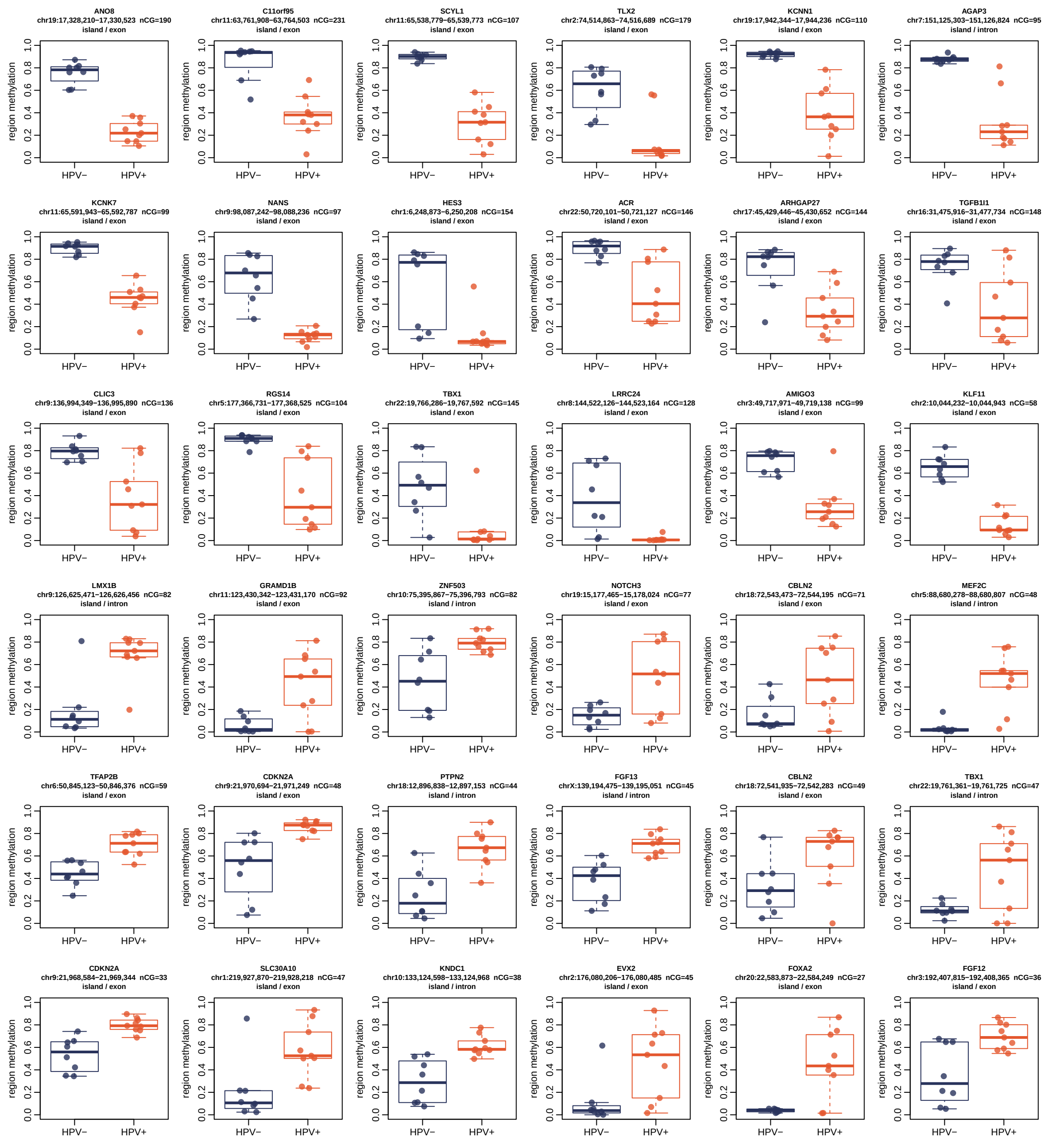
ANO8
chr19:17,328,210−17,330,523 nCG=190
island / exon
C11orf95
chr11:63,761,908−63,764,503 nCG=231
island / exon
SCYL1
chr11:65,538,779−65,539,773 nCG=107
island / exon
TLX2
chr2:74,514,863−74,516,689 nCG=179
island / exon
KCNN1
chr19:17,942,344−17,944,236 nCG=110
island / exon
AGAP3
chr7:151,125,303−151,126,824 nCG=95
island / intron
KCNK7
chr11:65,591,943−65,592,787 nCG=99
island / exon
NANS
chr9:98,087,242−98,088,236 nCG=97
island / exon
HES3
chr1:6,248,873−6,250,208 nCG=154
island / exon
ACR
chr22:50,720,101−50,721,127 nCG=146
island / exon
ARHGAP27
chr17:45,429,446−45,430,652 nCG=144
island / exon
TGFB1I1
chr16:31,475,916−31,477,734 nCG=148
island / exon
CLIC3
chr9:136,994,349−136,995,890 nCG=136
island / exon
RGS14
chr5:177,366,731−177,368,525 nCG=104
island / exon
TBX1
chr22:19,766,286−19,767,592 nCG=145
island / exon
LRRC24
chr8:144,522,126−144,523,164 nCG=128
island / exon
AMIGO3
chr3:49,717,971−49,719,138 nCG=99
island / exon
KLF11
chr2:10,044,232−10,044,943 nCG=58
island / exon
LMX1B
chr9:126,625,471−126,626,456 nCG=82
island / intron
GRAMD1B
chr11:123,430,342−123,431,170 nCG=92
island / exon
ZNF503
chr10:75,395,867−75,396,793 nCG=82
island / intron
NOTCH3
chr19:15,177,465−15,178,024 nCG=77
island / exon
CBLN2
chr18:72,543,473−72,544,195 nCG=71
island / exon
MEF2C
chr5:88,680,278−88,680,807 nCG=48
island / intron
TFAP2B
chr6:50,845,123−50,846,376 nCG=59
island / exon
CDKN2A
chr9:21,970,694−21,971,249 nCG=48
island / exon
PTPN2
chr18:12,896,838−12,897,153 nCG=44
island / intron
FGF13
chrX:139,194,475−139,195,051 nCG=45
island / intron
CBLN2
chr18:72,541,935−72,542,283 nCG=49
island / exon
TBX1
chr22:19,761,361−19,761,725 nCG=47
island / intron
CDKN2A
chr9:21,968,584−21,969,344 nCG=33
island / exon
SLC30A10
chr1:219,927,870−219,928,218 nCG=47
island / exon
KNDC1
chr10:133,124,598−133,124,968 nCG=38
island / intron
EVX2
chr2:176,080,206−176,080,485 nCG=45
island / exon
FOXA2
chr20:22,583,873−22,584,249 nCG=27
island / exon
FGF12
chr3:192,407,815−192,408,365 nCG=36
island / exon
region methylation
0.0
0.2
0.4
0.6
0.8
1.0
HPV−
HPV+