

ETABOLIC\_PROCESS,GOBP\_TRNA\_METABOLIC\_PROCESS

GSE18791\_CTRL\_VS\_NEWCASTLE\_VIRUS\_DC\_10H\_UP,GSE18791\_CTRL\_VS\_NEWCASTLE\_VIRUS\_DC\_10H\_UP  
GOBP\_RNA\_MODIFICATION,GOBP\_RNA\_MODIFICATION  
GSE46606\_UNSTIM\_VS\_CD40L\_IL2\_IL5\_DAY3\_STIMULATED\_BCELL\_DN,GSE46606\_UNSTIM\_VS\_CD40L\_IL2\_IL5\_DAY3\_STIMULATED\_BCELL\_DN  
GOBP\_TRNA\_PROCESSING,GOBP\_TRNA\_PROCESSING  
GOMF\_S\_ADENOSYLMETHIONINE\_DEPENDENT\_METHYLTRANSFERASE\_ACTIVITY,GOMF\_S\_ADENOSYLMETHIONINE\_DEPENDENT\_METHYLTRANSFERASE\_ACTIVITY  
GSE16451\_CTRL\_VS\_WEST\_EQUINE\_ENC\_VIRUS\_IMMATURE\_NEURON\_CELL\_LINE\_DN,GSE16451\_CTRL\_VS\_WEST\_EQUINE\_ENC\_VIRUS\_IMMATURE\_NEURON\_CELL\_LINE\_DN  
GOMF\_NUCLEASE\_ACTIVITY,GOMF\_NUCLEASE\_ACTIVITY  
MYC\_UP\_V1\_UP,MYC\_UP\_V1\_UP  
GOBP\_TRNA\_MODIFICATION,GOBP\_TRNA\_MODIFICATION  
GSE17974\_2H\_VS\_72H\_UNTREATED\_IN\_VITRO\_CD4\_TCELL\_UP,GSE17974\_2H\_VS\_72H\_UNTREATED\_IN\_VITRO\_CD4\_TCELL\_UP  
HP\_DEEPLY\_SET\_EYE,HP\_DEEPLY\_SET\_EYE  
GOBP\_RNA\_METHYLATION,GOBP\_RNA\_METHYLATION  
GOBP\_RNA\_PHOSPHODIESTER\_BOND\_HYDROLYSIS,GOBP\_RNA\_PHOSPHODIESTER\_BOND\_HYDROLYSIS  
HP\_MYOCLONIC\_SEIZURE,HP\_MYOCLONIC\_SEIZURE  
HP\_POSTNATAL\_MICROCEPHALY,HP\_POSTNATAL\_MICROCEPHALY  
HP\_EPILEPTIC\_SPASM,HP\_EPILEPTIC\_SPASM  
HP\_GENERALIZED\_ONSET\_MOTOR\_SEIZURE,HP\_GENERALIZED\_ONSET\_MOTOR\_SEIZURE  
HP\_INFANTILE\_MUSCULAR\_HYPOTONIA,HP\_INFANTILE\_MUSCULAR\_HYPOTONIA  
MIR877\_3P,MIR877\_3P  
GOMF\_ENDONUCLEASE\_ACTIVITY,GOMF\_ENDONUCLEASE\_ACTIVITY  
HP\_ABNORMALITY\_OF\_THE\_PERIVENTRICULAR\_WHITE\_MATTER,HP\_ABNORMALITY\_OF\_THE\_PERIVENTRICULAR\_WHITE\_MATTER  
GOMF\_RNA\_METHYLTRANSFERASE\_ACTIVITY,GOMF\_RNA\_METHYLTRANSFERASE\_ACTIVITY  
HP\_HYPERSARRHYTHMIA,HP\_HYPERSARRHYTHMIA  
GOMF\_TRNA\_BINDING,GOMF\_TRNA\_BINDING  
ELVIDGE\_HIF1A\_TARGETS\_UP,ELVIDGE\_HIF1A\_TARGETS\_UP  
HP\_DISTAL\_SENSORY\_IMPAIRMENT,HP\_DISTAL\_SENSORY\_IMPAIRMENT  
HP\_HETEROTROPIA,HP\_HETEROTROPIA  
GOBP\_MITOCHONDRIAL\_RNA\_METABOLIC\_PROCESS,GOBP\_MITOCHONDRIAL\_RNA\_METABOLIC\_PROCESS  
GSE34217\_MIR17\_92\_OVEREXPRESS\_VS\_WT\_ACT\_CD8\_TCELL\_DN,GSE34217\_MIR17\_92\_OVEREXPRESS\_VS\_WT\_ACT\_CD8\_TCELL\_DN  
GOMF\_RIBONUCLEASE\_ACTIVITY,GOMF\_RIBONUCLEASE\_ACTIVITY  
HP\_CHRONIC\_KIDNEY\_DISEASE,HP\_CHRONIC\_KIDNEY\_DISEASE  
HP\_HYPOPLASIA\_OF\_THE\_BRAINSTEM,HP\_HYPOPLASIA\_OF\_THE\_BRAINSTEM  
HP\_BILATERAL\_SINGLE\_TRANSVERSE\_PALMAR\_CREASES,HP\_BILATERAL\_SINGLE\_TRANSVERSE\_PALMAR\_CREASES  
HP\_NARROW\_FOREHEAD,HP\_NARROW\_FOREHEAD  
GSE5589\_WT\_VS\_IL10\_KO\_LPS\_AND\_IL6\_STIM\_MACROPHAGE\_45MIN\_DN,GSE5589\_WT\_VS\_IL10\_KO\_LPS\_AND\_IL6\_STIM\_MACROPHAGE\_45MIN\_DN  
REACTOME\_TRNA\_MODIFICATION\_IN\_THE\_NUCLEUS\_AND\_CYTOSOL,REACTOME\_TRNA\_MODIFICATION\_IN\_THE\_NUCLEUS\_AND\_CYTOSOL  
HP\_ABNORMAL\_SUBARACHNOID\_SPACE\_MORPHOLOGY,HP\_ABNORMAL\_SUBARACHNOID\_SPACE\_MORPHOLOGY  
HP\_ESODEVIATION,HP\_ESODEVIATION  
HP\_INFANTILE\_SPASMS,HP\_INFANTILE\_SPASMS  
HP\_CHOREOATHETOSIS,HP\_CHOREOATHETOSIS  
GOBP\_RRNA\_MODIFICATION,GOBP\_RRNA\_MODIFICATION  
HP\_GASTROSTOMY\_TUBE\_FEEDING\_IN\_INFANCY,HP\_GASTROSTOMY\_TUBE\_FEEDING\_IN\_INFANCY  
HP\_CNS\_HYPOMYELINATION,HP\_CNS\_HYPOMYELINATION  
GOMF\_ENDONUCLEASE\_ACTIVITY\_ACTIVE\_WITH\_EITHER\_RIBO\_OR\_DEOXYRIBONUCLEIC\_ACIDS\_AND\_PRODUCING\_5\_PHOSPHOMONOESTERS,GOMF\_ENDONUCLEASE\_ACTIVITY\_ACTIVE\_WITH\_EITHER\_RIBO\_OR\_DEOXYRIBONUCLEIC\_ACIDS\_AND\_PRODUCING\_5\_PHOSPHOMONOESTERS  
HP\_WIDENED\_SUBARACHNOID\_SPACE,HP\_WIDENED\_SUBARACHNOID\_SPACE  
HP\_GENERALIZED\_AMYTROPHY,HP\_GENERALIZED\_AMYTROPHY  
ELVIDGE\_HIF1A\_AND\_HIF2A\_TARGETS\_UP,ELVIDGE\_HIF1A\_AND\_HIF2A\_TARGETS\_UP  
HP\_ABNORMALITY\_OF\_OCULAR\_SMOOTH\_PURSUIT,HP\_ABNORMALITY\_OF\_OCULAR\_SMOOTH\_PURSUIT  
GOBP\_TRNA\_METHYLATION,GOBP\_TRNA\_METHYLATION  
GOBP\_NCRNA\_CATABOLIC\_PROCESS,GOBP\_NCRNA\_CATABOLIC\_PROCESS  
GOBP\_REGULATION\_OF\_MITOCHONDRIAL\_GENE\_EXPRESSION,GOBP\_REGULATION\_OF\_MITOCHONDRIAL\_GENE\_EXPRESSION  
HP\_ORAL\_PHARYNGEAL\_DYSPHAGIA,HP\_ORAL\_PHARYNGEAL\_DYSPHAGIA  
GSE5099\_MONOCYTE\_VS\_ALTERNATIVE\_M2\_MACROPHAGE\_UP,GSE5099\_MONOCYTE\_VS\_ALTERNATIVE\_M2\_MACROPHAGE\_UP  
HP\_NEPHROPATHY,HP\_NEPHROPATHY  
GOBP\_RNA\_POLYADENYLATION,GOBP\_RNA\_POLYADENYLATION  
HP\_SIMPLIFIED\_GYRAL\_PATTERN,HP\_SIMPLIFIED\_GYRAL\_PATTERN  
HP\_MULTIFOCAL\_EPILEPTIFORM\_DISCHARGES,HP\_MULTIFOCAL\_EPILEPTIFORM\_DISCHARGES  
HP\_NEPHROTIC\_SYNDROME,HP\_NEPHROTIC\_SYNDROME  
GSE36476\_YOUNG\_VS\_OLD\_DONOR\_MEMORY\_CD4\_TCELL\_16h\_TSST\_ACT\_UP,GSE36476\_YOUNG\_VS\_OLD\_DONOR\_MEMORY\_CD4\_TCELL\_16h\_TSST\_ACT\_UP  
ZNf708\_TARGET\_GENES,ZNF708\_TARGET\_GENES  
HP\_LARGE\_BASAL\_GANGLIA,HP\_LARGE\_BASAL\_GANGLIA  
HP\_ADDUCTED\_THUMB,HP\_ADDUCTED\_THUMB  
GOMF\_TRANSFERASE\_ACTIVITY\_TRANSFERRING\_ALKYL\_OR\_ARYL\_OTHER\_THAN\_METHYL\_GROUPS,GOMF\_TRANSFERASE\_ACTIVITY\_TRANSFERRING\_ALKYL\_OR\_ARYL\_OTHER\_THAN\_METHYL\_GROUPS  
HP\_ABNORMAL\_CIRCULATING\_ALBUMIN\_CONCENTRATION,HP\_ABNORMAL\_CIRCULATING\_ALBUMIN\_CONCENTRATION  
HP\_NASOGASTRIC\_TUBE\_FEEDING,HP\_NASOGASTRIC\_TUBE\_FEEDING  
HP\_INFECTION\_RELATED\_SEIZURE,HP\_INFECTION\_RELATED\_SEIZURE  
HOXC11\_TARGET\_GENES,HOXC11\_TARGET\_GENES  
MIR125A\_3P,MIR125A\_3P  
GOBP\_RRNA\_METHYLATION,GOBP\_RRNA\_METHYLATION  
HP\_DYSPLASTIC\_CORPUS\_CALLOSUM,HP\_DYSPLASTIC\_CORPUS\_CALLOSUM  
GOBP\_TRNA\_WOBBLE\_BASE\_MODIFICATION,GOBP\_TRNA\_WOBBLE\_BASE\_MODIFICATION  
GOBP\_NCRNA\_3\_END\_PROCESSING,GOBP\_NCRNA\_3\_END\_PROCESSING  
GOMF\_RNA\_POLYMERASE\_CORE\_ENZYME\_BINDING,GOMF\_RNA\_POLYMERASE\_CORE\_ENZYME\_BINDING  
HP\_PROFOUND\_GLOBAL\_DEVELOPMENTAL\_DELAY,HP\_PROFOUND\_GLOBAL\_DEVELOPMENTAL\_DELAY  
HP\_BILATERAL\_SENSORINEURAL\_HEARING\_IMPAIRMENT,HP\_BILATERAL\_SENSORINEURAL\_HEARING\_IMPAIRMENT  
GOMF\_CARBOXYLIC\_ESTER\_HYDROLASE\_ACTIVITY,GOMF\_CARBOXYLIC\_ESTER\_HYDROLASE\_ACTIVITY  
GOMF\_INTRAMOLECULAR\_TRANSFERASE\_ACTIVITY,GOMF\_INTRAMOLECULAR\_TRANSFERASE\_ACTIVITY  
GOBP\_POSITIVE\_REGULATION\_OF\_MITOCHONDRIAL\_TRANSLATION,GOBP\_POSITIVE\_REGULATION\_OF\_MITOCHONDRIAL\_TRANSLATION  
HP\_HORIZONTAL\_NYSTAGMUS,HP\_HORIZONTAL\_NYSTAGMUS  
HP\_BILATERAL\_TONIC\_CLONIC\_SEIZURE\_WITH\_GENERALIZED\_ONSET,HP\_BILATERAL\_TONIC\_CLONIC\_SEIZURE\_WITH\_GENERALIZED\_ONSET  
GOMF\_LIGASE\_ACTIVITY\_FORMING\_CARBON\_NITROGEN\_BONDS,GOMF\_LIGASE\_ACTIVITY\_FORMING\_CARBON\_NITROGEN\_BONDS  
HP\_IMPAIRED\_SMOOTH\_PURSUIT,HP\_IMPAIRED\_SMOOTH\_PURSUIT  
GOMF\_O\_METHYLTRANSFERASE\_ACTIVITY,GOMF\_O\_METHYLTRANSFERASE\_ACTIVITY  
GOMF\_HYDROLASE\_ACTIVITY\_ACTING\_ON\_CARBON\_NITROGEN\_BUT\_NOT\_PEPTIDE\_BONDS,GOMF\_HYDROLASE\_ACTIVITY\_ACTING\_ON\_CARBON\_NITROGEN\_BUT\_NOT\_PEPTIDE\_BONDS  
HP\_PERIVENTRICULAR\_LEUKOMALACIA,HP\_PERIVENTRICULAR\_LEUKOMALACIA  
GOMF\_TRANSFERASE\_ACTIVITY\_TRANSFERRING\_PENTOSYL\_GROUPS,GOMF\_TRANSFERASE\_ACTIVITY\_TRANSFERRING\_PENTOSYL\_GROUPS  
HP\_APLASIA\_HYPOPLASIA\_OF\_THE\_CEREBRAL\_WHITE\_MATTER,HP\_APLASIA\_HYPOPLASIA\_OF\_THE\_CEREBRAL\_WHITE\_MATTER  
KRIGE\_AMINO\_ACID\_DEPRIVATION,KRIGE\_AMINO\_ACID\_DEPRIVATION  
GOMF\_ADENYLYLTRANSFERASE\_ACTIVITY,GOMF\_ADENYLYLTRANSFERASE\_ACTIVITY  
MIR6854\_5P,MIR6854\_5P  
MODULE\_322,MODULE\_322  
HP\_NONIMMUNE\_HYDROPS\_FETALIS,HP\_NONIMMUNE\_HYDROPS\_FETALIS  
GOBP\_RNA\_SPLICING\_VIA\_ENDONUCLEOLYTIC\_CLEAVAGE\_AND\_LIGATION,GOBP\_RNA\_SPLICING\_VIA\_ENDONUCLEOLYTIC\_CLEAVAGE\_AND\_LIGATION  
HP\_DIFFUSE\_CEREBRAL\_ATROPHY,HP\_DIFFUSE\_CEREBRAL\_ATROPHY  
HP\_FINE\_HAIR,HP\_FINE\_HAIR  
HP\_HAMMERTOE,HP\_HAMMERTOE  
GOCC\_MRNA\_CLEAVAGE\_AND\_POLYADENYLATION\_SPECIFICITY\_FACTOR\_COMPLEX,GOCC\_MRNA\_CLEAVAGE\_AND\_POLYADENYLATION\_SPECIFICITY\_FACTOR\_COMPLEX  
HP\_STEPPAGE\_GAIT,HP\_STEPPAGE\_GAIT  
SIX4\_TARGET\_GENES,SIX4\_TARGET\_GENES  
HP\_ACUTE\_HEPATIC\_FAILURE,HP\_ACUTE\_HEPATIC\_FAILURE  
KEGG\_HISTIDINE\_METABOLISM,KEGG\_HISTIDINE\_METABOLISM  
GOMF\_TRANSFERASE\_ACTIVITY\_TRANSFERRING\_SULFUR\_CONTAINING\_GROUPS,GOMF\_TRANSFERASE\_ACTIVITY\_TRANSFERRING\_SULFUR\_CONTAINING\_GROUPS  
MIR3667\_3P,MIR3667\_3P  
REACTOME\_RRNA\_PROCESSING\_IN\_THE\_MITOCHONDRION,REACTOME\_RRNA\_PROCESSING\_IN\_THE\_MITOCHONDRION  
TERAMOTO\_OPN\_TARGETS\_CLUSTER\_7,TERAMOTO\_OPN\_TARGETS\_CLUSTER\_7  
HP\_RESPIRATORY\_FAILURE\_REQUIRING\_ASSISTED\_VENTILATION,HP\_RESPIRATORY\_FAILURE\_REQUIRING\_ASSISTED\_VENTILATION  
GOBP\_MITOCHONDRIAL\_TRNA\_PROCESSING,GOBP\_MITOCHONDRIAL\_TRNA\_PROCESSING  
REACTOME\_RNA\_POLYMERASE\_III\_TRANSCRIPTION\_TERMINATION,REACTOME\_RNA\_POLYMERASE\_III\_TRANSCRIPTION\_TERMINATION  
HP\_KERATOCONJUNCTIVITIS,HP\_KERATOCONJUNCTIVITIS  
KEGG\_SELENOAMINO\_ACID\_METABOLISM,KEGG\_SELENOAMINO\_ACID\_METABOLISM  
HP\_SPASTIC\_ATAXIA,HP\_SPASTIC\_ATAXIA  
GOBP\_PPTIDYL\_HISTIDINE\_MODIFICATION,GOBP\_PPTIDYL\_HISTIDINE\_MODIFICATION  
MIR1178\_5P,MIR1178\_5P  
HP\_ABNORMAL\_BLOOD\_TRANSITION\_ELEMENT\_CATION\_CONCENTRATION,HP\_ABNORMAL\_BLOOD\_TRANSITION\_ELEMENT\_CATION\_CONCENTRATION  
HP\_PSEUDOBULBAR\_SIGNS,HP\_PSEUDOBULBAR\_SIGNS  
BIOCARTA\_CPFS\_PATHWAY,BIOCARTA\_CPFS\_PATHWAY  
GOBP\_TRNA\_THREONYLCARBAMOYLADENOSINE\_METABOLIC\_PROCESS,GOBP\_TRNA\_THREONYLCARBAMOYLADENOSINE\_METABOLIC\_PROCESS  
HP\_BRITTLE\_HAIR,HP\_BRITTLE\_HAIR  
HP\_UPPER\_LIMB\_HYPERTONIA,HP\_UPPER\_LIMB\_HYPERTONIA  
REACTOME\_TRNA\_MODIFICATION\_IN\_THE\_MITOCHONDRION,REACTOME\_TRNA\_MODIFICATION\_IN\_THE\_MITOCHONDRION  
HP\_IMPAIRED\_OROPHARYNGEAL\_SWALLOW\_RESPONSE,HP\_IMPAIRED\_OROPHARYNGEAL\_SWALLOW\_RESPONSE  
CHESLER\_BRAIN\_HIGHEST\_GENETIC\_VARIANCE,CHESLER\_BRAIN\_HIGHEST\_GENETIC\_VARIANCE  
REACTOME\_SYNTHESIS\_OF\_DIPHTHAMIDE\_EEF2,REACTOME\_SYNTHESIS\_OF\_DIPHTHAMIDE\_EEF2  
HP\_DIFFUSE\_CEREBELLAR\_ATROPHY,HP\_DIFFUSE\_CEREBELLAR\_ATROPHY  
GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_BLACK\_DN,GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_BLACK\_DN  
HP\_MICROVESICULAR\_HEPATIC\_STEATOSIS,HP\_MICROVESICULAR\_HEPATIC\_STEATOSIS  
HP\_SIMPLE\_FEBRILE\_SEIZURE,HP\_SIMPLE\_FEBRILE\_SEIZURE  
GOMF\_SULFURTRANSFERASE\_ACTIVITY,GOMF\_SULFURTRANSFERASE\_ACTIVITY  
TOMIDA\_METASTASIS\_DN,TOMIDA\_METASTASIS\_DN  
DESCARTES\_MAIN\_FETAL\_HEMATOPOIETIC\_STEM\_CELLS,DESCARTES\_MAIN\_FETAL\_HEMATOPOIETIC\_STEM\_CELLS  
HP\_DECREASED\_MEAN\_CORPUSCULAR\_VOLUME,HP\_DECREASED\_MEAN\_CORPUSCULAR\_VOLUME  
GOBP\_SMALL\_RNA\_LOADING\_ONTO\_RISC,GOBP\_SMALL\_RNA\_LOADING\_ONTO\_RISC  
GOMF\_TRNA\_CYTOSINE\_METHYLTRANSFERASE\_ACTIVITY,GOMF\_TRNA\_CYTOSINE\_METHYLTRANSFERASE\_ACTIVITY  
HP\_SHIELD\_CHEST,HP\_SHIELD\_CHEST  
HP\_PAROXYSMAL\_DYSTONIA,HP\_PAROXYSMAL\_DYSTONIA  
GOMF\_HYDROLASE\_ACTIVITY\_ACTING\_ON\_CARBON\_NITROGEN\_BUT\_NOT\_PEPTIDE\_BONDS\_IN\_CYCLIC\_AMIDINES,GOMF\_HYDROLASE\_ACTIVITY\_ACTING\_ON\_CARBON\_NITROGEN\_BUT\_NOT\_PEPTIDE\_BONDS\_IN\_CYCLIC\_AMIDINES  
GOMF\_TRNA\_GUANINE\_METHYLTRANSFERASE\_ACTIVITY,GOMF\_TRNA\_GUANINE\_METHYLTRANSFERASE\_ACTIVITY  
HP\_GENERALIZED\_LIMB\_MUSCLE\_ATROPHY,HP\_GENERALIZED\_LIMB\_MUSCLE\_ATROPHY  
GOMF\_MRNA\_METHYLTRANSFERASE\_ACTIVITY,GOMF\_MRNA\_METHYLTRANSFERASE\_ACTIVITY  
HP\_RECURRENT\_GASTROENTERITIS,HP\_RECURRENT\_GASTROENTERITIS  
HP\_ACHILLES\_TENDON\_CONTRACTURE,HP\_ACHILLES\_TENDON\_CONTRACTURE  
GOBP\_CYTOPLASMIC\_TRANSLATIONAL\_ELONGATION,GOBP\_CYTOPLASMIC\_TRANSLATIONAL\_ELONGATION  
HP\_WIDENED\_CEREBELLAR\_SUBARACHNOID\_SPACE,HP\_WIDENED\_CEREBELLAR\_SUBARACHNOID\_SPACE  
GOBP\_PRIMARY\_MRNA\_PROCESSING,GOBP\_PRIMARY\_MRNA\_PROCESSING  
GOMF\_ADENOSINE\_DEAMINASE\_ACTIVITY,GOMF\_ADENOSINE\_DEAMINASE\_ACTIVITY  
HP\_HYPOPLASIA\_OF\_THE\_VENTRAL\_PONS,HP\_HYPOPLASIA\_OF\_THE\_VENTRAL\_PONS  
GOBP\_PROSTHETIC\_GROUP\_METABOLIC\_PROCESS,GOBP\_PROSTHETIC\_GROUP\_METABOLIC\_PROCESS  
REACTOME\_MOLYBDENUM\_COFACTOR\_BIOSYNTHESIS,REACTOME\_MOLYBDENUM\_COFACTOR\_BIOSYNTHESIS  
GOMF\_DEAMINASE\_ACTIVITY,GOMF\_DEAMINASE\_ACTIVITY  
HP\_LEG\_MUSCLE\_STIFFNESS,HP\_LEG\_MUSCLE\_STIFFNESS  
ZNf837\_TARGET\_GENES,ZNF837\_TARGET\_GENES  
HP\_MULTIPLE\_ROWS\_OF\_EYELASHES,HP\_MULTIPLE\_ROWS\_OF\_EYELASHES  
GOMF\_OXO\_ACID\_LYASE\_ACTIVITY,GOMF\_OXO\_ACID\_LYASE\_ACTIVITY  
GOMF\_RNA\_2\_O\_METHYLTRANSFERASE\_ACTIVITY,GOMF\_RNA\_2\_O\_METHYLTRANSFERASE\_ACTIVITY  
GOBP\_ENZYME\_DIRECTED\_RRNA\_PSEUDOURIDINE\_SYNTHESIS,GOBP\_ENZYME\_DIRECTED\_RRNA\_PSEUDOURIDINE\_SYNTHESIS  
ZNf418\_TARGET\_GENES,ZNF418\_TARGET\_GENES  
HARALAMBIEVA\_PBMC\_FLUARIX\_AGE\_50\_74YO\_CORR\_WITH\_28D\_MEM\_B\_CELL\_RESPONSE\_AT\_28DY\_LEUK\_MIGR\_MAPK\_ACT\_CYTOK\_SIG\_DIAB\_OF\_THE\_YNG\_NEGATIVE,HARALAMBIEVA\_PBMC\_FLUARIX\_AGE\_50\_74YO\_CORR\_WITH\_28D\_MEM\_B\_CELL\_RESPONSE\_AT\_28DY\_LEUK\_MIGR\_MAPK\_ACT\_CYTOK\_SIG\_DIAB\_OF\_THE\_YNG\_NEGATIVE  
GOCC\_ELONGATOR\_HOJENZYME\_COMPLEX,GOCC\_ELONGATOR\_HOJENZYME\_COMPLEX  
GOMF\_MRNA\_GUANINE\_METHYLTRANSFERASE\_ACTIVITY,GOMF\_MRNA\_GUANINE\_METHYLTRANSFERASE\_ACTIVITY  
HP\_DECREASED\_SERUM\_INSULIN\_LIKE\_GROWTH\_FACTOR\_1,HP\_DECREASED\_SERUM\_INSULIN\_LIKE\_GROWTH\_FACTOR\_1  
GOMF\_ENDONUCLEASE\_ACTIVITY\_ACTIVE\_WITH\_EITHER\_RIBO\_OR\_DEOXYRIBONUCLEIC\_ACIDS\_AND\_PRODUCING\_3\_PHOSPHOMONOESTERS,GOMF\_ENDONUCLEASE\_ACTIVITY\_ACTIVE\_WITH\_EITHER\_RIBO\_OR\_DEOXYRIBONUCLEIC\_ACIDS\_AND\_PRODUCING\_3\_PHOSPHOMONOESTERS  
HP\_ERYTHROCYTE\_INCLUSION\_BODIES,HP\_ERYTHROCYTE\_INCLUSION\_BODIES  
HP\_DECREASED\_CORNEAL\_REFLEX,HP\_DECREASED\_CORNEAL\_REFLEX  
HP\_ANISOCYTOSIS,HP\_ANISOCYTOSIS  
HP\_AXONAL\_REGENERATION,HP\_AXONAL\_REGENERATION  
HP\_IMPAIRED\_DISTAL\_TACTILE\_SENSATION,HP\_IMPAIRED\_DISTAL\_TACTILE\_SENSATION  
HP\_DECREASED\_SERUM\_IRON,HP\_DECREASED\_SERUM\_IRON  
HP\_METABOLIC\_ALKALOSIS,HP\_METABOLIC\_ALKALOSIS  
GOBP\_MRNA\_PSEUDOURIDINE\_SYNTHESIS,GOBP\_MRNA\_PSEUDOURIDINE\_SYNTHESIS  
HP\_INFANTILE\_ENCEPHALOPATHY,HP\_INFANTILE\_ENCEPHALOPATHY  
MIR1281,MIR1281  
HP\_ALKALOSIS,HP\_ALKALOSIS  
GOBP\_REGULATION\_OF\_GLUCAGON\_SECRETION,GOBP\_REGULATION\_OF\_GLUCAGON\_SECRETION  
ZNf292\_TARGET\_GENES,ZNF292\_TARGET\_GENES  
REACTOME\_SYNTHESIS\_OF\_WYBUTOSINE\_AT\_G37\_OF\_TRNA\_PHE,REACTOME\_SYNTHESIS\_OF\_WYBUTOSINE\_AT\_G37\_OF\_TRNA\_PHE  
HP\_TIP\_TOE\_GAIT,HP\_TIP\_TOE\_GAIT  
HP\_SMALL\_ANTERIOR\_FONTANELLE,HP\_SMALL\_ANTERIOR\_FONTANELLE  
GOBP\_IMPAIRED\_DISTAL\_PROPRIOCEPTION,HP\_IMPAIRED\_DISTAL\_PROPRIOCEPTION  
GOBP\_REGULATION\_OF\_MESODERM\_DEVELOPMENT,GOBP\_REGULATION\_OF\_MESODERM\_DEVELOPMENT  
HP\_MEDULLARY\_NEPHROCALCINOSIS,HP\_MEDULLARY\_NEPHROCALCINOSIS  
MIR554,MIR554  
GOBP\_REGULATION\_OF\_ATPASE\_COUPLED\_CALCIIUM\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY,GOBP\_REGULATION\_OF\_ATPASE\_COUPLED\_CALCIIUM\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY  
HP\_ERYTHROID\_HYPERPLASIA,HP\_ERYTHROID\_HYPERPLASIA  
HP\_DECREASED\_NUMBER\_OF\_LARGE\_PERIPHERAL\_MYELINATED\_NERVE\_FIBERS,HP\_DECREASED\_NUMBER\_OF\_LARGE\_PERIPHERAL\_MYELINATED\_NERVE\_FIBERS  
GAVIN\_IL2\_RESPONSIVE\_FOXP3\_TARGETS\_DN,GAVIN\_IL2\_RESPONSIVE\_FOXP3\_TARGETS\_DN  
MIR4681,MIR4681  
GOBP\_OXIDATIVE\_RNA\_DEMETHYLATION,GOBP\_OXIDATIVE\_RNA\_DEMETHYLATION  
GOMF\_OXIDATIVE\_RNA\_DEMETHYLASE\_ACTIVITY,GOMF\_OXIDATIVE\_RNA\_DEMETHYLASE\_ACTIVITY  
GOMF\_ENDORIBONUCLEASE\_ACTIVITY\_PRODUCING\_3\_PHOSPHOMONOESTERS,GOMF\_ENDORIBONUCLEASE\_ACTIVITY\_PRODUCING\_3\_PHOSPHOMONOESTERS  
GOMF\_ARGININE\_BINDING,GOMF\_ARGININE\_BINDING  
GOBP\_ADENOSINE\_TO\_INOSINE\_EDITING,GOBP\_ADENOSINE\_TO\_INOSINE\_EDITING  
GOMF\_TRNA\_SPECIFIC\_ADENOSINE\_DEAMINASE\_ACTIVITY,GOMF\_TRNA\_SPECIFIC\_ADENOSINE\_DEAMINASE\_ACTIVITY  
HP\_RIDGED\_NAIL,HP\_RIDGED\_NAIL  
WP\_DDX1\_AS\_A\_REGULATORY\_COMPONENT\_OF\_THE\_DROSHA\_MICROPROCESSOR,WP\_DDX1\_AS\_A\_REGULATORY\_COMPONENT\_OF\_THE\_DROSHA\_MICROPROCESSOR  
HP\_ABNORMALITY\_OF\_TASTE\_SENSATION,HP\_ABNORMALITY\_OF\_TASTE\_SENSATION  
GOBP\_OXIDATIVE\_DNA\_DEMETHYLATION,GOBP\_OXIDATIVE\_DNA\_DEMETHYLATION  
HP\_OVERBITE,HP\_OVERBITE  
MIR598\_3P,MIR598\_3P  
chr5q34,chr5q34  
MEBARK1\_HCC\_PROGENITOR\_WNT\_DN\_CTNNB1\_INDEPENDENT,MEBARK1\_HCC\_PROGENITOR\_WNT\_DN\_CTNNB1\_INDEPENDENT  
chr4q24,chr4q24