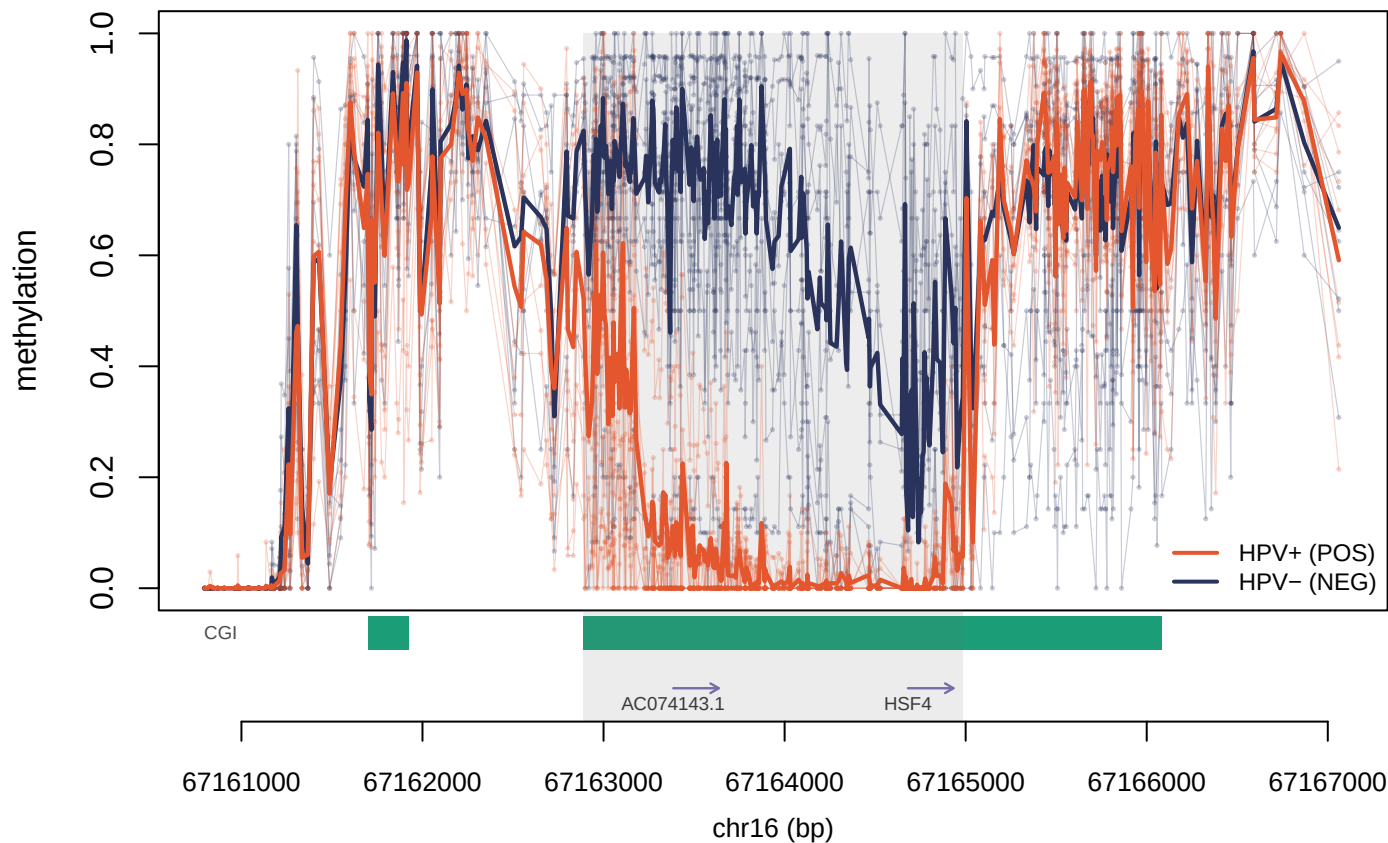
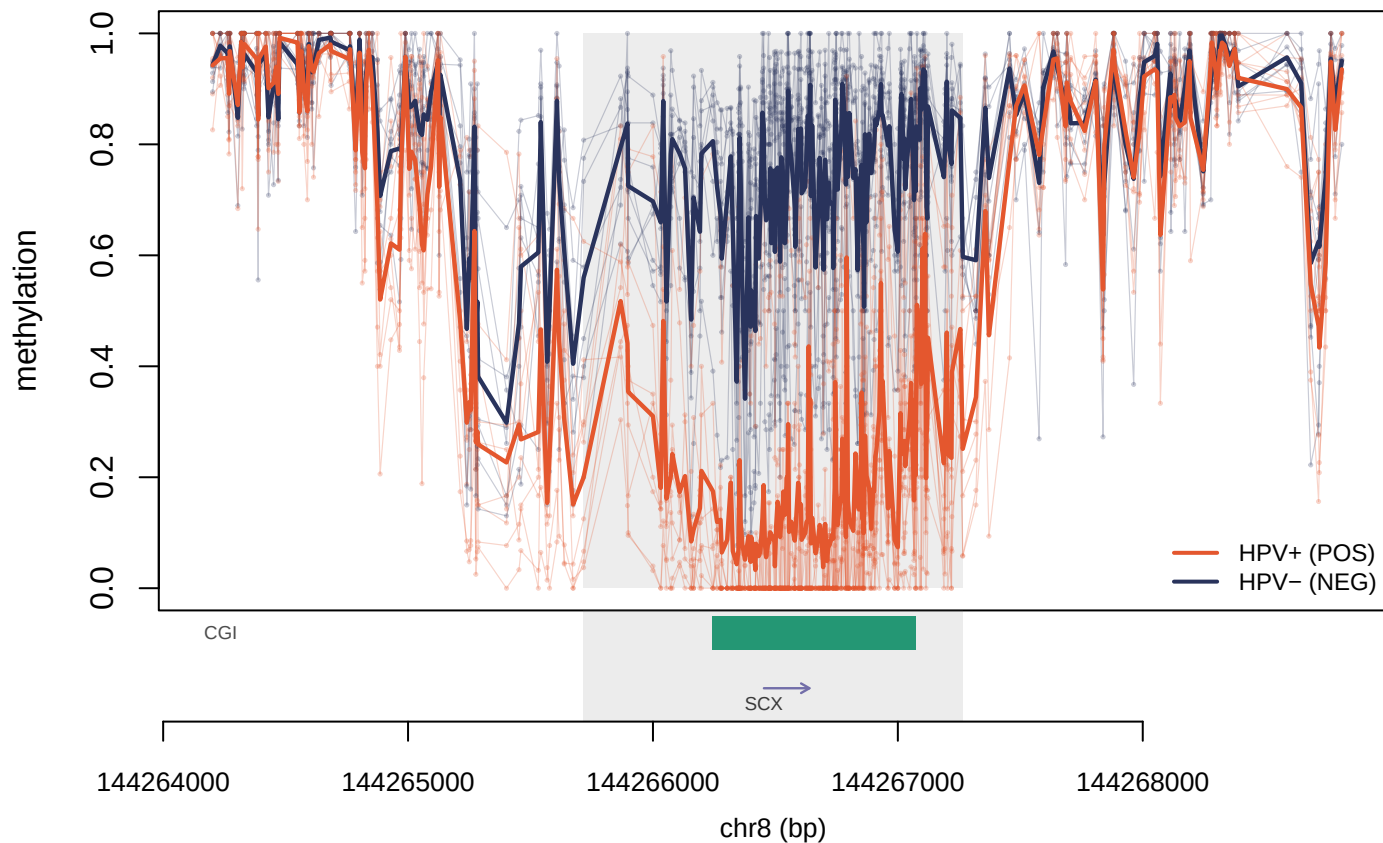


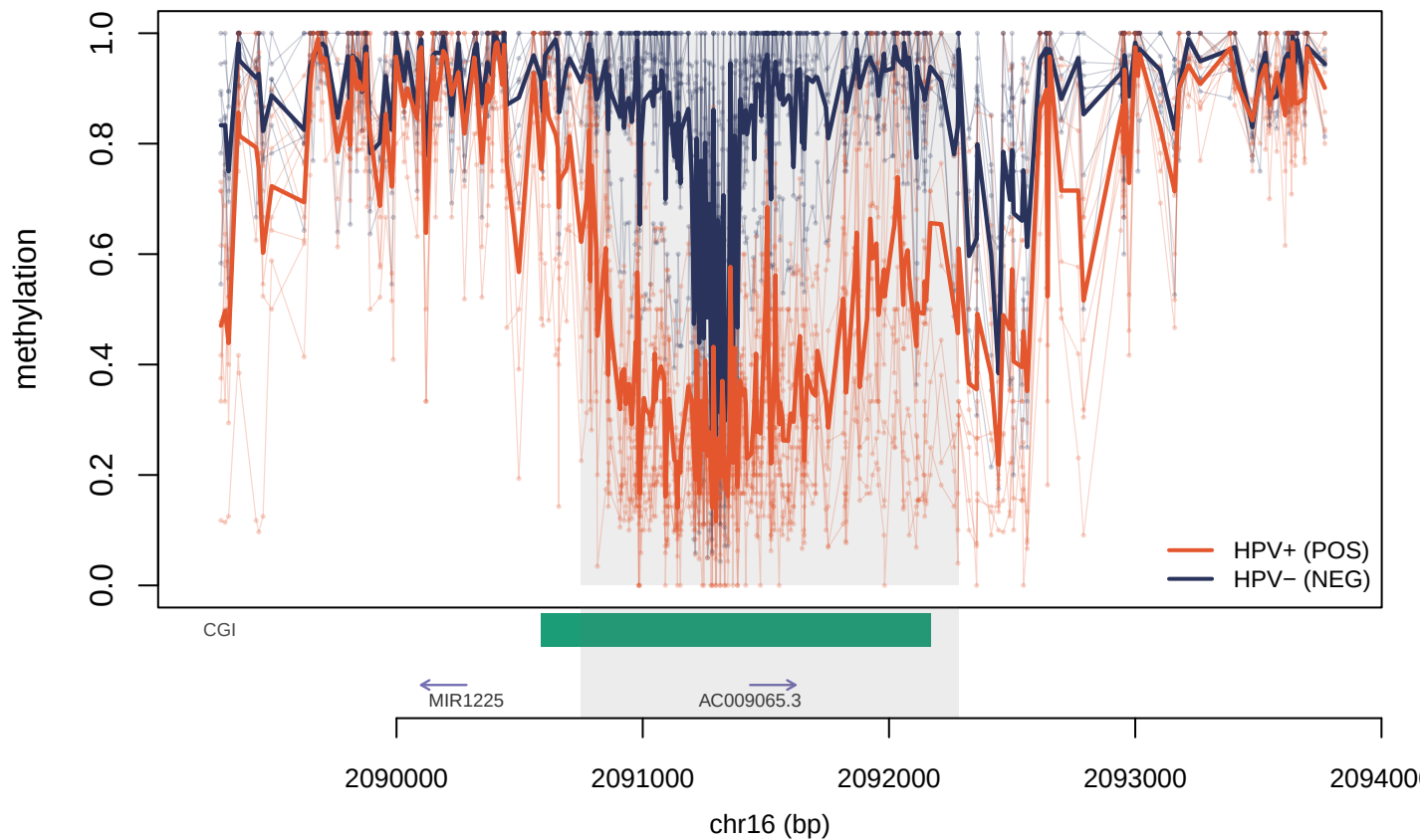
HSF4 chr16:67,162,890–67,164,984 nCG=205 island / promoter hypo (diff -0.49)



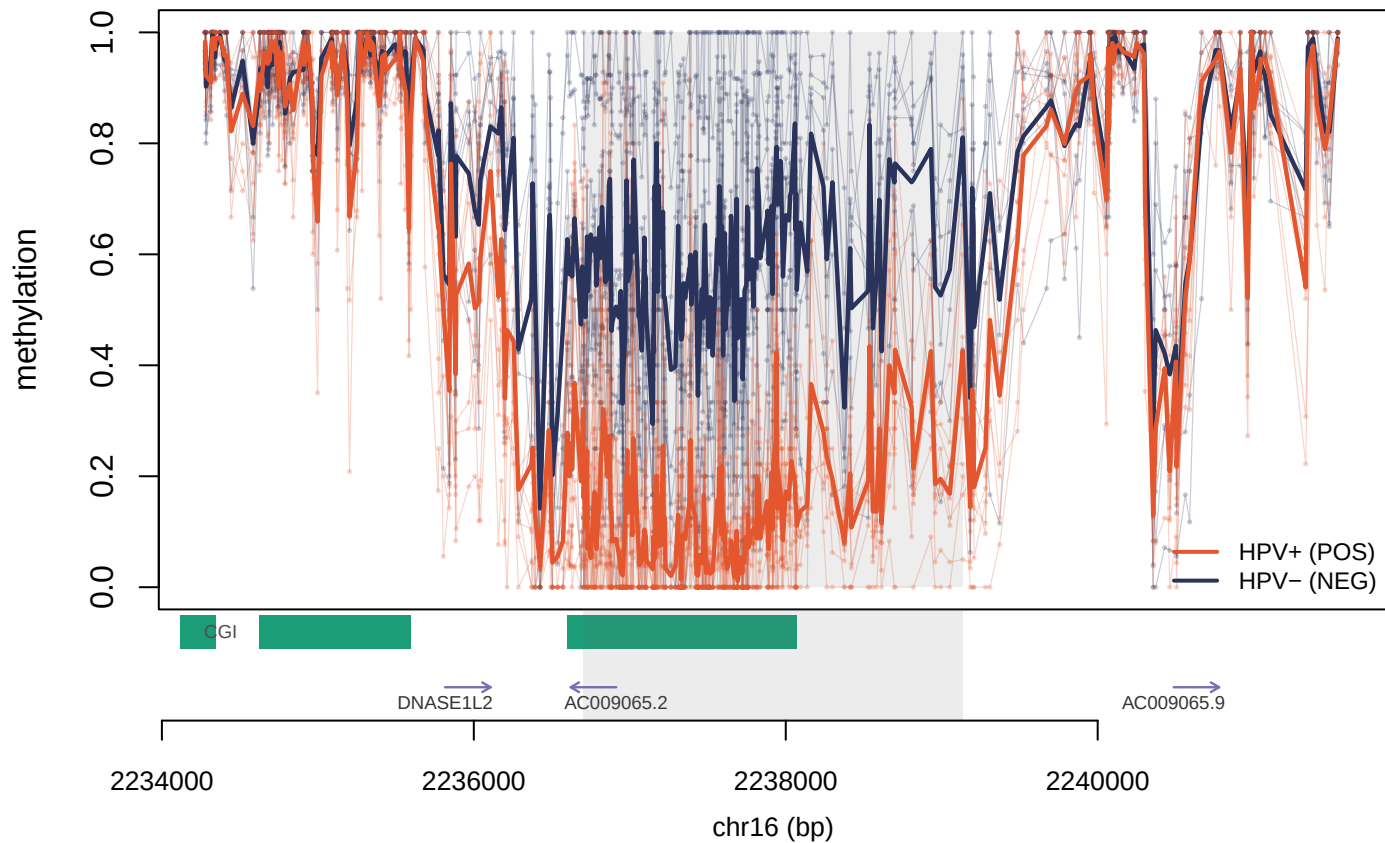
SCX chr8:144,265,717–144,267,265 nCG=162 island / promoter hypo (diff -0.53)



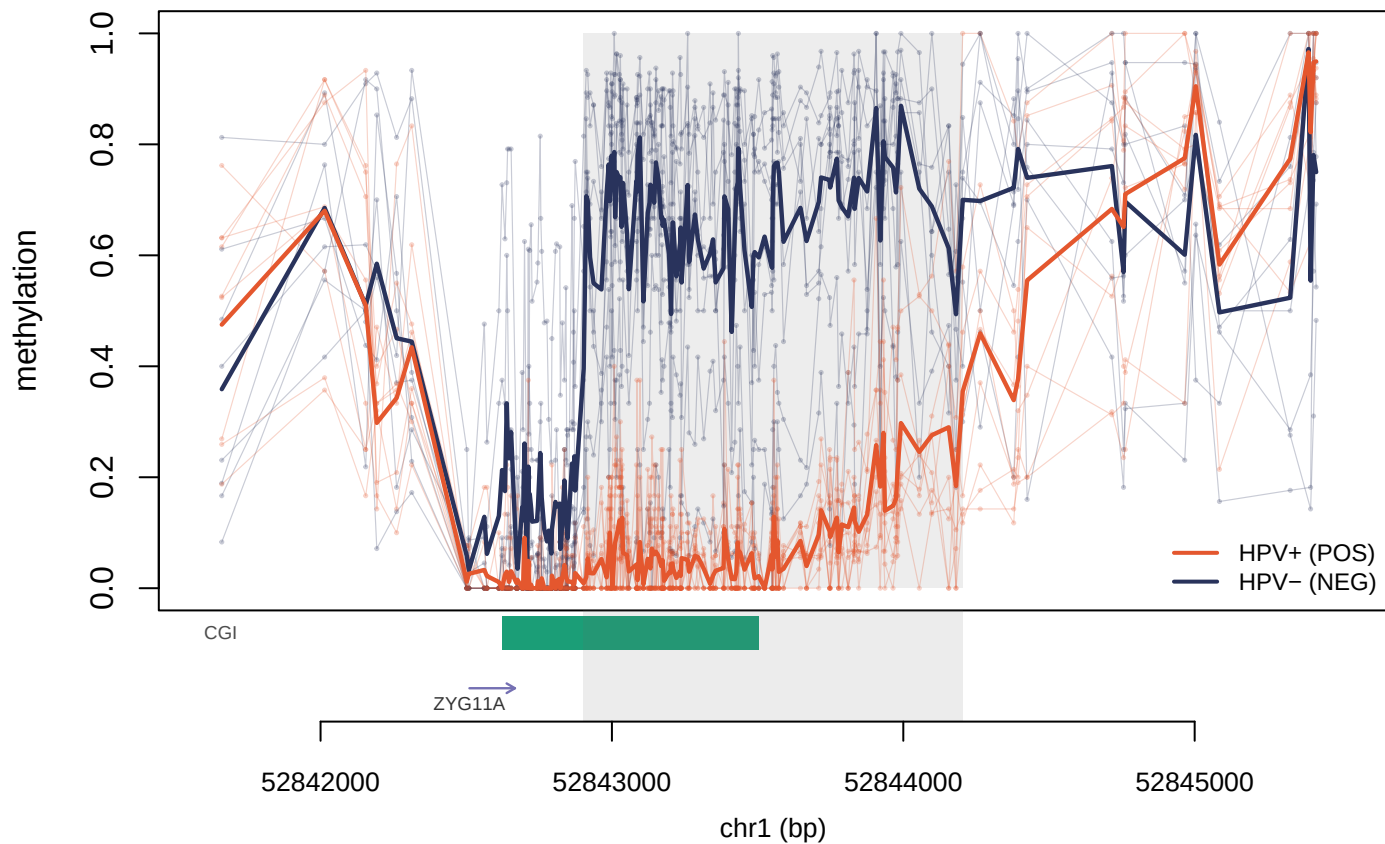
TSC2 chr16:2,090,750–2,092,283 nCG=155 island / promoter hypo (diff -0.48)



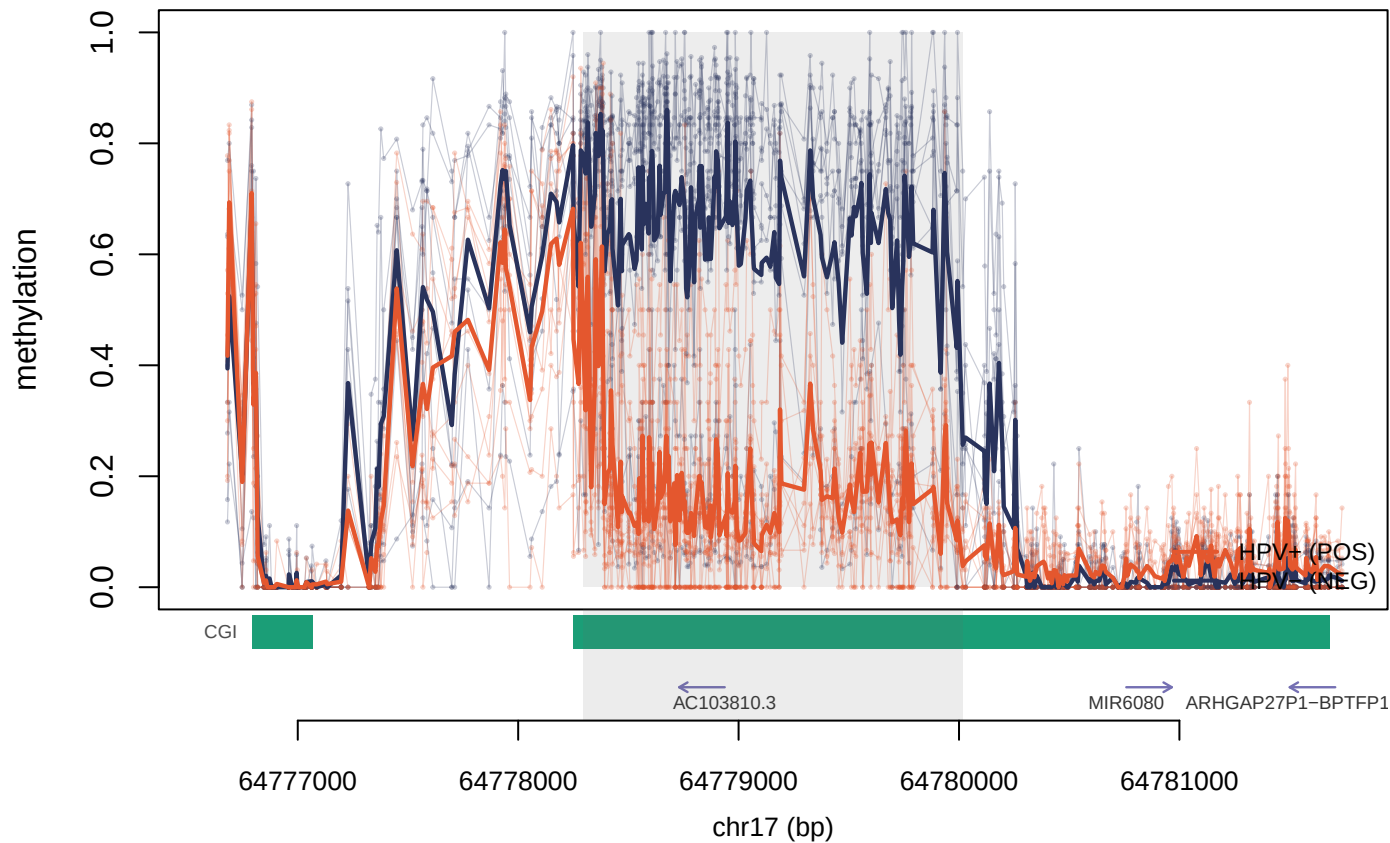
DNASE1L2 chr16:2,236,704–2,239,135 nCG=176 island / promoter hypo (diff -0.43)



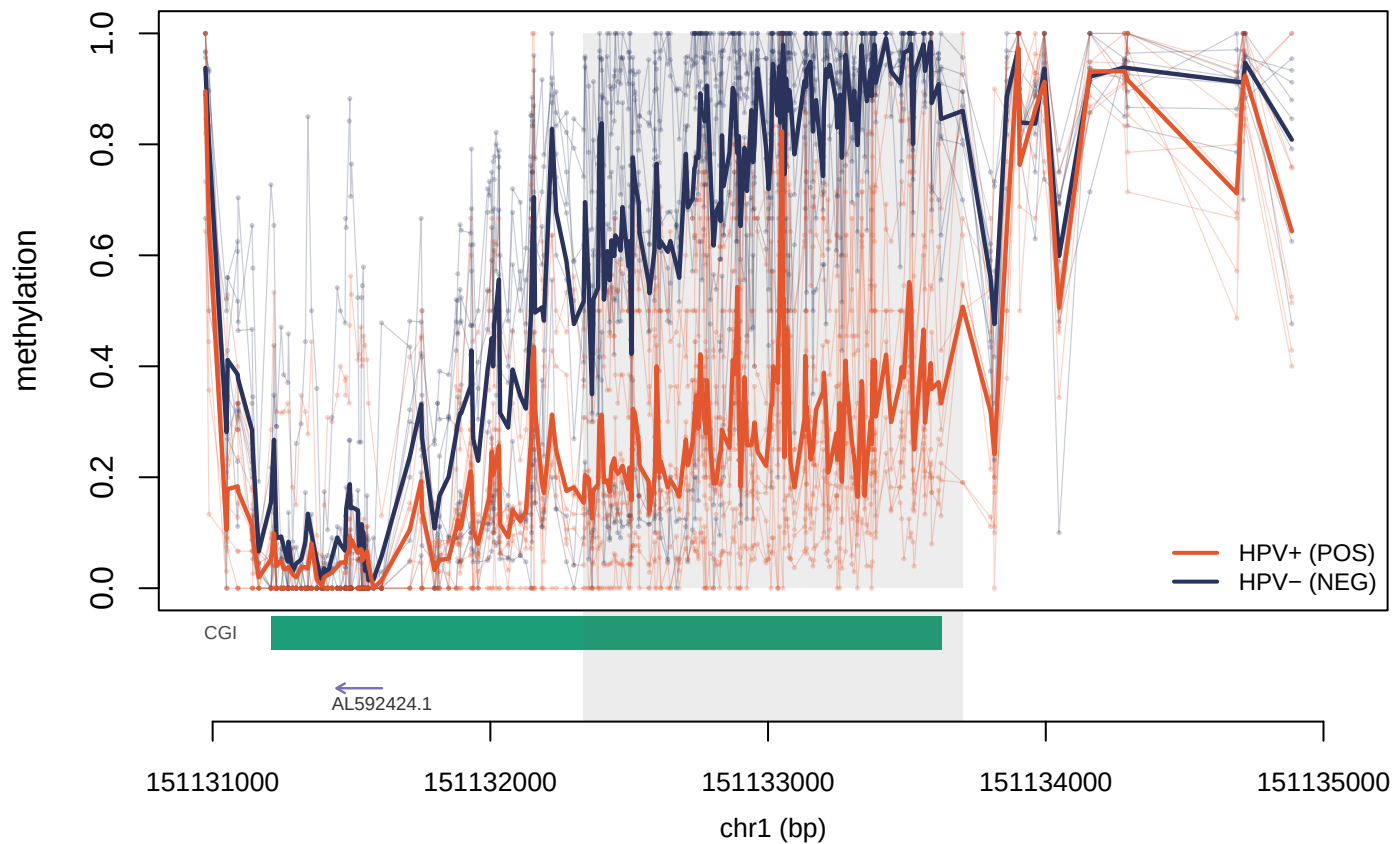
ZYG11A chr1:52,842,903–52,844,204 nCG=105 island / promoter hypo (diff -0.55)



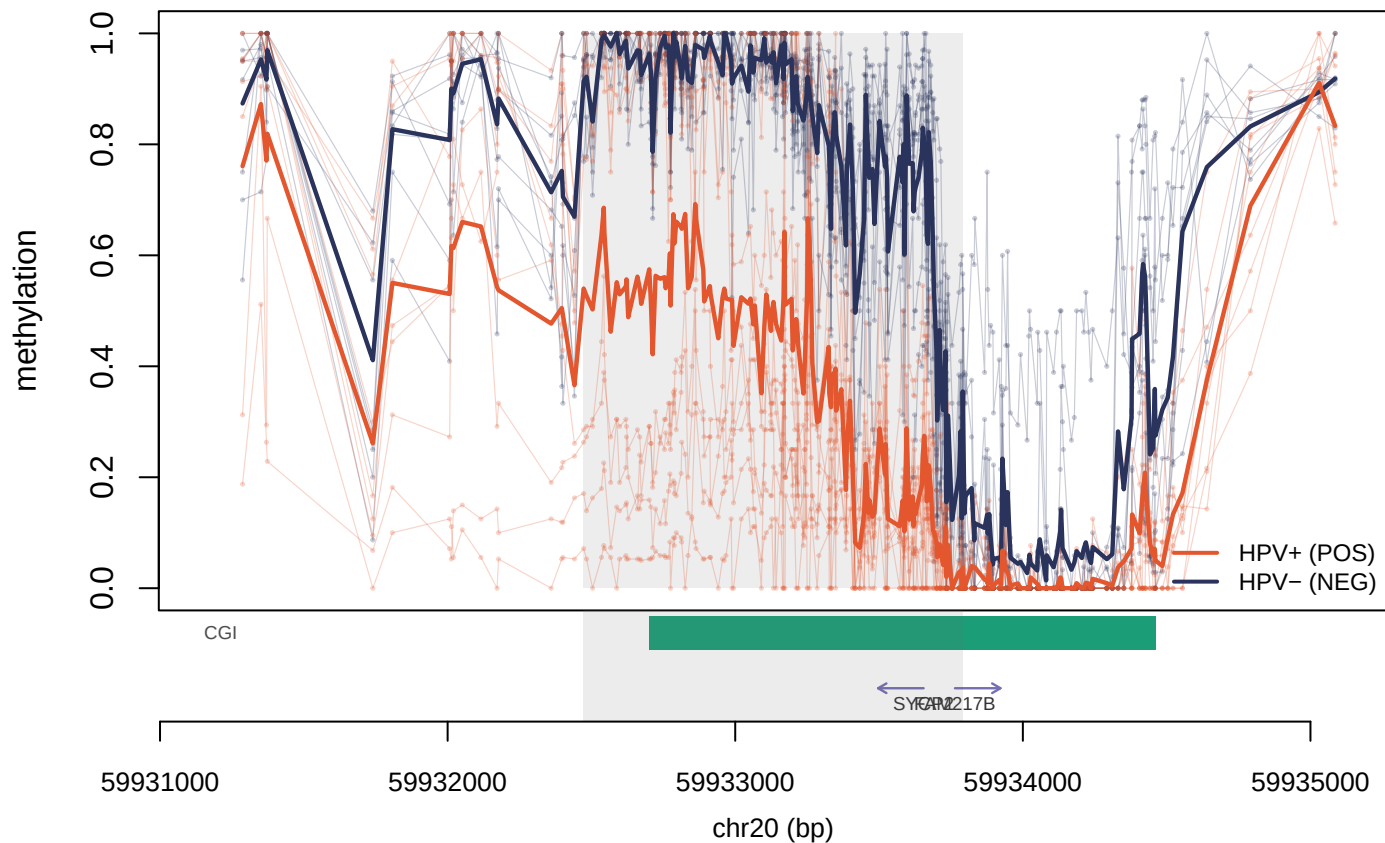
SMURF2 chr17:64,778,297-64,780,017 nCG=158 island / promoter hypo (diff -0.43)



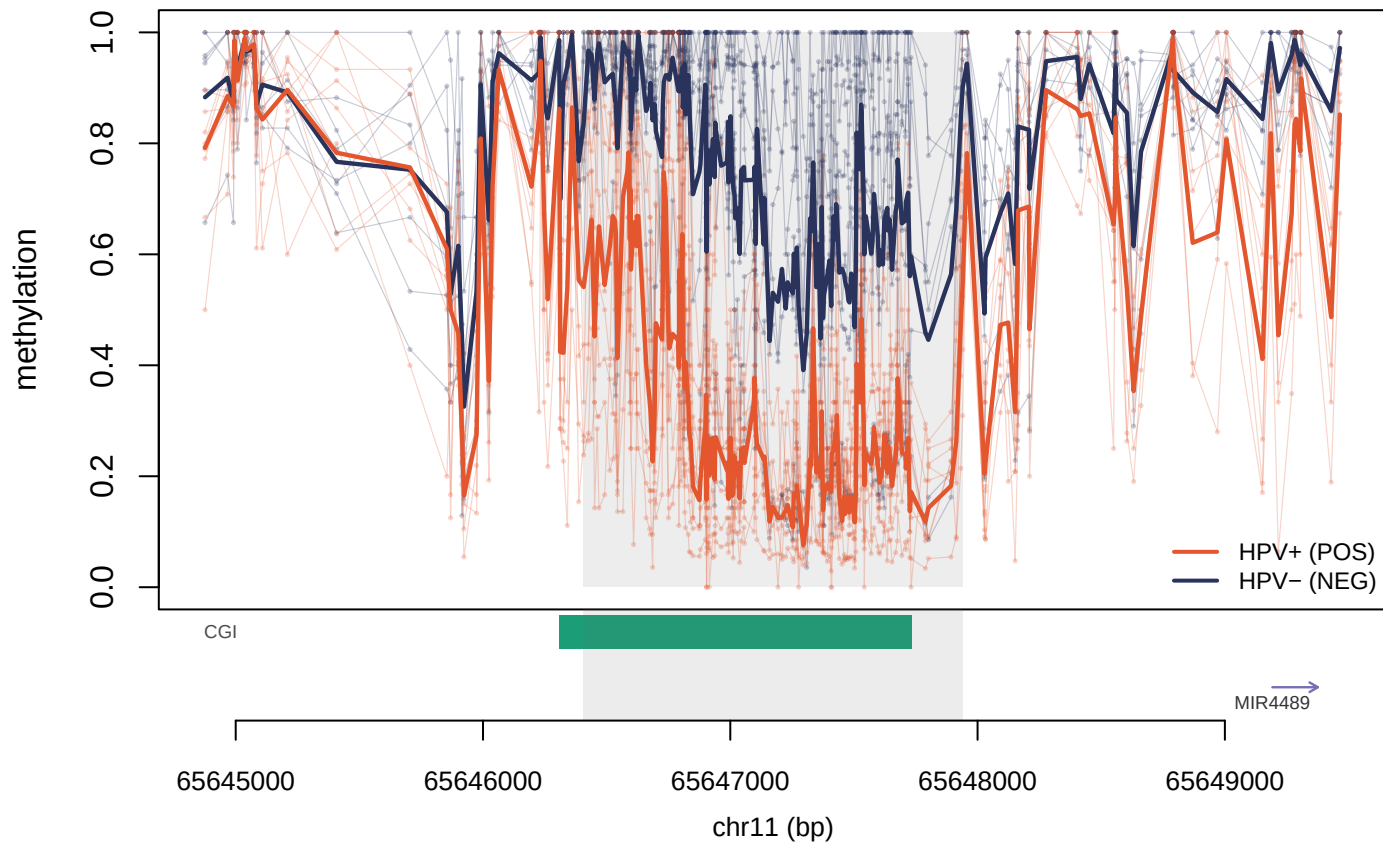
SEMA6C chr1:151,132,336–151,133,701 nCG=138 island / promoter hypo (diff -0.46)



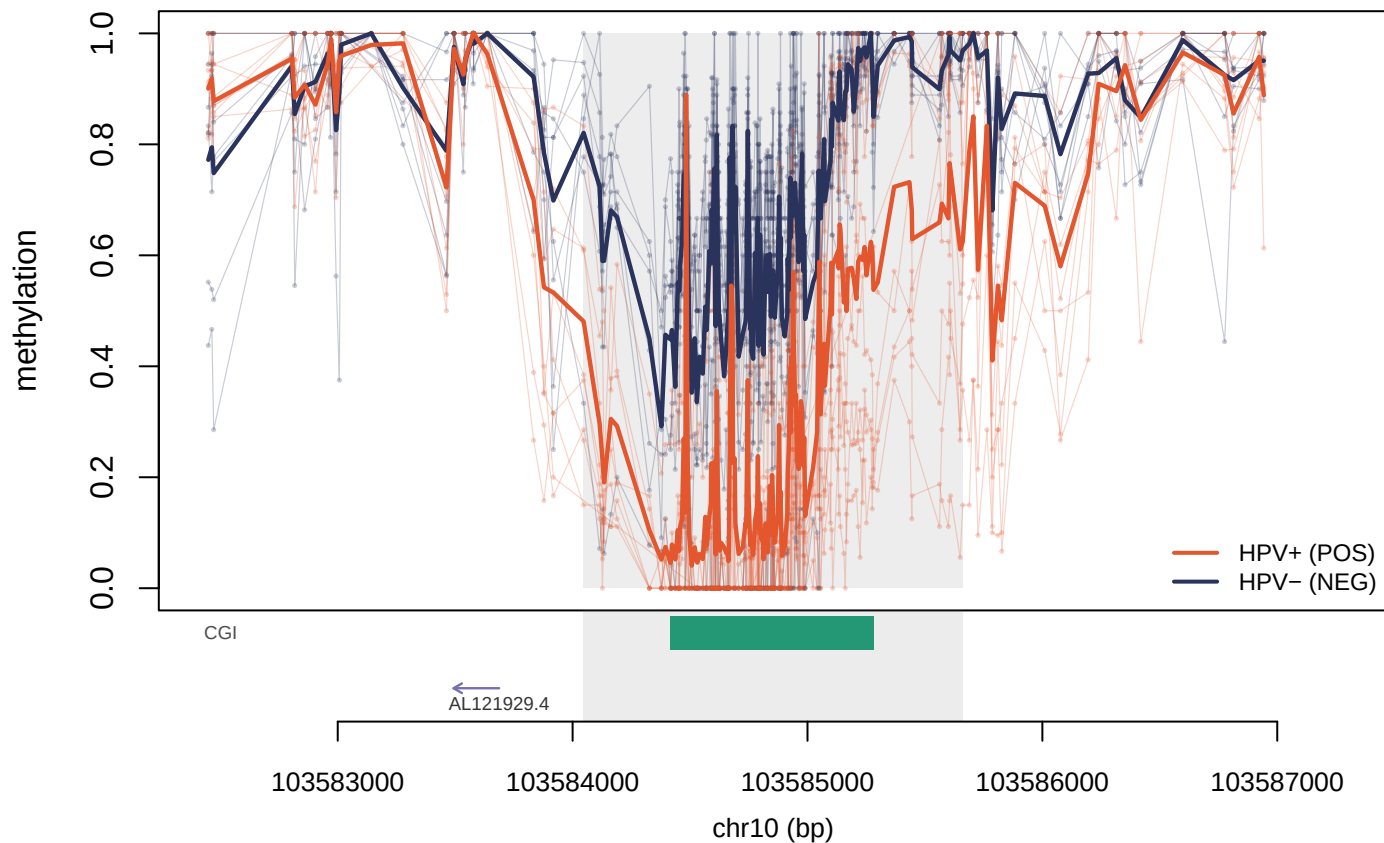
FAM217B chr20:59,932,473–59,933,791 nCG=141 island / promoter hypo (diff -0.43)



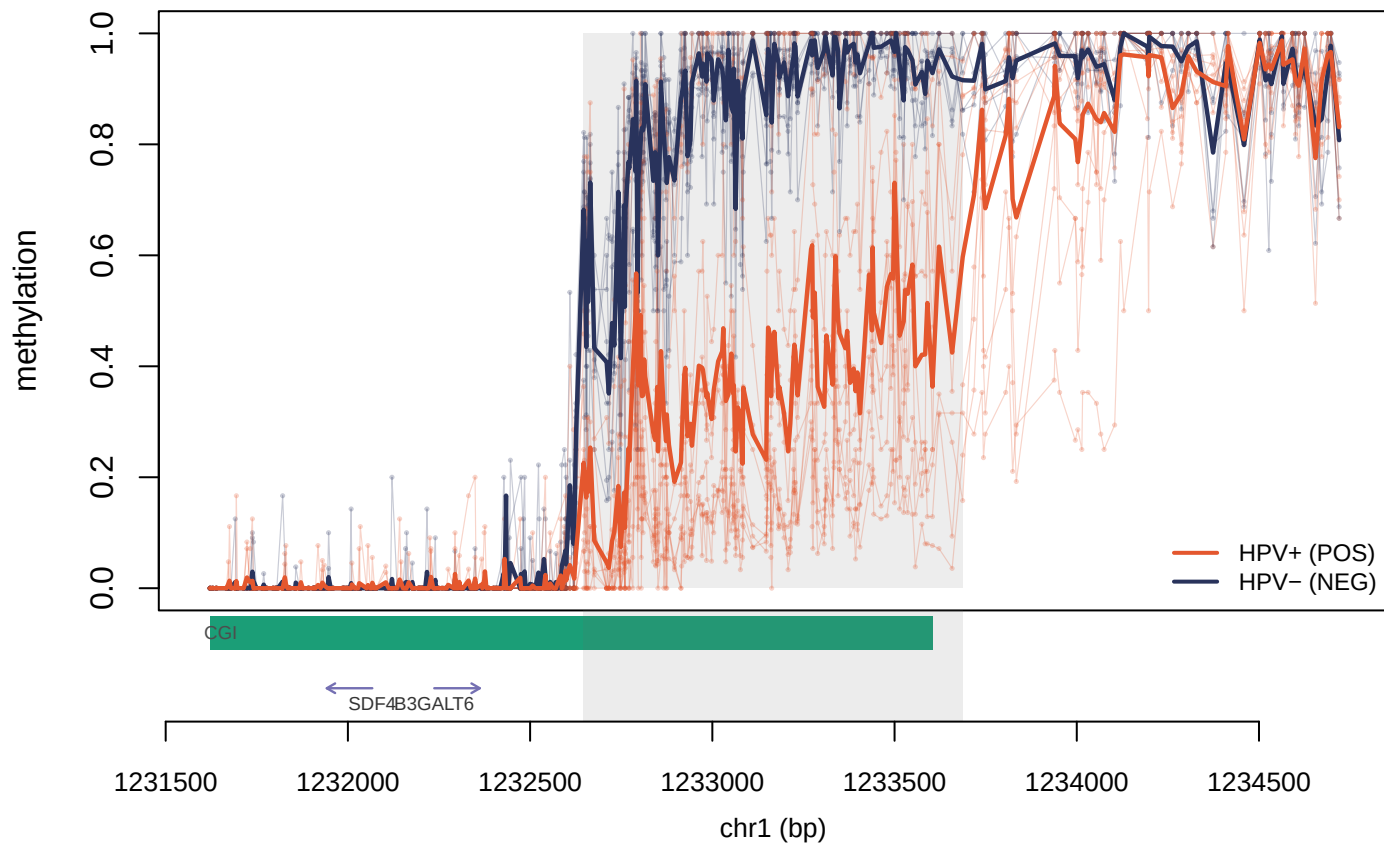
SIPA1 chr11:65,646,407–65,647,940 nCG=160 island / promoter hypo (diff -0.36)



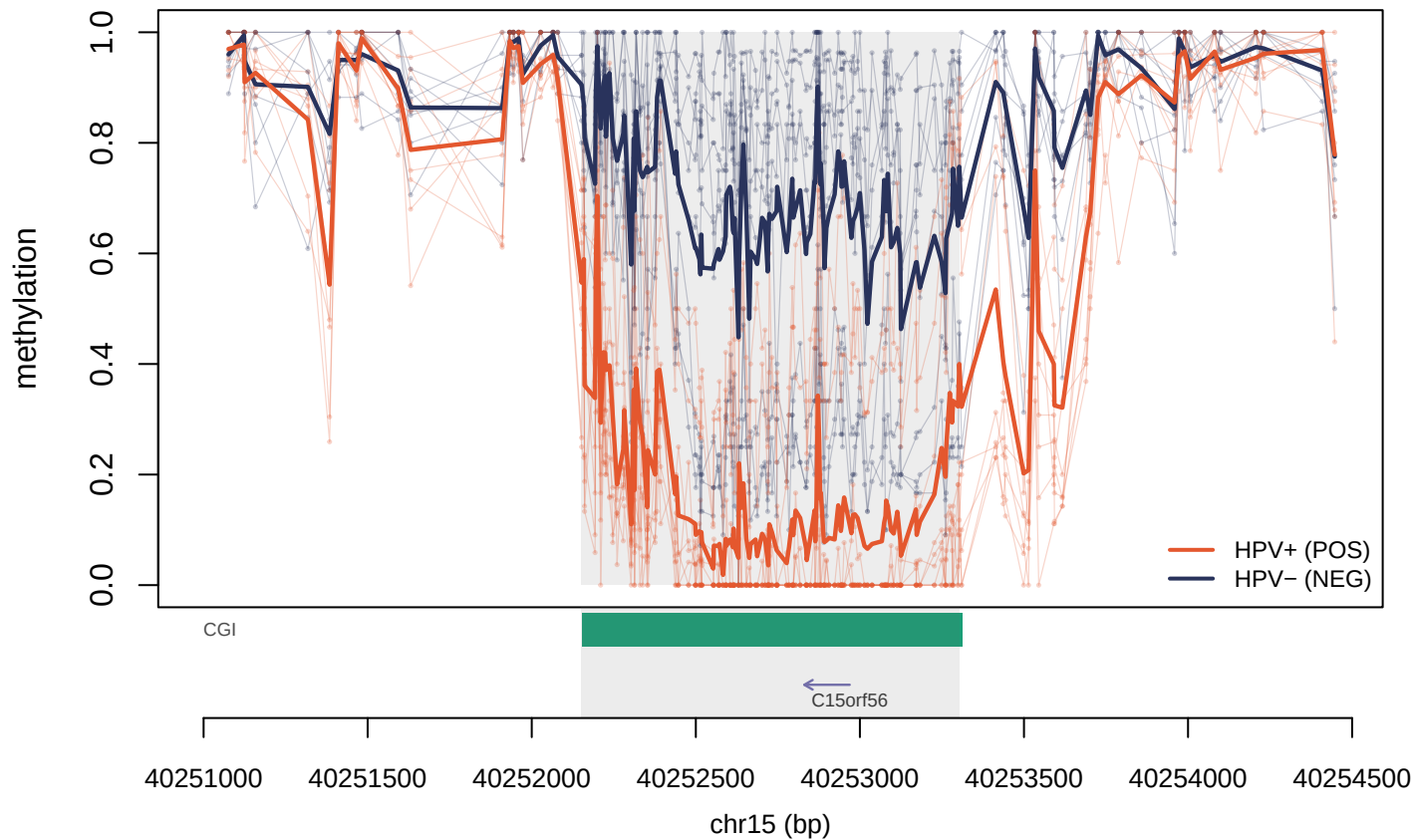
NEURL1 chr10:103,584,047–103,585,661 nCG=159 island / promoter hypo (diff -0.37)



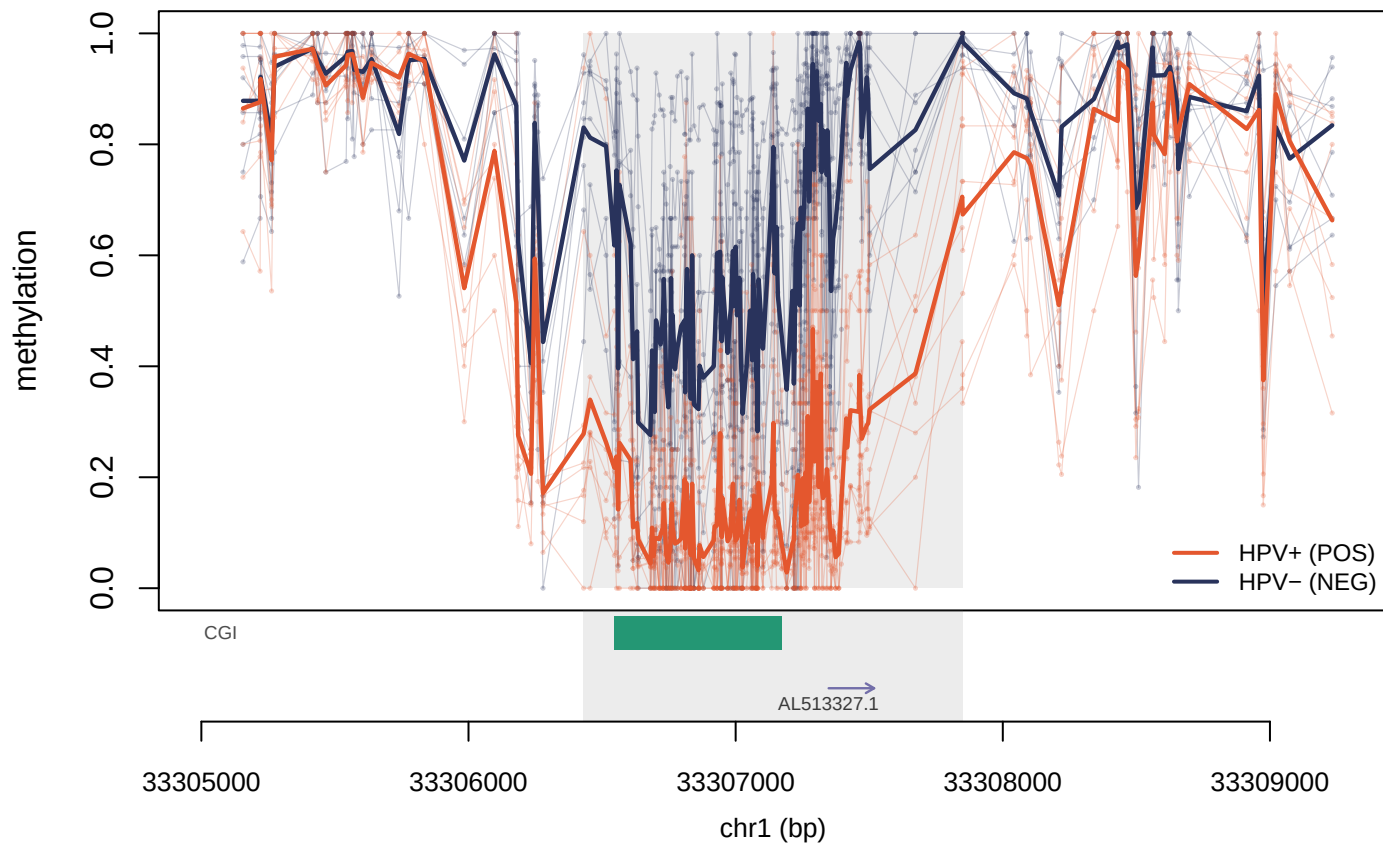
B3GALT6 chr1:1,232,647–1,233,687 nCG=128 island / promoter hypo (diff -0.47)



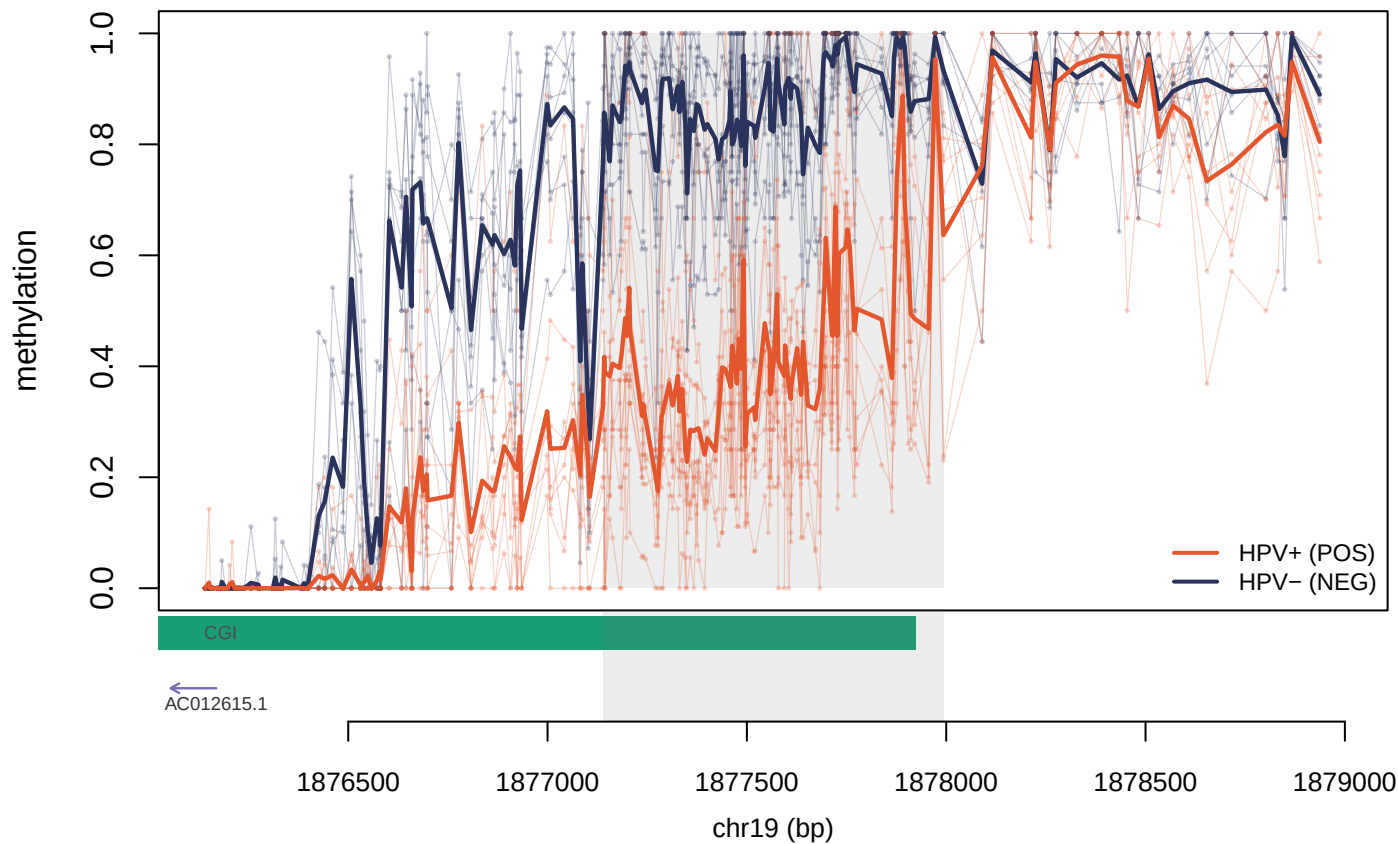
ANKRD63 chr15:40,252,152–40,253,303 nCG=122 island / promoter hypo (diff -0.49)



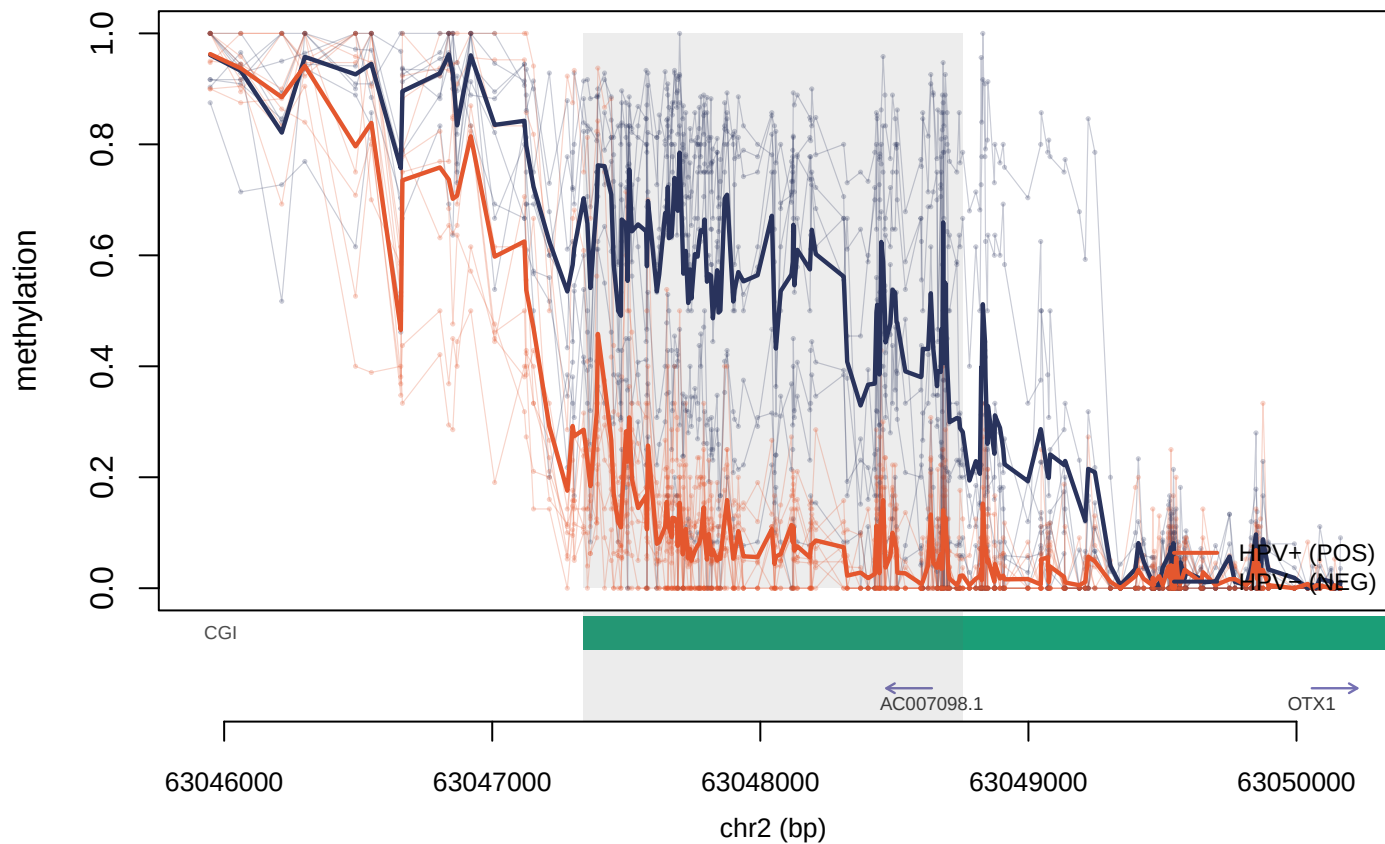
A3GALT2 chr1:33,306,431–33,307,850 nCG=129 island / promoter hypo (diff -0.40)



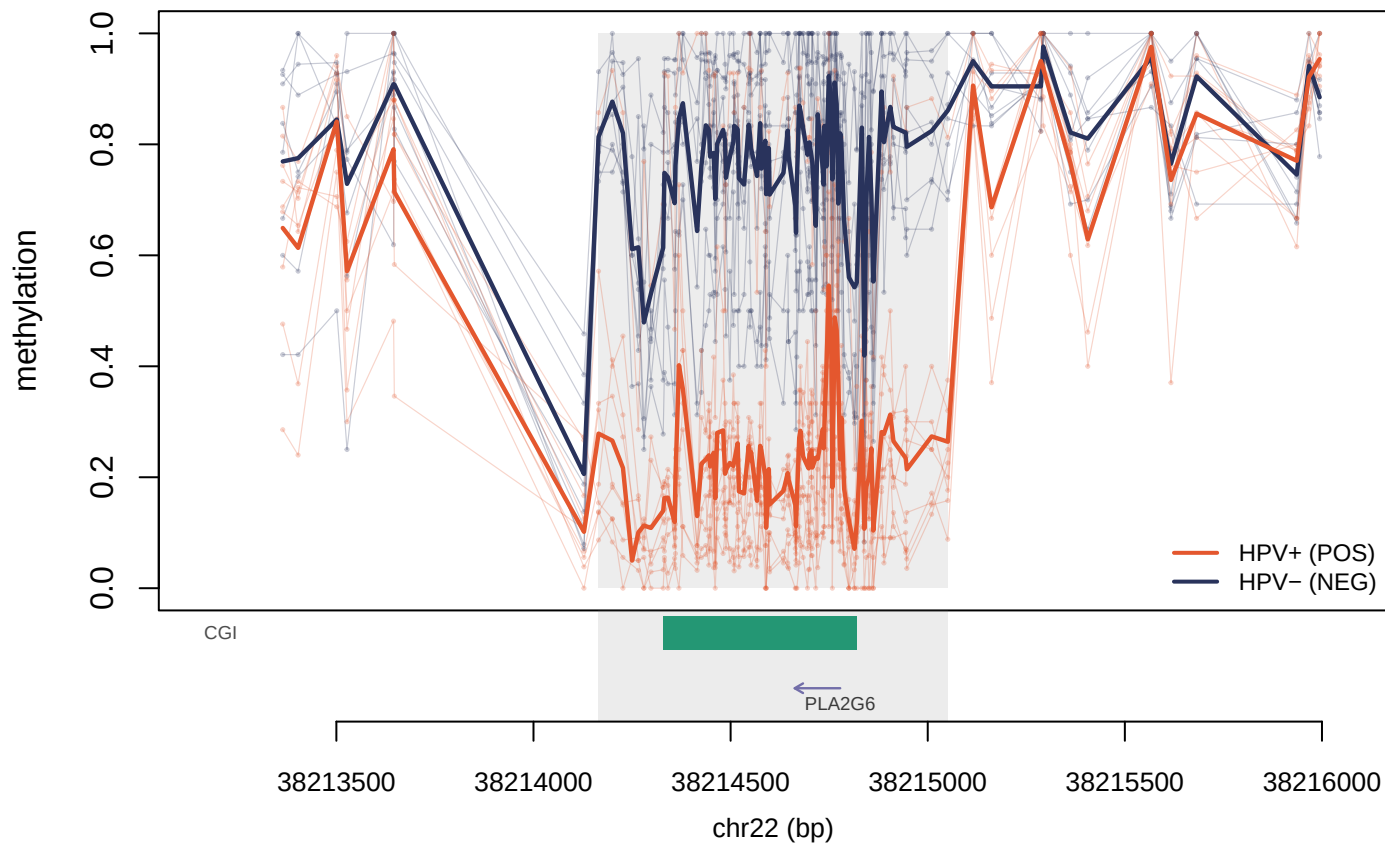
ABHD17A chr19:1,877,139–1,877,993 nCG=93 island / promoter hypo (diff -0.45)



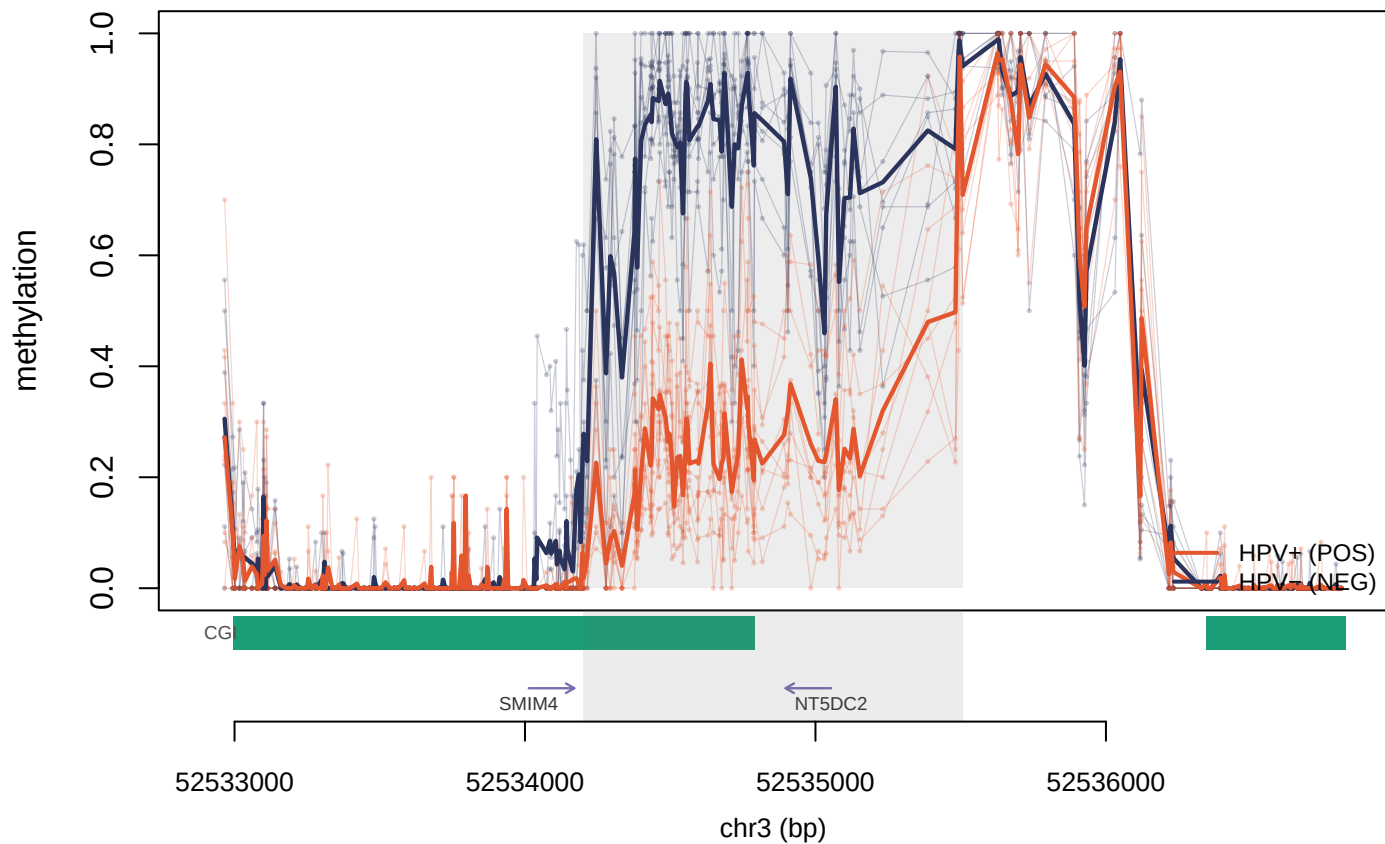
OTX1 chr2:63,047,341–63,048,755 nCG=111 island / promoter hypo (diff -0.42)



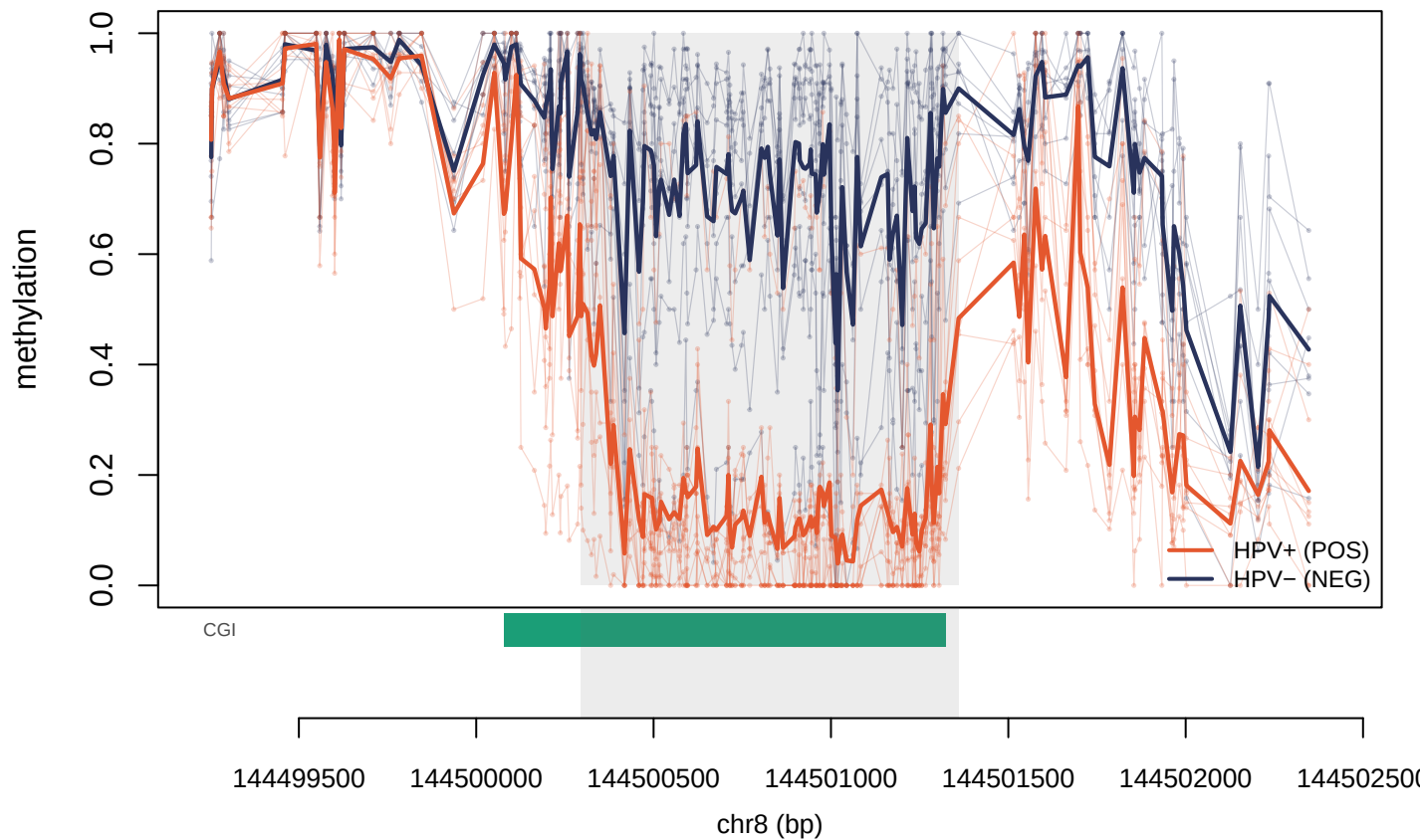
PLA2G6 chr22:38,214,165–38,215,051 nCG=88 island / promoter hypo (diff -0.52)



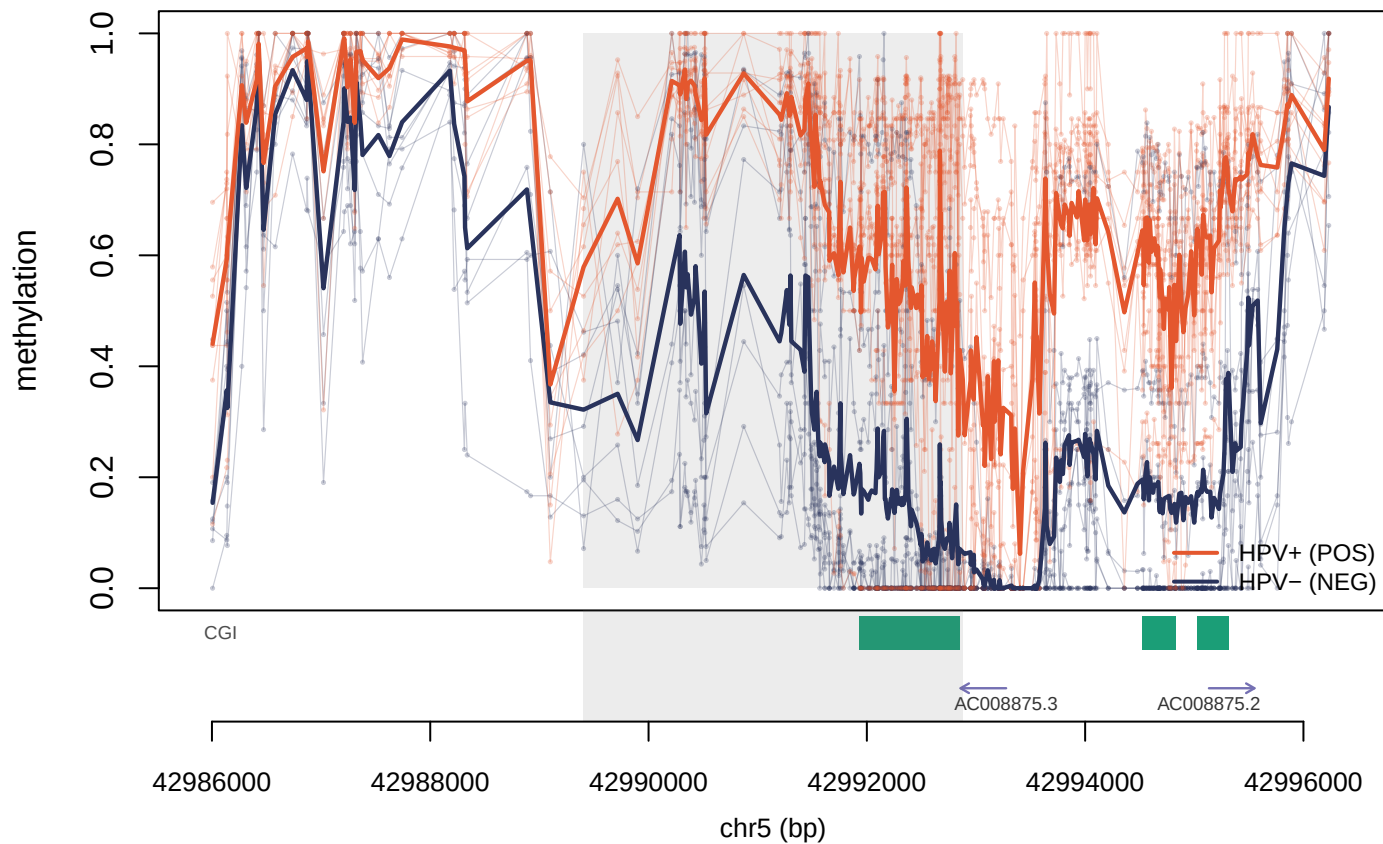
NT5DC2 chr3:52,534,202–52,535,507 nCG=68 island / promoter hypo (diff -0.49)



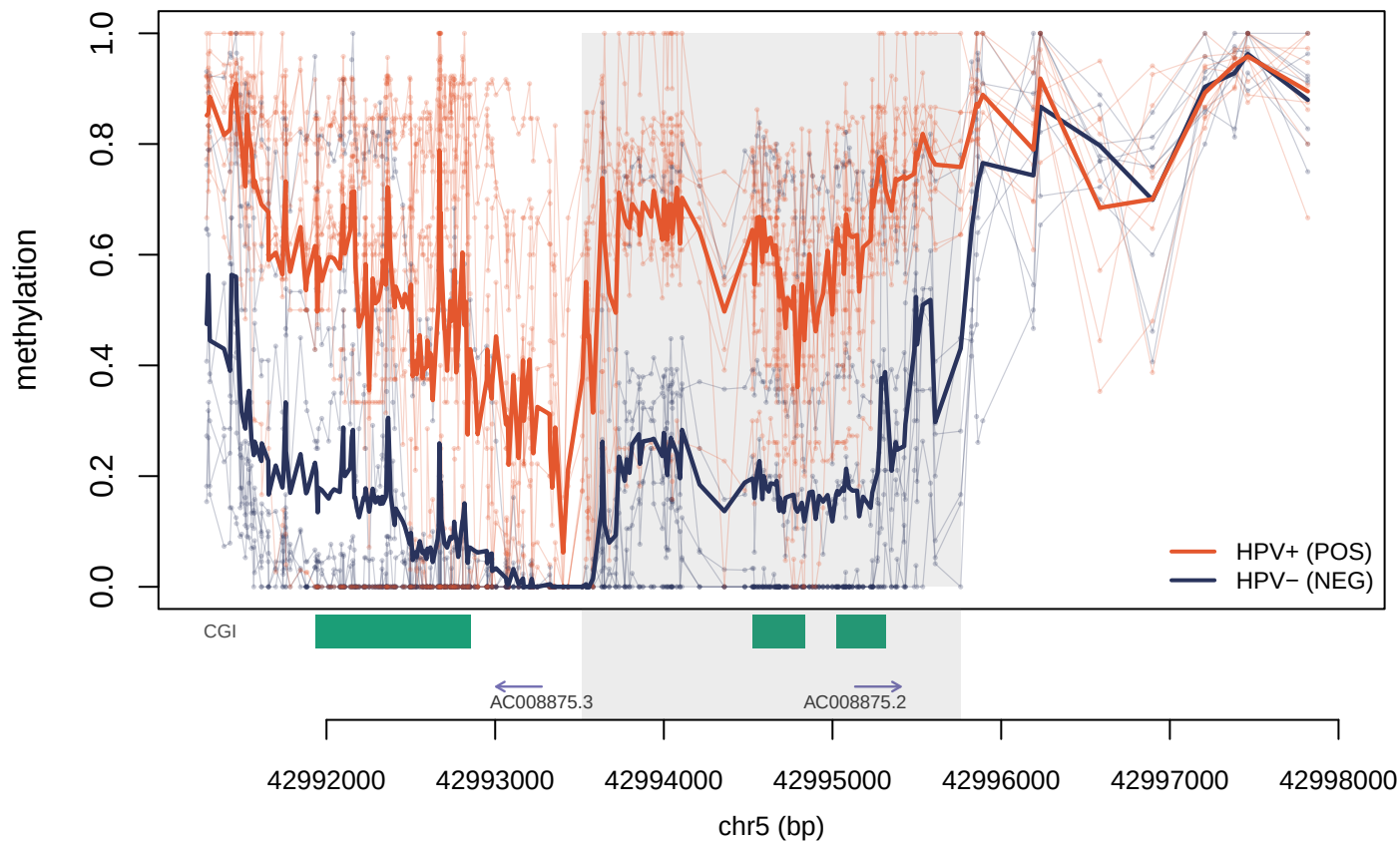
GPT chr8:144,500,296–144,501,360 nCG=95 island / promoter hypo (diff -0.49)



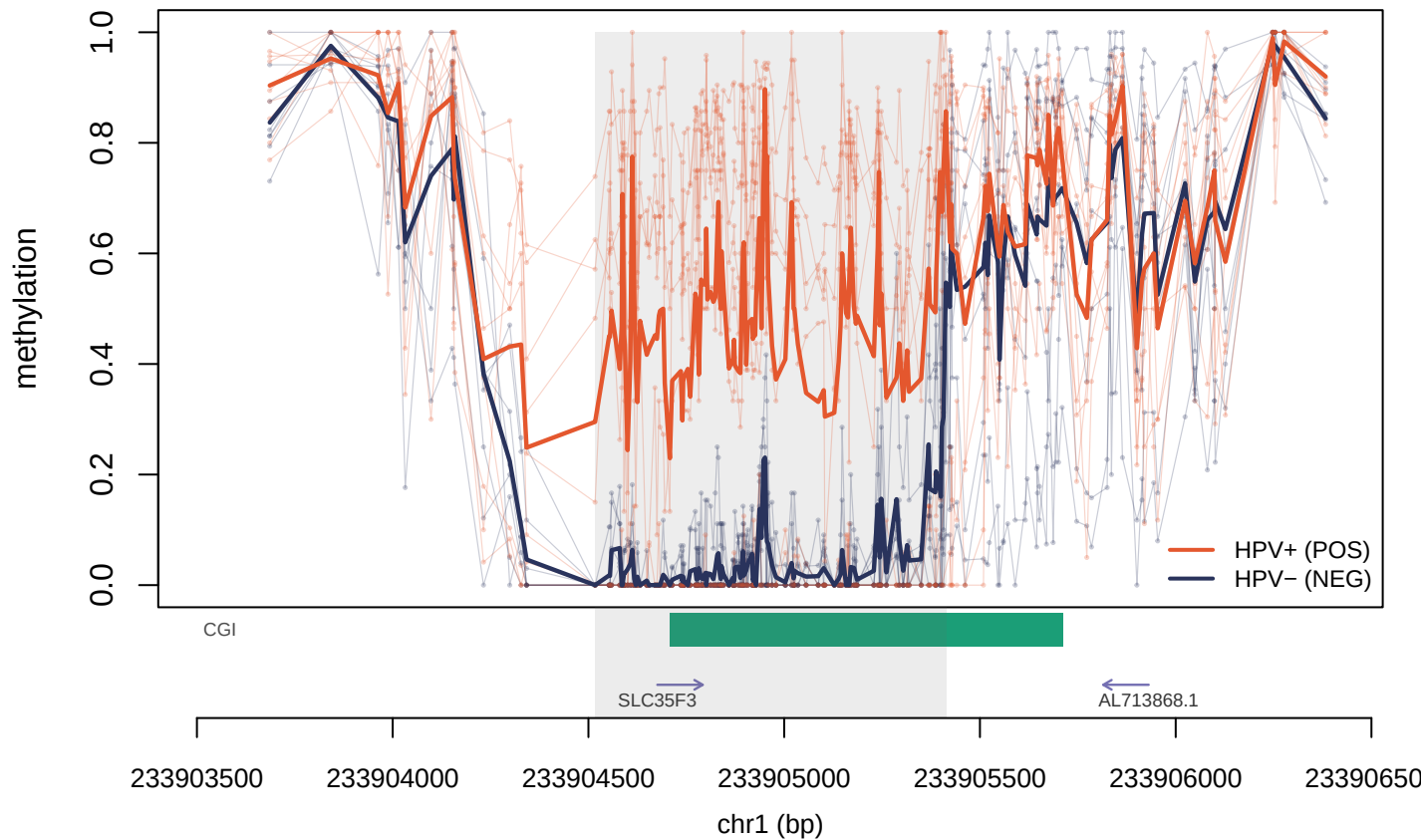
ANXA2R chr5:42,989,405–42,992,879 nCG=133 island / promoter hyper (diff +0.39)



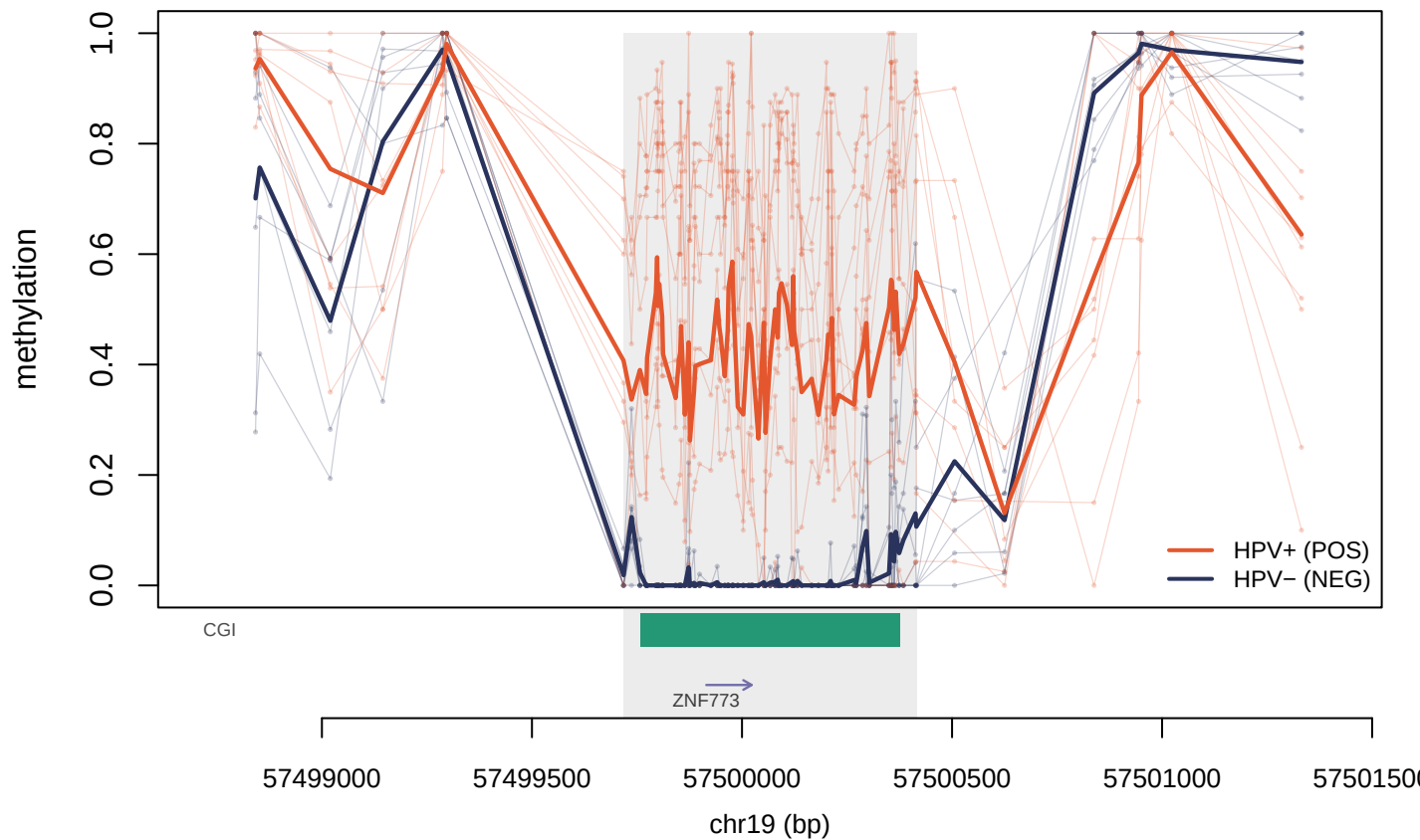
ANXA2R chr5:42,993,517–42,995,760 nCG=103 island / promoter hyper (diff +0.42)



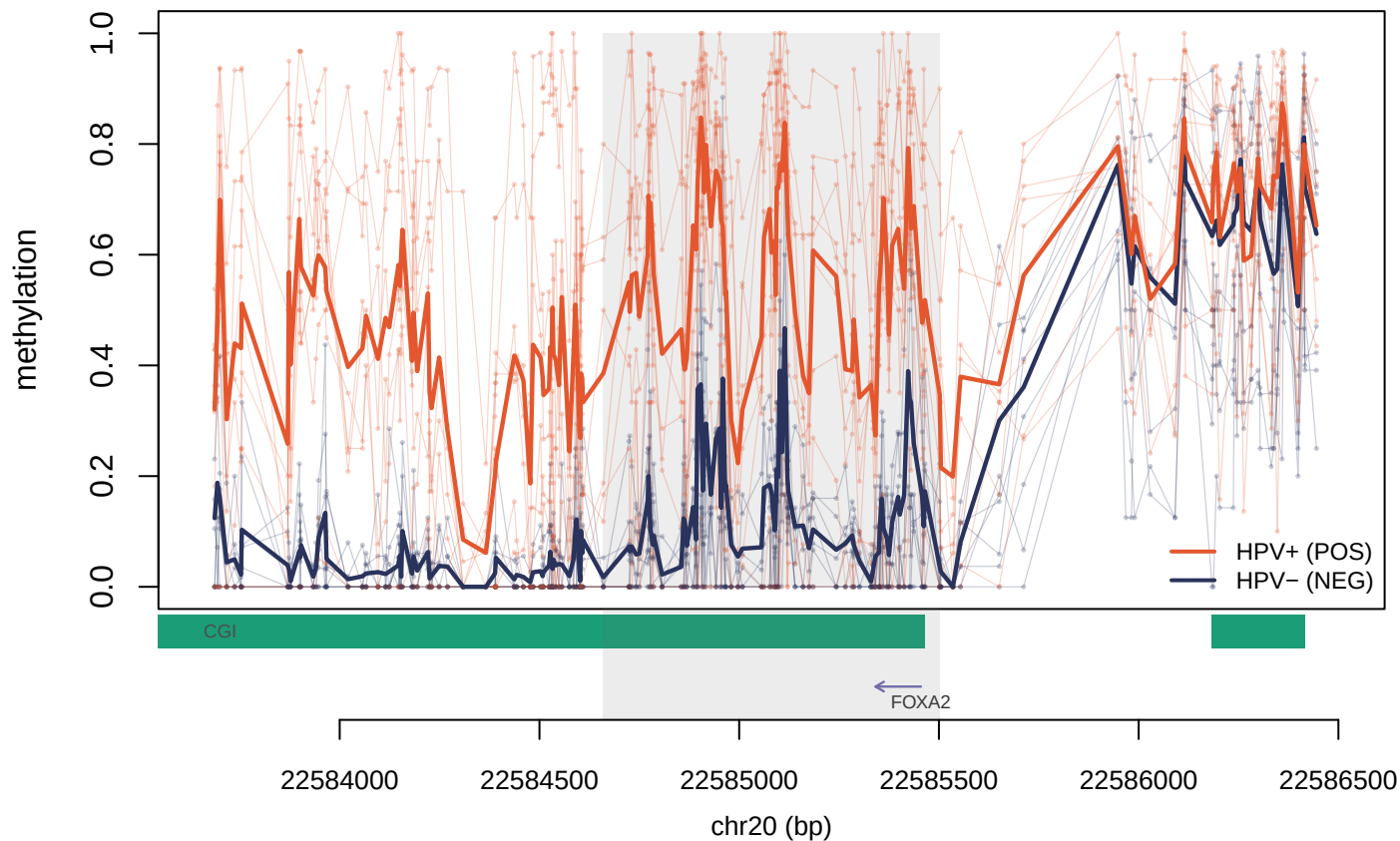
SLC35F3 chr1:233,904,517–233,905,413 nCG=117 island / promoter hyper (diff +0.33)



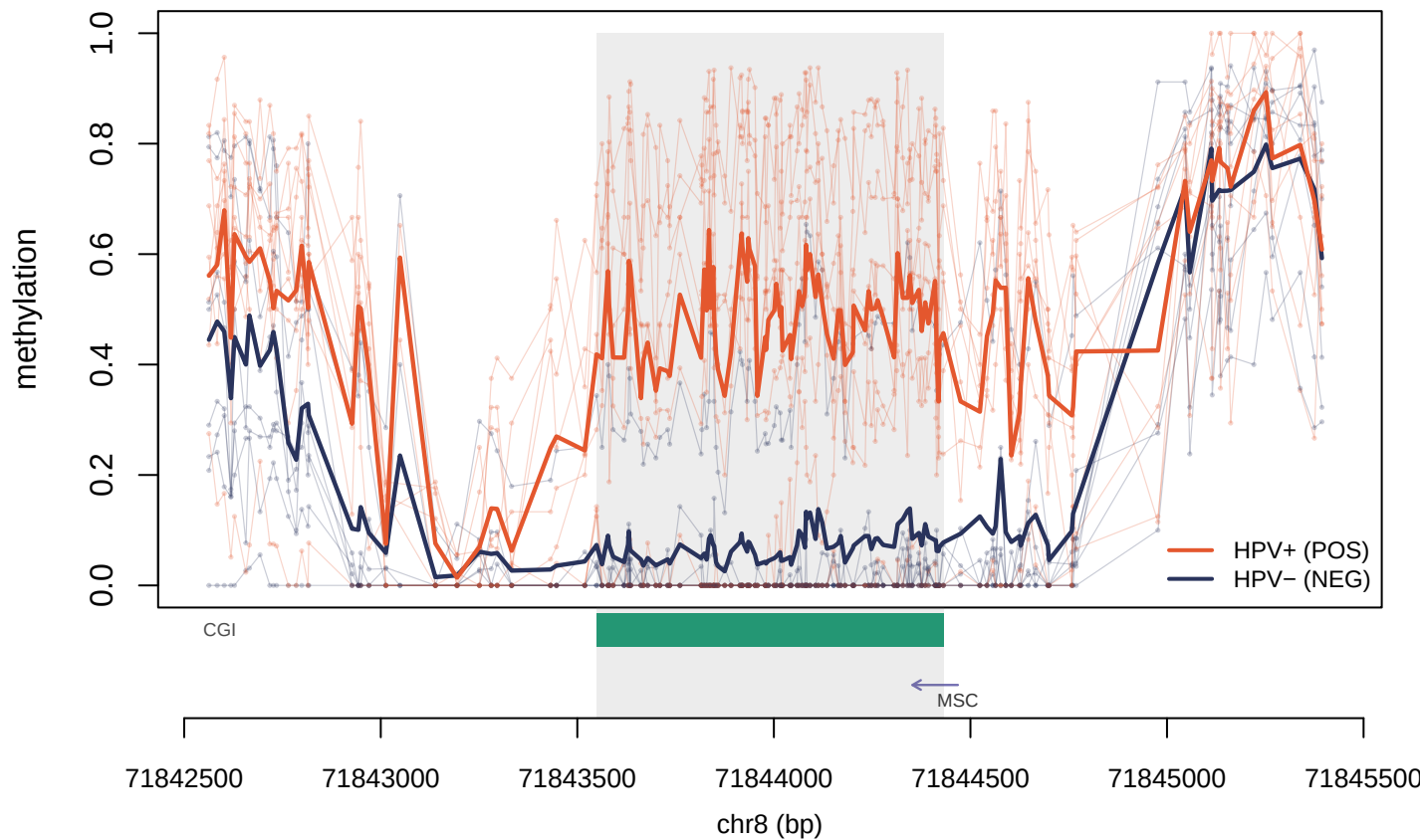
ZNF773 chr19:57,499,718–57,500,415 nCG=71 island / promoter hyper (diff +0.49)



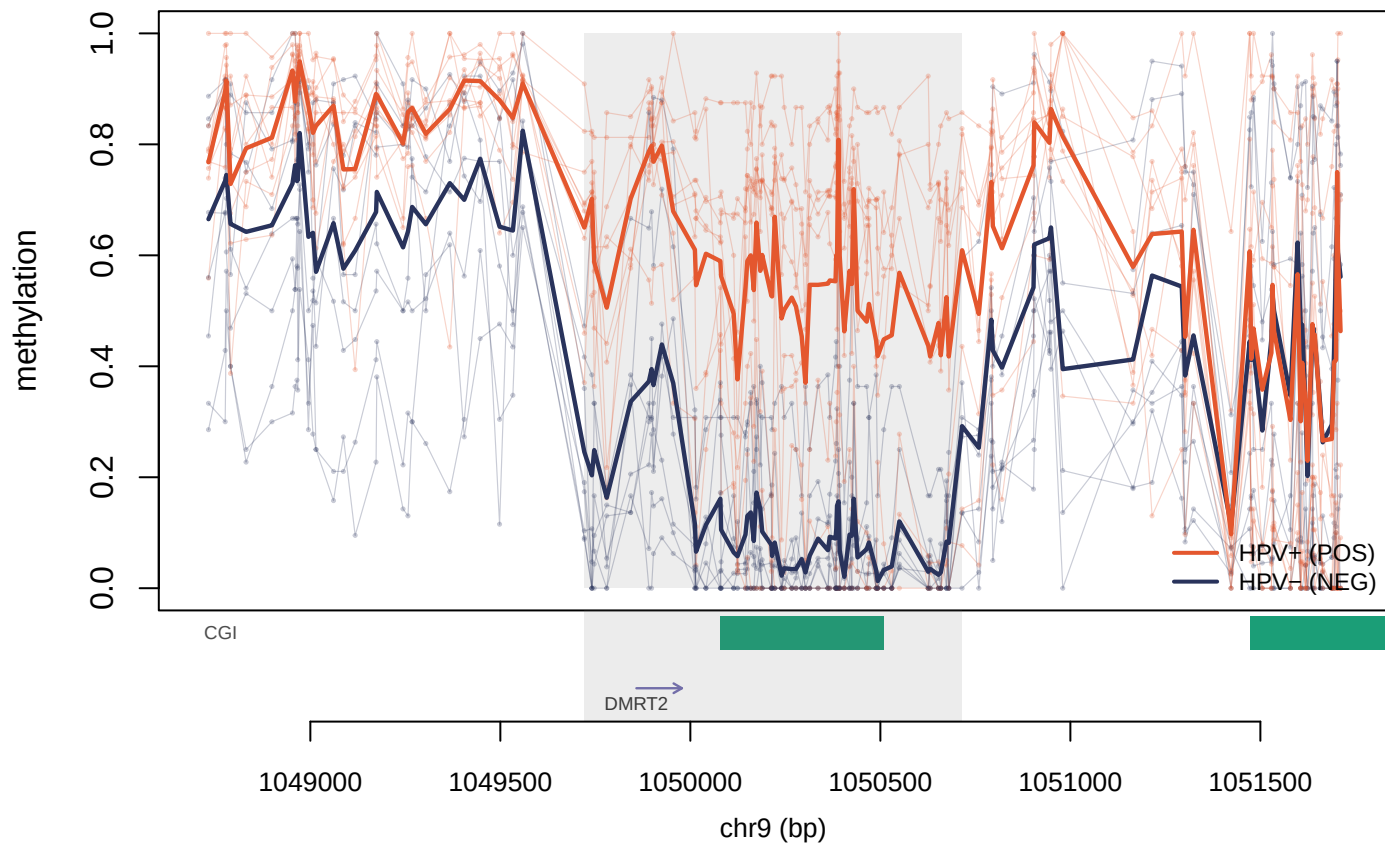
FOXA2 chr20:22,584,660–22,585,502 nCG=78 island / promoter hyper (diff +0.41)



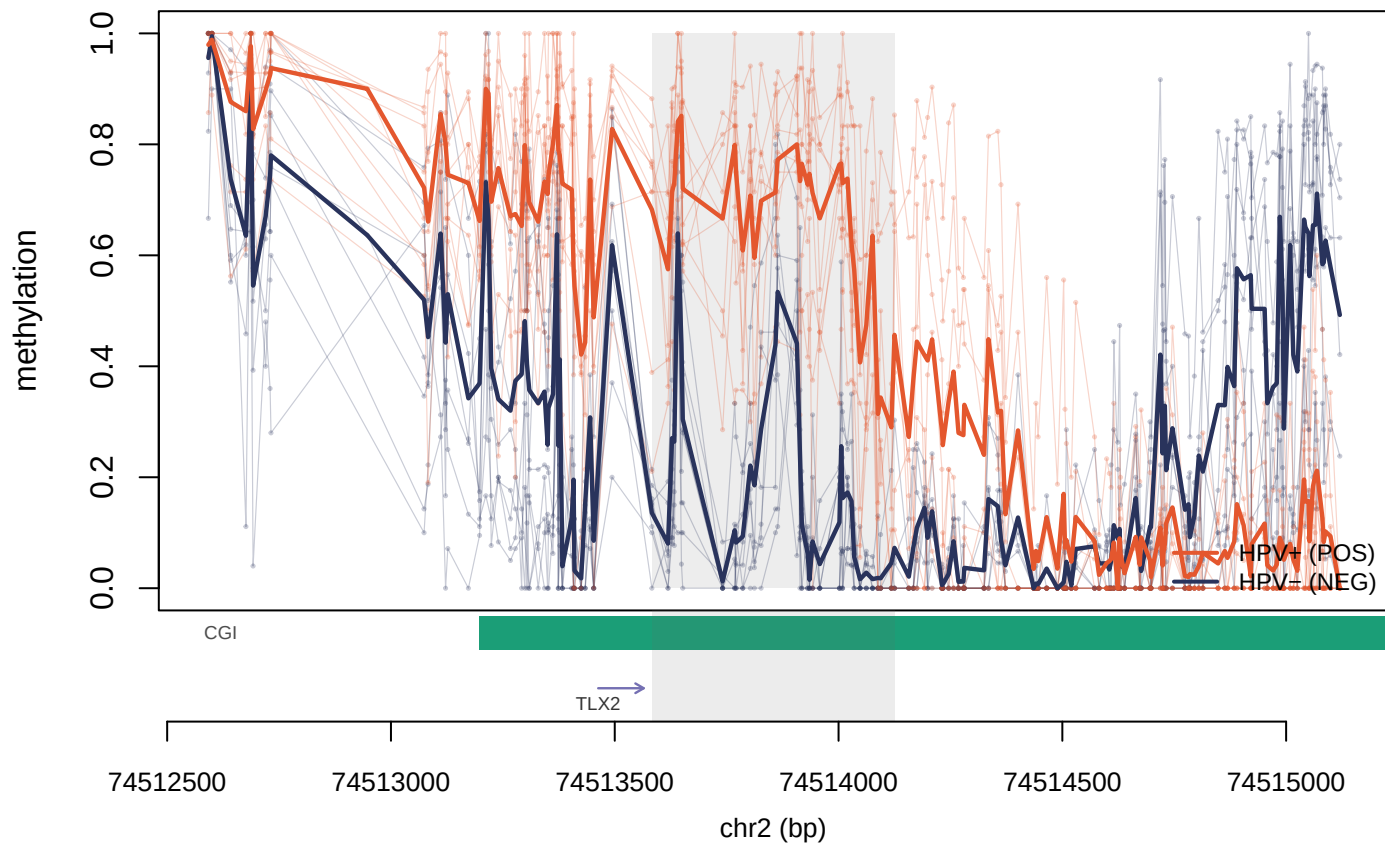
MSC chr8:71,843,549–71,844,431 nCG=88 island / promoter hyper (diff +0.38)



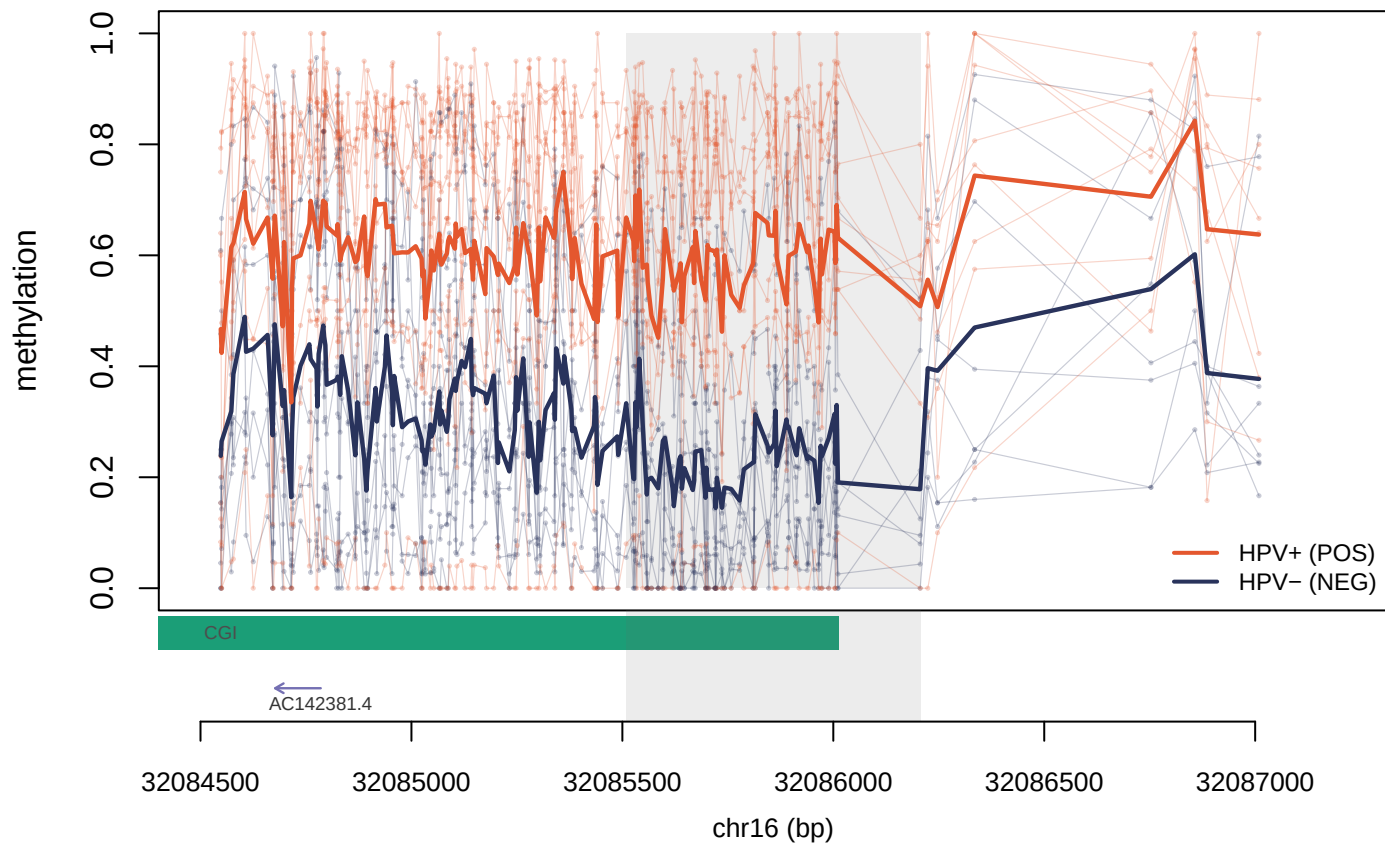
DMRT2 chr9:1,049,721–1,050,715 nCG=61 island / promoter hyper (diff +0.42)



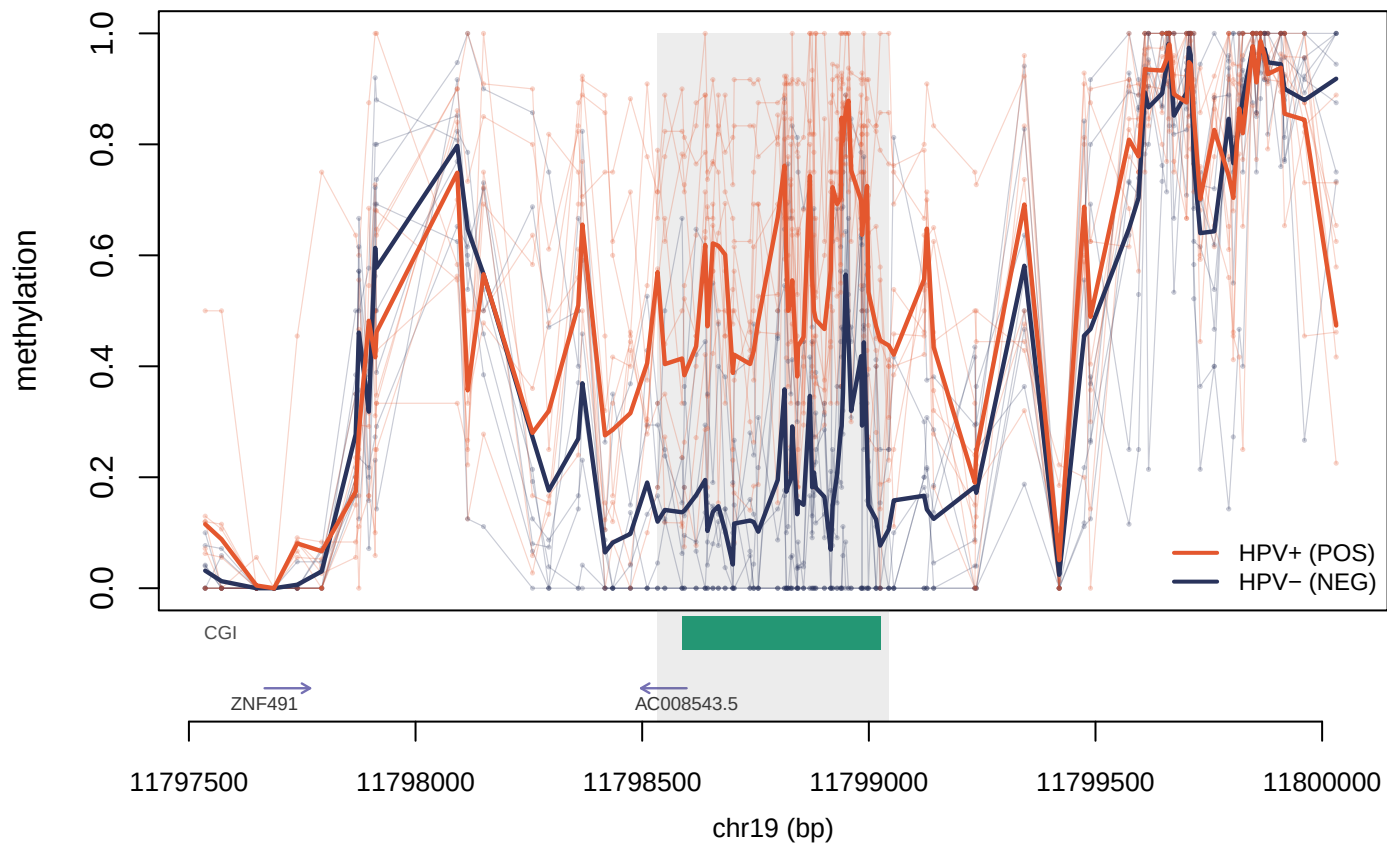
TLX2 chr2:74,513,583-74,514,125 nCG=36 island / promoter hyper (diff +0.47)



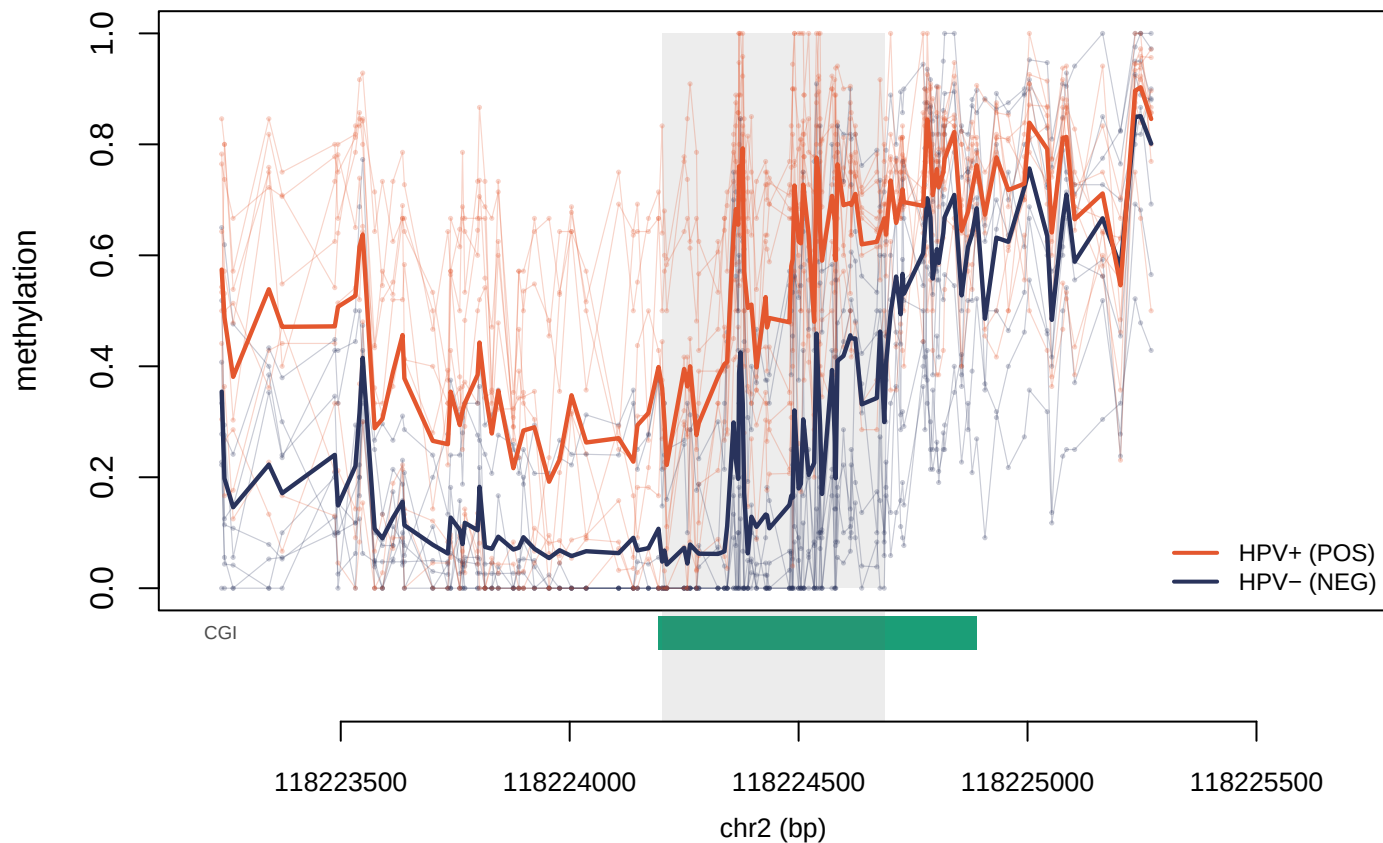
TP53TG3D chr16:32,085,509–32,086,206 nCG=61 island / promoter hyper (diff +0.37)



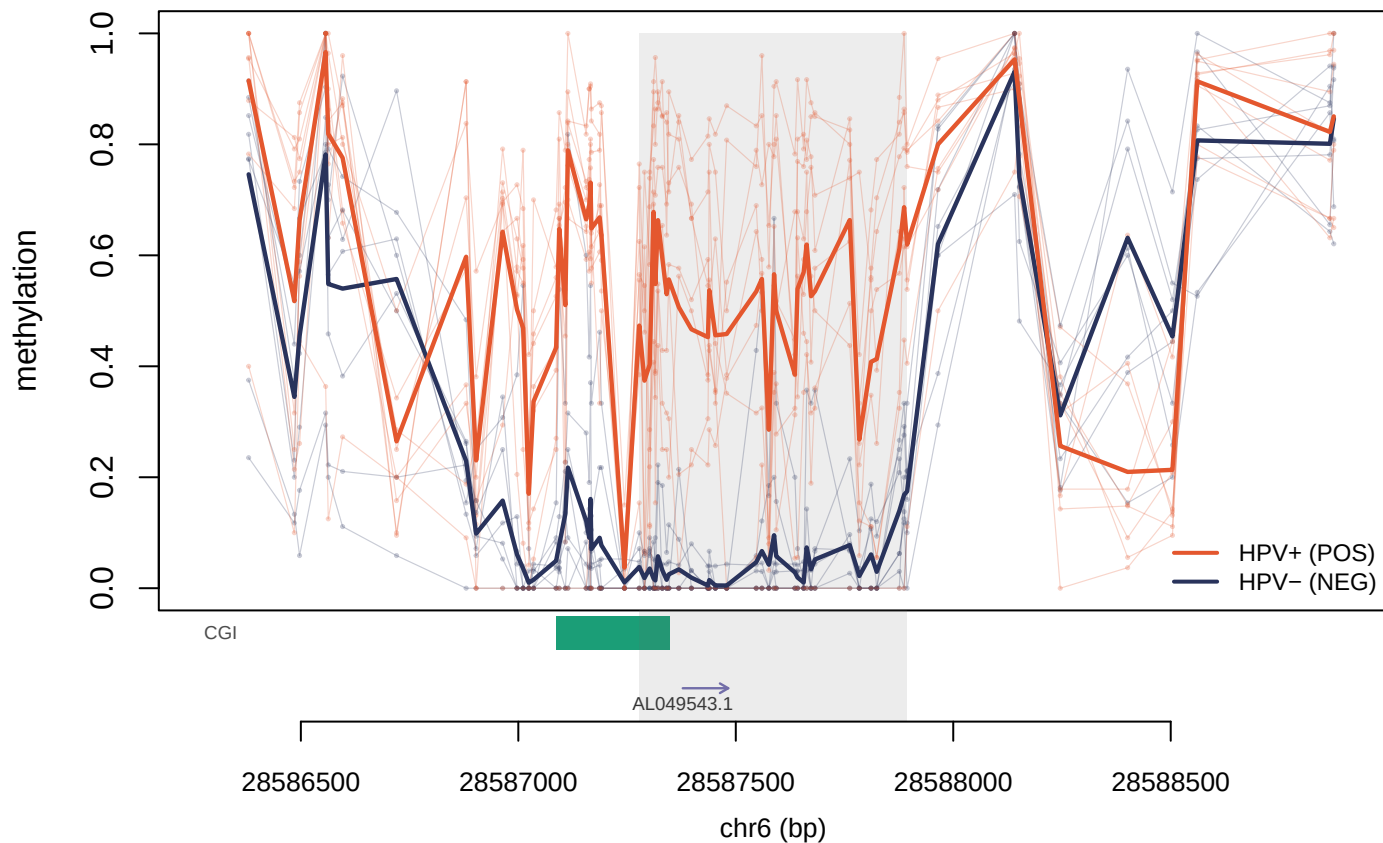
ZNF491 chr19:11,798,534–11,799,044 nCG=46 island / promoter hyper (diff +0.45)



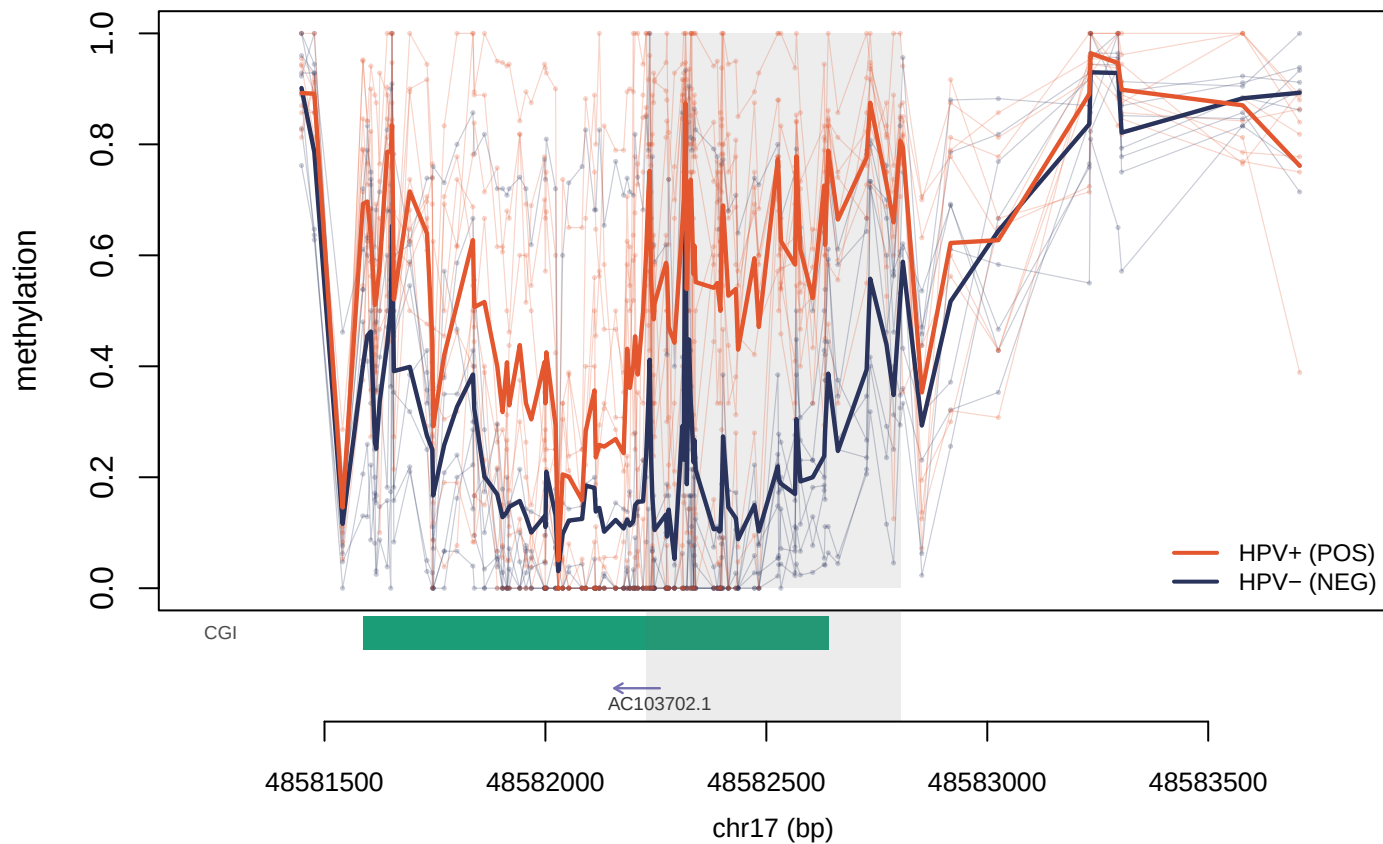
INSIG2 chr2:118,224,202–118,224,687 nCG=50 island / promoter hyper (diff +0.37)



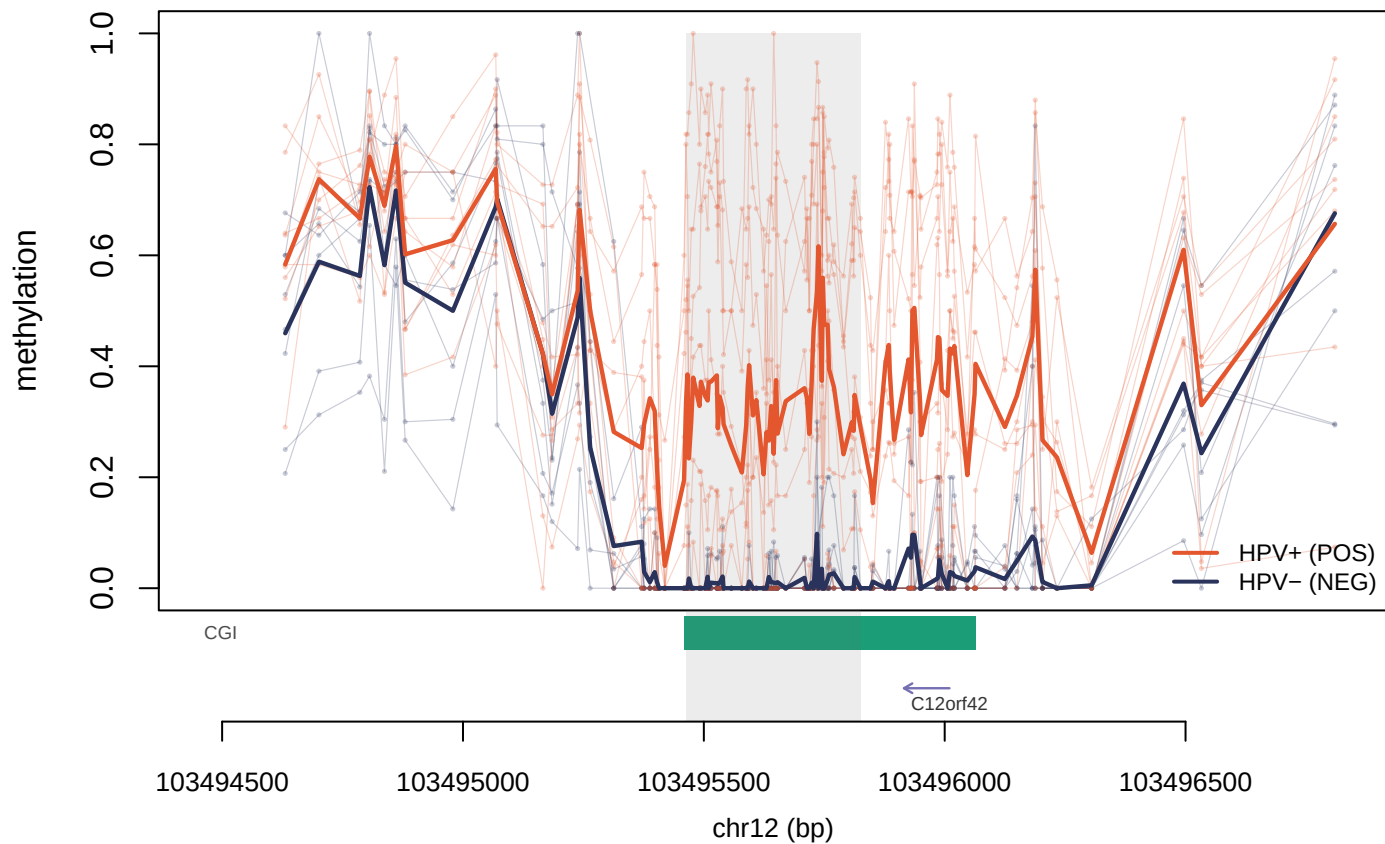
ZBED9 chr6:28,587,278–28,587,894 nCG=33 island / promoter hyper (diff +0.45)



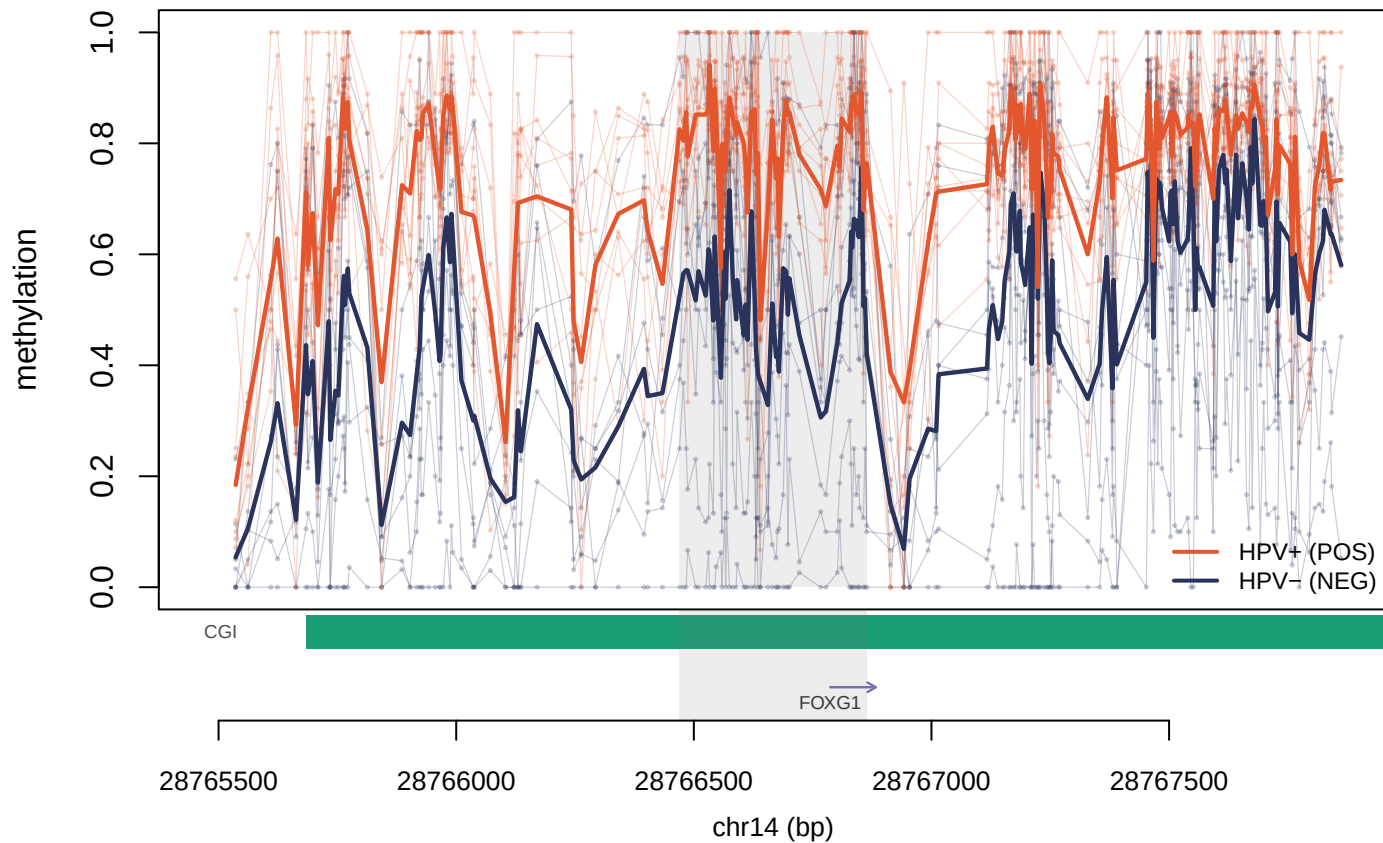
HOXB4 chr17:48,582,228–48,582,803 nCG=48 island / promoter hyper (diff +0.38)



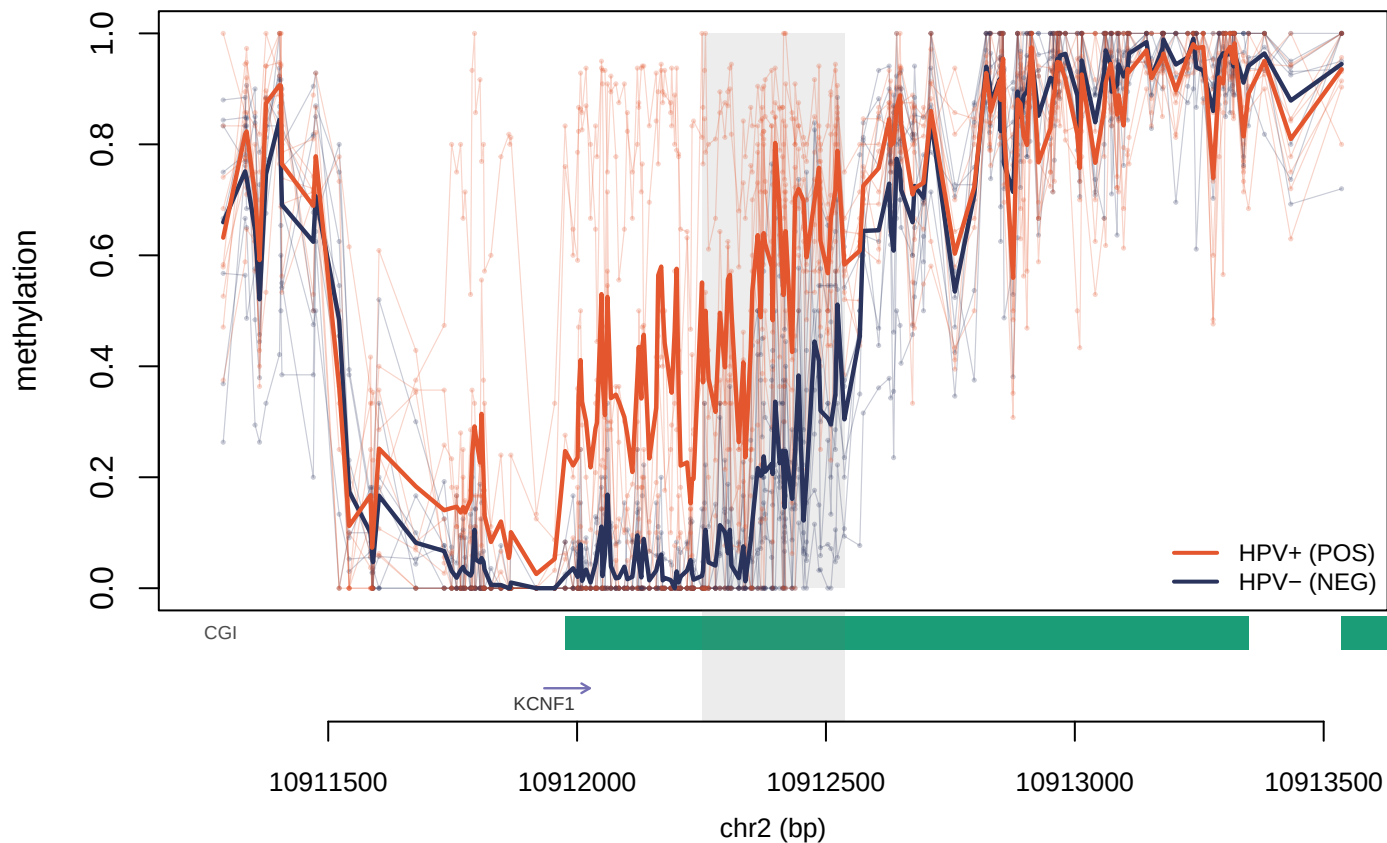
C12orf42 chr12:103,495,463–103,495,825 nCG=49 island / promoter hyper (diff +0.29)



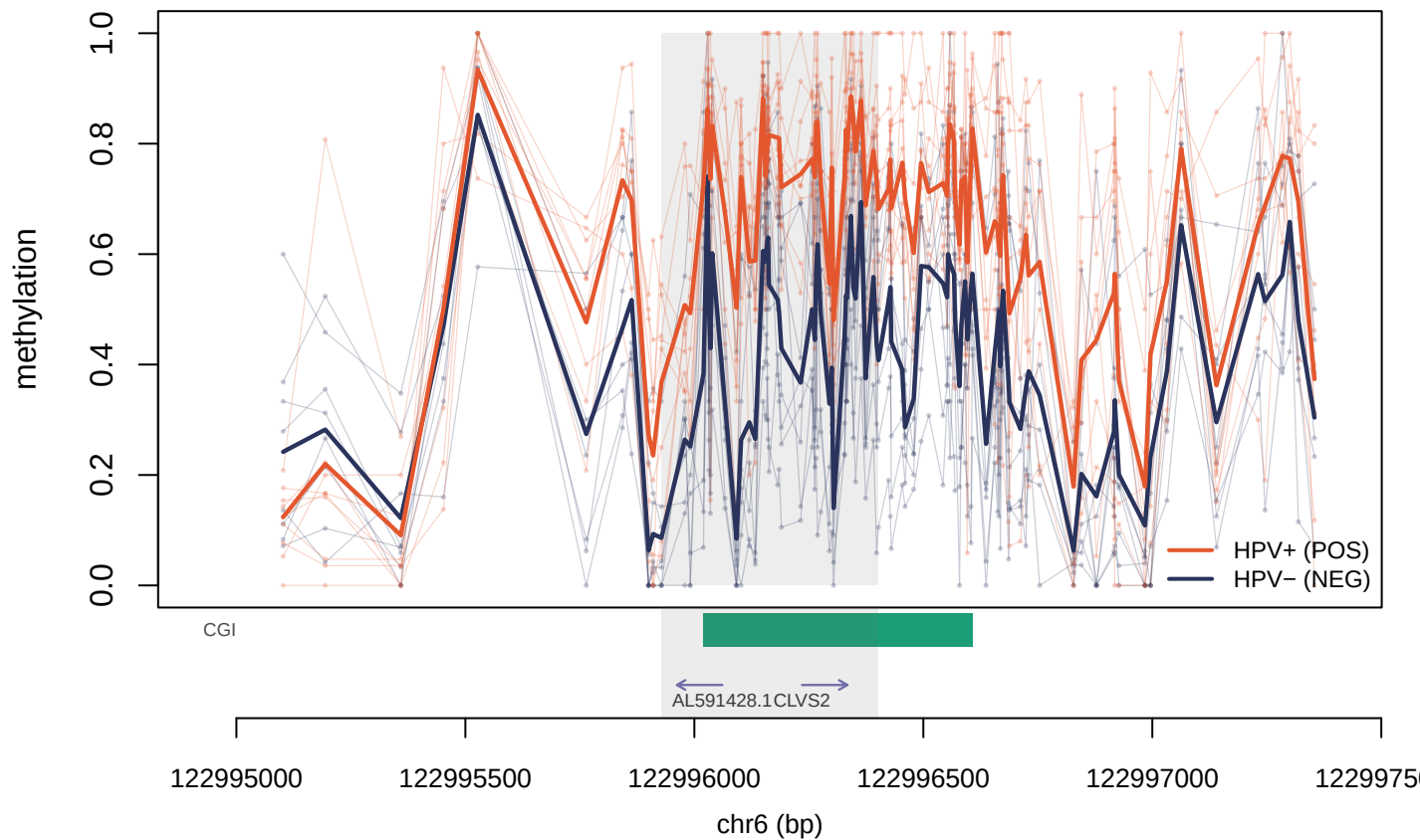
FOXG1 chr14:28,766,470–28,766,864 nCG=55 island / promoter hyper (diff +0.31)



KCNF1 chr2:10,912,251–10,912,537 nCG=42 island / promoter hyper (diff +0.32)



CLVS2 chr6:122,995,928–122,996,402 nCG=42 island / promoter hyper (diff +0.28)



ZNF69 chr19:11,887,962–11,888,510 nCG=42 island / promoter hyper (diff +0.37)

