Nerve sheath tumors pindle and epithelioid tumo Gliomas Glioneuronal tumors c2.WP OXYSTEROLS DERIVED FROM CHOLESTEROL c2.WP PROSTAGLANDIN SIGNALING c2.REACTOME FORMATION OF THE NEPHRIC DUCT c2.REACTOME PYRIMIDINE CATABOLISM c2.REACTOME CELLULAR HEXOSE TRANSPORT c2.FINETTI BREAST CANCER KINOME RED c2.SU SALIVARY GLAND
c2.KEGG MEDICUS REFERENCE GPCR PLCB ITPR SIGNALING PATHWAY
c2.WP DIFFERENTIATION OF WHITE AND BROWN ADIPOCYTE c2.KEGG STEROID HORMONE BIOSYNTHESIS c2.LI ESTROGENE MID E2 RESPONSE UP c2.BIOCARTA INFLAM PATHWAY c2.KEGG MEDICUS REFERENCE V D J RECOMBINATION c2.LEE NAIVE T LYMPHOCYTE c2.REACTOME FORMATION OF THE CORNIFIED ENVELOPE c2.REACTOME KERATINIZATION
c2.REACTOME TLR3 MEDIATED TICAM1 DEPENDENT PROGRAMMED CELL DEATH
c2.REACTOME DISPLACEMENT OF DNA GLYCOSYLASE BY APEX1 c2.KEGG HEMATOPOIETIC CELL LINEAGE c2.HUMMERICH BENIGN SKIN TUMOR DN c2.IGARASHI ATF4 TARGETS UP c2.KEGG MEDICUS REFERENCE BLOOD GROUP H O ANTIGEN TYPE 1 BIOSYNTHESIS c2.BIOCARTA CTL PATHWAY c2.WP TAMOXIFEN METABOLISM c2.REACTOME CLASS C 3 METABOTROPIC GLUTAMATE PHEROMONE RECEPTORS c2.REACTOME PHOSPHOLIPASE C MEDIATED CASCADE FGFR4 c2.MIKKELSEN ES LCP WITH H3K4ME3 AND H3K27ME3 c2.WP DEREGULATION OF RAB AND RAB EFFECTOR GENES IN BLADDER CANCER c2.BENPORATH PRC2 TARGETS c2.WP NUCLEOTIDE GPCRS c2.REACTOME NA CL DEPENDENT NEUROTRANSMITTER TRANSPORTERS
c2.REACTOME PHASE 1 INACTIVATION OF FAST NA CHANNELS c2.WP BIOGENIC AMINE SYNTHESIS c2.MIKKELSEN IPS ICP WITH H3K27ME3 c2.REACTOME RESOLUTION OF D LOOP STRUCTURES c2.KANG GLIS3 TARGETS c2.REACTOME ERBB2 ACTIVATES PTK6 SIGNALING
c2.REACTOME ERBB2 ACTIVATES PTK6 SIGNALING
c2.RICKMAN HEAD AND NECK CANCER E
c2.KEGG MEDICUS REFERENCE SKELETAL TYPE VGCC RYR SIGNALING
c2.BANG VERTEPORFIN ENDOMETRIAL CANCER CELLS DN
c2.REACTOME CASPASE ACTIVATION VIA DEATH RECEPTORS IN THE PRESENCE OF LIGAND c2.WP GLYOXYLATE METABOLISM c2.BIOCARTA NKT PATHWAY c2.REACTOME RESOLUTION OF D LOOP STRUCTURES THROUGH SYNTHESIS DEPENDENT STRAND ANNEALING SDSA c2.WP FOXP3 IN COVID19 c2.FERRANDO TAL1 NEIGHBORS c2.MEBARKI HCC PROGENITOR WNT DN BLOCKED BY FZD8CRD c2.WP DOPAMINERGIC NEUROGENESIS c2.WP SMALL LIGAND GPCRS c2.FUJIWARA PARK2 HEPATOCYTE PROLIFERATION UP c2.BASSO HAIRY CELL LEUKEMIA DN c2.PID CONE PATHWAY c2.BALLIF DEVELOPMENTAL DISABILITY P16 P12 DELETION c2.BIOCARTA ATRBRCA PATHWAY c2.WP GLUCOCORTICOID BIOSYNTHESIS c2.REACTOME ANDROGEN BIOSYNTHESIS c2.SCHLESINGER METHYLATED IN COLON CANCER c2.REACTOME TRP CHANNELS c2.BYSTRYKH HEMATOPOIESIS STEM CELL RUNX1 0 c2.STEGER ADIPOGENESIS UP c2.REACTOME ATORVASTATIN ADME c2.KEGG MEDICUS REFERENCE WNT SIGNALING MODULATION LGR RSPO c2.REACTOME NEPHRON DEVELOPMENT c2.REACTOME DEFECTIVE FACTOR IX CAUSES HEMOPHILIA B c2.REACTOME REGULATION BY C FLIP c2.NABA SECRETED FACTORS
c2.KEGG MEDICUS REFERENCE LEWIS X ANTIGEN BIOSYNTHESIS
c2.KEGG MEDICUS REFERENCE BMP SIGNALING PATHWAY BMP ANTAGONIST **• (1)** c2.REACTOME PD 1 SIGNALING
c2.REACTOME PD 1 SIGNALING
c2.FLORIO NEOCORTEX BASAL RADIAL GLIA UP
c2.REACTOME TNF RECEPTOR SUPERFAMILY TNFSF MEMBERS MEDIATING NON CANONICAL NF KB PATHWAY
c2.REACTOME AFLATOXIN ACTIVATION AND DETOXIFICATION c2.KORKOLA CORRELATED WITH POU5F1 c2.WP GPCRS OTHER c2.LOPEZ MESOTHELIOMA SURVIVAL OVERALL UP c2.WP LACTO SERIES SPHINGOLIPID METABOLISM c2.REACTOME INTERACTION WITH CUMULUS CELLS AND THE ZONA PELLUCIDA c2.REACTOME PHOSPHATE BOND HYDROLYSIS BY NUDT PROTEINS c2.WP NEPHROGENESIS c2.REACTOME TRIF MEDIATED PROGRAMMED CELL DEATH
c2.REACTOME METABOLIC DISORDERS OF BIOLOGICAL OXIDATION ENZYMES
c2.KEGG MEDICUS VARIANT DUPLICATION OR MUTATION ACTIVATED FLT3 TO JAK STAT SIGNALING PATHWAY c2.ONDER CDH1 TARGETS 3 DN c2.WP KININKALLIKREIN PATHWAY c2.REACTOME THE RETINOID CYCLE IN CONES DAYLIGHT VISION c2.REACTOME THE RETINOID CYCLE IN CONES DATLIGHT VISION
c2.REACTOME FERTILIZATION
c2.MIKKELSEN MEF HCP WITH H3 UNMETHYLATED
c2.FUNG IL2 SIGNALING 2
c2.REACTOME DEFECTIVE RIPK1 MEDIATED REGULATED NECROSIS c2.MIKKELSEN IPS LCP WITH H3K4ME3 AND H3K27ME3 c2.MIKKELSEN IPS WITH HCP H3K27ME3 c2.KEGG HISTIDINE METABOLISM
c2.ANDERSEN CHOLANGIOCARCINOMA CLASS2
c2.KIM BIPOLAR DISORDER OLIGODENDROCYTE DENSITY CORR DN c2.AIGNER ZEB1 TARGETS
c2.KHETCHOUMIAN TRIM24 TARGETS DN
c2.REACTOME TRANSPORT OF ORGANIC ANIONS c2.WP OVERVIEW OF PROINFLAMMATORY AND PROFIBROTIC MEDIATORS c2.KORKOLA CHORIOCARCINOMA DN c2.REACTOME ABACAVIR ADME c2.MEISSNER BRAIN HCP WITH H3K4ME2 c2.WP EV RELEASE FROM CARDIAC CELLS AND THEIR FUNCTIONAL EFFECTS c2.WP ACE INHIBITOR PATHWAY c2.MIKKELSEN NPC ICP WITH H3K27ME3 c2.HOLLERN EMT BREAST TUMOR DN c2.KEGG TYPE I DIABETES MELLITUS c2.WP TGIF DISRUPTION OF SHH SIGNALING c2.BLANCO MELO RESPIRATORY SYNCYTIAL VIRUS INFECTION A594 CELLS DN
c2.LIU OVARIAN CANCER TUMORS AND XENOGRAFTS XDGS UP c2.WP SELECTIVE EXPRESSION OF CHEMOKINE RECEPTORS DURING TCELL POLARIZATION c2.MIKKELSEN MEF HCP WITH H3K27ME3 c2.WEBER METHYLATED HCP IN FIBROBLAST UP c2.KEGG MEDICUS REFERENCE REGULATION OF NLRP3 INFLAMMASOME SIGNALING PATHWAY NLRP3 INHIBITION c2.MEISSNER ES ICP WITH H3K4ME3 AND H3K27ME3 c2.KEGG MEDICUS REFERENCE NLRC4 INFLAMMASOME SIGNALING PATHWAY c2.CHEMELLO SOLEUS VS EDL MYOFIBERS UP c2.KEGG MEDICUS REFERENCE NON CANONICAL INFLAMMASOME SIGNALING PATHWAY c2.REACTOME MRNA EDITING C TO U CONVERSION c2.MIKKELSEN ES HCP WITH H3K27ME3
c2.HUMMERICH MALIGNANT SKIN TUMOR DN
c2.KEGG MEDICUS REFERENCE SEROTONIN METABOLISM • c2.LIU CDX2 TARGETS UP

c2.LIU CDX2 TARGETS UP

c2.KEGG MEDICUS REFERENCE DOUBLE HOLLIDAY JUNCTION RESOLUTION

c2.WP GENES ASSOCIATED WITH THE DEVELOPMENT OF RHEUMATOID ARTHRITIS

c2.MIKKELSEN MCV6 LCP WITH H3K27ME3

c2.WP NEUROTRANSMITTER DISORDERS c2.WP NEUROTRANSMITTER DISORDERS
c2.WANG NFKB TARGETS
c2.MEISSNER NPC HCP WITH H3K4ME2 AND H3K27ME3
c2.BIOCARTA RHODOPSIN PATHWAY
c2.SHIN B CELL LYMPHOMA CLUSTER 1 c2.REACTOME ANTIMICROBIAL PEPTIDES c2.LOPEZ MESOTHELIOMA SURVIVAL WORST VS BEST DN c2.MIKKELSEN MCV6 ICP WITH H3K27ME3 c2.BIOCARTA THITH2 PATHWAY c2.WP HFE EFFECT ON HEPCIDIN PRODUCTION c2.REACTOME FRUCTOSE CATABOLISM c2.WORSCHECH TUMOR REJECTION UP c2.BIOCARTA THELPER PATHWAY c2.REACTOME VOLTAGE GATED POTASSIUM CHANNELS c2.REACTOME IRAK4 DEFICIENCY TLR2 4 c2.SU LIVER c2.REACTOME TFAP2A ACTS AS A TRANSCRIPTIONAL REPRESSOR DURING RETINOIC ACID INDUCED CELL DIFFERENTIATION c2.REACTOME ACYL CHAIN REMODELLING OF PI c2.REACTOME GPCR LIGAND BINDING c2.WP ACETAMINOPHEN IN ANALGESIA AND ANTIPYRESIS c2.KUROZUMI RESPONSE TO ONCOCYTIC VIRUS c2.REACTOME FGFR3B LIGAND BINDING AND ACTIVATION c2.LOPEZ EPITHELIOID MESOTHELIOMA c2.BIOCARTA TCYTOTOXIC PATHWAY c2.VILIMAS NOTCH1 TARGETS DN c2.BIOCARTA TCRA PATHWAY c2.GUTIERREZ WALDENSTROEMS MACROGLOBULINEMIA 1 DN c2.REACTOME REGULATION OF TLR BY ENDOGENOUS LIGAND c2.REACTOME SULFIDE OXIDATION TO SULFATE c2.REACTOME FASL CD95L SIGNALING c2.KEGG MEDICUS PATHOGEN SHIGELLA IPAH7.8 TO NLRP3 INFLAMMASOME SIGNALING PATHWAY c2.REN MIF TARGETS DN c2.KEGG MEDICUS VARIANT MUTATION ACTIVATED SMO TO HEDGEHOG SIGNALING PATHWAY c2.KEGG MEDICUS VARIANT MUTATION INACTIVATED PTCH1 TO HEDGEHOG SIGNALING PATHWAY c2.REACTOME INTERLEUKIN 18 SIGNALING c2.REACTOME SYNTHESIS OF LIPOXINS LX c2.BLANCO MELO COVID19 SARS COV 2 INFECTION CALU3 CELLS DN c2.RICKMAN HEAD AND NECK CANCER D c2.REACTOME RRNA MODIFICATION IN THE MITOCHONDRION c2.KEGG MEDICUS REFERENCE SIALYL LEWIS X ANTIGEN BIOSYNTHESIS c2.KEGG MEDICUS REFERENCE TRANSPORT OF CALCIUM c2.REACTOME ACETYLCHOLINE INHIBITS CONTRACTION OF OUTER HAIR CELLS c2.REACTOME PHOSPHOLIPASE C MEDIATED CASCADE FGFR2 c2.KEGG ASTHMA c2.REACTOME SCAVENGING OF HEME FROM PLASMA
c2.BIOCARTA GRANULOCYTES PATHWAY
c2.WP CYTOKINES AND INFLAMMATORY RESPONSE c2.WP CYTOKINES AND INFLAMMATORY RESPONSE
c2.GRAHAM CML DIVIDING VS NORMAL DIVIDING DN
c2.REACTOME FORMATION OF INTERMEDIATE MESODERM
c2.REACTOME ORGANIC CATION ANION ZWITTERION TRANSPORT
c2.MEBARKI HCC PROGENITOR WNT DN CTNNB1 INDEPENDENT
c2.KEGG MEDICUS REFERENCE CA2 ENTRY LIGAND GATED CA2 CHANNEL
c2.MATHEW FANCONI ANEMIA GENES
c2.SAENZ DETOX PATHWAY AND CARCINOGENESIS DN c2.MIKKELSEN MEF LCP WITH H3K27ME3 c2.BIOCARTA BARD1 PATHWAY c2.KEGG NEUROACTIVE LIGAND RECEPTOR INTERACTION c2.BIOCARTA MSP PATHWAY c2.REACTOME MULTIFUNCTIONAL ANION EXCHANGERS c2.KUROZUMI RESPONSE TO ONCOCYTIC VIRUS AND CYCLIC RGD c2.DEBOSSCHER NFKB TARGETS REPRESSED BY GLUCOCORTICOIDS c2.HAHTOLA CTCL PATHOGENESIS c2.SHIN B CELL LYMPHOMA CLUSTER 6 c2.MEISSNER BRAIN HCP WITH H3K4ME2 AND H3K27ME3 c2.WP EFFECTS OF NITRIC OXIDE c2.REACTOME PEPTIDE LIGAND BINDING RECEPTORS c2.WP IDO METABOLIC PATHWAY c2.BIOCARTA ALTERNATIVE PATHWAY c2.REACTOME VLDL CLEARANCE c2.KORKOLA EMBRYONAL CARCINOMA c2.VALK AML WITH T 8 21 TRANSLOCATION c2.CROSBY E2F4 TARGETS
c2.BIOCARTA EICOSANOID PATHWAY
c2.REACTOME ALTERNATIVE COMPLEMENT ACTIVATION 2 c2.TUOMISTO TUMOR SUPPRESSION BY COL13A1 DN c2.BIOCARTA NO2IL12 PATHWAY
c2.WEBER METHYLATED HCP IN SPERM UP
c2.KEGG MEDICUS VARIANT MUTATION CAUSED ABERRANT PDYN TO TRANSPORT OF CALCIUM c2.BIOCARTA BLYMPHOCYTE PATHWAY c2.WP SARSCOV2 AND ACE2 RECEPTOR MOLECULAR MECHANISMS c2.REACTOME FGFR3 LIGAND BINDING AND ACTIVATION c2.REACTOME SCAVENGING BY CLASS B RECEPTORS c2.REACTOME CLASS A 1 RHODOPSIN LIKE RECEPTORS c2.REACTOME CD22 MEDIATED BCR REGULATION c2.REACTOME DEFECTIVE C1GALT1C1 CAUSES TNPS c2.MEISSNER NPC ICP WITH H3 UNMETHYLATED c2.BIOCARTA CYTOKINE PATHWAY c2.KONDO COLON CANCER HCP WITH H3K27ME3 c2.KEGG LINOLEIC ACID METABOLISM c2.RUNNE GENDER EFFECT UP c2.WP PEPTIDE GPCRS c2.GALI TP53 TARGETS APOPTOTIC DN c2.MIKKELSEN NPC WITH LCP H3K27ME3 c2.REACTOME FGFR2C LIGAND BINDING AND ACTIVATION c2.MEISSNER BRAIN HCP WITH H3 UNMETHYLATED TOME SYNTHESIS OF WYBUTOSINE AT G37 OF TRNA PHI c2.REACTOME SYNTHESIS OF WYBUTOSINE AT GS7 OF TRNA PHE
c2.REACTOME CHEMOKINE RECEPTORS BIND CHEMOKINES
c2.REACTOME ERYTHROCYTES TAKE UP OXYGEN AND RELEASE CARBON DIOXIDE
c2.REACTOME TNFS BIND THEIR PHYSIOLOGICAL RECEPTORS
c2.WP TRYPTOPHAN CATABOLISM LEADING TO NAD PRODUCTION
c2.MIKKELSEN ES LCP WITH H3K27ME3 c2.MIKKELSEN ÉS LCP WITH H3K27ME3
c2.ZHOU PANCREATIC BETA CELL
c2.REACTOME INTESTINAL ABSORPTION
c2.REACTOME BUTYROPHILIN BTN FAMILY INTERACTIONS
c2.REACTOME FGFR2B LIGAND BINDING AND ACTIVATION
c2.WP PATHOGENESIS OF SARSCOV2 MEDIATED BY NSP9NSP10 COMPLEX
c2.REACTOME PHASE 3 RAPID REPOLARISATION
c2.MIKKELSEN MEF ICP WITH H3K27ME3
c2.REACTOME FORMATION OF LATERAL PLATE MESODERM
c2.WP PANCREATIC CANCER SUBTYPES
c2.KORKOLA CHORIOCARCINOMA
c2.KEGG MEDICUS REFERENCE ORGANIZATION OF THE INNER KINETOCHORE c2.KEGG MEDICUS REFERENCE ORGANIZATION OF THE INNER KINETOCHORE c2.KEGG INTESTINAL IMMUNE NETWORK FOR IGA PRODUCTION c2.REACTOME METABOLISM OF INGESTED SEMET SEC MESEC INTO H2SE c2.REACTOME DEFECTIVE F9 ACTIVATION
c2.FARMER BREAST CANCER CLUSTER 1
c2.KEGG MEDICUS REFERENCE WNT SIGNALING MODULATION WNT INHIBITOR c2.KEGG PRIMARY IMMUNODEFICIENCY
c2.KEGG PRIMARY IMMUNODEFICIENCY
c2.WP UREA CYCLE AND RELATED DISEASES
c2.ZHOU PANCREATIC ENDOCRINE PROGENITOR
c2.MEISSNER NPC HCP WITH H3K27ME3
c2.NAKAMURA LUNG CANCER DIFFERENTIATION MARKERS
c2.FIGUEROA AML METHYLATION CLUSTER 2 DN c2.WP CATALYTIC CYCLE OF MAMMALIAN FLAVINCONTAINING MONOOXYGENASES FMOS
c2.MEISSNER BRAIN HCP WITH H3K27ME3
c2.SALVADOR MARTIN PEDIATRIC TBD ANTI TNF THERAPY NONRESPONDER PRE TREATMENT UP c2.WP GPCRS CLASS A RHODOPSINLIKE c2.BIOCARTA STEM PATHWAY c2.REACTOME UREA CYCLE c2.KEGG MEDICUS REFERENCE TESTOSTERONE BIOSYNTHESIS c2.RAY ALZHEIMERS DISEASE c2.WP GASTRIC ACID PRODUCTION c2.BIOCARTA EOSINOPHILS PATHWAY c2.SU KIDNEY c2.MARTINELLI IMMATURE NEUTROPHIL UP c2.REACTOME TANDEM PORE DOMAIN POTASSIUM CHANNELS
c2.REACTOME ERYTHROCYTES TAKE UP CARBON DIOXIDE AND RELEASE OXYGEN
c2.REACTOME DISEASES OF BASE EXCISION REPAIR c2.WP CODEINE AND MORPHINE METABOLISM c2.WEBER METHYLATED HCP IN FIBROBLAST DN c2.REACTOME HISTIDINE CATABOLISM c2.NIKOLSKY BREAST CANCER 19Q13.1 AMPLICON
c2.WP COMPLEMENTMEDIATED INFLAMMATION OF PULMONARY ALVEOLUS IN COVID19 HYPOTHETICAL PATHWAY
c2.REACTOME METAL SEQUESTRATION BY ANTIMICROBIAL PROTEINS **(0)** (0) c2.SMID BREAST CANCER RELAPSE IN LIVER UP c2.REACTOME EICOSANOID LIGAND BINDING RECEPTORS c2.KEGG MEDICUS REFERENCE WNT SIGNALING MODULATION WNT ACYLATION c2.REACTOME EICOSANOIDS c2.WP NONCLASSICAL ROLE OF VITAMIN D c2.REACTOME LIPID PARTICLE ORGANIZATION c2.REACTOME LIPID PARTICLE ORGANIZATION
c2.REACTOME ADRENOCEPTORS
c2.REACTOME CA2 ACTIVATED K CHANNELS
c2.REACTOME DEFECTIVE CSF2RB CAUSES SMDP5
c2.KEGG MATURITY ONSET DIABETES OF THE YOUNG
c2.WP ENTEROHEPATIC CIRCULATION OF BILE ACIDS
c2.REACTOME HIGHLY CALCIUM PERMEABLE NICOTINIC ACETYLCHOLINE RECEPTORS
c2.KEGG MEDICUS REFERENCE TNFSF10 RIPK1 3 SIGNALING PATHWAY
c2.WP MINERALOCORTICOID BIOSYNTHESIS
c2.REACTOME SEROTONIN AND MELATONIN BIOSYNTHESIS
c2.KEGG MEDICUS REFERENCE TURES CYCLE c2.KEGG MEDICUS REFERENCE UREA CYCLE c2.WP SULFATASE AND AROMATASE PATHWAY c2.WEBER METHYLATED HCP IN SPERM DN c2.REACTOME REGULATION OF GENE EXPRESSION IN ENDOCRINE COMMITTED NEUROG3 PROGENITOR CELLS c2.REACTOME PHASE 4 RESTING MEMBRANE POTENTIAL c2.KEGG MEDICUS REFERENCE ALTERNATIVE PATHWAY OF COMPLEMENT CASCADE C3B BREAKDOWN KEGG MEDICUS REFERENCE ALTERNATIVE PATHWAY OF COMPLEMENT CASCADE C3B BREAKDOWN

c2.SHIN B CELL LYMPHOMA CLUSTER 9

c2.KEGG MEDICUS ENV FACTOR PHIP TO DNA ADDUCTS

c2.WP ARYLAMINE METABOLISM

c2.REACTOME FGFR2 LIGAND BINDING AND ACTIVATION

c2.KEGG MEDICUS REFERENCE DNA DEGRADATION BY EXTRACELLULAR ENDOLYSOSOMAL DNASE

c2.WP COVID19 ADVERSE OUTCOME PATHWAY

c2.BIOCARTA ACETAMINOPHEN PATHWAY

c2.WP CONTROL OF IMMUNE TOLERANCE BY VASOACTIVE INTESTINAL PEPTIDE

c2.REACTOME DIGESTION AND ABSORPTION

c2.REACTOME LTC4 CYSLTR MEDIATED IL4 PRODUCTION

c2.VANDESLUIS NORMAL EMBRYOS UP

c2.BIOCARTA IL17 PATHWAY c2.BIOCARTA IL17 PATHWAY
c2.KEGG MEDICUS ENV FACTOR E2 TO NUCLEAR INITIATED ESTROGEN SIGNALING PATHWAY
c2.WP STEROID HORMONE PRECURSOR BIOSYNTHESIS c2.REACTOME PEPTIDE HORMONE BIOSYNTHESIS
c2.REACTOME HIGHLY CALCIUM PERMEABLE POSTSYNAPTIC NICOTINIC ACETYLCHOLINE RECEPTORS
c2.KEGG MEDICUS REFERENCE HEDGEHOG SIGNALING PATHWAY HH LIGAND SECRETION **00)** 00 c2.KEGG MEDICUS VARIANT MUTATION INACTIVATED ERBB4 TO NRG ERBB4 PI3K SIGNALING PATHWAY c2.FUNG IL2 TARGETS WITH STAT5 BINDING SITES T1 c2.KEGG MEDICUS REFERENCE NAD BIOSYNTHESIS c2.WP MALE STEROID HORMONES IN CARDIOMYOCYTE ENERGY METABOLISM c2.WP DRUG INDUCTION OF BILE ACID PATHWAY c2.BIOCARTA IL5 PATHWAY c2.REACTOME DIGESTION OF DIETARY LIPID c2.BIOCARTA DC PATHWAY c2.REACTOME SODIUM COUPLED SULPHATE DI AND TRI CARBOXYLATE TRANSPORTERS c2.WP MULTIPLE EPIPHYSEAL DYSPLASIA AND PSEUDOACHONDROPLASIA GENES c2.JI CARCINOGENESIS BY KRAS AND STK11 UP c2.WP SECRETION OF HYDROCHLORIC ACID IN PARIETAL CELLS c2.REACTOME HIGHLY SODIUM PERMEABLE POSTSYNAPTIC ACETYLCHOLINE NICOTINIC RECEPTORS c2.REACTOME PRESYNAPTIC NICOTINIC ACETYLCHOLINE RECEPTORS c2.MATZUK CUMULUS EXPANSION c2.REACTOME DEFECTIVE GALNT3 CAUSES HFTC
c2.REACTOME MUSCARINIC ACETYLCHOLINE RECEPTORS
c2.REACTOME ACETYLCHOLINE BINDING AND DOWNSTREAM EVENTS c2.MIKKELSEN IPS LCP WITH H3K27ME3 c2.WP MONOAMINE GPCRS c2.KOHN EMT EPITHELIAL c2.REACTOME THYROXINE BIOSYNTHESIS
c2.REACTOME OREXIN AND NEUROPEPTIDES FF AND QRFP BIND TO THEIR RESPECTIVE RECEPTORS
c2.REACTOME AMINE LIGAND BINDING RECEPTORS c2.KINNEY DNMT1 METHYLATION TARGETS c2.REACTOME DIGESTION c2.REACTOME METABOLISM OF AMINE DERIVED HORMONES c2.BIOCARTA TCAPOPTOSIS PATHWAY c2.REACTOME BETA DEFENSINS c2.REACTOME FATTY ACIDS c2.REACTOME MISCELLANEOUS SUBSTRATES
c2.KEGG MEDICUS REFERENCE COMMON PATHWAY OF COMPLEMENT CASCADE MAC FORMATION c2.REACTOME BIOSYNTHESIS OF MARESINS c2.UNTERMAN PROGRESSIVE VS STABLE IPF CD8T UP c2.REACTOME HORMONE LIGAND BINDING RECEPTORS c2.REACTOME MET RECEPTOR ACTIVATION c2.TESAR ALK TARGETS HUMAN ES 5D UP c2.REACTOME FICOLINS BIND TO REPETITIVE CARBOHYDRATE STRUCTURES ON THE TARGET CELL SURFACE c2.UNTERMAN IPF VS CTRL NK CELL UP c2.REACTOME SPERM MOTILITY AND TAXES c2.REACTOME GLUCOCORTICOID BIOSYNTHESIS c2.WP NAD BIOSYNTHESIS II FROM TRYPTOPHAN c2.REACTOME DEFENSINS c2.REACTOME SEROTONIN RECEPTORS c2.KEGG MEDICUS REFERENCE STEROID HORMONE BIOSYNTHESIS PROGESTERONE TO CORTISOL CORTISONE c2.WP COVID19 THROMBOSIS AND ANTICOAGULATION c2.WEBER METHYLATED ICP IN SPERM DN c2.UNTERMAN IPF VS CTRL CD8T DN c2.REACTOME VITAMINS c2.AUJLA IL22 AND IL17A SIGNALING c2.BIOCARTA BBCELL PATHWAY c2.WEBER METHYLATED LCP IN FIBROBLAST DN c2.WEBER METHYLATED LCP IN SPERM DN c2.WP EICOSANOID METABOLISM VIA CYTOCHROME P450 MONOOXYGENASES PATHWAY c2.TESAR ALK AND JAK TARGETS MOUSE ES D4 DN c2.REACTOME PREDNISONE ADME c2.REACTOME PROSTANOID LIGAND RECEPTORS c2.NAKAMURA BRONCHIAL AND BRONCHIOLAR EPITHELIA c2.REACTOME TACHYKININ RECEPTORS BIND TACHYKININS c2.BIOCARTA ASBCELL PATHWAY c2.WEBER METHYLATED ICP IN FIBROBLAST (0)00 c2.REACTOME LECTIN PATHWAY OF COMPLEMENT ACTIVATION c2.CURSONS NATURAL KILLER CELLS c2.WEBER METHYLATED LCP IN FIBROBLAST UP c2.WEBER METHYLATED LCP IN SPERM UP c2.WP GLP1 SECRETION FROM INTESTINE TO PORTAL VEIN c2.REACTOME MINERALOCORTICOID BIOSYNTHESIS c2.BRUNEAU SEPTATION ATRIAL c2.KEGG GRAFT VERSUS HOST DISEASE c2.REACTOME OPSINS c2.KEGG ALLOGRAFT REJECTION c2.KEGG MEDICUS REFERENCE LECTIN PATHWAY OF COAGULATION CASCADE PROTHROMBIN TO THROMBIN c2.REACTOME ALPHA DEFENSINS c2.REACTOME ABACAVIR TRANSMEMBRANE TRANSPORT c2.KEGG AUTOIMMUNE THYROID DISEASE c2.REACTOME DOPAMINE RECEPTORS c2.NIKOLSKY BREAST CANCER 7P15 AMPLICON c2.REACTOME CIPROFLOXACIN ADME c2.REACTOME ORGANIC ANION TRANSPORT c2.REACTOME CYP2E1 REACTIONS c2.KEGG MEDICUS REFERENCE LECTIN PATHWAY OF COAGULATION CASCADE FIBRINOGEN TO FIBRIN c2.REACTOME MELANIN BIOSYNTHESIS c2.REACTOME DIGESTION OF DIETARY CARBOHYDRATE
c2.WEBER METHYLATED ICP IN SPERM UP c2.REACTOME VASOPRESSIN LIKE RECEPTORS c2.REACTOME RELAXIN RECEPTORS
c2.REACTOME REGULATION OF GENE EXPRESSION IN EARLY PANCREATIC PRECURSOR CELLS c2.REACTOME BIOSYNTHESIS OF MARESIN LIKE SPMS c2.KEGG MEDICUS PATHOGEN SARS COV 2 S N TO LECTIN PATHWAY OF COAGULATION CASCADE c2.BERGER MBD2 TARGETS c3.RBMX TARGET GENES c3.PER1 TARGET GENES c3.AGTGCGT MIR521 c3.RPA2 TARGET GENES c3.RBM17 TARGET GENES c3.FOXF1 TARGET GENES c3.HSF4 TARGET GENES c3.PSMB5 TARGET GENES c3.MDM2 TARGET GENES c3.NR2E3 TARGET GENES c3.NEUROD2 TARGET GENES c3.TCGATGG MIR213 c3.AACGGTT MIR451 c3.TCCGTCC MIR184 c3.CARM1 TARGET GENES c3.ATCMNTCCGY UNKNOWN c3.GGTAACC MIR4095P c3.AGGAGTG MIR483 c3.NR4A2 TARGET GENES c3.SNAPC2 TARGET GENES c3.ADCYAP1 TARGET GENES c3.CGTCTTA MIR208 c3.ZMYND11 TARGET GENES c3.KCCGNSWTTT UNKNOWN c3.CRGAARNNNCGA UNKNOWN c3.GCCATNTTG YY1 Q6 c3.ACAWNRNSRCGG UNKNOWN c3.HJURP TARGET GENES c3.CGGAARNGGCNG UNKNOWN c3.MAP2K1 TARGET GENES c3.ACACTAC MIR1423P c3.SOX11 TARGET GENES c3.AACTGAC MIR223 c3.ZNF622 TARGET GENES c3.GACTGTT MIR212 MIR132 c3.TUBG1 TARGET GENES c3.MAPK3 TARGET GENES c3.CTR9 TARGET GENES c3.ZNF302 TARGET GENES c3.ACCATTT MIR522 c3.ATAGGAA MIR202 c3.ZNF19 TARGET GENES c3 TMTCGCGANR UNKNOWN c3.GGCAGTG MIR3243P c3.GGCNKCCATNK UNKNOWN C3 GTGTTGA MIR509 c3.GAGACTG MIR452 c3.ZZZ3 TARGET GENES c3.ARHGAP35 TARGET GENES c3.AGTCAGC MIR345 c3.GTATGAT MIR154 MIR487 c3.ATGAAGG MIR205 c3.GGCNNMSMYNTTG UNKNOWN c3.GTCGATC MIR3695P c3.ATAACCT MIR154 c3.CTTGTAT MIR381 c3.TCTGATC MIR383 c3.ACTGTAG MIR139 c3.TCTGATA MIR361 c3.YGCANTGCR UNKNOWN c3.EGFR TARGET GENES c3.NRF2 01 c3.TAGAACC MIR182 c3.ZNF445 TARGET GENES c3.CAGTGTT MIR141 MIR200A c3.SOX15 TARGET GENES c3.GTF2A2 TARGET GENES c3.MED16 TARGET GENES c3.ATGCAGT MIR217 c3.GTAAACC MIR2995F c3.YY1 02 c3.TAF9B TARGET GENES c3.GCGCCTT MIR525 MIR524 c3.CAGTATT MIR200B MIR200C MIR429 c3.TTTGCAG MIR518A2 c3.TGTATGA MIR4853P c3.GCGSCMNTTT UNKNOWN c3.GCACCTT MIR18A MIR18B c3.MTHFD1 TARGET GENES c3.MSX2 TARGET GENES c3.CTAGGAA MIR384 c3.GTTAAAG MIR302B c3.ACTGCAG MIR173P c3.CATTTCA MIR203 c3.RRCCGTTA UNKNOWN c3.ACATTCC MIR1 MIR206 c3.ATAAGCT MIR21 c3.CCAWNWWNNNGGC UNKNOWN c3.YY1 Q6 c3.TAATGTG MIR323 c3.TAGGTCA MIR192 MIR215 c3.CCCACAT MIR2993P c3.ATGTTAA MIR302C c3.CCATCCA MIR432 c3.PRMT5 TARGET GENES patho_cat_name c3.TTTTGAG MIR373 c3.GTGCCAT MIR183 c3.MED25 TARGET GENES Nerve sheath tumors c3.GTF2E2 TARGET GENES Spindle and epithelioid tumors c3.STN1 TARGET GENES c3.GTGACTT MIR224 c3.FOXR2 TARGET GENES Gliomas c3.TCTATGA MIR376A MIR376B c3.GATGKMRGCG UNKNOWN Glioneuronal tumors c3.TTGCCAA MIR182 c3.ELK1 02 c3.TAXCREB 01 c3.GCATTTG MIR105 c3.CTGTTAC MIR194 c3.AAGGGAT MIR188 c3.AHR 01 c3.ATXN7L3 TARGET GENES c3.CTTTGTA MIR524 c3.AGTCTTA MIR499 c3.TTTGTAG MIR520D c3.GCTTGAA MIR498 c3.FOXE1 TARGET GENES c3.GACAGGG MIR339 c3.AAGWWRNYGGC UNKNOWN c3.PGBD5 TARGET GENES c3.PPARA TARGET GENES c3.CCAGGTT MIR490 c3.NPM1 TARGET GENES c3.GTTTGTT MIR495 c3.TGAGATT MIR216 c3.CATGTAA MIR496 c3.SREBP1 01 c3.ACACTCC MIR122A c3.HTF 01 c3.GTTATAT MIR410 c3.ATGCACG MIR517B c3.EMX1 TARGET GENES c3.GGAANCGGAANY UNKNOWN c3.ATTCTTT MIR186 c3.LET 7I 3P c3.RUVBL1 TARGET GENES c3.SETD7 TARGET GENES c3.SRF 01 c3.GCAAAAA MIR129 c3.CTTTGCA MIR527 c3.TTGGGAG MIR150 c3.GTAGGCA MIR189 c3.AAAGGGA MIR204 MIR211 c3 GTACTGT MIR101 c3.ACAACTT MIR382 c3.ATGGYGGA UNKNOWN c3.ZNF746 TARGET GENES c3.AHRARNT 02 c3.ATGTTTC MIR494 c3.TAGCTTT MIR9 c3.ATGTACA MIR493 c3.CGCAAAA MIR450 c3.ZNF776 TARGET GENES c3.ZA UNIPROT Q9UM89 UNREVIEWED GENES TARGET GENES c3.GTAAGAT MIR200A c3.ATCTTGC MIR31 c3.GTGTCAA MIR514 c3.GGCACTT MIR519E c3.ATCATGA MIR433 c3.GCTGAGT MIR5125P c3.NRF1 Q6 c3.GGCNRNWCTTYS UNKNOWN c3.ETF Q6 c3.GGGATGC MIR3245P c3.ACACTGG MIR199A MIR199B c3.YBX1 TARGET GENES c3.ATGCTGG MIR338 c3.TGCTTTG MIR330 c3.TTGCACT MIR130A MIR301 MIR130B c3.SNRNP70 TARGET GENES c3.TAANNYSGCG UNKNOWN c3.TCCAGAT MIR5165P c3.RUVBL2 TARGET GENES c3.TFCP2 TARGET GENES c3.GGCCAGT MIR193A MIR193B c3.ACAGGGT MIR10A MIR10B c3.NFE2L1 TARGET GENES c3.PGM3 TARGET GENES c3.USF 01 c3.YRTCANNRCGC UNKNOWN c3.GCGNNANTTCC UNKNOWN c3.GTATTAT MIR3693P c3.CCGNMNNTNACG UNKNOWN c3.CAGCTTT MIR320 c3.TTTGCAC MIR19A MIR19B c3.YY1 01 c3.RCGCANGCGY NRF1 Q6 c3.ZNF236 TARGET GENES c3.ARNT 02 c3.AACATTC MIR4093P c3.CTACTGT MIR199A c3.GABP B c3.TGAATGT MIR181A MIR181B MIR181C MIR181D c3.ZNF585B TARGET GENES c3.AGCATTA MIR155 c3.MSX1 TARGET GENES c3.KAT2A TARGET GENES c3.MXD1 TARGET GENES c3.GGCAGAC MIR346 c3.KAT5 TARGET GENES c3.ACTTTAT MIR1425P c3.TGCGCANK UNKNOWN c3.CMYB 01 c3.DYRK1A TARGET GENES c3.CTATGCA MIR153 c3.AGGAAGC MIR5163P c3.CACGTTT MIR302A c3.SALL4 TARGET GENES c3.NEUROG3 TARGET GENES c3.AACTGGA MIR145 c3.KDM1B TARGET GENES c3.XBP1 01 c3.ACTGCCT MIR34B c3.GTCTTCC MIR7 c3.SVIL TARGET GENES c3.TGCACGA MIR517A MIR517C c3.CTACTAG MIR325 c3.TGTGTGA MIR377 c3.TCATCTC MIR143 c3.SRPK1 TARGET GENES c3.TOX4 TARGET GENES c3.CAGTCAC MIR134 c3.AGCGCAG MIR191 c3.HMGB2 TARGET GENES c3.TACAATC MIR508 c3.GATA2 01 c3.CETS1P54 01 c3.UBE2I TARGET GENES c3.CCTGAGT MIR510 c3.ZNF318 TARGET GENES c3.CREB3L2 TARGET GENES c3.AGGTGCA MIR500 c3.RAG1 TARGET GENES c3.HSF2 TARGET GENES c3.ZFP37 TARGET GENES c3.TTGGAGA MIR5155P MIR519E c3.TCCCCAC MIR491 c3.ATACTGT MIR144 c3.IRF5 TARGET GENES c3.GGCACAT MIR455 c3.TGCACTG MIR148A MIR152 MIR148B c3.CAATGCA MIR33 c3.WIZ TARGET GENES c3.AGCTCCT MIR28 c3.TACTTGA MIR26A MIR26B c3.ZNF711 TARGET GENES c3.CCACACA MIR147 c3.CIITA TARGET GENES c3.CHAF1A TARGET GENES c3.GTGGTGA MIR197 c3.ZSCAN18 TARGET GENES c3.ATF5 TARGET GENES c3.BANP TARGET GENES c3.GAGCCAG MIR149 c3.TGCCTTA MIR124A c3.ATM TARGET GENES c3.NR1I2 TARGET GENES c3.GGCAGCT MIR22 c3.THRAP3 TARGET GENES c3.NR5A1 TARGET GENES c3.CTCTAGA MIR526C MIR518F MIR526A c3.SMTTTTGT UNKNOWN c3.TCCCRNNRTGC UNKNOWN c3.TATCTGG MIR488 c3.TGCACTT MIR519C MIR519B MIR519A c3.AAGCCAT MIR135A MIR135B c3.AAAGACA MIR511 c3.GKCGCNNNNNNTGAYG UNKNOWN c3.EPC1 TARGET GENES c3.THRA TARGET GENES c3.GUCY1B1 TARGET GENES c3.RBM15 TARGET GENES c3.LAMB3 TARGET GENES c3.GCAAGGA MIR502 c3.SNIP1 TARGET GENES c3.GTGCAAA MIR507 c3.ACATATC MIR190 c3.ZNF532 TARGET GENES c3.DBP TARGET GENES c3.KLF7 TARGET GENES c3.UBN1 TARGET GENES c3.CTCCAAG MIR432 c3.E2F1 Q3 01 c3.SRF Q5 01 c3.GGGACCA MIR133A MIR133B c3.DLX6 TARGET GENES c3.GTGCAAT MIR25 MIR32 MIR92 MIR363 MIR367 c3.GTACAGG MIR486 c3.AUTS2 TARGET GENES c3.ZNF112 TARGET GENES c3.IRX3 TARGET GENES \mathbb{S} c3.ZFHX3 TARGET GENES c3.PPARGC1A TARGET GENES c3.ZNF586 TARGET GENES c3.LET 7B 3P c3.TEAD2 TARGET GENES c3.AAGCACT MIR520F c3.GCACTTT MIR175P MIR20A MIR106A MIR106B MIR20B MIR519D c3.LET 7C 3P c3.GAGCCTG MIR484 c3.NGFIC 01 c3.TGGTGCT MIR29A MIR29B MIR29C c3.NCOA4 TARGET GENES c3.FOXG1 TARGET GENES c3.YGTCCTTGR UNKNOWN c3.DNMT3A TARGET GENES c3.TATTATA MIR374 c3.ZNF740 TARGET GENES c3.ZMYM2 TARGET GENES c3.AAAGGAT MIR501 c3.GGATCCG MIR127 c3.NAB2 TARGET GENES c3.ZNF507 TARGET GENES c3.NPAT TARGET GENES c3.GTGTGAG MIR342 c3.ATTACAT MIR3803P c3.AGGCACT MIR5153P c3.CCTGTGA MIR513 c3.TTCNRGNNNNTTC HSF Q6 c3.AATGGAG MIR136 c3.CREB3L4 TARGET GENES c3.TFEB TARGET GENES c3.TCCATTKW UNKNOWN c3.TAXCREB 02 c3.CIC TARGET GENES c3.CHAF1B TARGET GENES c3.CCAWYNNGAAR UNKNOWN c3.GTGACGY E4F1 Q6 c3.TBX1 TARGET GENES c3.MAX 01 c3.ATGTCAC MIR489 c3.HOXA10 TARGET GENES c3.AATGTGA MIR23A MIR23B c3.ZNF626 TARGET GENES c3.CAGCAGG MIR370 c3.CREB 02 c3.GR 01 c3.AAGCAAT MIR137 c3.FEV TARGET GENES c3.GTGCCTT MIR506 c3.ACCAAAG MIR9 c3.CACTGTG MIR128A MIR128B c3.TBX3 TARGET GENES c3.GTGCCAA MIR96 c3.GAGCTGG MIR337 c3.AAGCACA MIR218 c3.TCTCTCC MIR185 c3.E4F1 Q6 c3.USF Q6 01 c3.SETD1A TARGET GENES c3.TGTTTAC MIR30A5P MIR30C MIR30D MIR30B MIR30E5P c3.AP1FJ Q2 c3.ZNF282 TARGET GENES c3.CDH4 TARGET GENES c3.HOXC11 TARGET GENES c3.USF 02 c3.SFMBT1 TARGET GENES c3.ACCAATC MIR509 c3.NKX2 2 TARGET GENES c3.MGGAAGTG GABP B c3.BARHL1 TARGET GENES c3.ZFP36L1 TARGET GENES c3.F2F1 Q4 c3.AAACCAC MIR140 c3.GGGCATT MIR365 c3.GGTGTGT MIR329 c3.TGANNYRGCA TCF11MAFG 01 c3.CACCAGC MIR138 c3.ATGTAGC MIR221 MIR222 c3.CGTSACG PAX3 B c3.MYCMAX 03 c3.IRX2 TARGET GENES c3.HOXA7 TARGET GENES c3.STAT1 02 c3.PAX8 TARGET GENES c3.LET 7F 2 3P c3.TGCAAAC MIR452 c3.CACBINDINGPROTEIN Q6 c3.CACTTTG MIR520G MIR520H c3.ASH1L TARGET GENES c3.NR1D1 TARGET GENES c3.RBL1 TARGET GENES c3.ATF6 TARGET GENES c3.OVOL3 TARGET GENES c3.RAX2 TARGET GENES c3.ATF3 Q6 c3.CCTGCTG MIR214 c3.PAX4 03 c3.WRNIP1 TARGET GENES c3.ZNF175 TARGET GENES c3.CTCTGGA MIR520A MIR525 c3.DMRT1 TARGET GENES c3.TASOR TARGET GENES c3.ZSCAN5DP TARGET GENES c3.GGGNRMNNYCAT UNKNOWN c3.HLF 01 c3.SP1 Q6 c3.NKX2 3 TARGET GENES c3.MZF1 TARGET GENES c3.ZNF513 TARGET GENES c3.BRCA2 TARGET GENES c3.ERCA2 TARGET GENES c3.E4BP4 01 c3.HOXD1 TARGET GENES c3.TCF11MAFG 01 c3.PTPRA TARGET GENES c3.AP2ALPHA 01 c3.PRKDC TARGET GENES c3.SOX3 TARGET GENES c3.CDX1 TARGET GENES c3.ZNF785 TARGET GENES c3.SRF Q4 c3.ZNF837 TARGET GENES c3.AHRR TARGET GENES c3.AEBP2 TARGET GENES c3.CREBP1 01 c3.ZNF354B TARGET GENES c3.LET 7G 3P c3.BARX1 TARGET GENES c3.CREB Q2 c3.FOXO1 02 c3.GAANYNYGACNY UNKNOWN c3.ZNF250 TARGET GENES c3.FOXQ1 TARGET GENES c3.CAGCACT MIR5123F c3.AP1 Q6 01 c3.SUPT16H TARGET GENES c3.AGGGCAG MIR18A c3.ATF 01 c3.ARID5B TARGET GENES c3.NUFIP1 TARGET GENES c3.FOXC1 TARGET GENES c3.ATF4 Q2 c3.SKIL TARGET GENES c3.EVI1 06 c3.PAX4 01 c3.COBLL1 TARGET GENES c3.ARNT2 TARGET GENES c3.TCTGGAC MIR198 c3.YYCATTCAWW UNKNOWN c3.HES4 TARGET GENES c3.ARNT 01 c3.HOXA1 TARGET GENES c3.ERR1 Q2 c3.ZNF524 TARGET GENES c3.ZNF589 TARGET GENES c3.E2F Q6 01 c3.RFX7 TARGET GENES c3.SP1 Q4 01 c3.EGR3 01 c3.ZSCAN23 TARGET GENES c3.TGCTGCT MIR15A MIR16 MIR15B MIR195 MIR424 MIR497 c3.AGCACTT MIR93 MIR302A MIR302B MIR302C MIR302D MIR372 MIR373 MIR520E MIR520A MIR526B MIR520B MIR520C MIR520D c3.ZNF843 TARGET GENES c3.BRF1 TARGET GENES c3.HOX13 01 c3.ZNF610 TARGET GENES c3.PITX1 TARGET GENES c3.MCRS1 TARGET GENES c3.INSM2 TARGET GENES c3.EWSR1 TARGET GENES c3.CCCNNNNNNAAGWT UNKNOWN c3.AAGTCCA MIR422B MIR422A c3.ZNF669 TARGET GENES c3.TET1 TARGET GENES c3.MAFG TARGET GENES c3.TCF11 01 c3.DACH1 TARGET GENES c3.MAZ Q6 c3.TCOF1 TARGET GENES c3.ZNF350 TARGET GENES c3.SF1 Q6 c3.CAGCCTC MIR4855P c3.CAAGGAT MIR362 c3.ZNF561 TARGET GENES c3.BPTF TARGET GENES c3.ZNF138 TARGET GENES c3.AGTTCTC MIR146A MIR146B c3.SOX9 B1 c3.KMCATNNWGGA UNKNOWN c3.ZNF581 TARGET GENES c3.MCAATNNNNNGCG UNKNOWN c3.ELF5 TARGET GENES c3.HOXB6 TARGET GENES c3.ATATGCA MIR448 c3.CREB Q3 c3.ZNF184 TARGET GENES c3.FOXD2 TARGET GENES c3.ZNF407 TARGET GENES c3.ACCGAGC MIR423 c3.NCOA2 TARGET GENES c3.HOXB4 TARGET GENES c3.YGCGYRCGC UNKNOWN c3.MAZR 01 c3.FOXO4 TARGET GENES c3.ZNF416 TARGET GENES c3.BACH1 01 c3.POU2AF1 TARGET GENES c3.TCANNTGAY SREBP1 01 c3.ATACCTC MIR202 c3.ZNF391 TARGET GENES c3.TEL2 Q6 c3.NCOA6 TARGET GENES c3.BCL6B TARGET GENES c3.ATGCTGC MIR103 MIR107 c3.AP2 Q6 01 c3.NRF2 Q4 c3.PTF1BETA Q6 c3.HOXC6 TARGET GENES c3.RPA1 TARGET GENES c3.LXR Q3 c3.CREB Q4 c3.HIF1 Q5 c3.ZNF592 TARGET GENES c3.ZBTB12 TARGET GENES c3.CTGAGCC MIR24 c3.CREBP1 Q2 c3.ZSCAN5B TARGET GENES c3.HDGF TARGET GENES c3.SRSF9 TARGET GENES c3.SUMO1 TARGET GENES c3.PR 02 c3.TERT TARGET GENES c3.BARX2 TARGET GENES c3.STAT1 03 c3.NKX2 5 TARGET GENES c3.ELK1 01 c3.ALKBH3 TARGET GENES c3.GLI1 TARGET GENES c3.USF2 Q6 c3.LCORL TARGET GENES c3.RREB1 01 c3.GCTNWTTGK UNKNOWN c3.ZNF766 TARGET GENES c3.ACTGTGA MIR27A MIR27B c3.ZNF85 TARGET GENES c3.F10 TARGET GENES c3.GGAMTNNNNNTCCY UNKNOWN c3.TACGGGT MIR99A MIR100 MIR99B c3.DLX4 TARGET GENES c3.MZF1 01 c3.ZNF830 TARGET GENES c3.GTCTACC MIR379 c3.SUPT20H TARGET GENES c3.MEF2D TARGET GENES c3.HHEX TARGET GENES c3.PAX7 TARGET GENES c3.ATOH8 TARGET GENES c3.NR0B1 TARGET GENES c3.ZIC2 01 c3.GACAATC MIR219 c3.CACTGCC MIR34A MIR34C MIR449 c3.GGCKCATGS UNKNOWN c3.PAF1 TARGET GENES c3.EGR Q6 c3.NFY 01 c3.AP1 Q4 c3.MYBL1 TARGET GENES c3.PAX3 B c3.SETBP1 TARGET GENES c3.ZNF169 TARGET GENES c3.AP1 Q6 c3.CCCAGAG MIR326 c3.AGTCTAG MIR151 c3.NKX22 01 c3.ZBTB1 TARGET GENES c3.SP1 Q2 01 c3.TGACCTTG SF1 Q6 c3.CEBPZ TARGET GENES c3.ZNF257 TARGET GENES c3.RORA TARGET GENES c3.1 HX9 TARGET GENES c3.ZBTB44 TARGET GENES c3.ZSCAN31 TARGET GENES c3.ZNF30 TARGET GENES c3.DIDO1 TARGET GENES -0.3 0.0 -0.3-0.30.3 0.3 -0.30.0 0.3 0.0 z_score