

7\_POS\_VS\_NEG\_NK\_CELL\_UP, GSE23695\_CD57\_POS\_VS\_NEG\_NK\_CELL\_UP

GSE22935\_UNSTIM\_VS\_12H\_MBOVIS\_BCG\_STIM\_MYD88\_KO\_MACROPHAGE\_DN, GSE22935\_UNSTIM\_VS\_12H\_MBOVIS\_BCG\_STIM\_MYD88\_KO\_MACROPHAGE\_DN  
GSE17721\_PAM3CSK4\_VS\_CPG\_1H\_BMDC\_UP, GSE17721\_PAM3CSK4\_VS\_CPG\_1H\_BMDC\_UP  
GSE3039\_NKT\_CELL\_VS\_B2\_BCELL\_UP, GSE3039\_NKT\_CELL\_VS\_B2\_BCELL\_UP  
HP\_HIGHLY\_ARCHED\_EYEBROW, HP\_HIGHLY\_ARCHED\_EYEBROW  
MIR6852\_5P, MIR6852\_5P  
HP\_HEPATIC\_FIBROSIS, HP\_HEPATIC\_FIBROSIS  
HP\_ABNORMALITY\_OF\_THE\_HALLUX, HP\_ABNORMALITY\_OF\_THE\_HALLUX  
HP\_HAND\_POLYDACTYLY, HP\_HAND\_POLYDACTYLY  
GSE13306\_TREG\_VS\_TCONV\_LAMINA\_PROPRIA\_DN, GSE13306\_TREG\_VS\_TCONV\_LAMINA\_PROPRIA\_DN  
GSE17721\_LPS\_VS\_PAM3CSK4\_2H\_BMDC\_DN, GSE17721\_LPS\_VS\_PAM3CSK4\_2H\_BMDC\_DN  
HP\_RECURRENT\_FRACTURES, HP\_RECURRENT\_FRACTURES  
GSE18281\_PERIMEDULLARY\_CORTICAL\_REGION\_VS\_WHOLE\_MEDULLA\_THYMUS\_UP, GSE18281\_PERIMEDULLARY\_CORTICAL\_REGION\_VS\_WHOLE\_MEDULLA\_THYMUS\_UP  
GOMF\_REPRESSING\_TRANSCRIPTION\_FACTOR\_BINDING, GOMF\_REPRESSING\_TRANSCRIPTION\_FACTOR\_BINDING  
MIR4709\_3P, MIR4709\_3P  
HP\_ECTRODACTYLY, HP\_ECTRODACTYLY  
MIR15B\_3P, MIR15B\_3P  
MIR3923, MIR3923  
HP\_AGANGLIONIC\_MEGACOLON, HP\_AGANGLIONIC\_MEGACOLON  
GOBP\_AORTA\_DEVELOPMENT, GOBP\_AORTA\_DEVELOPMENT  
GOMF\_RNA\_POLYMERASE\_II\_REPRESSING\_TRANSCRIPTION\_FACTOR\_BINDING, GOMF\_RNA\_POLYMERASE\_II\_REPRESSING\_TRANSCRIPTION\_FACTOR\_BINDING  
HP\_JOINT\_SUBLUXATION, HP\_JOINT\_SUBLUXATION  
HP\_ABSENT\_TOE, HP\_ABSENT\_TOE  
HP\_ENCEPHALOCELE, HP\_ENCEPHALOCELE  
HP\_11\_PAIRS\_OF\_RIBS, HP\_11\_PAIRS\_OF\_RIBS  
CCANNAGRKGGC\_UNKNOWN, CCANNAGRKGGC\_UNKNOWN  
HP\_MICROSCOPIC\_HEMATURIA, HP\_MICROSCOPIC\_HEMATURIA  
HP\_CONGENITAL\_HEPATIC\_FIBROSIS, HP\_CONGENITAL\_HEPATIC\_FIBROSIS  
GENTLES\_LEUKEMIC\_STEM\_CELL\_UP, GENTLES\_LEUKEMIC\_STEM\_CELL\_UP  
HP\_MENINGOCELE, HP\_MENINGOCELE  
PEDERSEN\_METASTASIS\_BY\_ERBB2\_ISOFORM\_3, PEDERSEN\_METASTASIS\_BY\_ERBB2\_ISOFORM\_3  
HP\_ABNORMALITY\_OF\_SECONDARY\_SEXUAL\_HAIR, HP\_ABNORMALITY\_OF\_SECONDARY\_SEXUAL\_HAIR  
HP\_EPIPHYSEAL\_DYSPLASIA, HP\_EPIPHYSEAL\_DYSPLASIA  
TONKS\_TARGETS\_OF\_RUNX1\_RUNX1T1\_FUSION\_SUSTAINED\_IN\_MONOCYTE\_UP, TONKS\_TARGETS\_OF\_RUNX1\_RUNX1T1\_FUSION\_SUSTAINED\_IN\_MONOCYTE\_UP  
HP\_DEVIATION\_OF\_THE\_2ND\_FINGER, HP\_DEVIATION\_OF\_THE\_2ND\_FINGER  
HP\_CALVARIAL\_SKULL\_DEFECT, HP\_CALVARIAL\_SKULL\_DEFECT  
HP\_ABNORMAL\_2ND\_FINGER\_MORPHOLOGY, HP\_ABNORMAL\_2ND\_FINGER\_MORPHOLOGY  
GOBP\_SUBSTRATE\_DEPENDENT\_CELL\_MIGRATION, GOBP\_SUBSTRATE\_DEPENDENT\_CELL\_MIGRATION  
MIR6787\_5P, MIR6787\_5P