EXCITATORY_SYNAPSE_ASSEMBLY
 MODULE_512, MODULE_512
WP_MATRIX_METALLOPROTEINASES, WP_MATRIX_METALLOPROTEINASES
GOBP REGULATION OF ENDOTHELIAL CELL DIFFERENTIATION, GOBP REGULATION OF ENDOTHELIAL CELL DIFFERENTIATION
GOBP_POSITIVE_REGULATION_OF_OSSIFICATION, GOBP_POSITIVE_REGULATION_OF_OSSIFICATION
GOMF_ANKYRIN_BINDING, GOMF_ANKYRIN_BINDING
 GOBP_MEMBRANE_DEPOLARIZATION_DURING_ACTION_POTENTIAL, GOBP_MEMBRANE_DEPOLARIZATION_DURING_ACTION_POTENTIAL
 GOBP_MEMBRANE_REPOLARIZATION, GOBP_MEMBRANE_REPOLARIZATION
GOBP CARDIAC MUSCLE CELL MEMBRANE REPOLARIZATION, GOBP CARDIAC MUSCLE CELL MEMBRANE REPOLARIZATION
 MODULE 178, MODULE 178
GOBP_COLLAGEN_ACTIVATED_SIGNALING_PATHWAY, GOBP_COLLAGEN_ACTIVATED_SIGNALING_PATHWAY
CUI_DEVELOPING_HEART_LEFT_ATRIAL_CARDIOMYOCYTE, CUI_DEVELOPING_HEART_LEFT_ATRIAL_CARDIOMYOCYTE
HP_EMG_IMPAIRED_NEUROMUSCULAR_TRANSMISSION, HP_EMG_IMPAIRED_NEUROMUSCULAR_TRANSMISSION
ITO_PTTG1_TARGETS_UP, ITO_PTTG1_TARGETS_UP
MODULE_14, MODULE_14
WP MIRNA TARGETS IN ECM AND MEMBRANE RECEPTORS, WP MIRNA TARGETS IN ECM AND MEMBRANE RECEPTORS
LIU_VAV3_PROSTATE_CARCINOGENESIS_DN, LIU_VAV3_PROSTATE_CARCINOGENESIS_DN
GOBP_EXPORT_ACROSS_PLASMA_MEMBRANE, GOBP_EXPORT_ACROSS_PLASMA_MEMBRANE
HP_ST_SEGMENT_ELEVATION, HP_ST_SEGMENT_ELEVATION
HP_FLAT_CAPITAL_FEMORAL_EPIPHYSIS, HP_FLAT_CAPITAL_FEMORAL_EPIPHYSIS
HP_ABNORMAL_ST_SEGMENT, HP_ABNORMAL_ST_SEGMENT
GOBP MEMBRANE DEPOLARIZATION DURING CARDIAC MUSCLE CELL ACTION POTENTIAL, GOBP MEMBRANE DEPOLARIZATION DURING CARDIAC MUSCLE CELL ACTION POTENTIAL
HP_IRREGULAR_EPIPHYSES, HP_IRREGULAR_EPIPHYSES
GOMF_STRUCTURAL_MOLECULE_ACTIVITY_CONFERRING_ELASTICITY, GOMF_STRUCTURAL_MOLECULE_ACTIVITY_CONFERRING_ELASTICITY
LIM_MAMMARY_LUMINAL_PROGENITOR_DN, LIM_MAMMARY_LUMINAL_PROGENITOR_DN
HP_EMG_DECREMENTAL_RESPONSE_OF_COMPOUND_MUSCLE_ACTION_POTENTIAL_TO_REPETITIVE_NERVE_STIMULATION, HP_EMG_DECREMENTAL_RESPONSE_OF_COMPOUND_MUSCLE_ACTION_POTENTIAL_1
GOBP_AV_NODE_CELL_TO_BUNDLE_OF_HIS_CELL_COMMUNICATION, GOBP_AV_NODE_CELL_TO_BUNDLE_OF_HIS_CELL_COMMUNICATION
GOBP_POSITIVE_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
ROY WOUND BLOOD VESSEL DN, ROY WOUND BLOOD VESSEL DN
GNF2 MYL2, GNF2 MYL2
HP FIRST DEGREE ATRIOVENTRICULAR BLOCK, HP FIRST DEGREE ATRIOVENTRICULAR BLOCK
HP MIDLINE_DEFECT_OF_THE_NOSE, HP_MIDLINE_DEFECT_OF_THE_NOSE
GOBP_REGULATION_OF_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY
REACTOME_CROSSLINKING_OF_COLLAGEN_FIBRILS, REACTOME_CROSSLINKING_OF_COLLAGEN_FIBRILS
HP SICK SINUS SYNDROME, HP SICK SINUS SYNDROME
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KANG_AR_TARGETS_UP, KANG_AR_TARGETS_UP

BIOCARTA_PLATELETAPP_PATHWAY, BIOCARTA_PLATELETAPP_PATHWAY

GOBP_POTASSIUM_ION_HOMEOSTASIS, GOBP_POTASSIUM_ION_HOMEOSTASIS

GOBP_EMBRYONIC_SKELETAL_JOINT_MORPHOGENESIS, GOBP_EMBRYONIC_SKELETAL_JOINT_MORPHOGENESIS

MIKKELSEN_MEF_ICP_WITH_H3K4ME3_AND_H3K27ME3, MIKKELSEN_MEF_ICP_WITH_H3K4ME3_AND_H3K27ME3

GOBP_POSITIVE_REGULATION_OF_KERATINOCYTE_PROLIFERATION, GOBP_POSITIVE_REGULATION_OF_KERATINOCYTE_PROLIFERATION

GOBP CALCIUM ION TRANSMEMBRANE TRANSPORT VIA HIGH VOLTAGE GATED CALCIUM CHANNEL, GOBP CALCIUM ION TRANSMEMBRANE TRANSPORT VIA HIGH VOLTAGE GATED CALCIUM CHANN

GOBP_REGULATION_OF_ATRIAL_CARDIAC_MUSCLE_CELL_MEMBRANE_DEPOLARIZATION, GOBP_REGULATION_OF_ATRIAL_CARDIAC_MUSCLE_CELL_MEMBRANE_DEPOLARIZATION

GOBP_EMBRYONIC_FORELIMB_MORPHOGENESIS, GOBP_EMBRYONIC_FORELIMB_MORPHOGENESIS

GOBP_PULMONARY_VALVE_MORPHOGENESIS, GOBP_PULMONARY_VALVE_MORPHOGENESIS

HP_HYPOGLYCEMIC_SEIZURES, HP_HYPOGLYCEMIC_SEIZURES

BIOCARTA_AMI_PATHWAY, BIOCARTA_AMI_PATHWAY

HP_ACETABULAR_SPURS, HP_ACETABULAR_SPURS

HP_CRYPTOPHTHALMOS, HP_CRYPTOPHTHALMOS HP ATRIAL STANDSTILL, HP ATRIAL STANDSTILL

BIOCARTA_INTRINSIC_PATHWAY, BIOCARTA_INTRINSIC_PATHWAY

HP_SMALL_EPIPHYSES, HP_SMALL_EPIPHYSES

CULAR ATRIAL CARDIOMYOCYTE, CUI DEVELOPING HEART TRABECULAR ATRIAL CARDIOMYOCYTE