

MORF\_TPR, MORF\_TPR

MORF\_BECN1, MORF\_BECN1  
GSE360\_CTRL\_VS\_T\_GONDII\_MAC\_UP, GSE360\_CTRL\_VS\_T\_GONDII\_MAC\_UP  
MORF\_CSNK1D, MORF\_CSNK1D  
GO\_SWI\_SNF\_SUPERFAMILY\_TYPE\_COMPLEX, GO\_SWI\_SNF\_SUPERFAMILY\_TYPE\_COMPLEX  
MORF\_SS18, MORF\_SS18  
GSE17721\_POLYIC\_VS\_PAM3CSK4\_12H\_BMDC\_UP, GSE17721\_POLYIC\_VS\_PAM3CSK4\_12H\_BMDC\_UP  
MORF\_PSMF1, MORF\_PSMF1  
GO\_4\_IRON\_4\_SULFUR\_CLUSTER\_BINDING, GO\_4\_IRON\_4\_SULFUR\_CLUSTER\_BINDING  
GSE22033\_WT\_VS\_PPARG\_KO\_MEF\_DN, GSE22033\_WT\_VS\_PPARG\_KO\_MEF\_DN  
MORF\_ERCC2, MORF\_ERCC2  
MORF\_ARAF1, MORF\_ARAF1  
GO\_IRE1\_MEDIATED\_UNFOLDED\_PROTEIN\_RESPONSE, GO\_IRE1\_MEDIATED\_UNFOLDED\_PROTEIN\_RESPONSE  
GSE22611\_NOD2\_VS\_MUTANT\_NOD2\_TRANSDUCEd\_HEK293T\_CELL\_UP, GSE22611\_NOD2\_VS\_MUTANT\_NOD2\_TRANSDUCEd\_HEK293T\_CELL\_UP  
GO\_PROTEIN\_MONOUBIQUITINATION, GO\_PROTEIN\_MONOUBIQUITINATION  
GO\_NEGATIVE\_REGULATION\_OF\_MAPK\_CASCADE, GO\_NEGATIVE\_REGULATION\_OF\_MAPK\_CASCADE  
KEGG\_INSULIN\_SIGNALING\_PATHWAY, KEGG\_INSULIN\_SIGNALING\_PATHWAY  
GO\_DSRNA\_FRAGMENTATION, GO\_DSRNA\_FRAGMENTATION  
AAAGACA\_MIR511, AAAGACA\_MIR511  
POMEROY\_MEDULLOBLASTOMA\_DESMOPLASIC\_VS\_CLASSIC\_UP, POMEROY\_MEDULLOBLASTOMA\_DESMOPLASIC\_VS\_CLASSIC\_UP  
MORF\_RUNX1, MORF\_RUNX1  
CHOW\_RASSF1\_TARGETS\_UP, CHOW\_RASSF1\_TARGETS\_UP  
MORF\_PRKACA, MORF\_PRKACA  
GO\_REGULATION\_OF\_PHOSPHOPROTEIN\_PHOSPHATASE\_ACTIVITY, GO\_REGULATION\_OF\_PHOSPHOPROTEIN\_PHOSPHATASE\_ACTIVITY  
CCANNAGRKGGC\_UNKNOWN, CCANNAGRKGGC\_UNKNOWN  
GSE29949\_CD8\_POS\_DC\_SPLEEN\_VS\_MONOCYTE\_BONE\_MARROW\_UP, GSE29949\_CD8\_POS\_DC\_SPLEEN\_VS\_MONOCYTE\_BONE\_MARROW\_UP  
GO\_RESPONSE\_TO\_DSRNA, GO\_RESPONSE\_TO\_DSRNA  
GO\_REGULATION\_OF\_PROTEIN\_PHOSPHATASE\_TYPE\_2A\_ACTIVITY, GO\_REGULATION\_OF\_PROTEIN\_PHOSPHATASE\_TYPE\_2A\_ACTIVITY  
GO\_NEGATIVE\_REGULATION\_OF\_BINDING, GO\_NEGATIVE\_REGULATION\_OF\_BINDING  
ST\_G\_ALPHA\_I\_PATHWAY, ST\_G\_ALPHA\_I\_PATHWAY  
YANAGIHARA\_ESX1\_TARGETS, YANAGIHARA\_ESX1\_TARGETS  
GO\_NEGATIVE\_REGULATION\_OF\_SMALL\_GTPASE\_MEDIATED\_SIGNAL\_TRANSDUCTION, GO\_NEGATIVE\_REGULATION\_OF\_SMALL\_GTPASE\_MEDIATED\_SIGNAL\_TRANSDUCTION  
GO\_POLY\_A\_SPECIFIC\_RIBONUCLEASE\_ACTIVITY, GO\_POLY\_A\_SPECIFIC\_RIBONUCLEASE\_ACTIVITY  
GO\_SIGNAL\_RELEASE, GO\_SIGNAL\_RELEASE  
MORF\_NF1, MORF\_NF1  
GO\_HORMONE\_TRANSPORT, GO\_HORMONE\_TRANSPORT  
GO\_FORELIMB\_MORPHOGENESIS, GO\_FORELIMB\_MORPHOGENESIS  
PID\_ERBB2\_ERBB3\_PATHWAY, PID\_ERBB2\_ERBB3\_PATHWAY  
GTCNYYATGR\_UNKNOWN, GTCNYYATGR\_UNKNOWN  
GO\_NEGATIVE\_REGULATION\_OF\_PROTEIN\_BINDING, GO\_NEGATIVE\_REGULATION\_OF\_PROTEIN\_BINDING  
MEDINA\_SMARCA4\_TARGETS, MEDINA\_SMARCA4\_TARGETS  
GO\_MESENCHYME\_MORPHOGENESIS, GO\_MESENCHYME\_MORPHOGENESIS  
REACTOME\_RAF\_MAP\_KINASE\_CASCADE, REACTOME\_RAF\_MAP\_KINASE\_CASCADE  
GO\_REGULATION\_OF\_GLYCOGEN\_METABOLIC\_PROCESS, GO\_REGULATION\_OF\_GLYCOGEN\_METABOLIC\_PROCESS  
REACTOME\_ANTIGEN\_PRESENTATION\_FOLDING\_ASSEMBLY\_AND\_PEPTIDE\_LOADING\_OF\_CLASS\_I\_MHC, REACTOME\_ANTIGEN\_PRESENTATION\_FOLDING\_ASSEMBLY\_AND\_PEPTIDE\_LOADING\_OF\_CLASS\_I\_MHC  
GSE2770\_IL12\_VS\_TGFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_6H\_UP, GSE2770\_IL12\_VS\_TGFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_6H\_UP