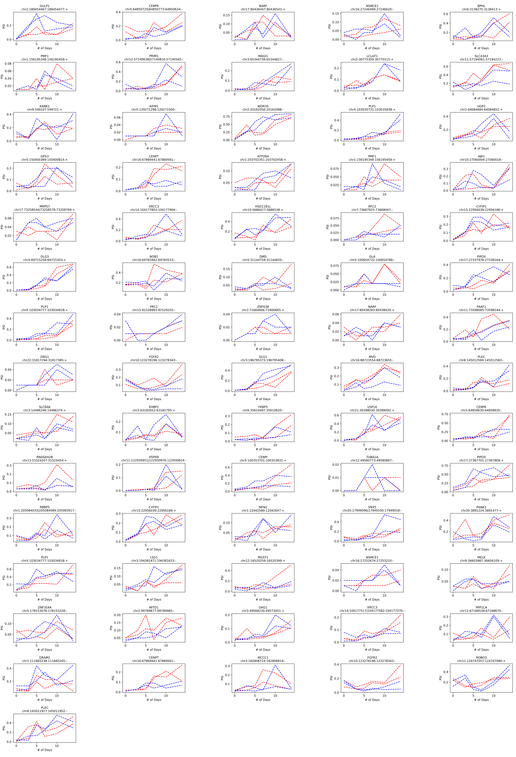
GLUT1
chr2:190956867:190956877 +
CDKN
chr5:94903720:94903737:94905824 -
NANP
chr17:80420947:80420970 +
NGWC1
chr16:2154490:2154492:- -
EPN1
chr6:3138270:3138313 +
PRR
chr1:156620548:156620559 +
PRKR
chr12:371464:371465:371470:371470 -
MAGI1
chr3:65917:65919:65927 -
LCLAT1
chr2:3037253:3037255:3037255 +
SLC43A1
chr17:1370000:1370023:1370023 -
NANP
chr6:3450287:3450292 -
APRR1
chr9:12917280:12917280 -
NDRG4
chr2:20122036:20122036 -
PLPL
chr4:1030570:1030570:1030 +
GDF1
chr12:6409849:6409850:1 -
GRTP1
chr5:150450854:150450854 +
CNNM2
chr10:6780854:6780854:- -
ATP5D4
chr1:2025670:2025670:2025 +
PMP
chr1:156230158:156230159 +
BIN1
chr10:127860854:127860858 -
WIPF3
chr17:75258940:75258978:75258789 +
ATF7
chr4:1041780:1041779:1041 -
PDCD10
chr3:9089427:9089 +
RC1
chr7:3960925:3960927 -
CYP2R1
chr11:14350239:14350239 +
DLG3
chr4:88532028:88532028 +
NR3C
chr16:8819600:8819633 -
DNG
chr9:3154370:3154820 -
CLA
chr4:1000427:1000427 -
NPDC
chr17:2191878:2191924 +
PCF1
chr8:1026270:1026269 +
PRL7
chr3:5:5:5:5:5:5:5:5:5:5 -
ZNF638
chr17:75400000:75400001 +
NANP
chr17:80420:80420:80420 +
PAAT
chr11:7100000:7100144 +
CNN1
chr19:15041:15041:15041:15041 -
FUNC
chr10:1101:1101:1101:1101 -
PLCL
chr3:1386:1386:1386:1386 -
PRX
chr16:4872:4872:4872:4872 -
PRX
chr4:14431:2587:14431:2587 -
SLC16A9
chr17:1049000:1049074 +
ENAH
chr9:130200:130200:130 +
TAAR9
chr6:2963:2963:2963:2963 -
USP14
chr21:30300:30300:30300 +
CDHR
chr5:64000:64000:64000 -
MAGOH
chr12:5329207:5329209 +
PPP1R1A
chr1:1155000:1155000:1155000 -
ZNF
chr9:1003:0:0:0:0:1003:1003 -
FUBGL4
chr12:4858073:4858080 -
PRTG
chr17:2736:2792:2736:2836 +
RNMT
chr1:2050845:2050845:2050845:2 -
CYP2F
chr15:2200200:2200200 +
MPN2
chr1:1204200:1204204 +
DAH
chr20:1749000:1749000:1749000 -
PGBD1
chr20:3862200:3862477 +
CYP2D6
chr22:42300:42300:42300 +
PPP1R1
chr1:1204200:1204200 +
DDX5
chr19:1000:0:0:0:0:0:0:0:0:0 -
PPP1R1
chr19:4862000:4862000 +
CYP2D6
chr22:42300:42300:42300 +
PPP1R1