

TANT\_NOD2\_TRANSDUCED\_HEK293T\_CELL\_DN, GSE22611\_UNSTIM\_VS\_2H\_MDP\_STIM\_MUTANT\_NOD2\_TRANSDUCED\_HEK293T\_CELL\_DN

GOBP\_GOLGI\_ORGANIZATION, GOBP\_GOLGI\_ORGANIZATION  
GSE14769\_20MIN\_VS\_360MIN\_LPS\_BMDM\_DN, GSE14769\_20MIN\_VS\_360MIN\_LPS\_BMDM\_DN  
ACACTGG\_MIR199A\_MIR199B, ACACTGG\_MIR199A\_MIR199B  
GSE11924\_TH1\_VS\_TH2\_CD4\_TCELL\_UP, GSE11924\_TH1\_VS\_TH2\_CD4\_TCELL\_UP  
GSE26928\_NAIVE\_VS\_CXCR5\_POS\_CD4\_TCELL\_DN, GSE26928\_NAIVE\_VS\_CXCR5\_POS\_CD4\_TCELL\_DN  
GSE3039\_CD4\_TCELL\_VS\_B1\_BCELL\_UP, GSE3039\_CD4\_TCELL\_VS\_B1\_BCELL\_UP  
GOBP\_REGULATION\_OF\_TOR\_SIGNALING, GOBP\_REGULATION\_OF\_TOR\_SIGNALING  
MIR1245B\_3P, MIR1245B\_3P  
GSE22589\_HEALTHY\_VS\_SIV\_INFECTED\_DC\_DN, GSE22589\_HEALTHY\_VS\_SIV\_INFECTED\_DC\_DN  
MIR623, MIR623  
GSE2585\_THYMIC\_MACROPHAGE\_VS\_MTEC\_DN, GSE2585\_THYMIC\_MACROPHAGE\_VS\_MTEC\_DN  
MIR2467\_3P, MIR2467\_3P  
ZHANG\_TARGETS\_OF\_EWSR1\_FLI1\_FUSION, ZHANG\_TARGETS\_OF\_EWSR1\_FLI1\_FUSION  
MIR10397\_5P, MIR10397\_5P  
KMCATNNWGGA\_UNKNOWN, KMCATNNWGGA\_UNKNOWN  
KEGG\_WNT\_SIGNALING\_PATHWAY, KEGG\_WNT\_SIGNALING\_PATHWAY  
MIR4288, MIR4288  
MIR6775\_3P, MIR6775\_3P  
HP\_PATELLAR\_DISLOCATION, HP\_PATELLAR\_DISLOCATION  
MIR499B\_5P, MIR499B\_5P  
CHUNG\_BLISTER\_CYTOTOXICITY\_DN, CHUNG\_BLISTER\_CYTOTOXICITY\_DN  
HP\_GASTROINTESTINAL\_INFLAMMATION, HP\_GASTROINTESTINAL\_INFLAMMATION  
MIR4656, MIR4656  
MIR550B\_3P, MIR550B\_3P  
MIR4749\_3P, MIR4749\_3P  
FONTAINE\_FOLLICULAR\_THYROID\_ADENOMA\_UP, FONTAINE\_FOLLICULAR\_THYROID\_ADENOMA\_UP  
CAATGCA\_MIR33, CAATGCA\_MIR33  
GOBP\_CARDIAC\_CHAMBER\_DEVELOPMENT, GOBP\_CARDIAC\_CHAMBER\_DEVELOPMENT  
MIR5699\_3P, MIR5699\_3P  
MIR4421, MIR4421  
MIR6734\_5P, MIR6734\_5P  
GAANYNYGACNY\_UNKNOWN, GAANYNYGACNY\_UNKNOWN  
EVI1\_05, EVI1\_05  
GOBP\_LEARNING, GOBP\_LEARNING  
GOBP\_CARDIAC\_CHAMBER\_MORPHOGENESIS, GOBP\_CARDIAC\_CHAMBER\_MORPHOGENESIS  
HP\_ANTINUCLEAR\_ANTIBODY\_POSITIVITY, HP\_ANTINUCLEAR\_ANTIBODY\_POSITIVITY  
GOBP\_BONE\_CELL\_DEVELOPMENT, GOBP\_BONE\_CELL\_DEVELOPMENT  
HP\_INSOMNIA, HP\_INSOMNIA  
GOBP\_ORGANELLE\_INHERITANCE, GOBP\_ORGANELLE\_INHERITANCE  
GOBP\_RETINA\_DEVELOPMENT\_IN\_CAMERA\_TYPE\_EYE, GOBP\_RETINA\_DEVELOPMENT\_IN\_CAMERA\_TYPE\_EYE  
MIR4513, MIR4513  
MIR4525, MIR4525  
MIR5010\_5P, MIR5010\_5P  
FONTAINE\_PAPILLARY\_THYROID\_CARCINOMA\_DN, FONTAINE\_PAPILLARY\_THYROID\_CARCINOMA\_DN  
GOBP\_NEURON\_DEATH\_IN\_RESPONSE\_TO\_OXIDATIVE\_STRESS, GOBP\_NEURON\_DEATH\_IN\_RESPONSE\_TO\_OXIDATIVE\_STRESS  
GOBP\_REGULATION\_OF\_HORMONE\_BIOSYNTHETIC\_PROCESS, GOBP\_REGULATION\_OF\_HORMONE\_BIOSYNTHETIC\_PROCESS  
MODULE\_366, MODULE\_366  
SCHAEFFER\_PROSTATE\_DEVELOPMENT\_AND\_CANCER\_BOX1\_UP, SCHAEFFER\_PROSTATE\_DEVELOPMENT\_AND\_CANCER\_BOX1\_UP  
HP\_FRONTAL\_LOBE\_DEMENTIA, HP\_FRONTAL\_LOBE\_DEMENTIA  
GOBP\_ESTABLISHMENT\_OF\_LYMPHOCYTE\_POLARITY, GOBP\_ESTABLISHMENT\_OF\_LYMPHOCYTE\_POLARITY  
GOBP\_REGULATION\_OF\_OXIDATIVE\_STRESS\_INDUCED\_NEURON\_DEATH, GOBP\_REGULATION\_OF\_OXIDATIVE\_STRESS\_INDUCED\_NEURON\_DEATH  
GOBP\_HEART TRABECULA MORPHOGENESIS, GOBP\_HEART TRABECULA MORPHOGENESIS  
GOBP\_POSITIVE\_REGULATION\_OF\_ATP\_METABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_ATP\_METABOLIC\_PROCESS  
HP\_DENTAL\_ENAMEL\_PITS, HP\_DENTAL\_ENAMEL\_PITS  
GOBP\_POSITIVE\_REGULATION\_OF\_TRANSCRIPTION\_BY\_RNA\_POLYMERASE\_III, GOBP\_POSITIVE\_REGULATION\_OF\_TRANSCRIPTION\_BY\_RNA\_POLYMERASE\_III  
GOBP\_HEART TRABECULA FORMATION, GOBP\_HEART TRABECULA FORMATION  
GOBP TRABECULA FORMATION, GOBP TRABECULA FORMATION  
HP\_CHILDHOOD\_ONSET\_SENSORINEURAL\_HEARING\_IMPAIRMENT, HP\_CHILDHOOD\_ONSET\_SENSORINEURAL\_HEARING\_IMPAIRMENT  
GOBP\_REGULATION\_OF\_THYMOCYTE\_APOPTOTIC\_PROCESS, GOBP\_REGULATION\_OF\_THYMOCYTE\_APOPTOTIC\_PROCESS  
GOBP\_EMBRYONIC\_RETINA\_MORPHOGENESIS\_IN\_CAMERA\_TYPE\_EYE, GOBP\_EMBRYONIC\_RETINA\_MORPHOGENESIS\_IN\_CAMERA\_TYPE\_EYE  
GOBP\_INOSITOL\_PHOSPHATE\_BIOSYNTHETIC\_PROCESS, GOBP\_INOSITOL\_PHOSPHATE\_BIOSYNTHETIC\_PROCESS  
REACTOME\_CASPASE\_ACTIVATION\_VIA\_DEATH\_RECEPTORS\_IN\_THE\_PRESENCE\_OF\_LIGAND, REACTOME\_CASPASE\_ACTIVATION\_VIA\_DEATH\_RECEPTORS\_IN\_THE\_PRESENCE\_OF\_LIGAND  
GOBP\_ARF\_PROTEIN\_SIGNAL\_TRANSDUCTION, GOBP\_ARF\_PROTEIN\_SIGNAL\_TRANSDUCTION  
GOBP\_POSITIVE\_REGULATION\_OF\_HORMONE\_METABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_HORMONE\_METABOLIC\_PROCESS  
REACTOME\_DEFECTIVE\_RIPK1\_MEDIATED\_REGULATED\_NECROSIS, REACTOME\_DEFECTIVE\_RIPK1\_MEDIATED\_REGULATED\_NECROSIS  
GOCC\_RIPOPOTOSOME, GOCC\_RIPOPOTOSOME  
GOBP\_NEURAL\_FOLD\_FORMATION, GOBP\_NEURAL\_FOLD\_FORMATION  
GOBP\_POSITIVE\_REGULATION\_OF\_HORMONE\_BIOSYNTHETIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_HORMONE\_BIOSYNTHETIC\_PROCESS  
GOBP\_REGULATION\_OF\_MESENCHYMAL\_STEM\_CELL\_DIFFERENTIATION, GOBP\_REGULATION\_OF\_MESENCHYMAL\_STEM\_CELL\_DIFFERENTIATION  
GOBP\_MESENCHYMAL\_CELL\_APOPTOTIC\_PROCESS, GOBP\_MESENCHYMAL\_CELL\_APOPTOTIC\_PROCESS  
REACTOME\_PHYSIOLOGICAL\_FACTORS, REACTOME\_PHYSIOLOGICAL\_FACTORS  
GOBP\_POSITIVE\_REGULATION\_OF\_GLYCOLYTIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_GLYCOLYTIC\_PROCESS  
GOBP\_NEGATIVE\_REGULATION\_OF\_MESENCHYMAL\_CELL\_APOPTOTIC\_PROCESS, GOBP\_NEGATIVE\_REGULATION\_OF\_MESENCHYMAL\_CELL\_APOPTOTIC\_PROCESS  
GOBP\_IMMUNE\_RESPONSE\_INHIBITING\_SIGNAL\_TRANSDUCTION, GOBP\_IMMUNE\_RESPONSE\_INHIBITING\_SIGNAL\_TRANSDUCTION  
GOBP\_POSITIVE\_REGULATION\_OF\_NUCLEOTIDE\_METABOLIC\_PROCESS, GOBP\_POSITIVE\_REGULATION\_OF\_NUCLEOTIDE\_METABOLIC\_PROCESS  
FIGUEROA\_AML\_METHYLATION\_CLUSTER\_7\_DN, FIGUEROA\_AML\_METHYLATION\_CLUSTER\_7\_DN  
GOBP\_MORPHOGENESIS\_OF\_AN\_EPITHELIAL\_FOLD, GOBP\_MORPHOGENESIS\_OF\_AN\_EPITHELIAL\_FOLD  
GOBP\_REGULATION\_OF\_ALDOSTERONE\_METABOLIC\_PROCESS, GOBP\_REGULATION\_OF\_ALDOSTERONE\_METABOLIC\_PROCESS  
GOBP\_NEGATIVE\_REGULATION\_OF\_INTERLEUKIN\_5\_PRODUCTION, GOBP\_NEGATIVE\_REGULATION\_OF\_INTERLEUKIN\_5\_PRODUCTION  
GOBP\_ALDOSTERONE\_METABOLIC\_PROCESS, GOBP\_ALDOSTERONE\_METABOLIC\_PROCESS  
GOBP\_MINERALOCORTICOID\_METABOLIC\_PROCESS, GOBP\_MINERALOCORTICOID\_METABOLIC\_PROCESS