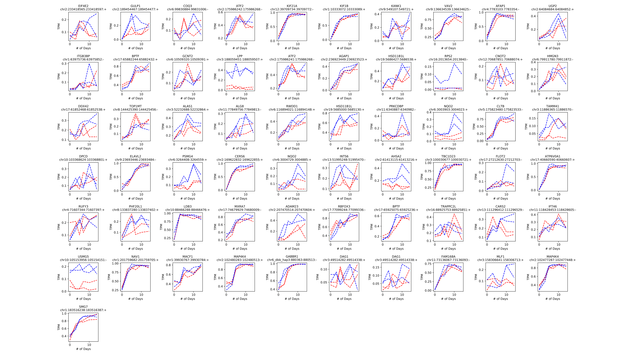
DFNB2
chr2:210418562..210418597 +
GJA1
chr2:108494467..108543477 +
CX26
chr13:20189282..20189286 +
ATP1
chr1:173988516..173988568 -
KIF21A
chr12:39956716..39978772 -
AIFM
chr1:51378422..51418389 +
KDM6
chrX:44822528..44859000 +
MID1
chrX:10464728..10624425 -
ATAT1
chr6:74032278..74032354 +
USP7
chr16:8408468..8468452 +
FOXL2
chr3:138777600..138777636 +
BPP
chrX:153611944..153685432 +
SOX3
chrX:139551140..139552941 +
ATP7
chrX:77747103..77789628 -
ADAP1
chr7:27852676..27852723 +
HSD11B1
chr1:207000132..207000196 +
AP2
chr19:45728512..45728543 -
CNOT3
chr19:7007175..7048878 +
DMD
chrX:79839602..79917872 -
FOXA2
chr20:22547..36480 +
SOX10
chr22:38380290..38380324 +
ZEB2
chr2:144512453..144512495 +
ITPA
chr20:3132..3149 +
ALX1
chr12:84520..84539 +
RCN
chr11:17394737..17394813 -
RBFOX1
chr16:7189645..7189722 +
ANKRD11
chr16:89228434..89228474 +
PRKCSH
chr19:8696282..8696313 +
NOS1
chr12:117405307..117405350 +
CLTB
chr3:179623983..179623998 -
SMPD1
chr11:6382397..6382755 +
DRC7
chr22:29168822..29168883 +
ELANE
chr19:852..1040 +
PSMA8
chr20:62884..62900 +
CERS6
chr2:69922553..69962305 +
NOS3
chr7:150642736..150695400 +
MYO5A
chr15:52533415..52533435 +
RAB39B
chrX:154412442..154412478 +
TBC1D23
chr3:184150927..184150971 +
FLOT2
chr17:27213052..27217703 +
ATP6AP2
chrX:47995650..48040607 +
RUNX1
chr21:36160777..36160837 +
PHF6
chrX:133452615..133452655 +
ANKRD11
chr16:89228434..89228474 +
ATM
chr11:108093559..108093611 +
CHD7
chr8:61772353..61772374 +
SMARCA2
chr9:2046492..2046557 +
ZEB2
chr2:144512453..144512495 +
GNB1
chr1:1744854..1744894 +
PACS1
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ANKRD11
chr16:89228434..89228474 +
SOX10
chr22:38380290..38380324 +
SMARCA2
chr9:2046492..2046557 +
ATP6AP2
chrX:47995650..48040607 +
SOX3
chrX:139551140..139552941 +
ATP7A
chrX:77747103..77789628 +
RBFOX1
chr16:7189645..7189722 +
ATM
chr11:108093559..108093611 +
TRAPPC9
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CHD7
chr8:61772353..61772374 +
CHD7
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ANKRD11
chr16:89228434..89228474 +
SOX10
chr22:38380290..38380324