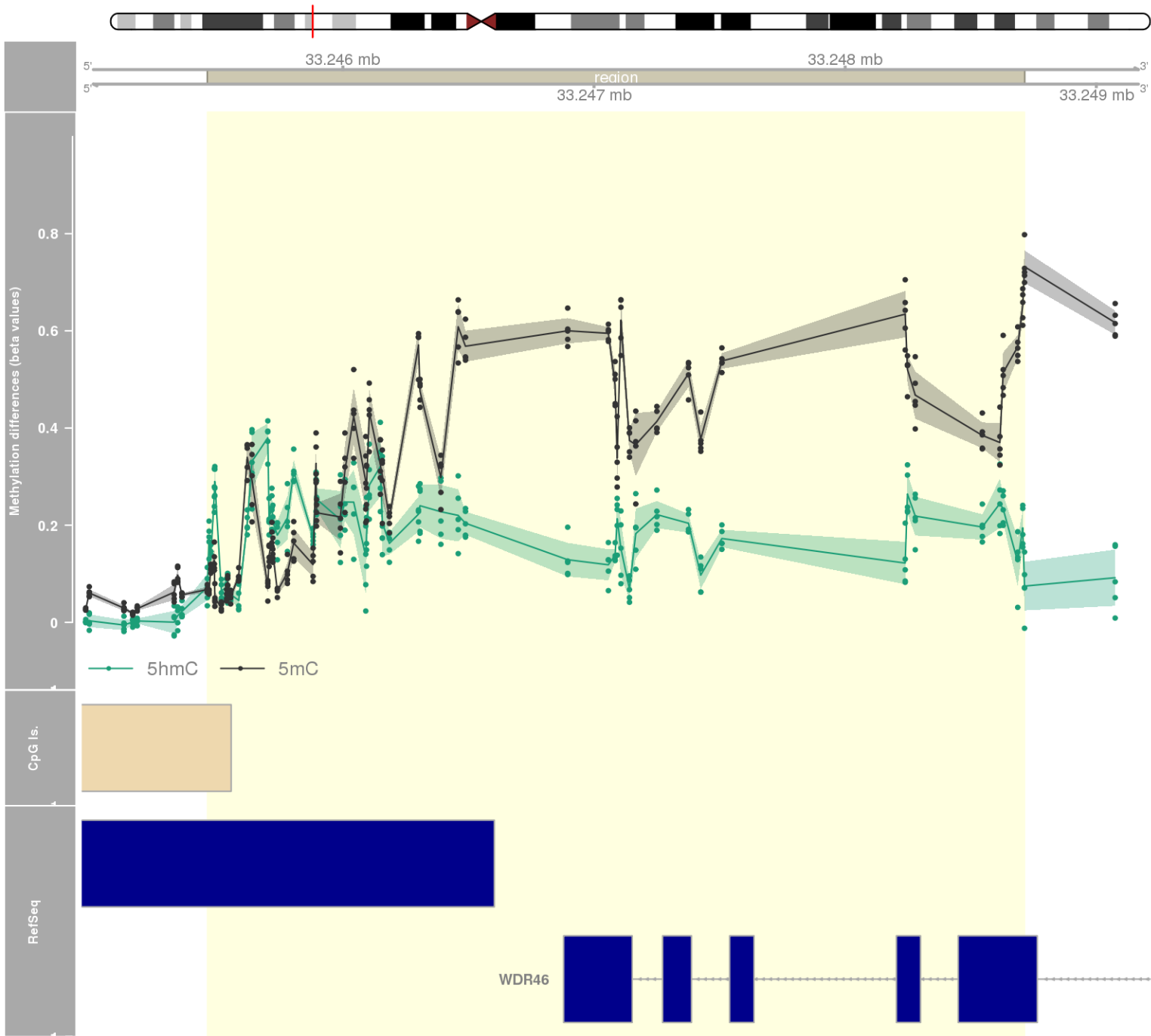
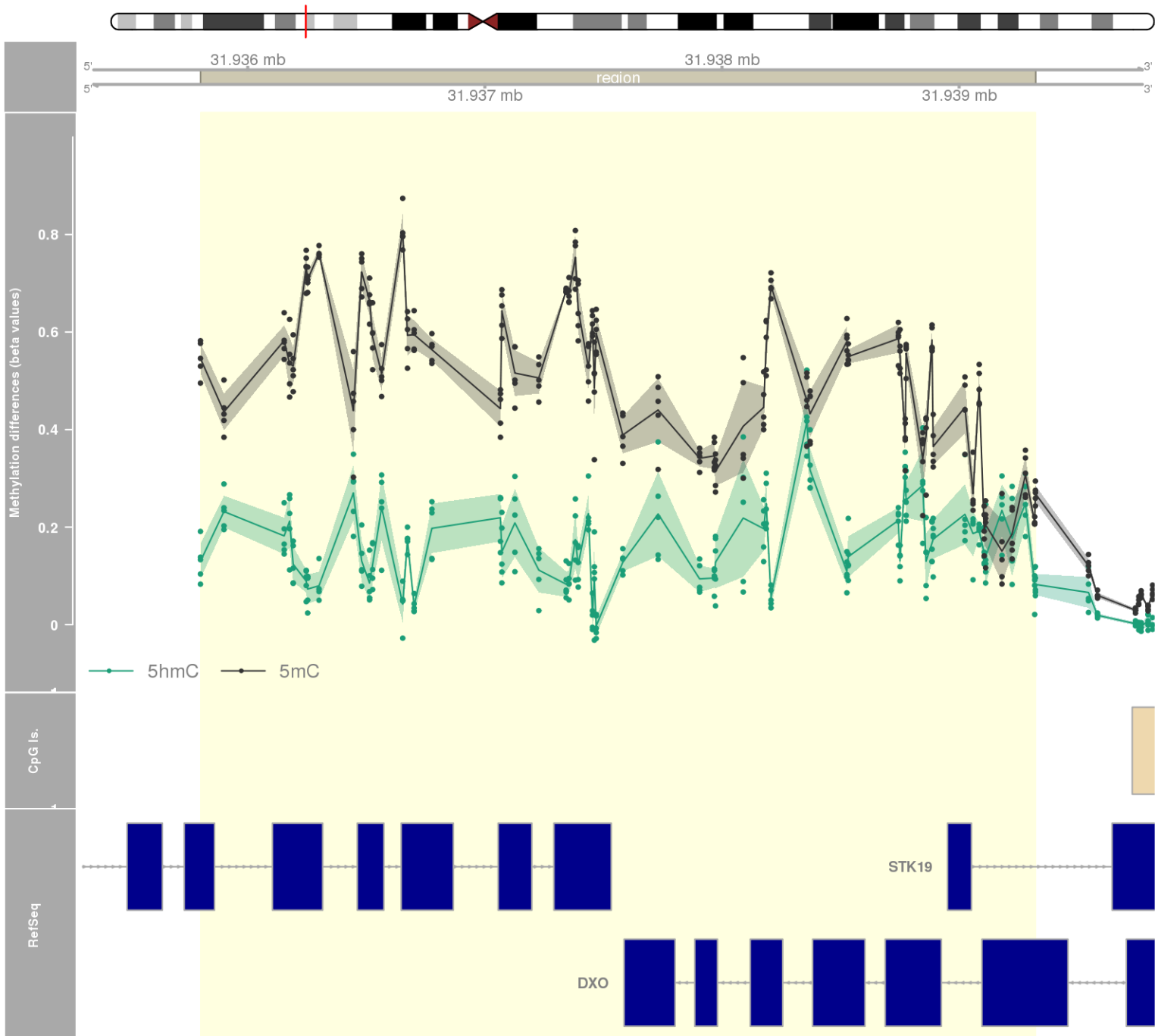


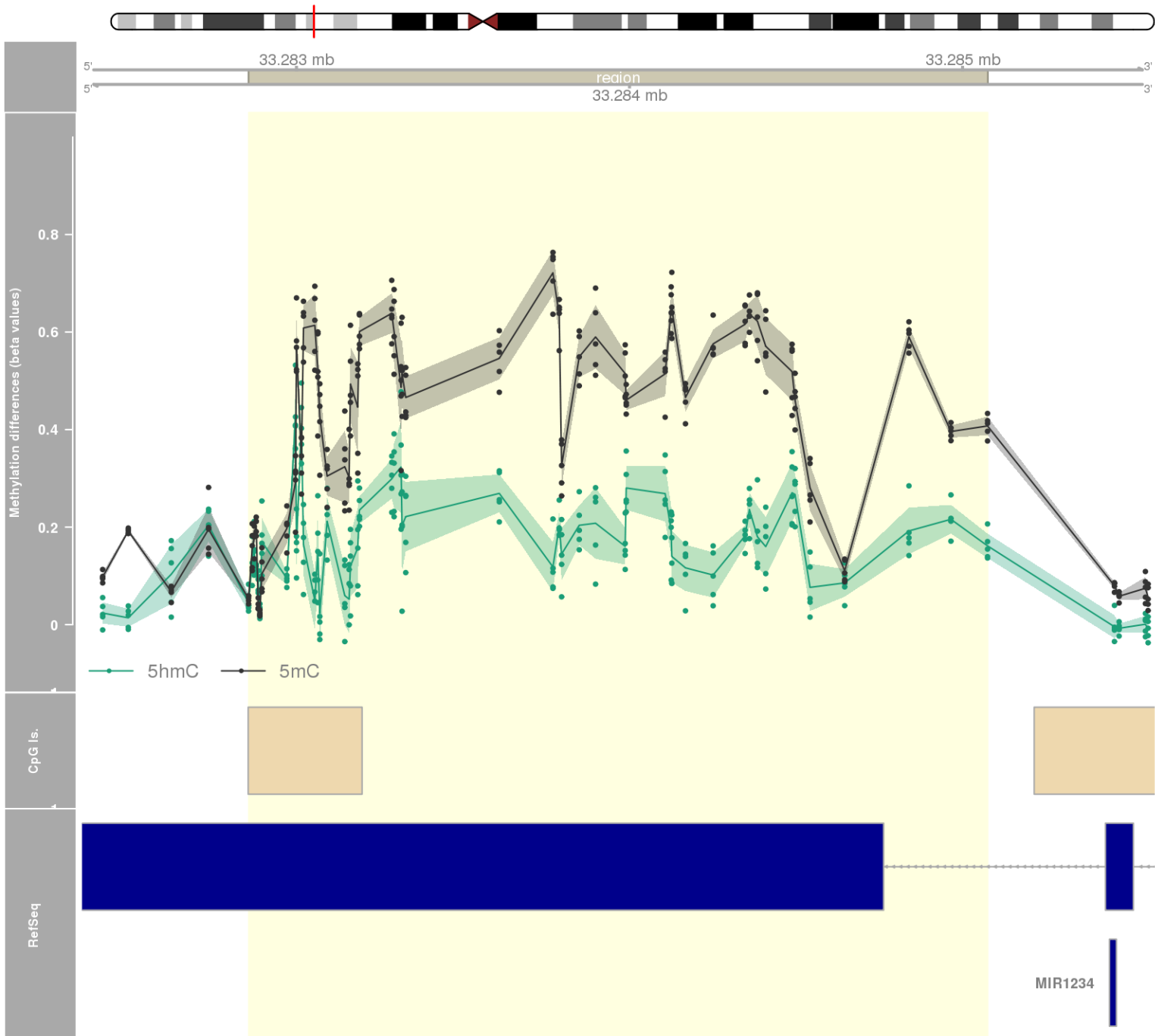
DMR 1 // chr6:33245460-33248714 // 3254 pb. (56 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0 // fwerArea: 0.173
- genes: B3GALT4 / WDR46 -



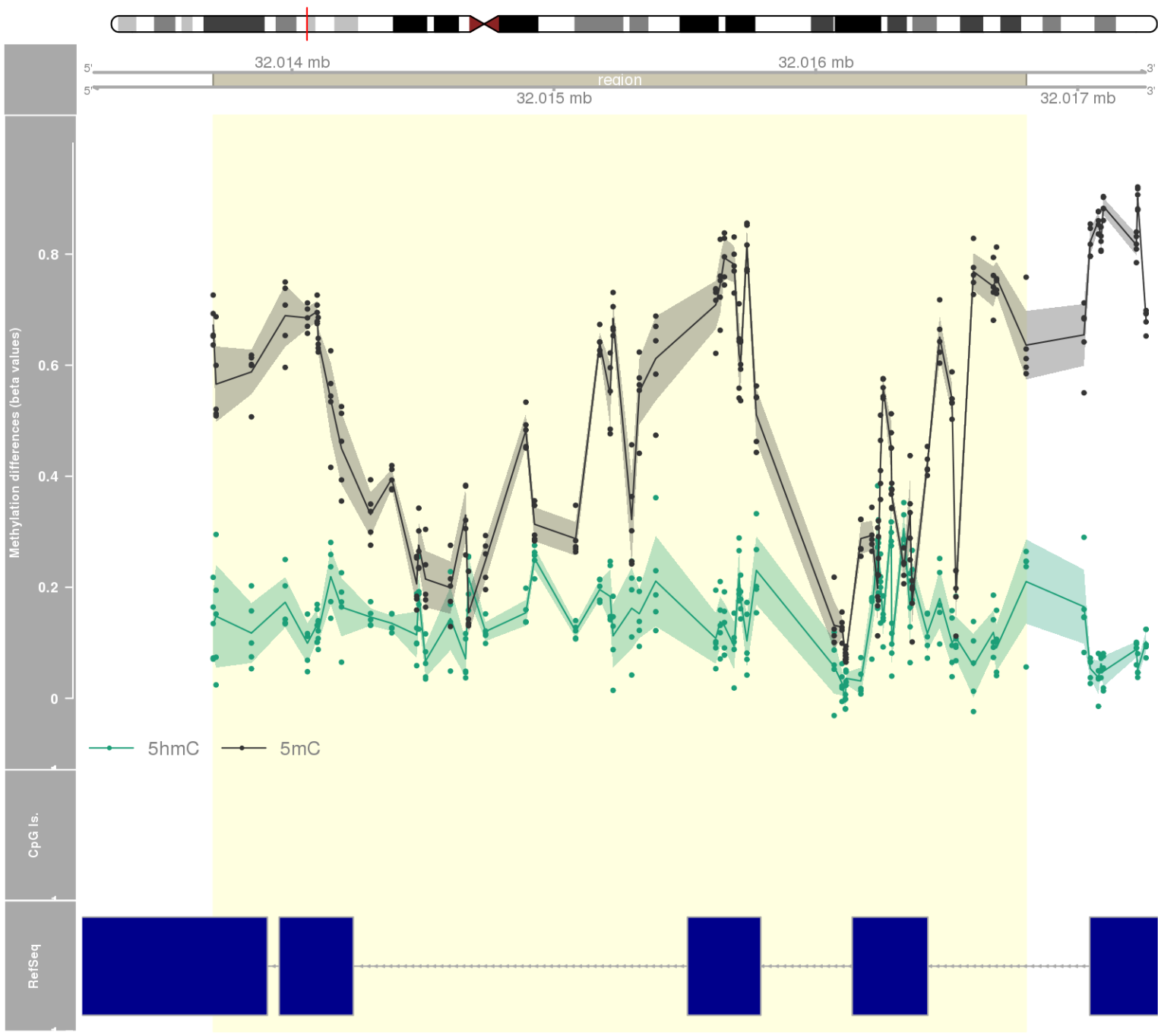
DMR 2 // chr6:31935801-31939322 // 3521 pb. (60 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0 // fwerArea: 0.173
- genes: SKIV2L / DXO / STK19 -



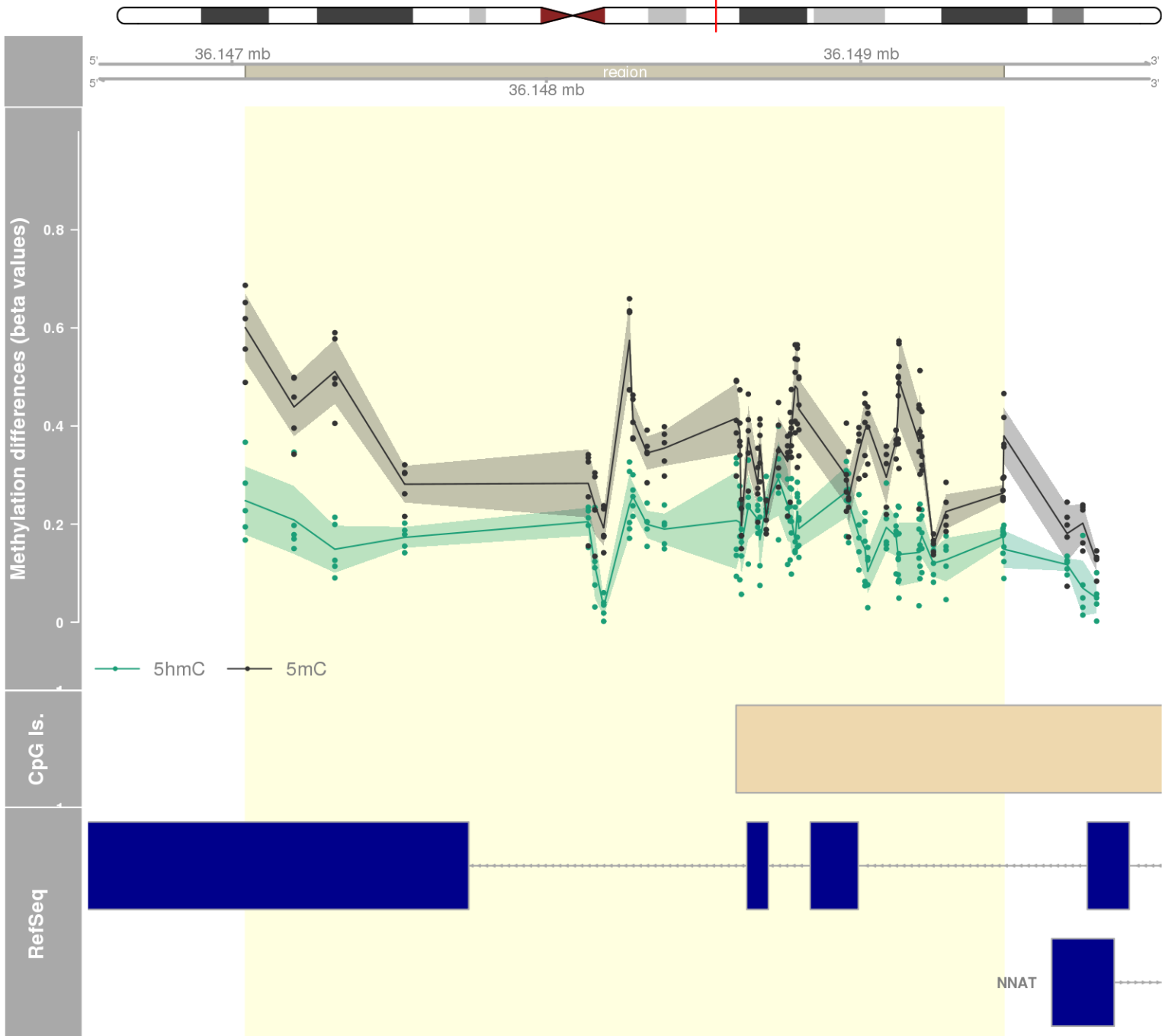
DMR 3 // chr6:33282856-33285075 // 2219 pb. (50 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0 // fwerArea: 0.173
- genes: ZBTB22 -



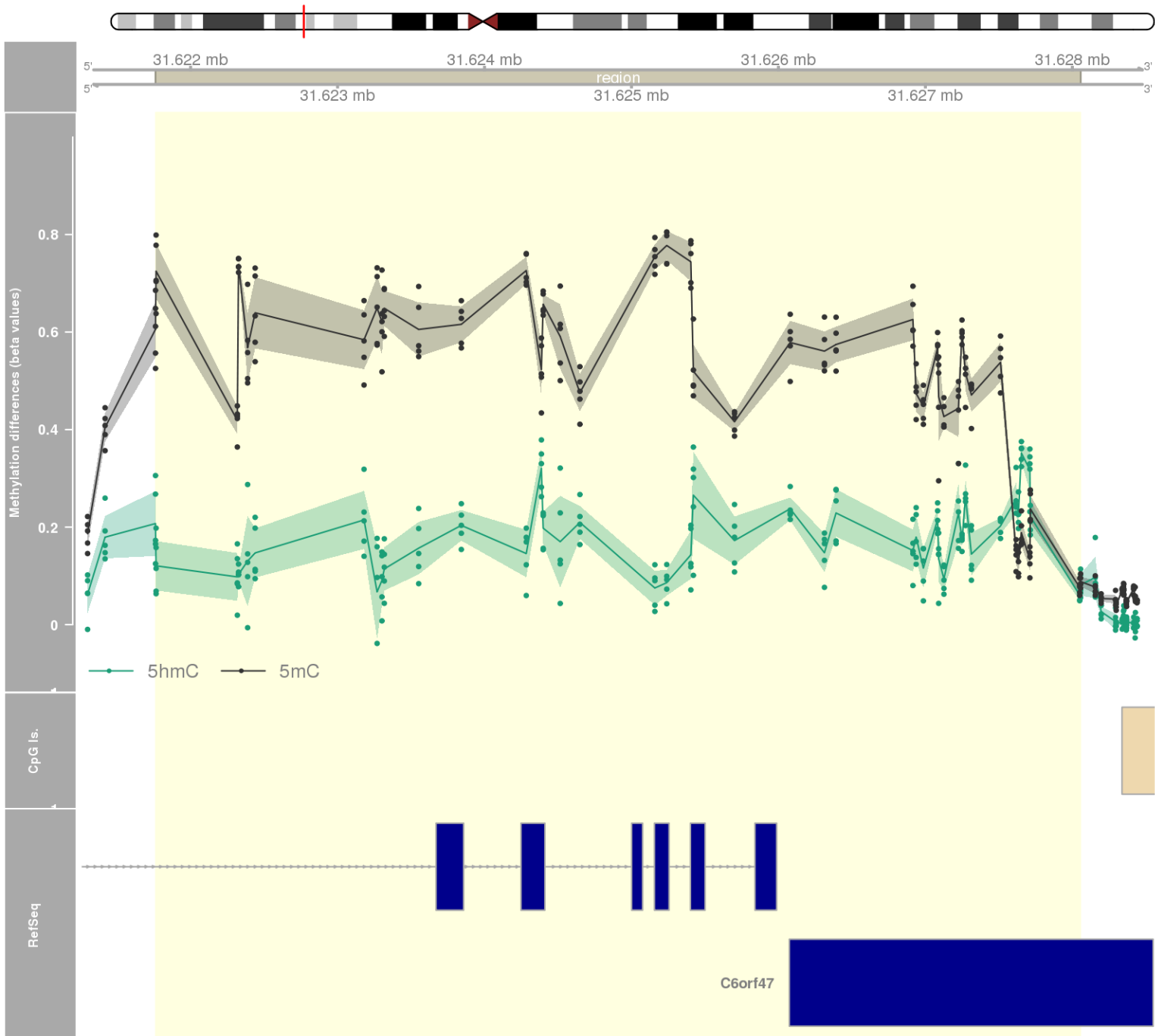
DMR 4 // chr6:32013699-32016803 // 3104 pb. (58 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0 // fwerArea: 0.173
- genes: TNXB -



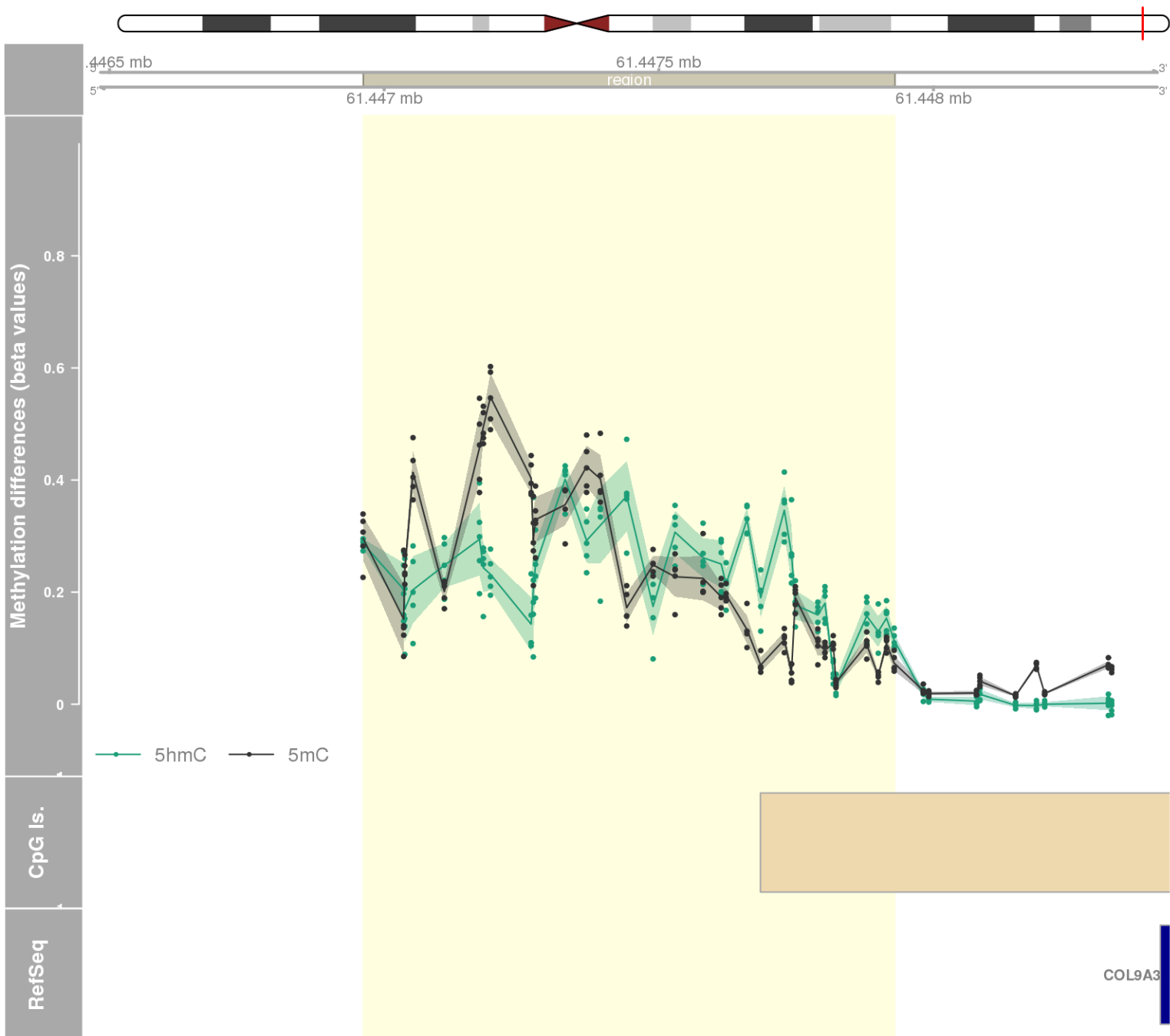
DMR 5 // chr20:36147042-36149455 // 2413 pb. (41 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0 // fwerArea: 0.173
- genes: BLCAP -



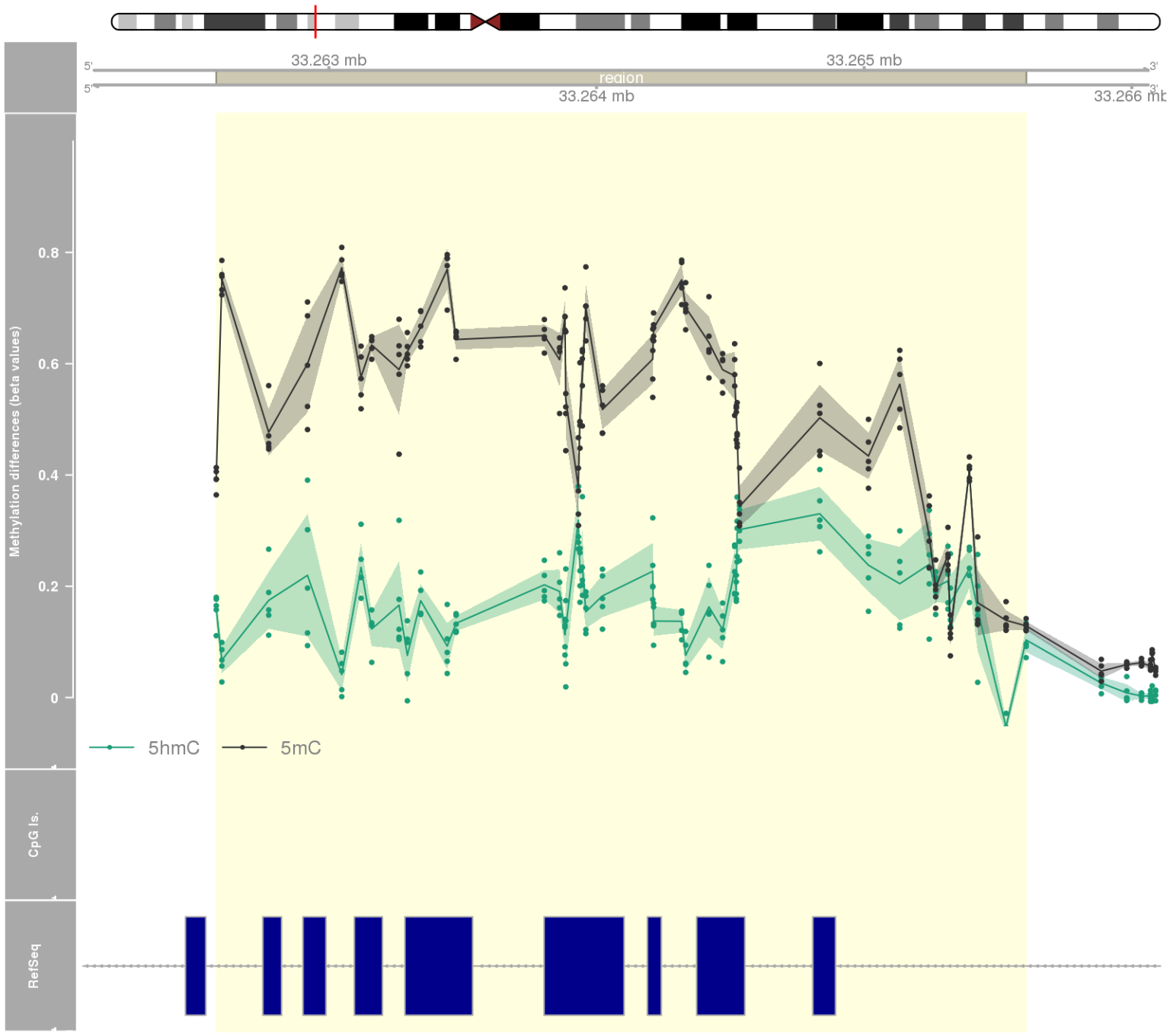
DMR 6 // chr6:31621761-31628056 // 6295 pb. (43 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0 // fwerArea: 0.173
- genes: C6orf47-AS1 / APOM / C6orf47 -



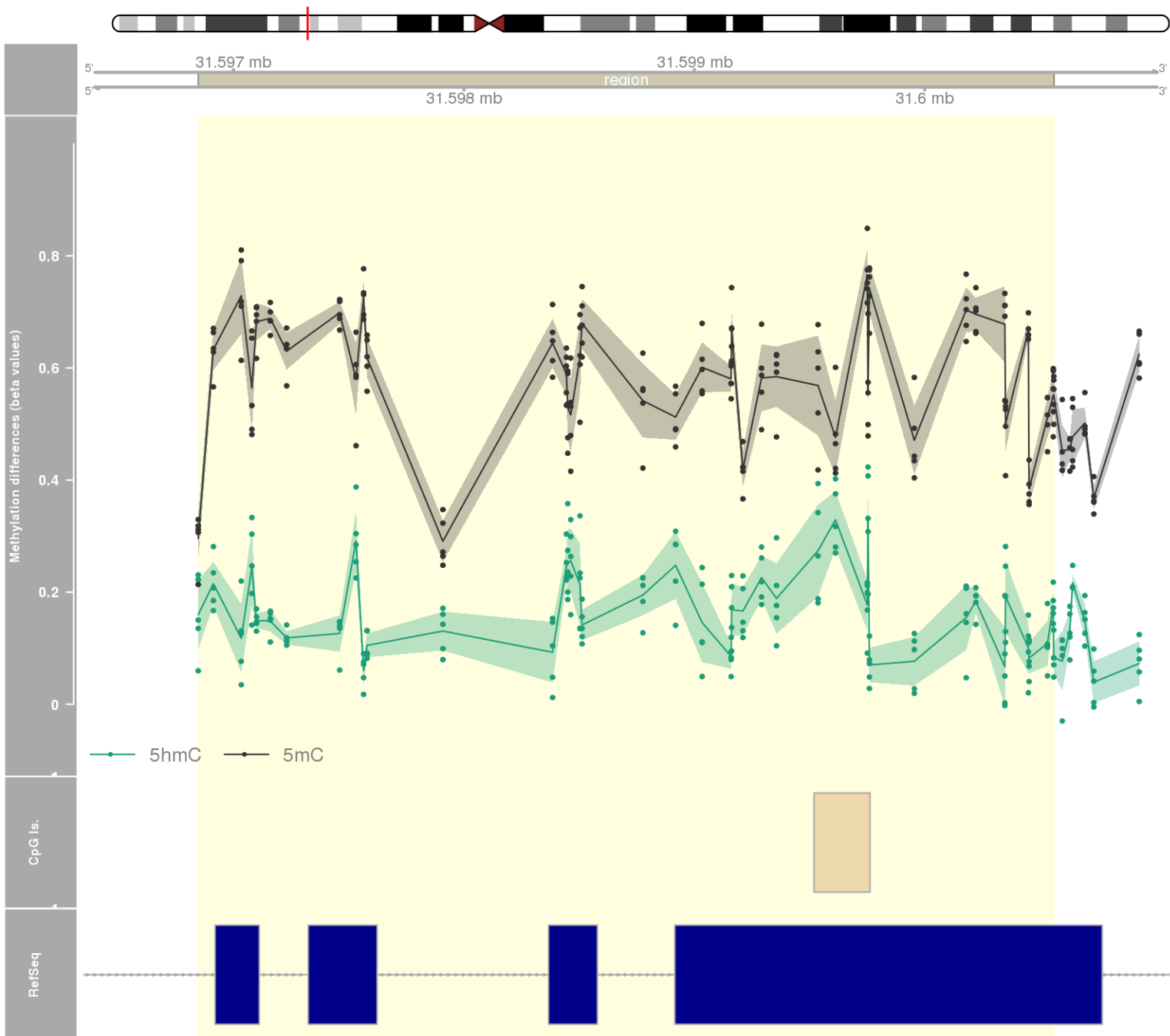
DMR 7 // chr20:61446962-61447929 // 967 pb. (33 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0 // fwerArea: 0.173
- genes: COL9A3 -



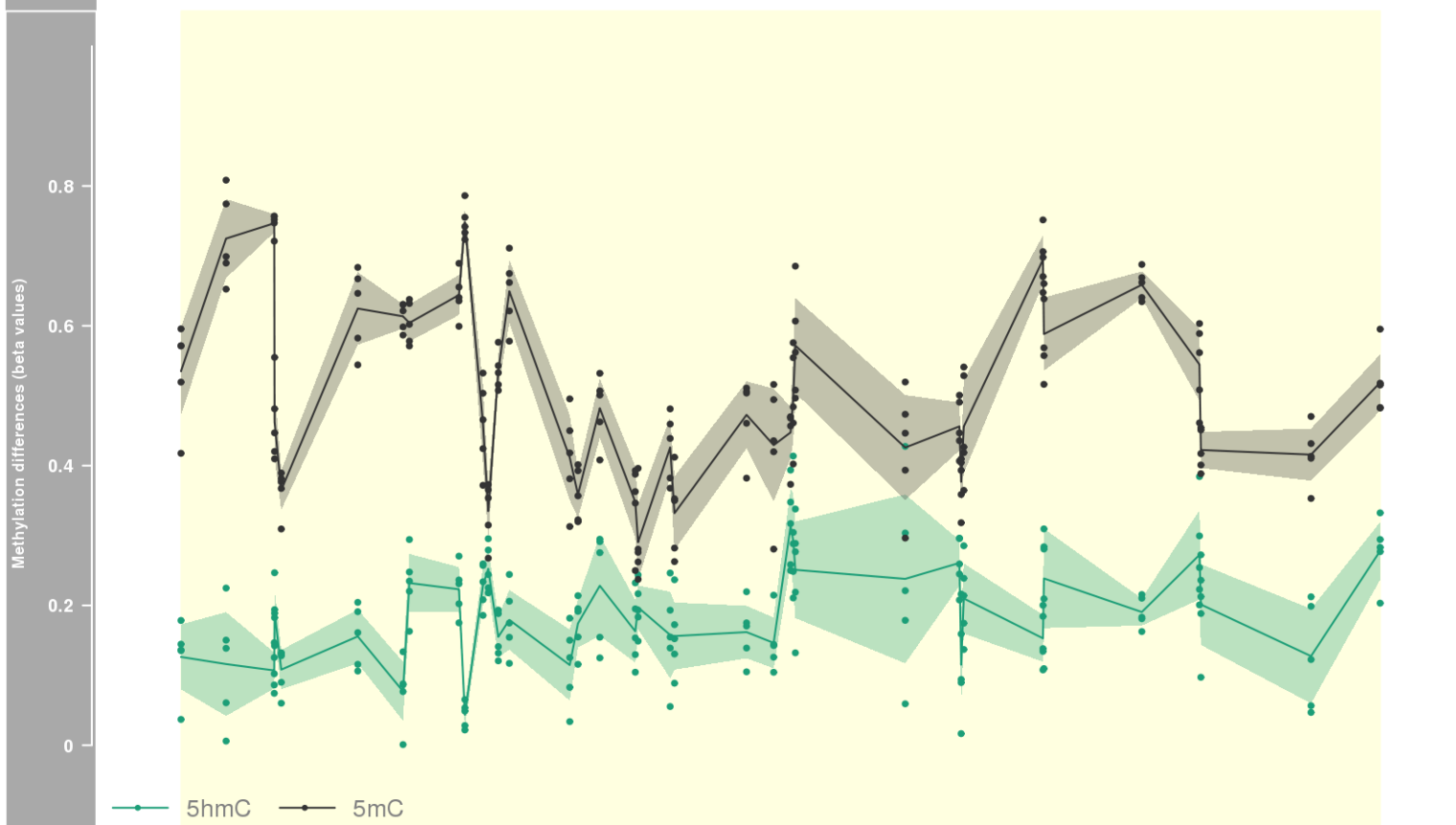
DMR 8 // chr6:33262579-33265605 // 3026 pb. (42 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: PFDN6 / RGL2 -



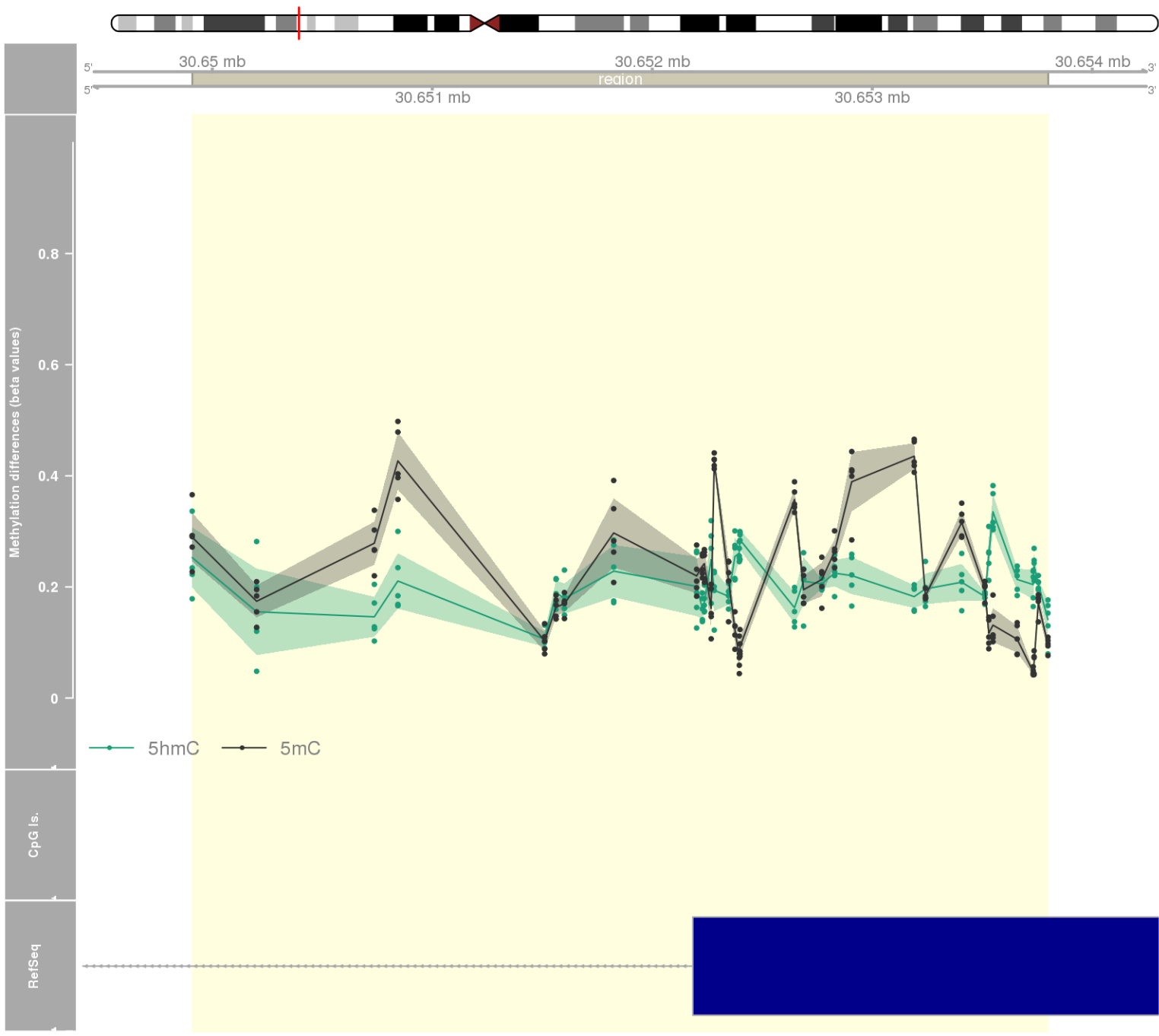
DMR 9 // chr6:31596847-31600560 // 3713 pb. (41 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: PRRC2A -



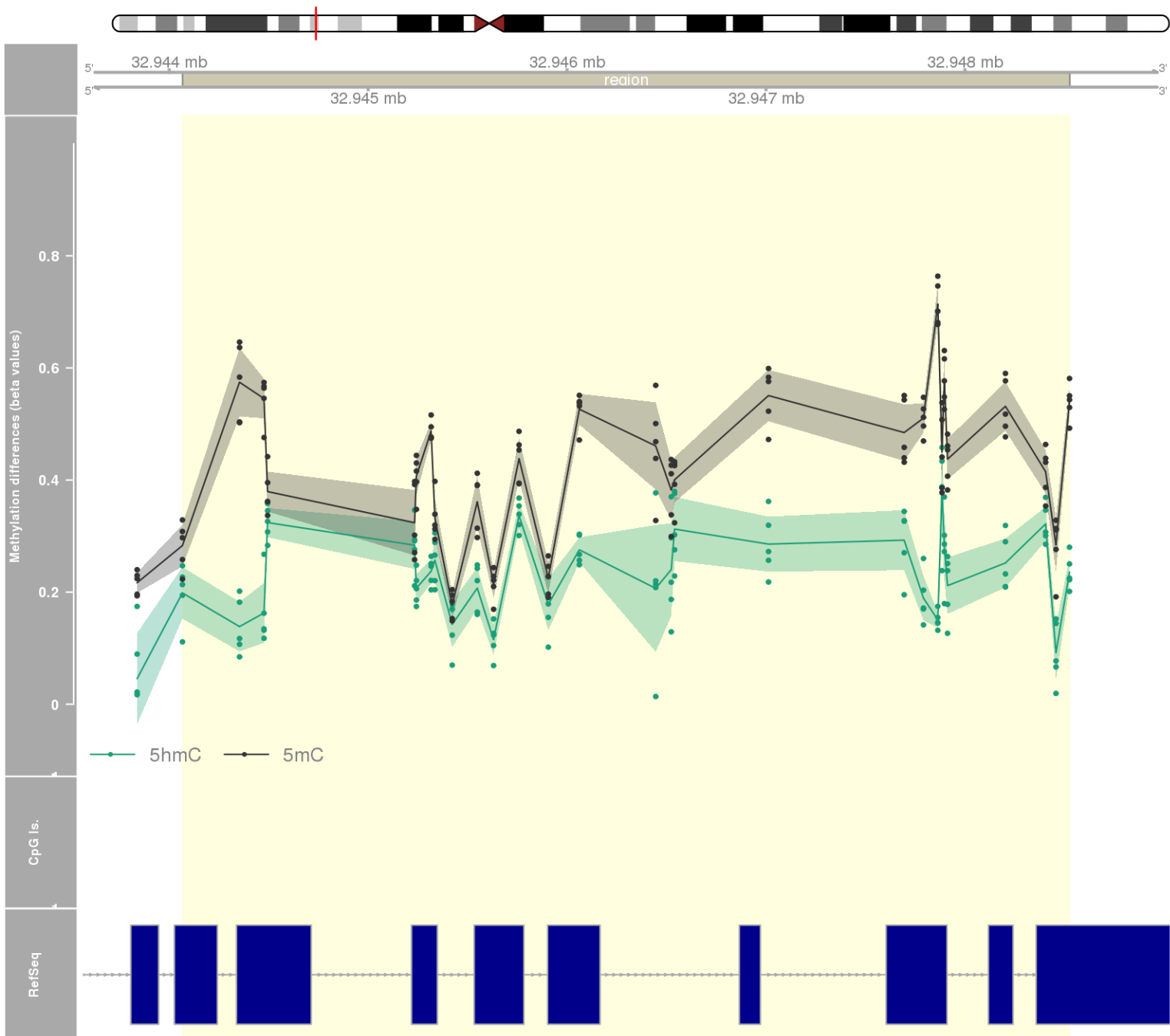
DMR 10 // chr6:30172142-30179881 // 7739 pb. (37 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: TRIM26 -



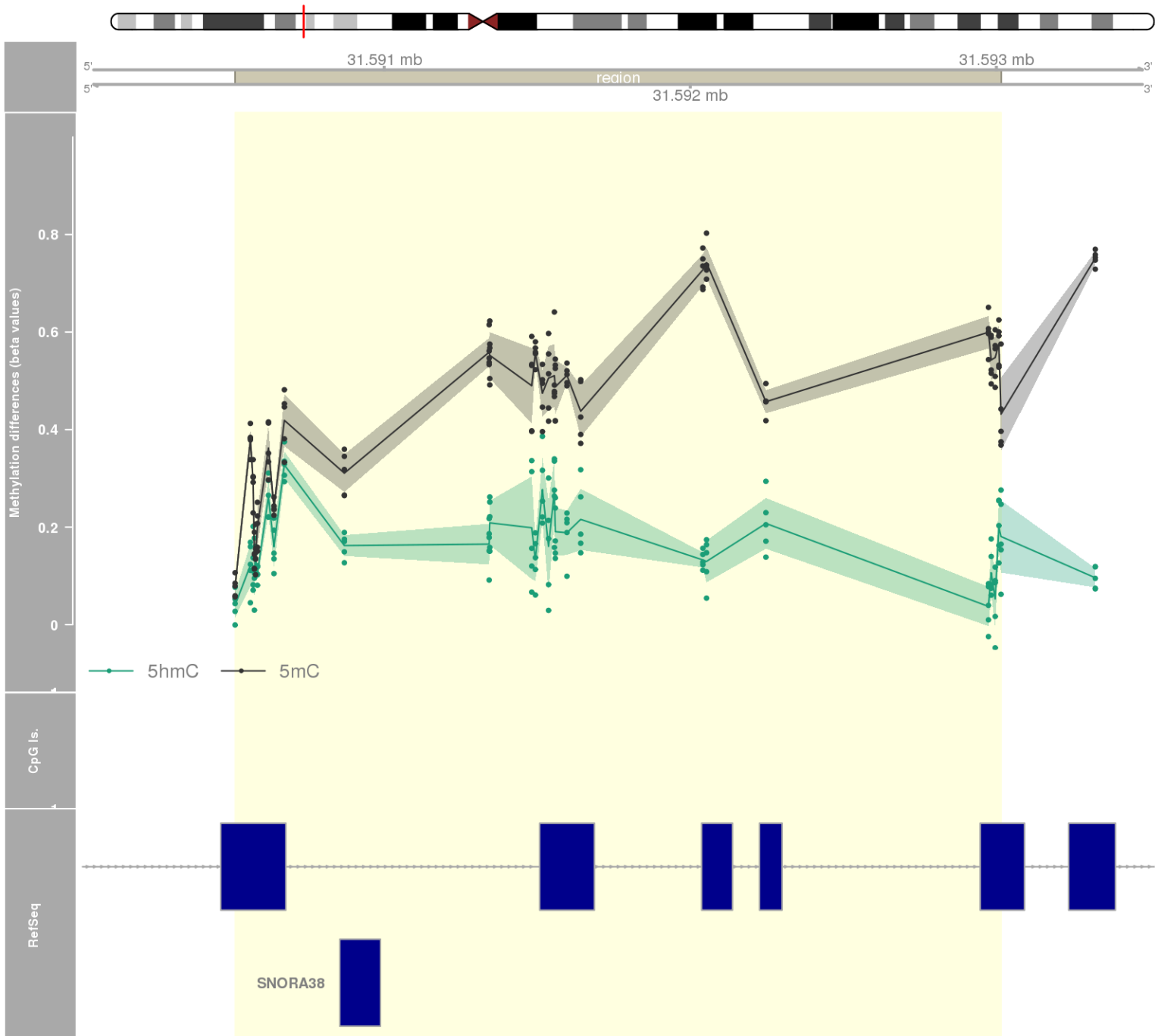
DMR 11 // chr6:30649909-30653799 // 3890 pb. (33 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: PPP1R18 -



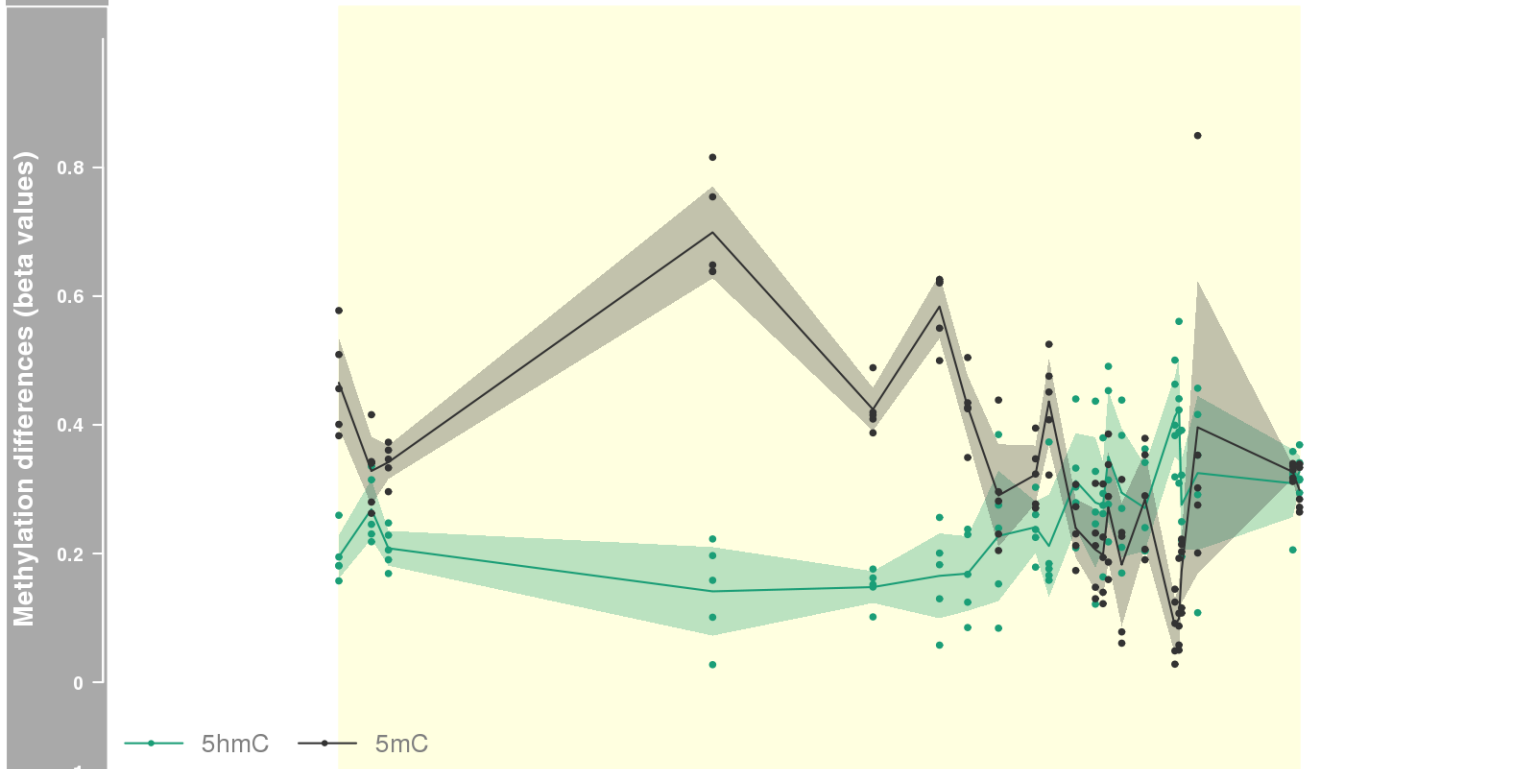
DMR 12 // chr6:32944067-32948526 // 4459 pb. (28 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: BRD2 -



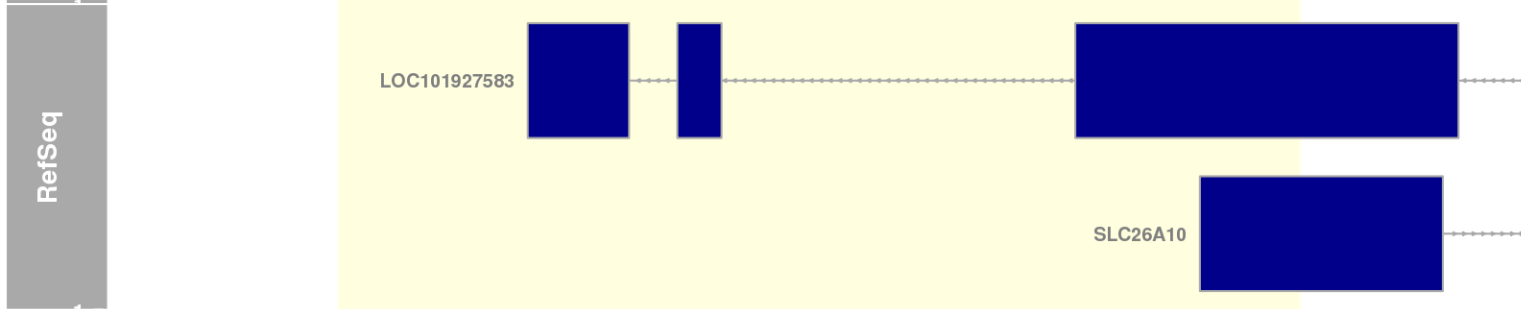
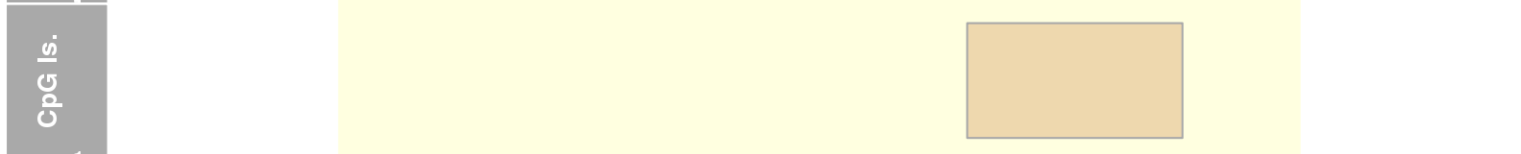
DMR 13 // chr6:31590513-31593015 // 2502 pb. (28 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: SNORA38 / PRRC2A -



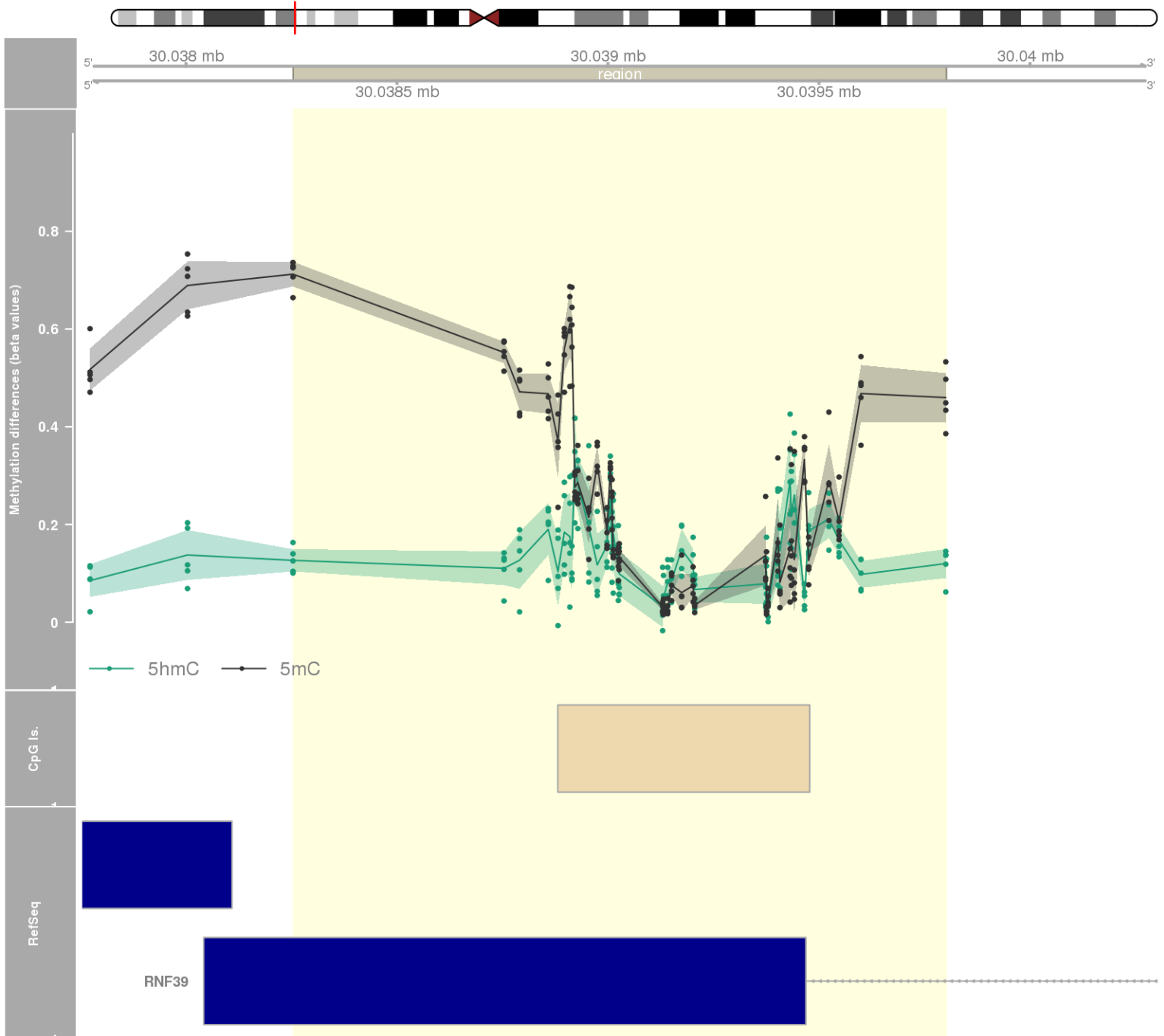
DMR 14 // chr12:58011764-58013915 // 2151 pb. (22 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: ARHGEF25 / SLC26A10 -



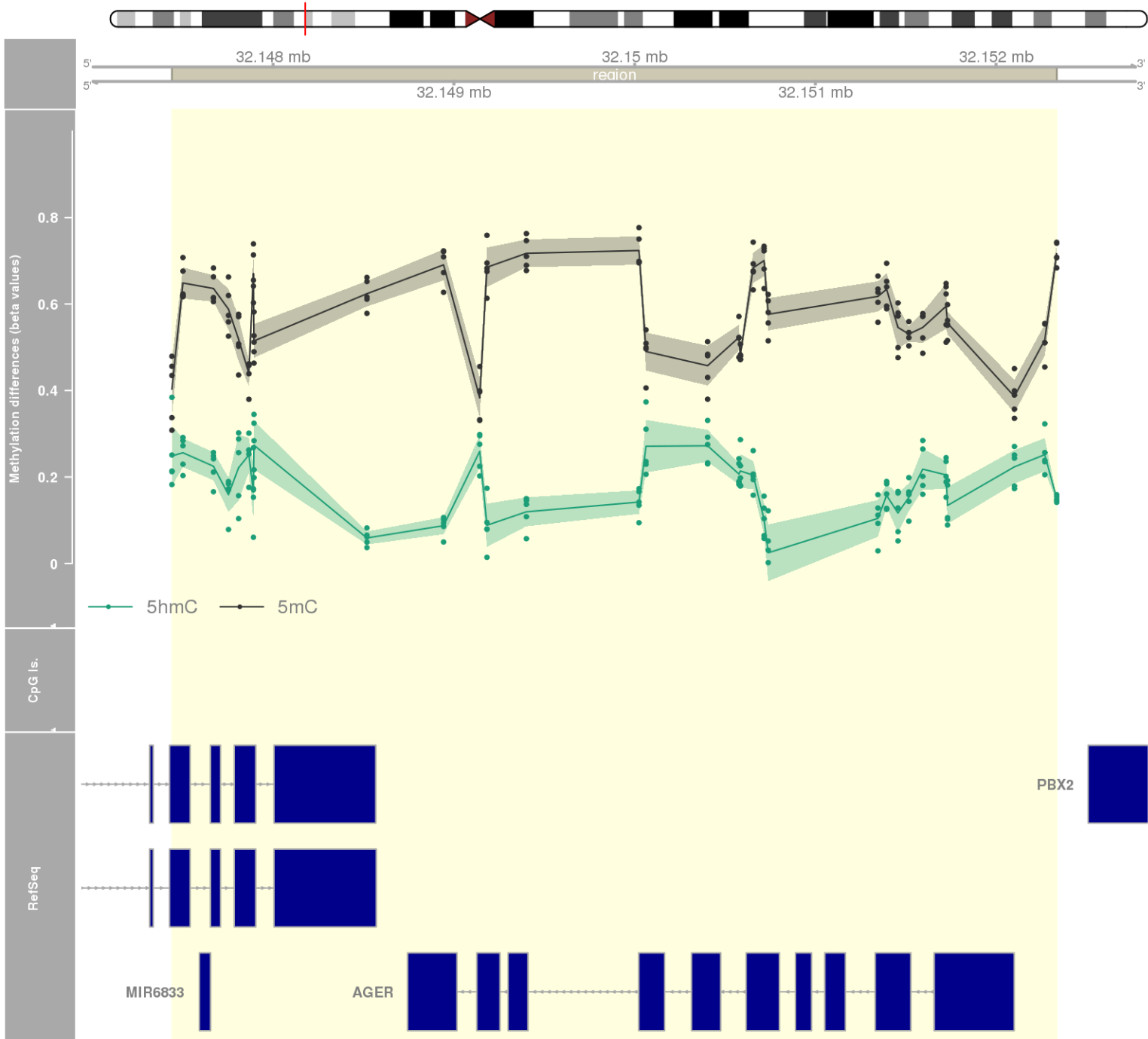
—●— 5hmC —●— 5mC



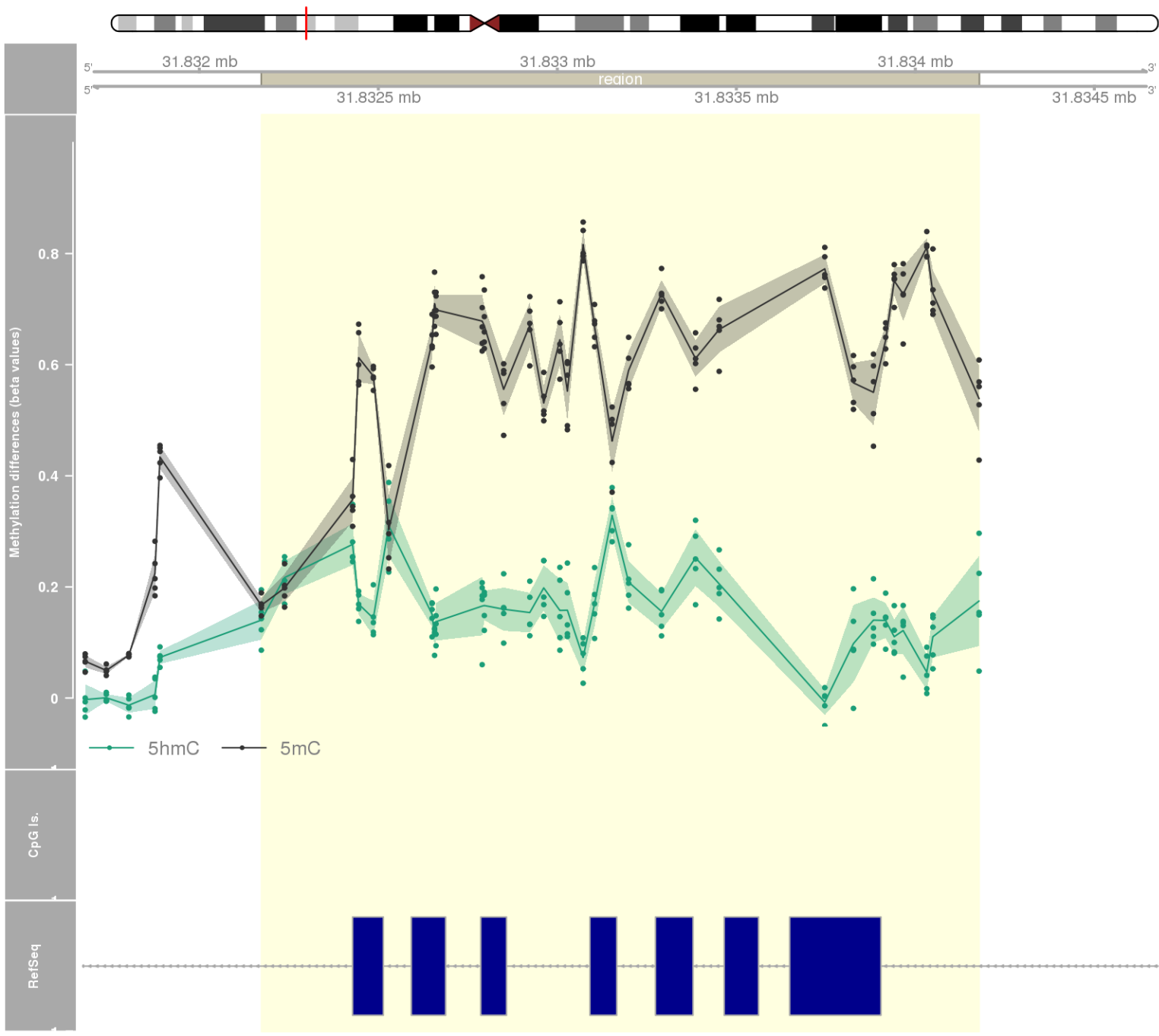
DMR 15 // chr6:30038254-30039801 // 1547 pb. (39 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: RNF39 -



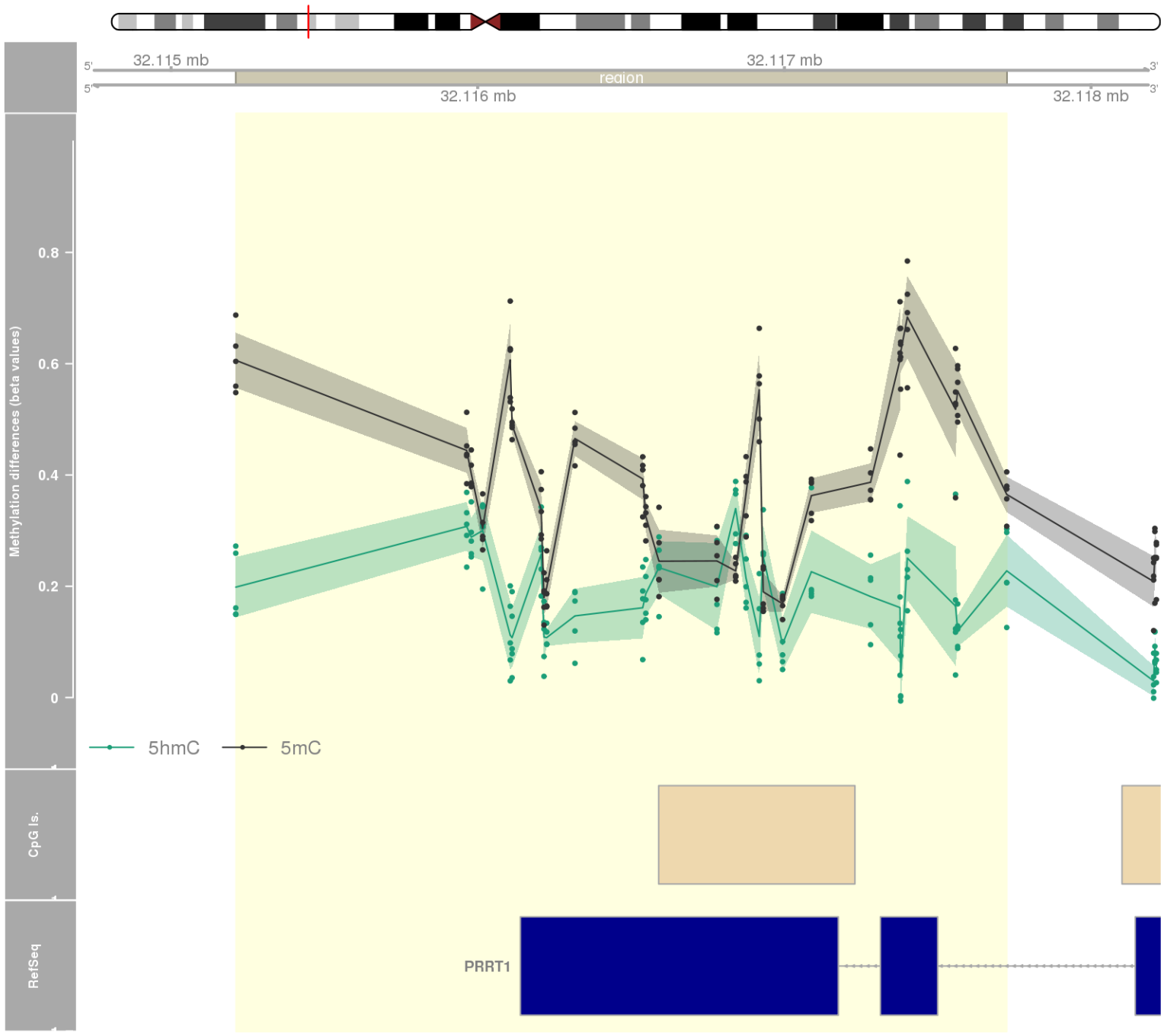
DMR 16 // chr6:32147440-32152335 // 4895 pb. (31 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: RNF5 / AGER -



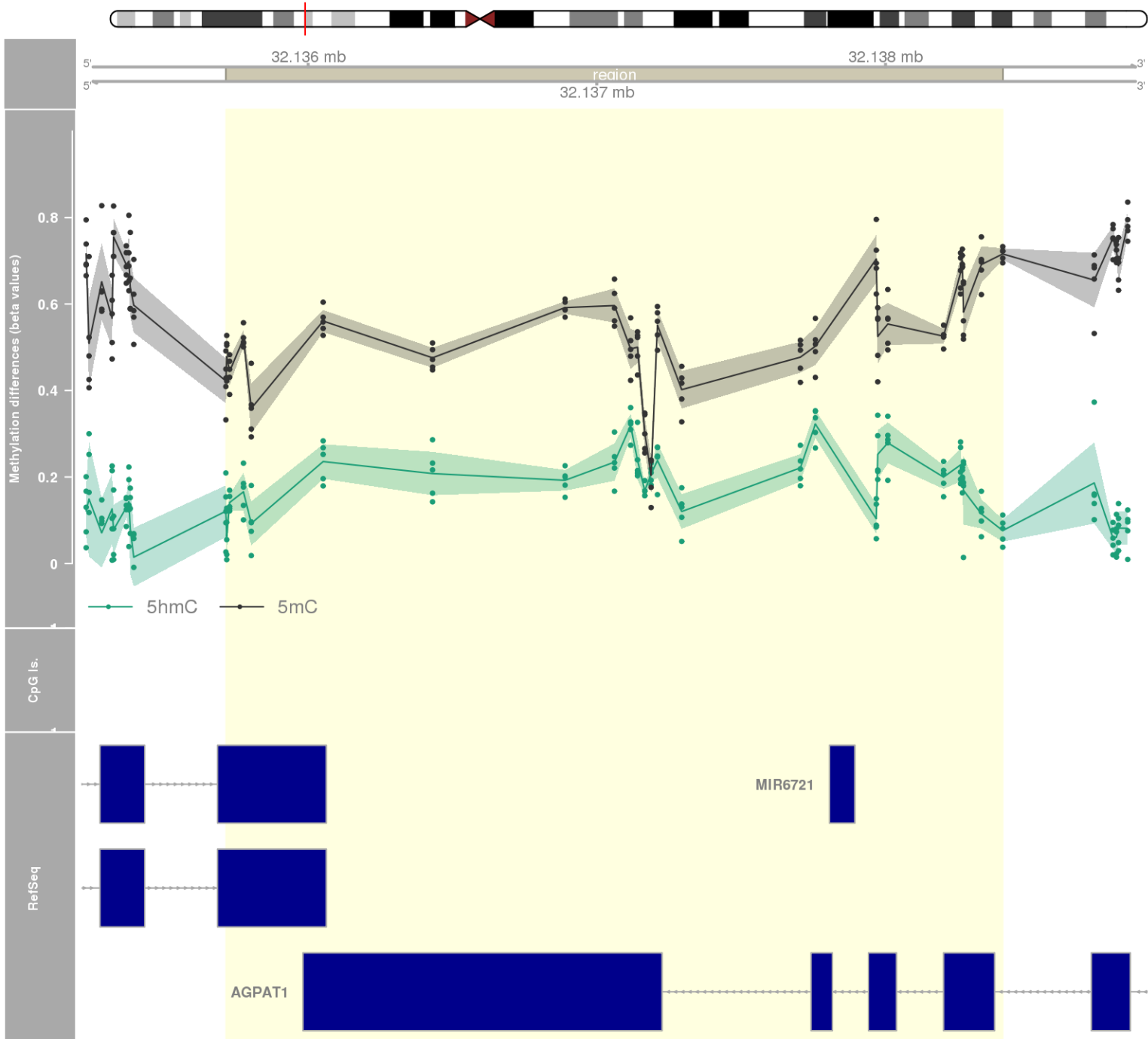
DMR 17 // chr6:31832173-31834178 // 2005 pb. (32 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: SLC44A4 -



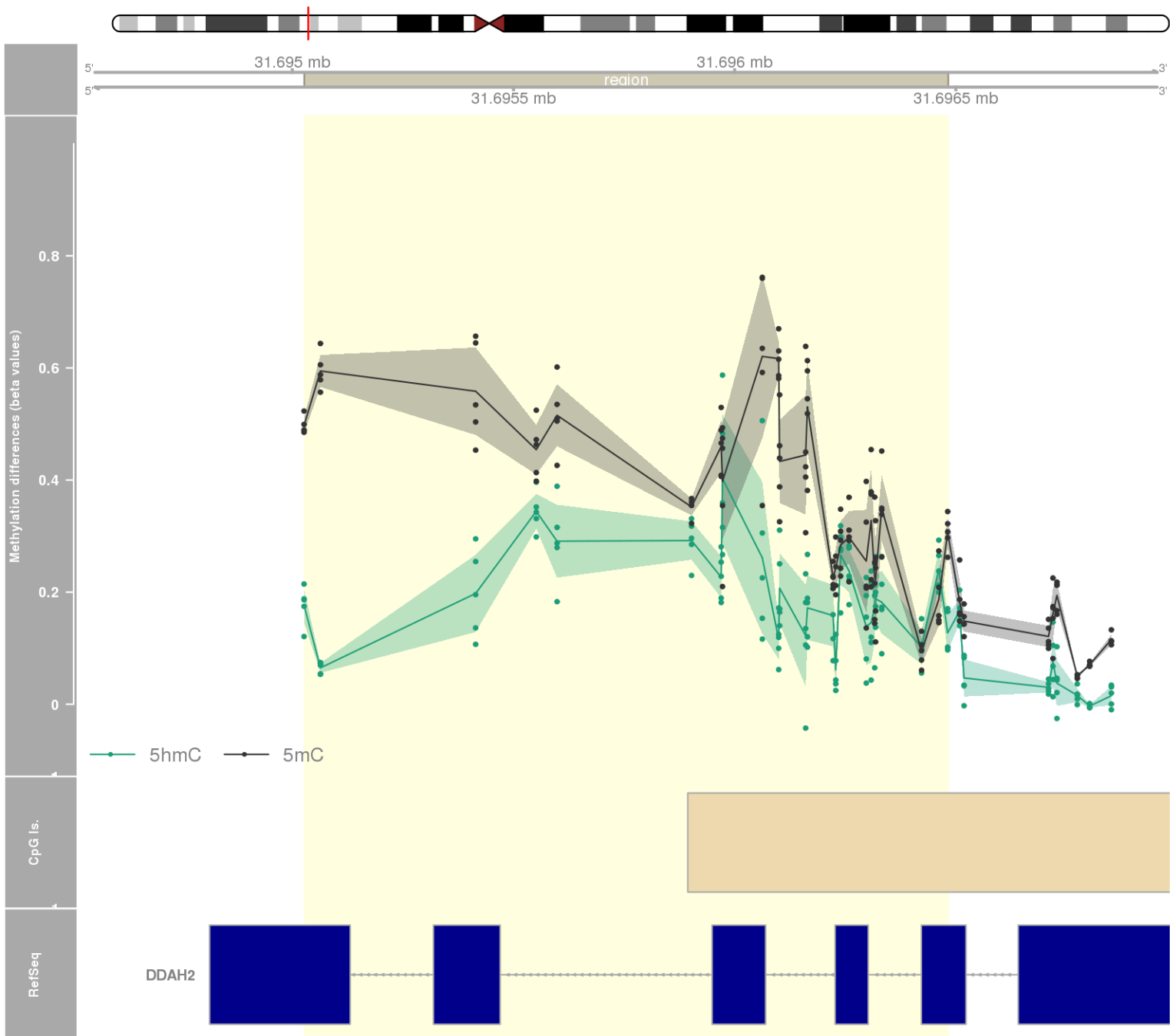
DMR 18 // chr6:32115211-32117725 // 2514 pb. (27 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: PRRT1 -



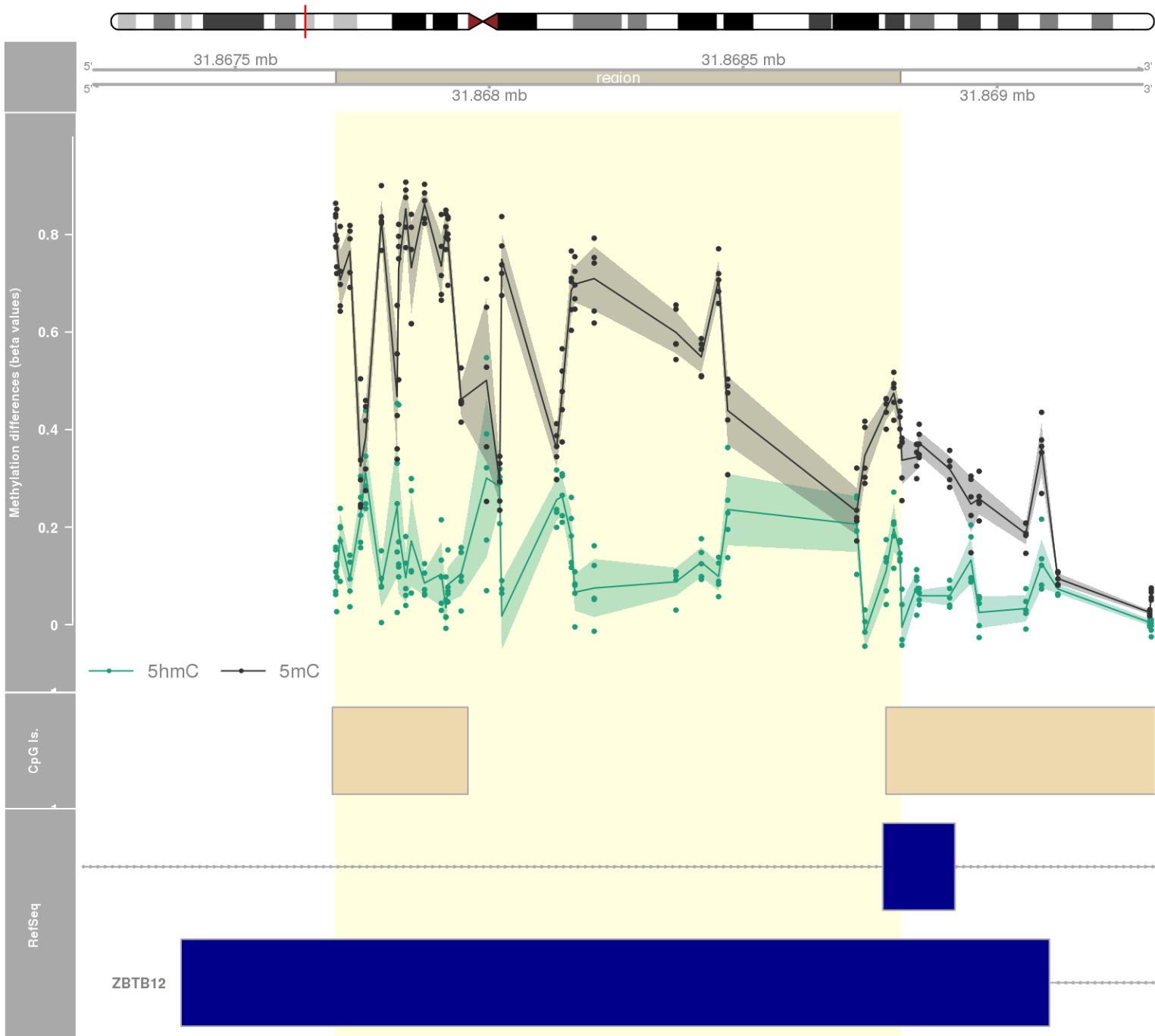
DMR 19 // chr6:32135715-32138406 // 2691 pb. (26 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: PPT2-EGFL8 / EGFL8 / AGPAT1 -



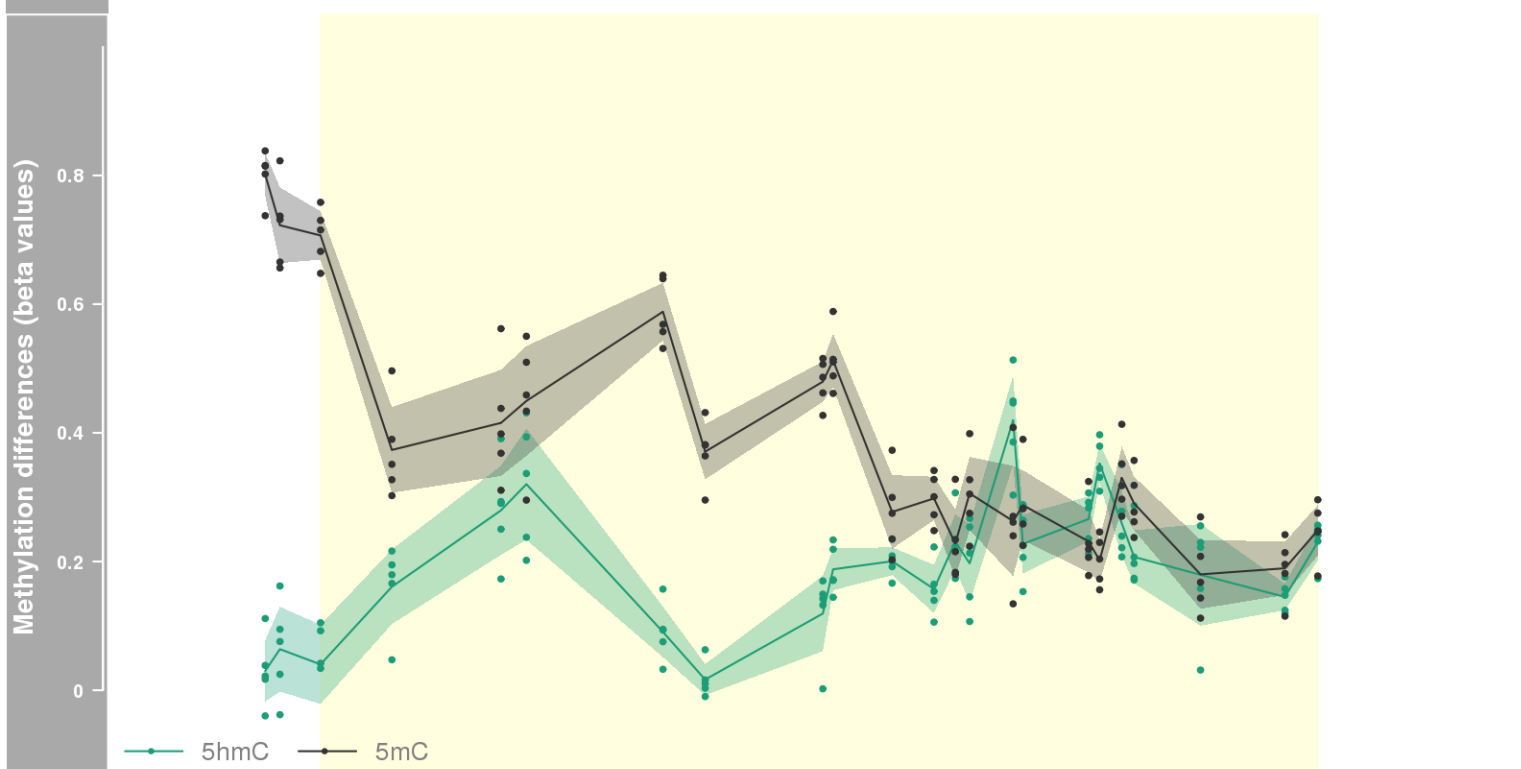
DMR 20 // chr6:31695027-31696482 // 1455 pb. (25 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: DDAH2 -



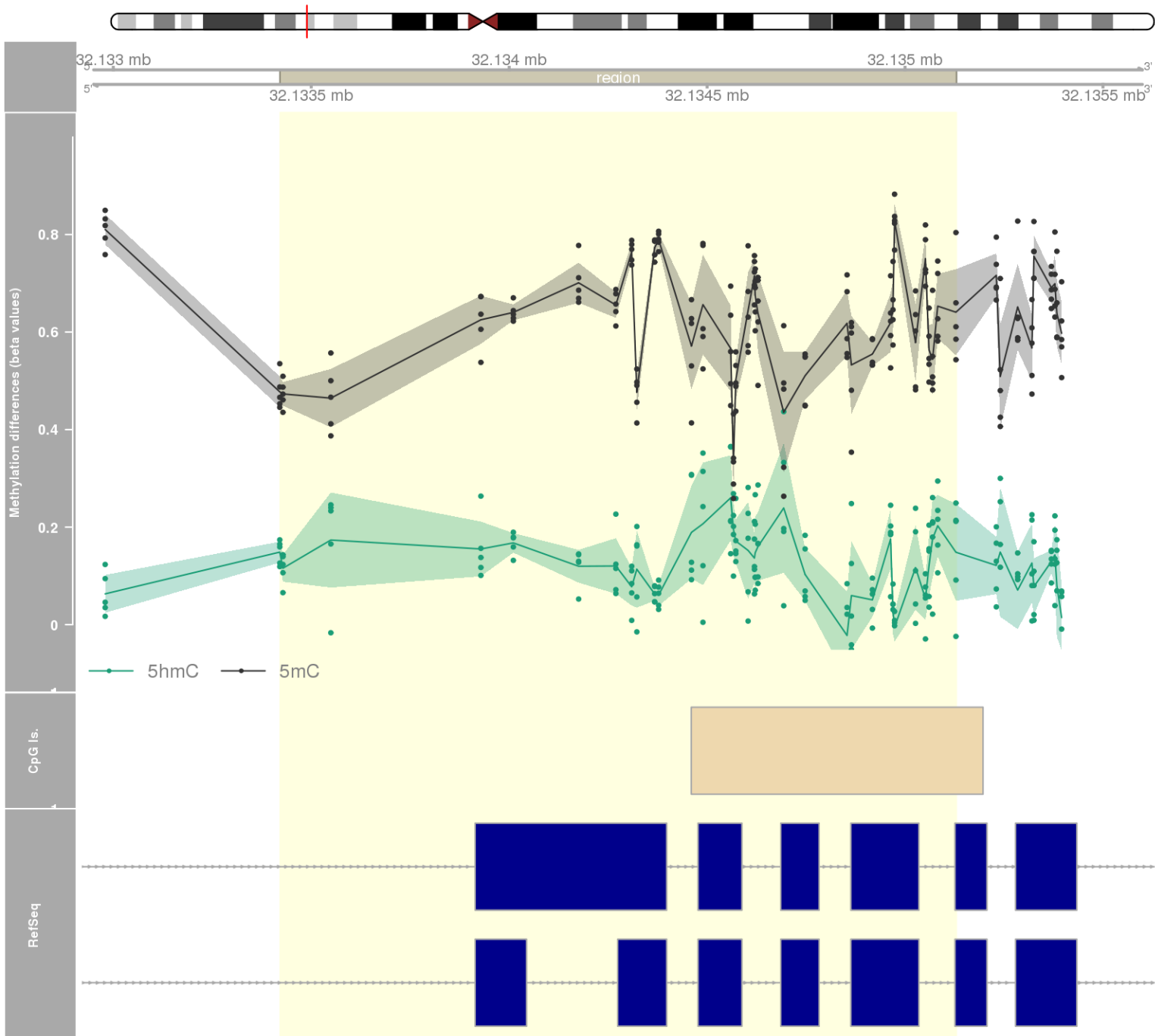
DMR 21 // chr6:31867698-31868809 // 1111 pb. (33 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: C2 / ZBTB12 -



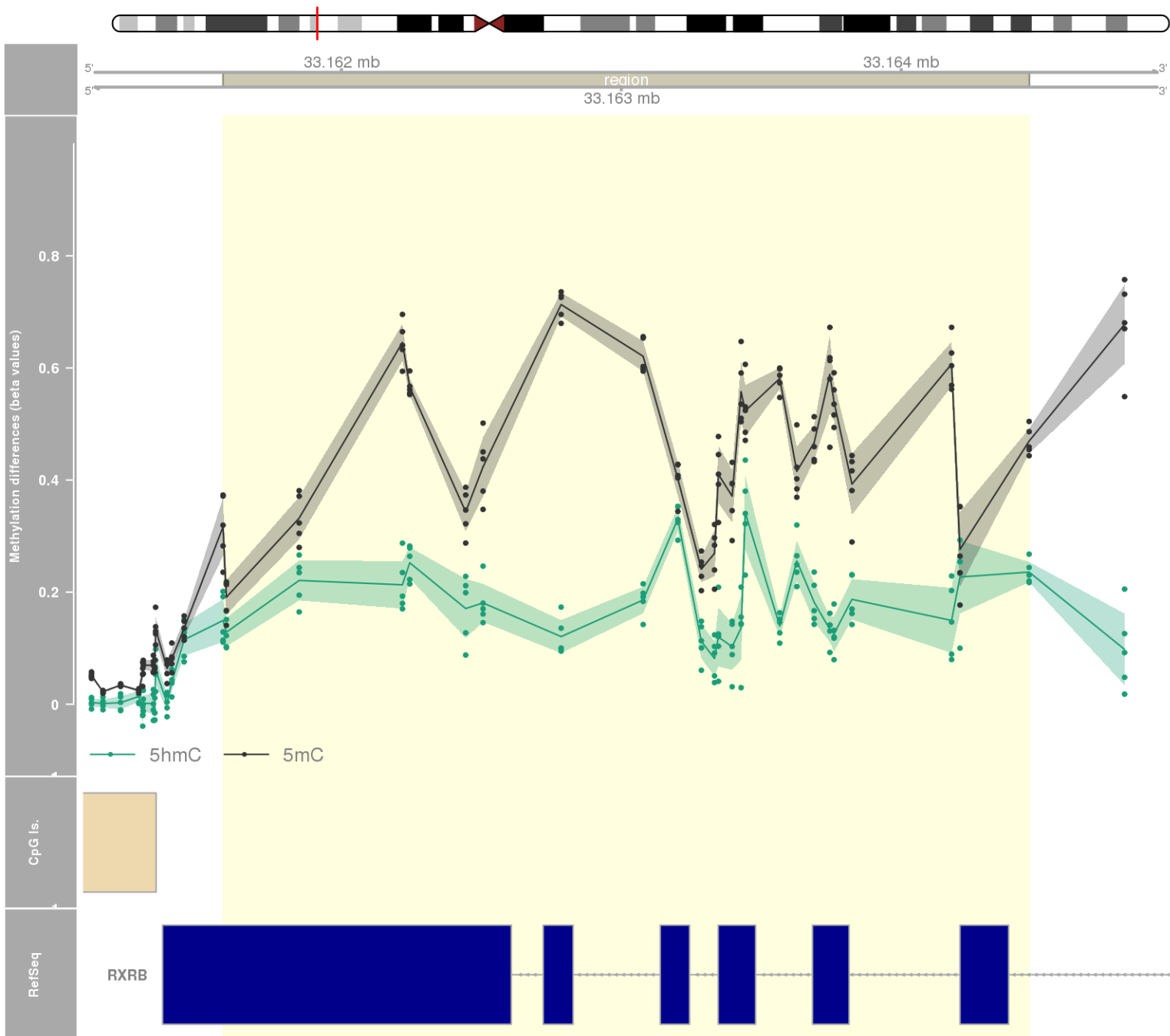
DMR 22 // chr17:7756708-7759140 // 2432 pb. (21 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.001 // fwerArea: 0.173
- genes: KDM6B / TMEM88 -



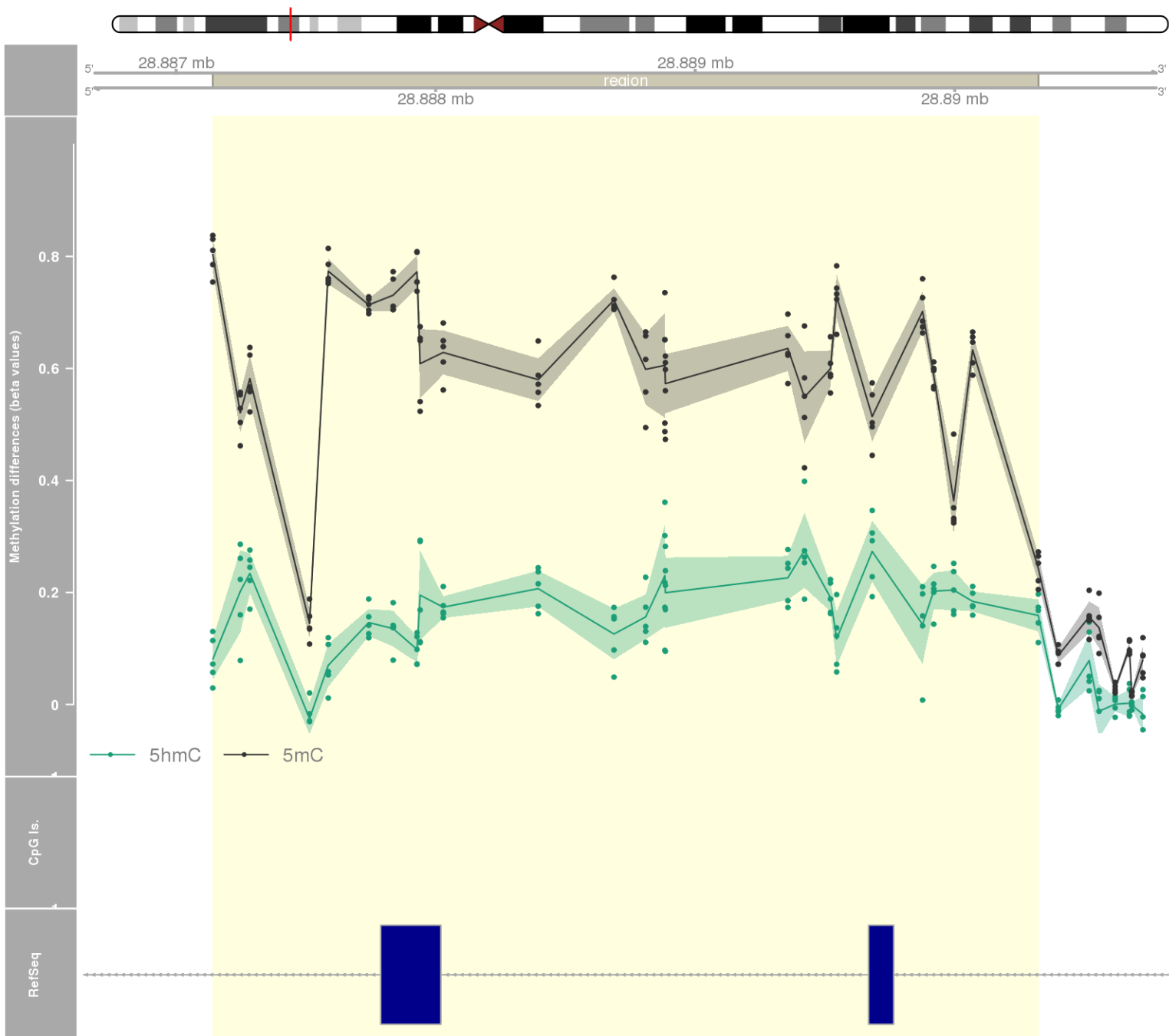
DMR 23 // chr6:32133421-32135129 // 1708 pb. (34 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: PPT2 / PPT2-EGFL8 / EGFL8 -



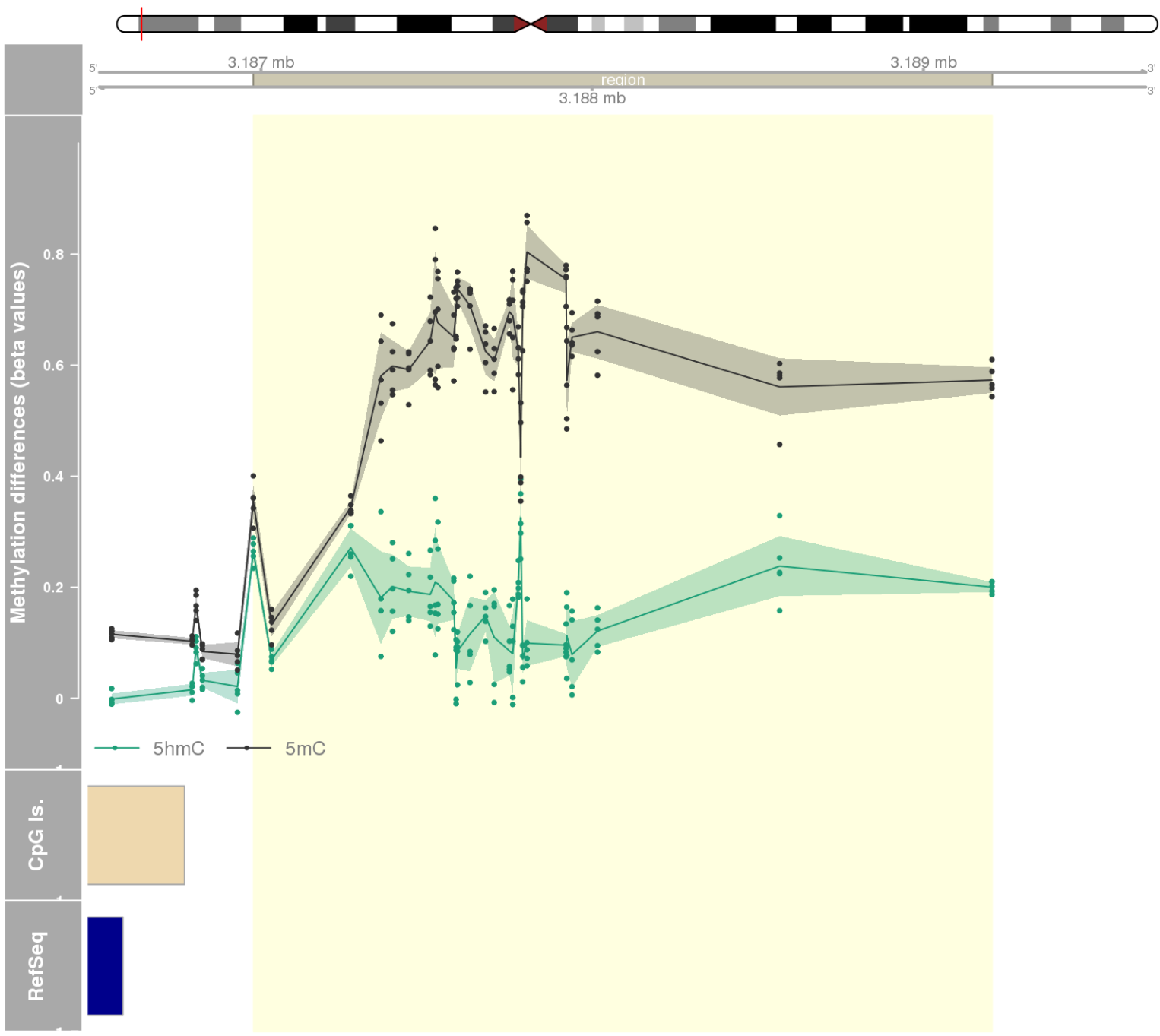
DMR 24 // chr6:33161577-33164457 // 2880 pb. (25 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: RXRB -



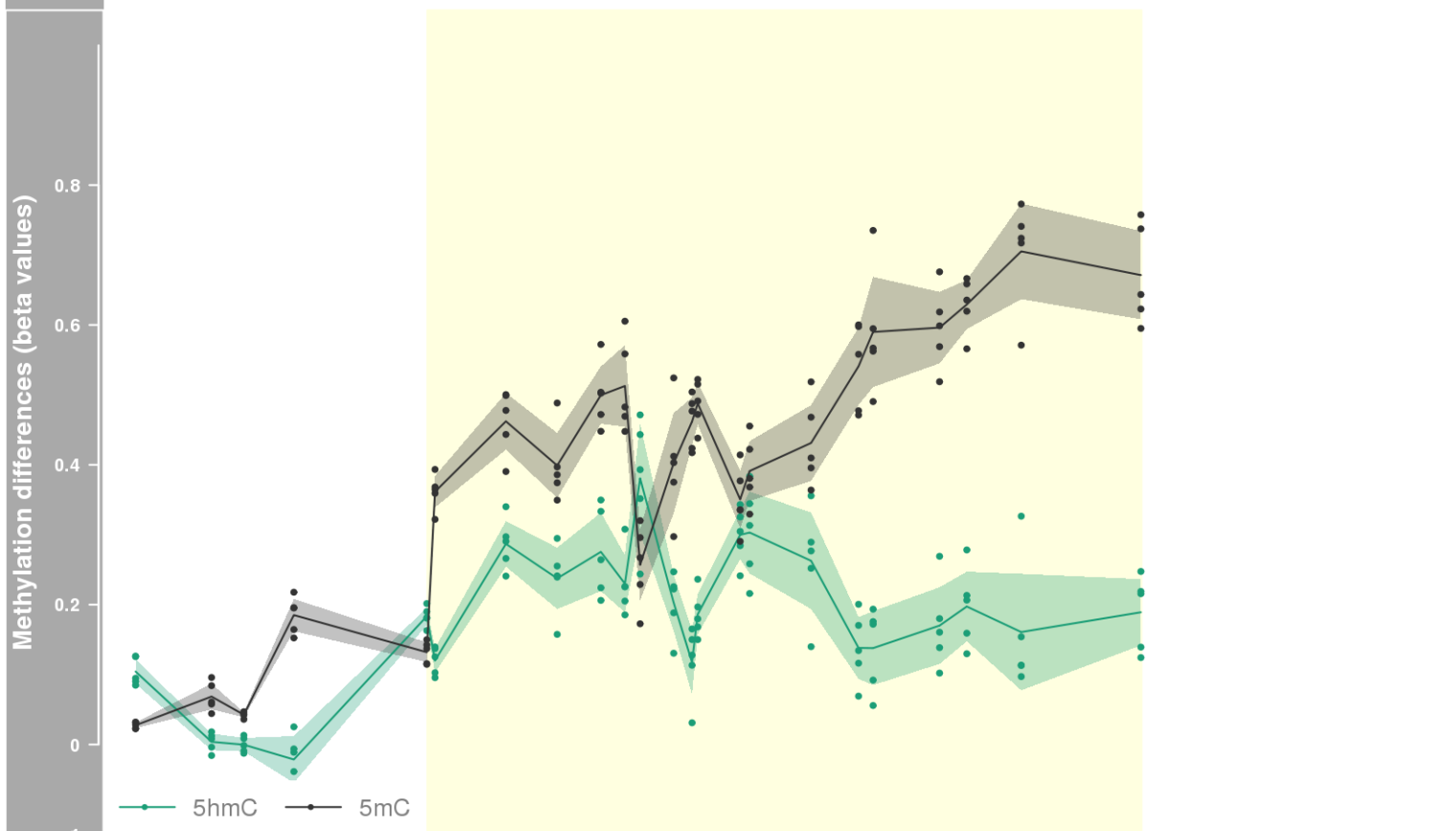
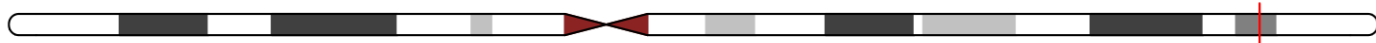
DMR 25 // chr6:28887141-28890322 // 3181 pb. (25 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: TRIM27 -



DMR 26 // chr11:3186977-3189207 // 2230 pb. (27 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: OSBPL5 -



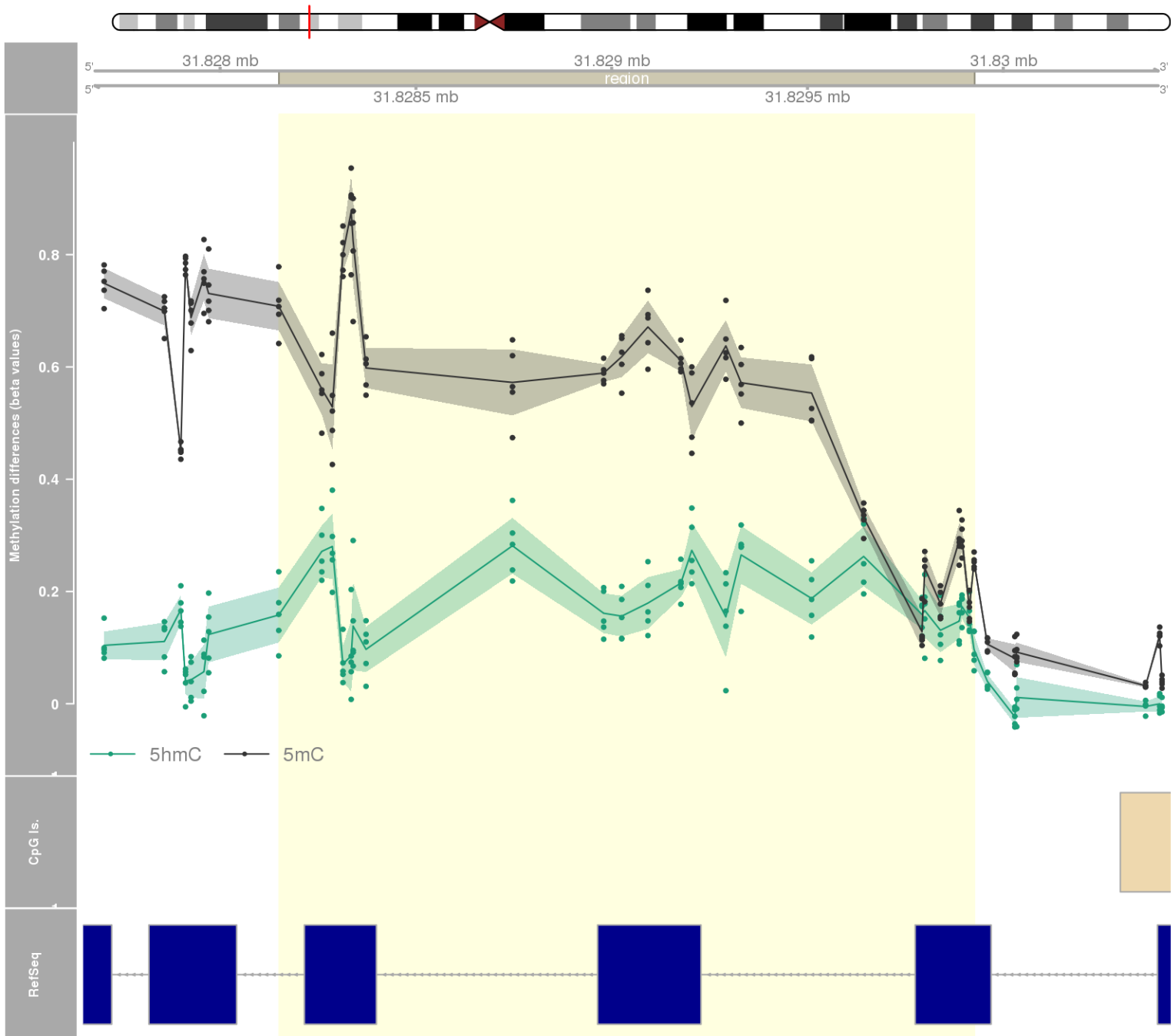
DMR 27 // chr20:57582581-57583709 // 1128 pb. (19 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173



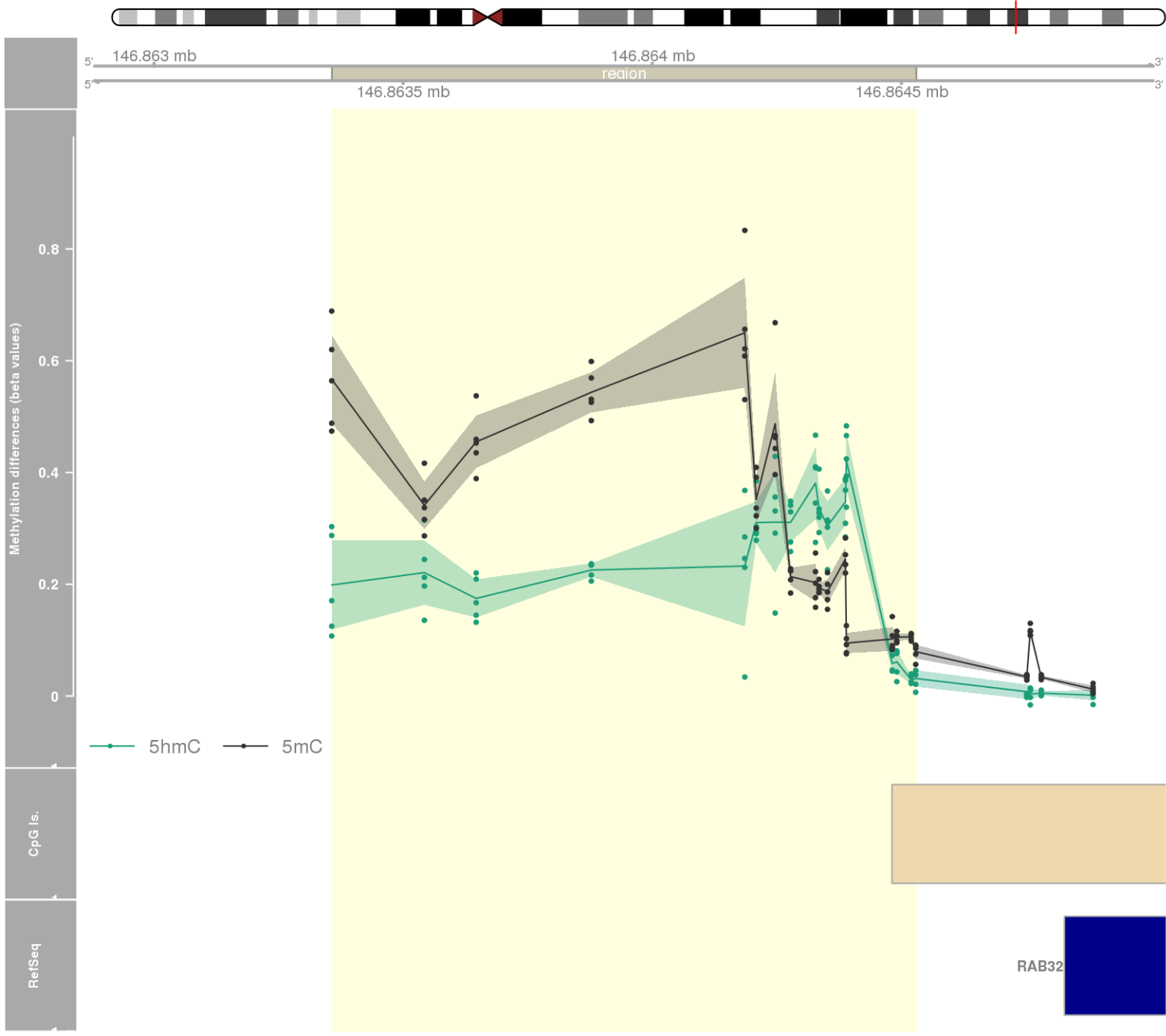
5hmC 5mC



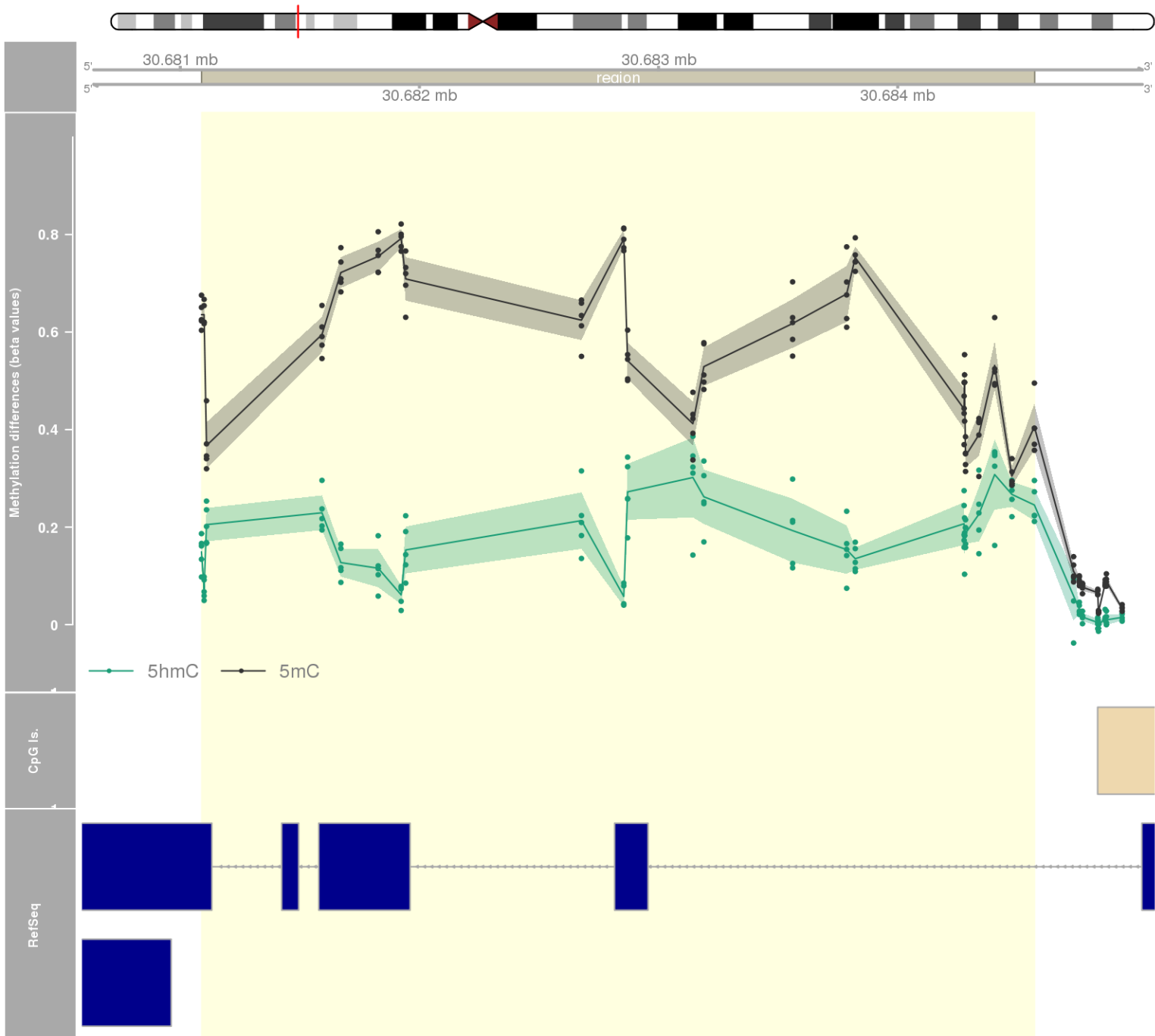
DMR 28 // chr6:31828150-31829926 // 1776 pb. (24 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: NEU1 -



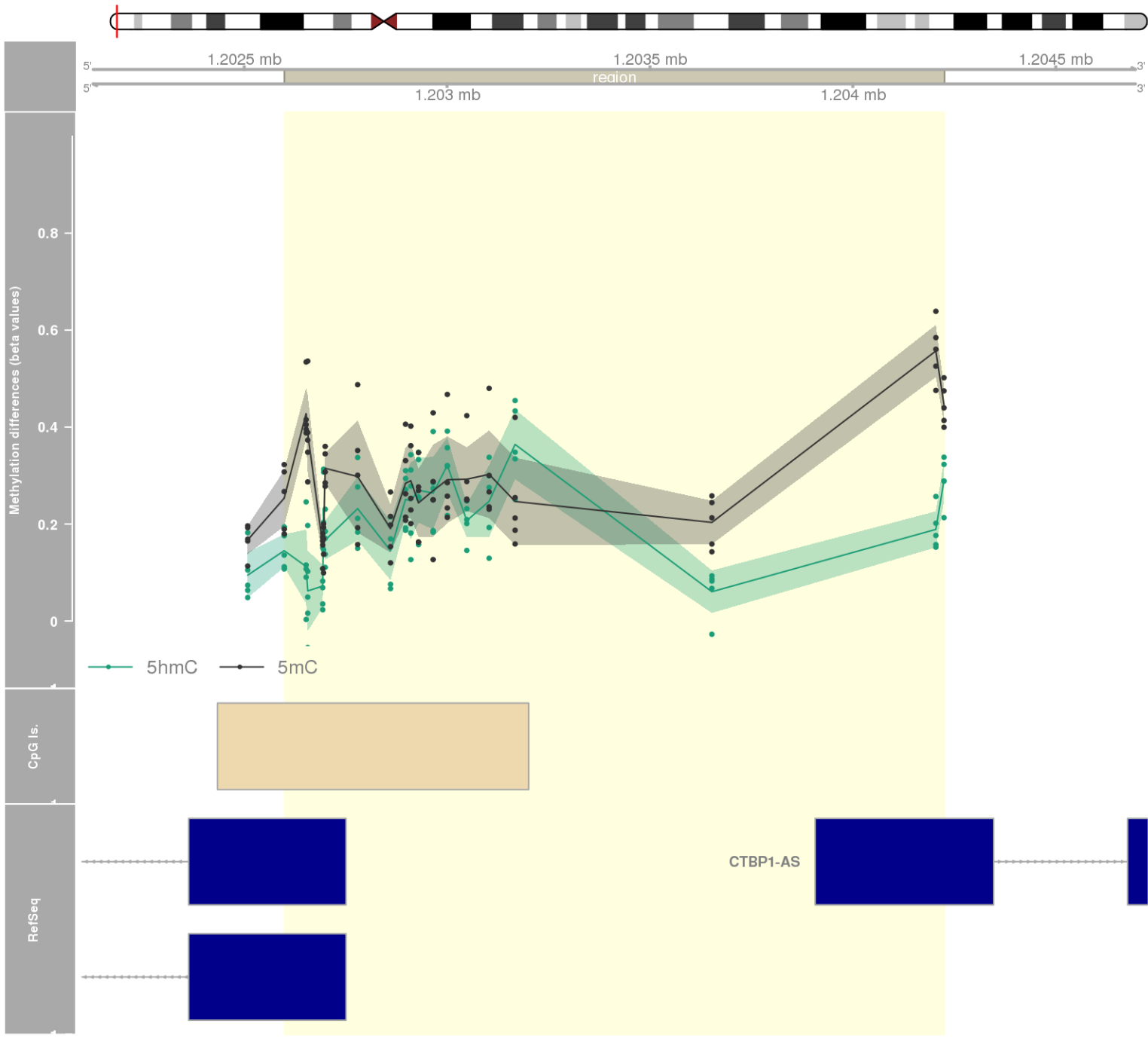
DMR 29 // chr6:146863357-146864529 // 1172 pb. (17 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173



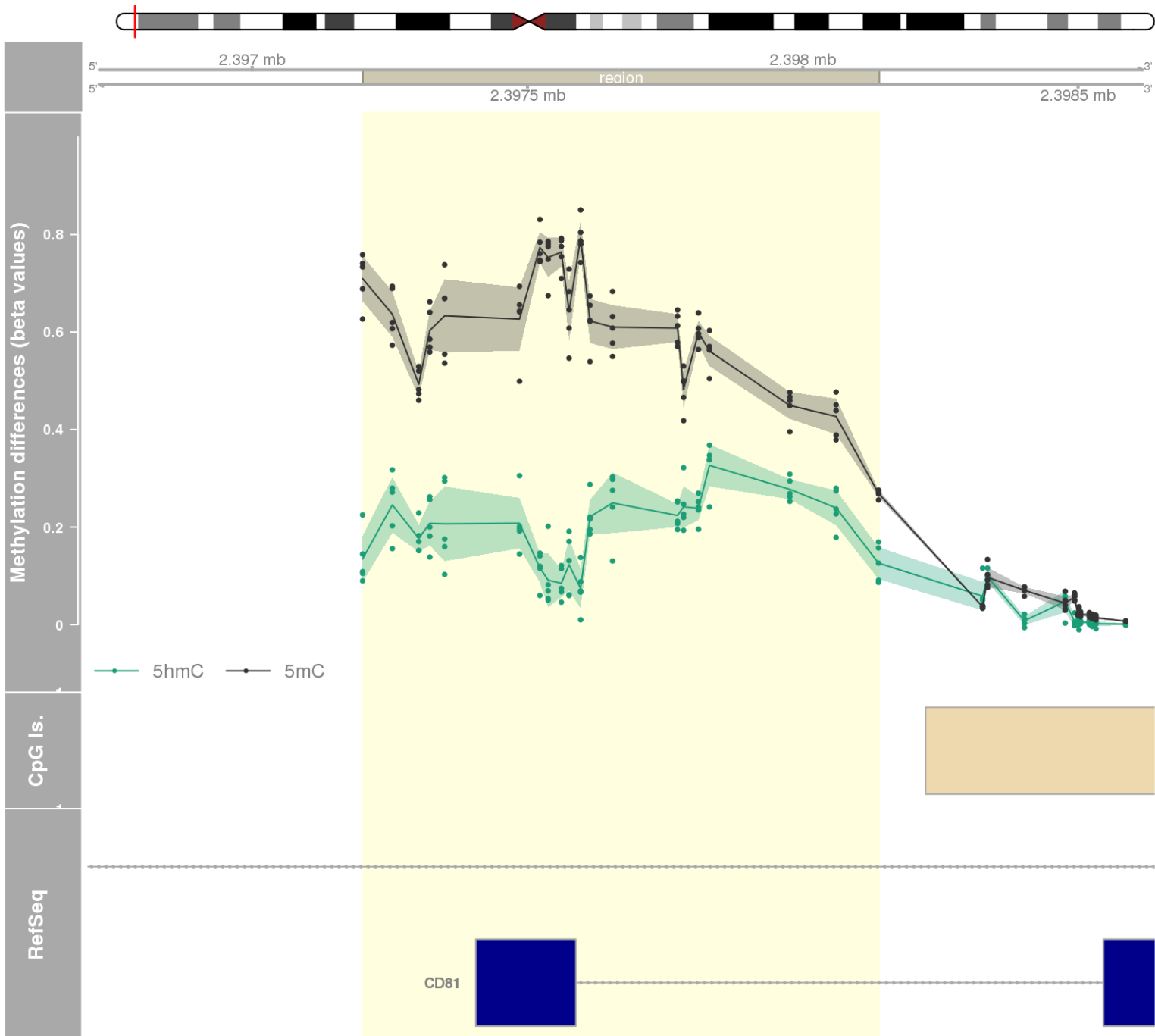
DMR 30 // chr6:30681089-30684572 // 3483 pb. (23 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: MDC1 -



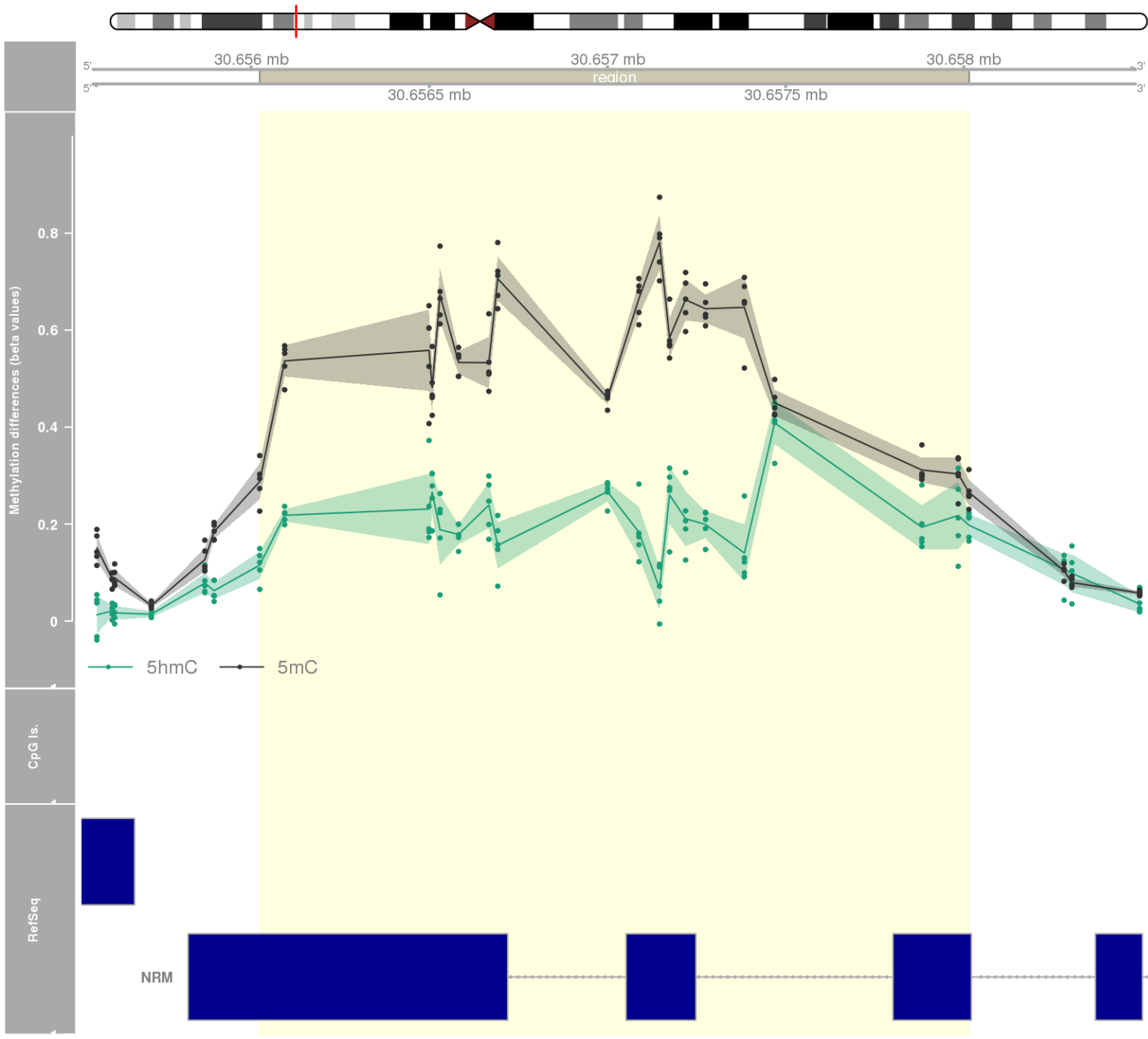
DMR 31 // chr4:1202599-1204225 // 1626 pb. (19 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: SPON2 -



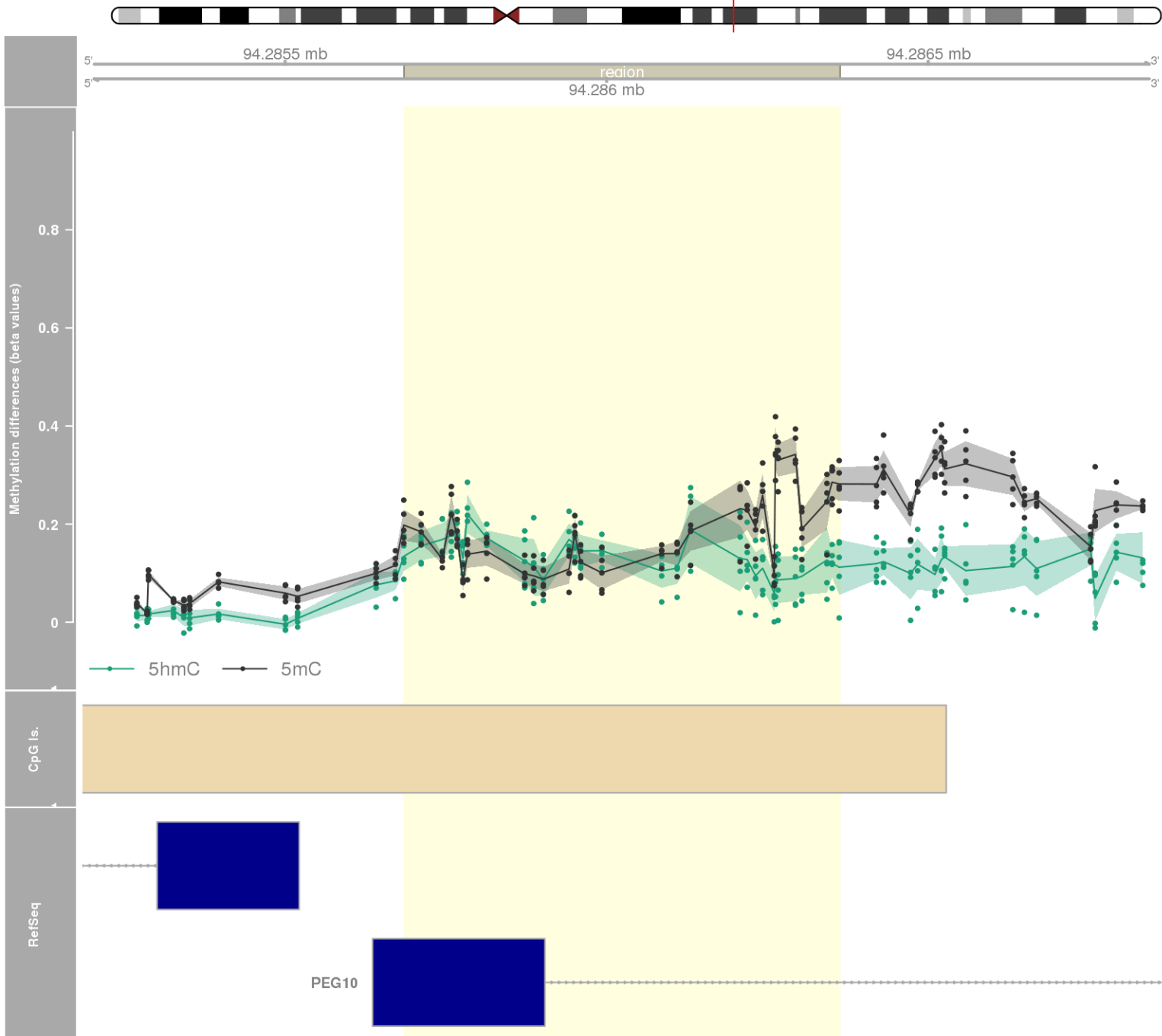
DMR 32 // chr11:2397201-2398138 // 937 pb. (20 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: CD81-AS1 / CD81 -



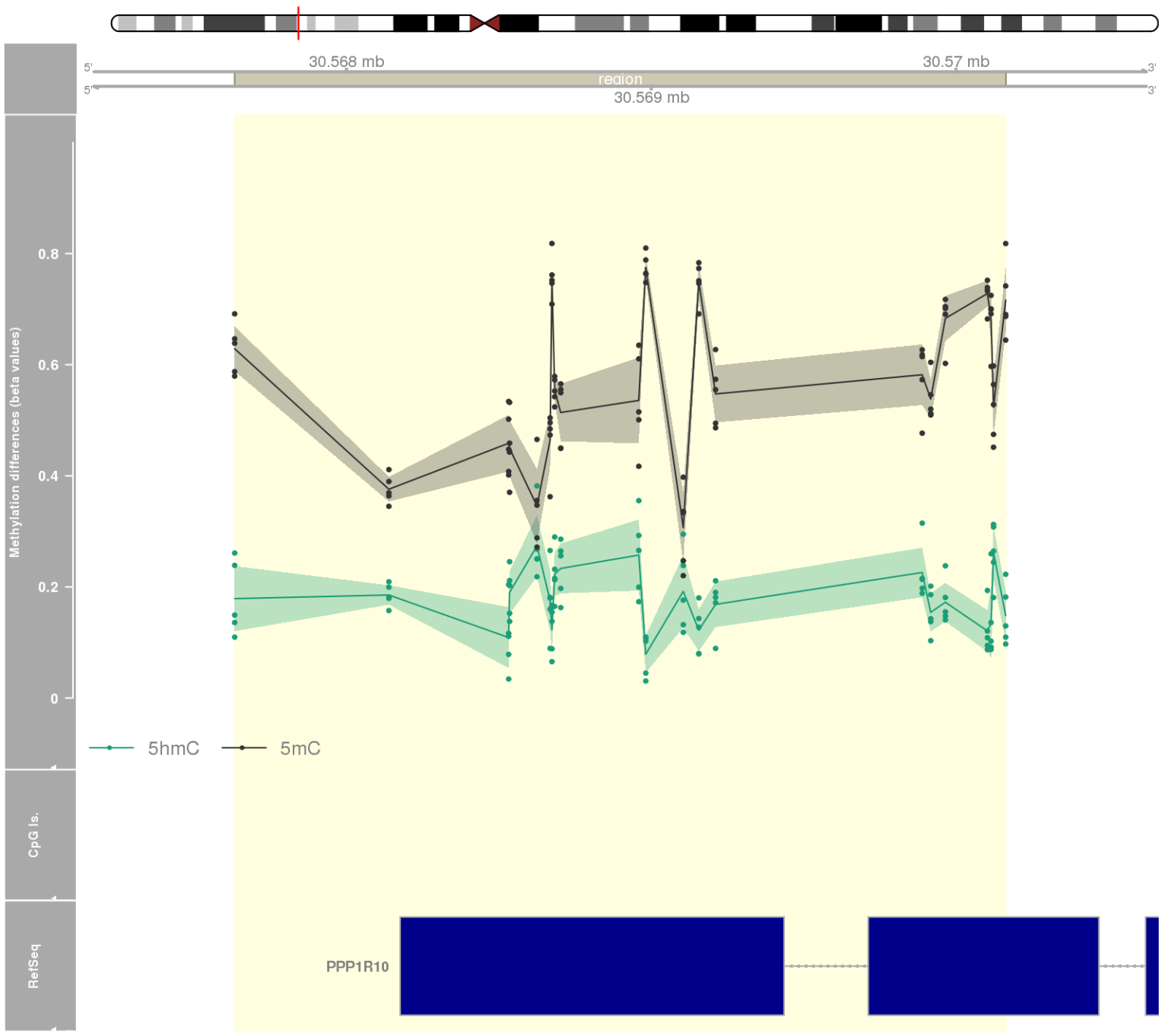
DMR 33 // chr6:30656024-30658014 // 1990 pb. (19 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: NRM -



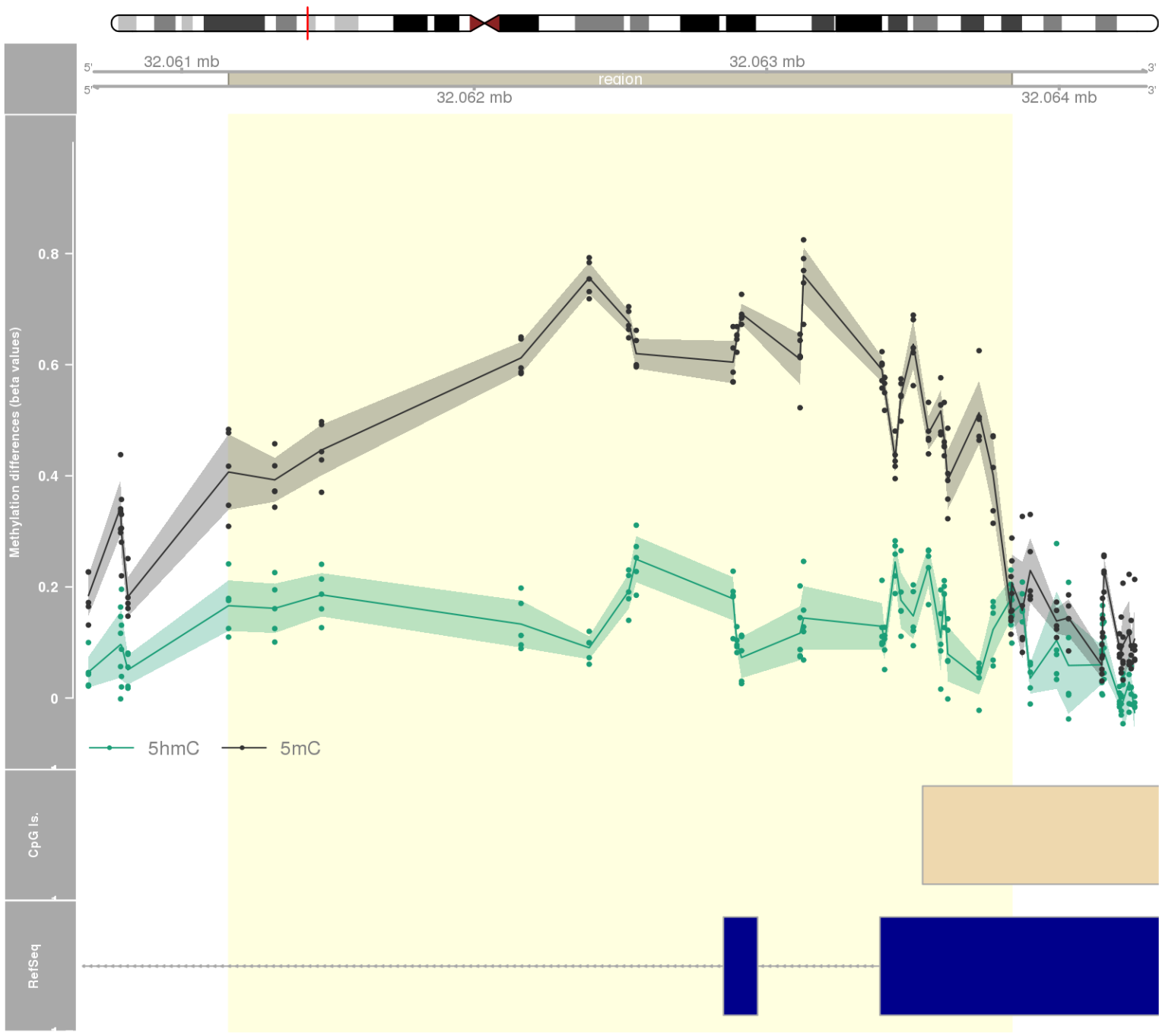
DMR 34 // chr7:94285685-94286362 // 677 pb. (30 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: PEG10 -



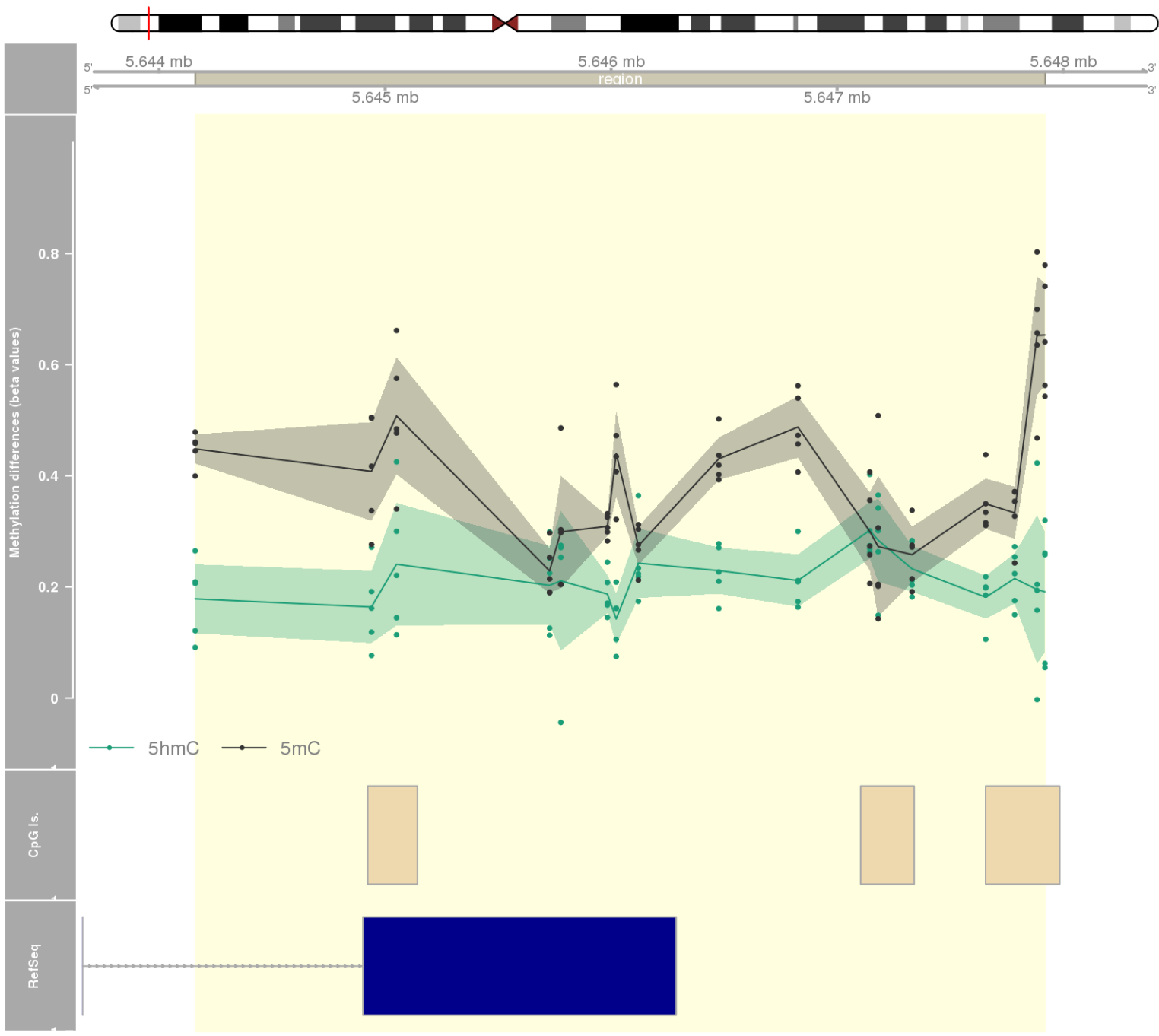
DMR 35 // chr6:30567633-30570163 // 2530 pb. (21 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: PPP1R10 -



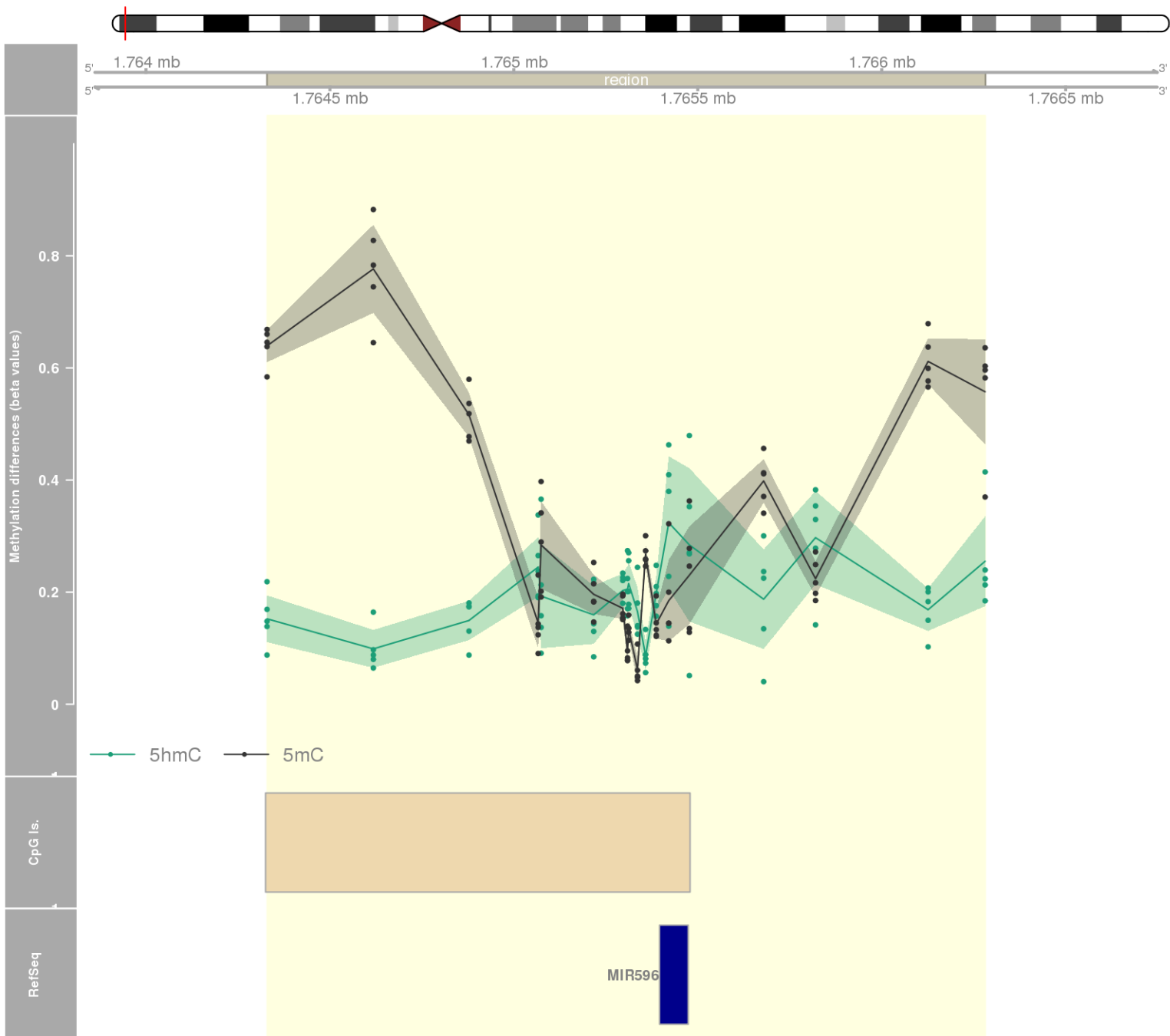
DMR 36 // chr6:32061160-32063838 // 2678 pb. (25 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: TNXB -



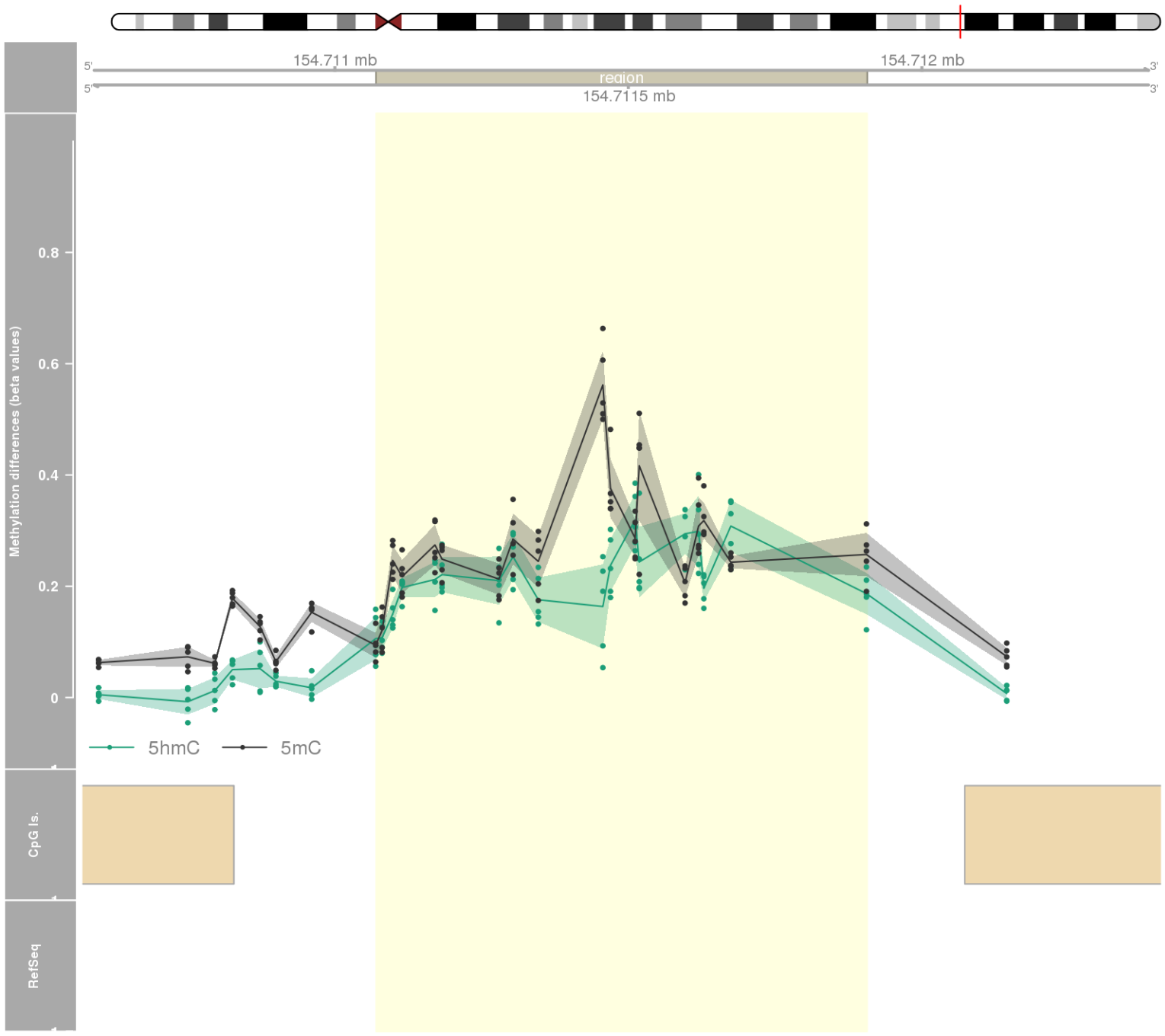
DMR 37 // chr7:5644160-5647920 // 3760 pb. (17 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.002 // fwerArea: 0.173
- genes: FSCN1 -



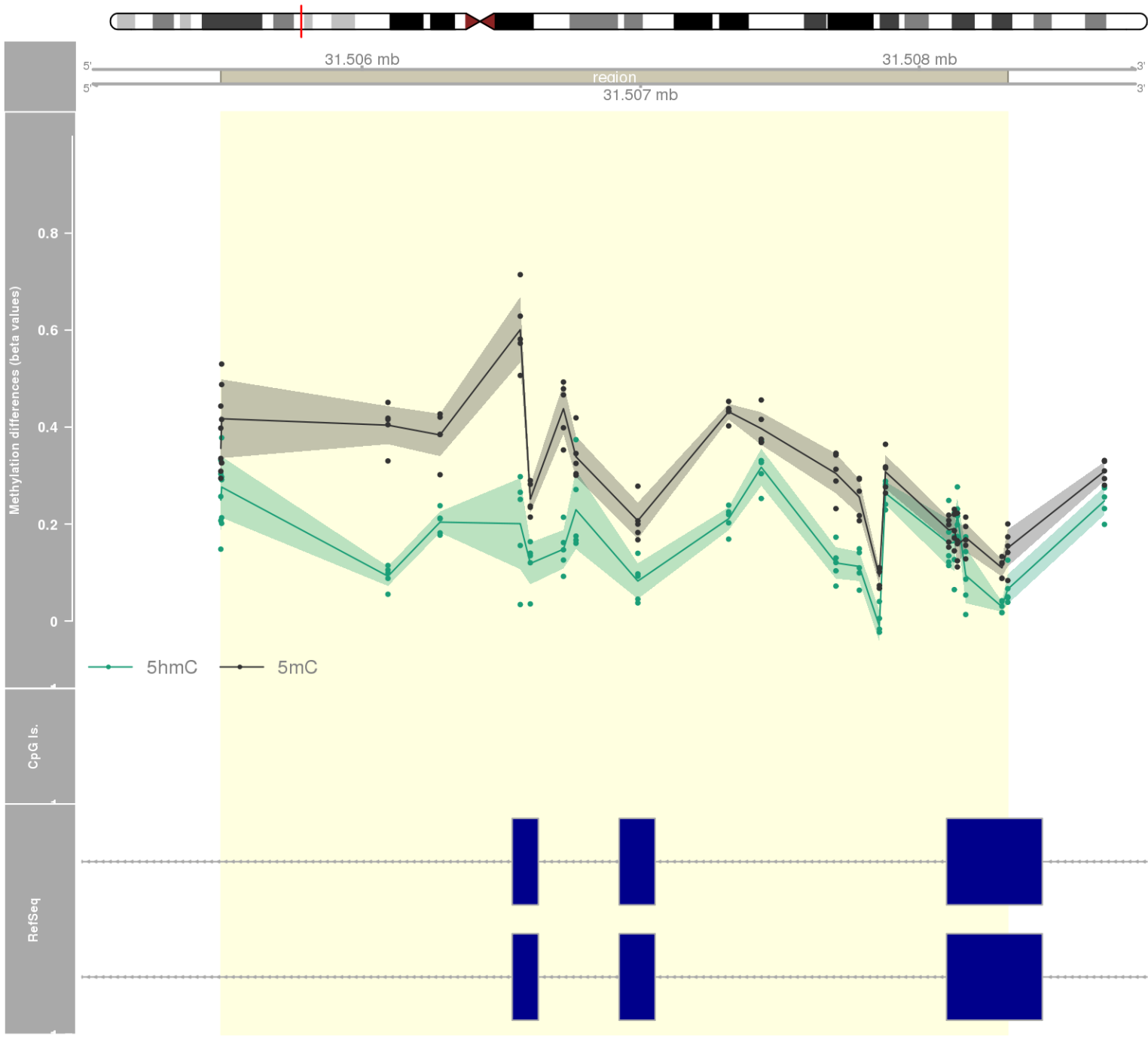
DMR 38 // chr8:1764329-1766281 // 1952 pb. (18 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.173
- genes: MIR596 -



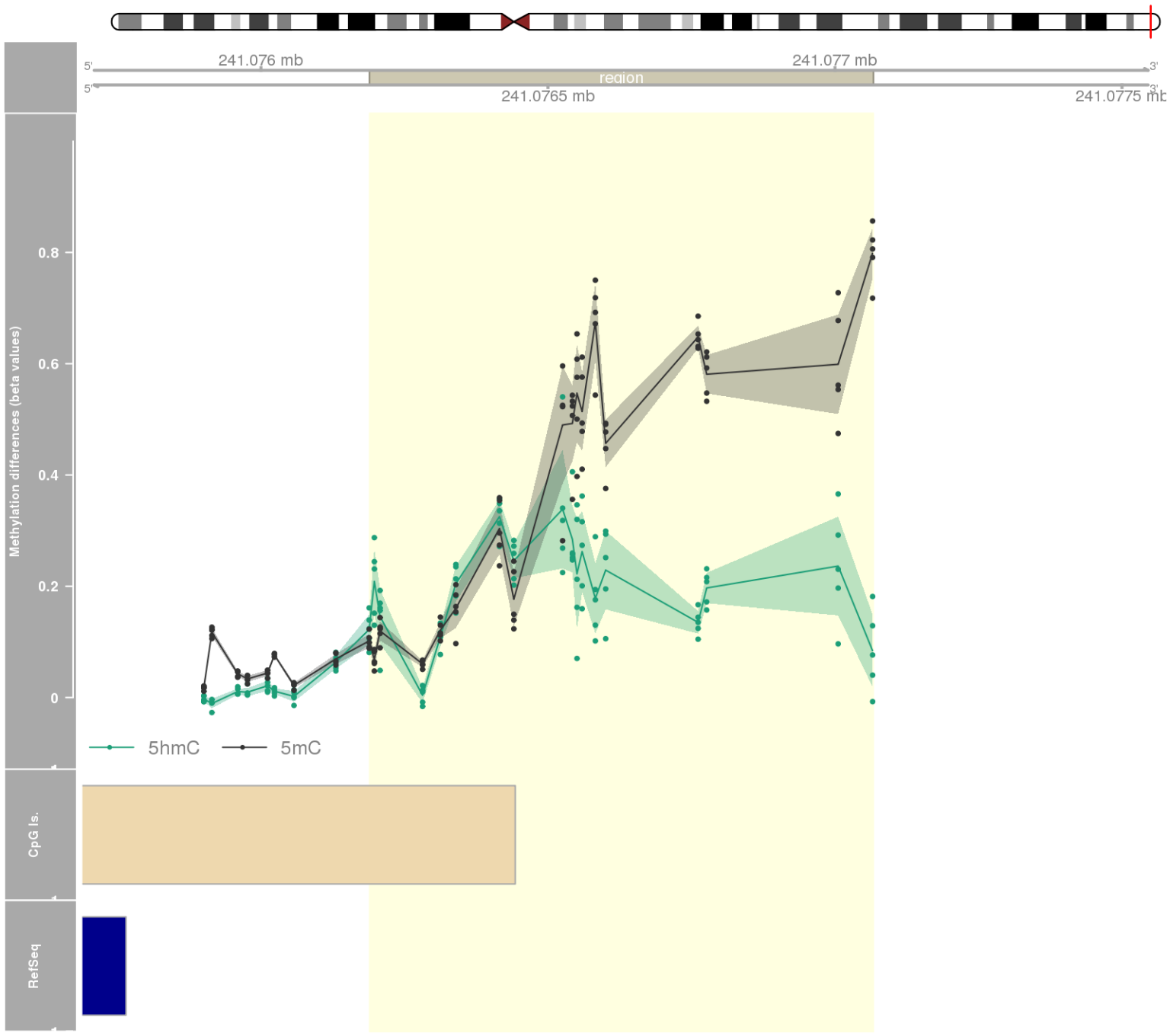
DMR 39 // chr4:154711070-154711906 // 836 pb. (18 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.173



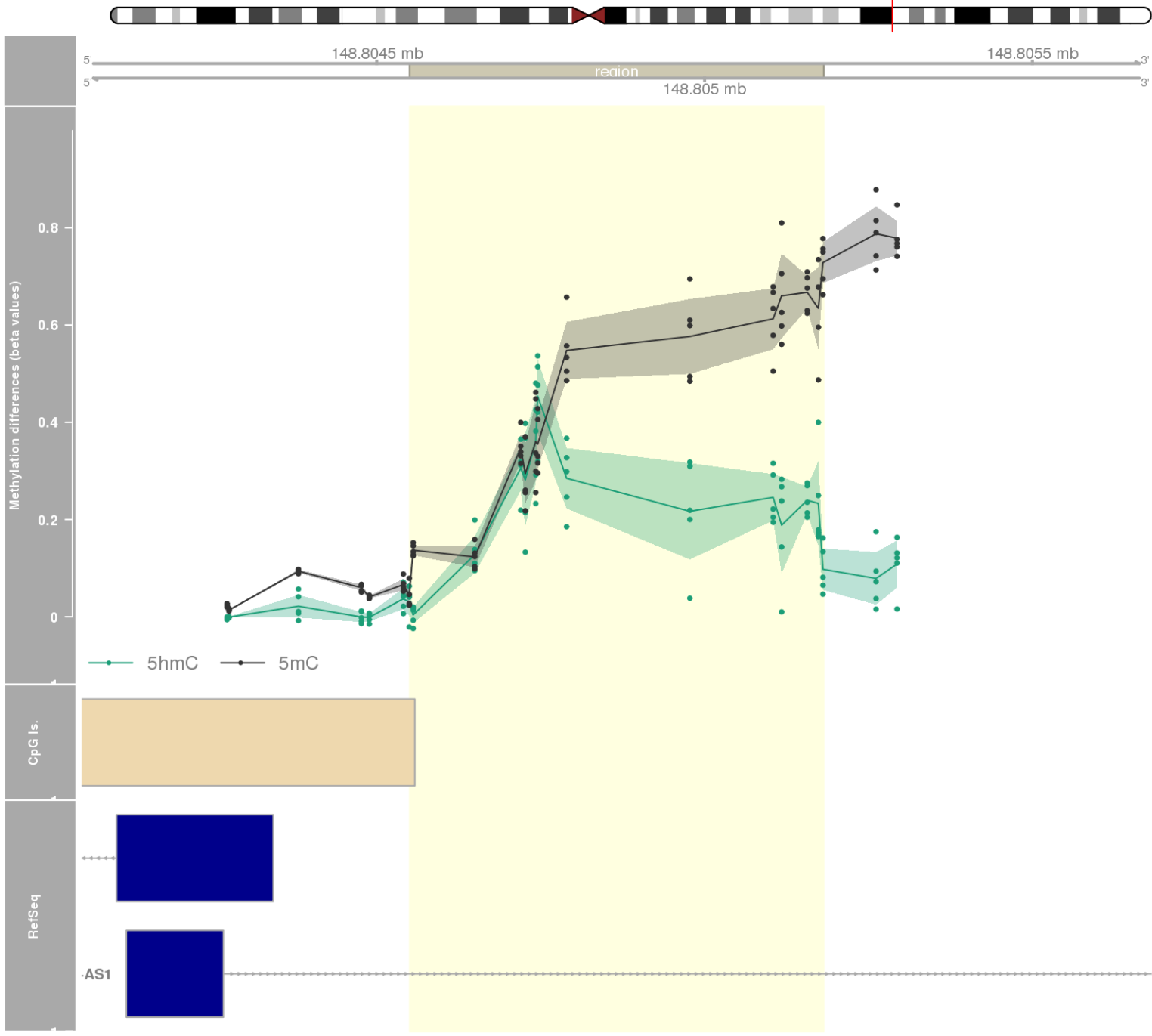
DMR 40 // chr6:31505493-31508318 // 2825 pb. (21 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.173
- genes: ATP6V1G2-DDX39B / DDX39B -



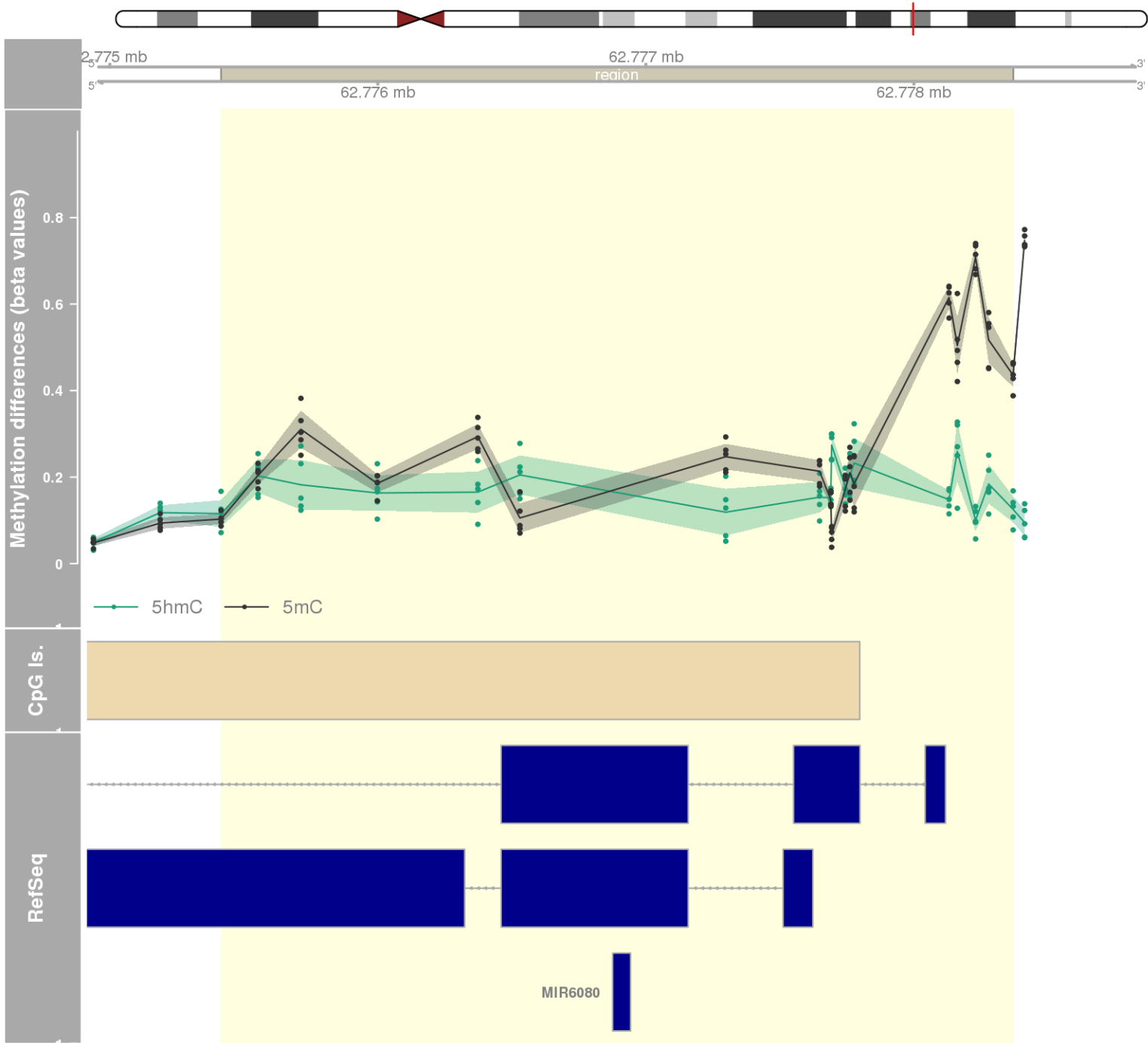
DMR 41 // chr2:241076189-241077066 // 877 pb. (18 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.173
- genes: MYEOV2 -



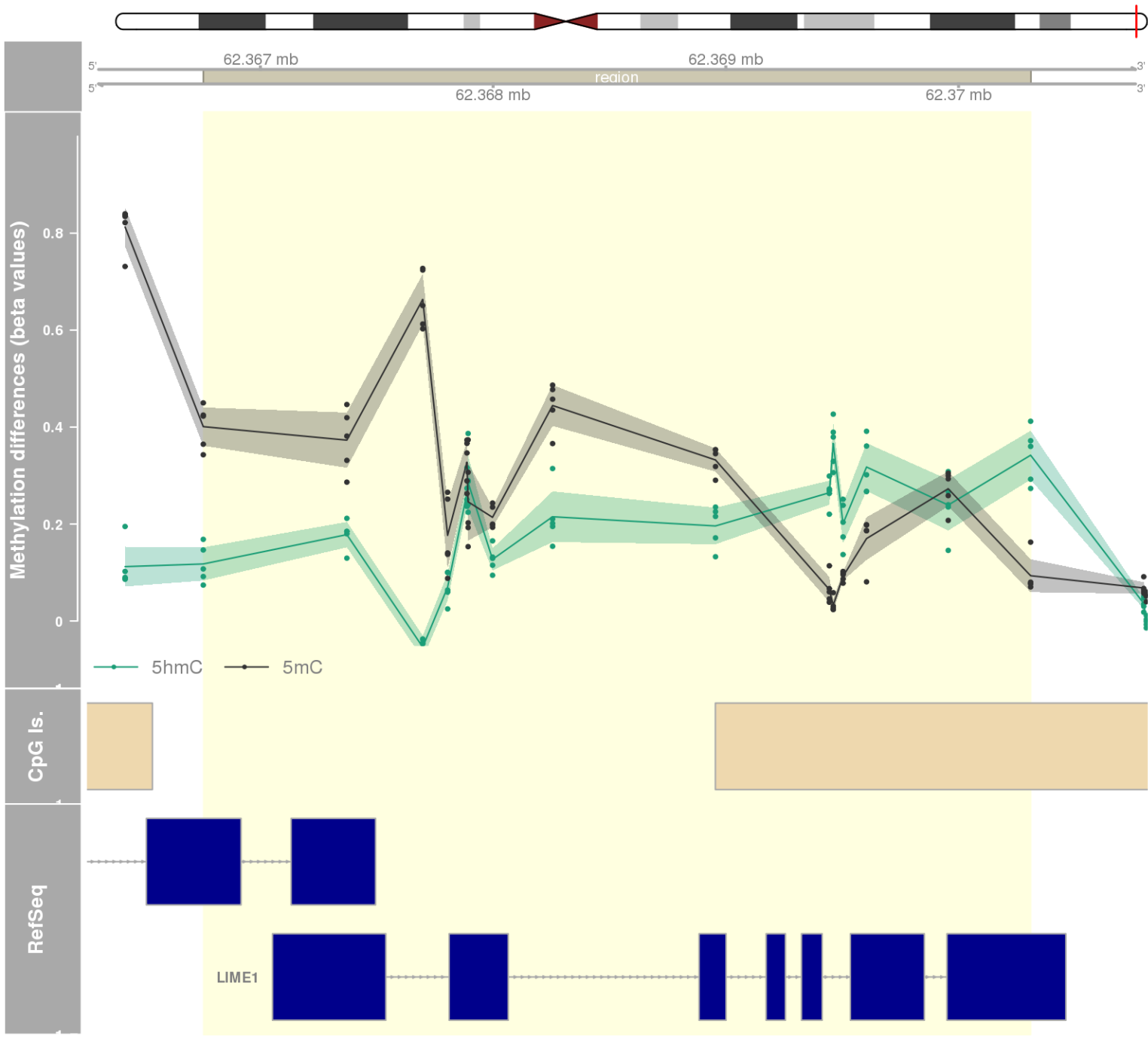
DMR 42 // chr3:148804550-148805181 // 631 pb. (14 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.523
- genes: HLTF-AS1 -



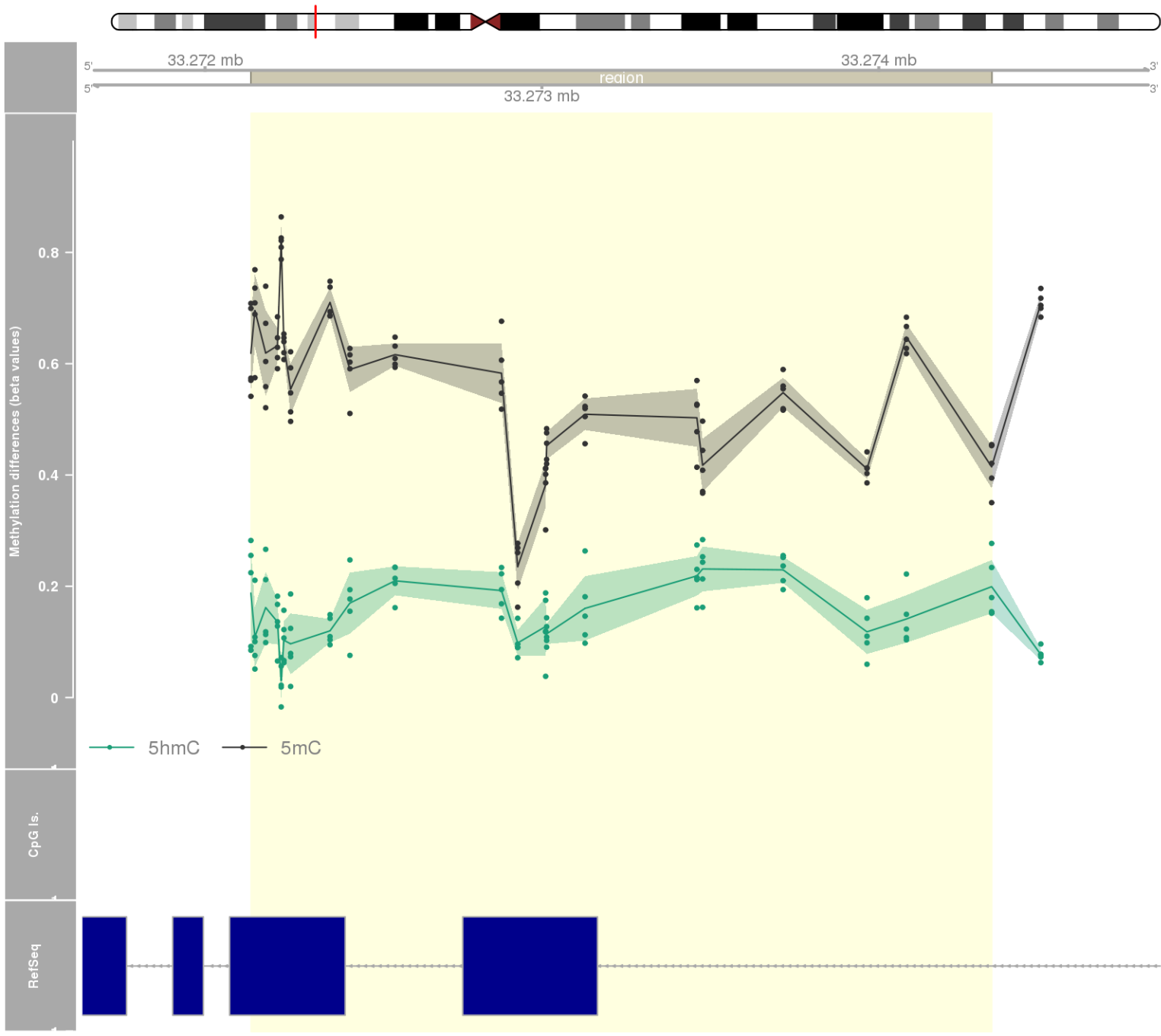
DMR 43 // chr17:62775415-62778370 // 2955 pb. (18 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.523
- genes: PLEKHM1P -



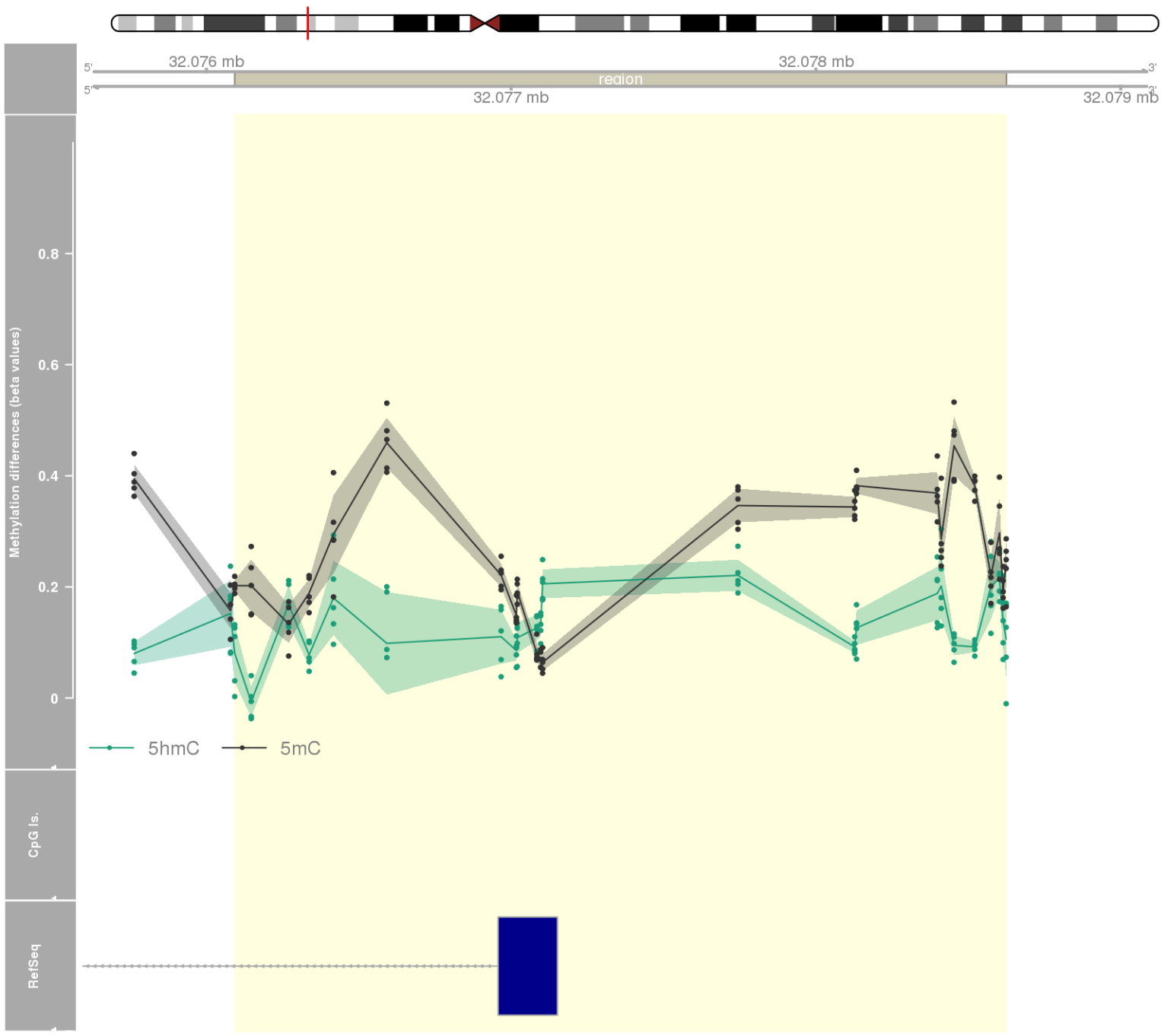
DMR 44 // chr20:62366755-62370310 // 3555 pb. (15 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.523
- genes: ZGPAT / LIME1 -



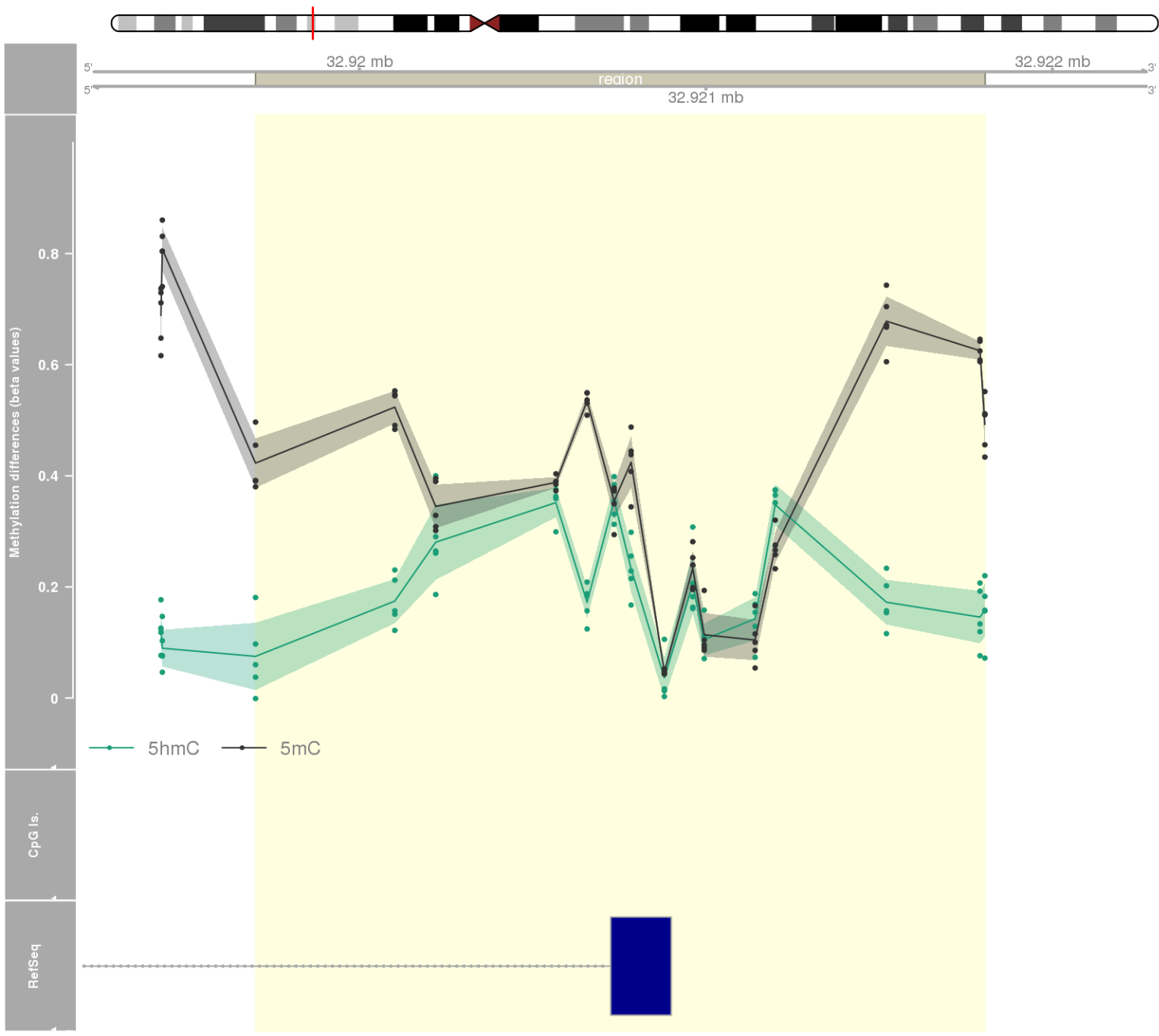
DMR 45 // chr6:33272136-33274335 // 2199 pb. (21 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.523
- genes: TAPBP -



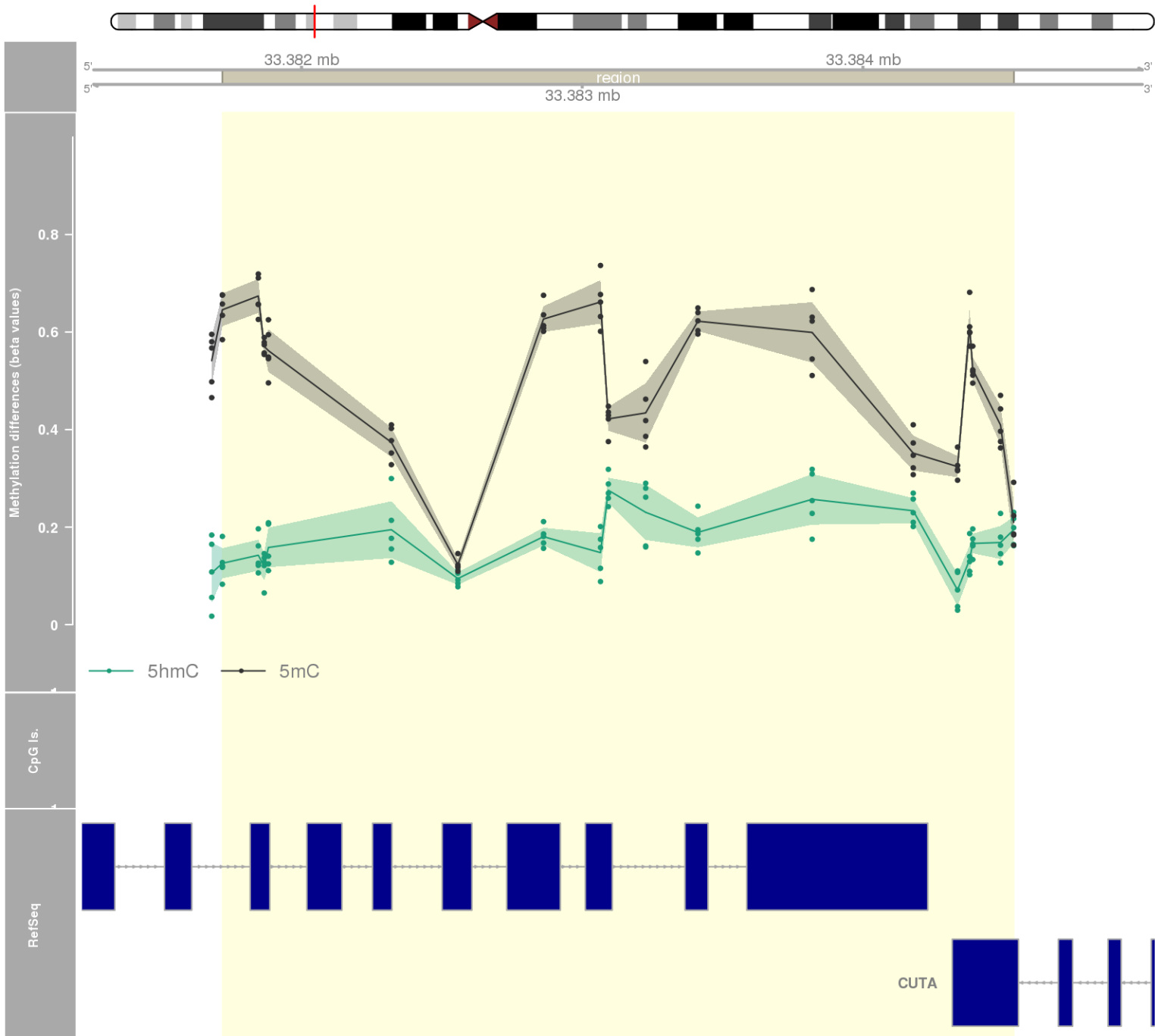
DMR 46 // chr6:32076093-32078624 // 2531 pb. (23 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.523
- genes: TNXB / ATF6B -



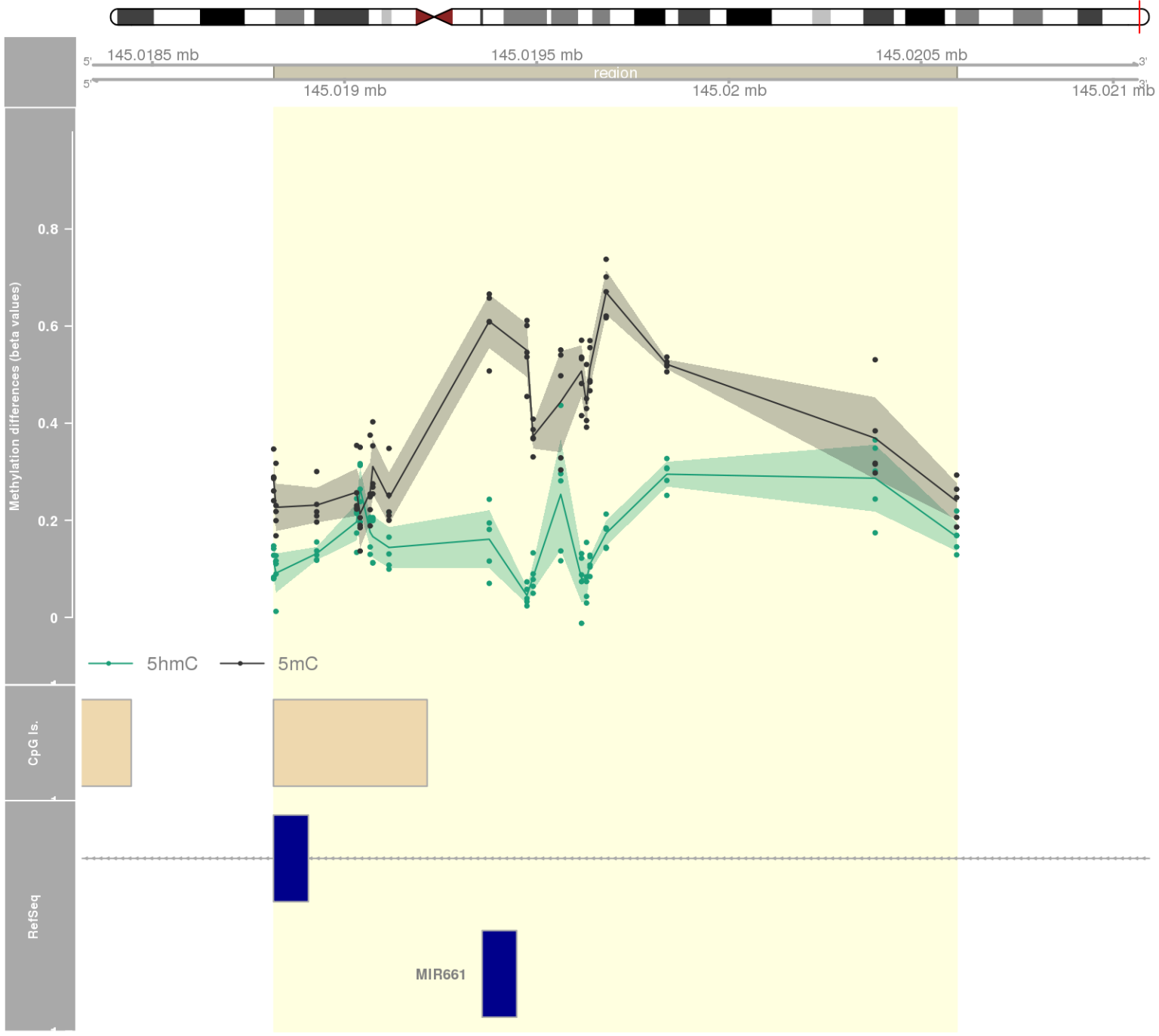
DMR 47 // chr6:32919700-32921805 // 2105 pb. (15 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.523
- genes: HLA-DMA -



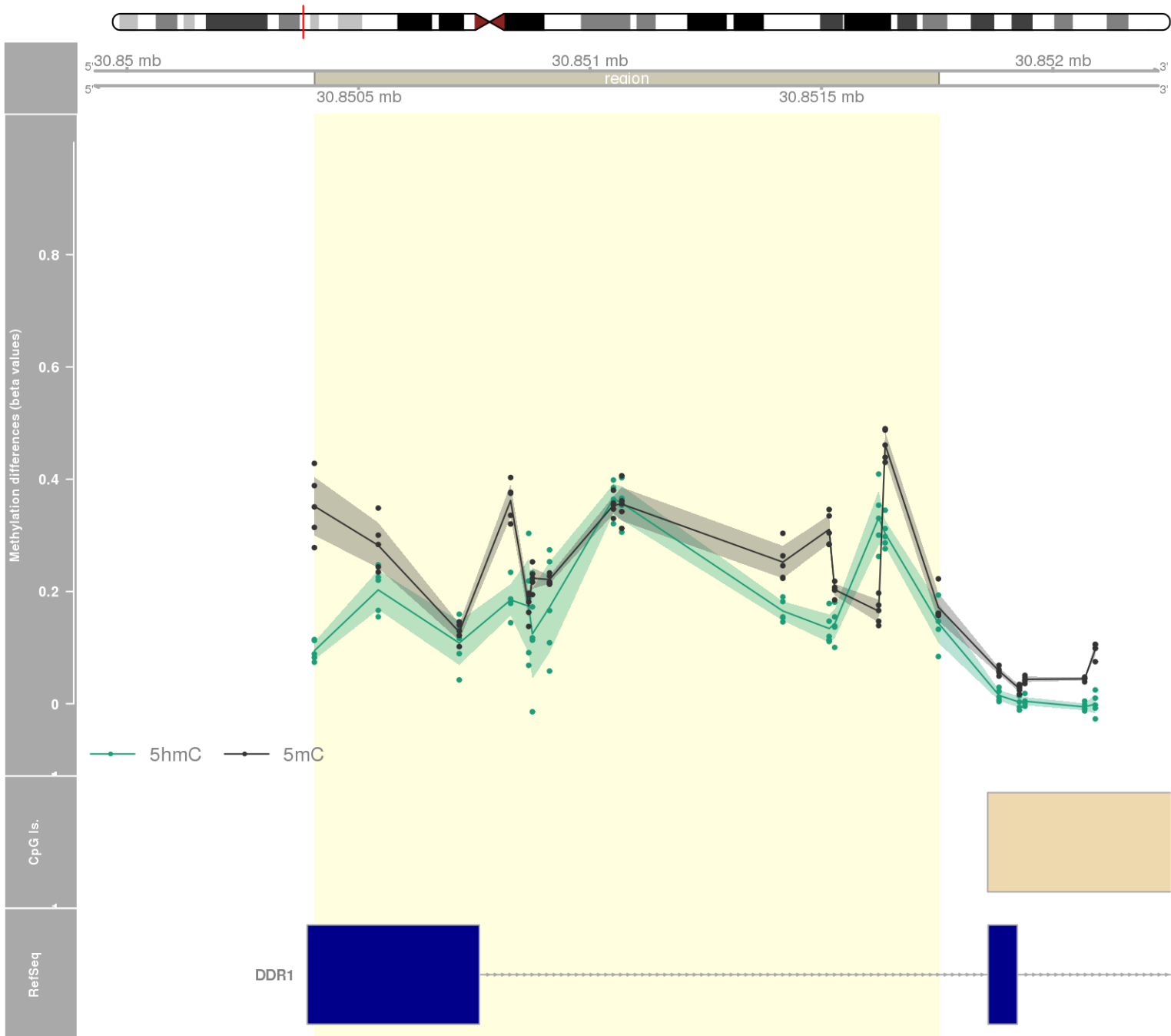
DMR 48 // chr6:33381718-33384537 // 2819 pb. (18 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.523
- genes: PHF1 / CUTA -



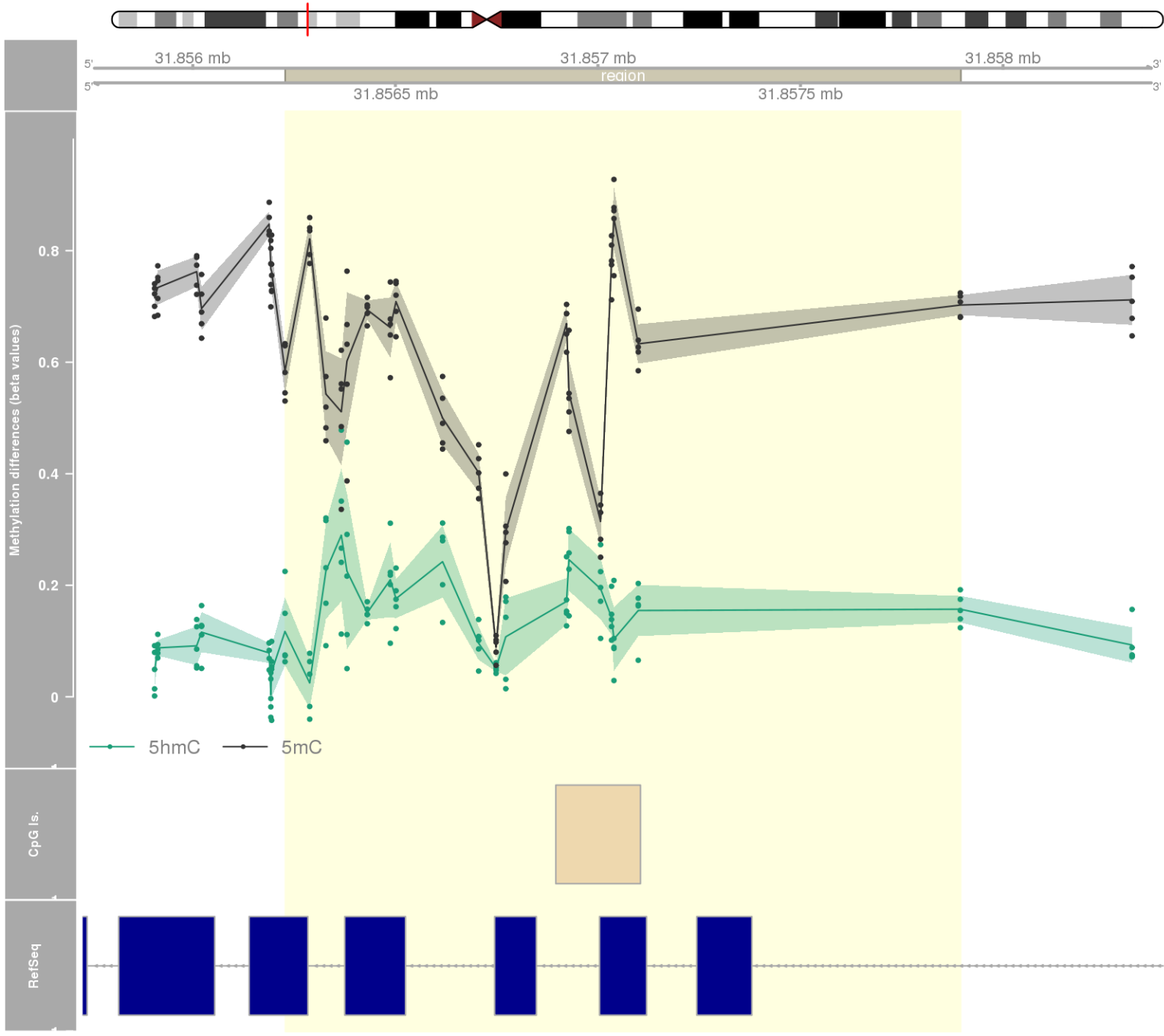
DMR 49 // chr8:145018816-145020593 // 1777 pb. (19 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.523
- genes: MIR661 / PLEC -



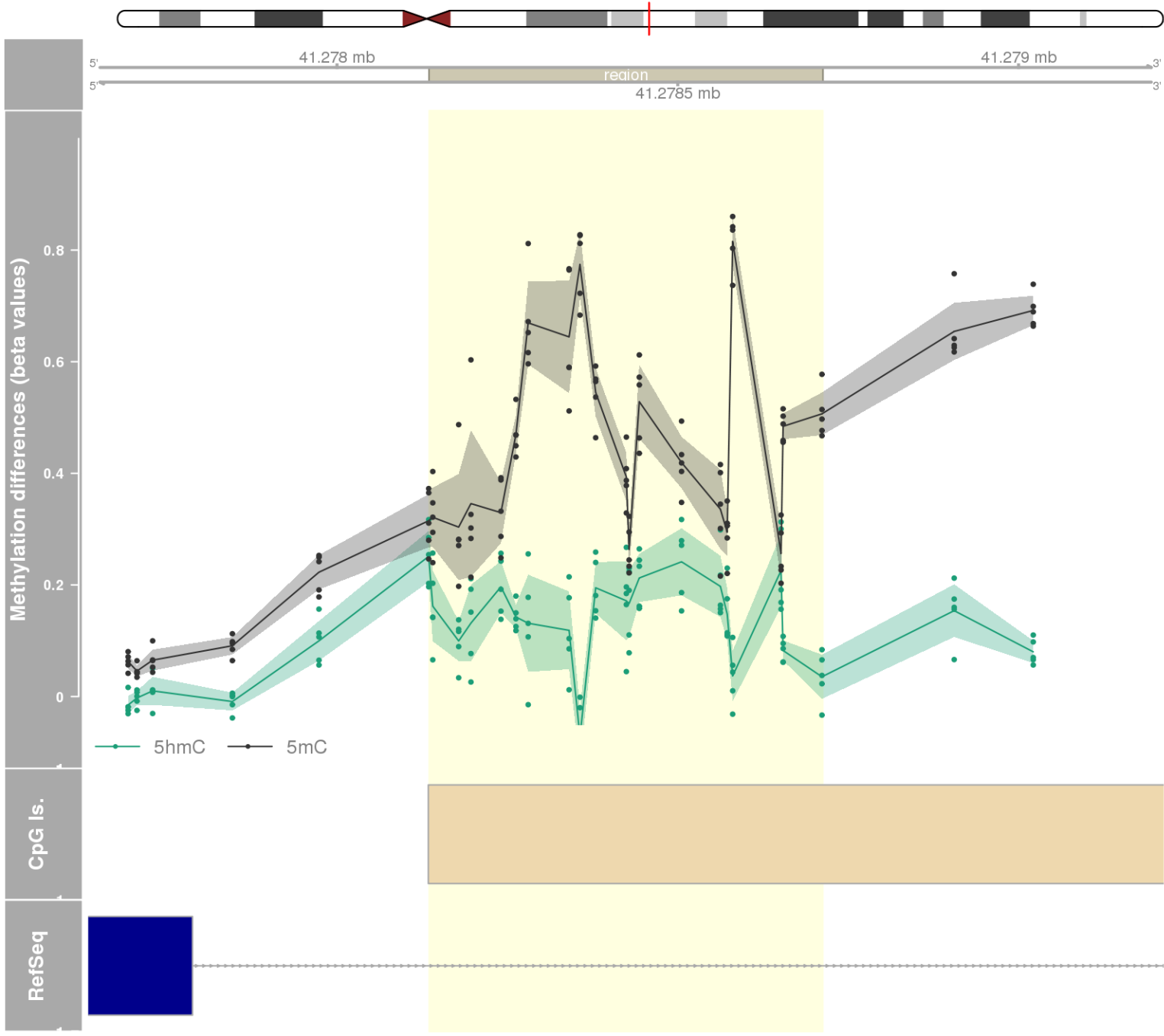
DMR 50 // chr6:30850405-30851753 // 1348 pb. (15 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.003 // fwerArea: 0.523
- genes: DDR1 -



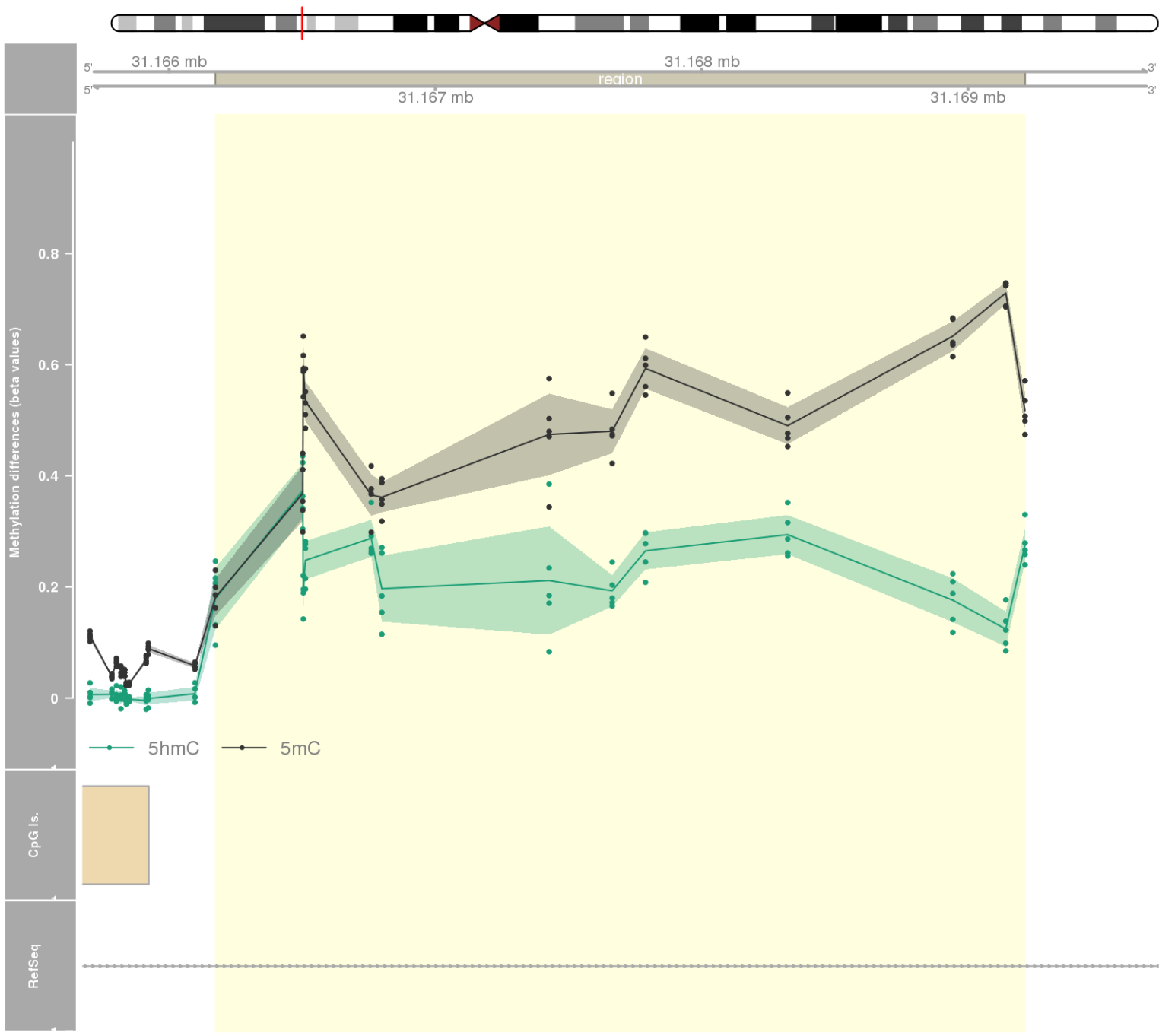
DMR 51 // chr6:31856228-31857895 // 1667 pb. (19 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.004 // fwerArea: 0.523
- genes: EHMT2 -



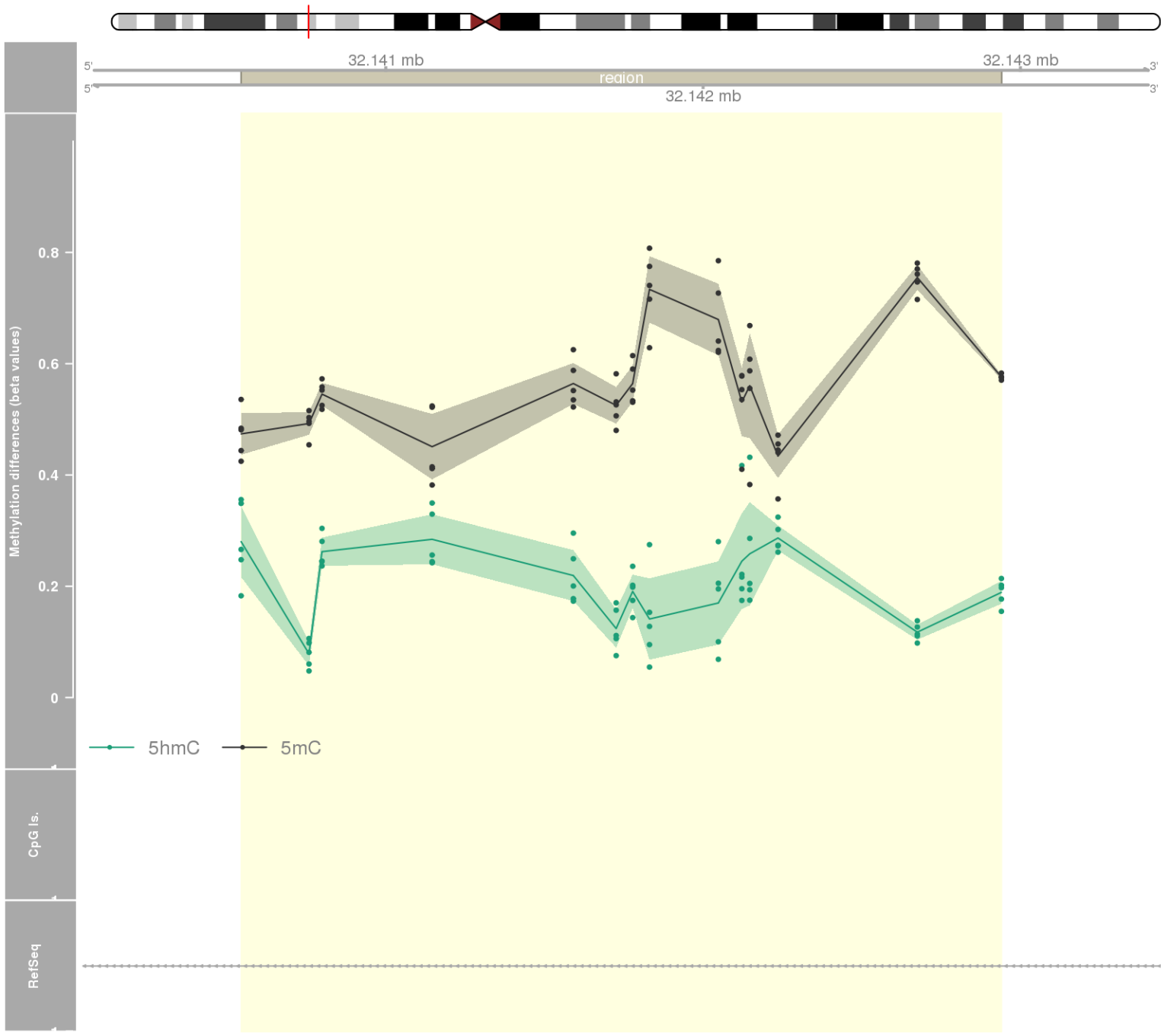
DMR 52 // chr17:41278135-41278712 // 577 pb. (20 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.004 // fwerArea: 0.523
- genes: NBR2 -



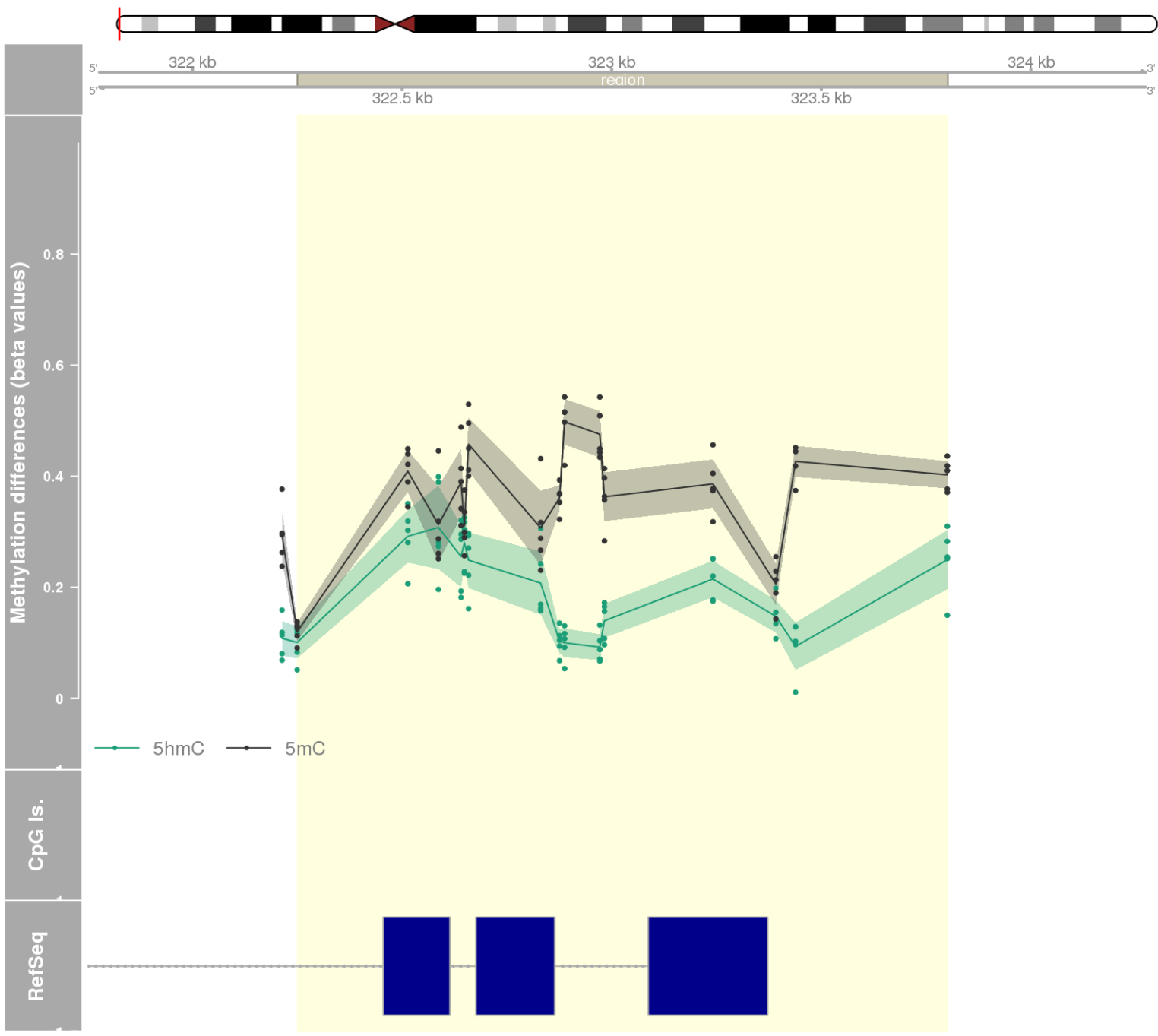
DMR 53 // chr6:31166174-31169214 // 3040 pb. (13 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.004 // fwerArea: 0.523
- genes: HCG27 -



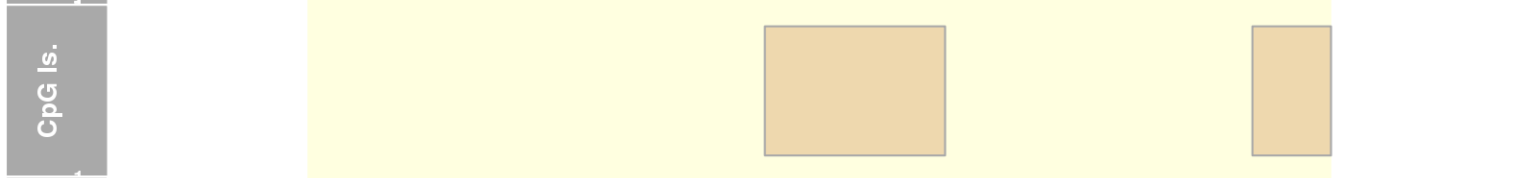
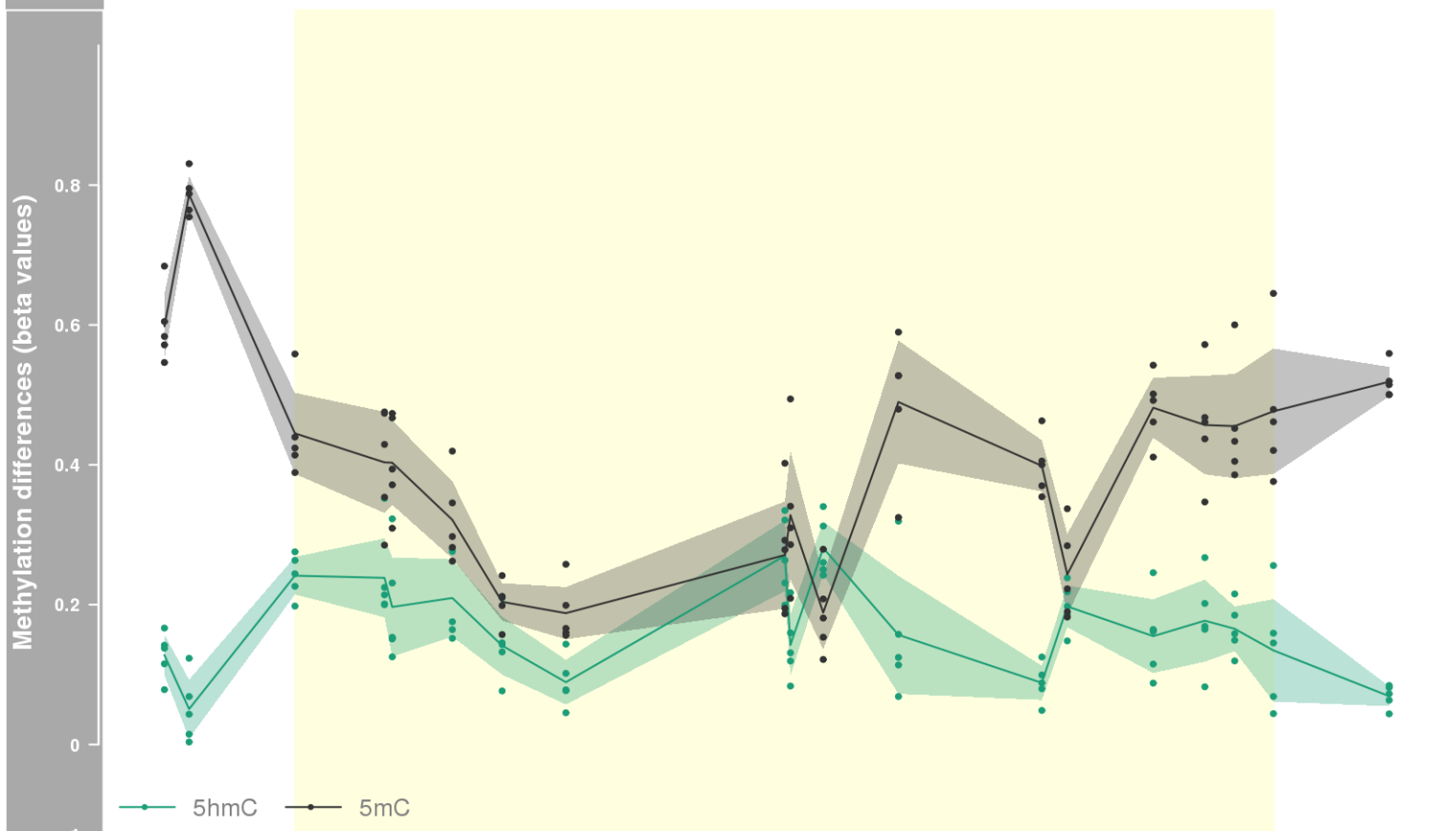
DMR 54 // chr6:32140544-32142940 // 2396 pb. (14 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.004 // fwerArea: 0.523
- genes: AGPAT1 -



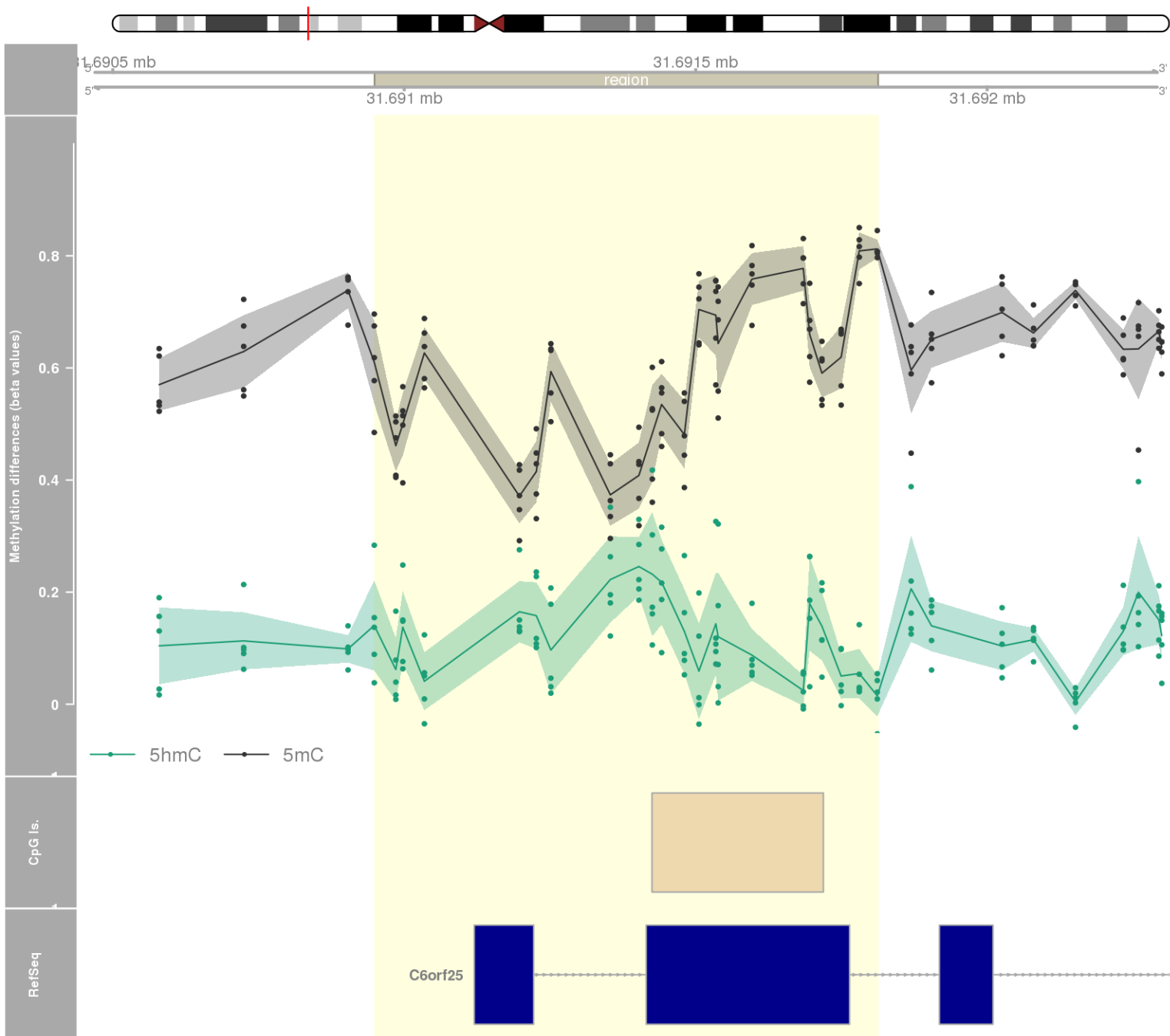
DMR 55 // chr12:322250-323801 // 1551 pb. (15 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.004 // fwerArea: 0.523
- genes: SLC6A12 -



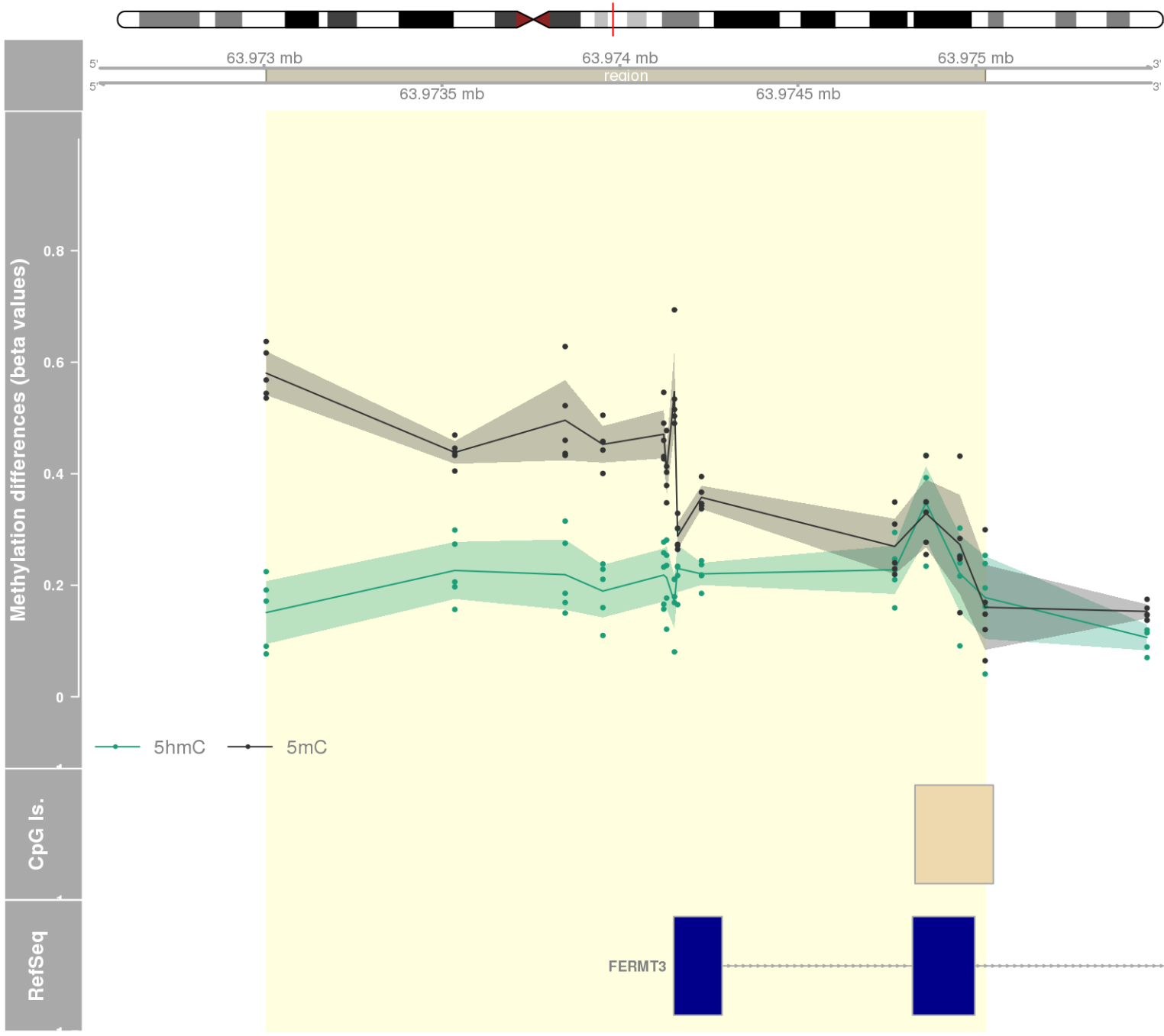
DMR 56 // chr11:1296823-1299477 // 2654 pb. (16 probes) // pvalue: 0.002 // fwer: 0.36 // pvalueArea: 0.004 // fwerArea: 0.523
- genes: TOLLIP -



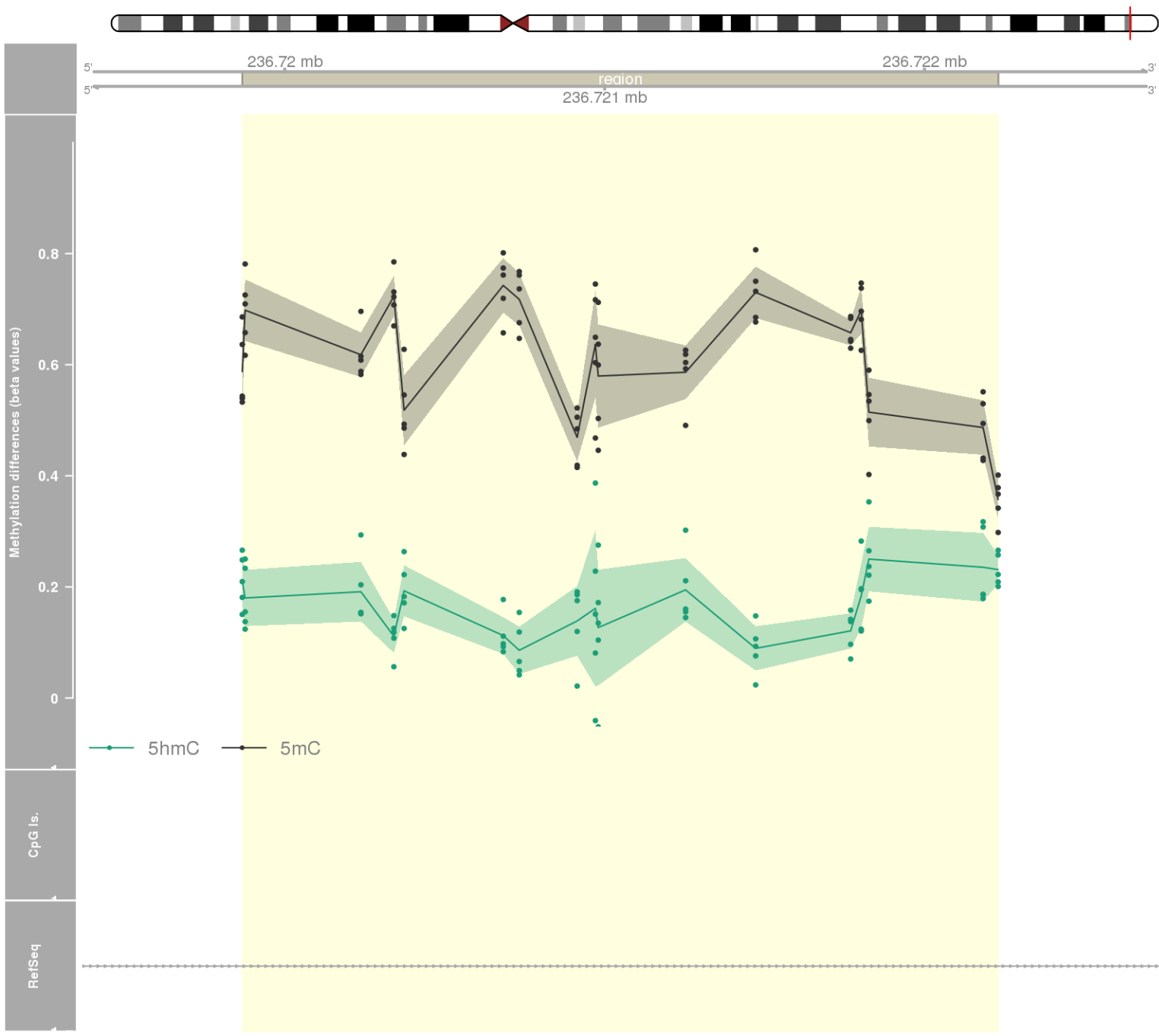
DMR 57 // chr6:31690949-31691812 // 863 pb. (22 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.004 // fwerArea: 0.523
- genes: C6orf25 -



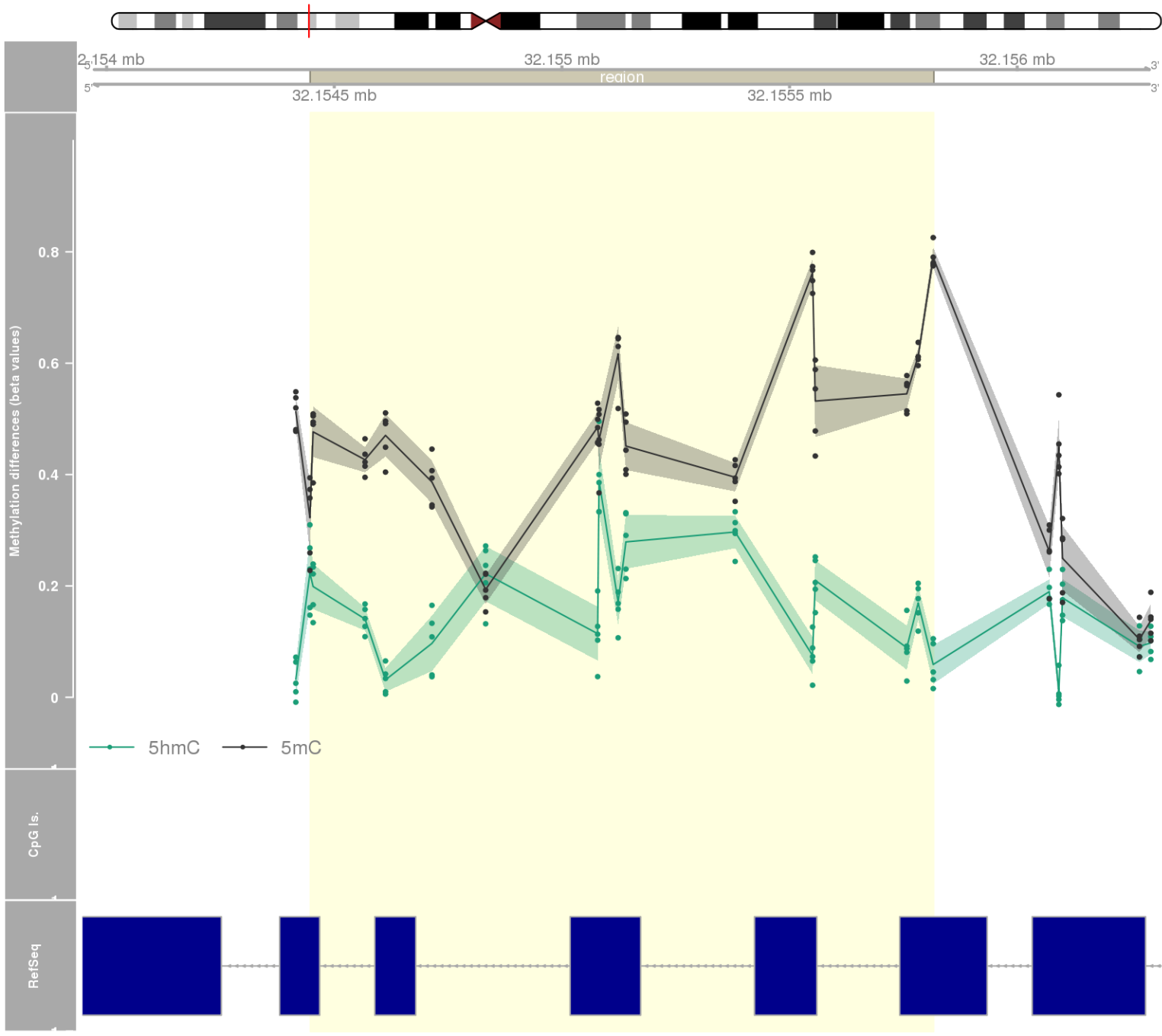
DMR 58 // chr11:63973006-63975026 // 2020 pb. (13 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.004 // fwerArea: 0.684
- genes: FERMT3 -



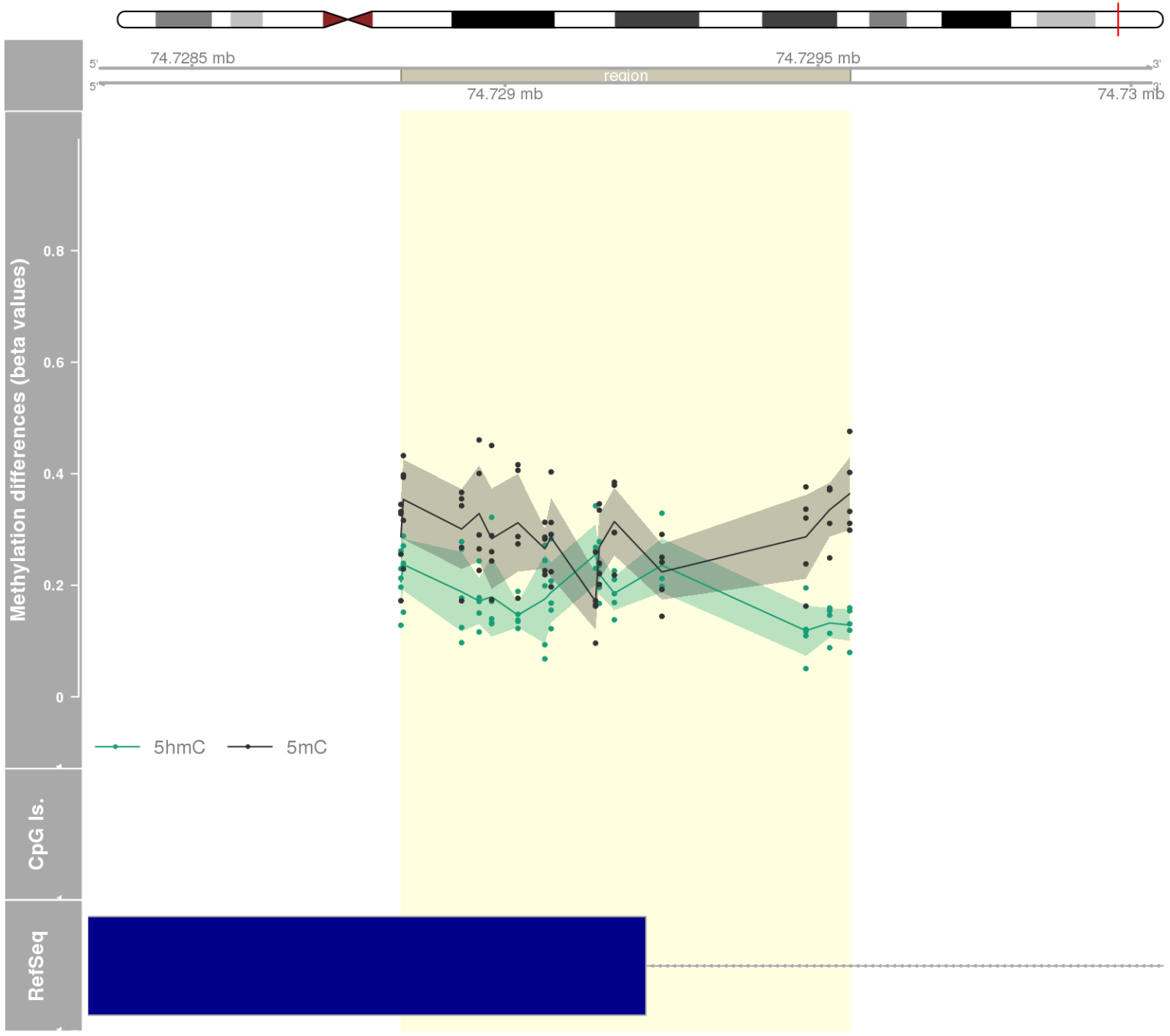
DMR 59 // chr2:236719867-236722230 // 2363 pb. (17 probes) // pvalue: 0.002 // fwer: 0.36 // pvalueArea: 0.004 // fwerArea: 0.684
- genes: AGAP1 -



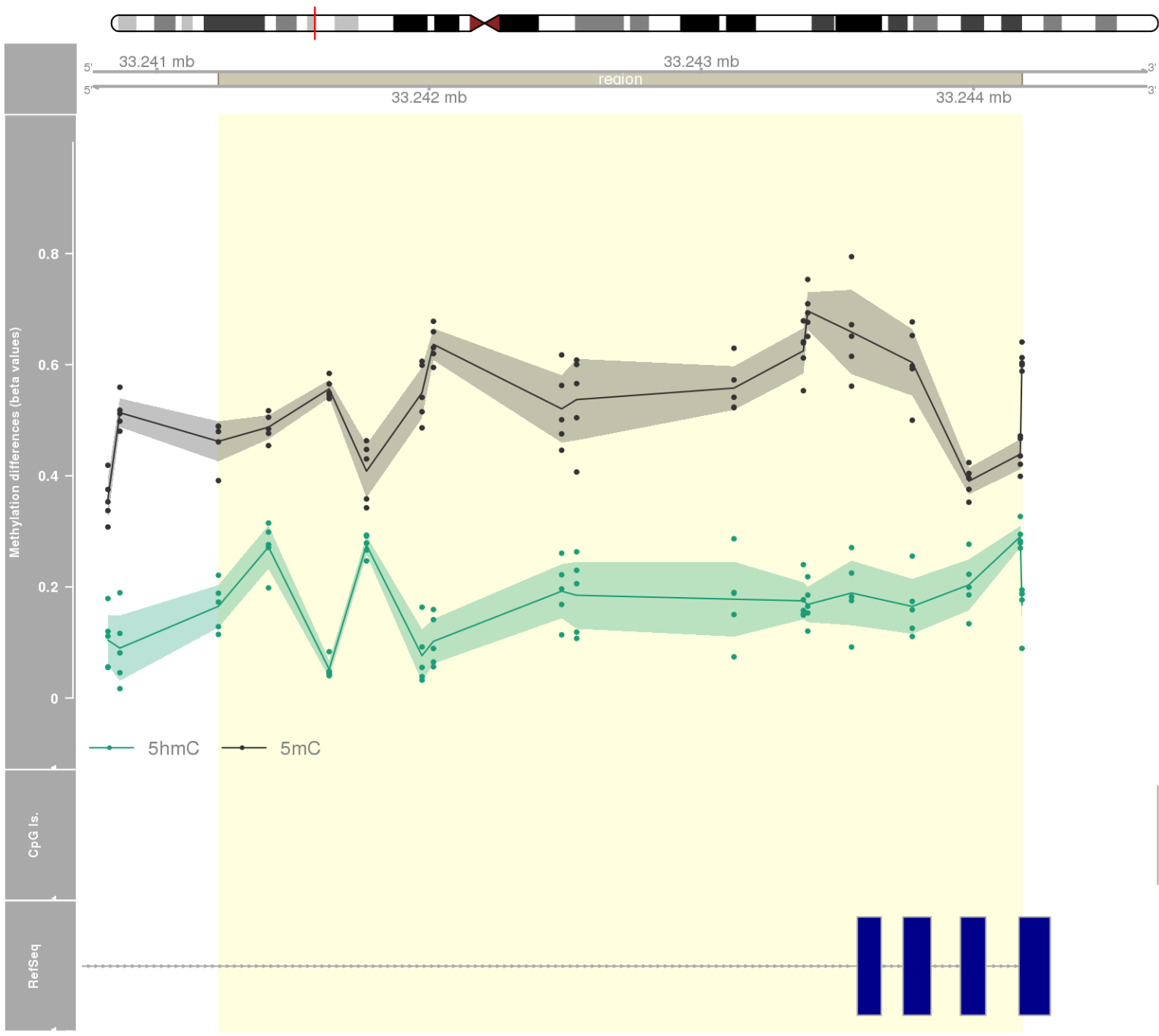
DMR 60 // chr6:32154447-32155816 // 1369 pb. (16 probes) // pvalue: 0.002 // fwer: 0.36 // pvalueArea: 0.004 // fwerArea: 0.684
- genes: PBX2 -



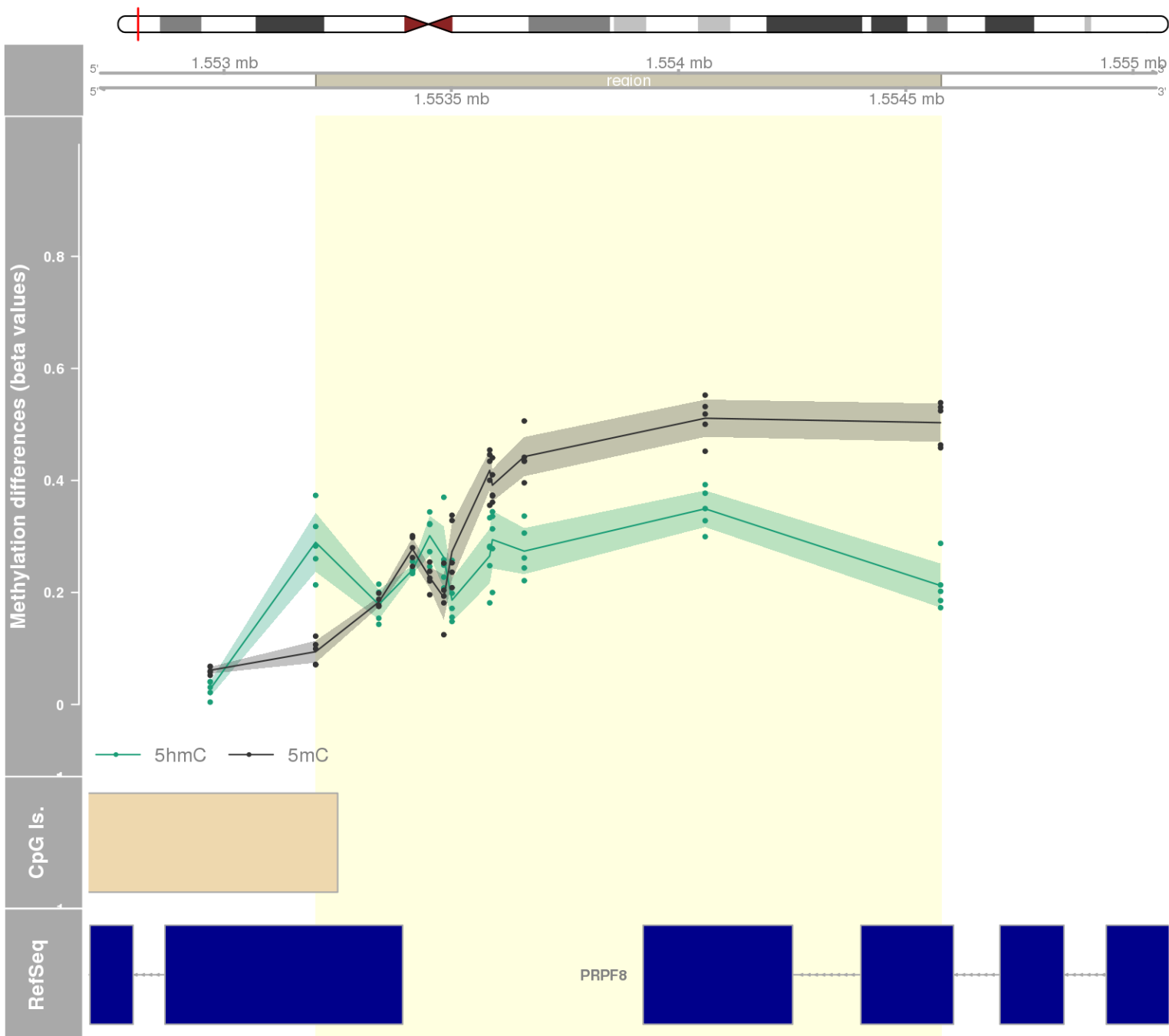
DMR 61 // chr18:74728834-74729551 // 717 pb. (15 probes) // pvalue: 0.002 // fwer: 0.36 // pvalueArea: 0.004 // fwerArea: 0.684
- genes: MBP -



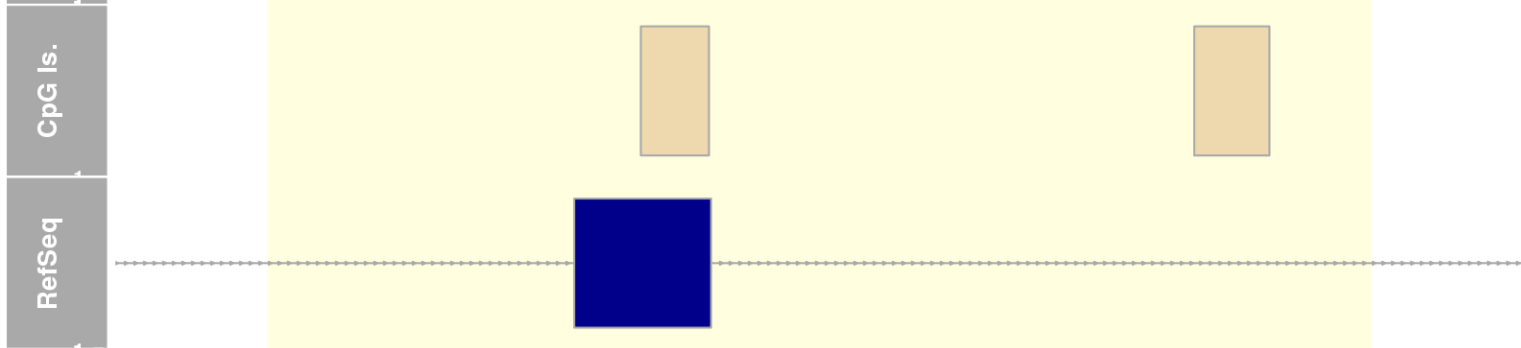
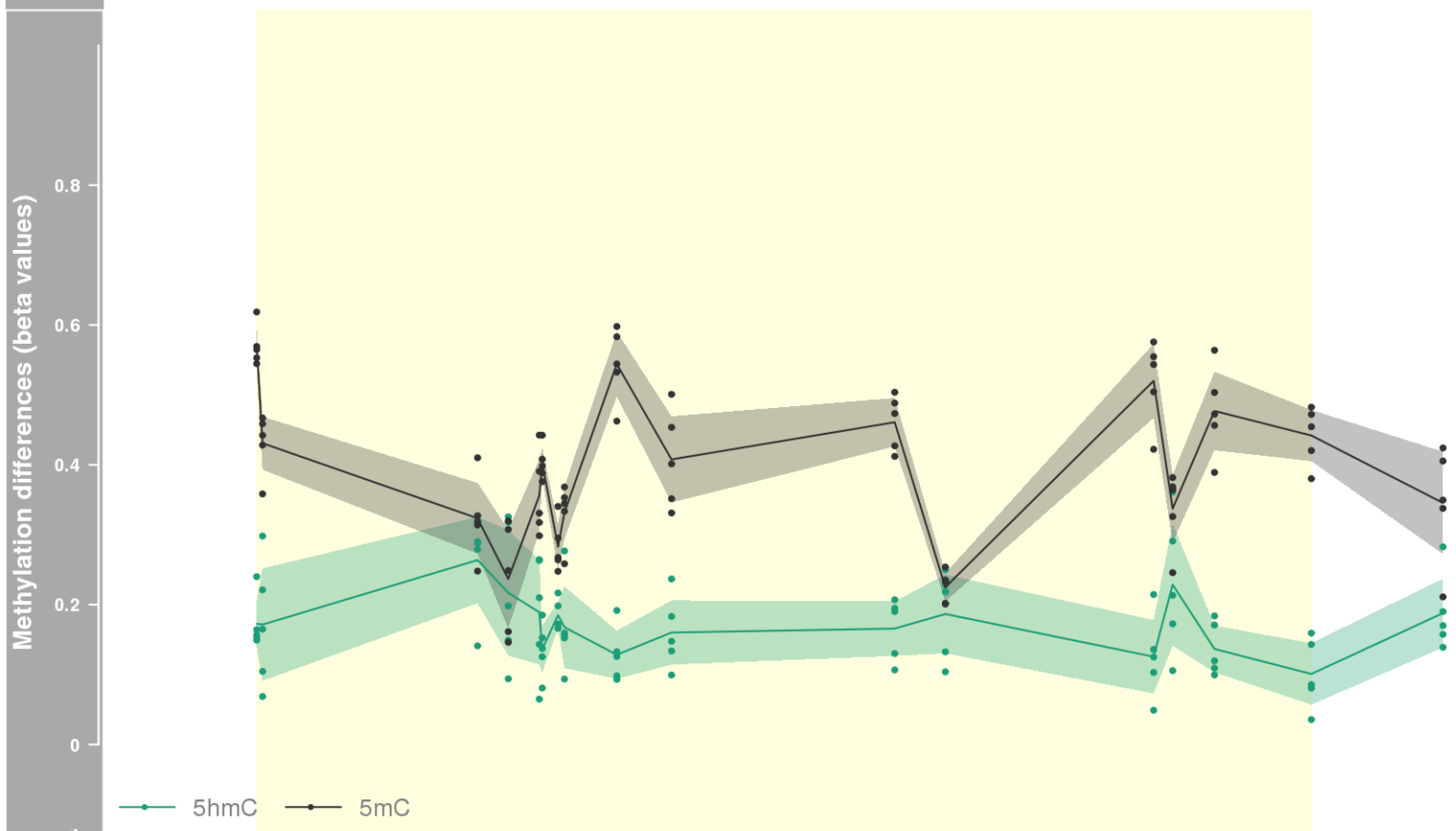
DMR 62 // chr6:33241226-33244178 // 2952 pb. (16 probes) // pvalue: 0.002 // fwer: 0.36 // pvalueArea: 0.004 // fwerArea: 0.684
- genes: RPS18 -



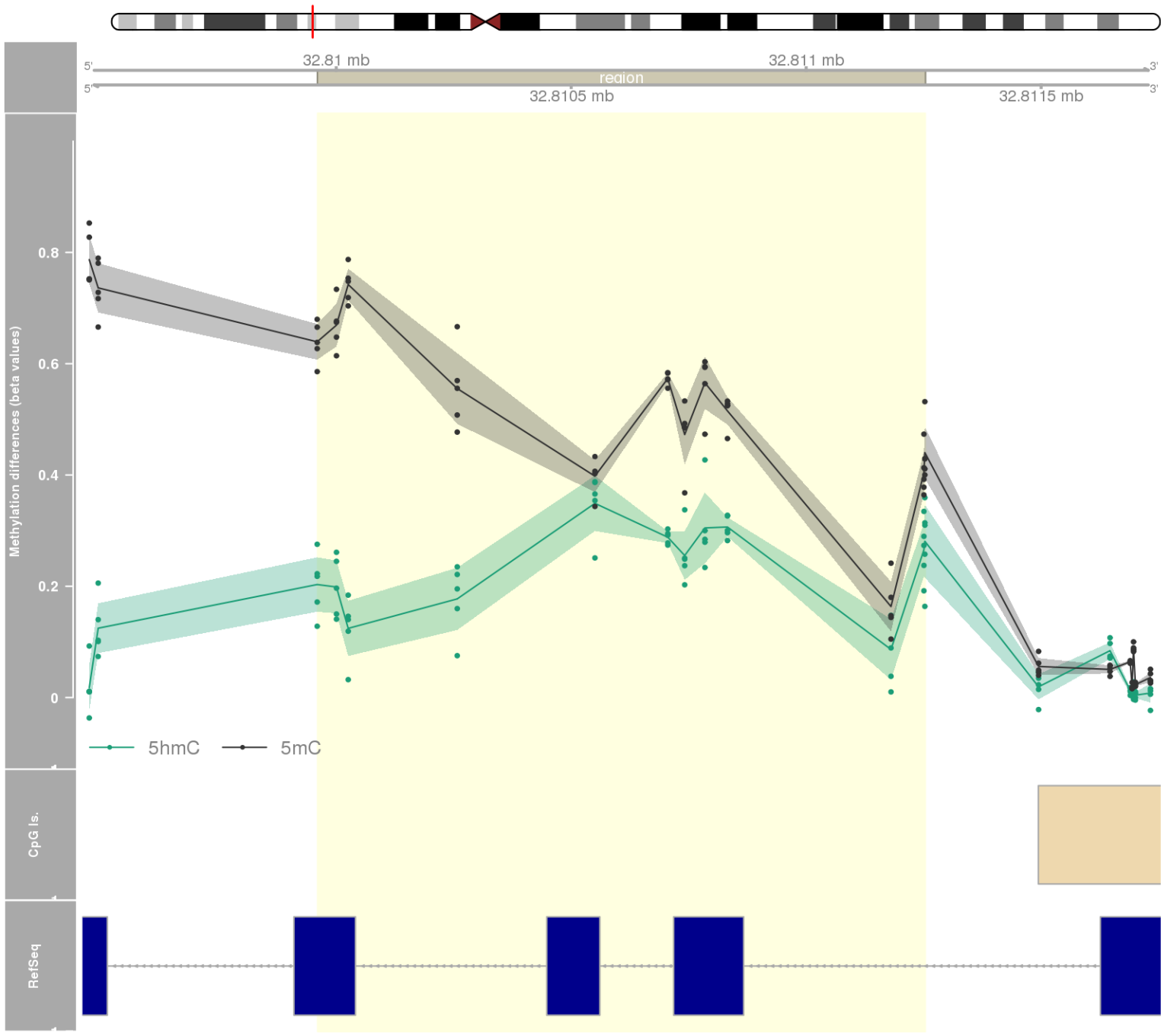
DMR 63 // chr17:1553202-1554577 // 1375 pb. (11 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.004 // fwerArea: 0.684
- genes: RILP / PRPF8 -



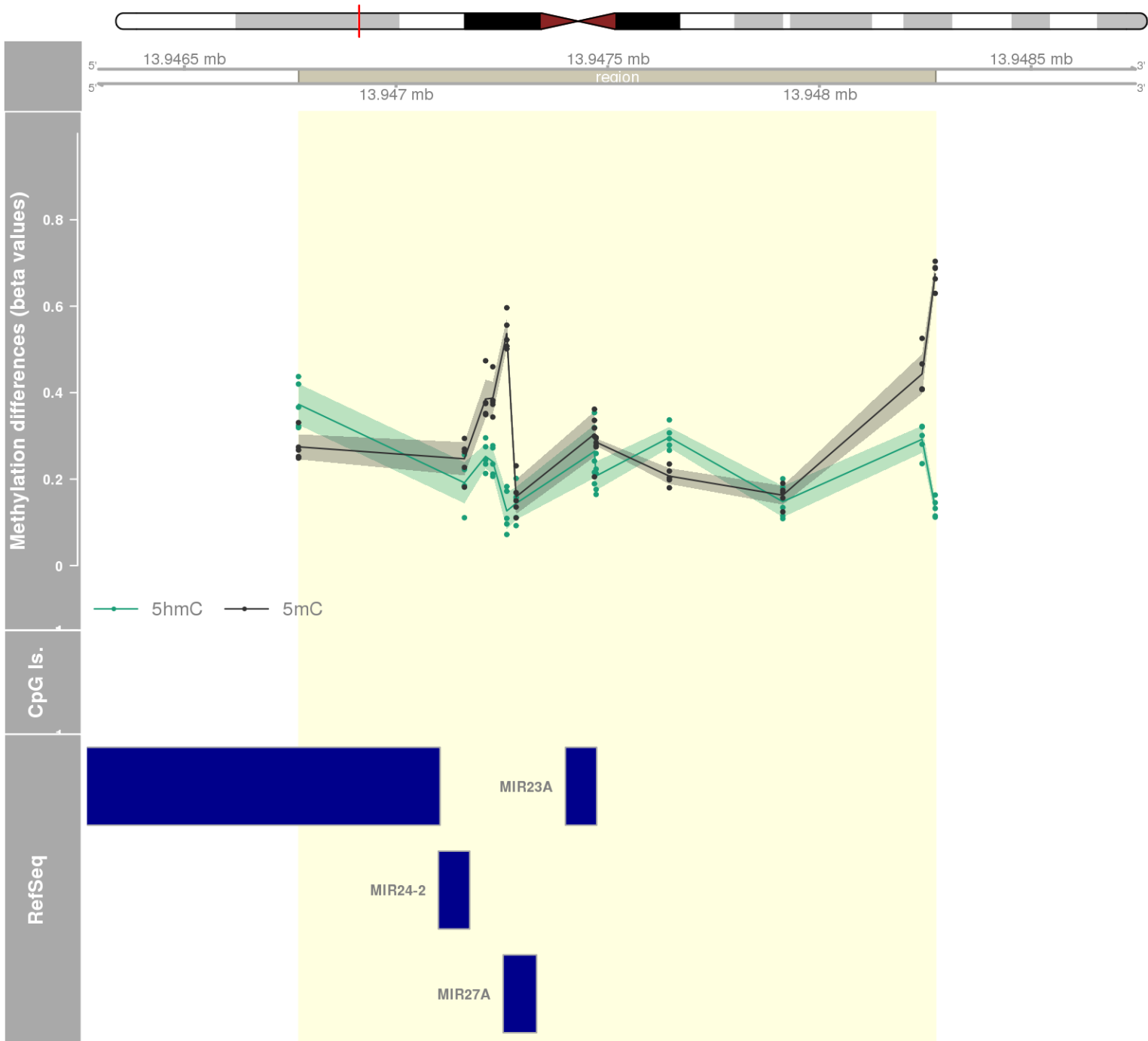
DMR 64 // chr13:113632624-113636229 // 3605 pb. (16 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: MCF2L -



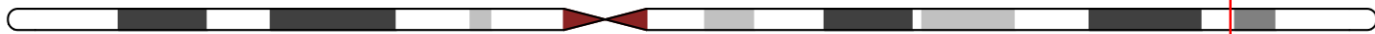
DMR 65 // chr6:32809960-32811253 // 1293 pb. (12 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: PSMB8 -



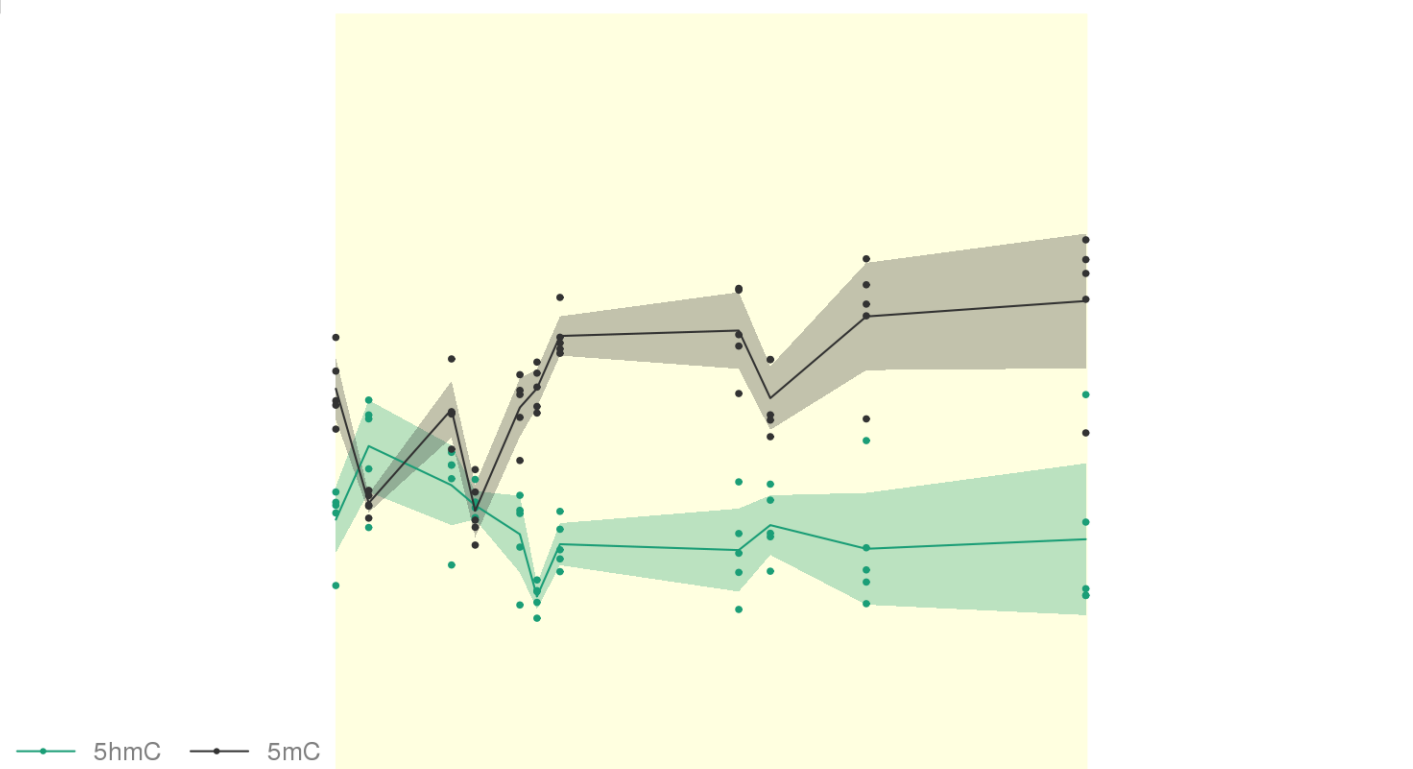
DMR 66 // chr19:13946770-13948274 // 1504 pb. (12 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: MIR24-2 / MIR27A / MIR23A -



DMR 67 // chr20:56286391-56287532 // 1141 pb. (11 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: PMEPA1 -



Methylation differences (beta values)



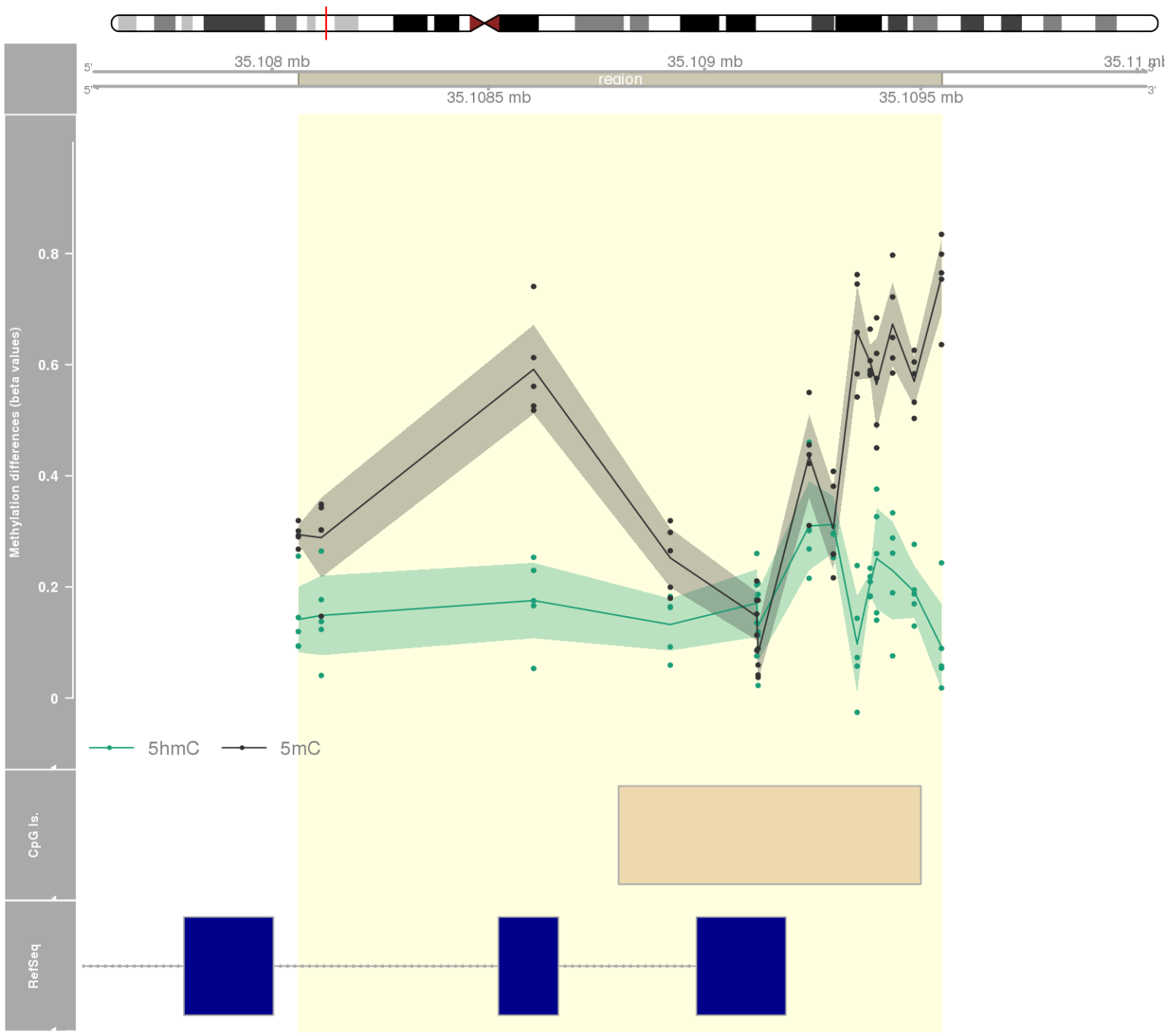
—●— 5hmC —●— 5mC

CpG Is.

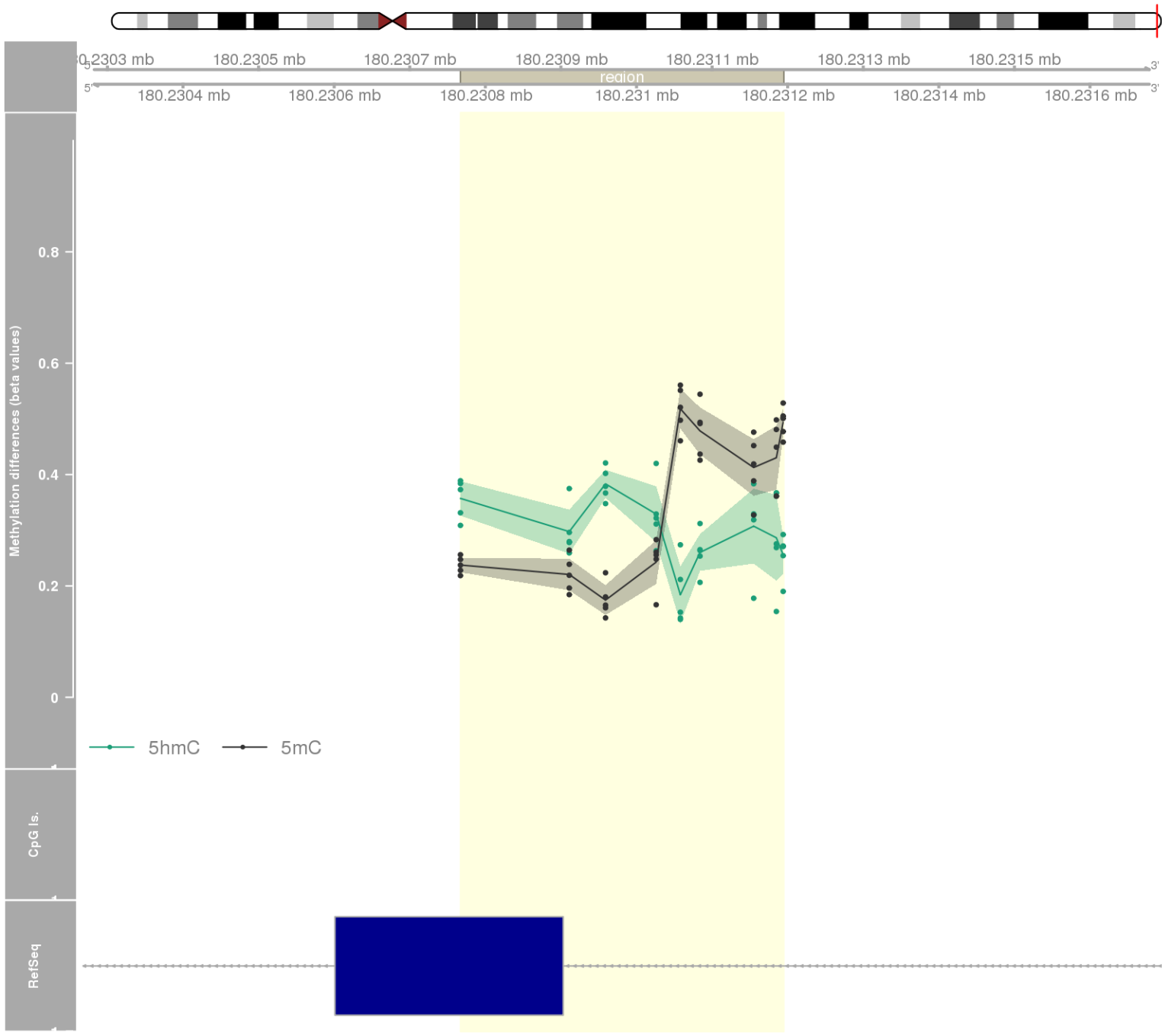
RefSeq



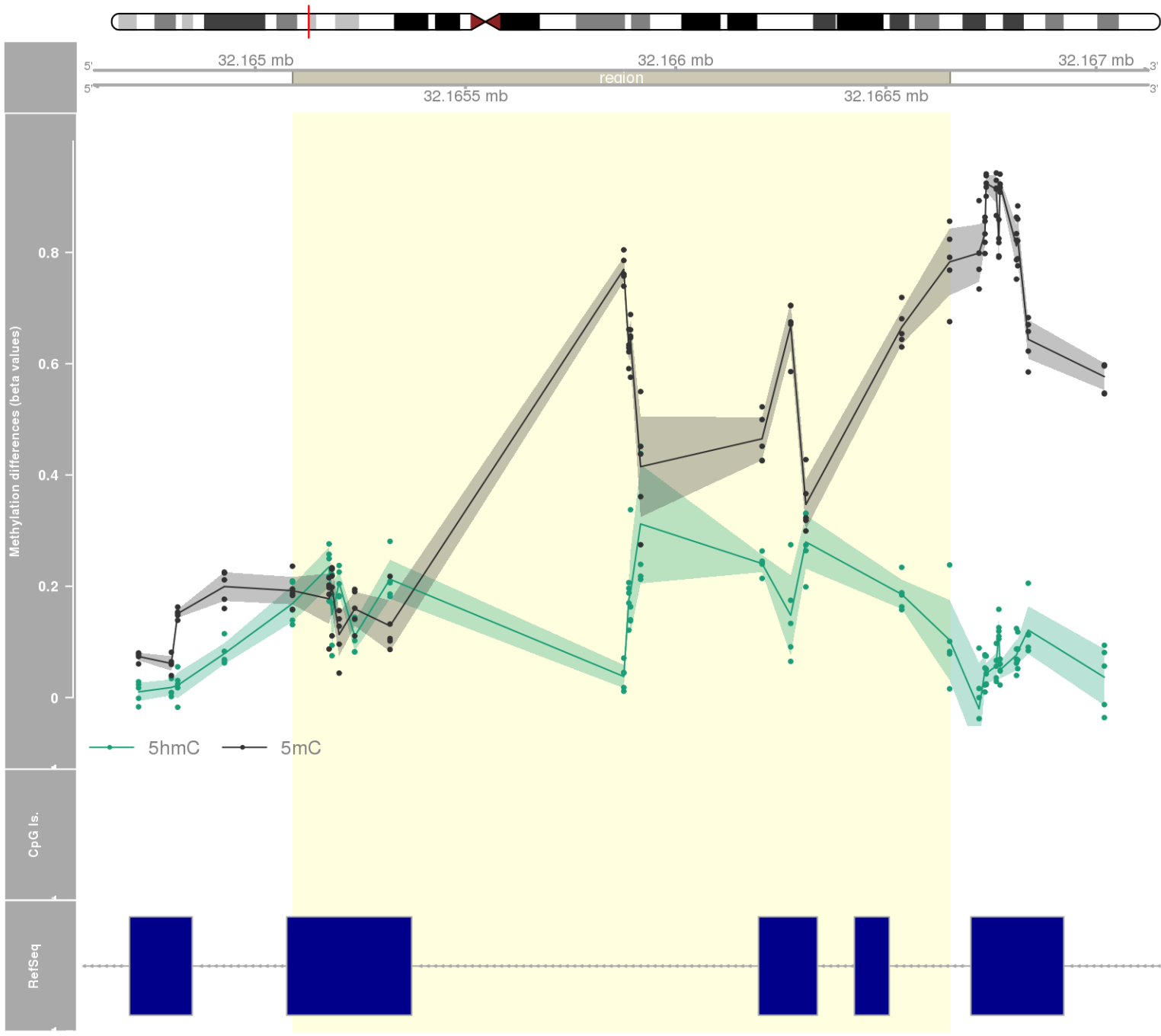
DMR 68 // chr6:35108061-35109548 // 1487 pb. (14 probes) // pvalue: 0.002 // fwer: 0.36 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: TCP11 -



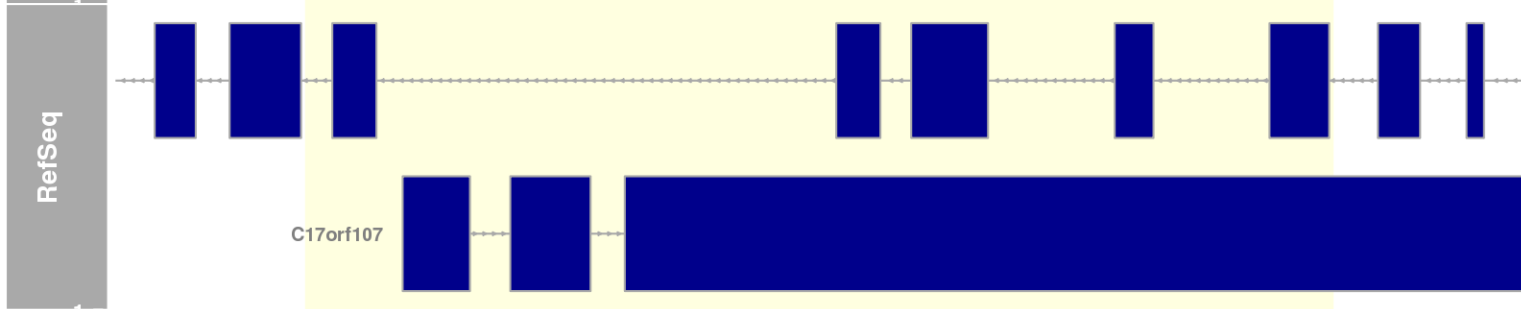
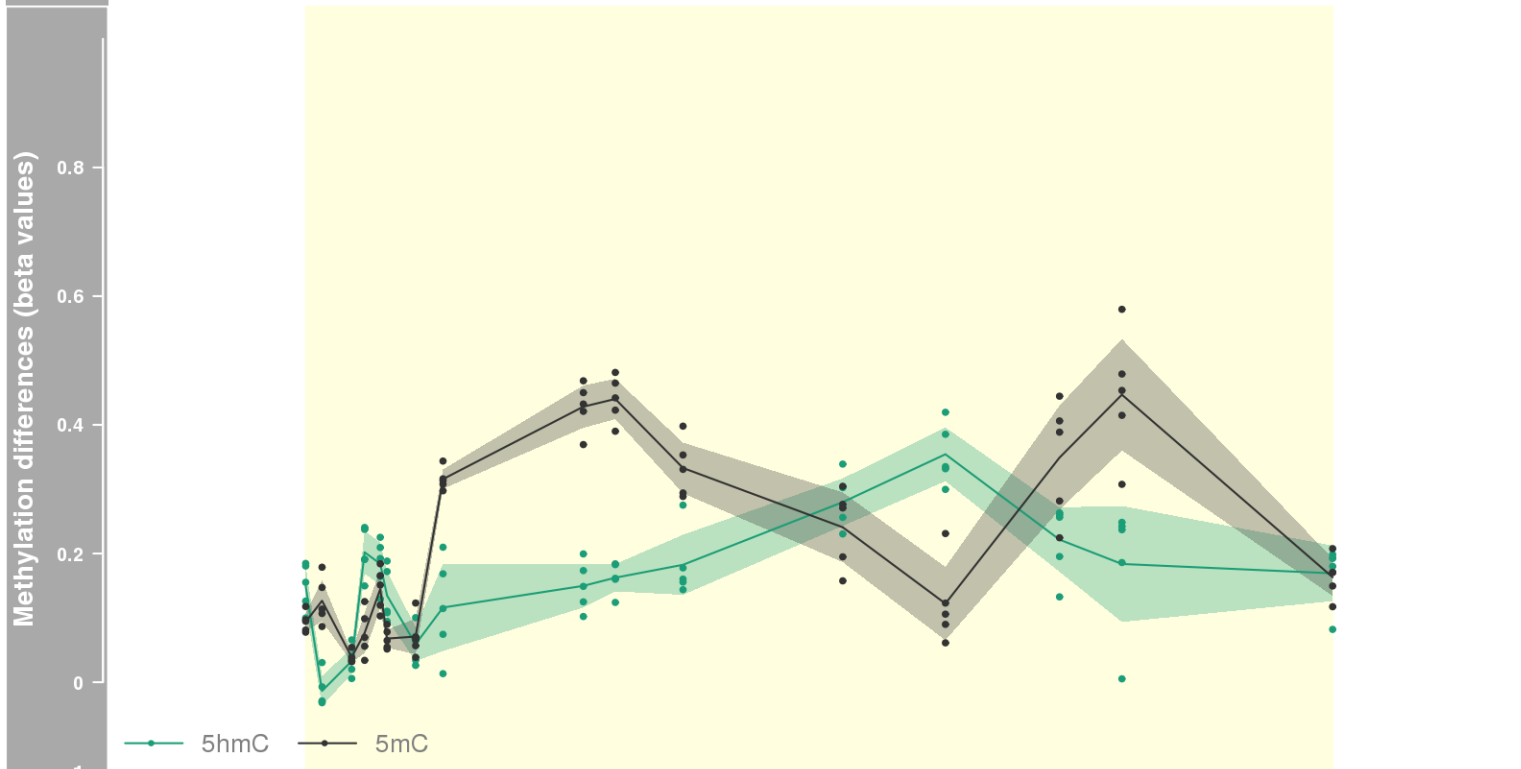
DMR 69 // chr5:180230767-180231194 // 427 pb. (9 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: MGAT1 -



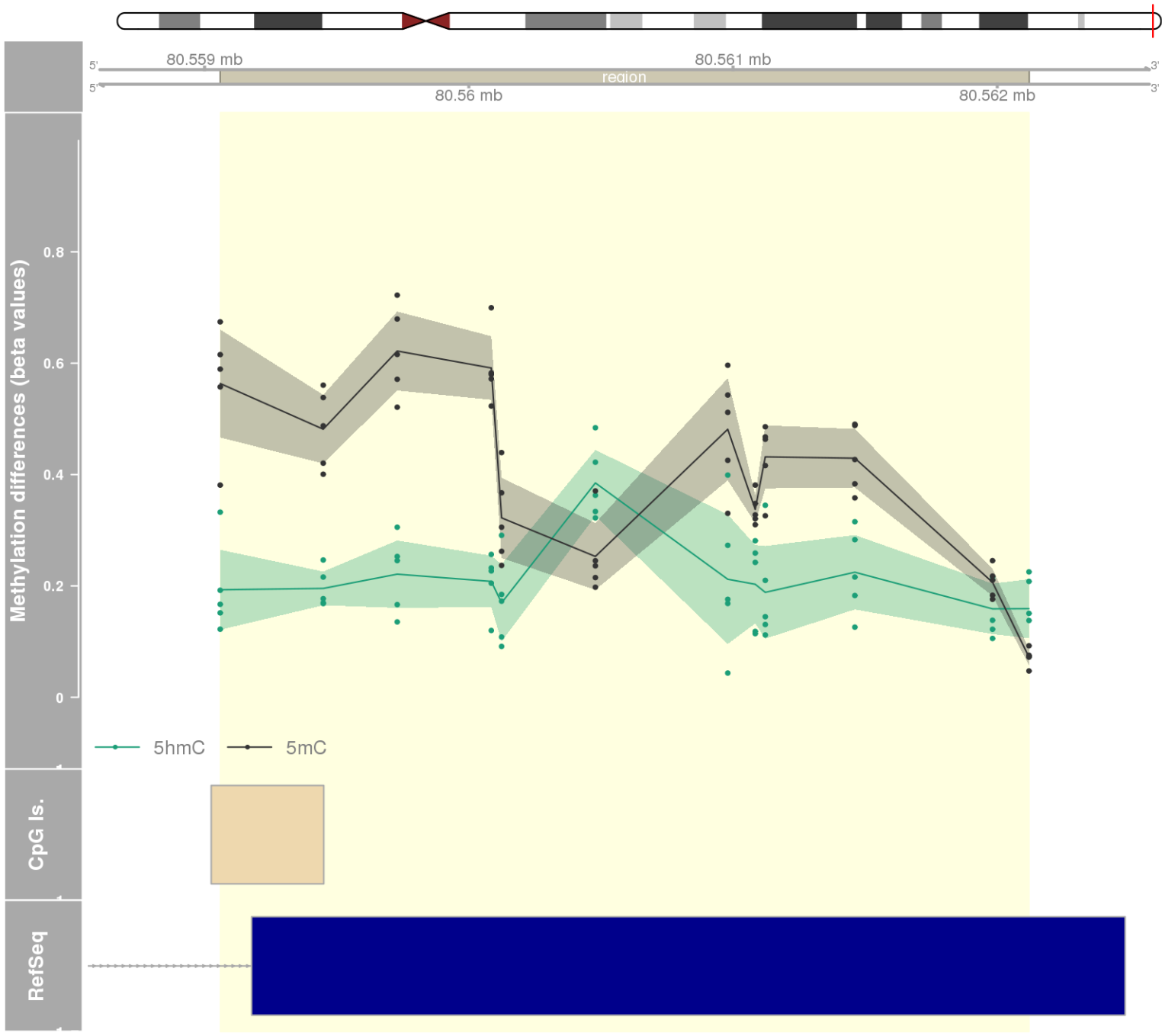
DMR 70 // chr6:32165089-3216652 // 1563 pb. (15 probes) // pvalue: 0.002 // fwer: 0.36 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: NOTCH4 -



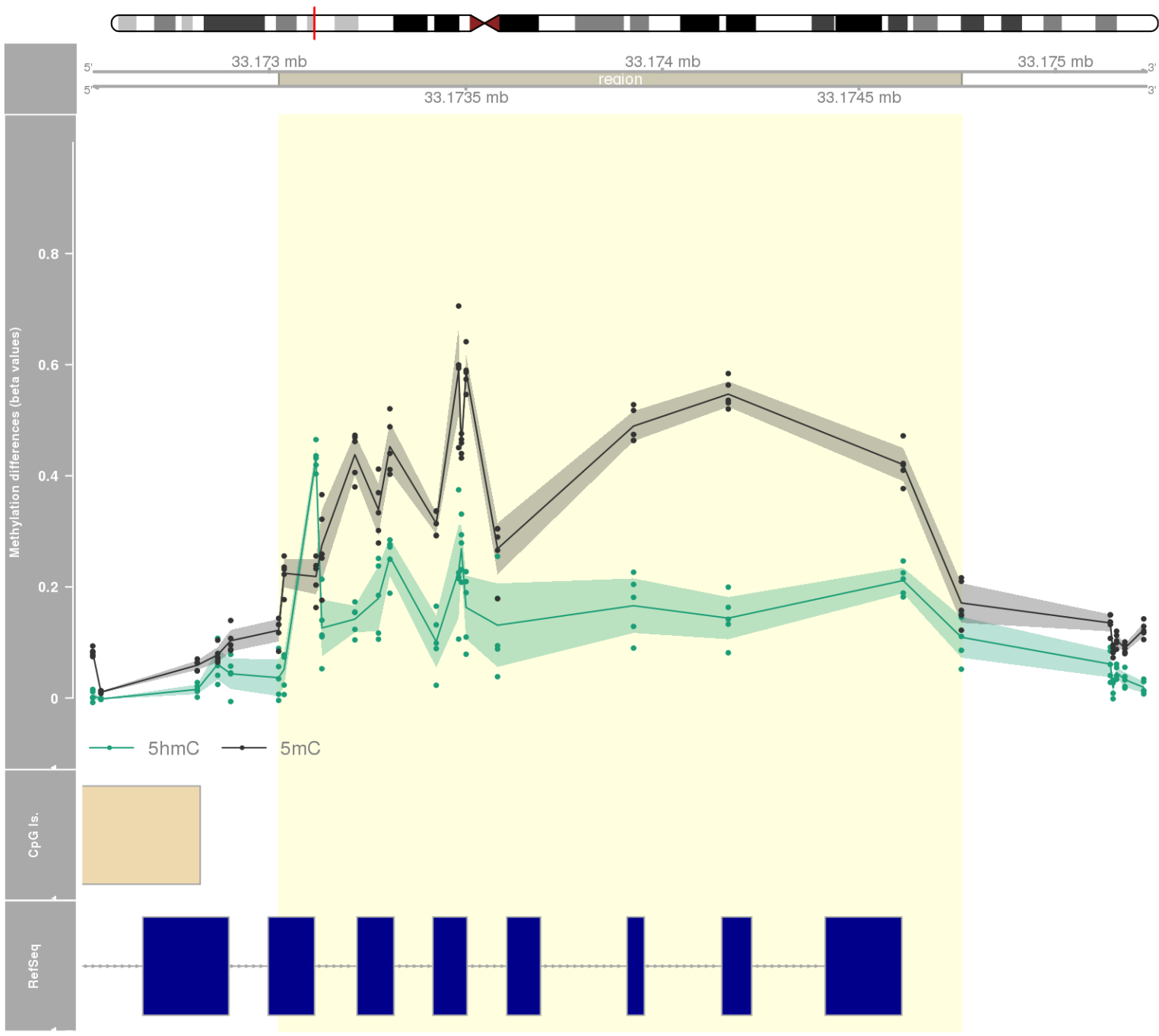
DMR 71 // chr17:4802692-4805392 // 2700 pb. (16 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: CHRNE / C17orf107 -



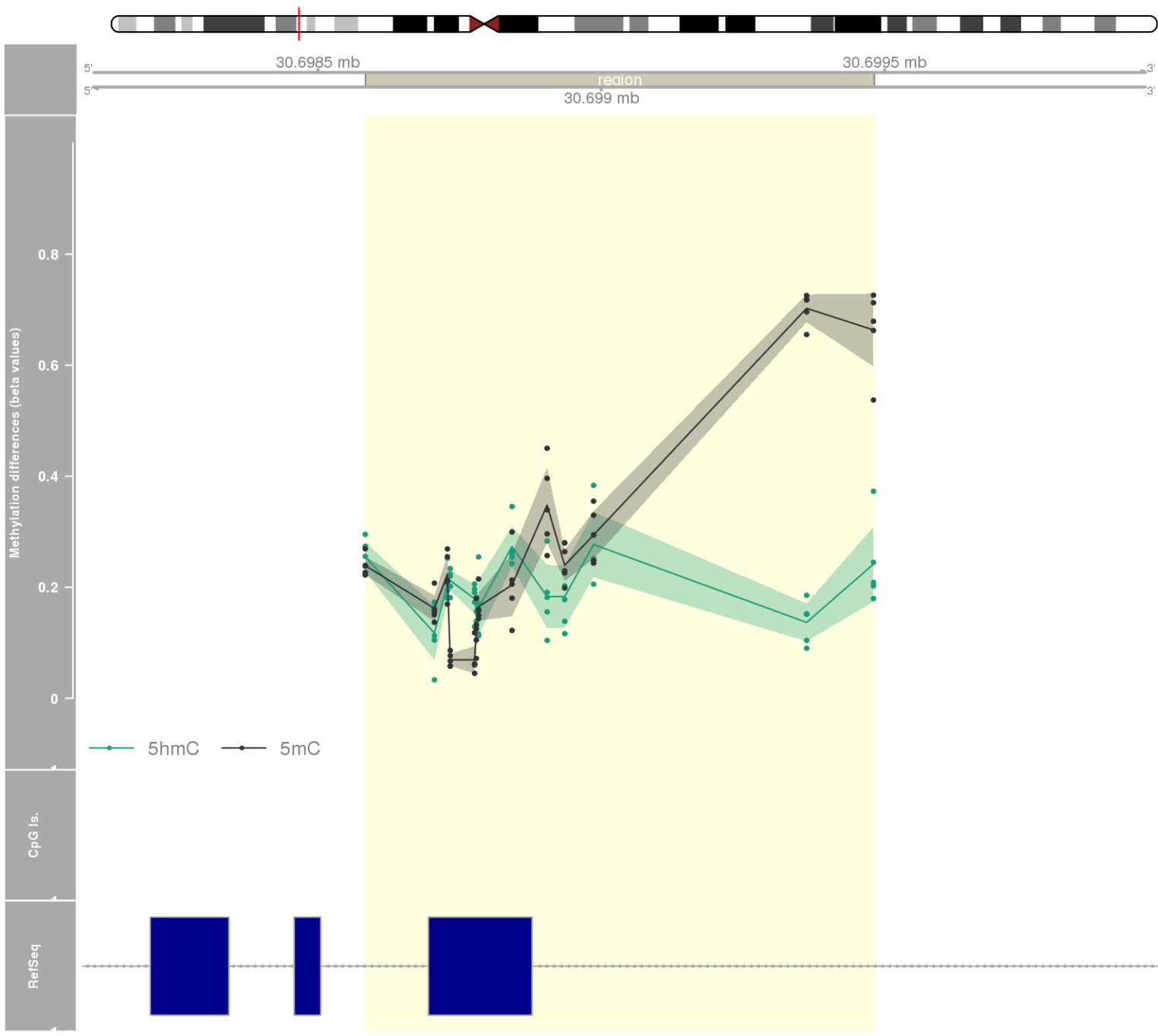
DMR 72 // chr17:80559059-80562120 // 3061 pb. (12 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: FOXK2 -



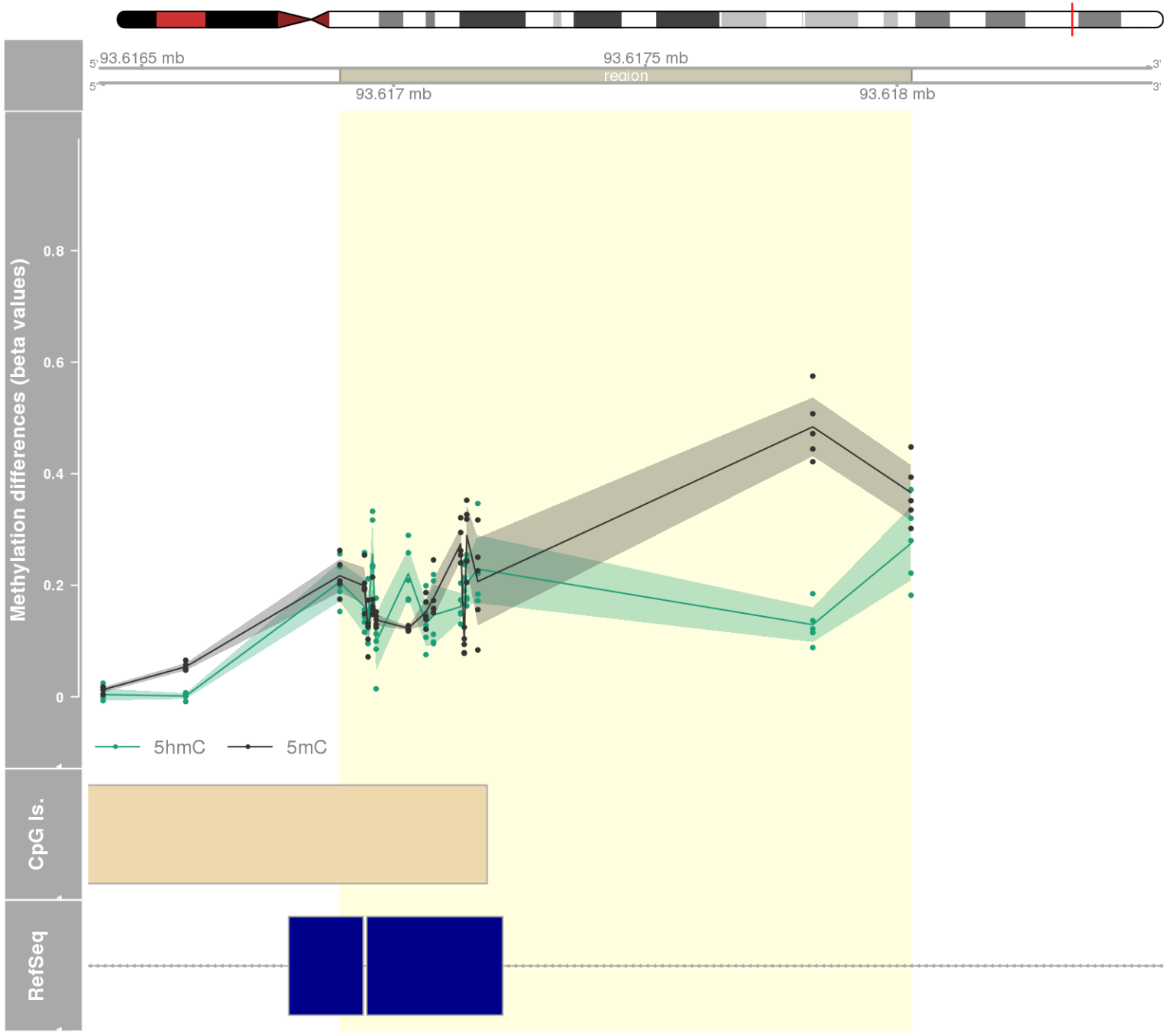
DMR 73 // chr6:33173024-33174761 // 1737 pb. (16 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: HSD17B8 -



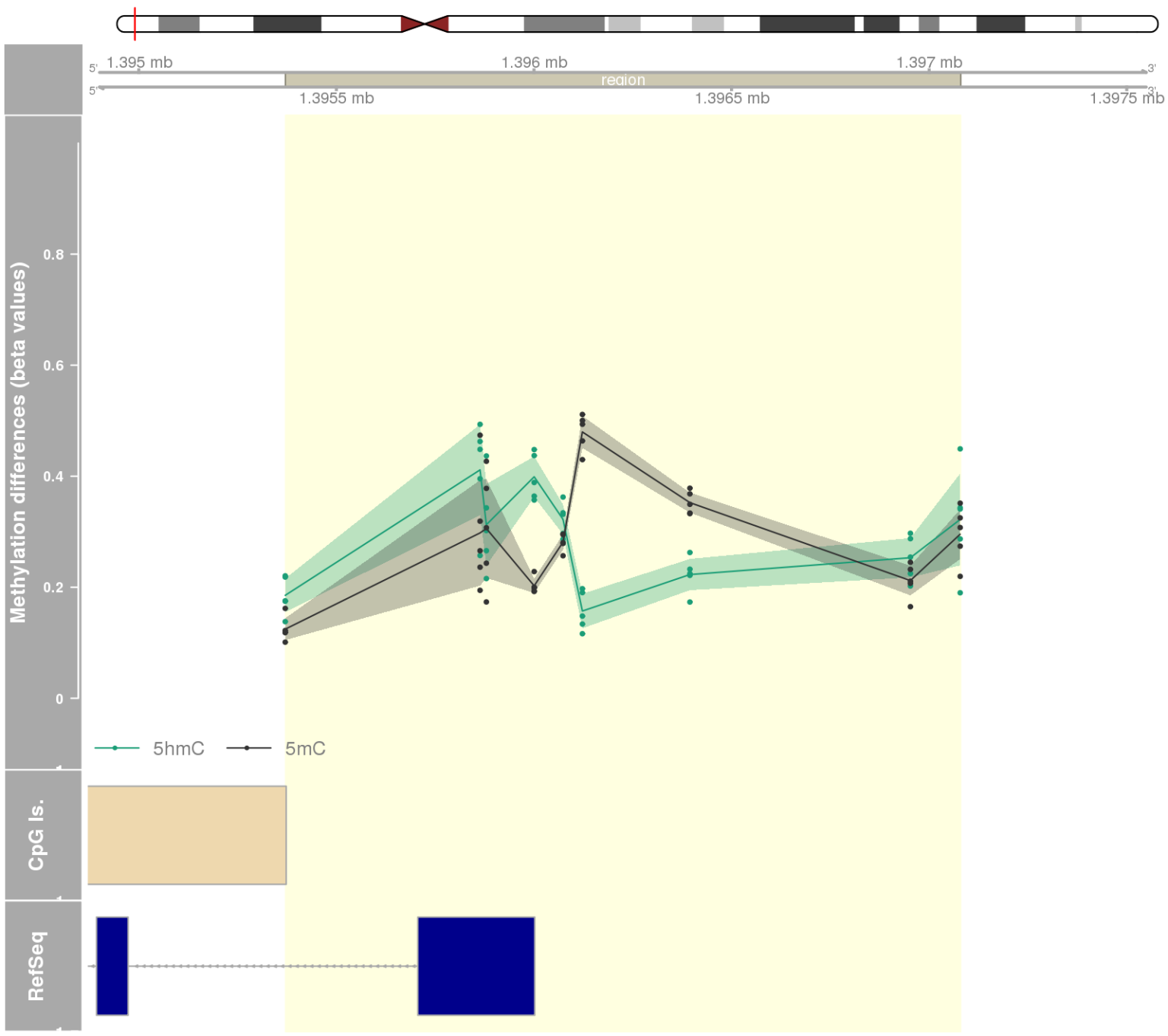
DMR 74 // chr6:30698584-30699481 // 897 pb. (13 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: FLOT1 -



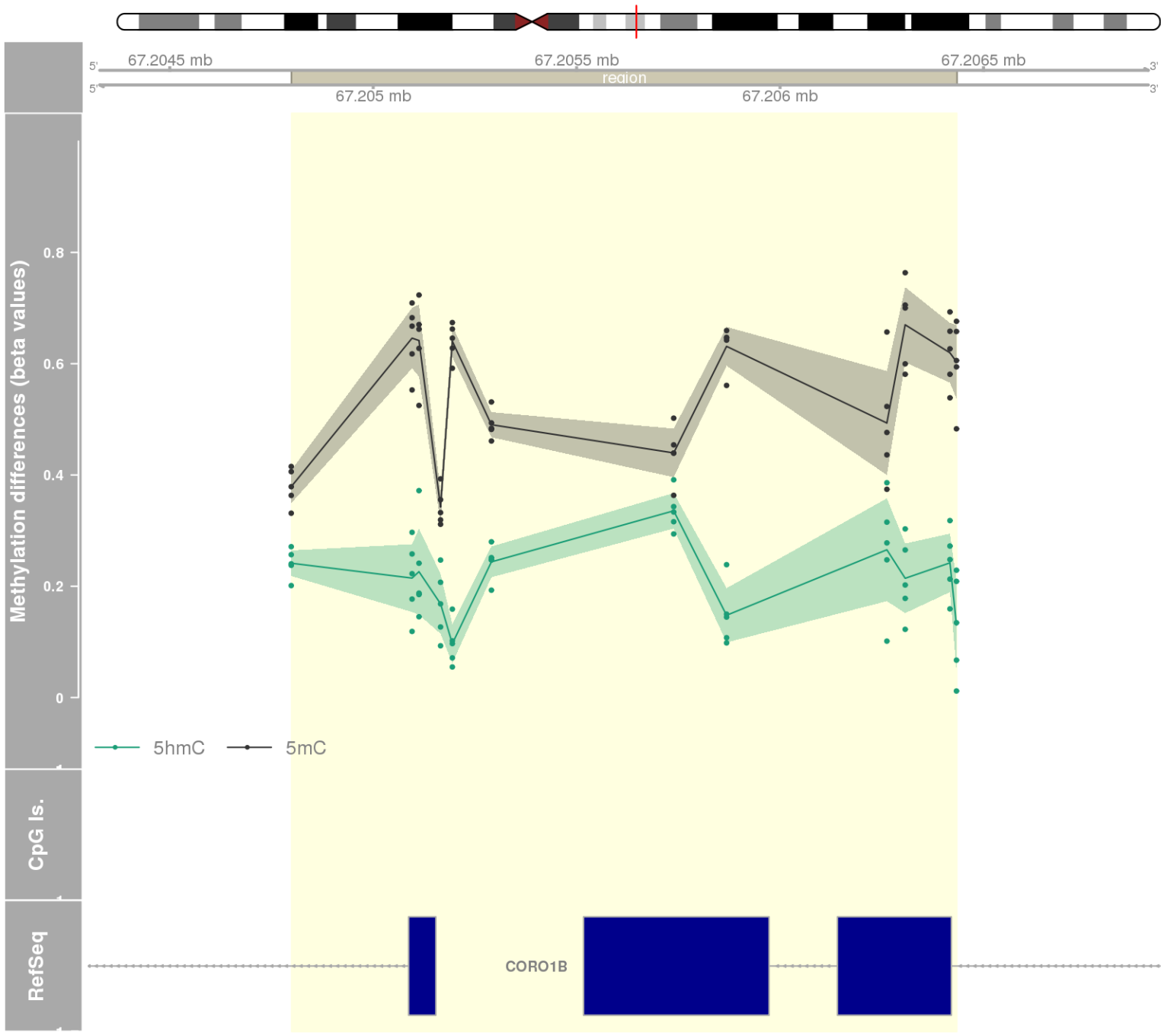
DMR 75 // chr15:93616894-93618028 // 1134 pb. (14 probes) // pvalue: 0.002 // fwer: 0.36 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: RGMA -



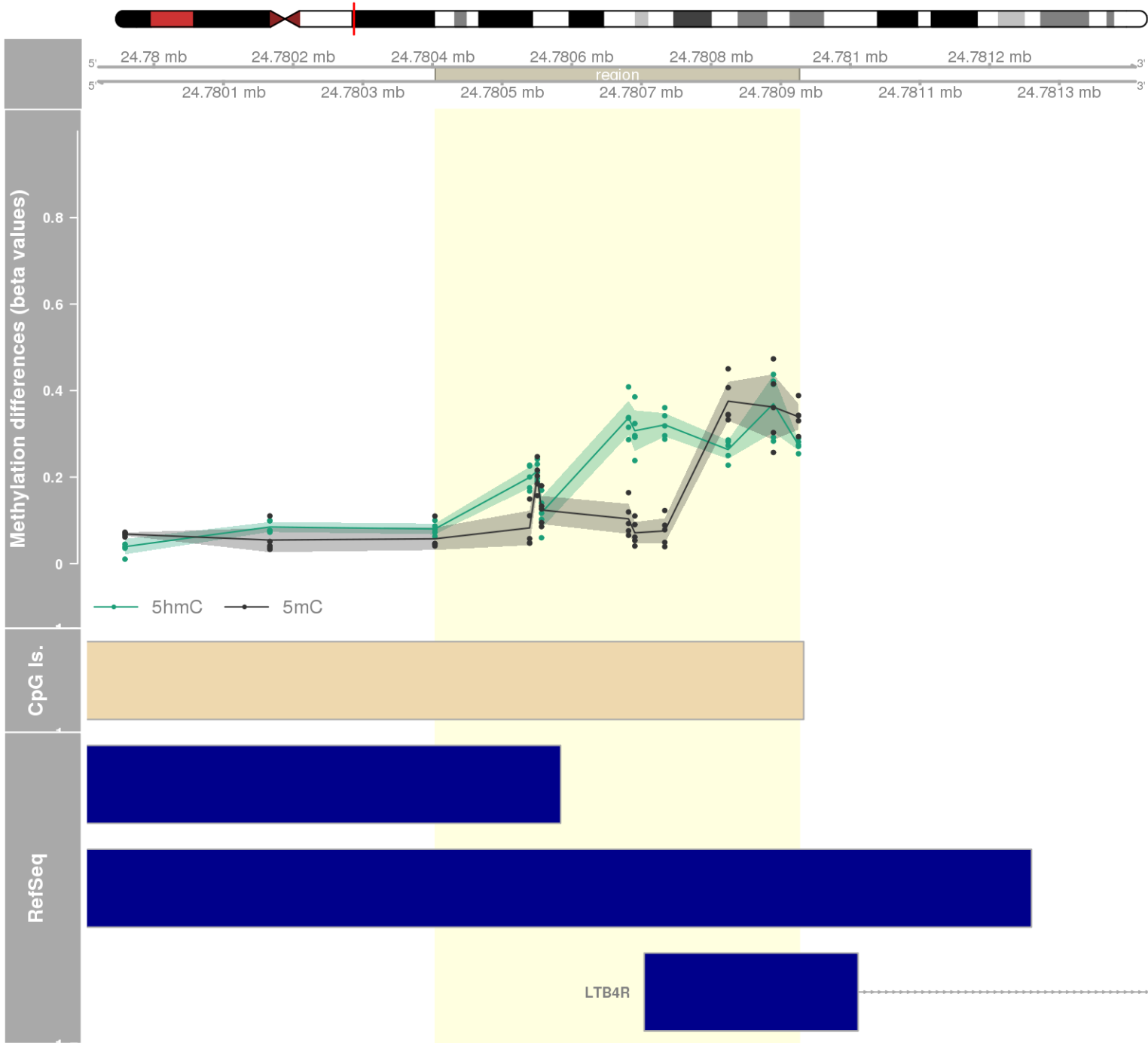
DMR 76 // chr17:1395371-1397079 // 1708 pb. (9 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: MYO1C -



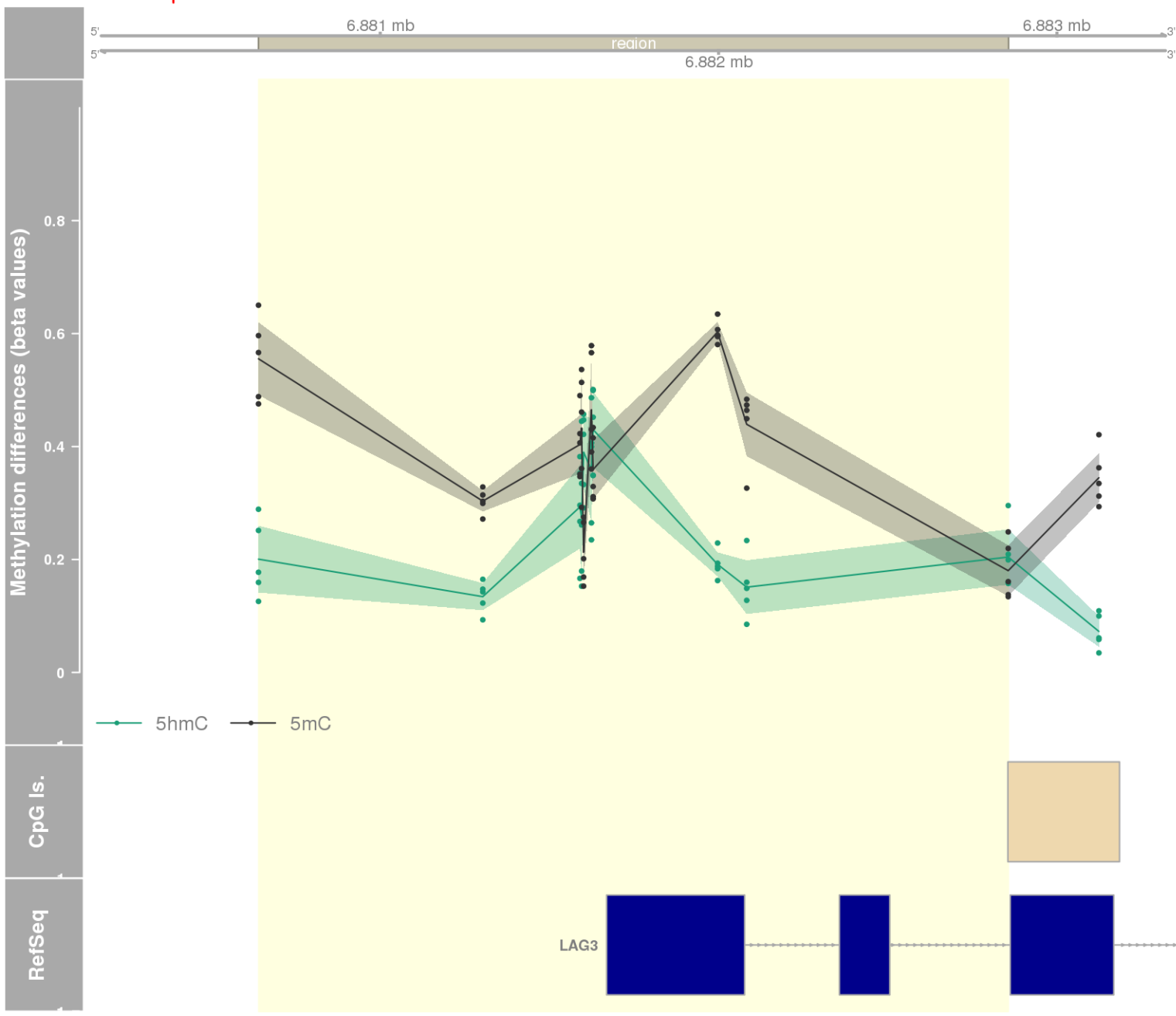
DMR 77 // chr11:67204799-67206434 // 1635 pb. (12 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: PTPRCAP / CORO1B -



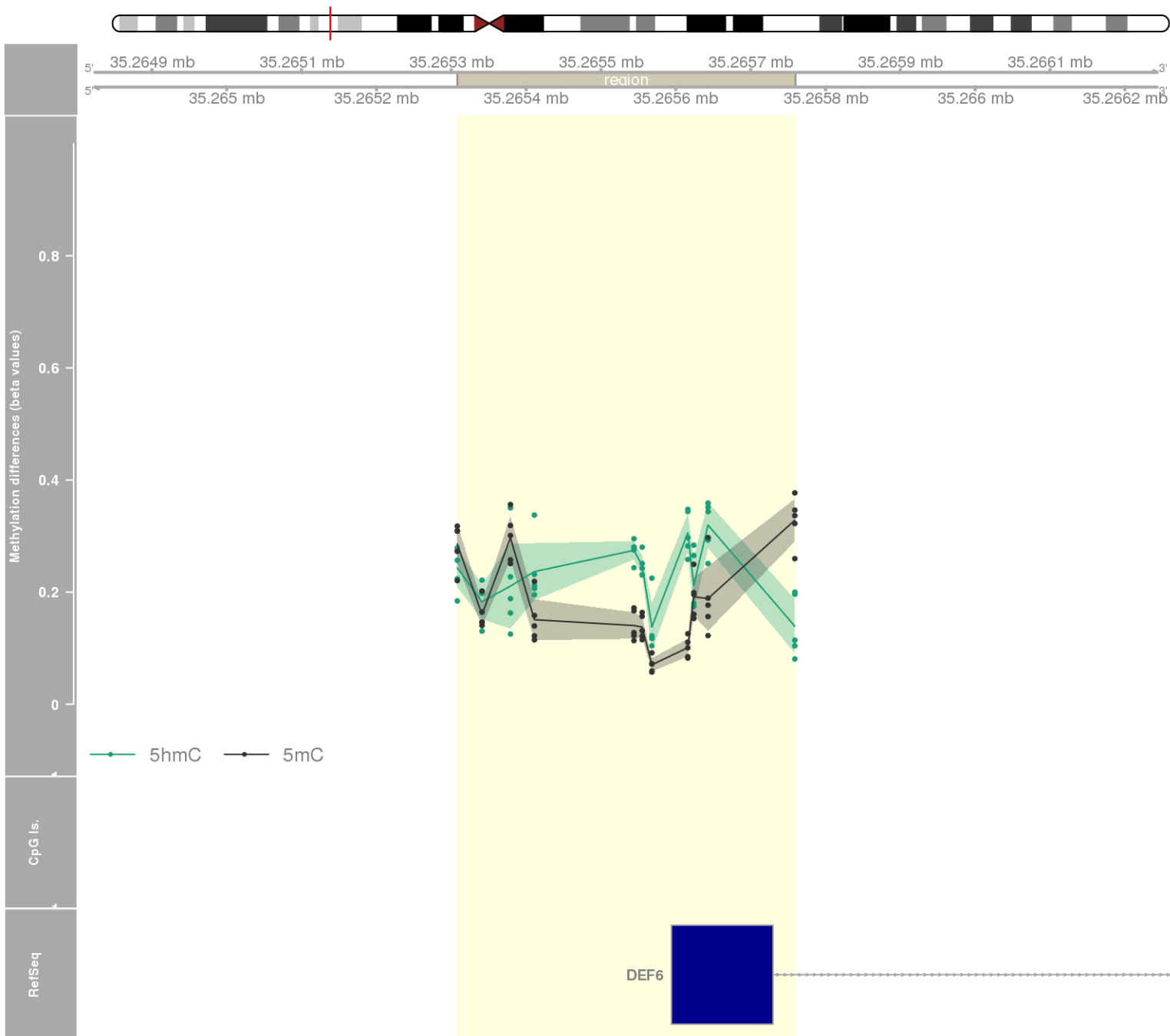
DMR 78 // chr14:24780404-24780926 // 522 pb. (10 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.005 // fwerArea: 0.684
- genes: CIDEB / LTB4R2 / LTB4R -



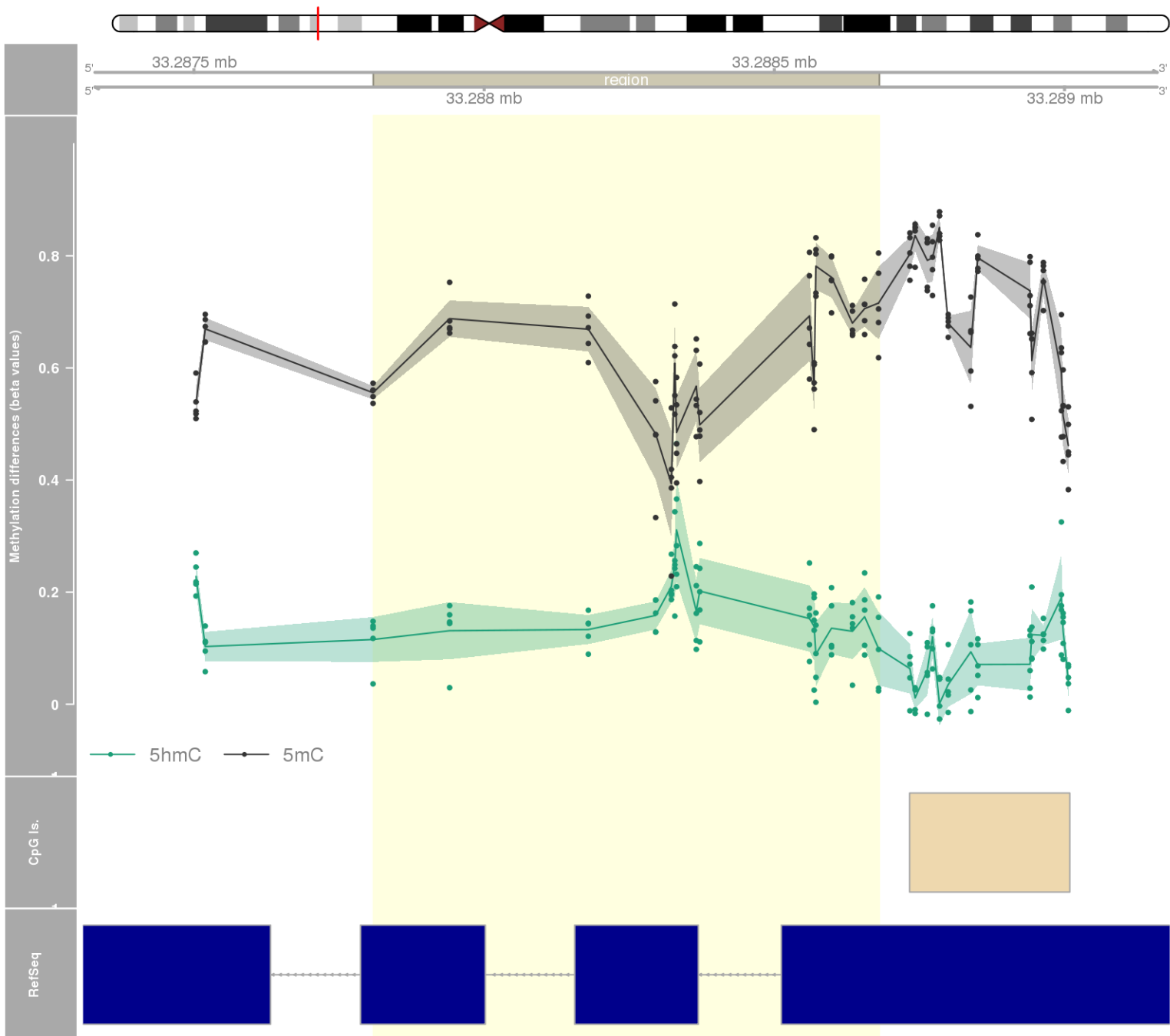
DMR 79 // chr12:6880640-6882856 // 2216 pb. (10 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.006 // fwerArea: 0.684
- genes: LAG3 -



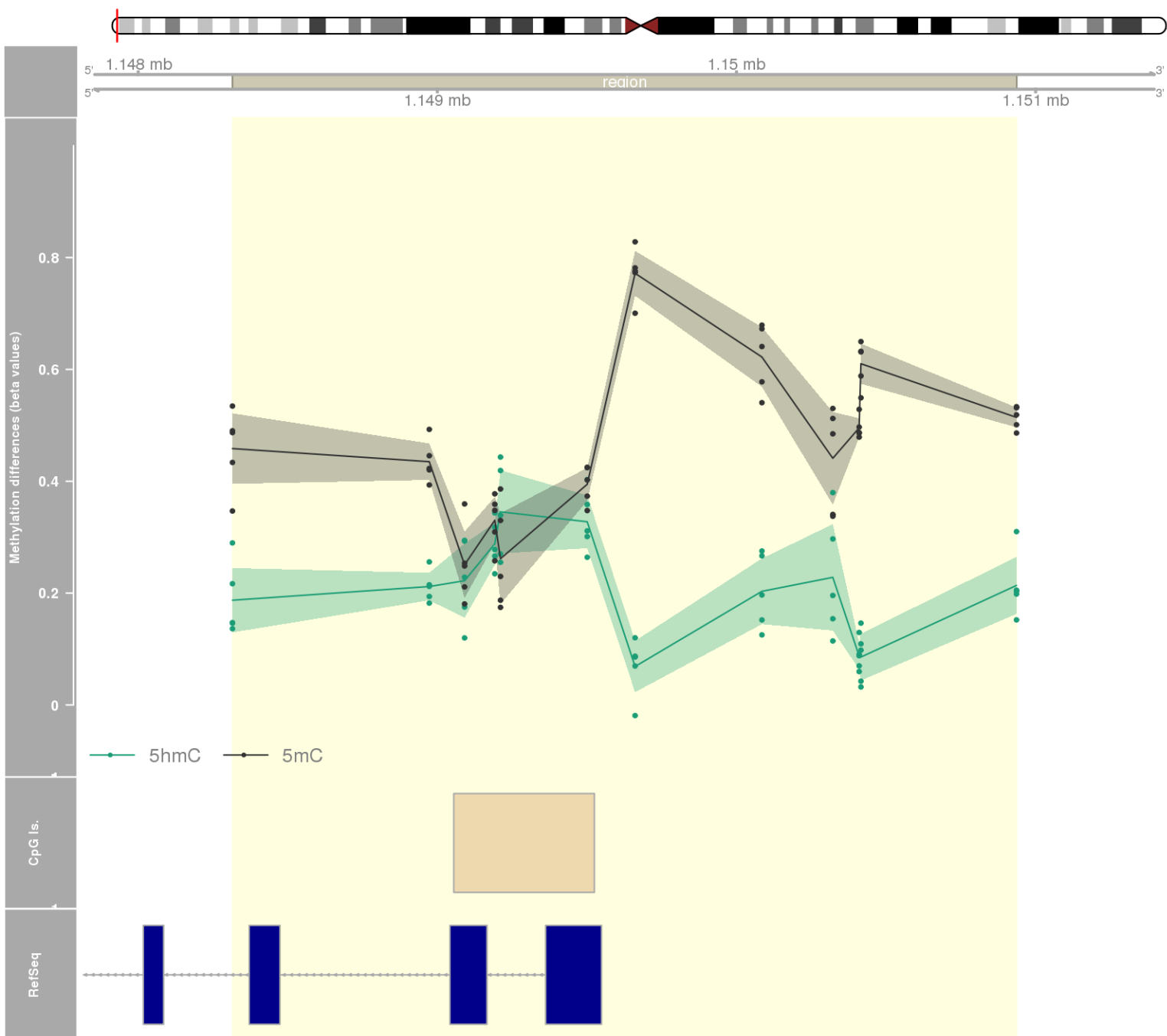
DMR 80 // chr6:35265308-35265759 // 451 pb. (11 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.006 // fwerArea: 0.684
- genes: DEF6 -



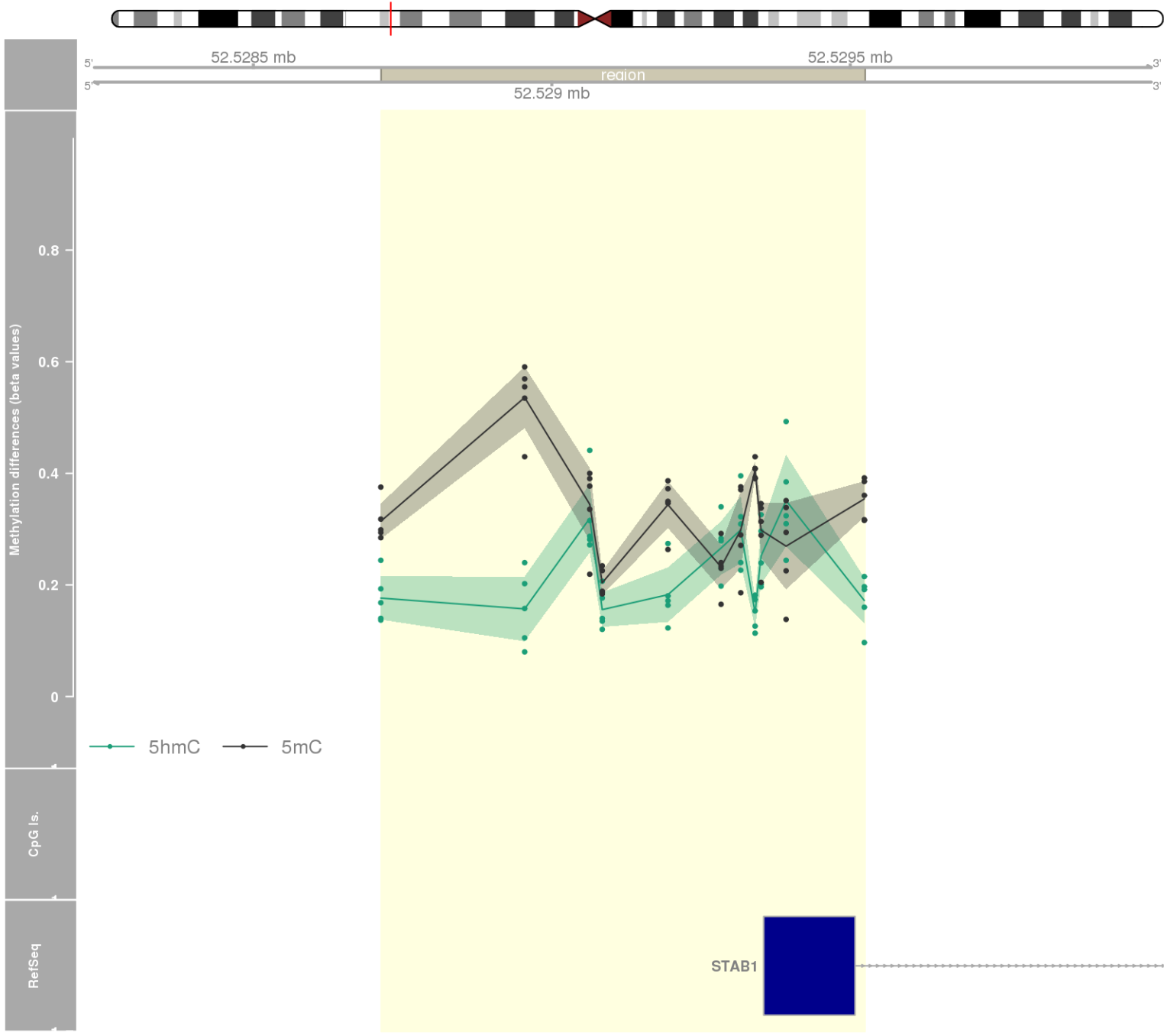
DMR 81 // chr6:33287809-33288680 // 871 pb. (16 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.006 // fwerArea: 0.684
- genes: DAXX -



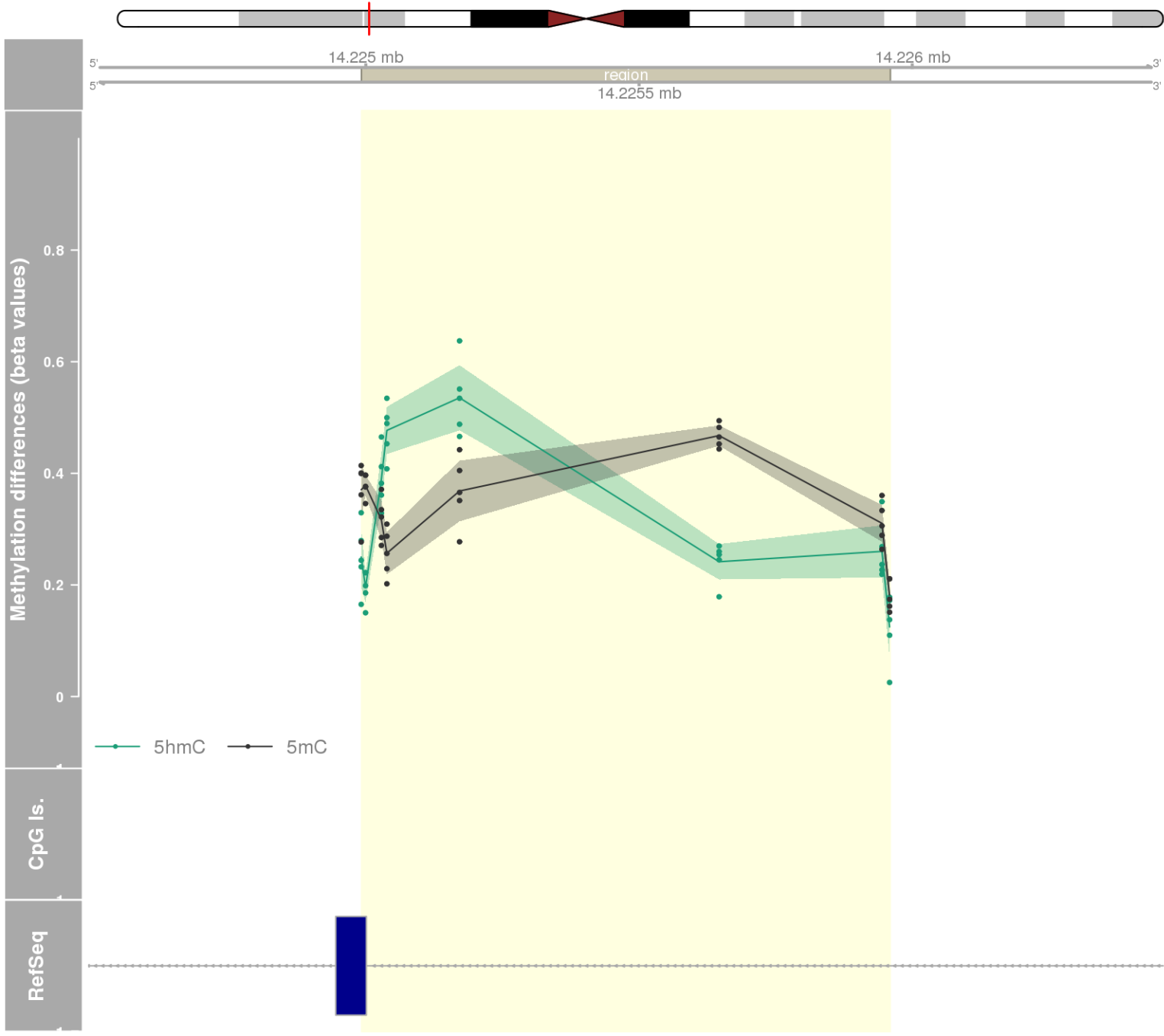
DMR 82 // chr1:1148315-1150936 // 2621 pb. (12 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.006 // fwerArea: 0.684
- genes: TNFRSF4 -



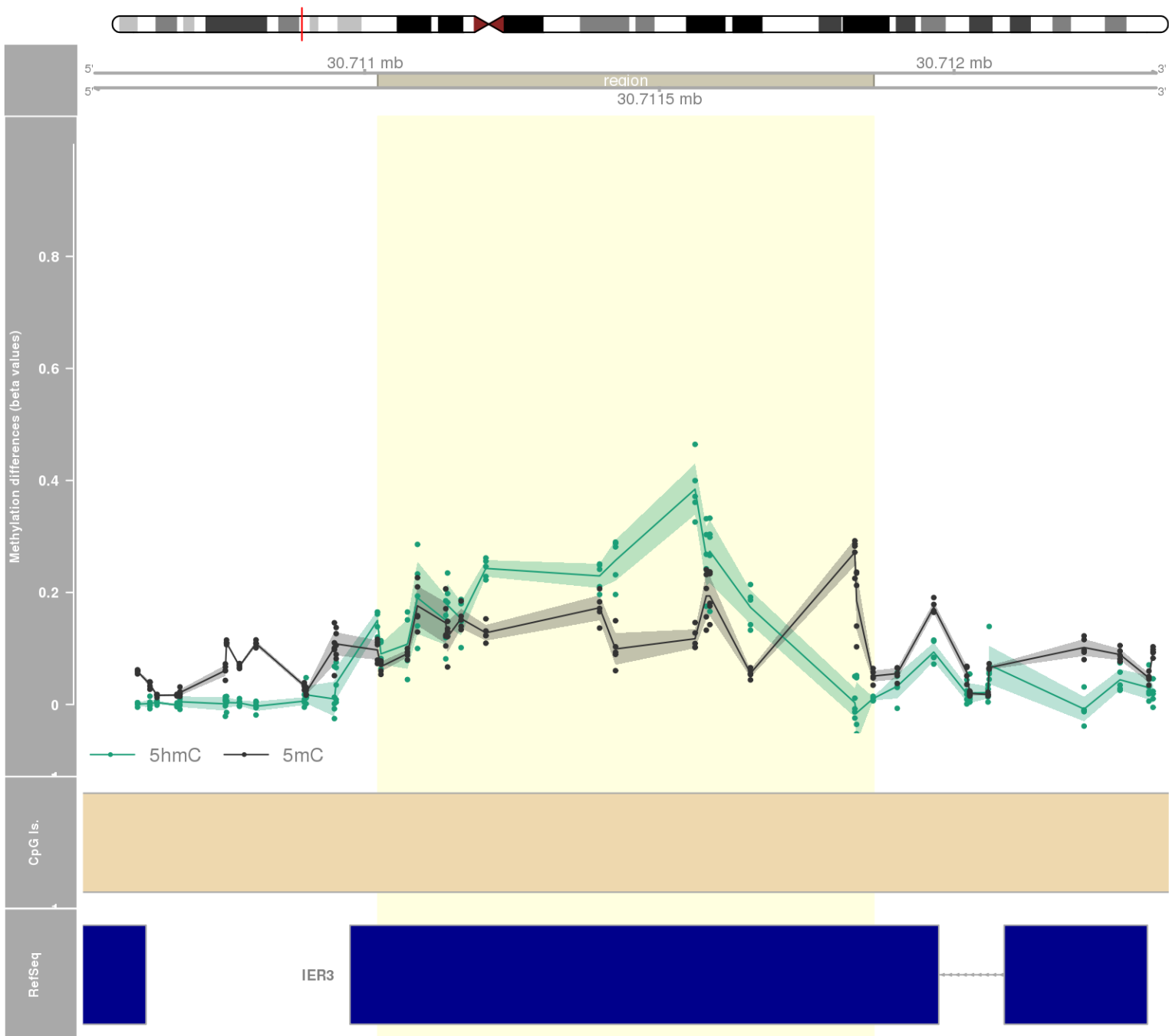
DMR 83 // chr3:52528714-52529524 // 810 pb. (11 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.006 // fwerArea: 0.684
- genes: STAB1 -



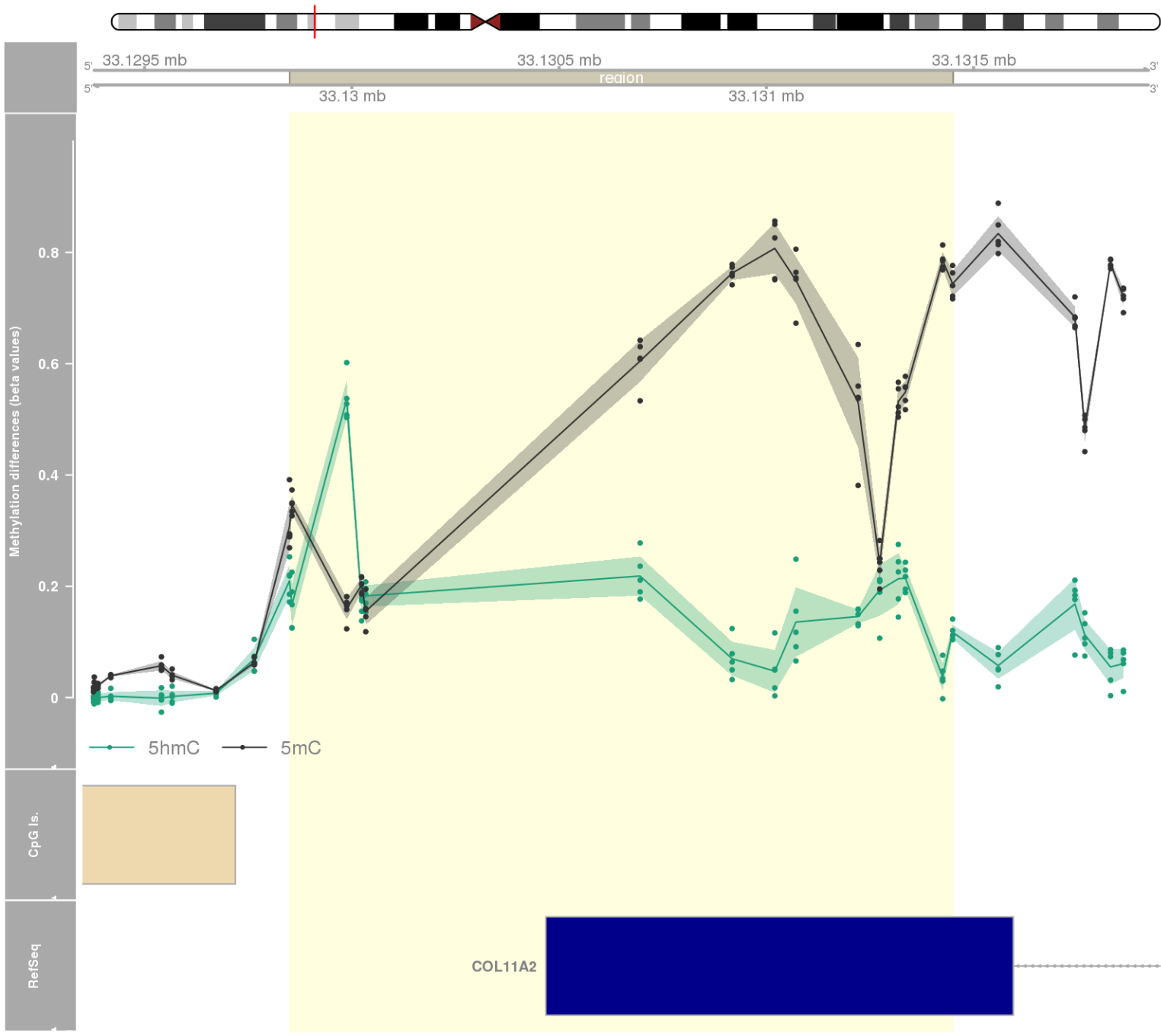
DMR 84 // chr19:14224992-14225959 // 967 pb. (8 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.006 // fwerArea: 0.846
- genes: PRKACA -



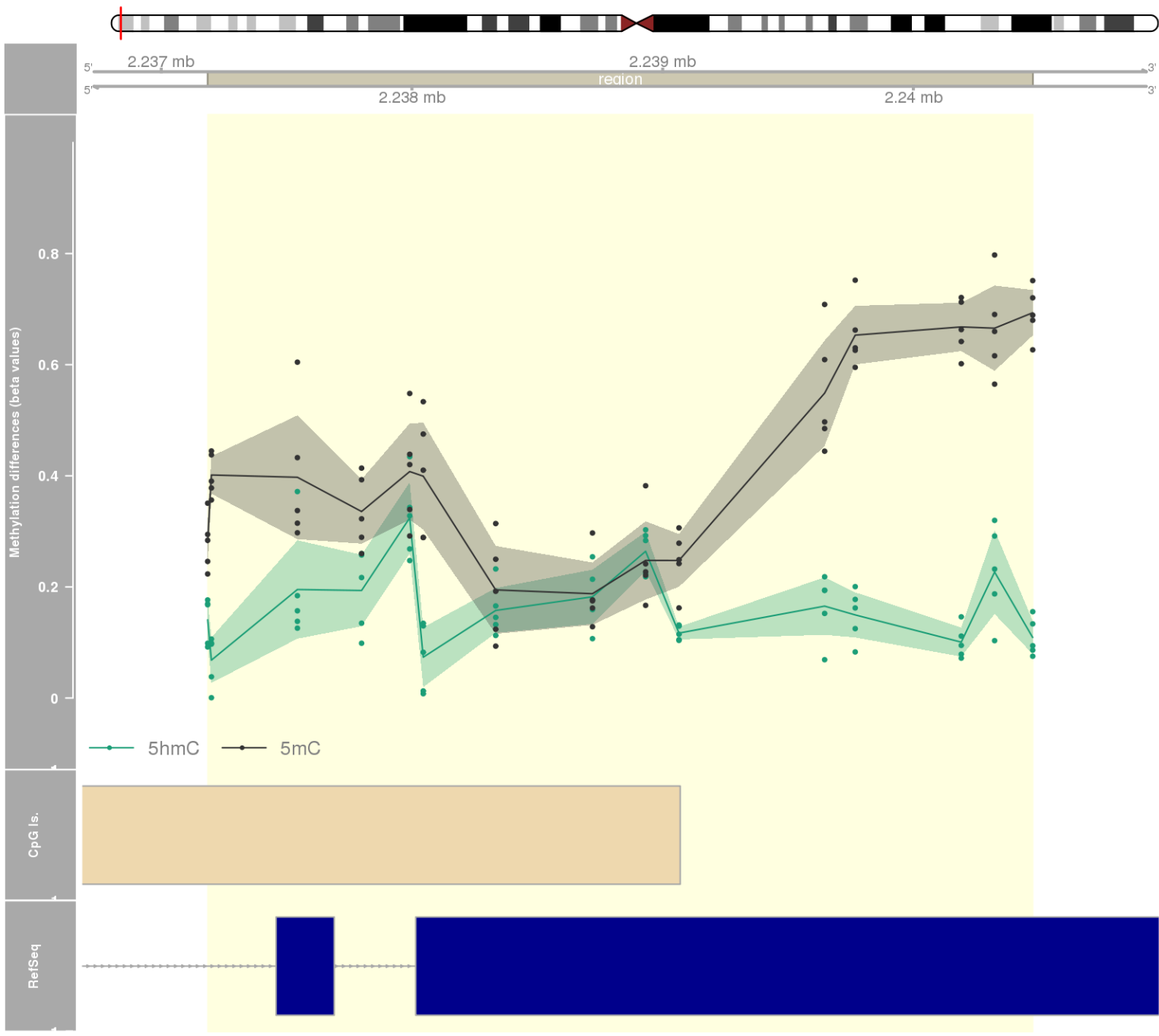
DMR 85 // chr6:30711022-30711863 // 841 pb. (17 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.006 // fwerArea: 0.846
- genes: IER3 -



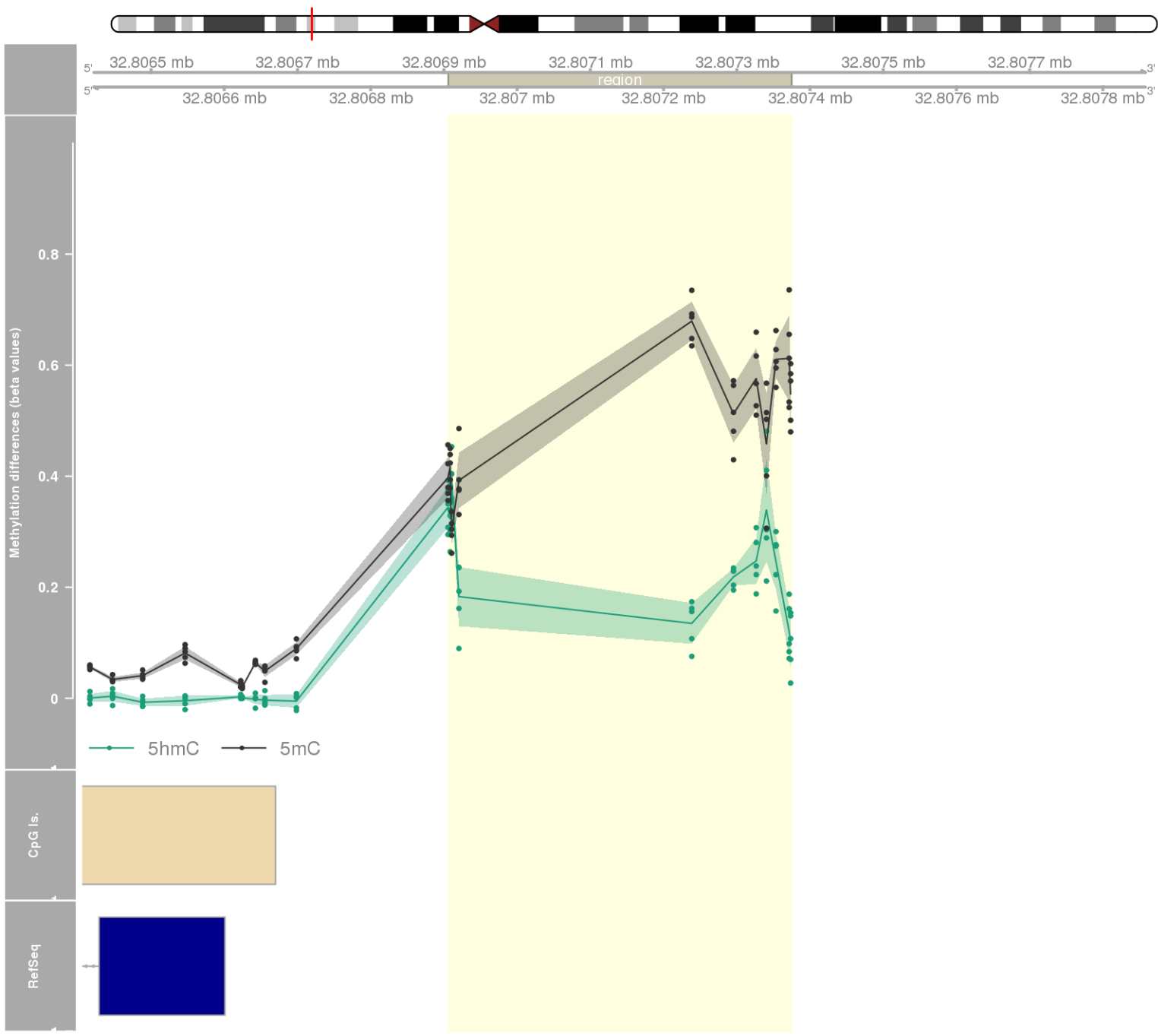
DMR 86 // chr6:33129850-33131450 // 1600 pb. (15 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.006 // fwerArea: 0.846
- genes: COL11A2 -



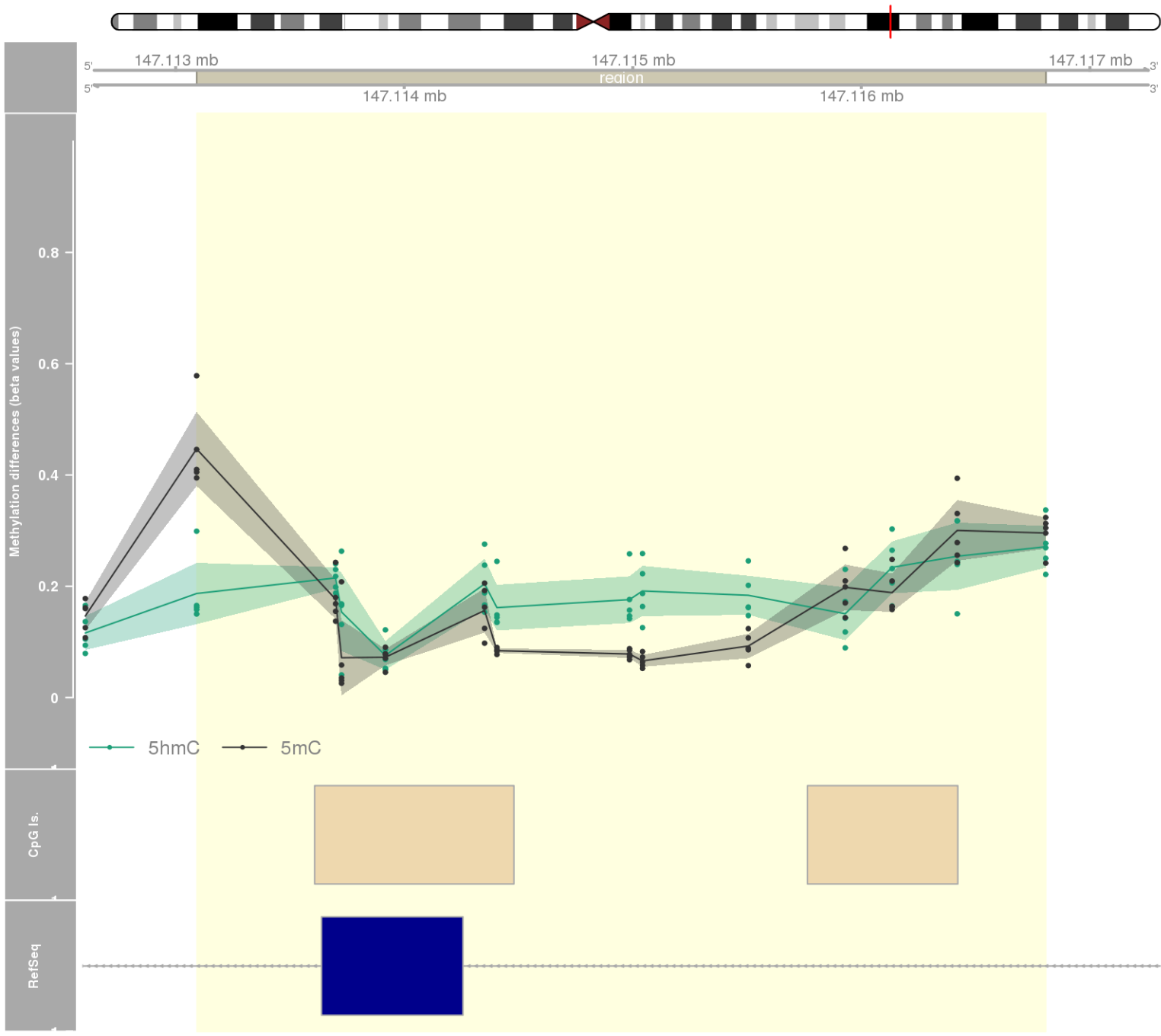
DMR 87 // chr1:2237185-2240476 // 3291 pb. (15 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.006 // fwerArea: 0.846
- genes: SKI -



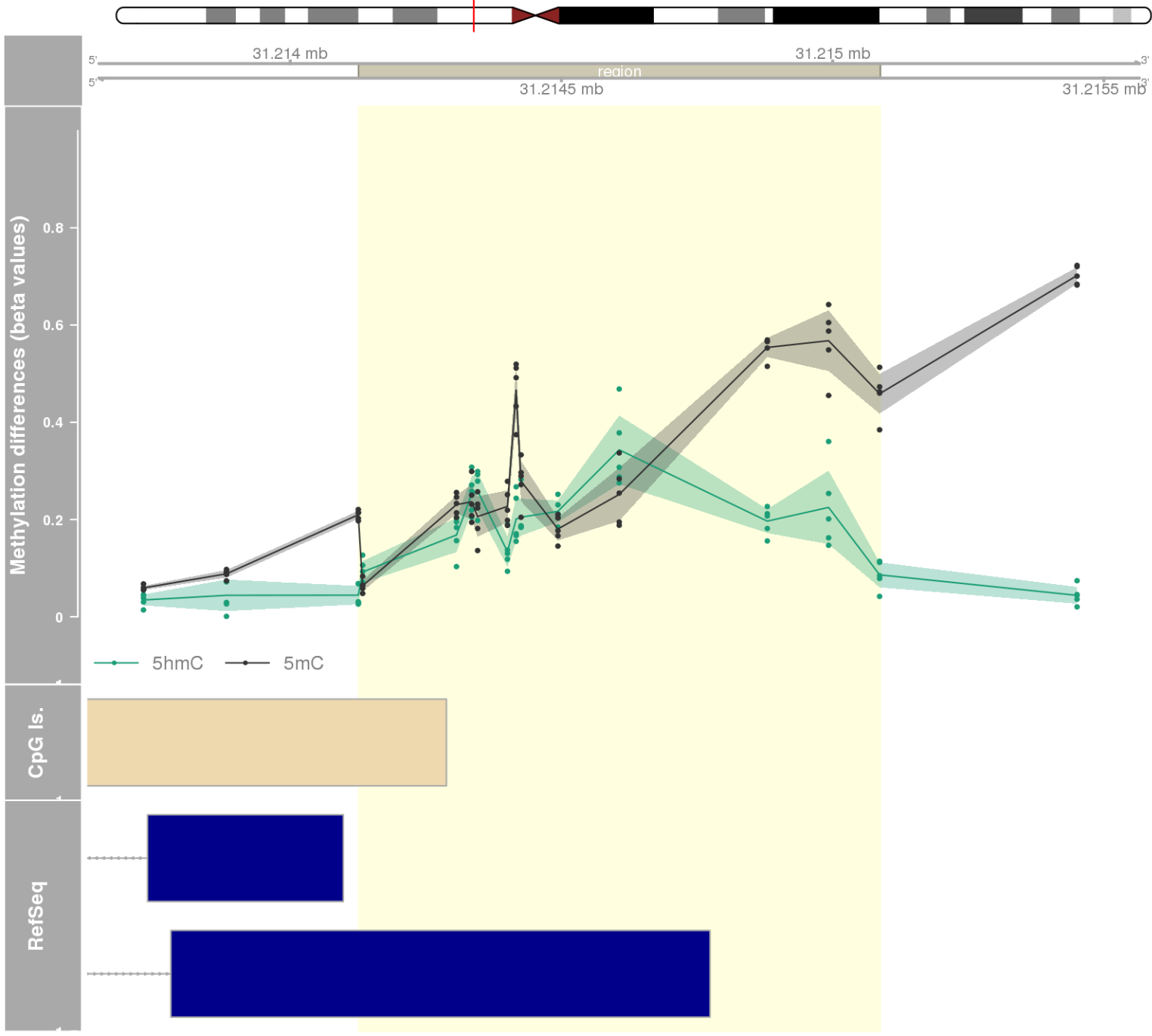
DMR 88 // chr6:32806906-32807374 // 468 pb. (11 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.006 // fwerArea: 0.846



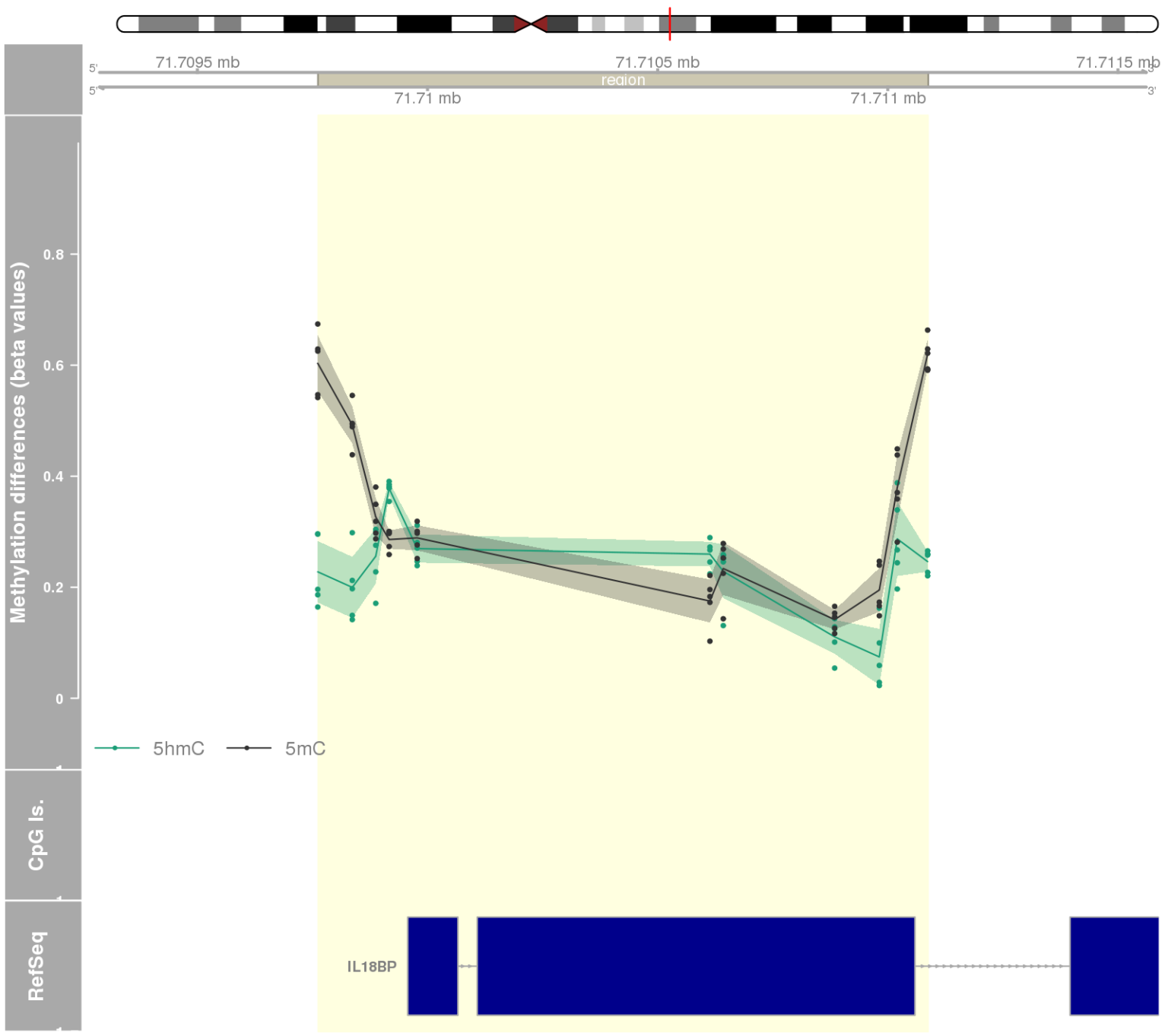
DMR 89 // chr3:147113092-147116807 // 3715 pb. (13 probes) // pvalue: 0.002 // fwer: 0.36 // pvalueArea: 0.006 // fwerArea: 0.846
- genes: ZIC4 / ZIC1 -



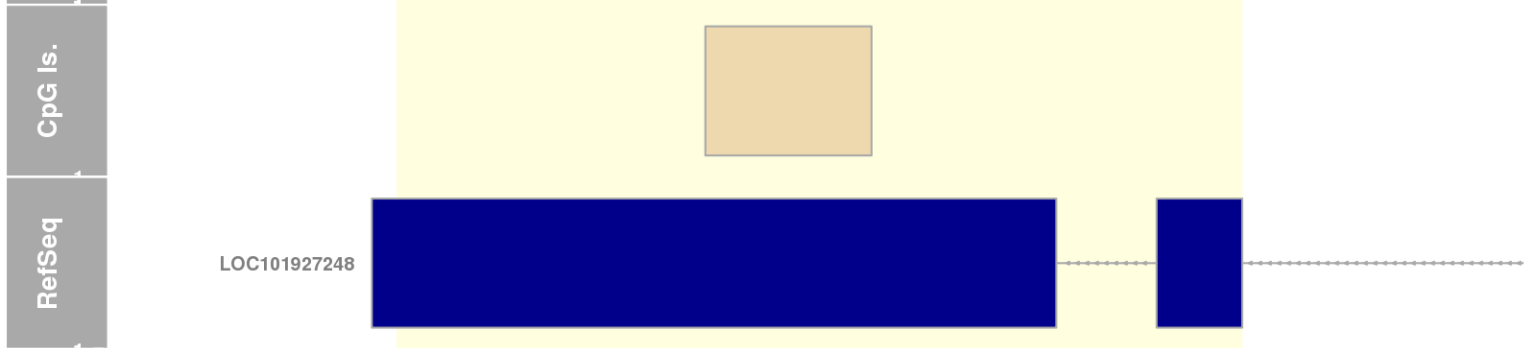
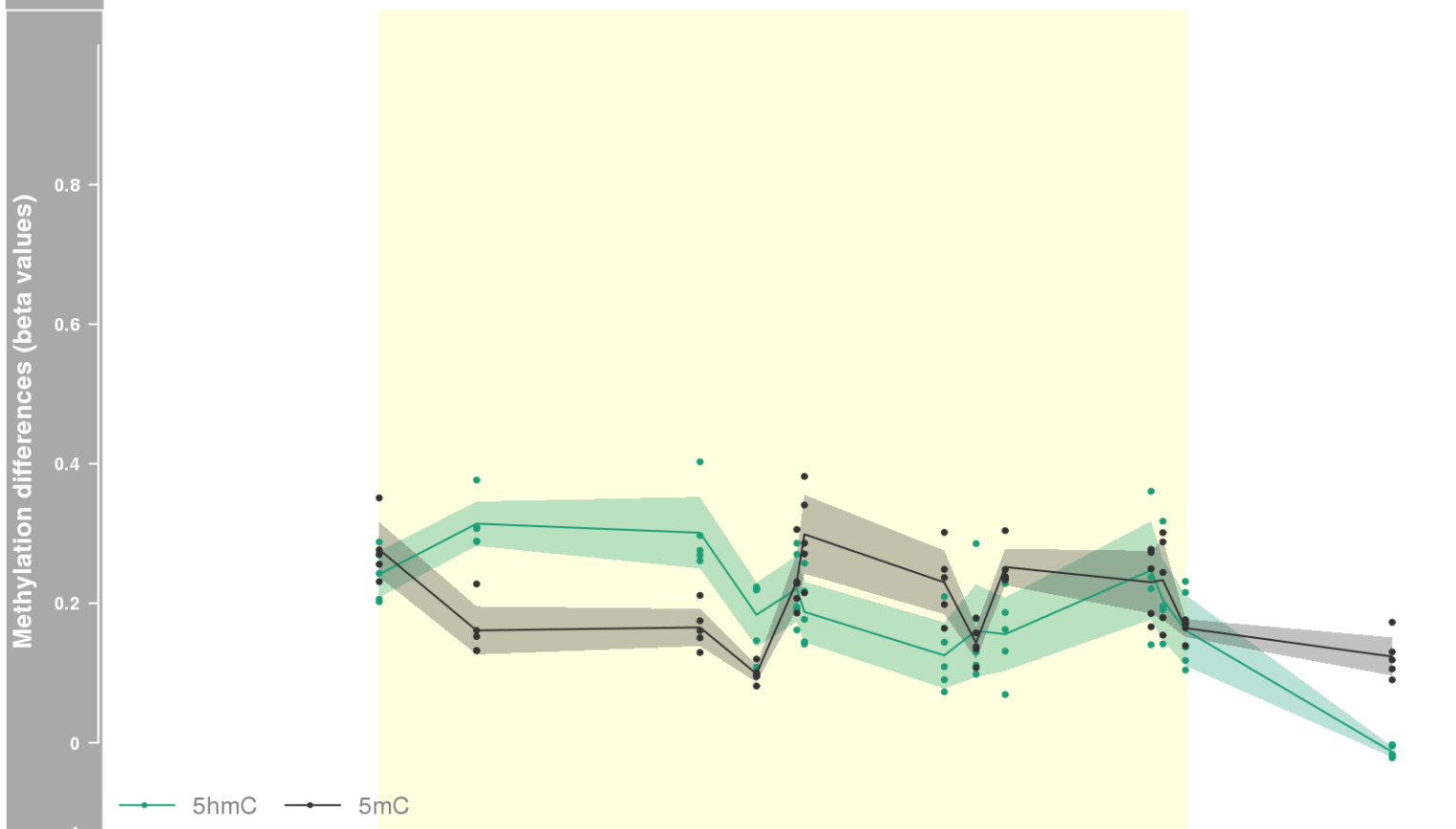
DMR 90 // chr16:31214126-31215087 // 961 pb. (13 probes) // pvalue: 0.002 // fwer: 0.36 // pvalueArea: 0.006 // fwerArea: 0.846
- genes: PYCARD / C16orf98 -



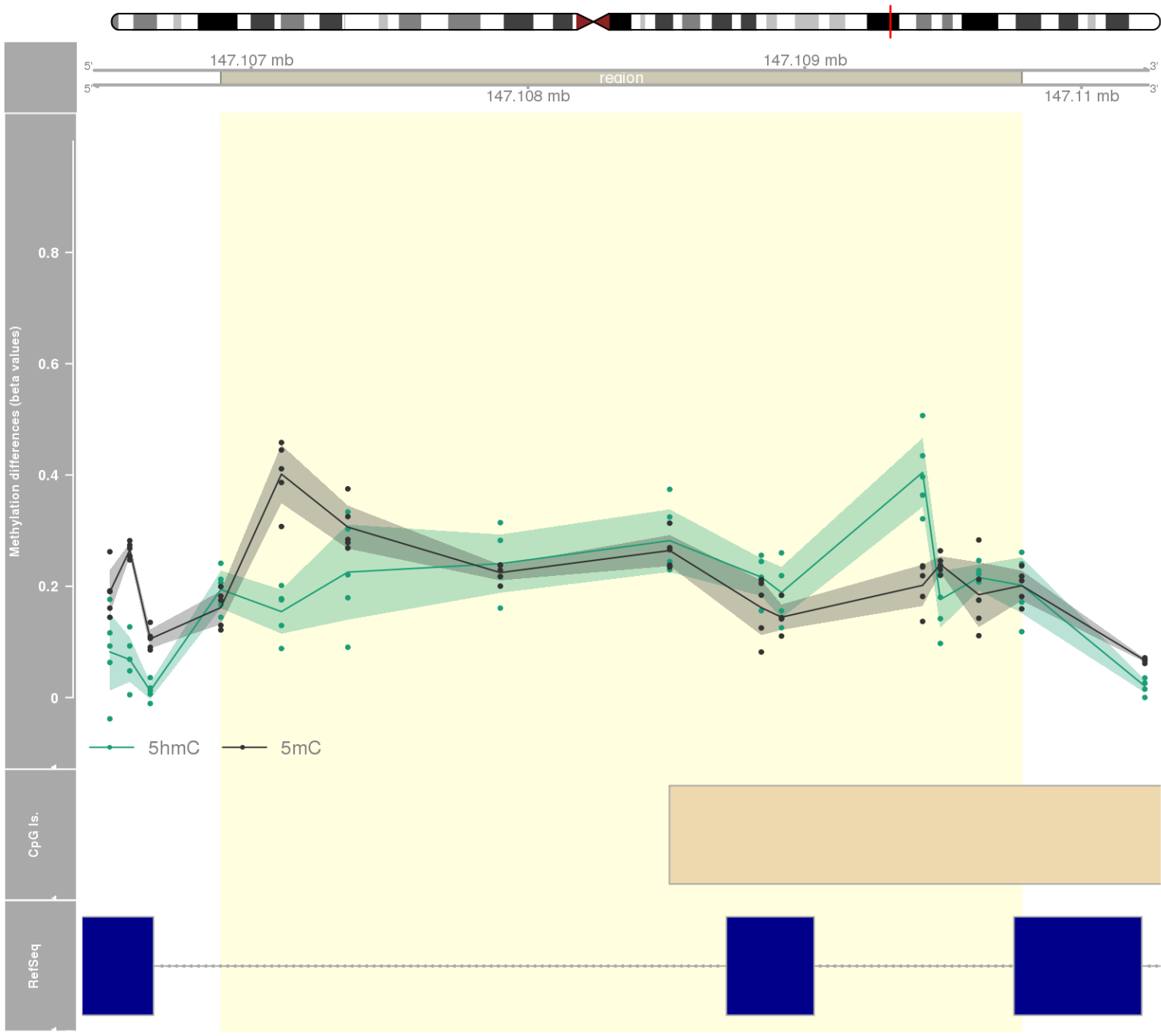
DMR 91 // chr11:71709762-71711087 // 1325 pb. (11 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.006 // fwerArea: 0.846
- genes: IL18BP -



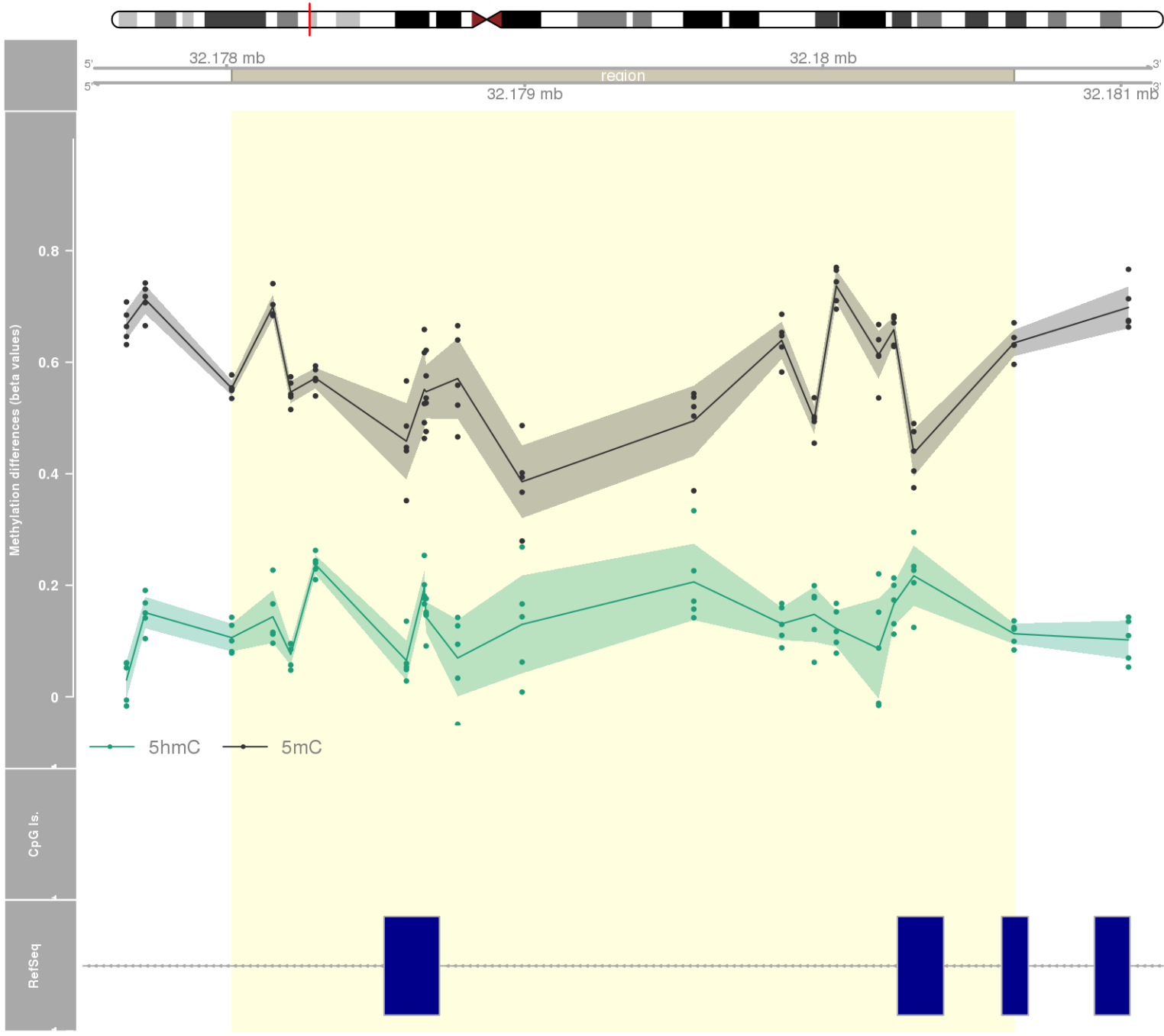
DMR 92 // chr13:95357042-95358540 // 1498 pb. (12 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.007 // fwerArea: 0.846



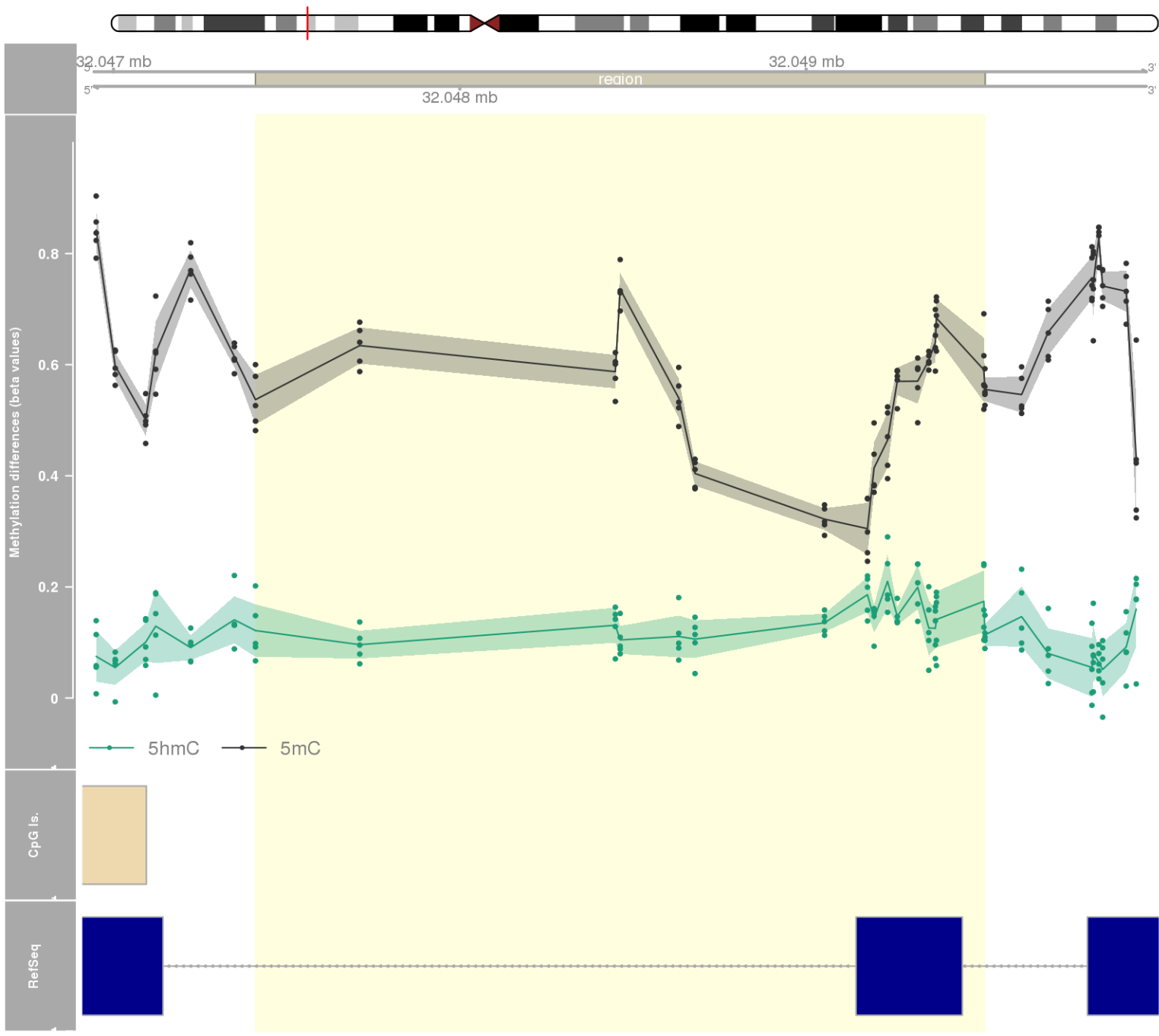
DMR 93 // chr3:147106890-147109784 // 2894 pb. (11 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.007 // fwerArea: 0.846
- genes: ZIC4 -



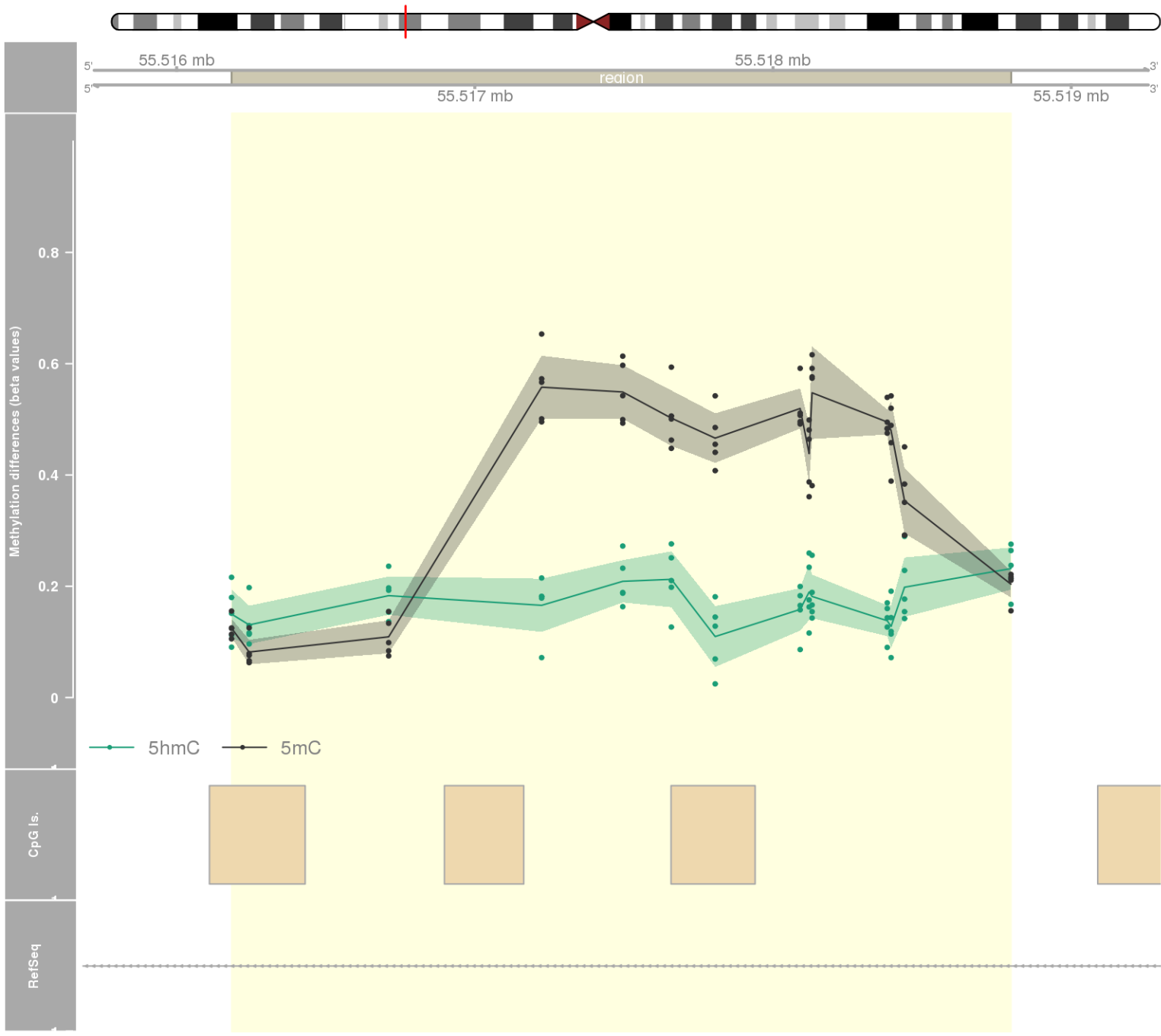
DMR 95 // chr6:32178017-32180641 // 2624 pb. (17 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.007 // fwerArea: 0.846
- genes: NOTCH4 -



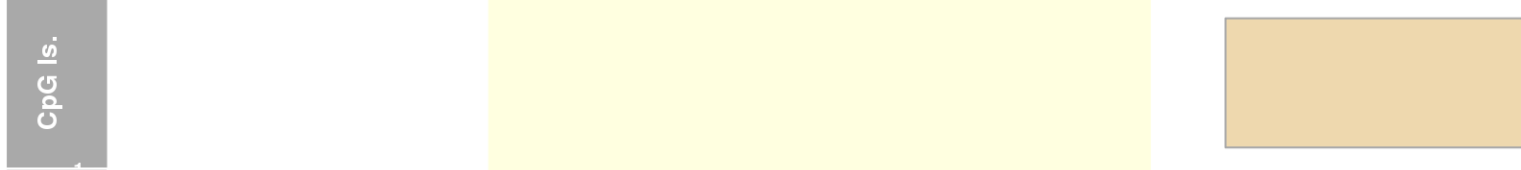
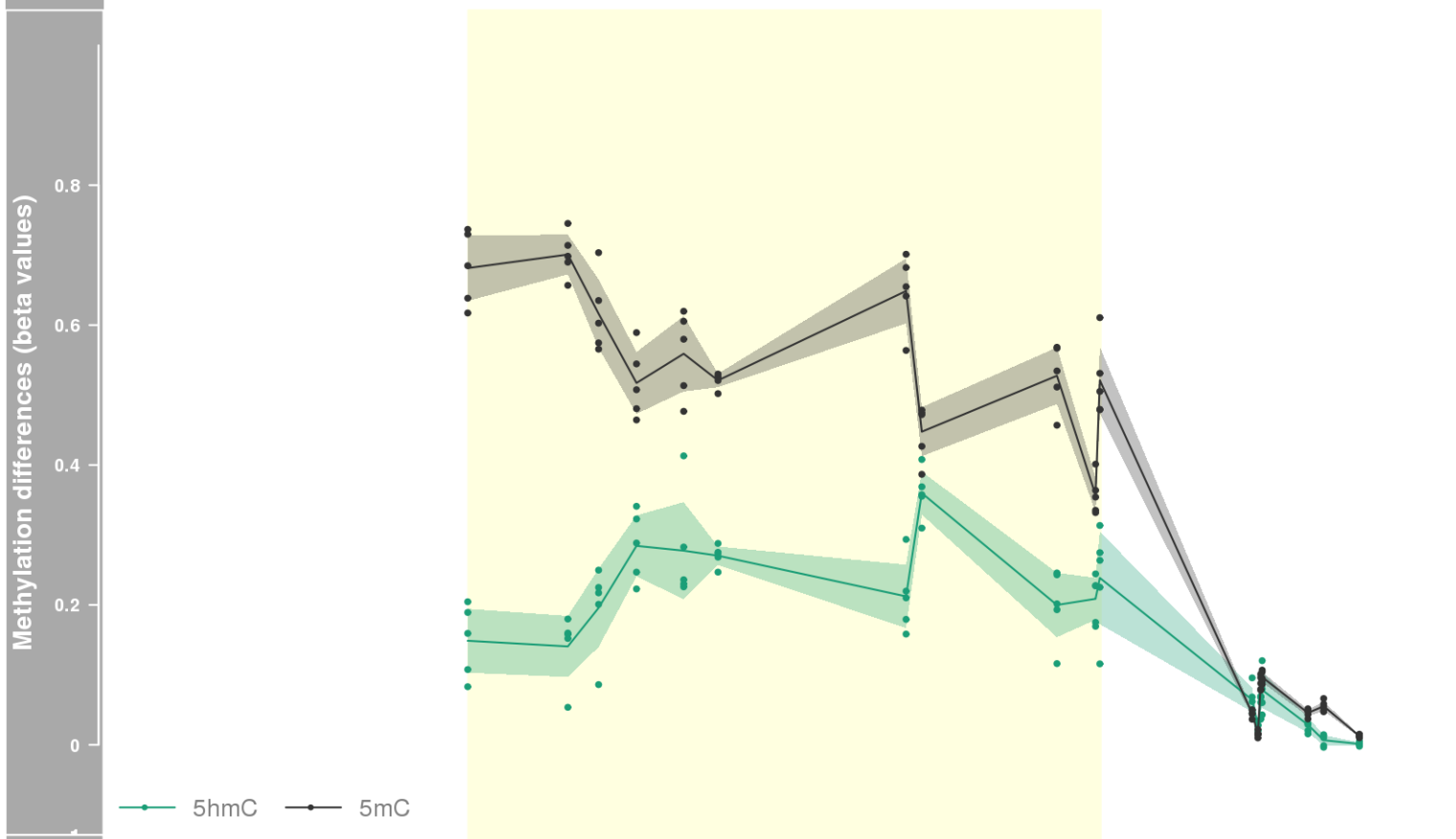
DMR 96 // chr6:32047410-32049516 // 2106 pb. (17 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.007 // fwerArea: 0.846
- genes: TNXB -



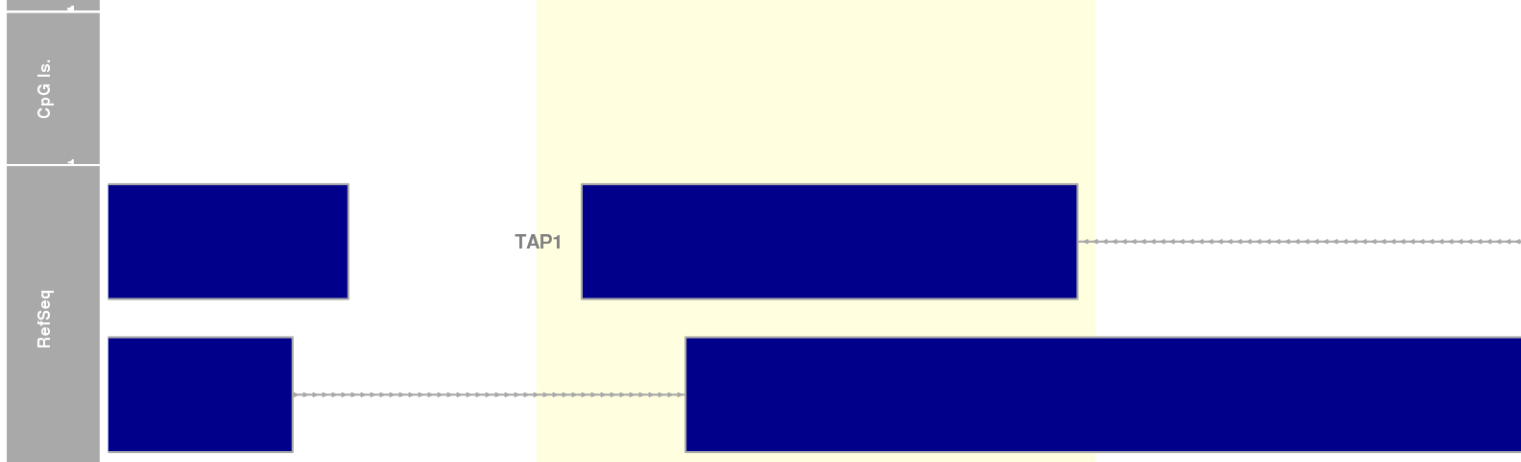
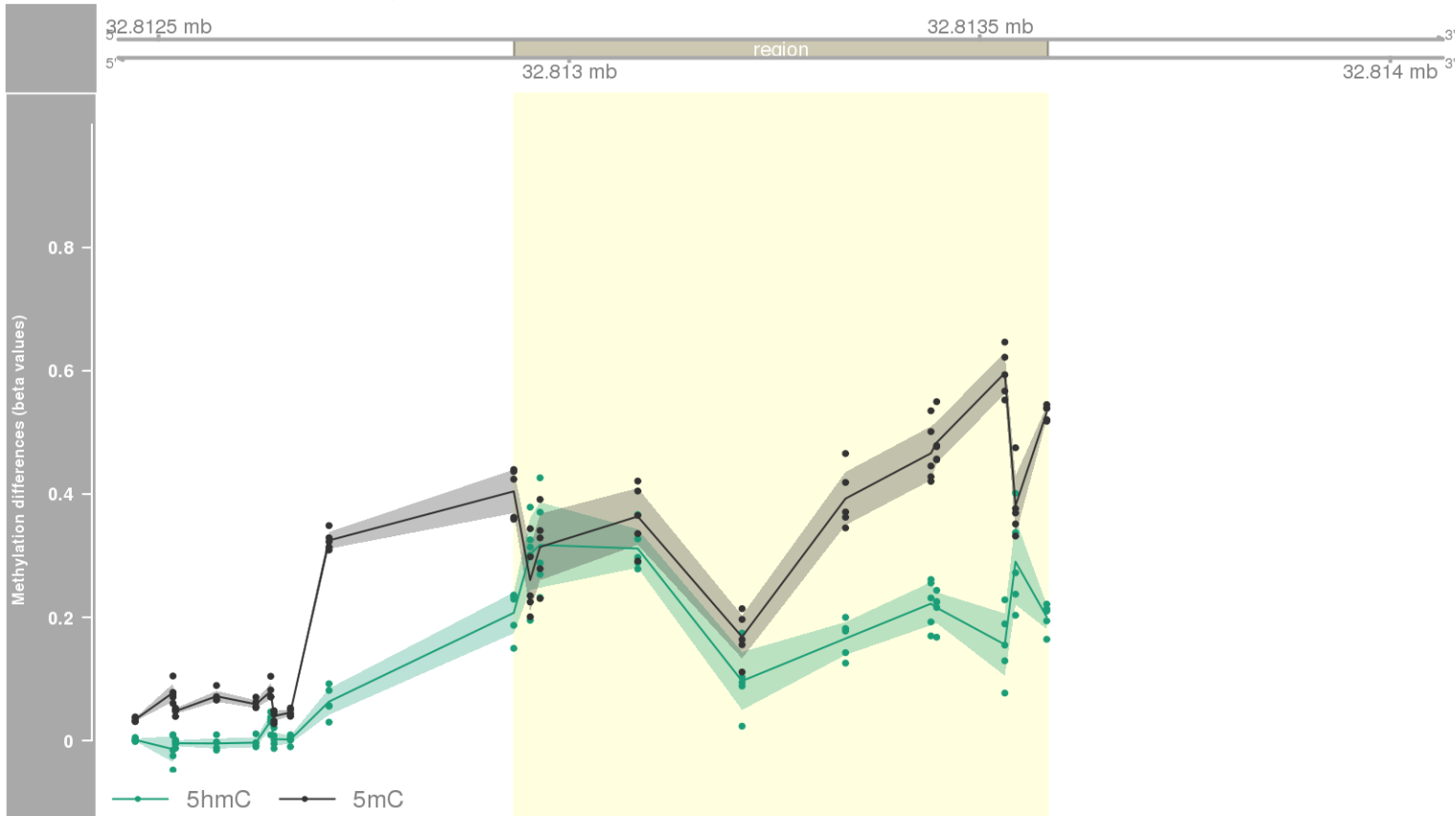
DMR 97 // chr3:55516184-55518798 // 2614 pb. (14 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.007 // fwerArea: 0.846
- genes: WNT5A -



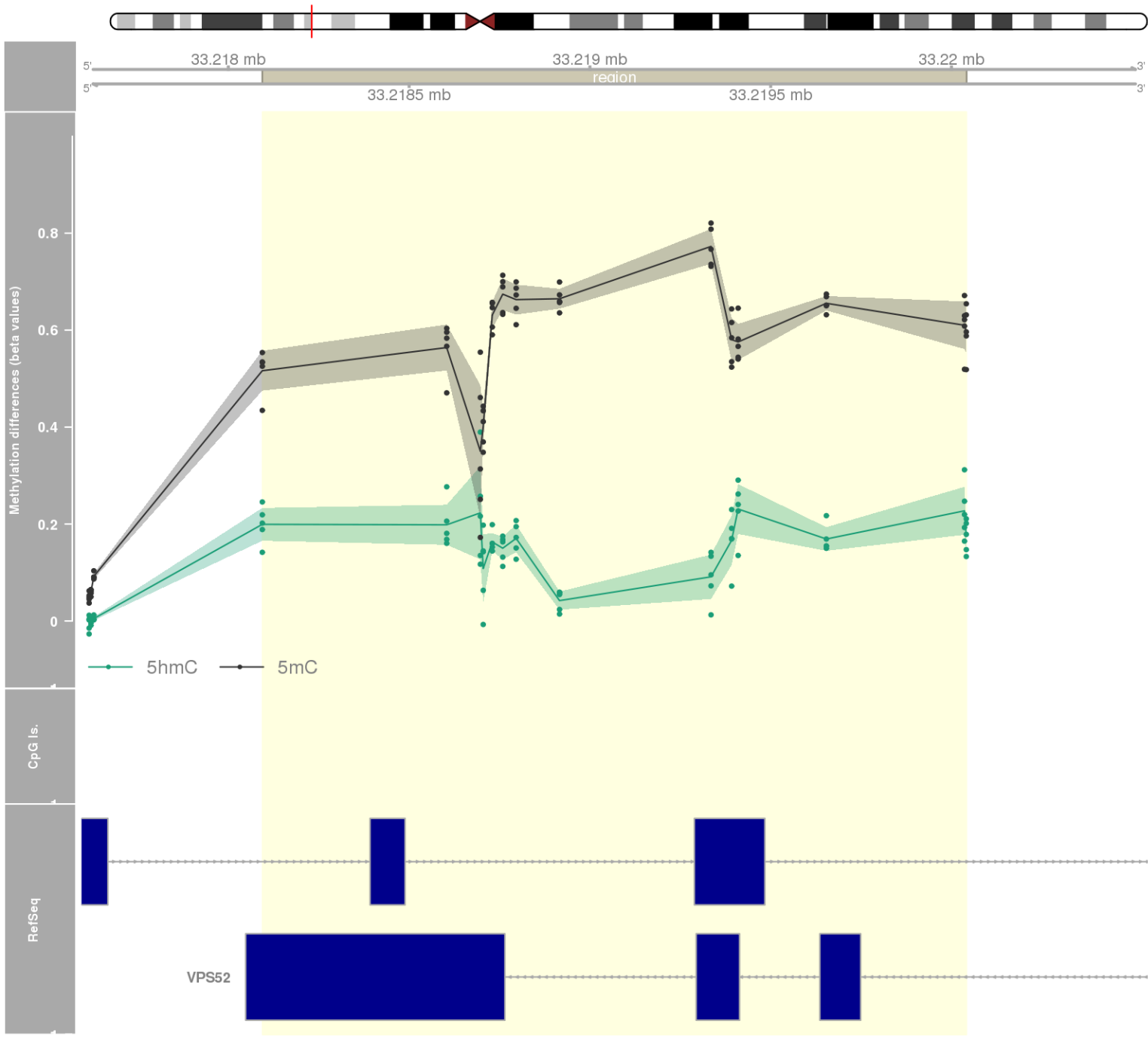
DMR 98 // chr10:131263962-131264846 // 884 pb. (11 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.007 // fwerArea: 0.846



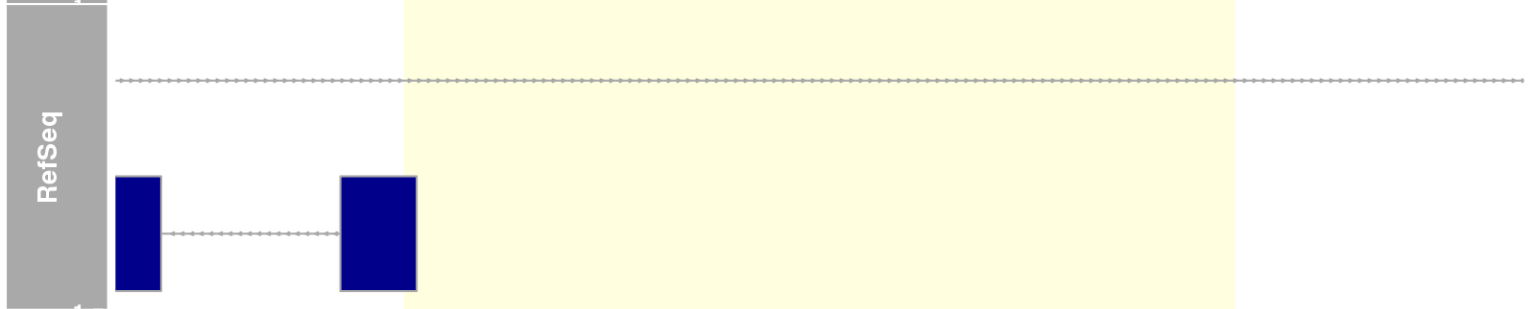
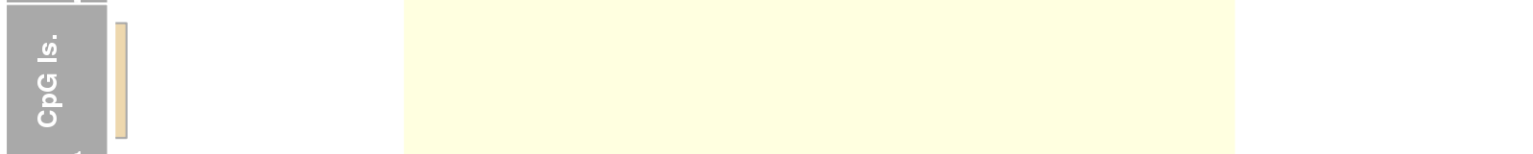
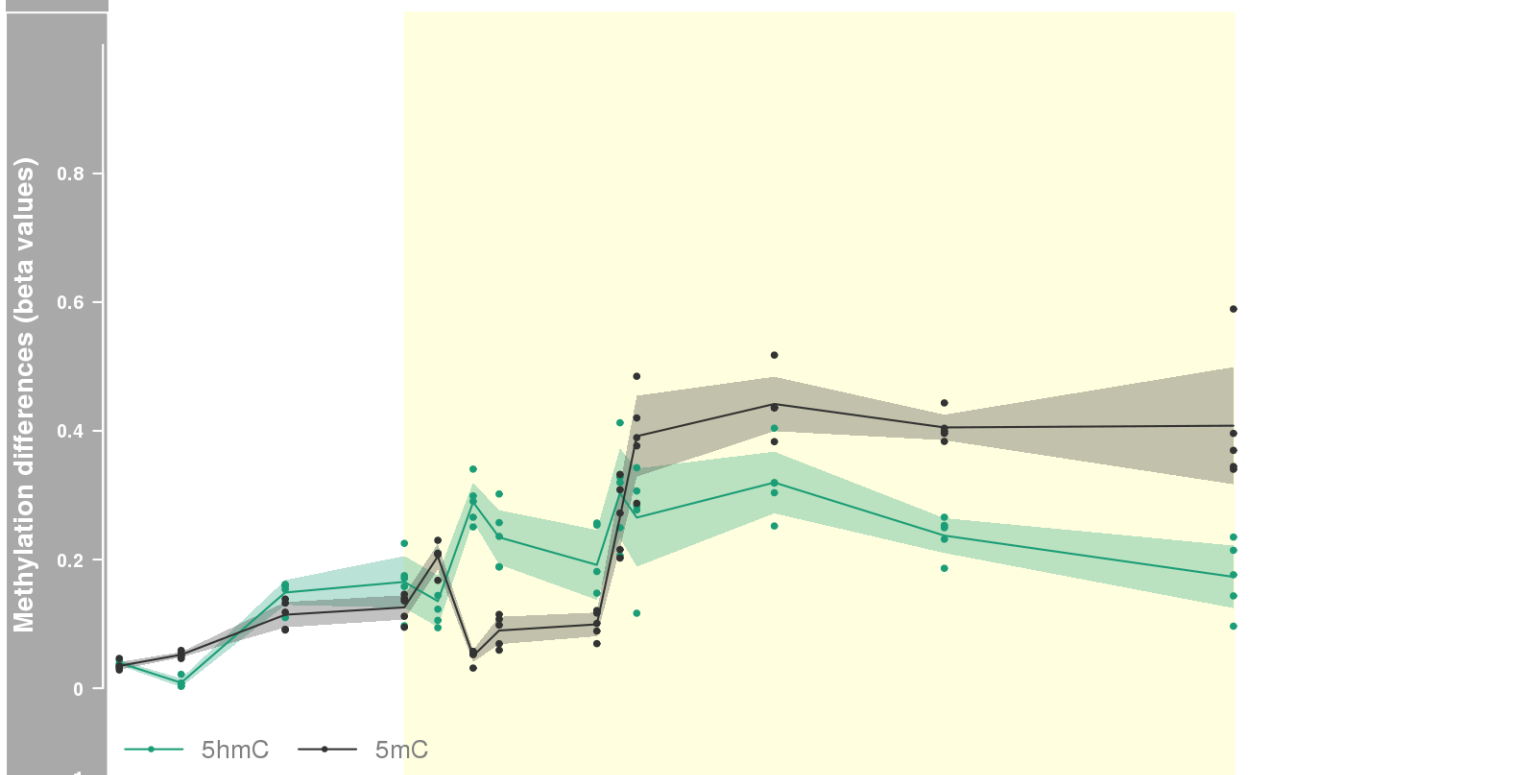
DMR 99 // chr6:32812933-32813582 // 649 pb. (11 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.007 // fwerArea: 0.846
- genes: TAPSAR1 / PSMB9 / TAP1 -



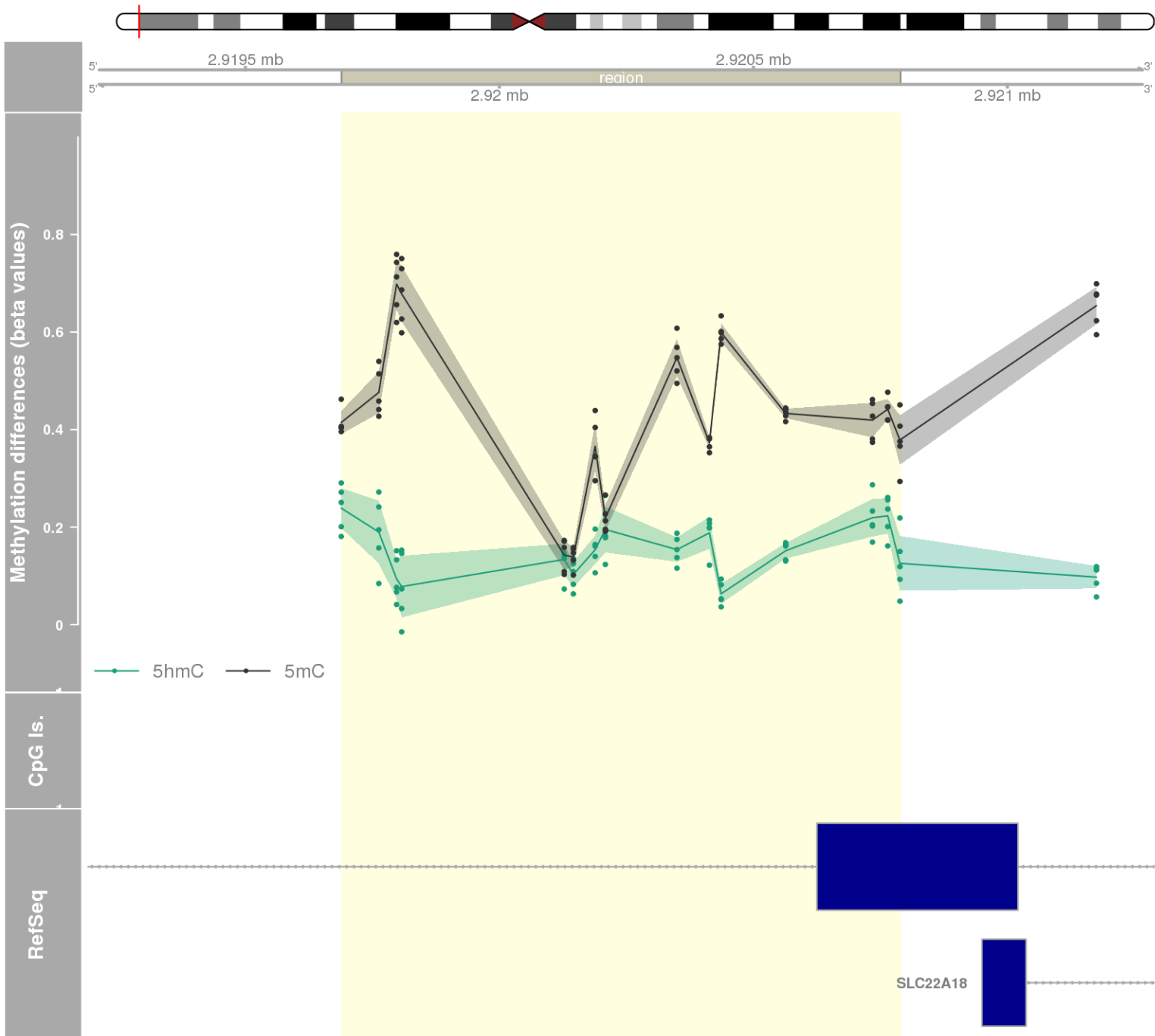
DMR 100 // chr6:33218094-33220041 // 1947 pb. (14 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.007 // fwerArea: 0.846
- genes: VPS52 / HCG25 -



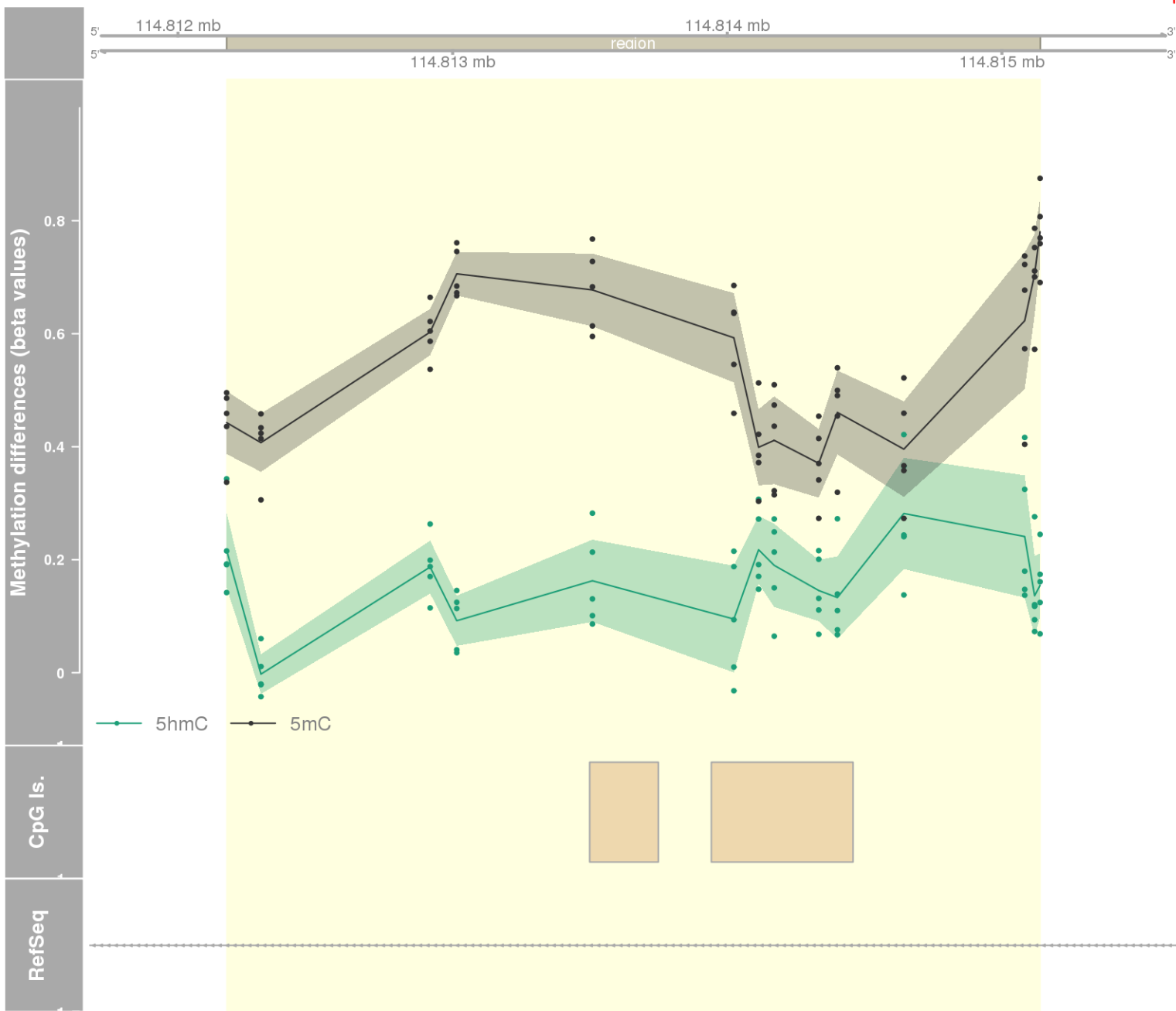
DMR 101 // chr11:67141628-67143062 // 1434 pb. (10 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.007 // fwerArea: 0.846
- genes: CLCF1 -



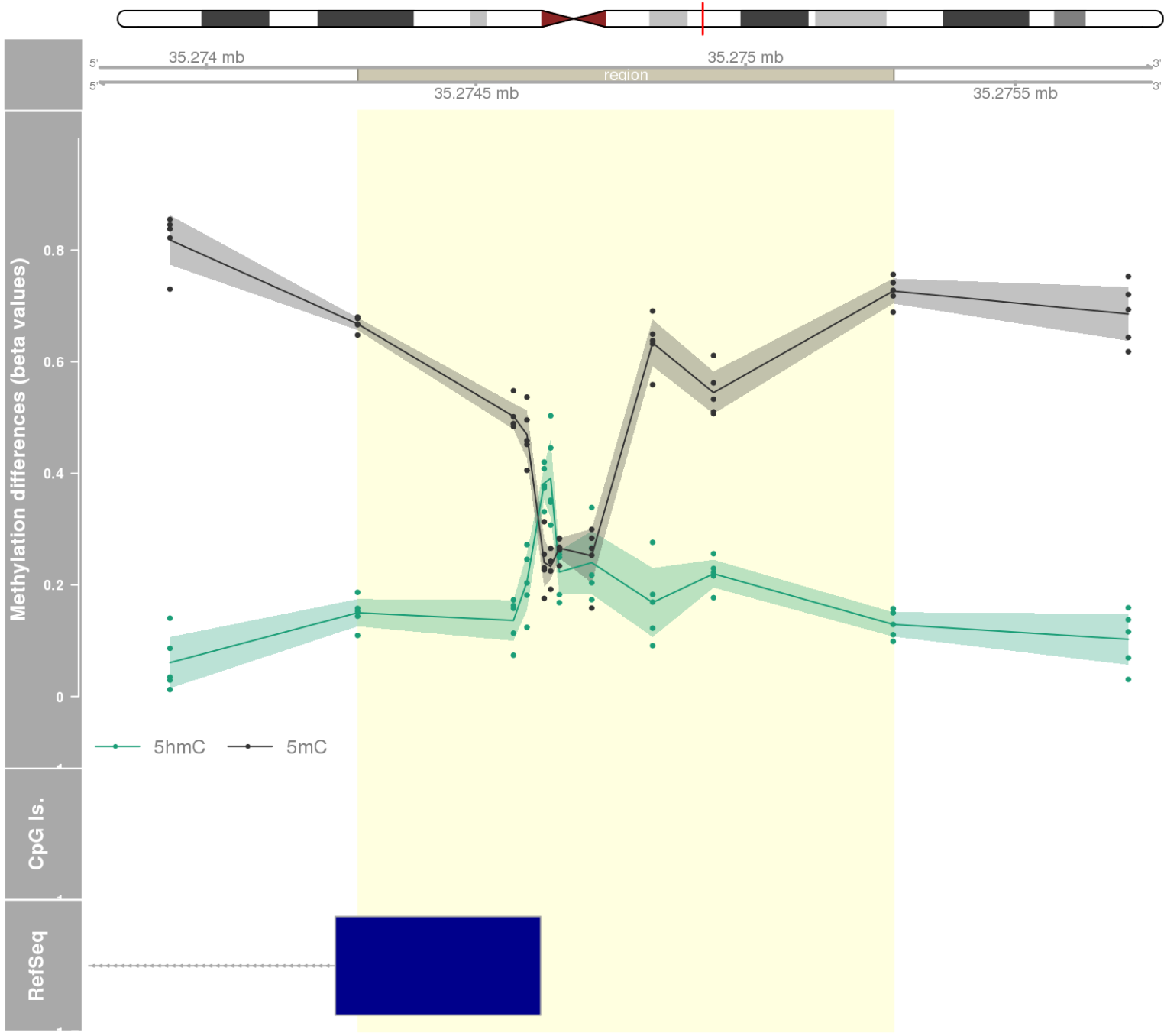
DMR 102 // chr11:2919689-2920789 // 1100 pb. (15 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.007 // fwerArea: 0.846
- genes: SLC22A18AS -



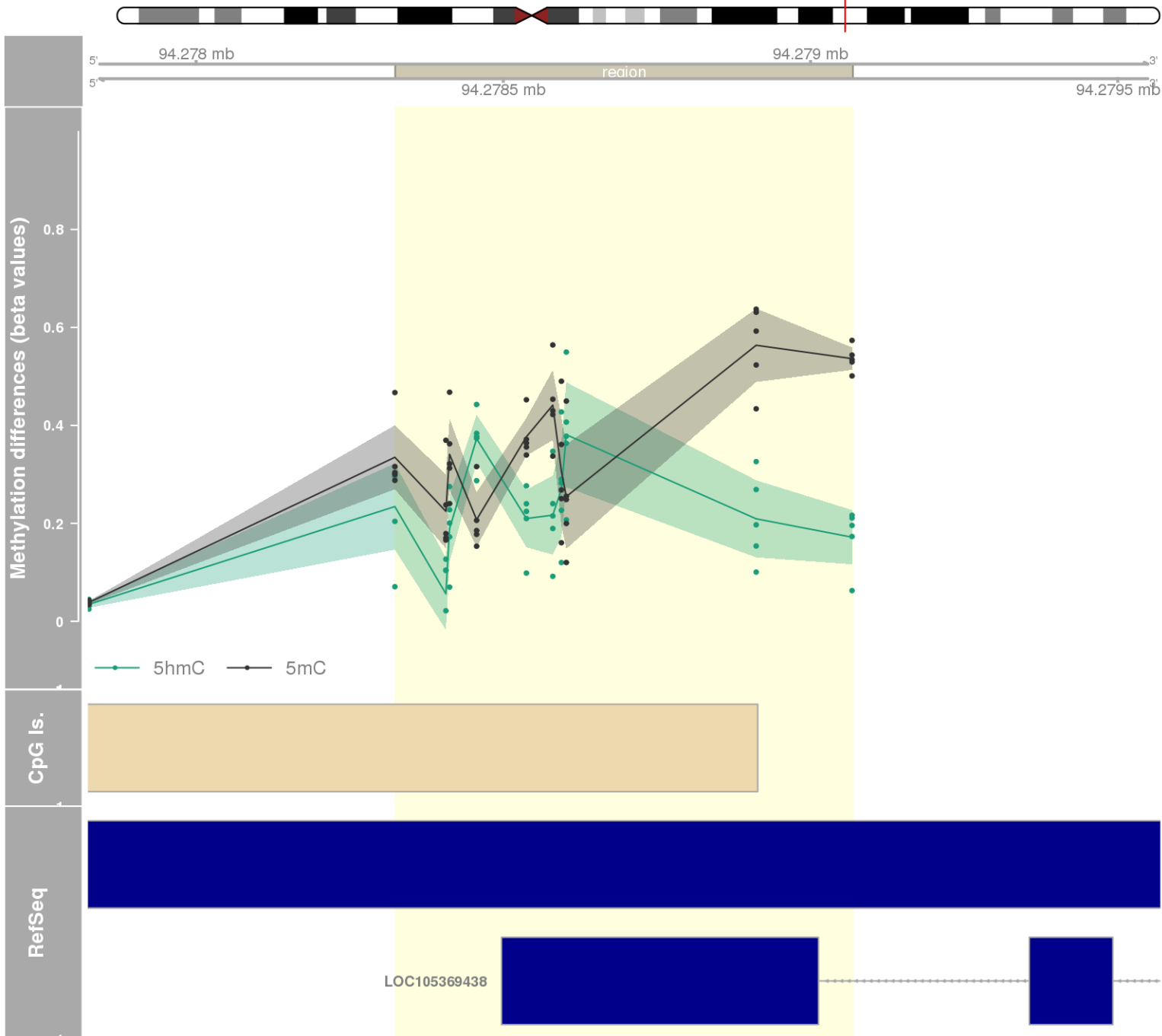
DMR 103 // chr13:114812177-114815139 // 2962 pb. (14 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.007 // fwerArea: 0.846
- genes: RASA3 -



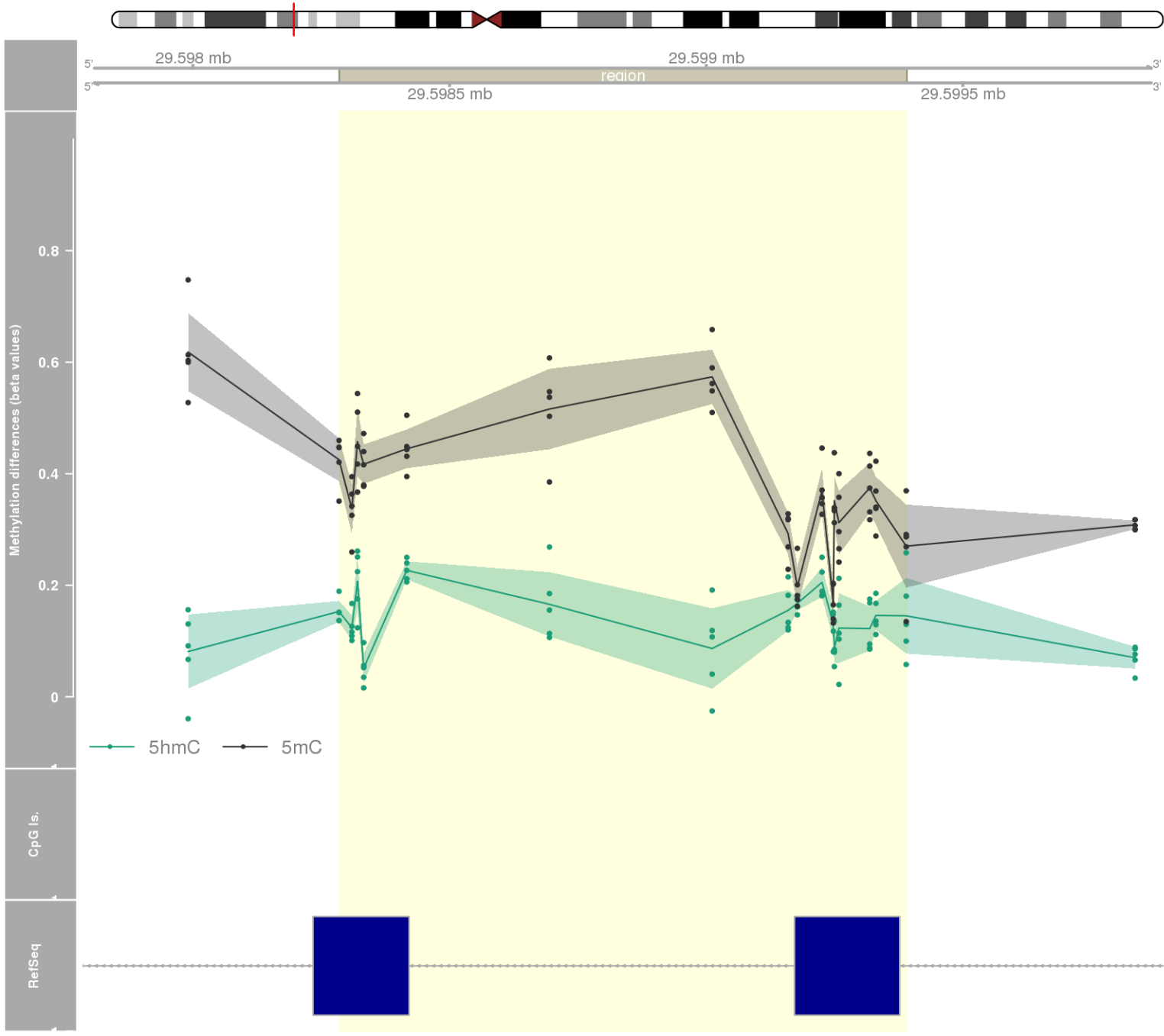
DMR 104 // chr20:35274281-35275274 // 993 pb. (10 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.007 // fwerArea: 0.846
- genes: SLA2 -



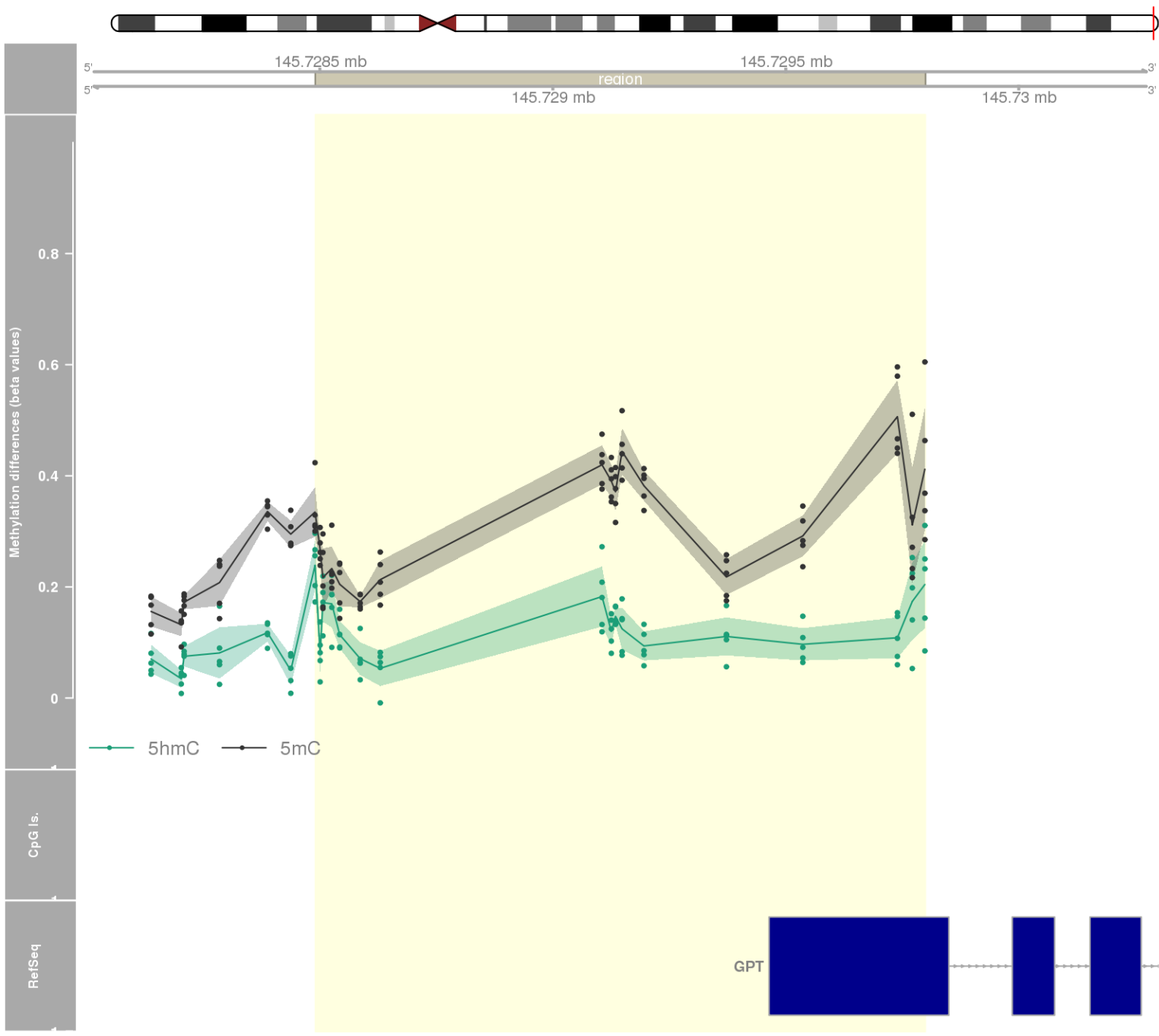
DMR 105 // chr11:94278324-94279068 // 744 pb. (10 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.008 // fwerArea: 0.846
- genes: PIWIL4 / FUT4 -



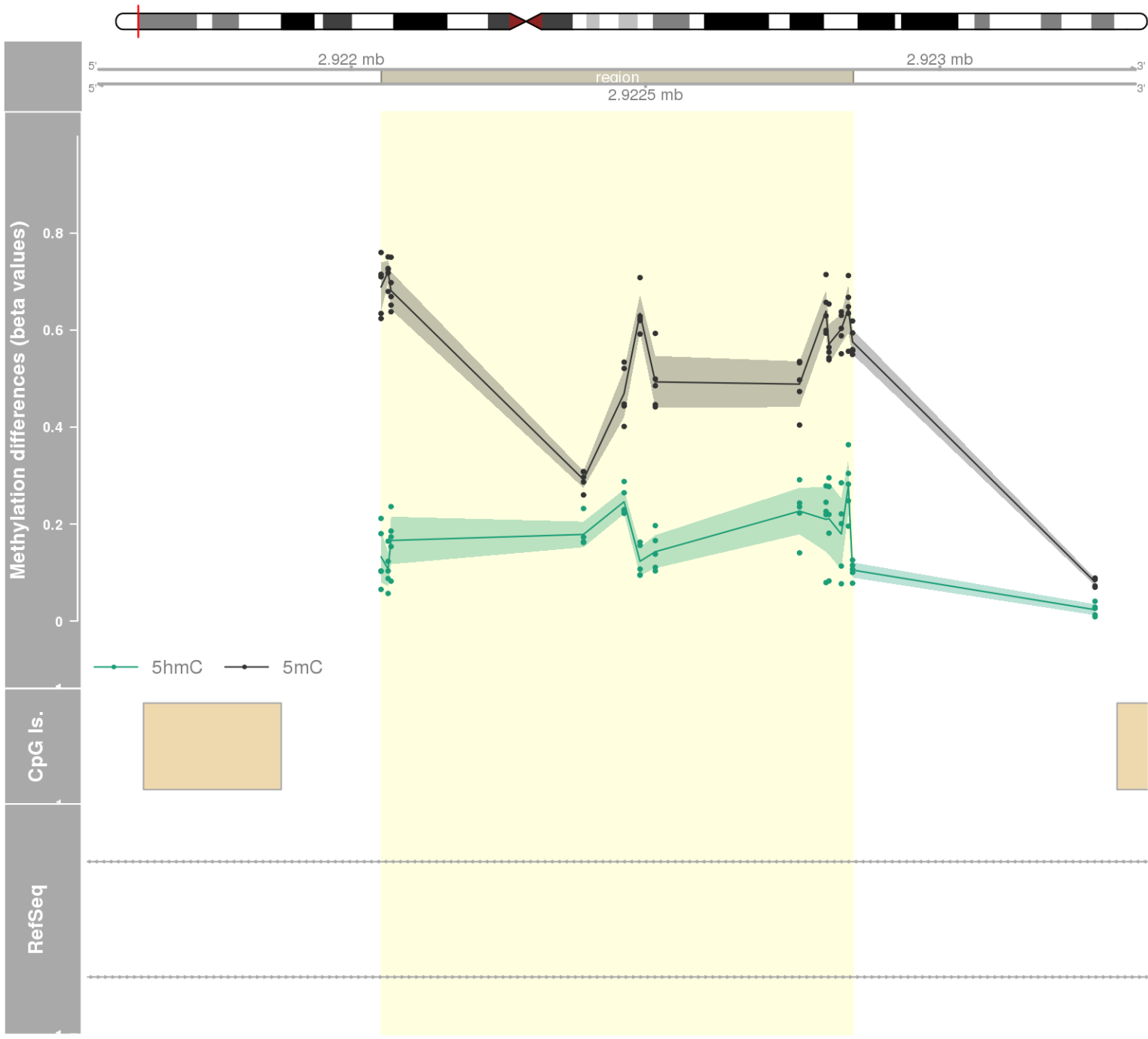
DMR 106 // chr6:29598285-29599390 // 1105 pb. (16 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.008 // fwerArea: 0.846
- genes: GABBR1 -



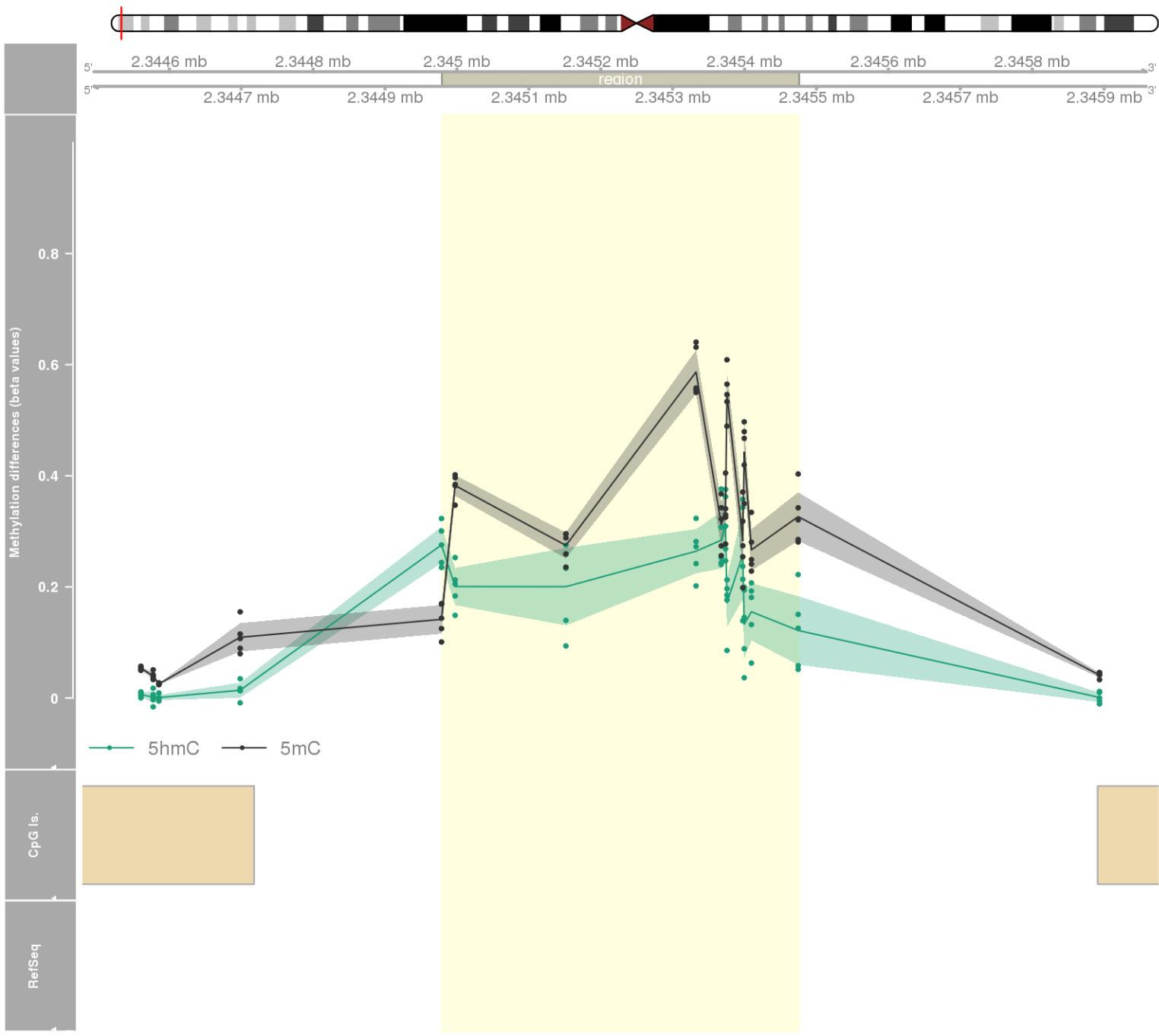
DMR 107 // chr8:145728490-145729799 // 1309 pb. (17 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.008 // fwerArea: 0.846
- genes: GPT -



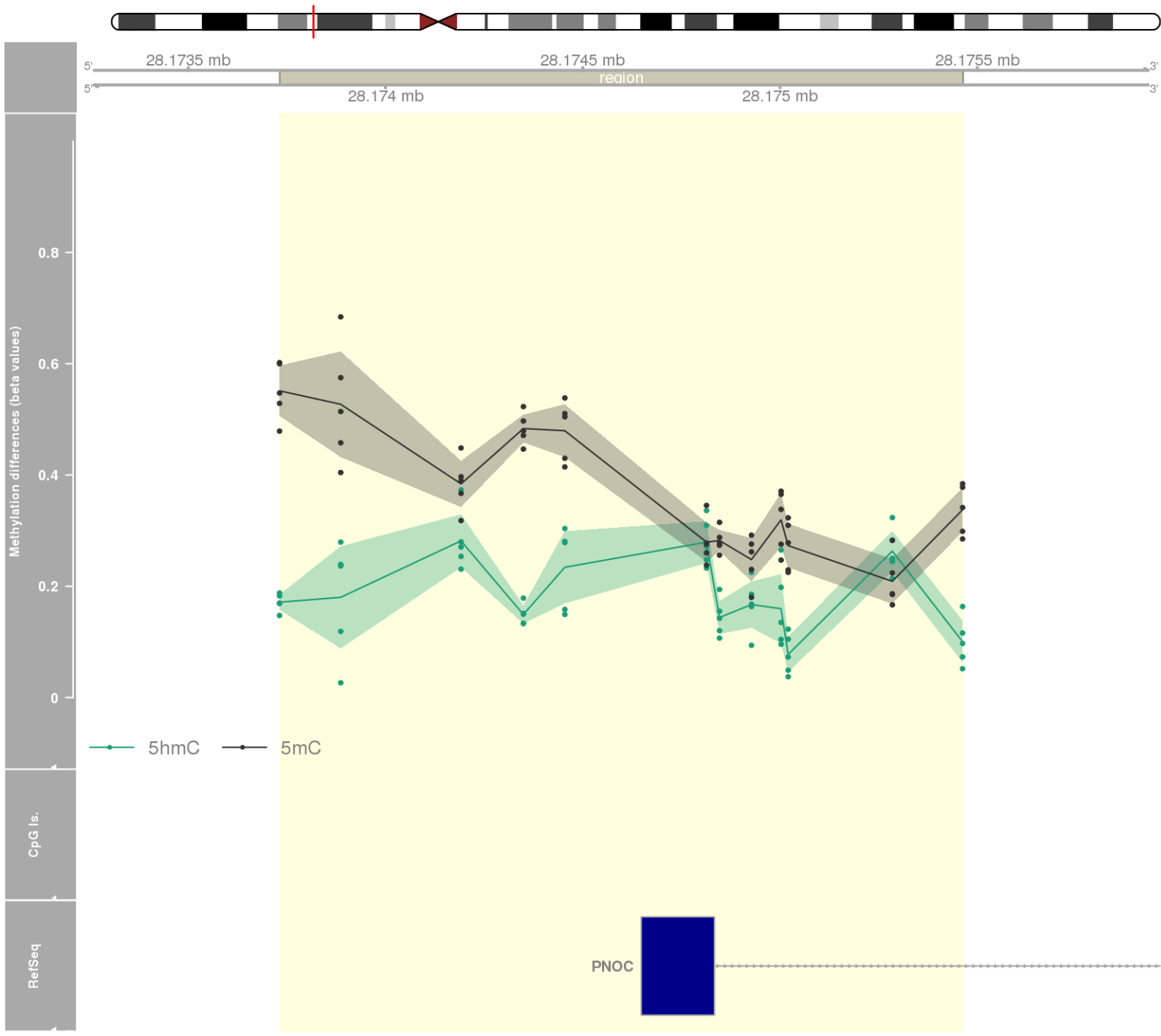
DMR 108 // chr11:2922051-2922852 // 801 pb. (13 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.008 // fwerArea: 0.846
- genes: SLC22A18AS / SLC22A18 -



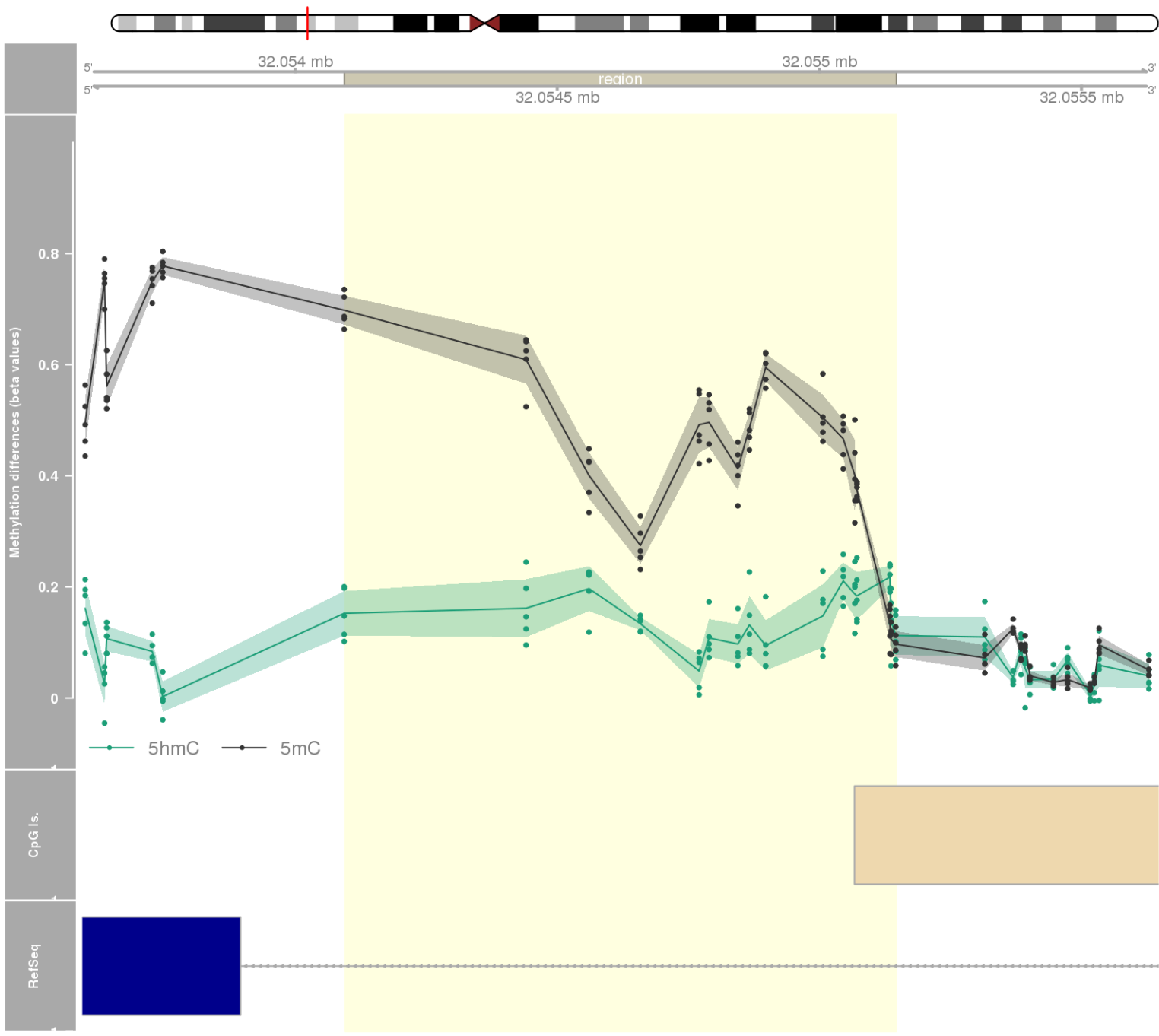
DMR 109 // chr1:2344979-2345475 // 496 pb. (11 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.008 // fwerArea: 0.846
- genes: PEX10 -



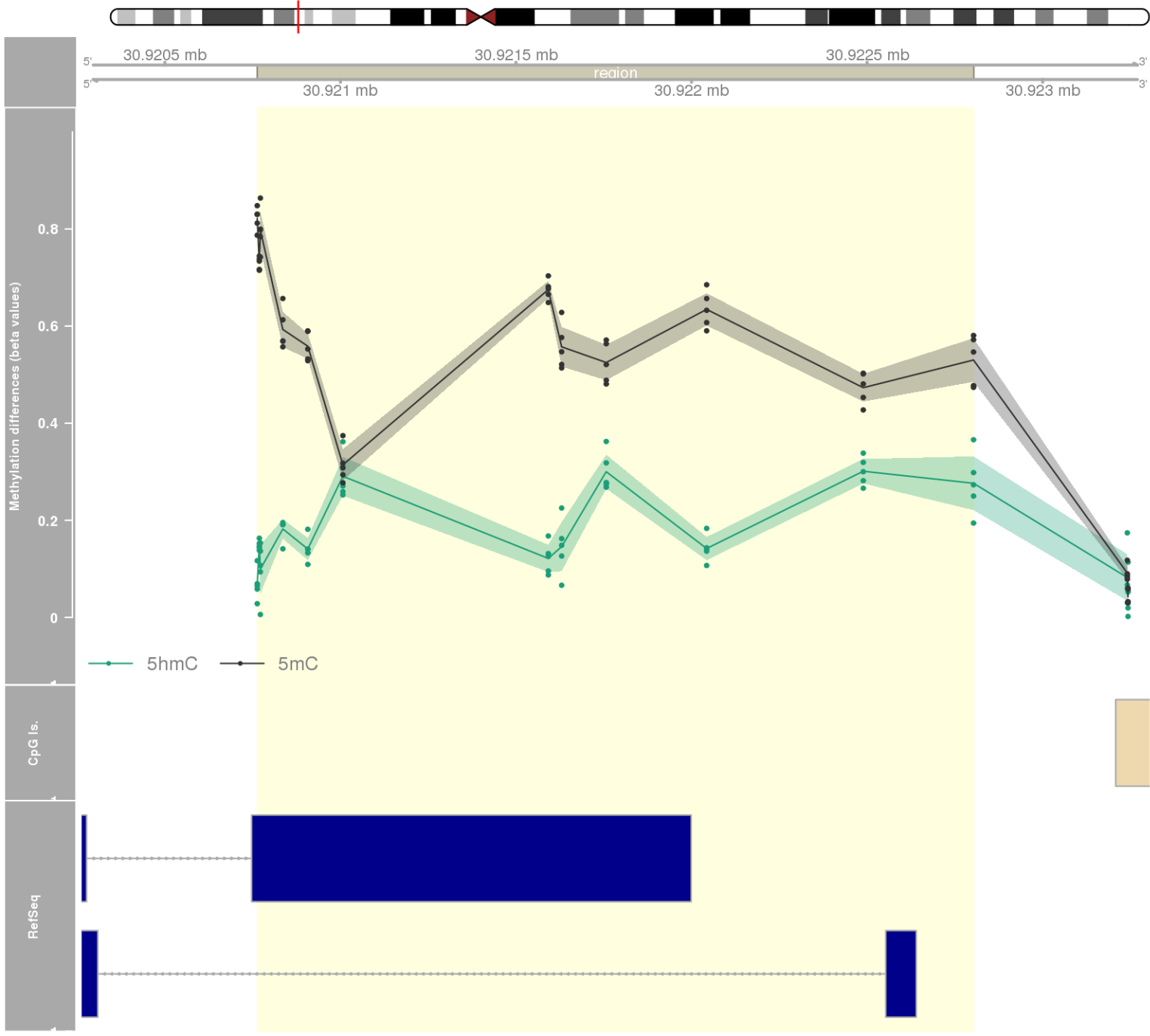
DMR 110 // chr8:28173732-28175463 // 1731 pb. (12 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.008 // fwerArea: 0.846
- genes: PNOC -



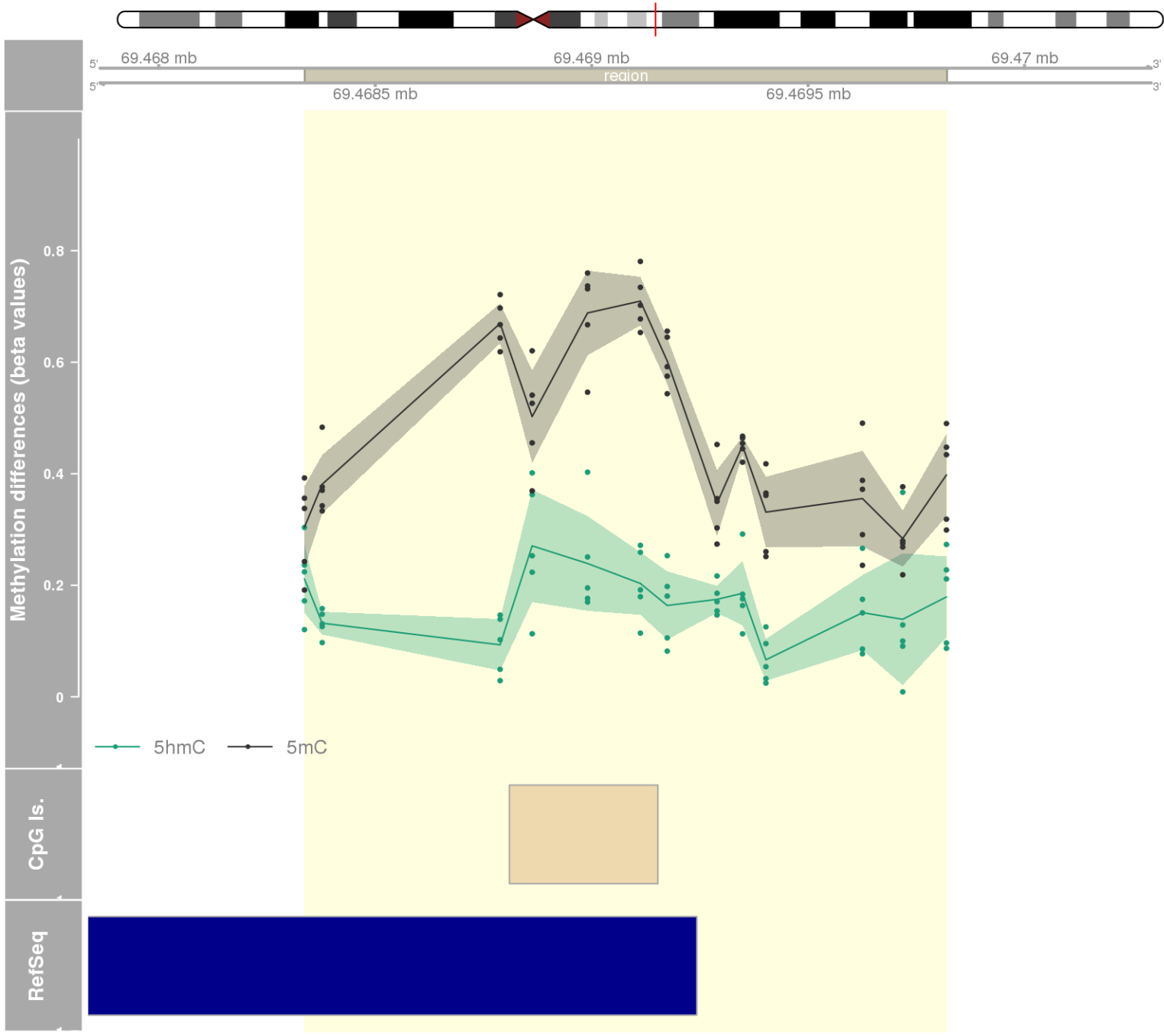
DMR 111 // chr6:32054094-32055146 // 1052 pb. (16 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.008 // fwerArea: 0.846
- genes: TNXB -



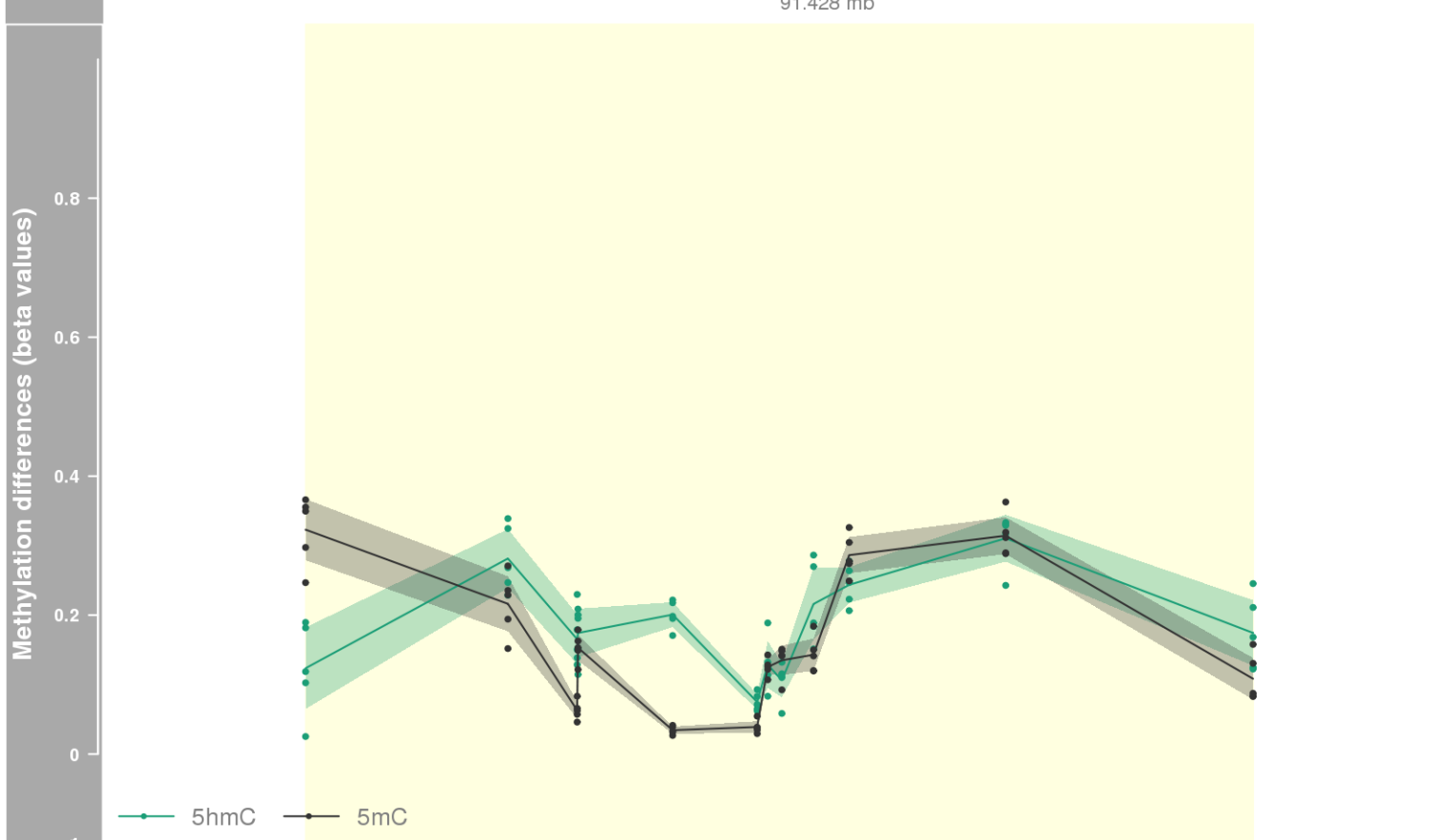
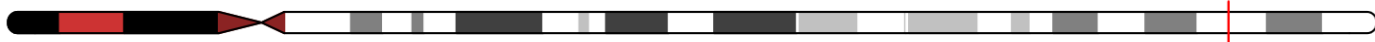
DMR 112 // chr6:30920763-30922803 // 2040 pb. (12 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.008 // fwerArea: 0.846
- genes: HCG21 / DPCR1 -



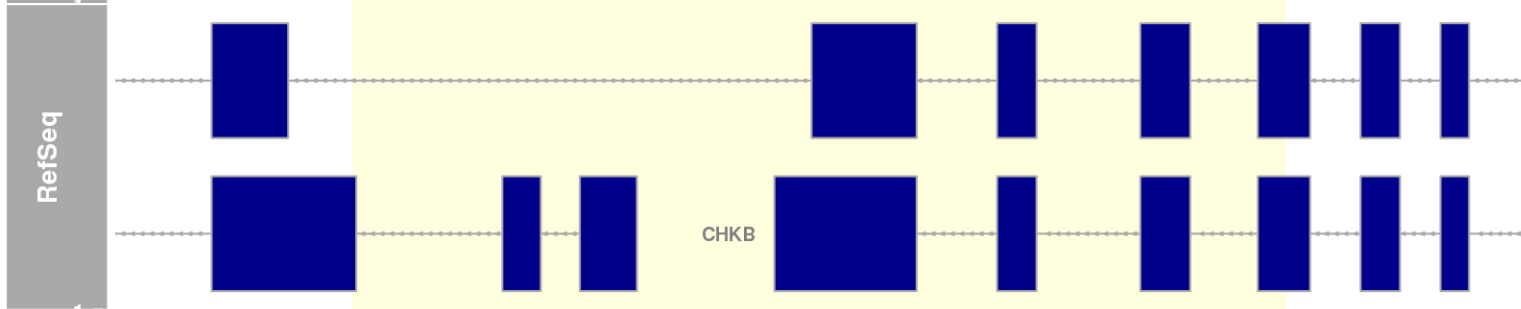
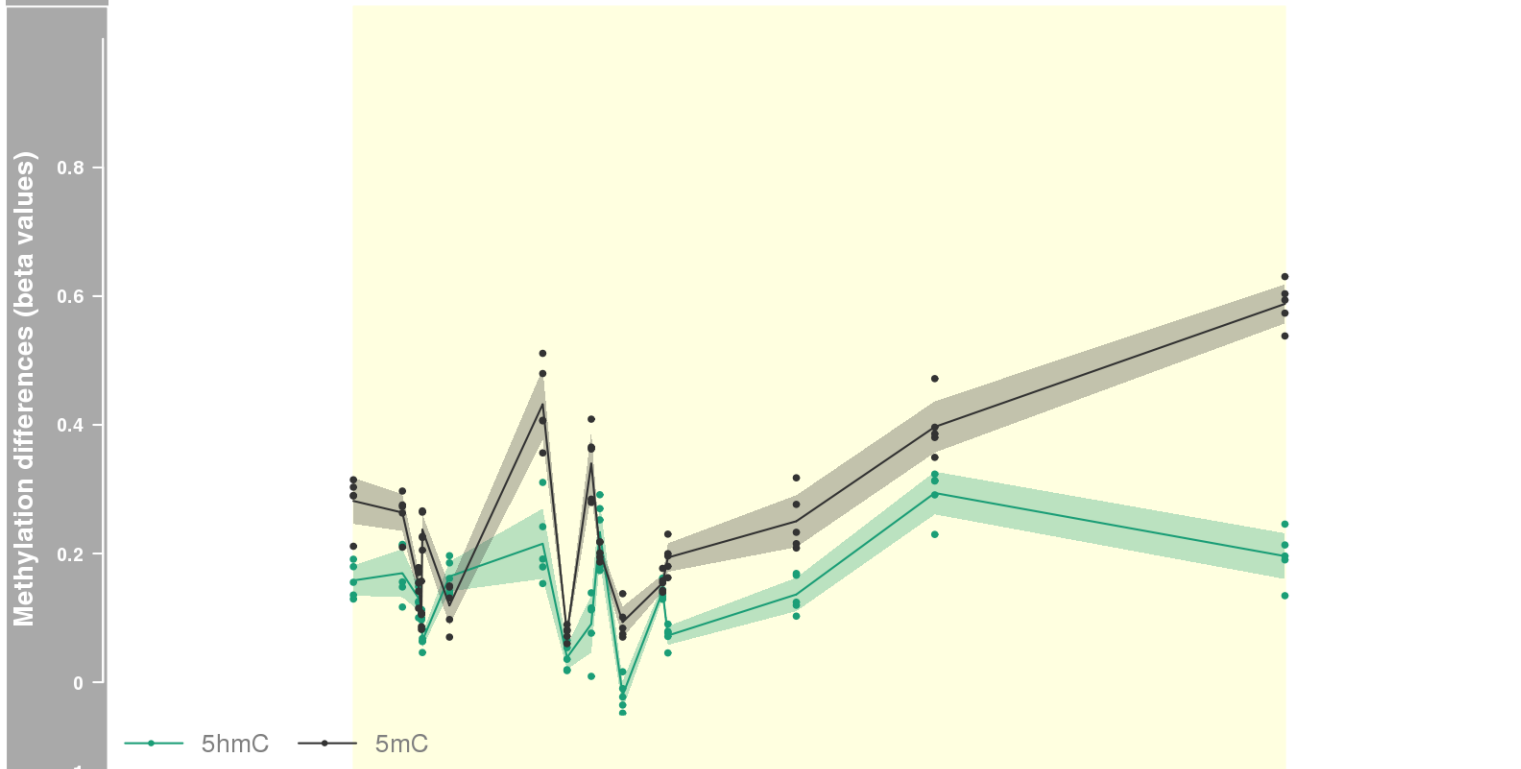
DMR 113 // chr11:69468337-69469820 // 1483 pb. (13 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.008 // fwerArea: 0.846
- genes: CCND1 / ORAOV1 -



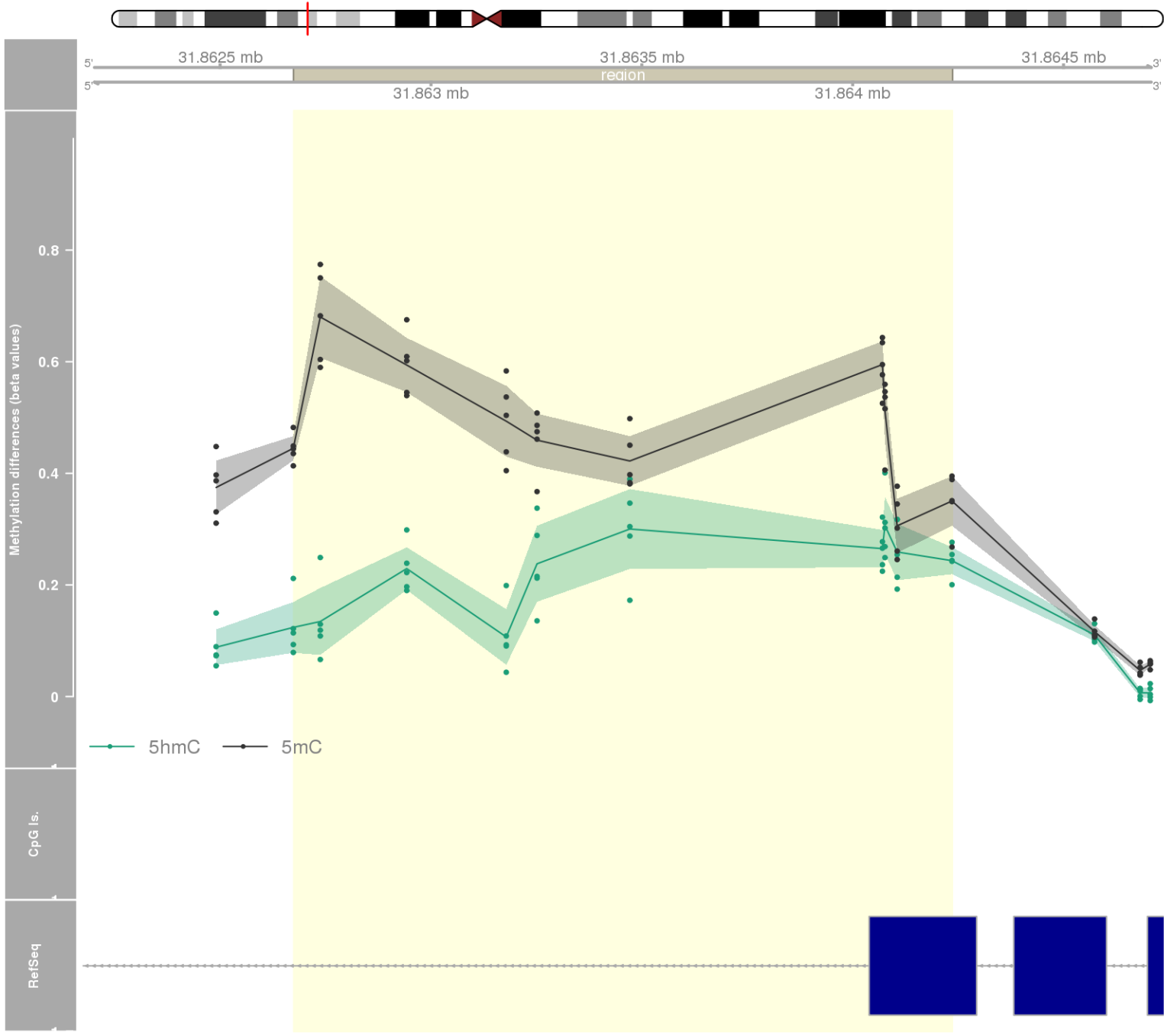
DMR 114 // chr15:91426667-91429088 // 2421 pb. (12 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.008 // fwerArea: 0.846
- genes: *FURIN* / *FES* -



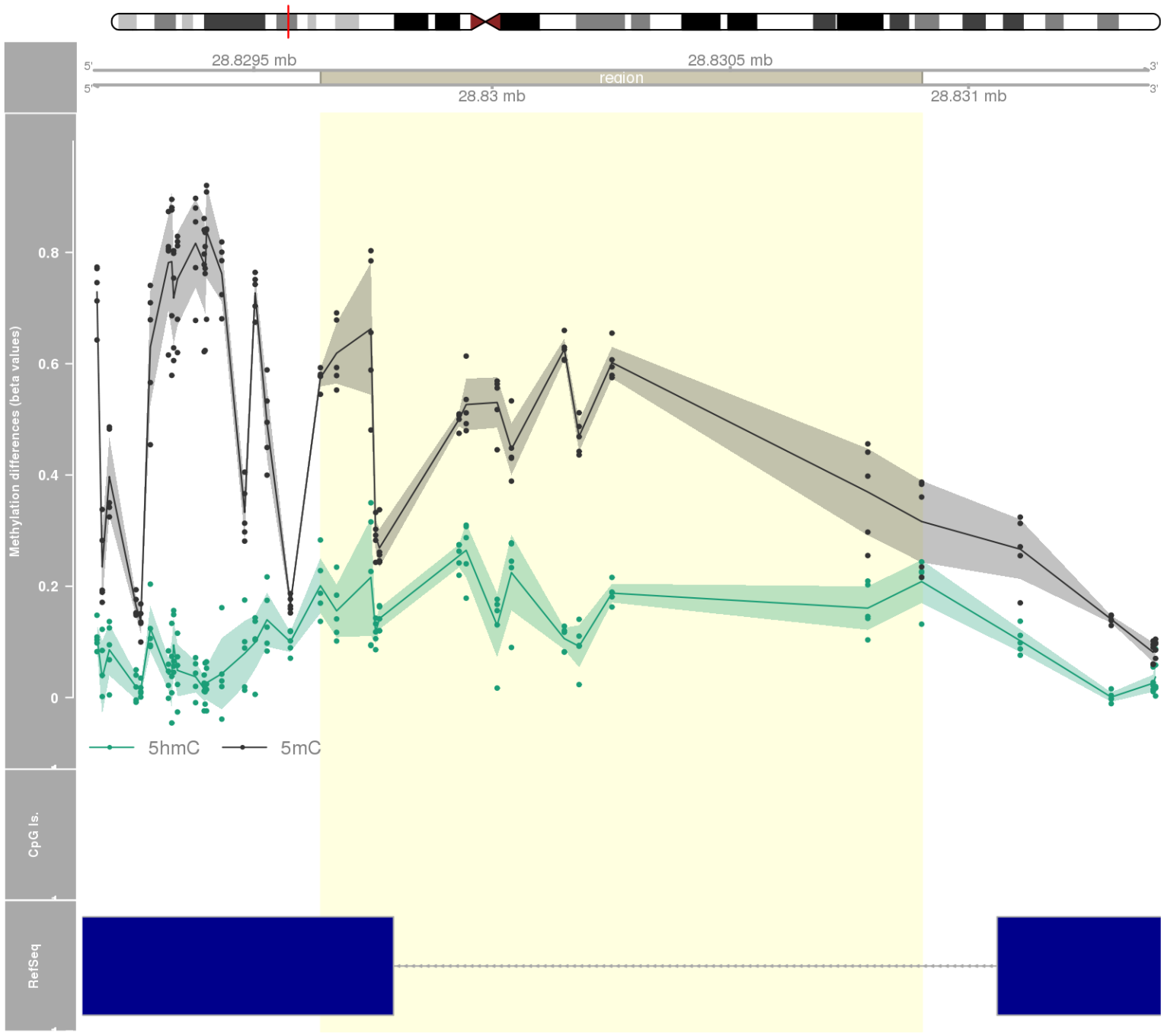
DMR 115 // chr22:51016501-51018459 // 1958 pb. (16 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.008 // fwerArea: 1
- genes: CPT1B / CHKB-CPT1B / CHKB -



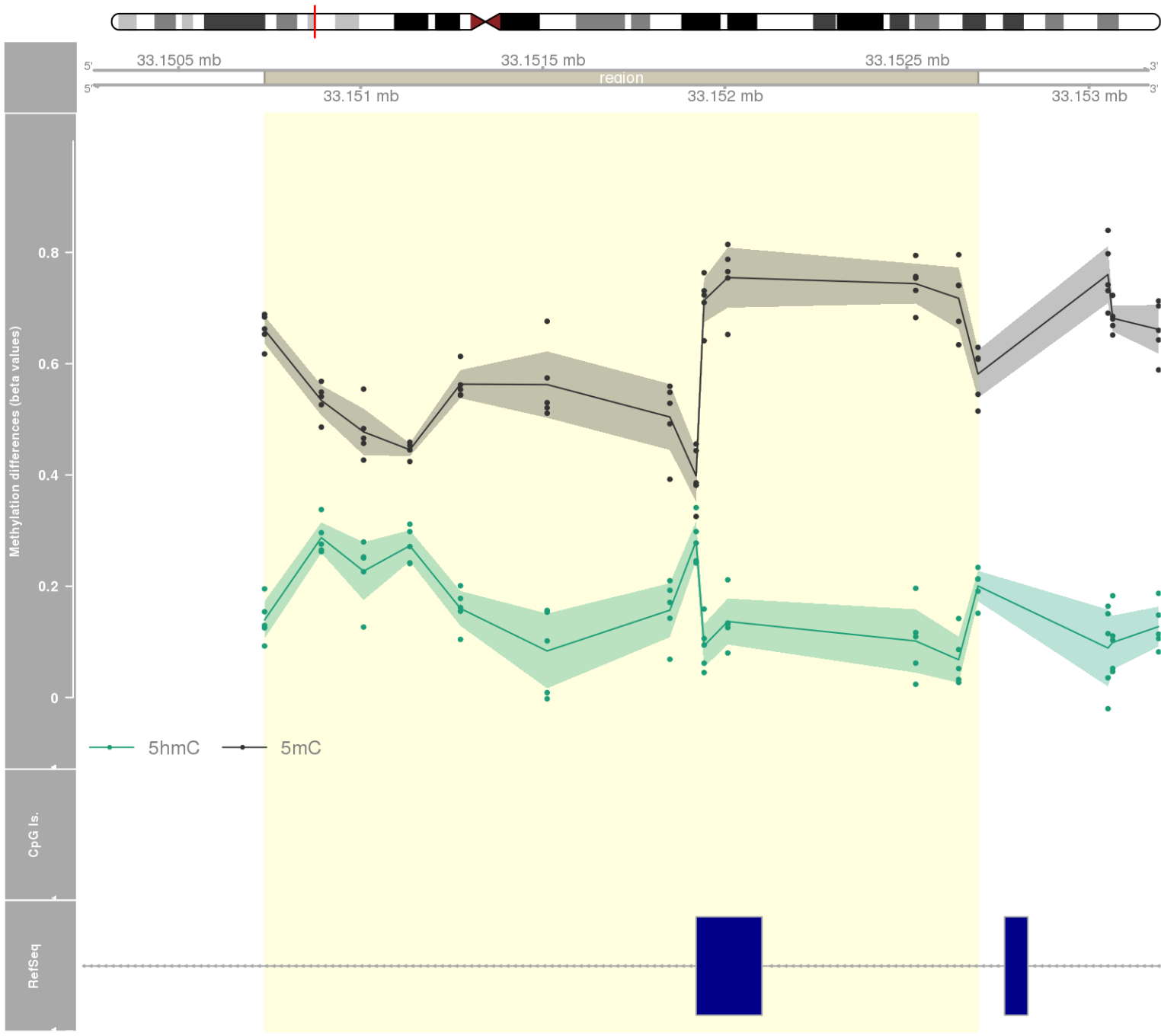
DMR 116 // chr6:31862674-31864236 // 1562 pb. (10 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.008 // fwerArea: 1
- genes: EHMT2 -



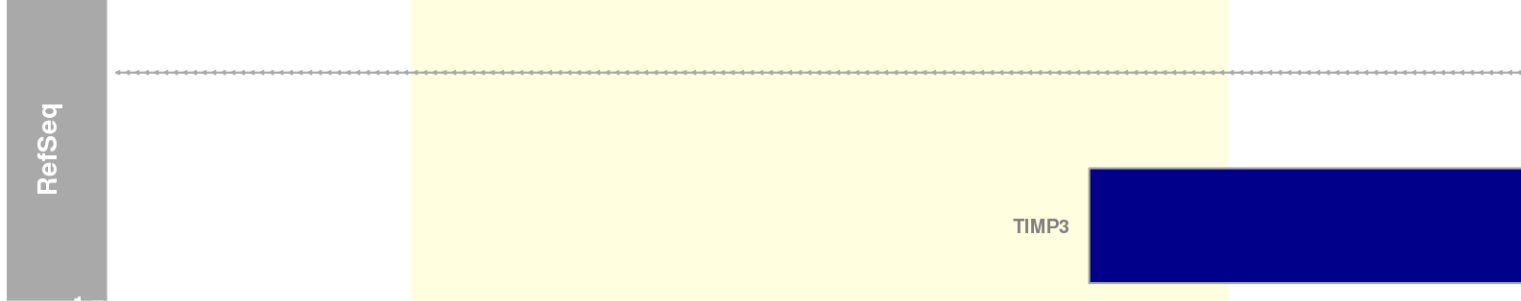
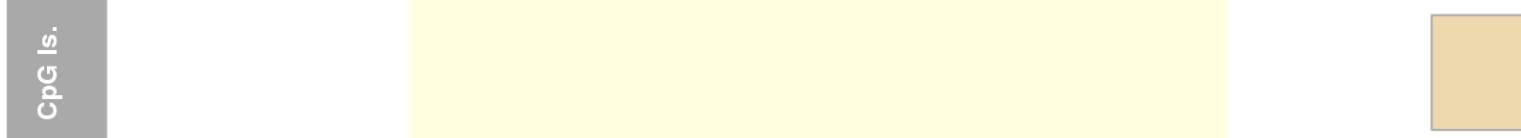
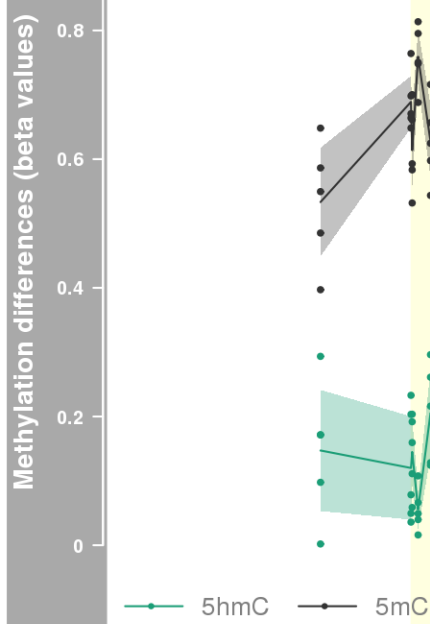
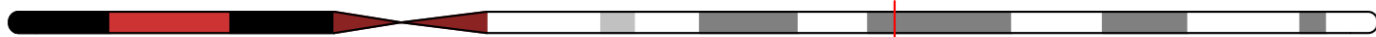
DMR 117 // chr6:28829640-28830902 // 1262 pb. (14 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.008 // fwerArea: 1
- genes: RPL13P -



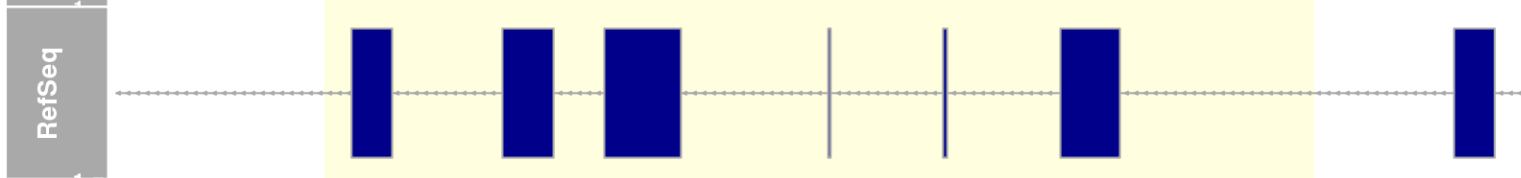
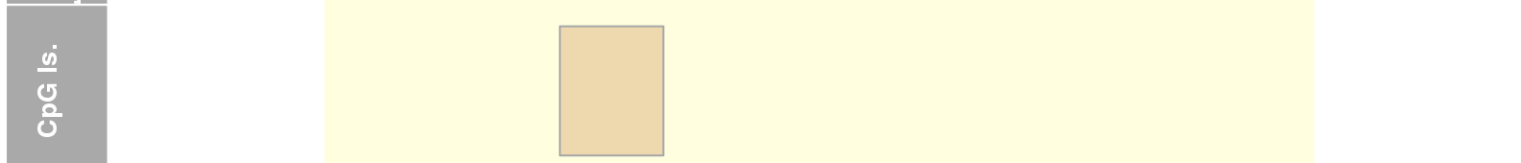
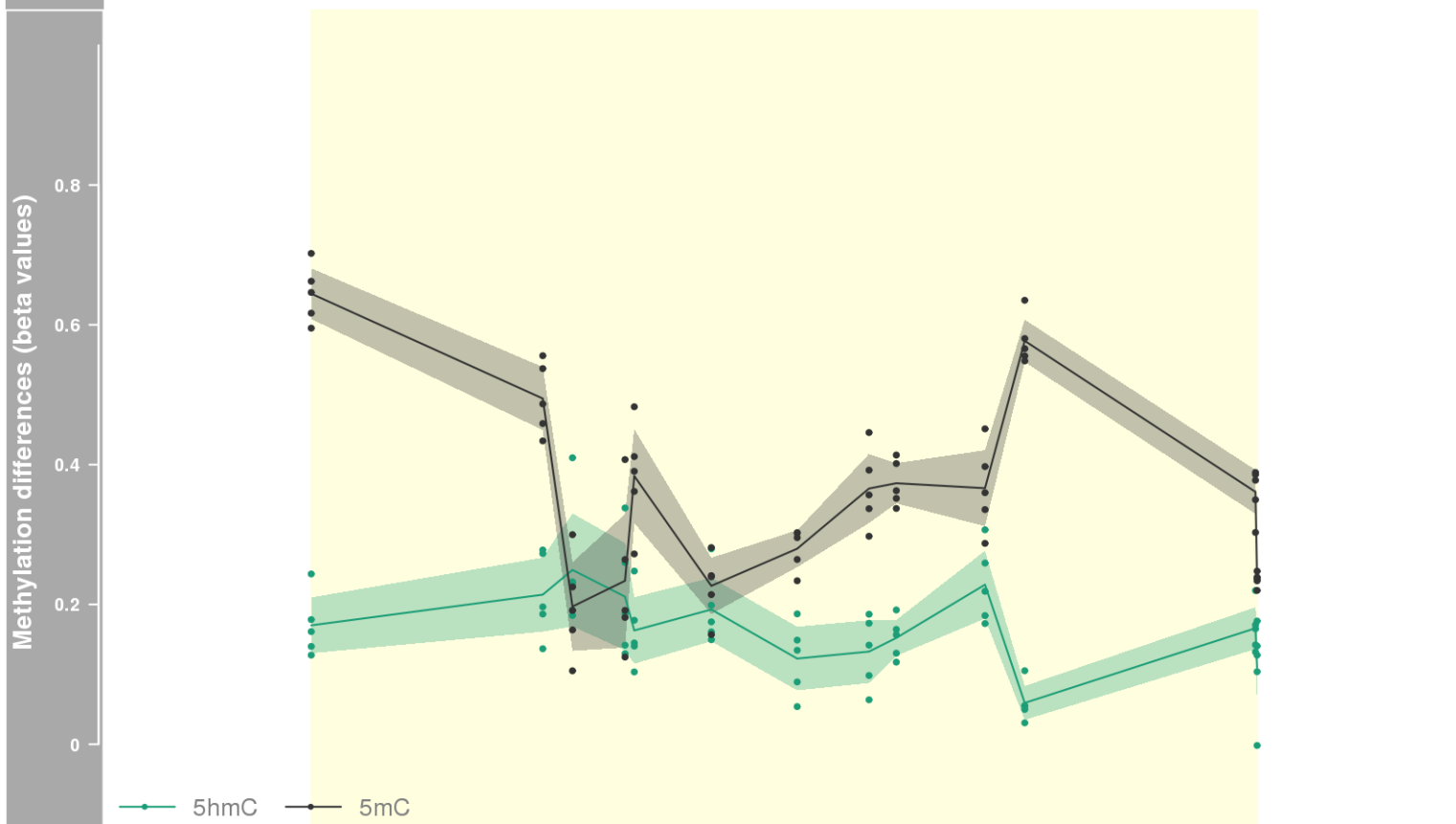
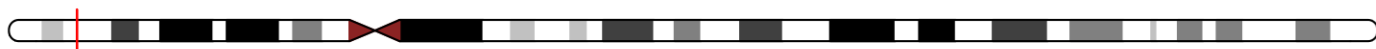
DMR 118 // chr6:33150736-33152695 // 1959 pb. (13 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.008 // fwerArea: 1
- genes: COL11A2 -



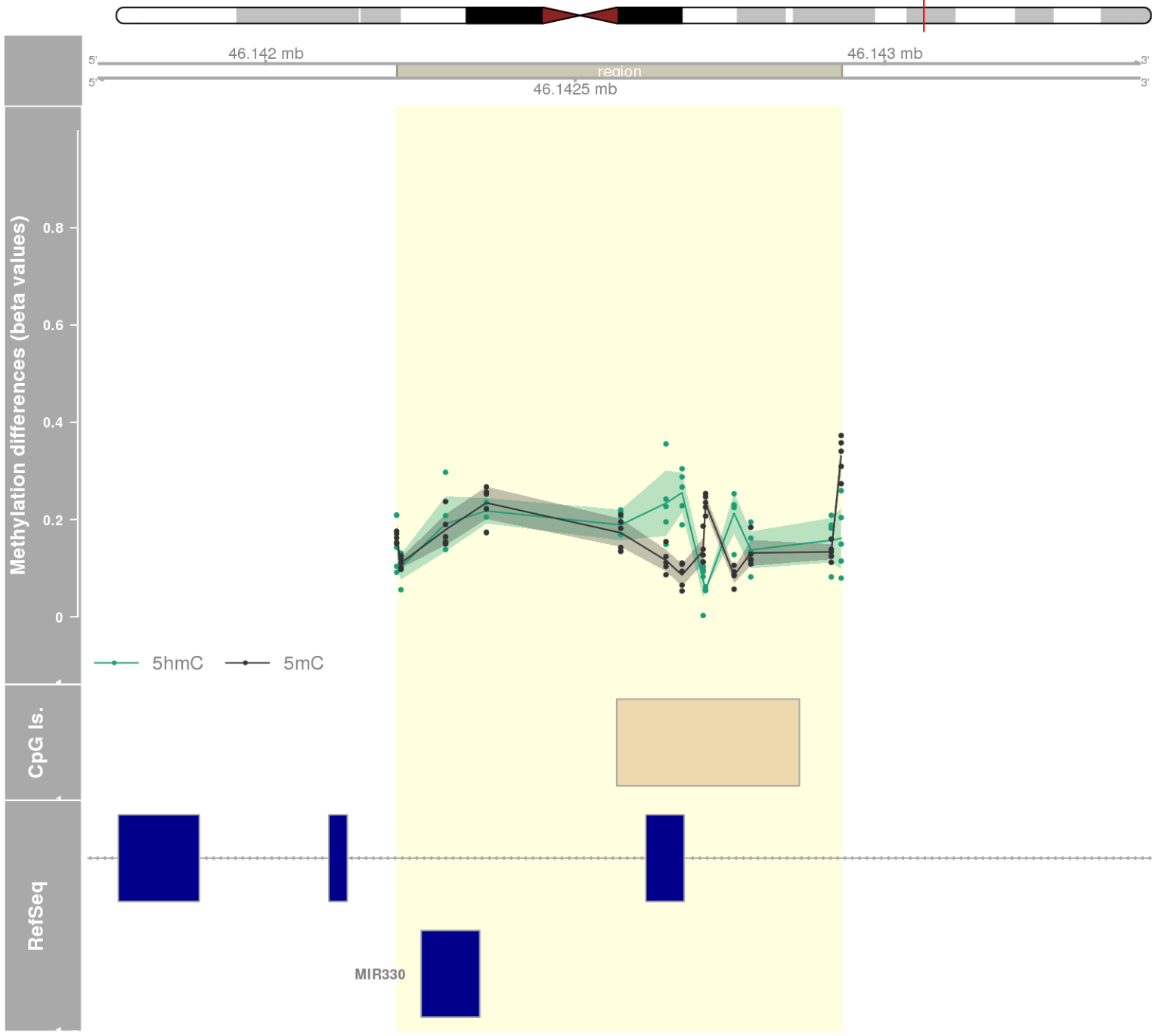
DMR 119 // chr22:33195655-33197034 // 1379 pb. (11 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.008 // fwerArea: 1
- genes: SYN3 -



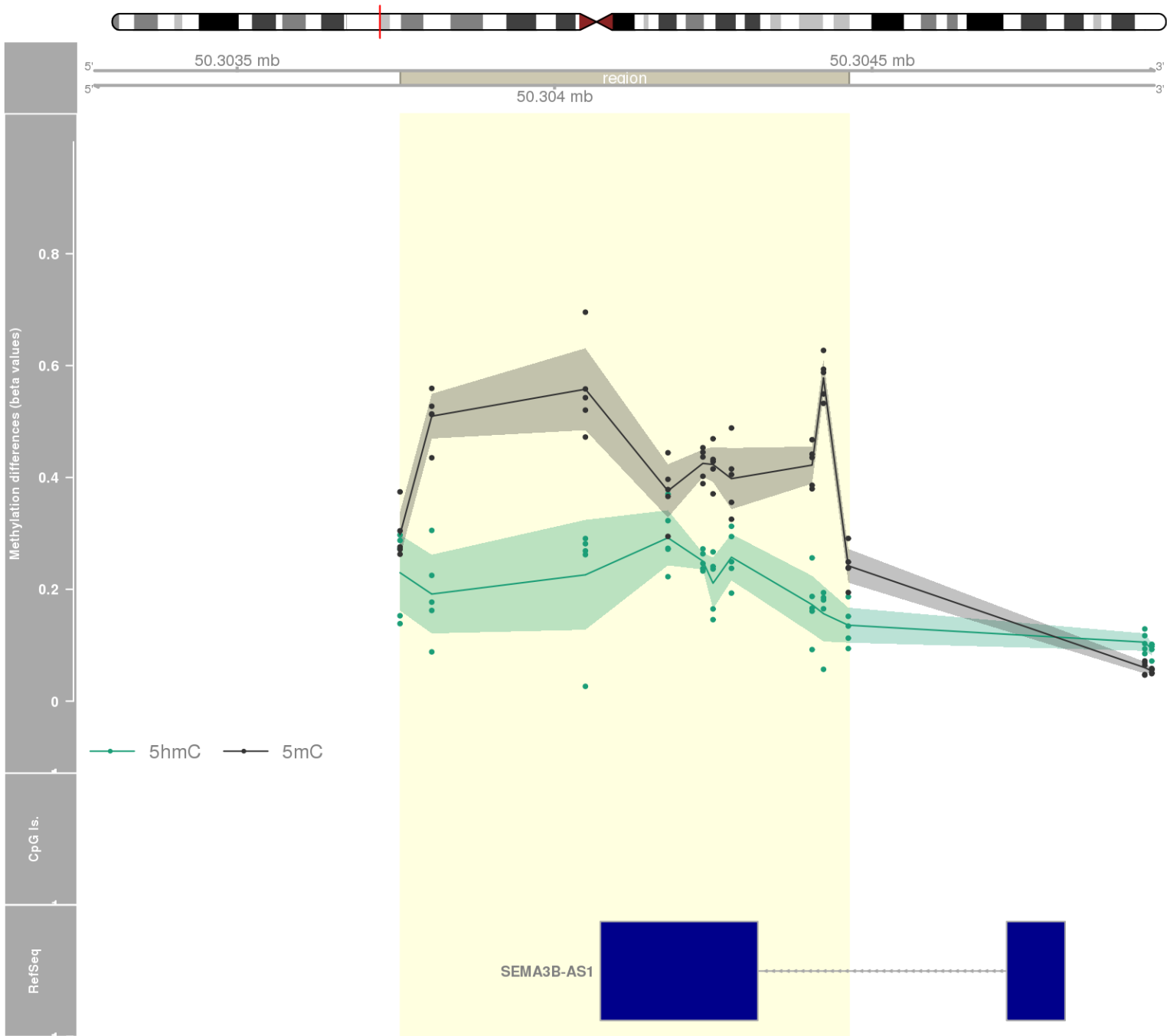
DMR 120 // chr12:6657167-6659524 // 2357 pb. (13 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.009 // fwerArea: 1
- genes: IFFO1 -



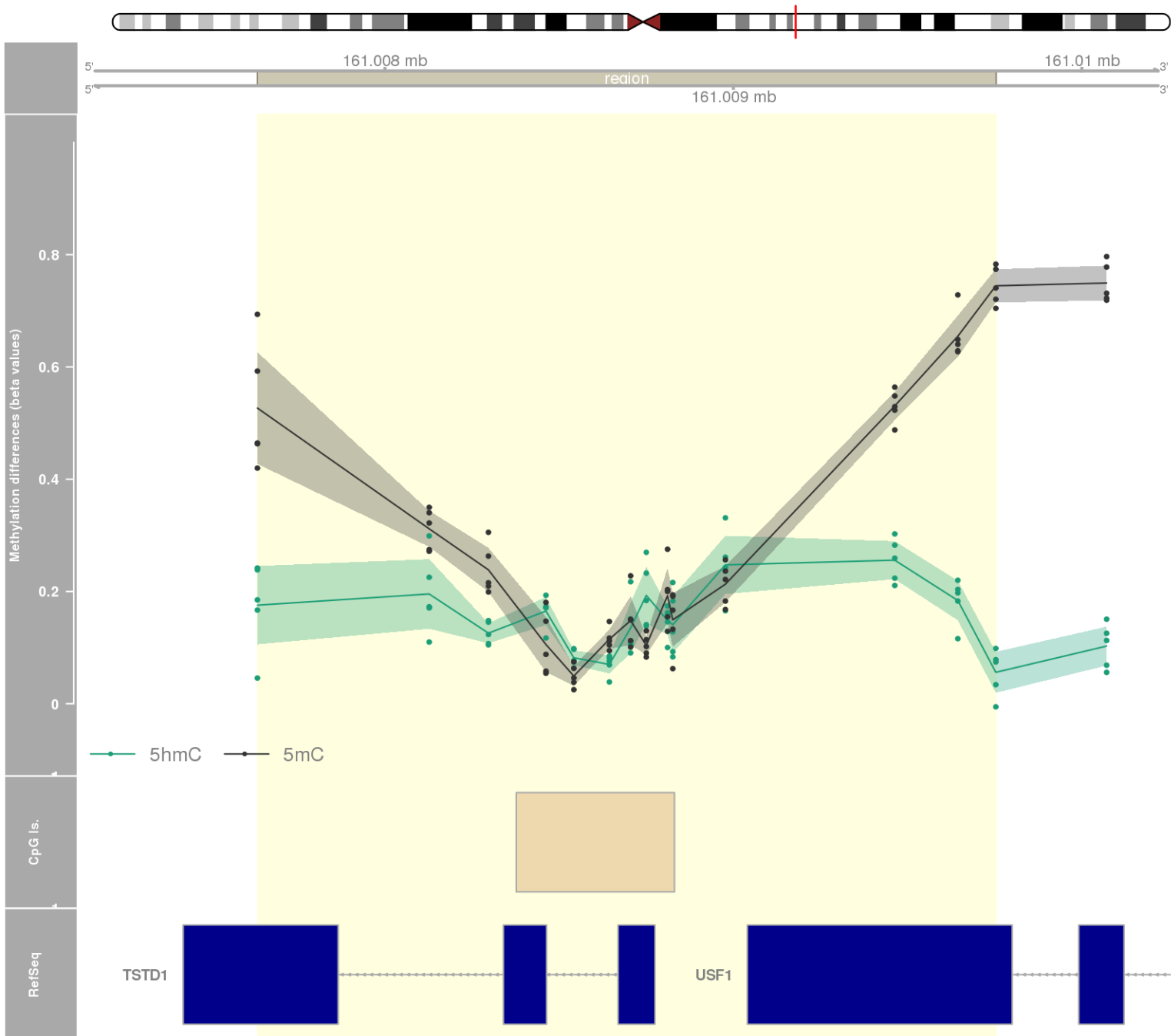
DMR 121 // chr19:46142212-46142930 // 718 pb. (13 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.009 // fwerArea: 1
- genes: MIR330 / EML2 -



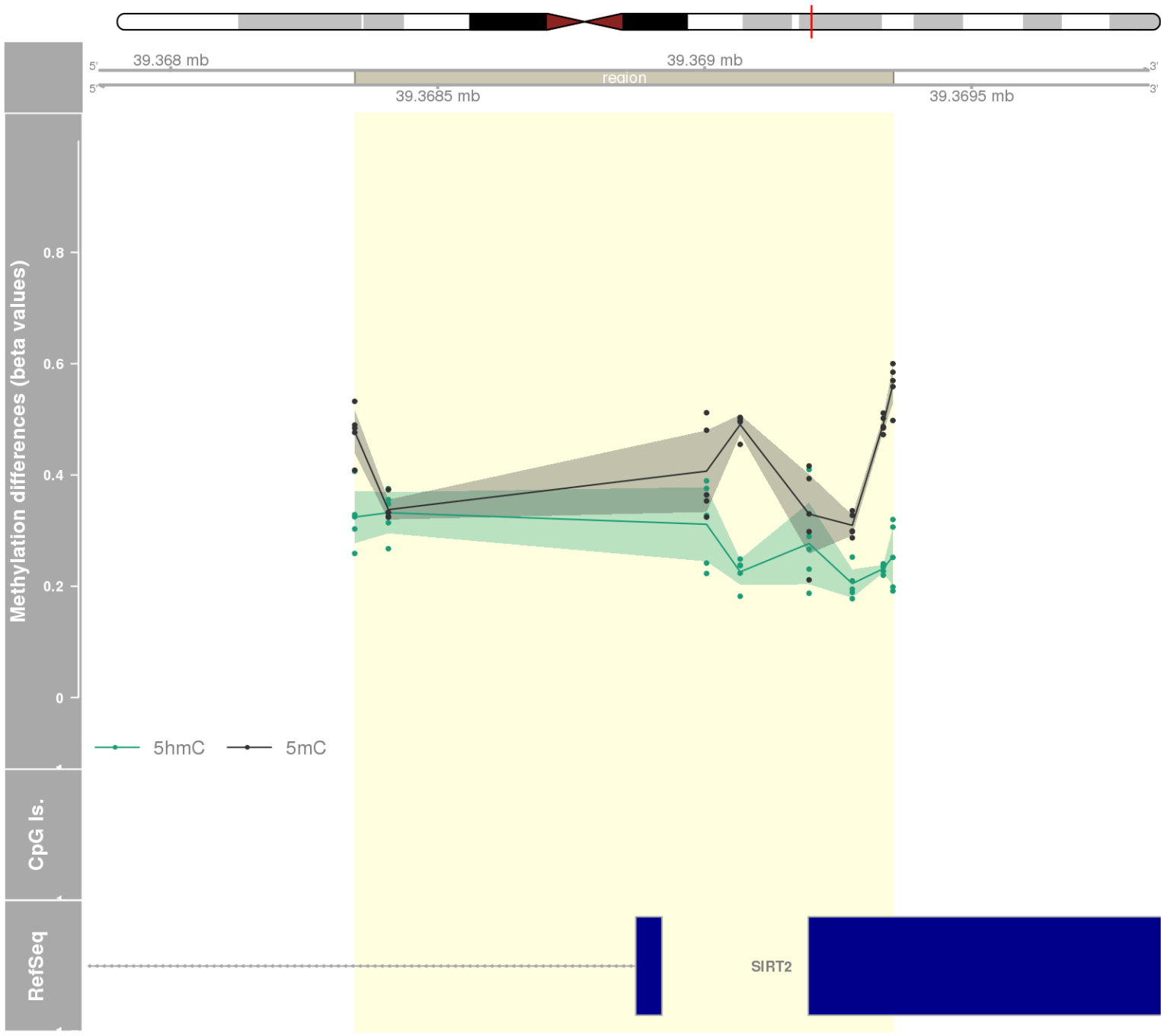
DMR 122 // chr3:50303757-50304463 // 706 pb. (10 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.009 // fwerArea: 1
- genes: SEMA3B-AS1 -



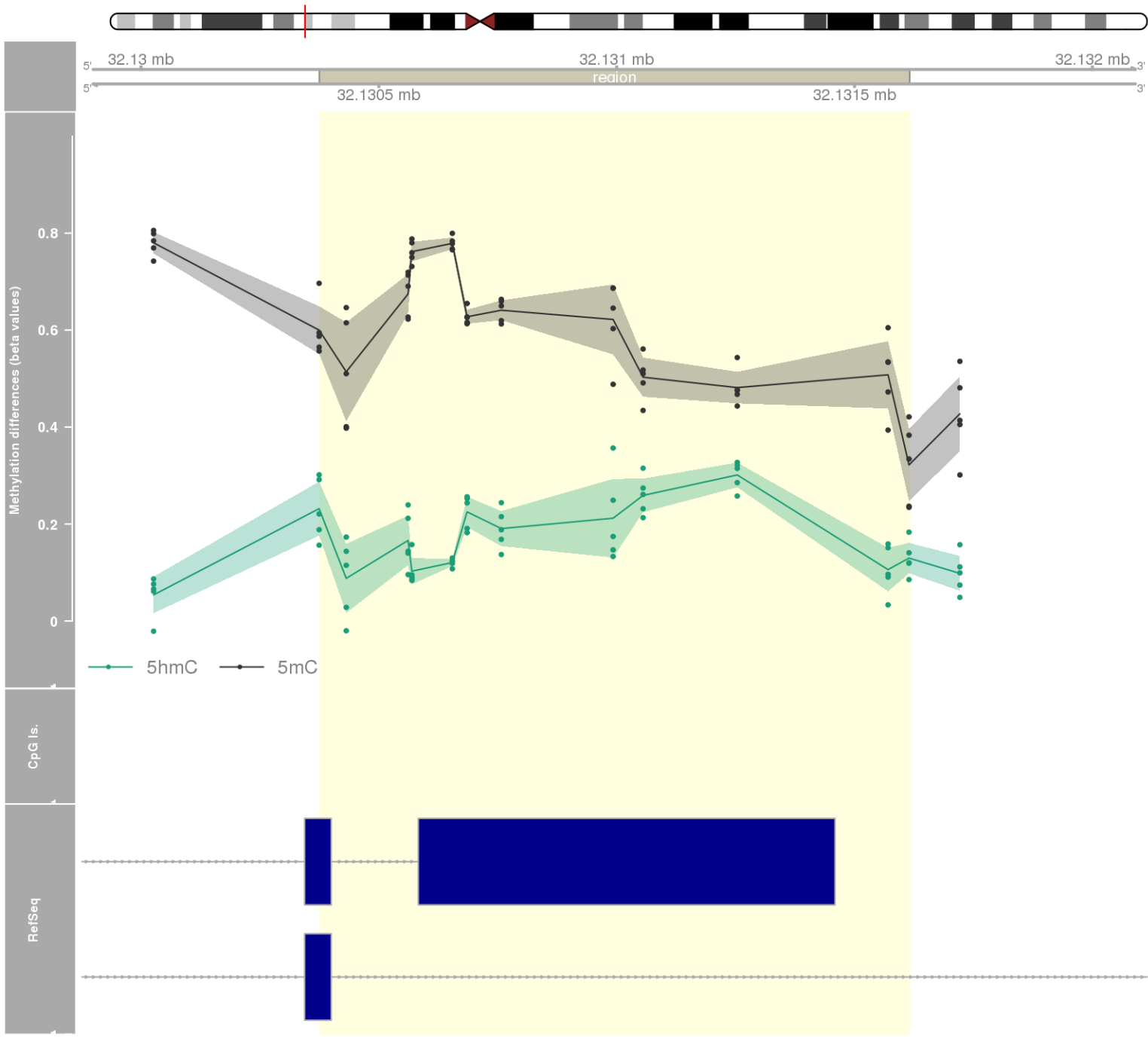
DMR 123 // chr1:161007634-161009753 // 2119 pb. (14 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.009 // fwerArea: 1
- genes: TSTD1 / USF1 -



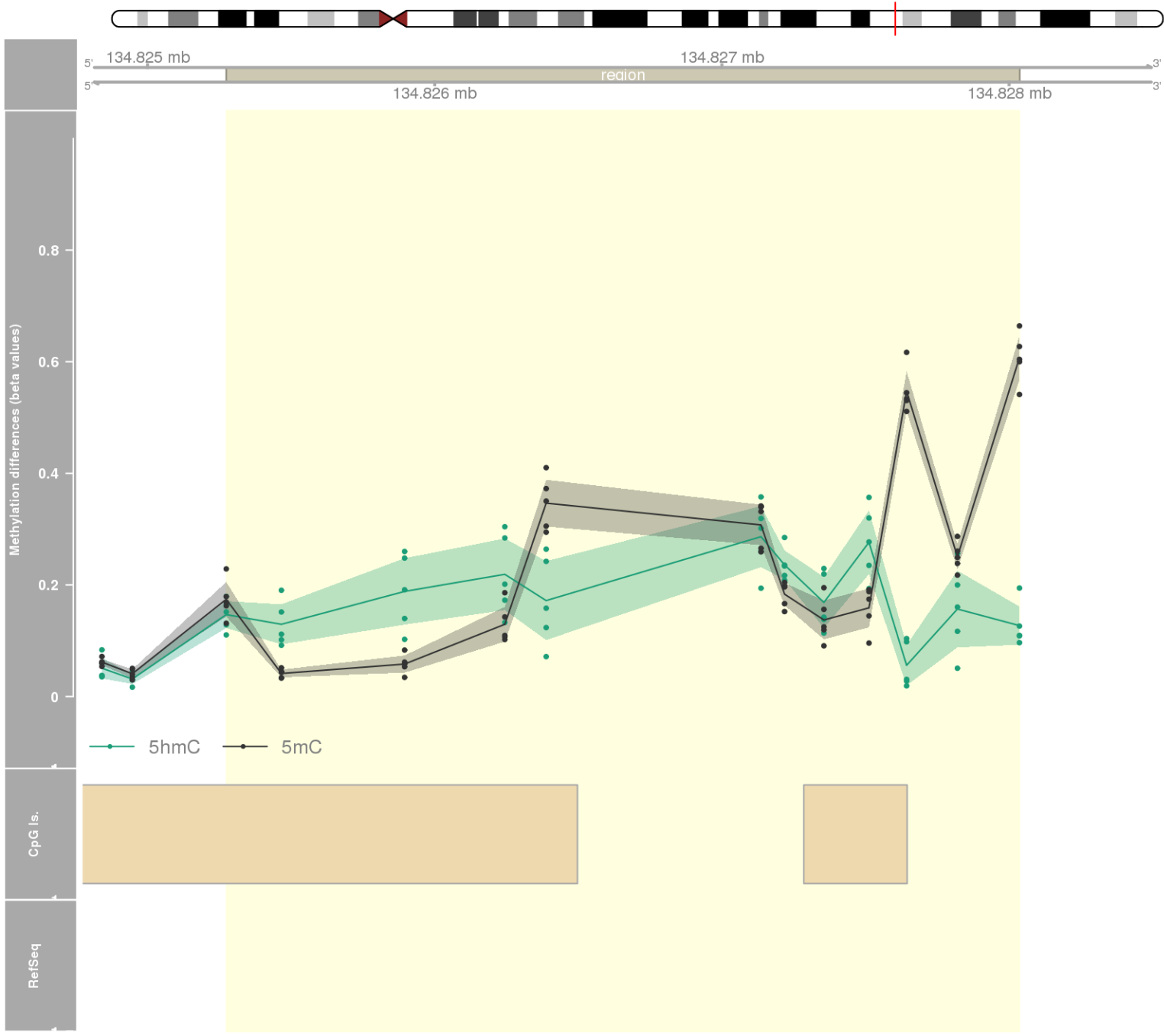
DMR 124 // chr19:39368345-39369353 // 1008 pb. (8 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.009 // fwerArea: 1
- genes: RINL / SIRT2 -



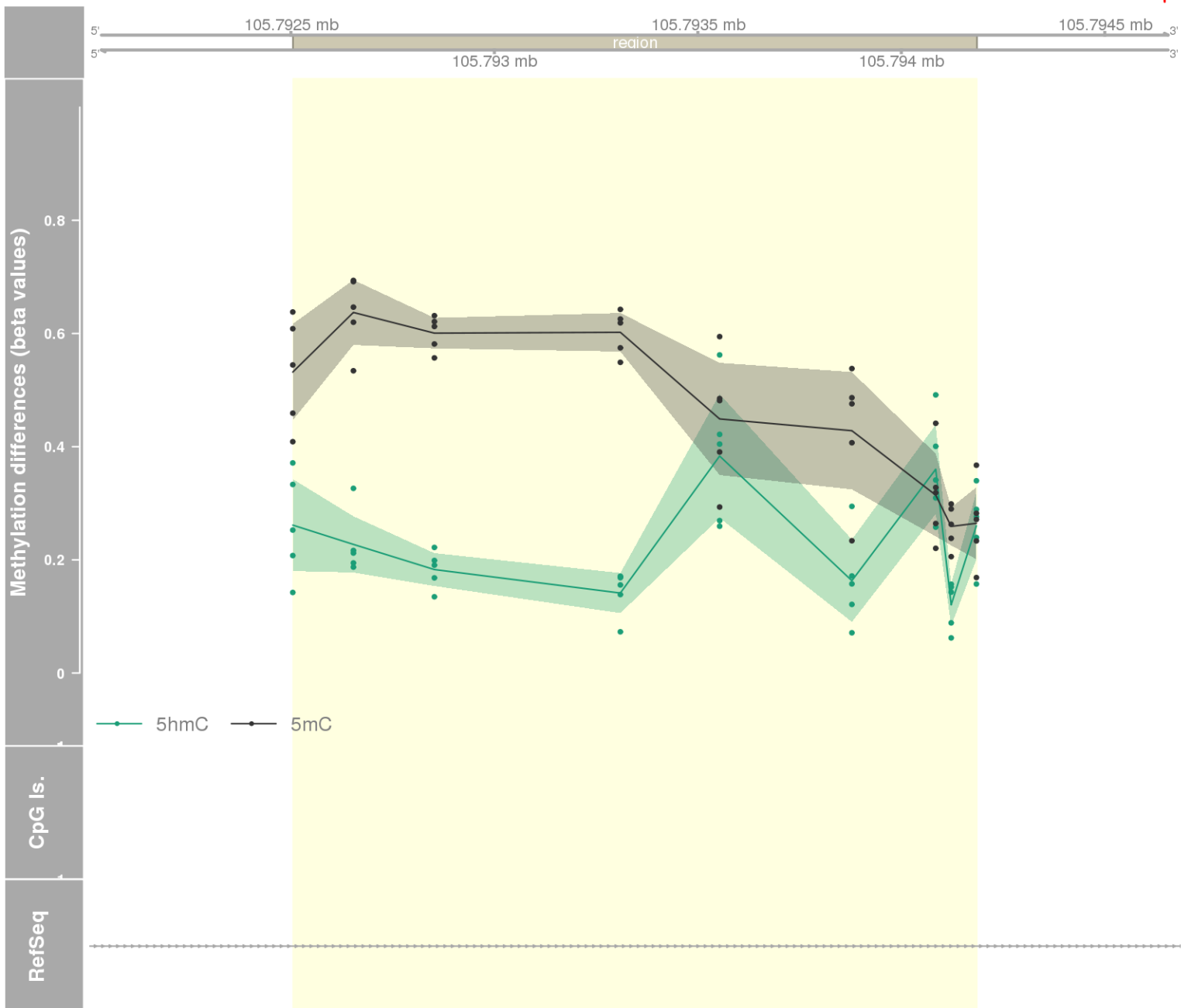
DMR 125 // chr6:32130375-32131615 // 1240 pb. (12 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.009 // fwerArea: 1
- genes: PPT2 / PPT2-EGFL8 -



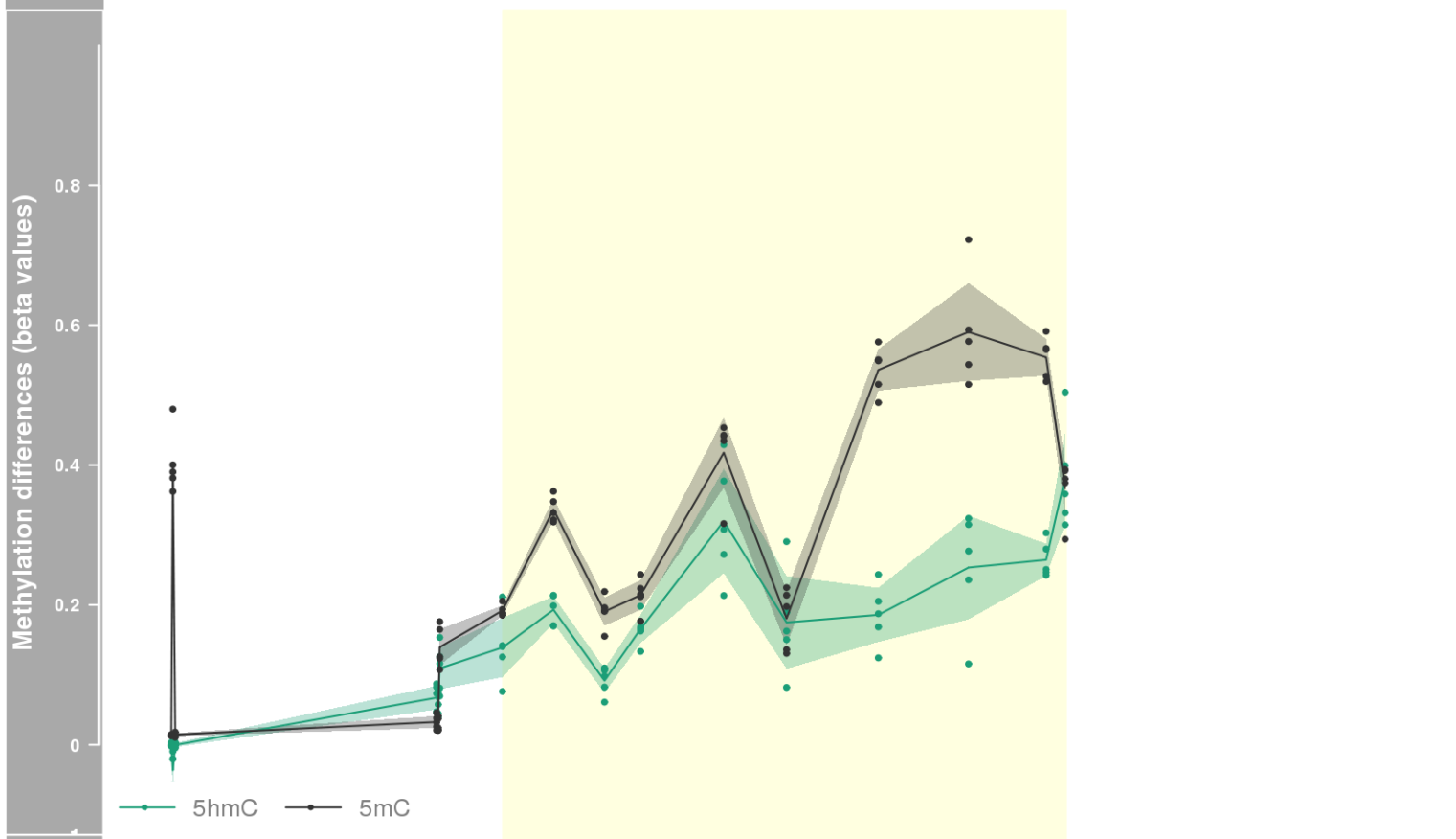
DMR 126 // chr5:134825274-134828036 // 2762 pb. (12 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.009 // fwerArea: 1



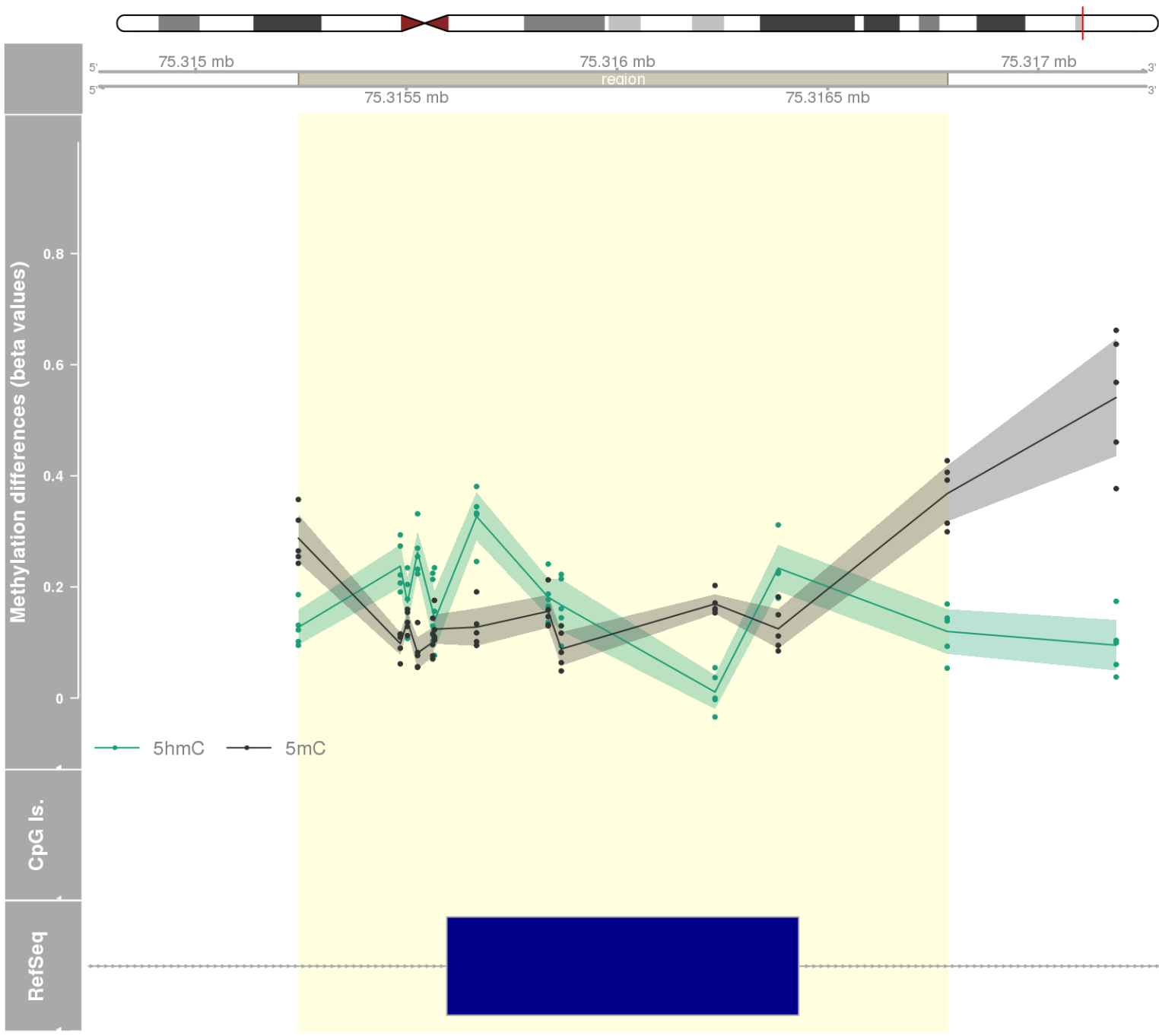
DMR 127 // chr14:105792505-105794185 // 1680 pb. (9 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.009 // fwerArea: 1
- genes: PACS2 -



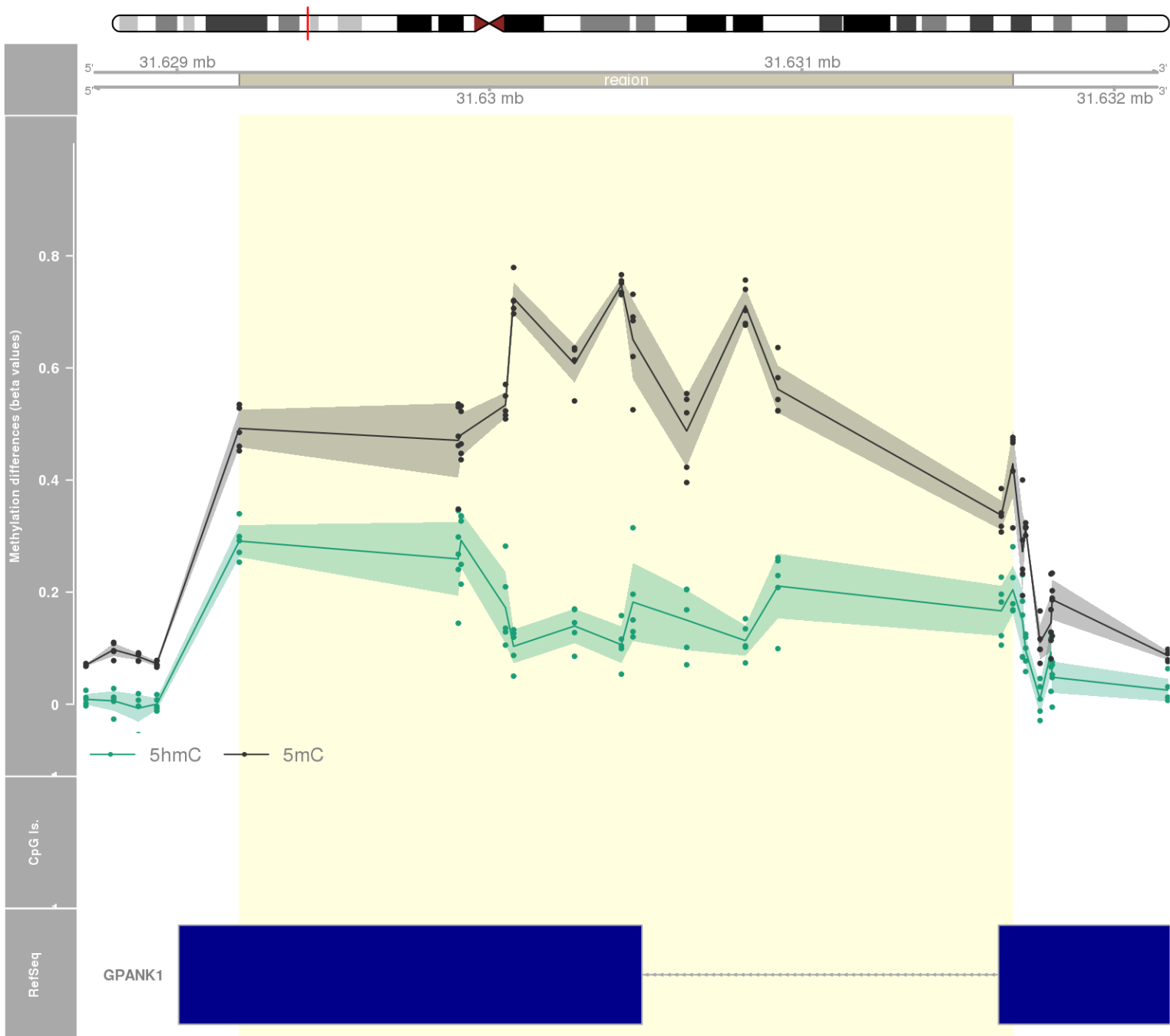
DMR 128 // chr11:2907754-2908471 // 717 pb. (10 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.009 // fwerArea: 1



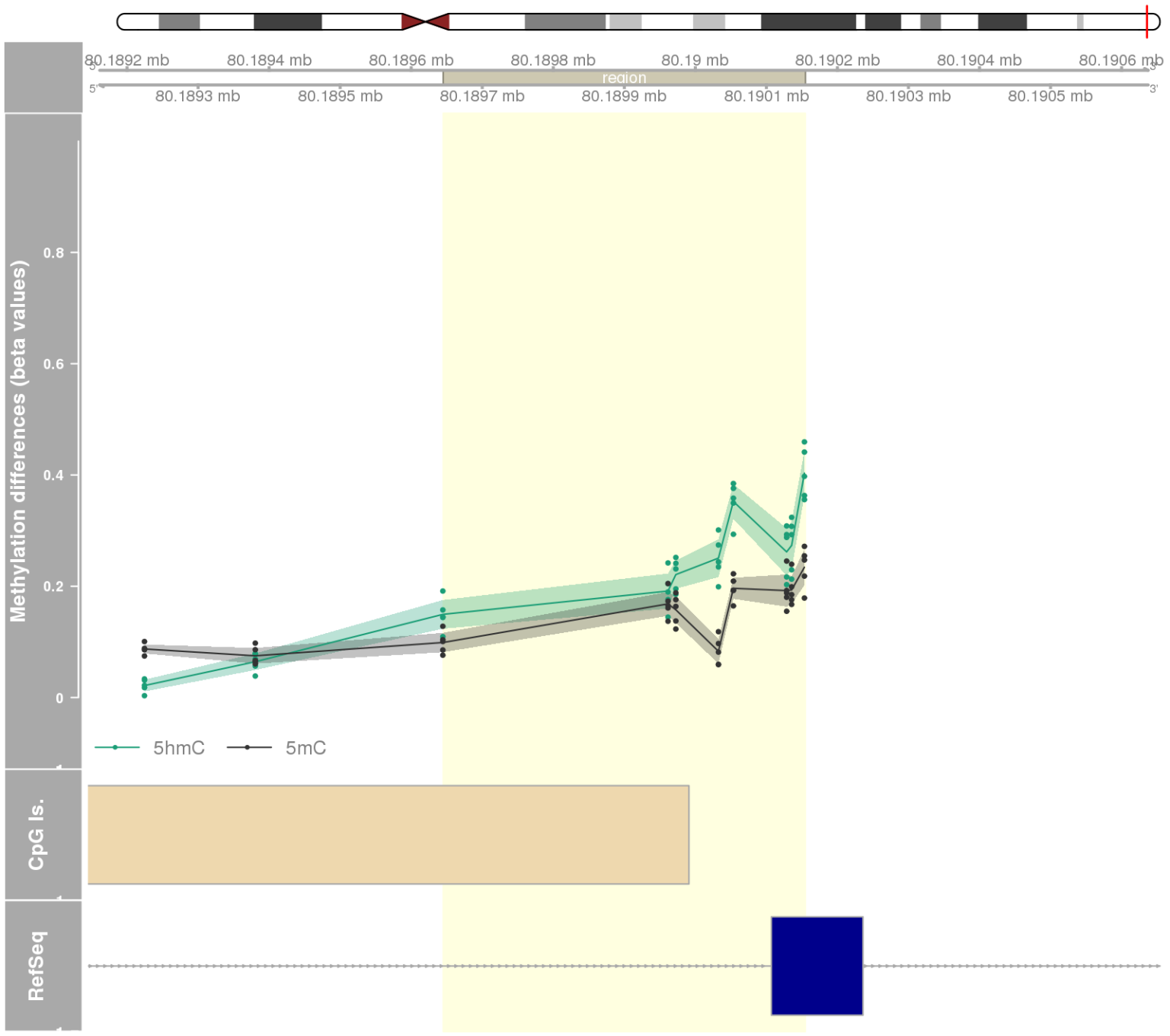
DMR 129 // chr17:75315244-75316784 // 1540 pb. (12 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.009 // fwerArea: 1
- genes: SEPT9 -



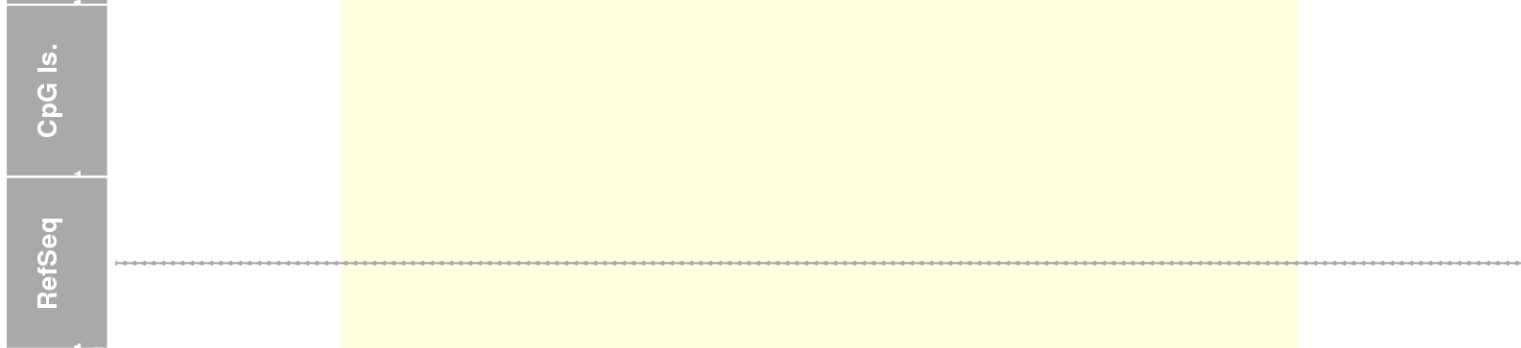
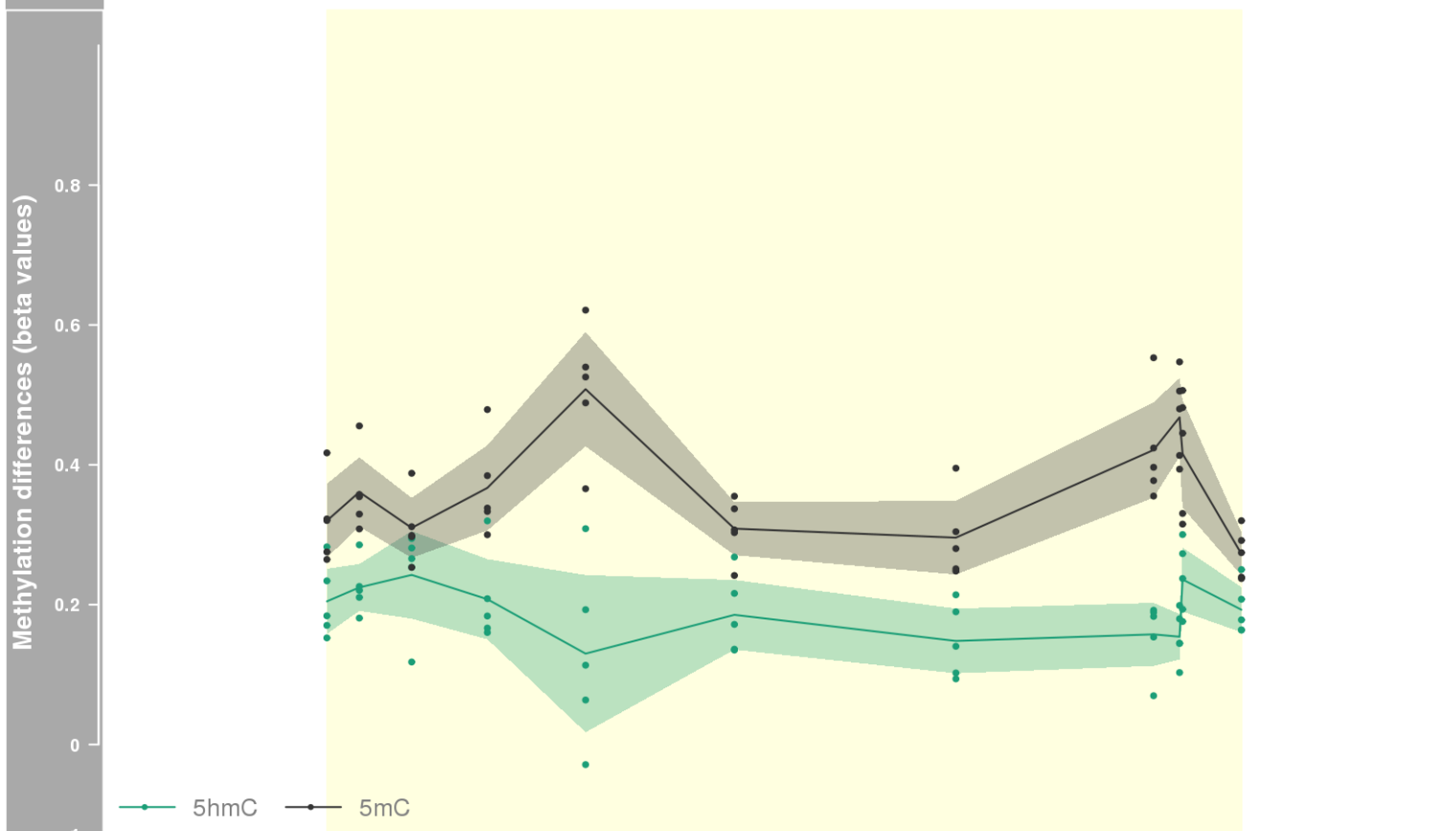
DMR 130 // chr6:31629199-31631675 // 2476 pb. (13 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.009 // fwerArea: 1
- genes: GPANK1 -



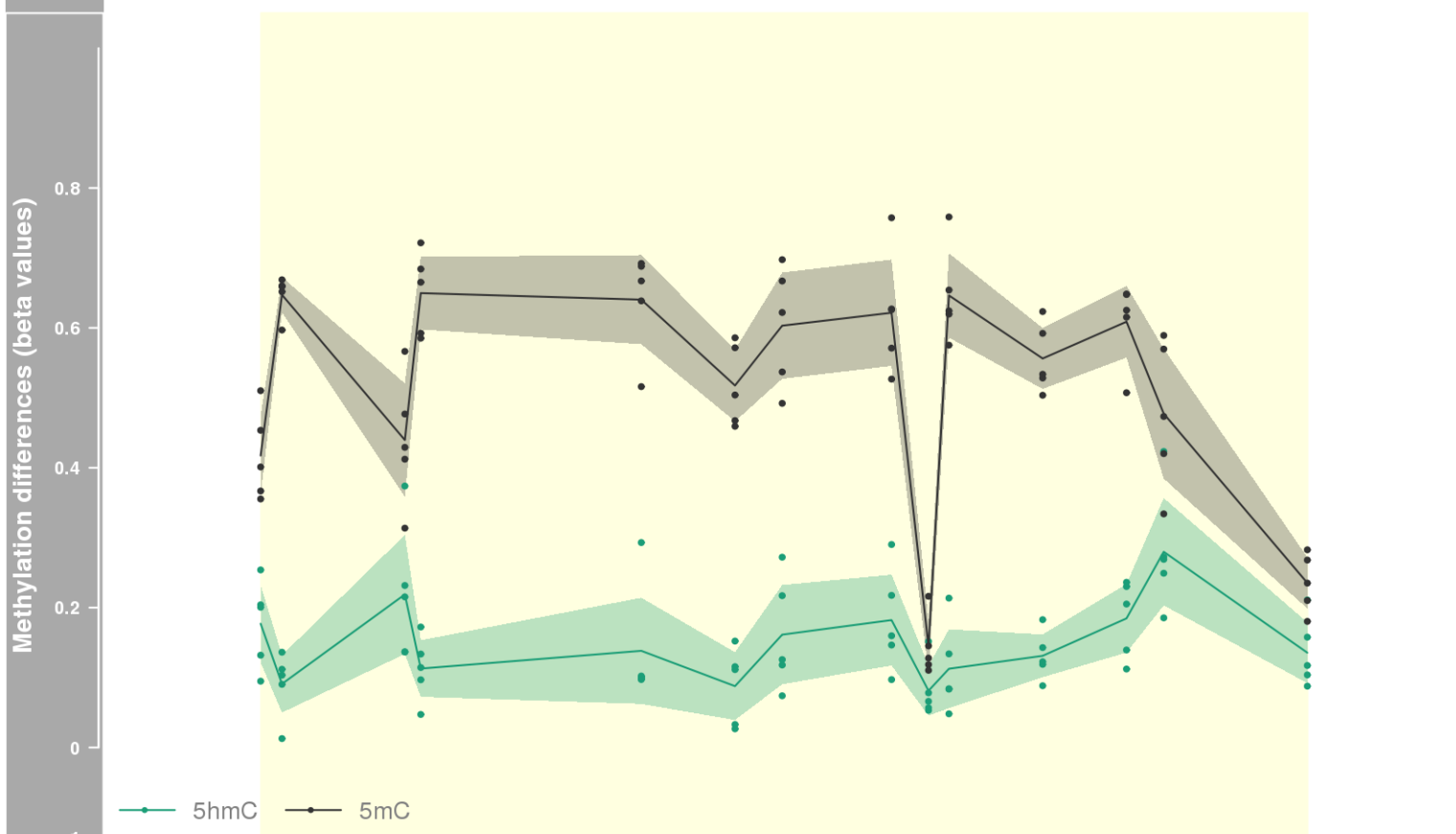
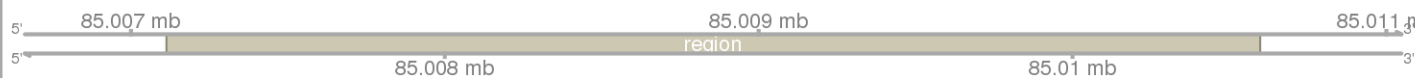
DMR 131 // chr17:80189645-80190154 // 509 pb. (8 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.009 // fwerArea: 1
- genes: SLC16A3 -



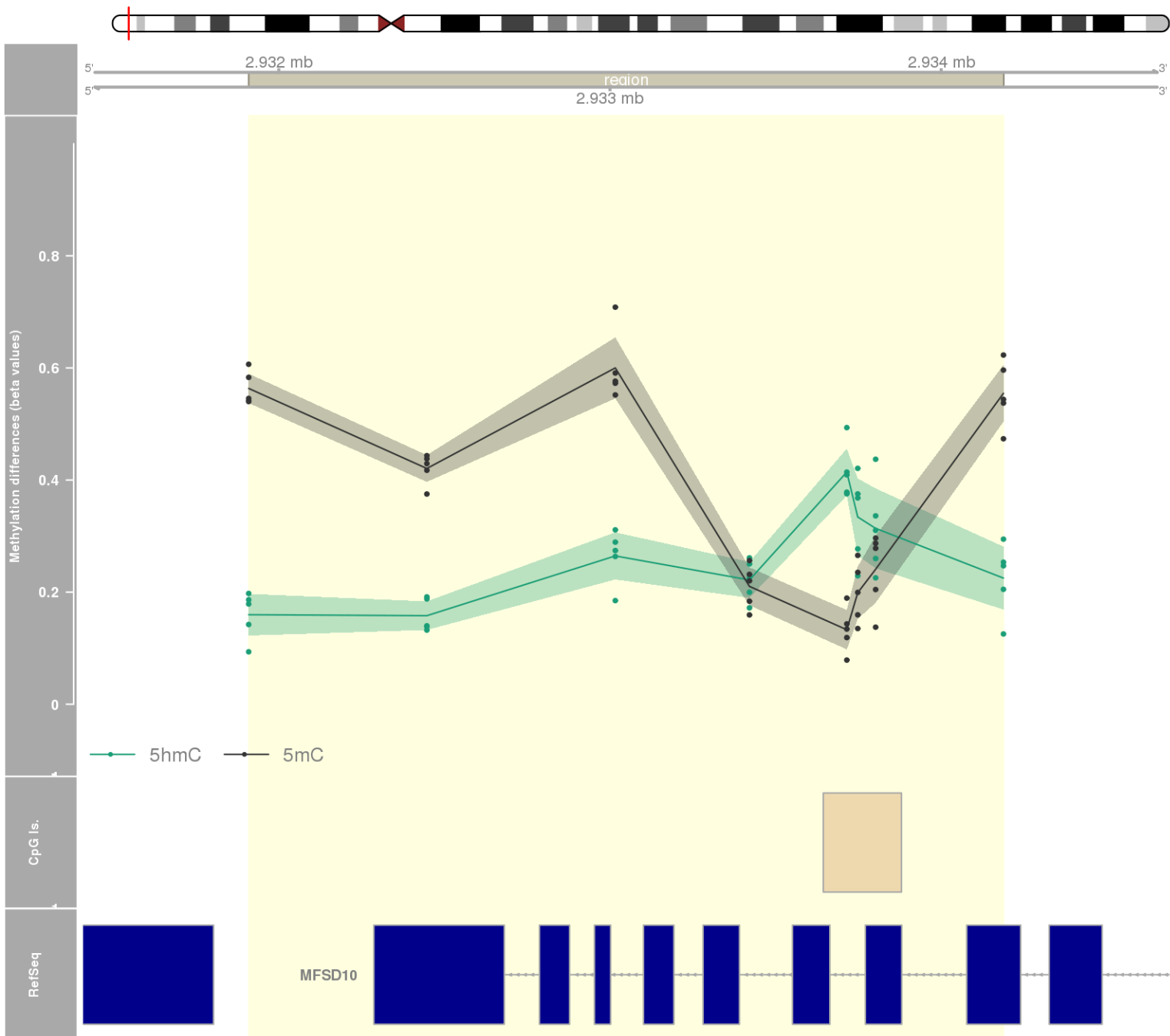
DMR 132 // chr11:1326508-1328622 // 2114 pb. (11 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.009 // fwerArea: 1
- genes: TOLLIP -



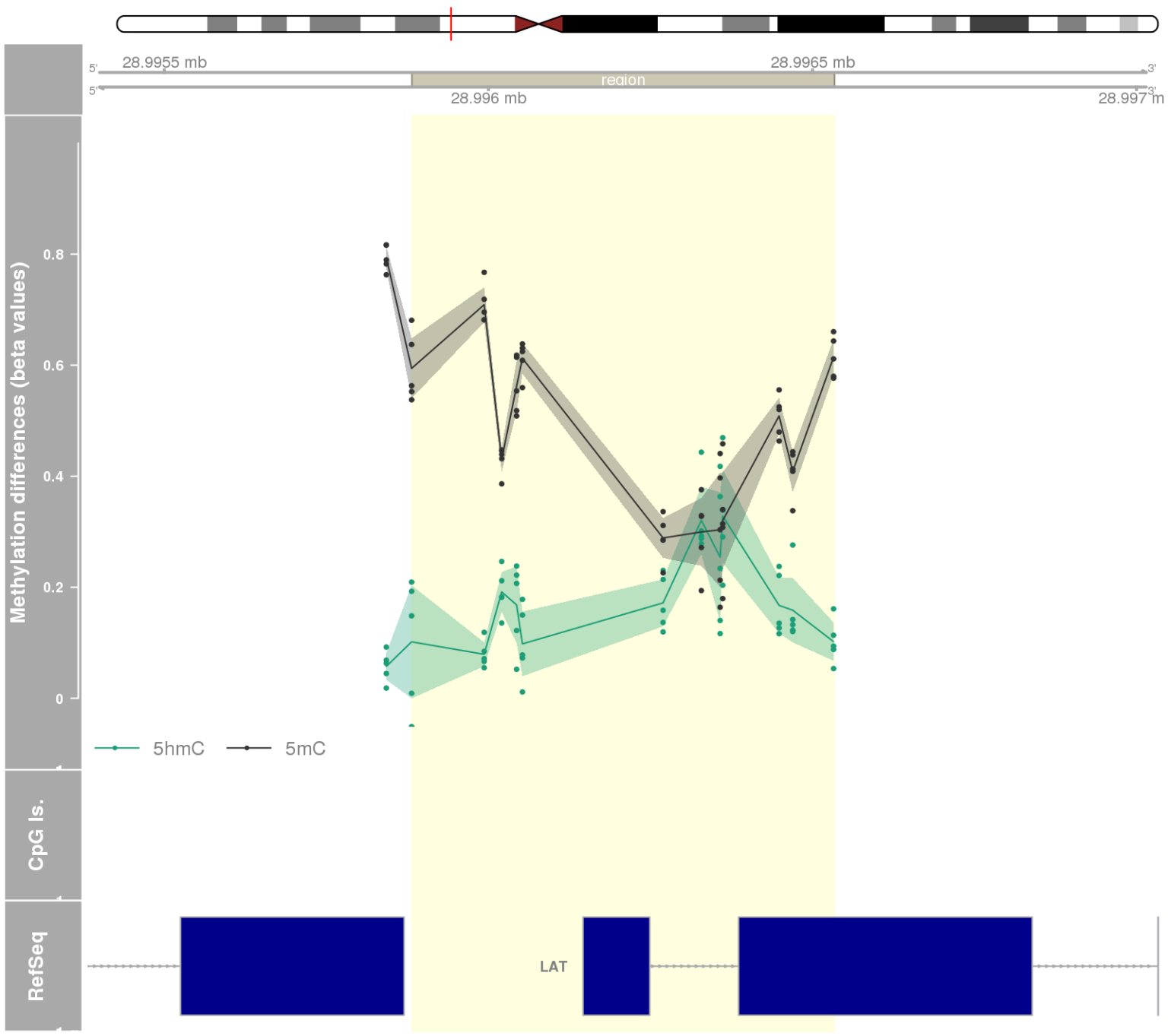
DMR 133 // chr16:85007113-85010597 // 3484 pb. (14 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.01 // fwerArea: 1
- genes: ZDHHC7 -



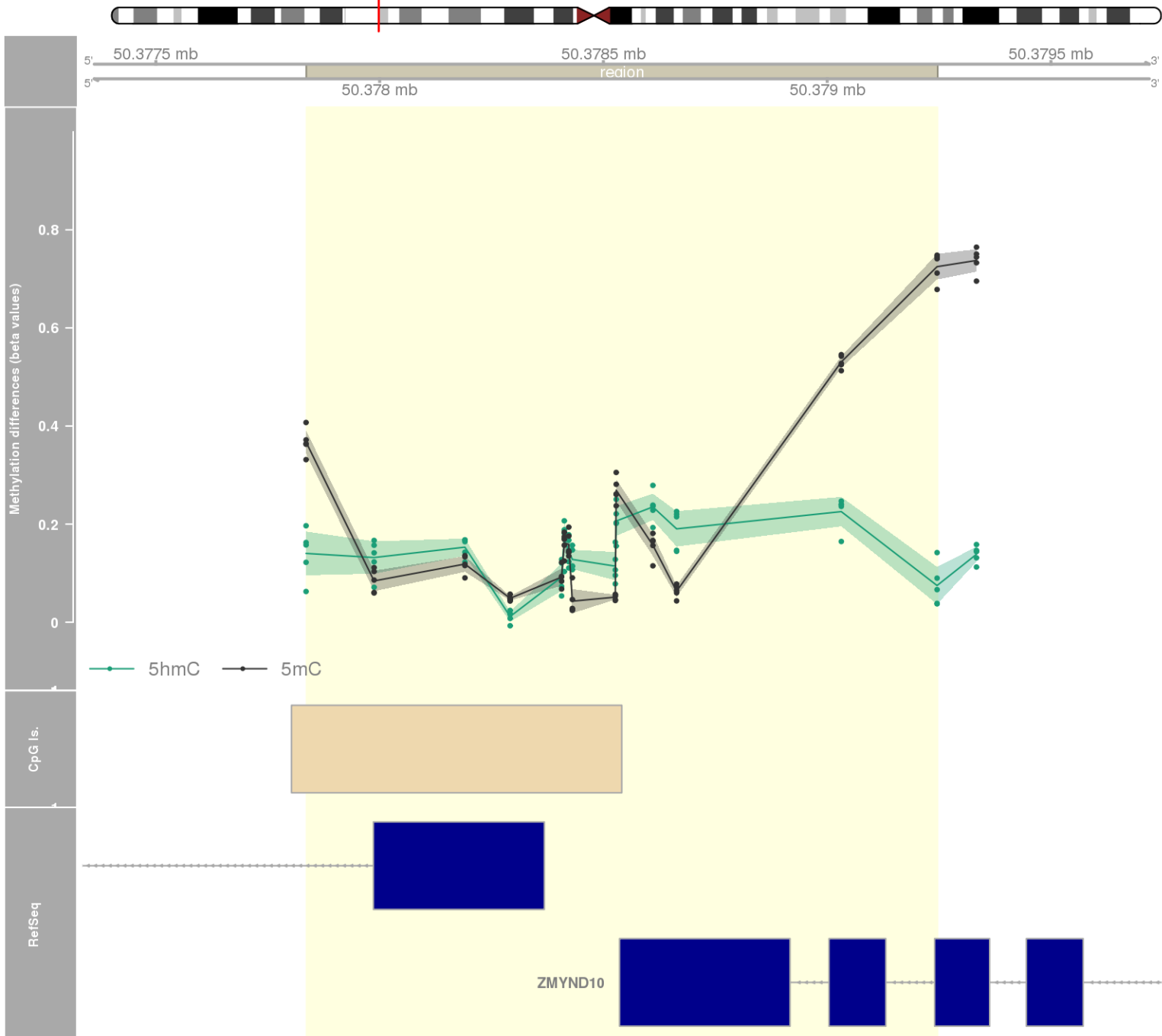
DMR 134 // chr4:2931909-2934188 // 2279 pb. (8 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.01 // fwerArea: 1
- genes: MFSD10 -



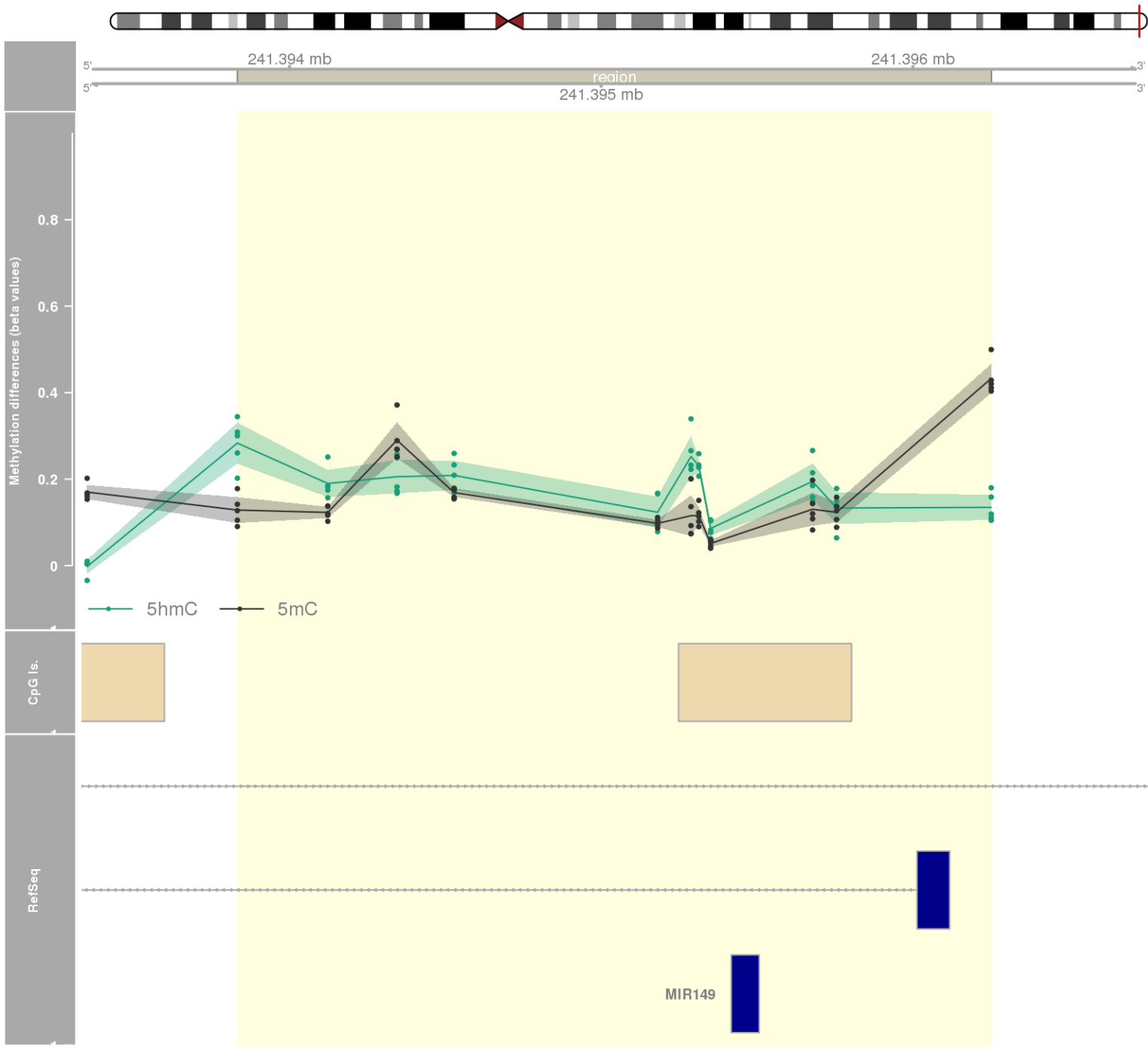
DMR 135 // chr16:28995882-28996533 // 651 pb. (12 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.01 // fwerArea: 1
- genes: LAT -



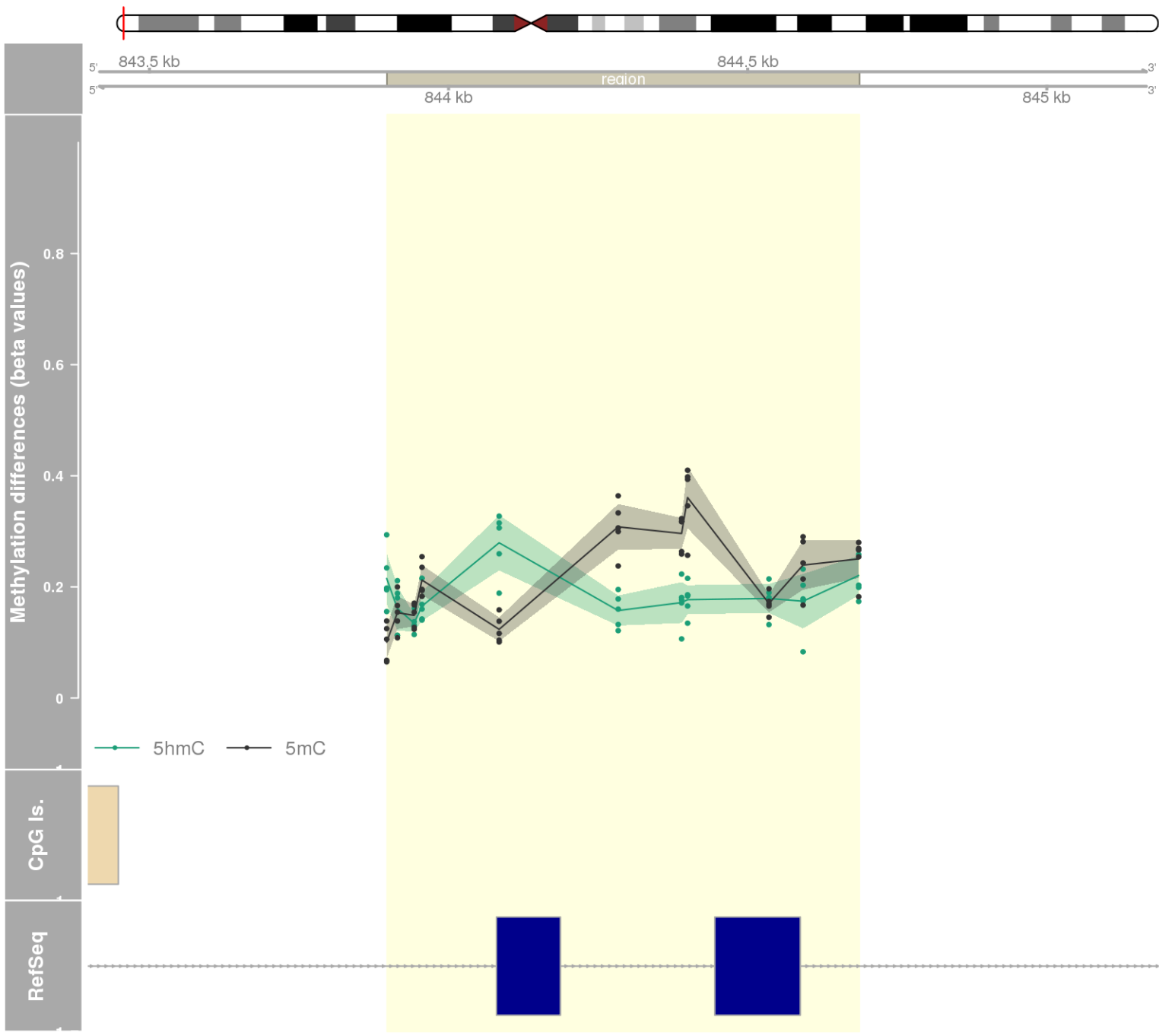
DMR 136 // chr3:50377836-50379246 // 1410 pb. (14 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.01 // fwerArea: 1
- genes: RASSF1 / ZMYND10 / ZMYND10-AS1 -



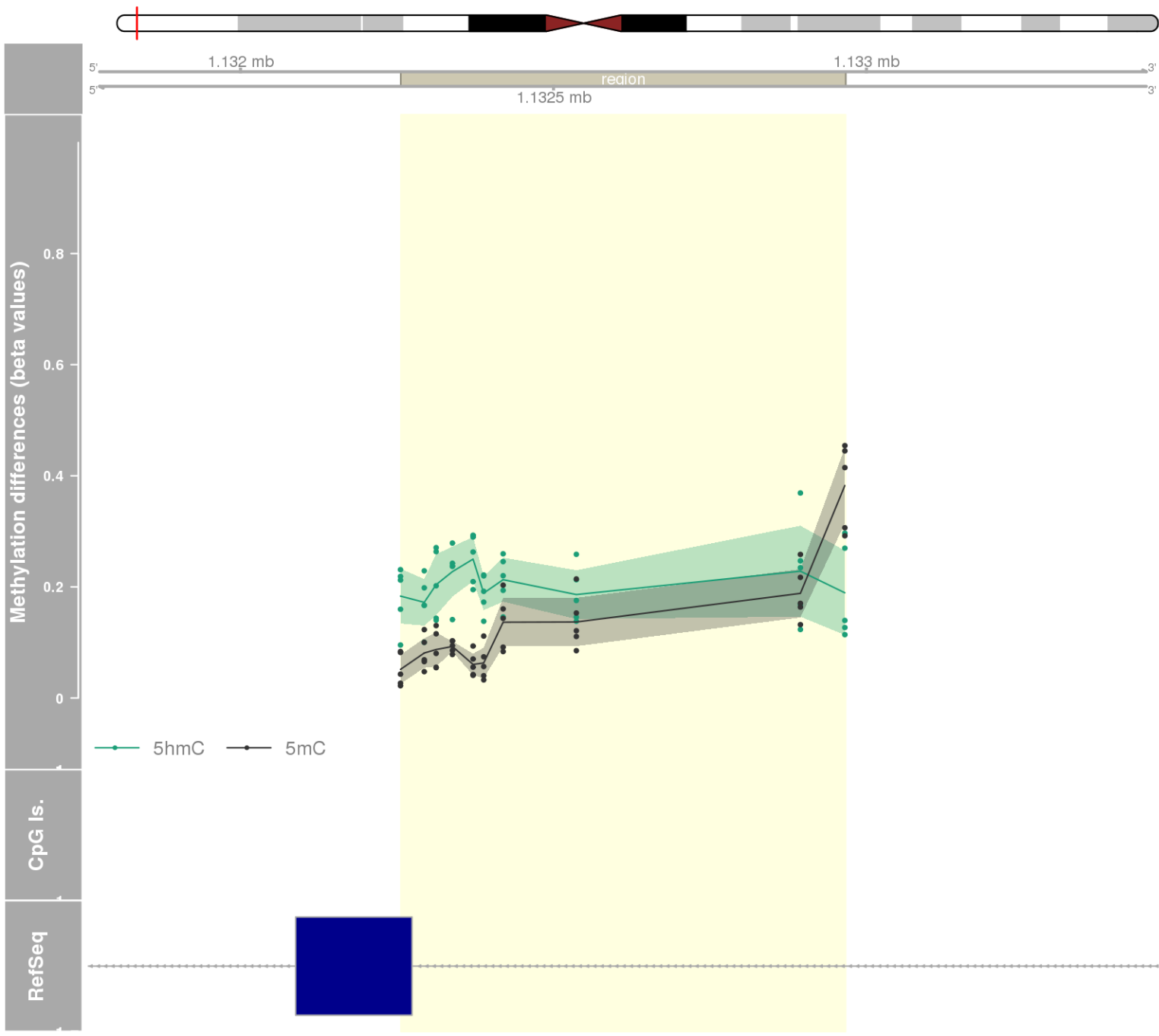
DMR 137 // chr2:241393833-241396251 // 2418 pb. (11 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.01 // fwerArea: 1
- genes: MIR149 / GPC1 -



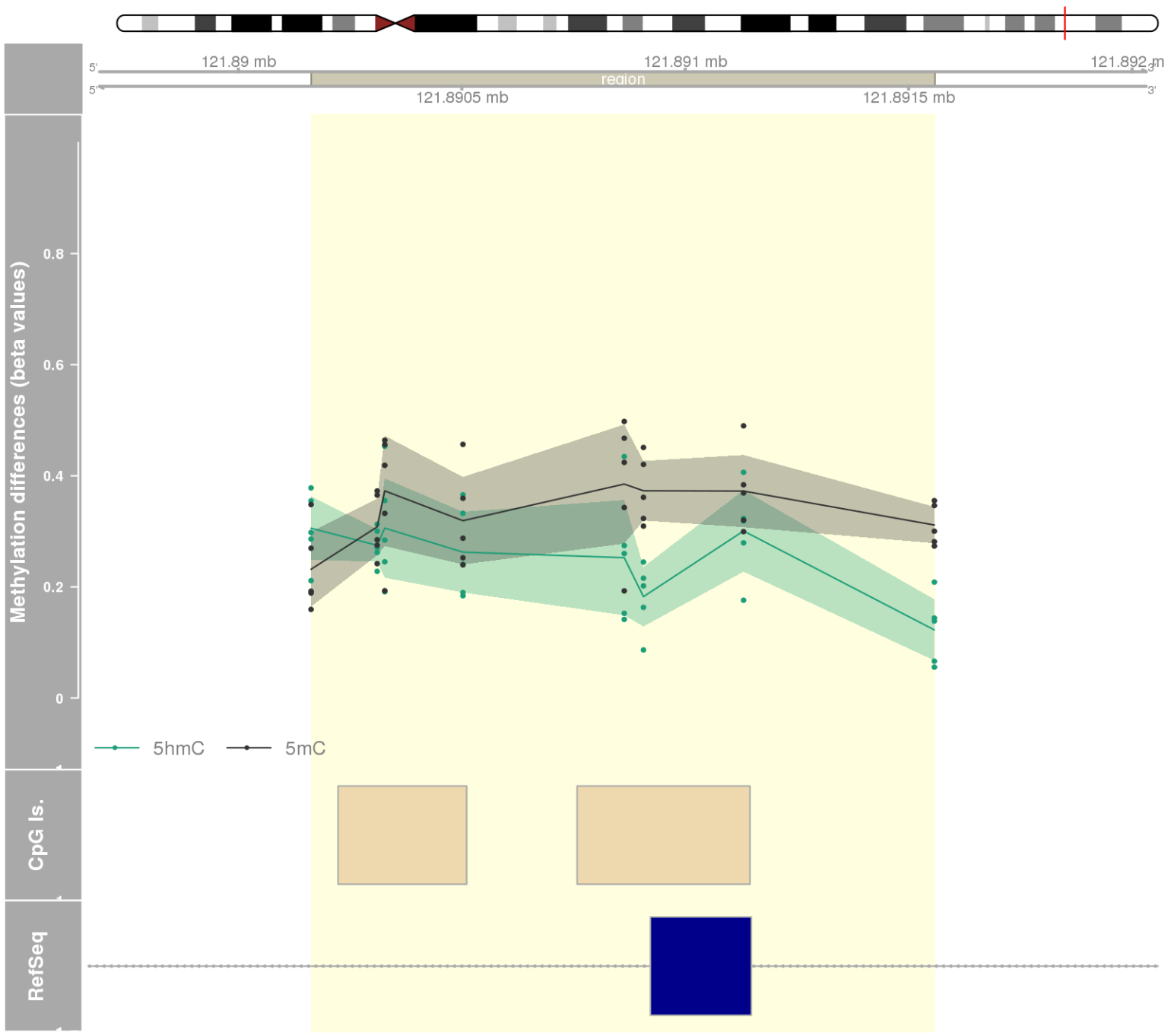
DMR 138 // chr11:843897-844686 // 789 pb. (11 probes) // pvalue: 0.003 // fwer: 0.36 // pvalueArea: 0.01 // fwerArea: 1
- genes: TSPAN4 -



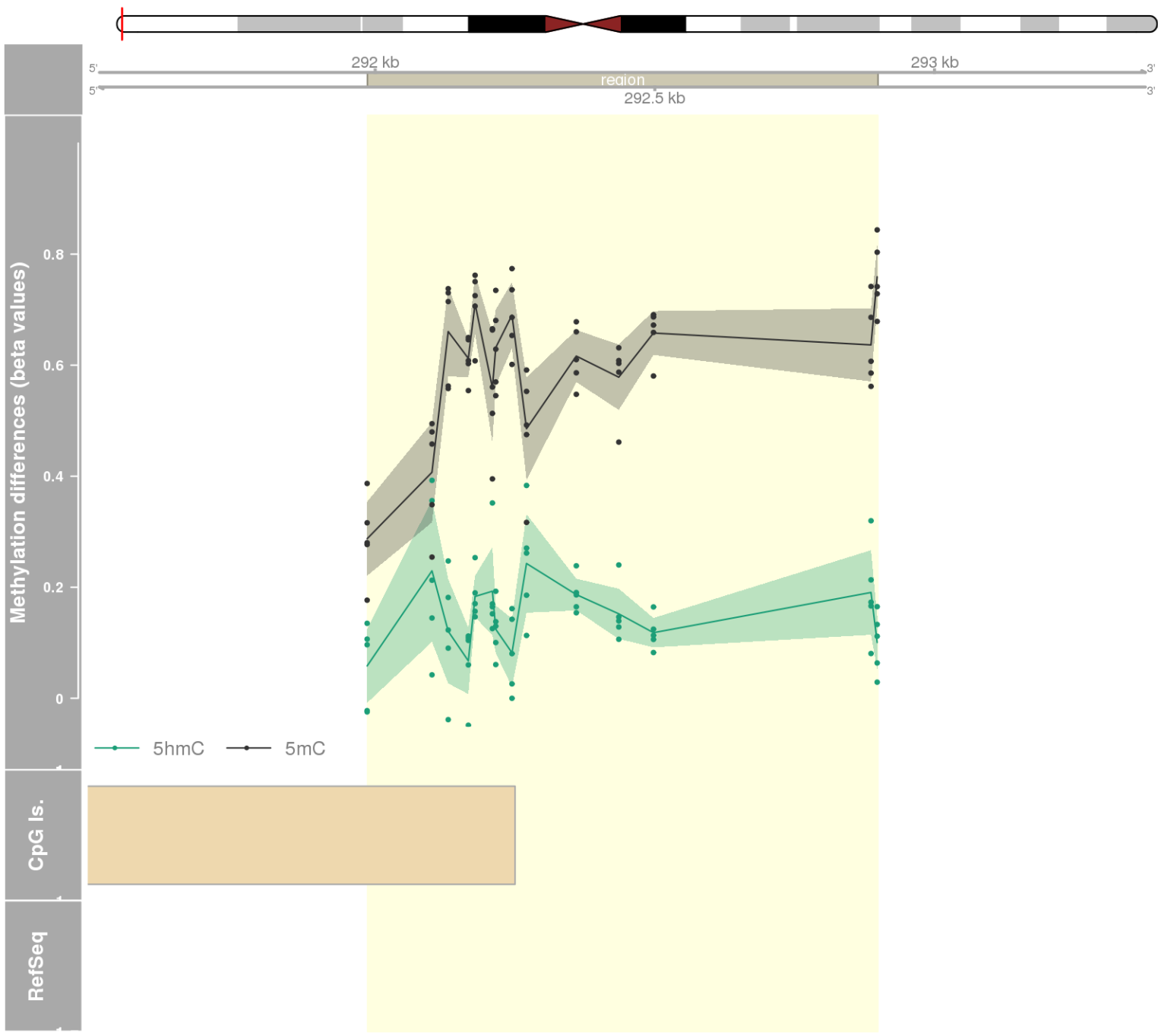
DMR 139 // chr19:1132256-1132966 // 710 pb. (10 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.01 // fwerArea: 1
- genes: SBNO2 -



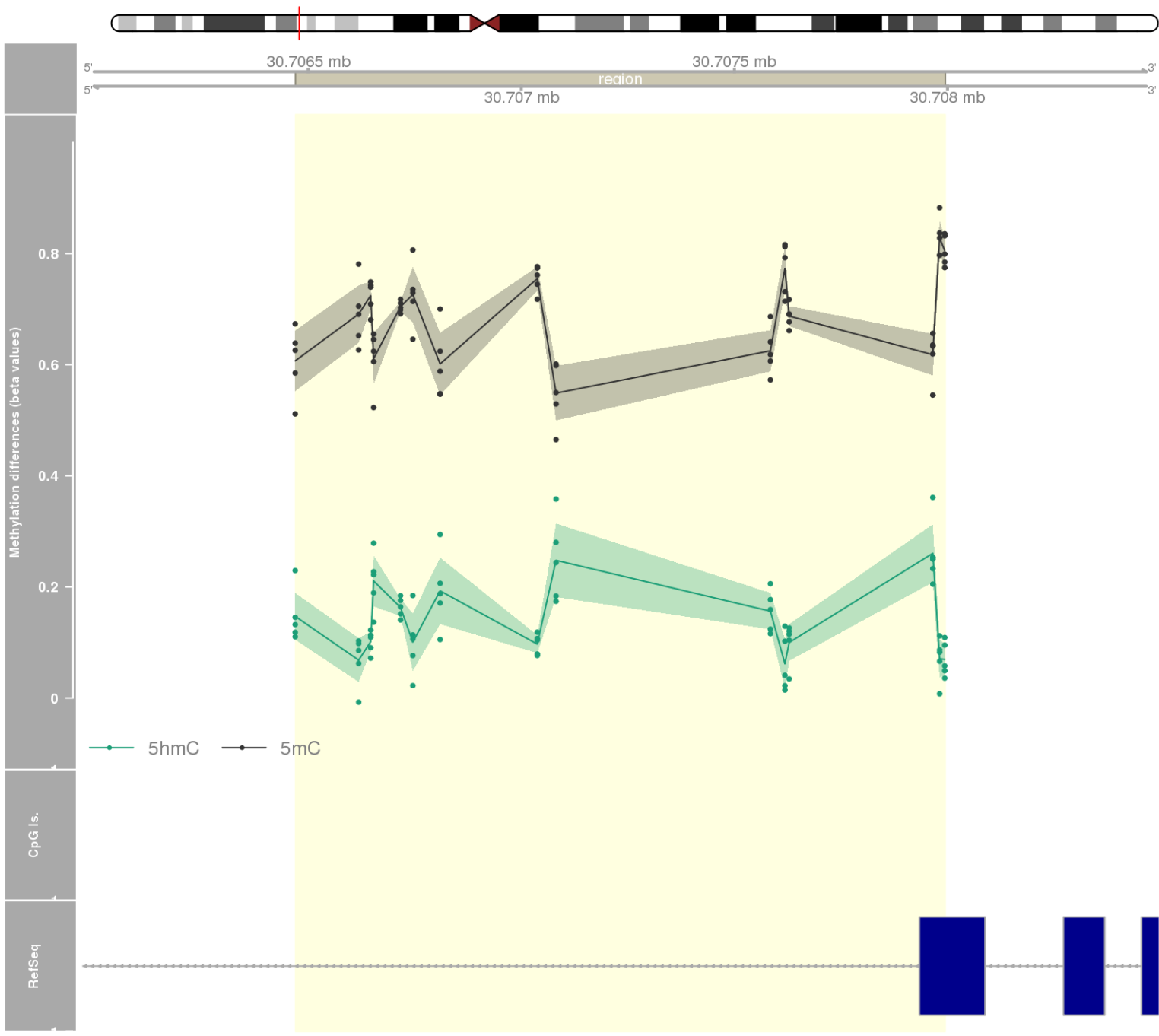
DMR 140 // chr12:121890163-121891558 // 1395 pb. (8 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.01 // fwerArea: 1
- genes: KDM2B -



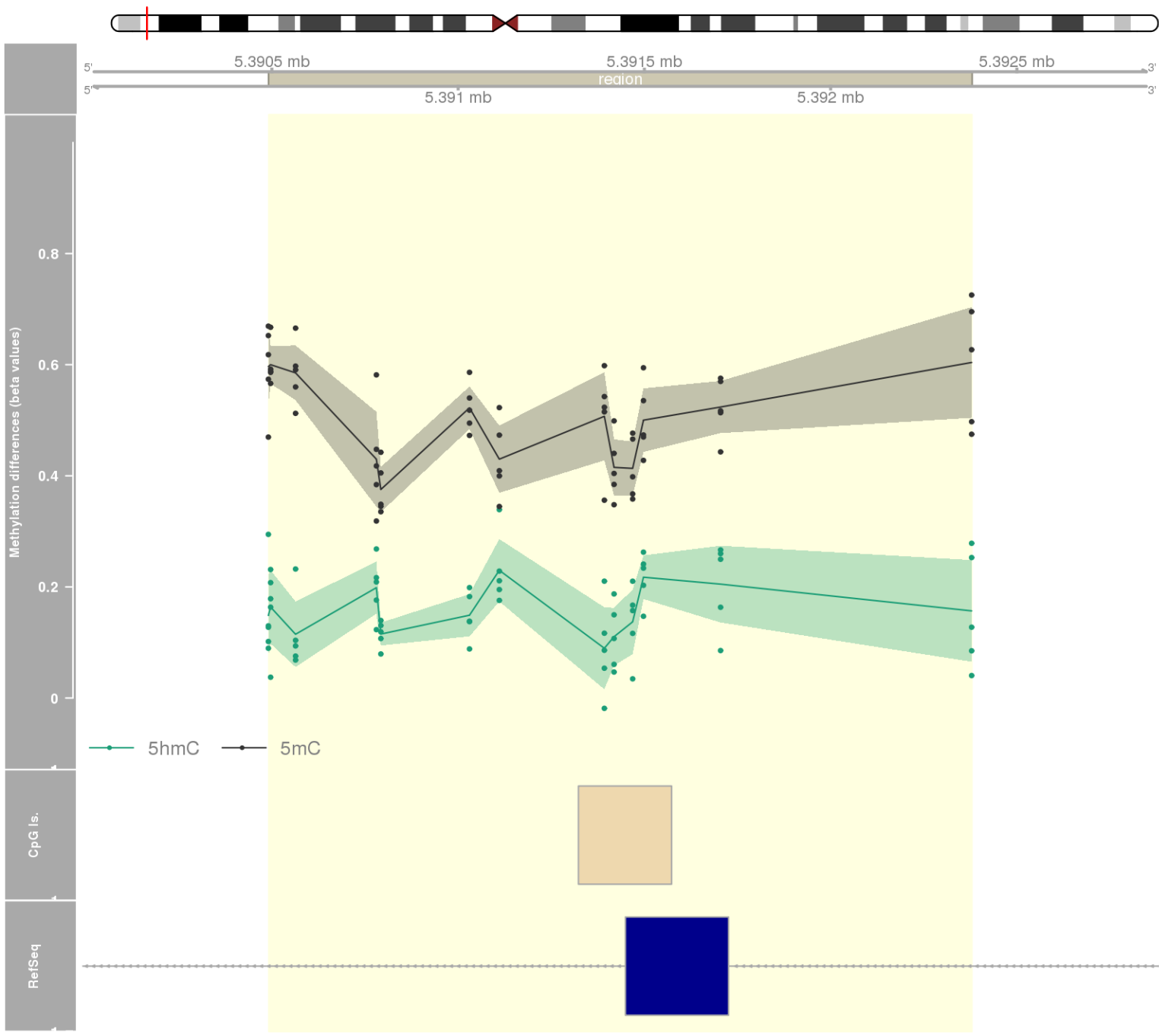
DMR 141 // chr19:291986-292898 // 912 pb. (14 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.01 // fwerArea: 1



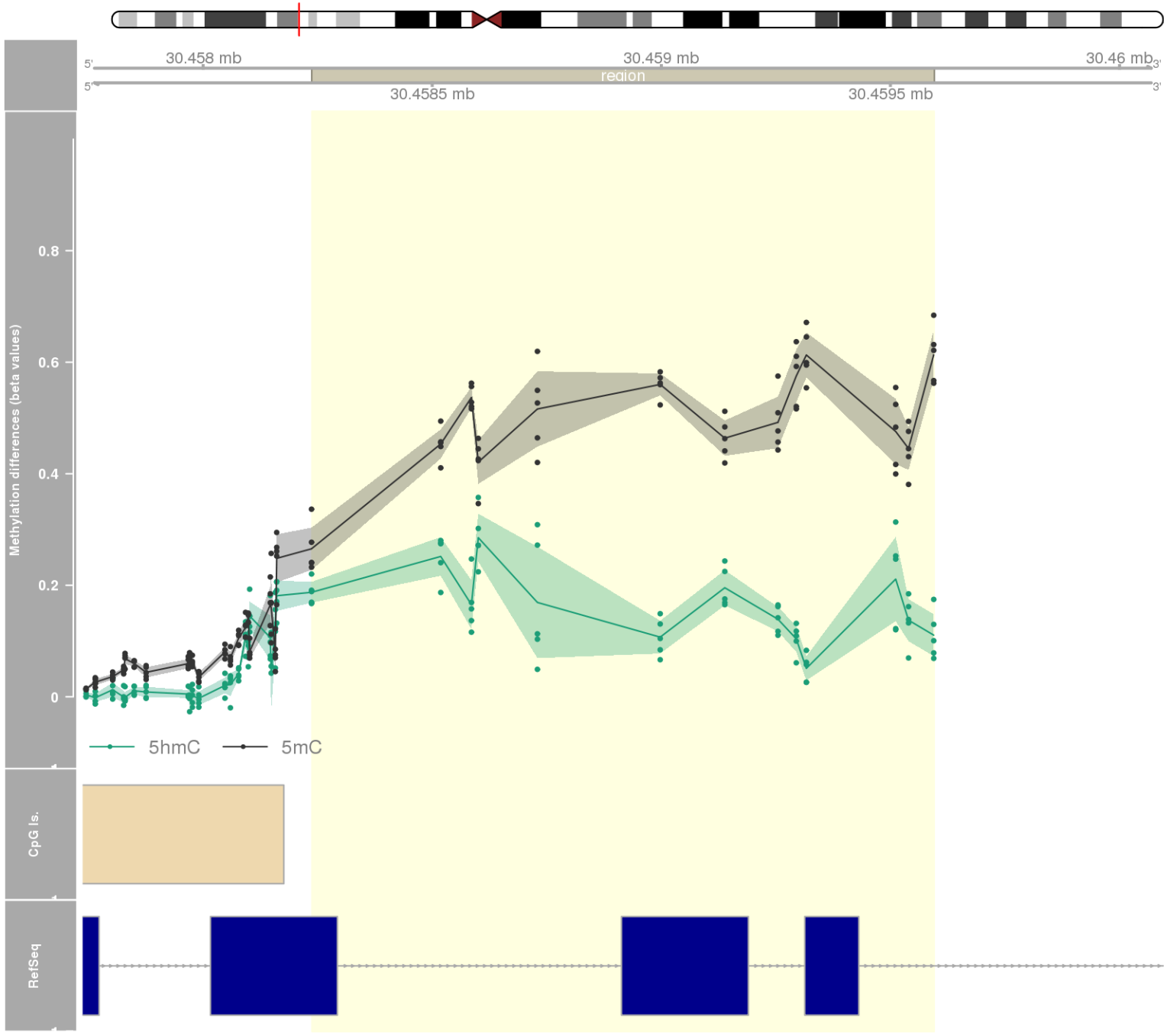
DMR 142 // chr6:30706470-30707994 // 1524 pb. (15 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.01 // fwerArea: 1
- genes: FLOT1 -



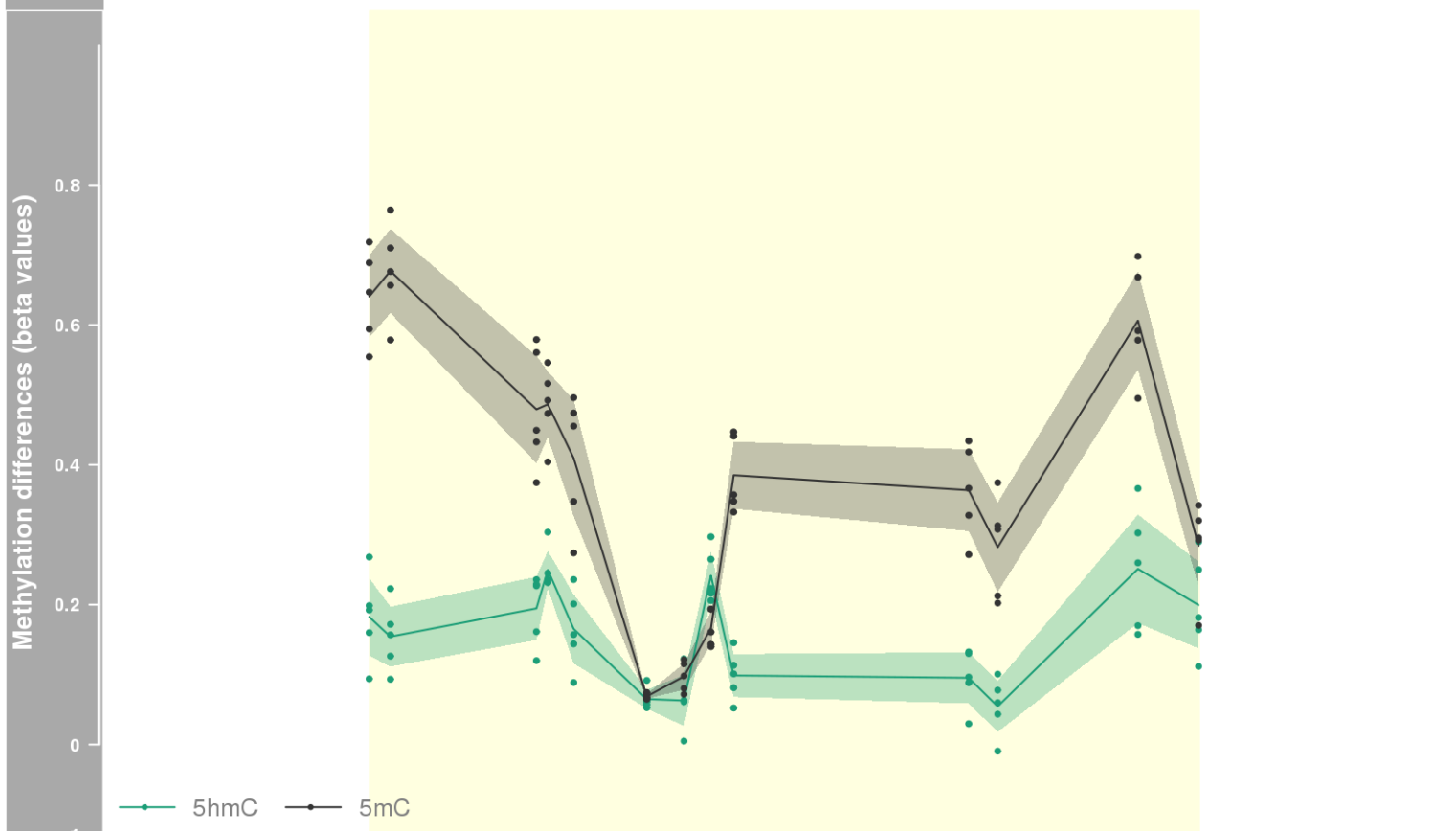
DMR 143 // chr7:5390491-5392379 // 1888 pb. (13 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.01 // fwerArea: 1
- genes: TNRC18 -



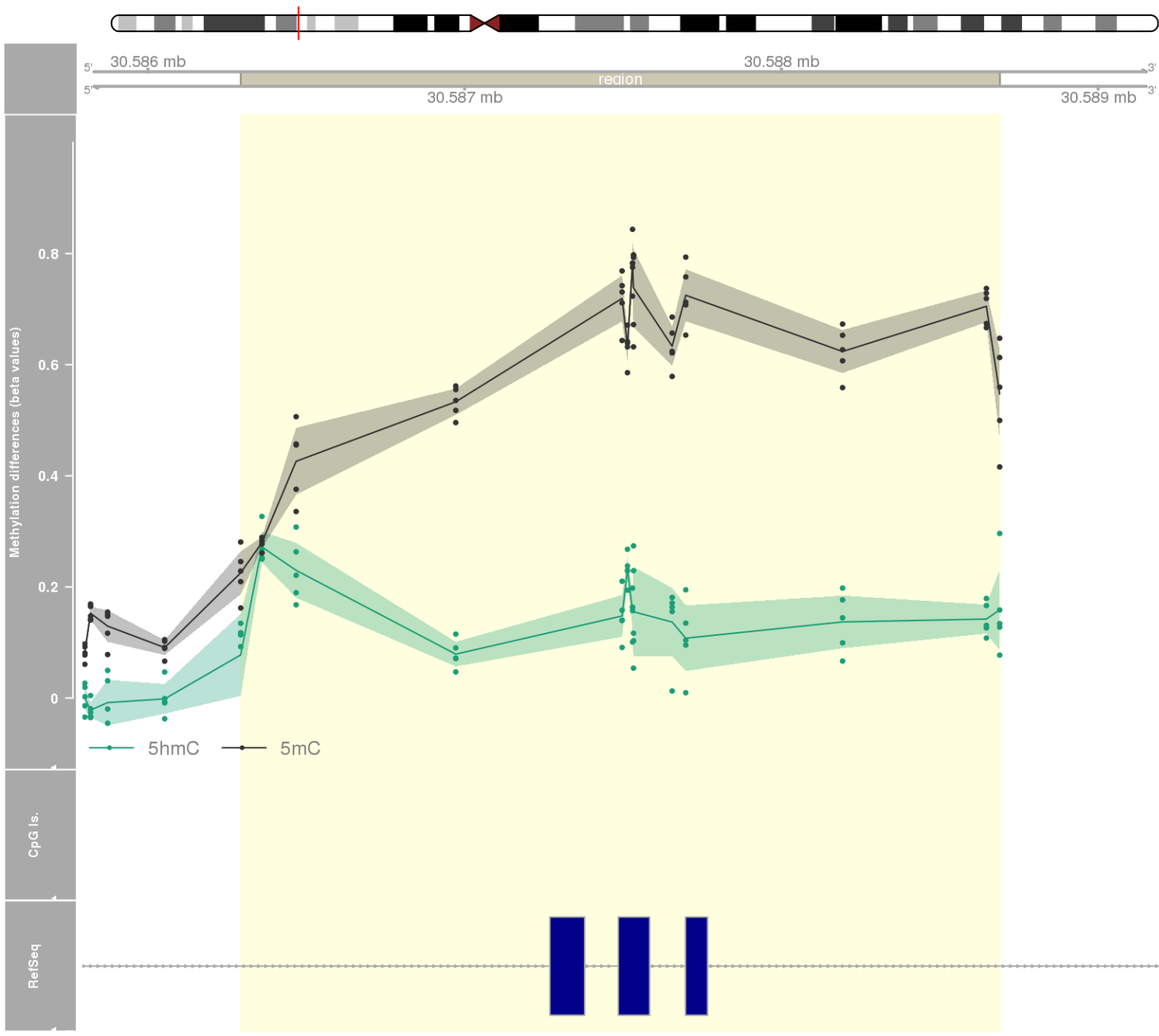
DMR 144 // chr6:30458237-30459595 // 1358 pb. (13 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.01 // fwerArea: 1
- genes: HLA-E -



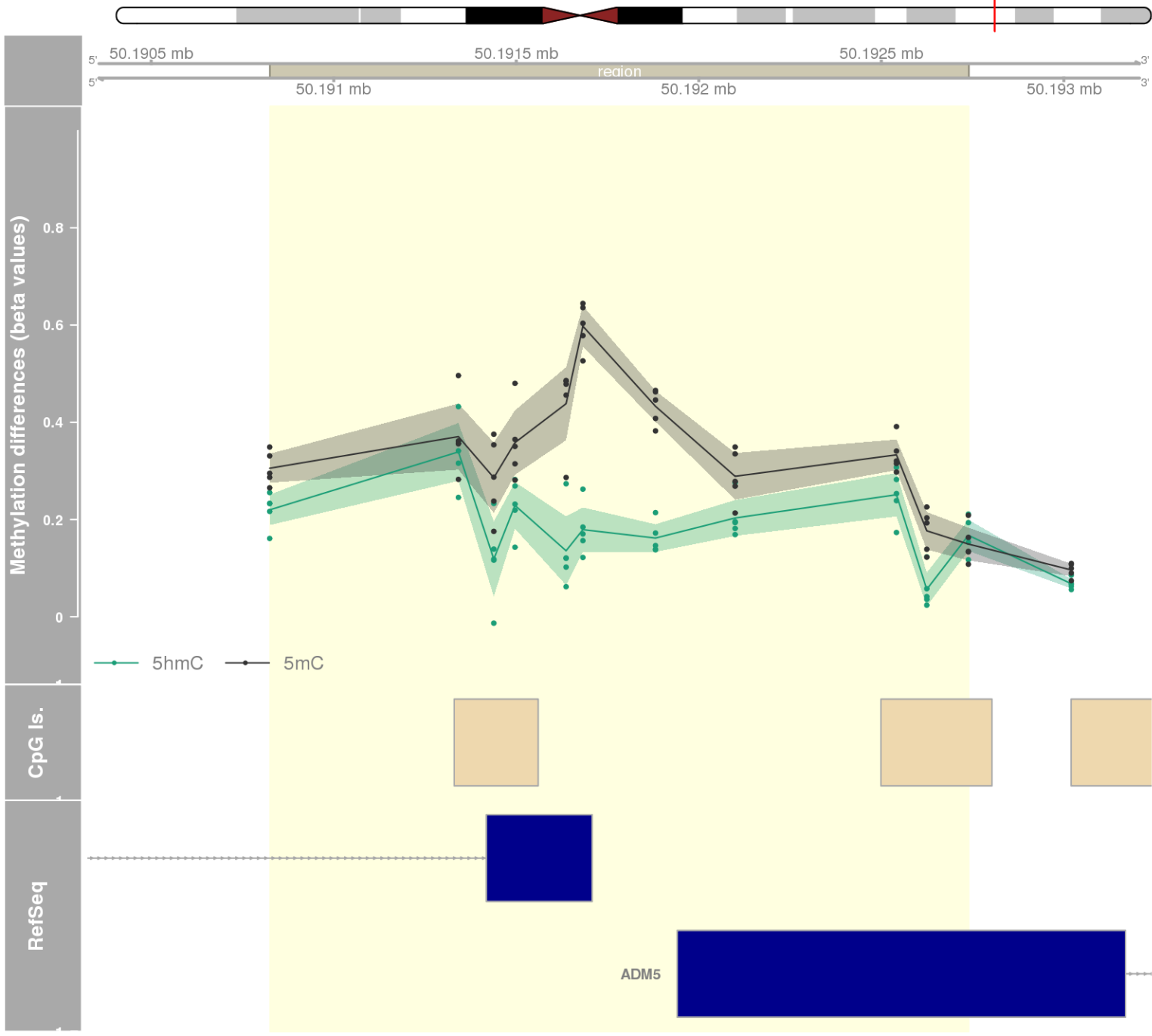
DMR 145 // chr12:133100010-133101612 // 1602 pb. (13 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.01 // fwerArea: 1
- genes: FBRSL1 -



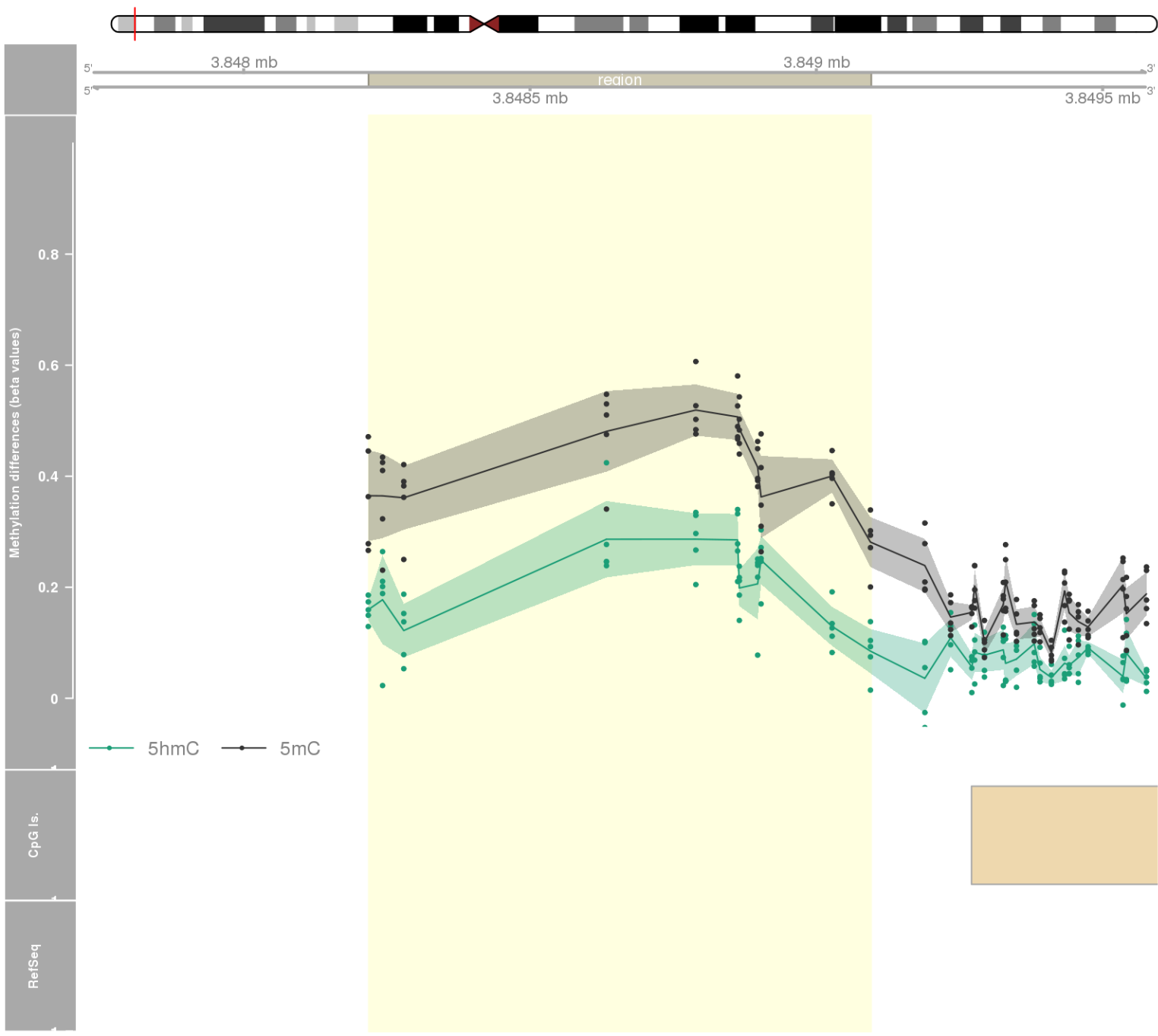
DMR 146 // chr6:30586293-30588689 // 2396 pb. (13 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.01 // fwerArea: 1
- genes: PPP1R10 / MRPS18B -



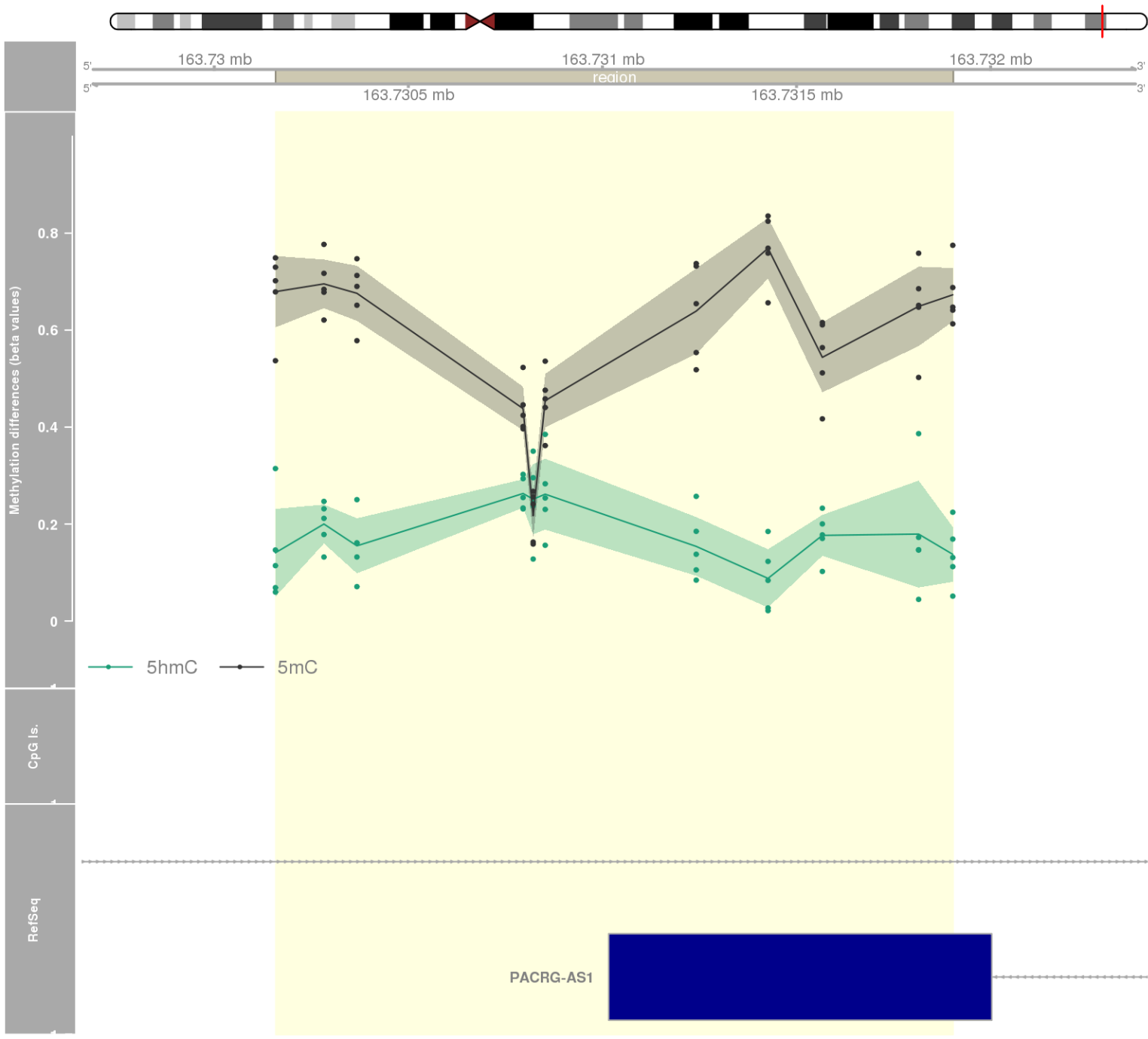
DMR 147 // chr19:50190825-50192739 // 1914 pb. (11 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.011 // fwerArea: 1
- genes: PRMT1 / ADM5 -



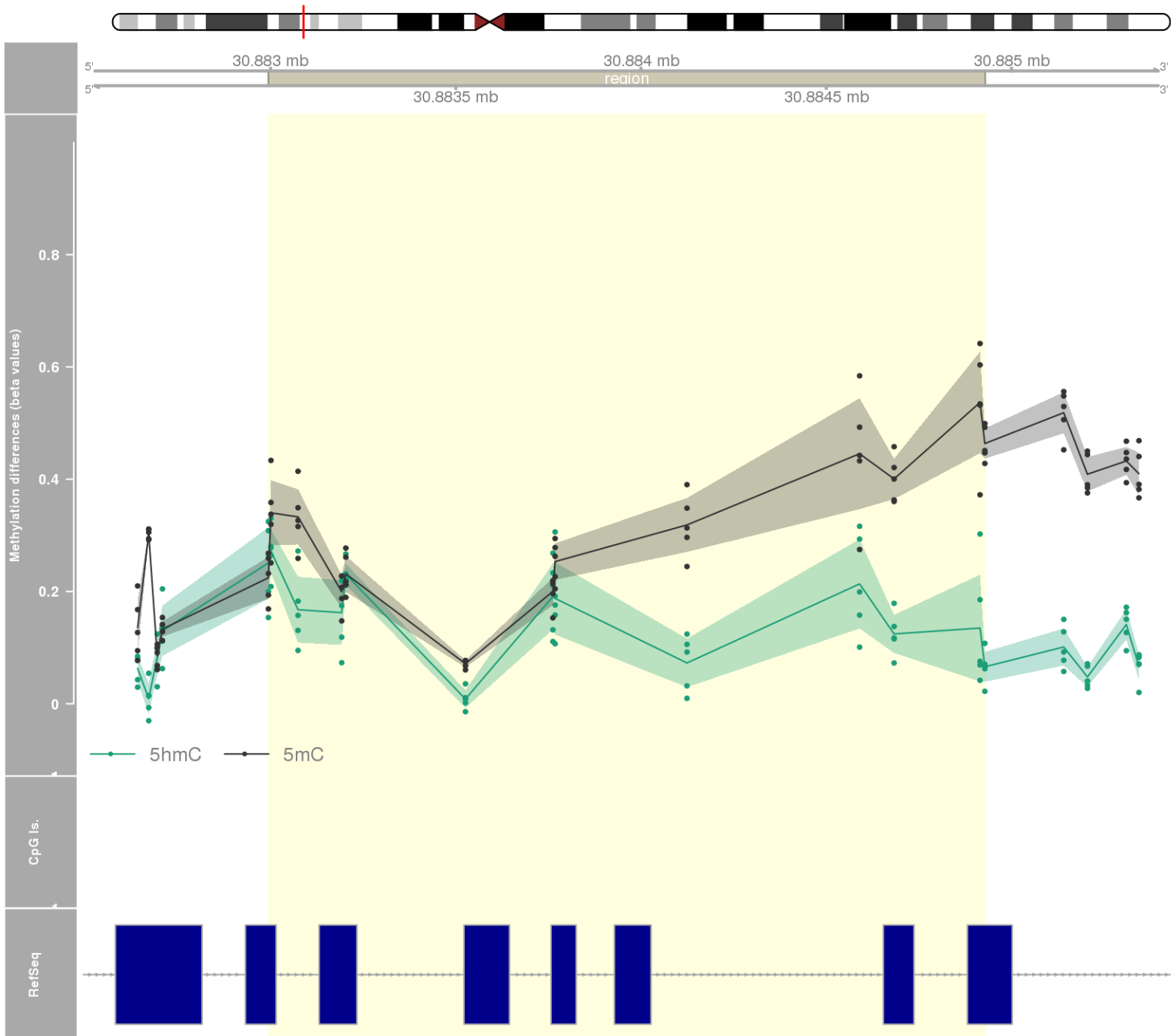
DMR 148 // chr6:3848218-3849095 // 877 pb. (11 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.011 // fwerArea: 1



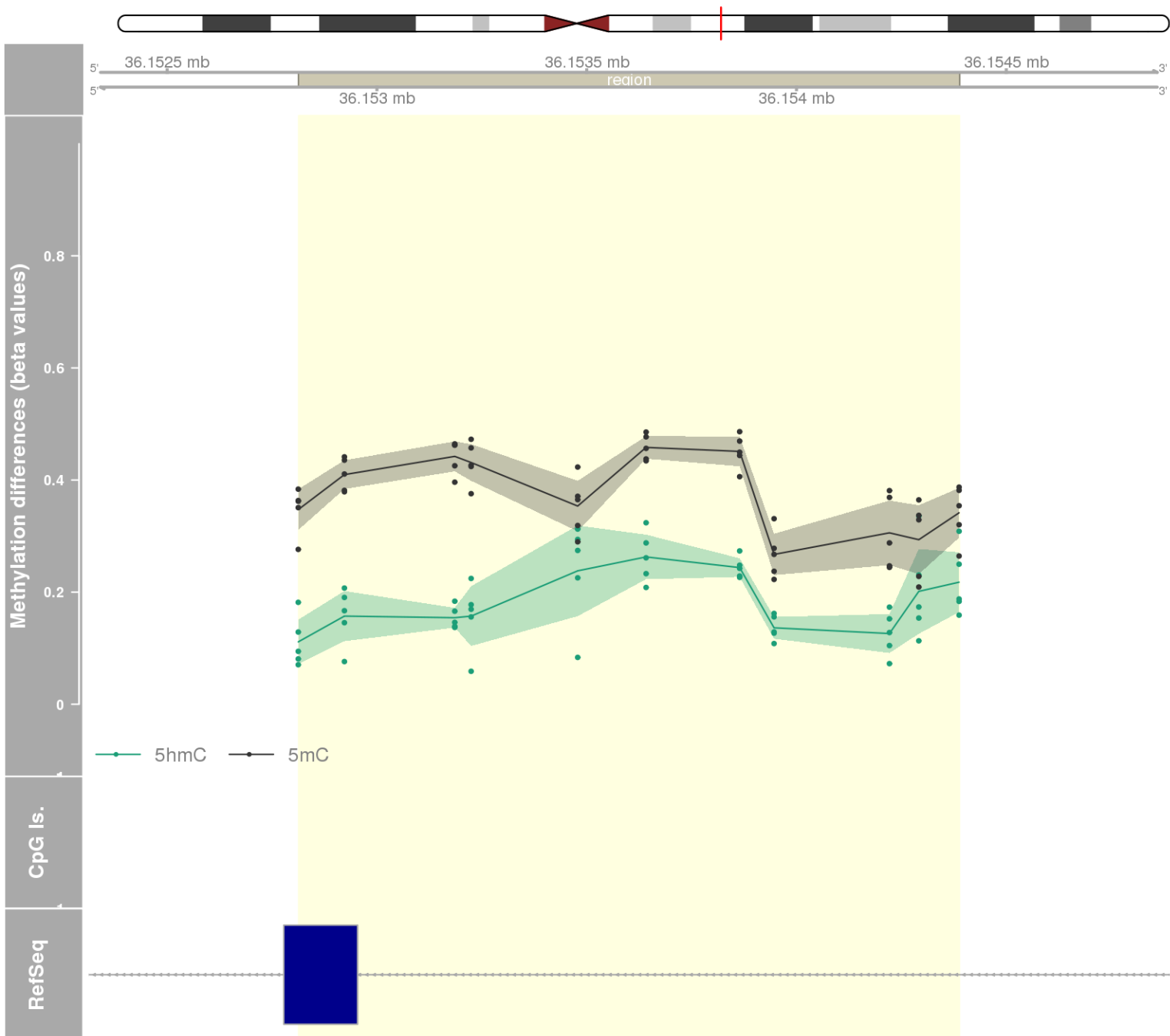
DMR 149 // chr6:163730158-163731903 // 1745 pb. (11 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.011 // fwerArea: 1
- genes: PACRG -



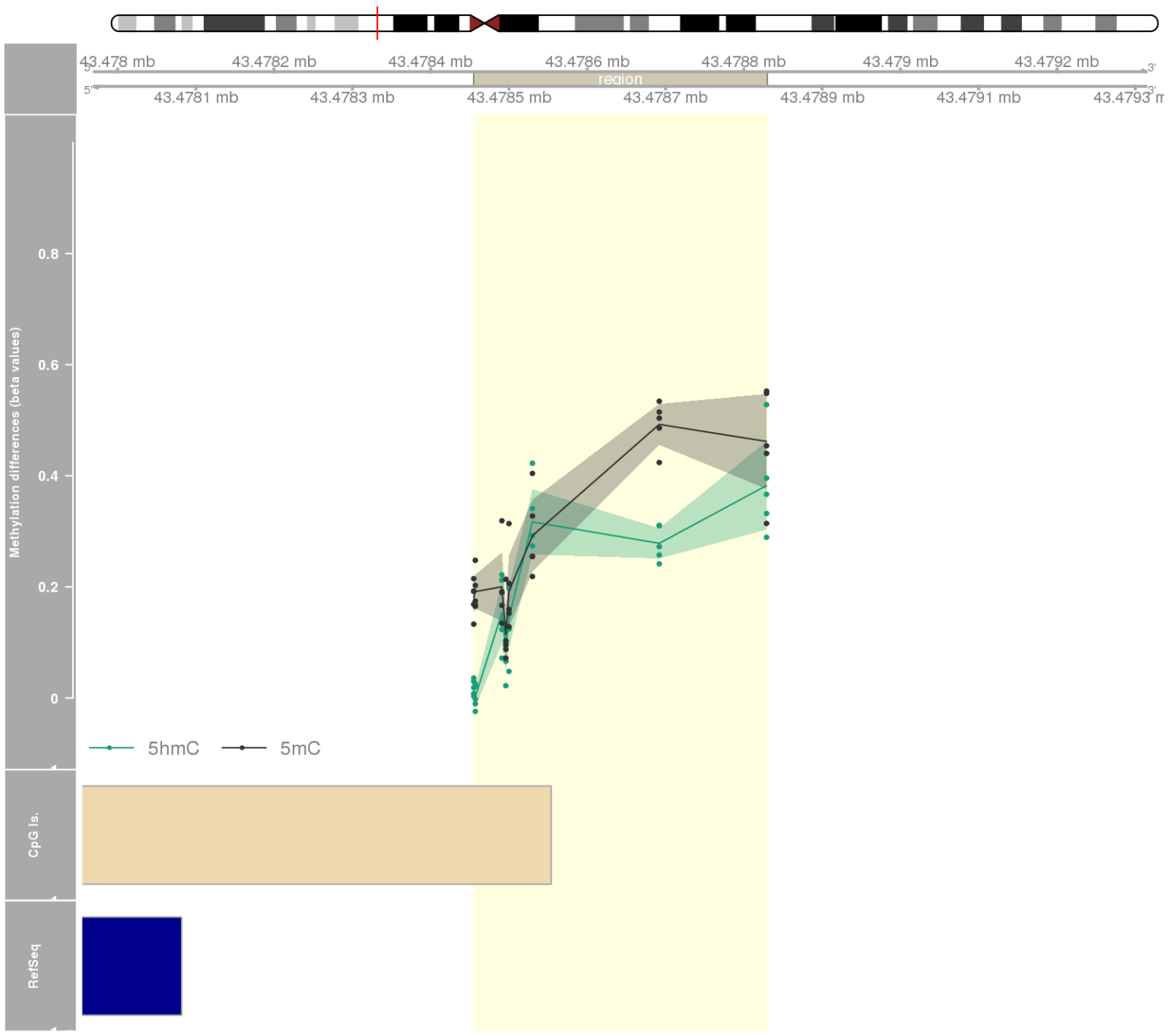
DMR 150 // chr6:30882994-30884928 // 1934 pb. (13 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.011 // fwerArea: 1
- genes: VARS2 -



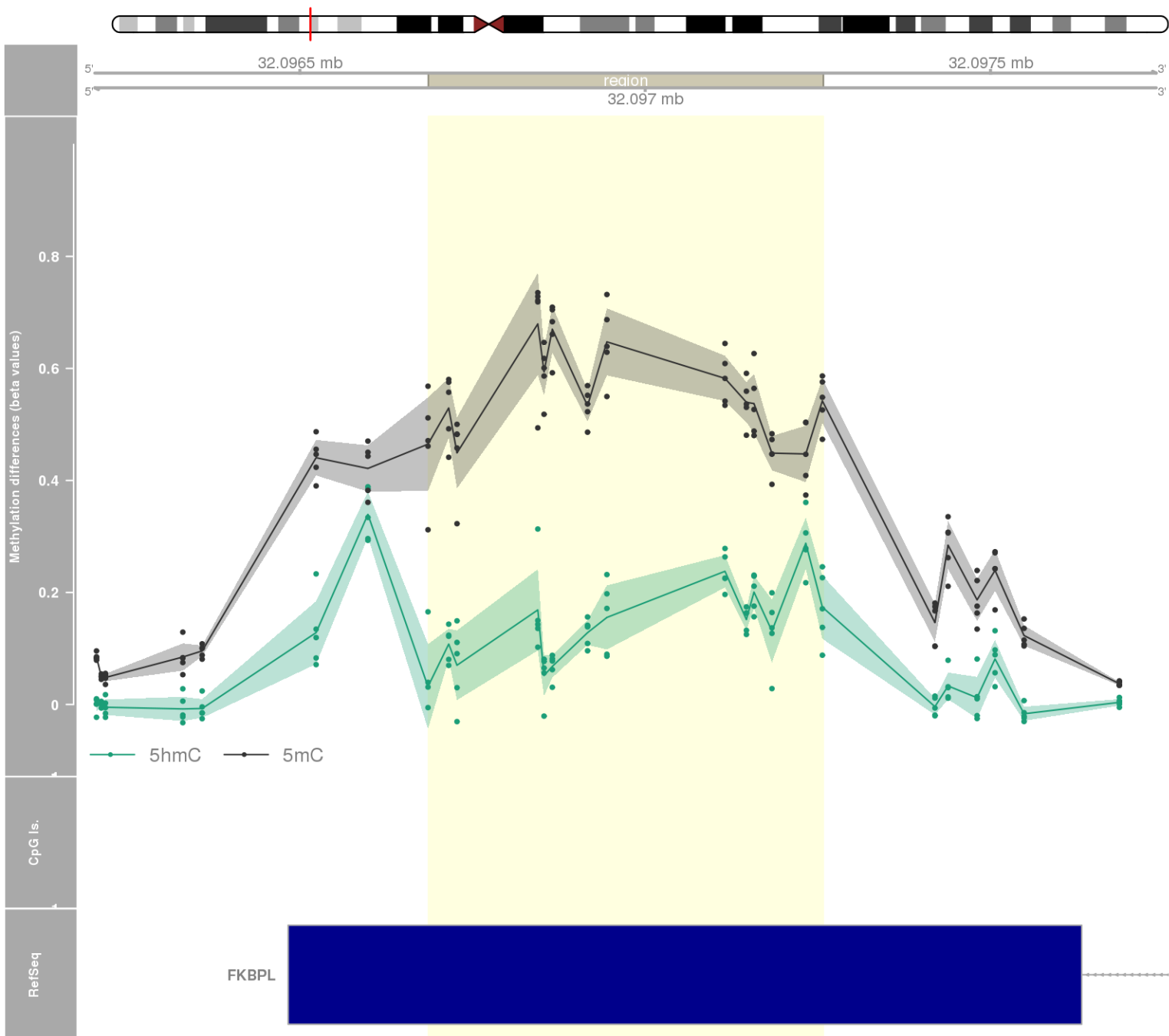
DMR 151 // chr20:36152813-36154388 // 1575 pb. (11 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.011 // fwerArea: 1
- genes: BLCAP -



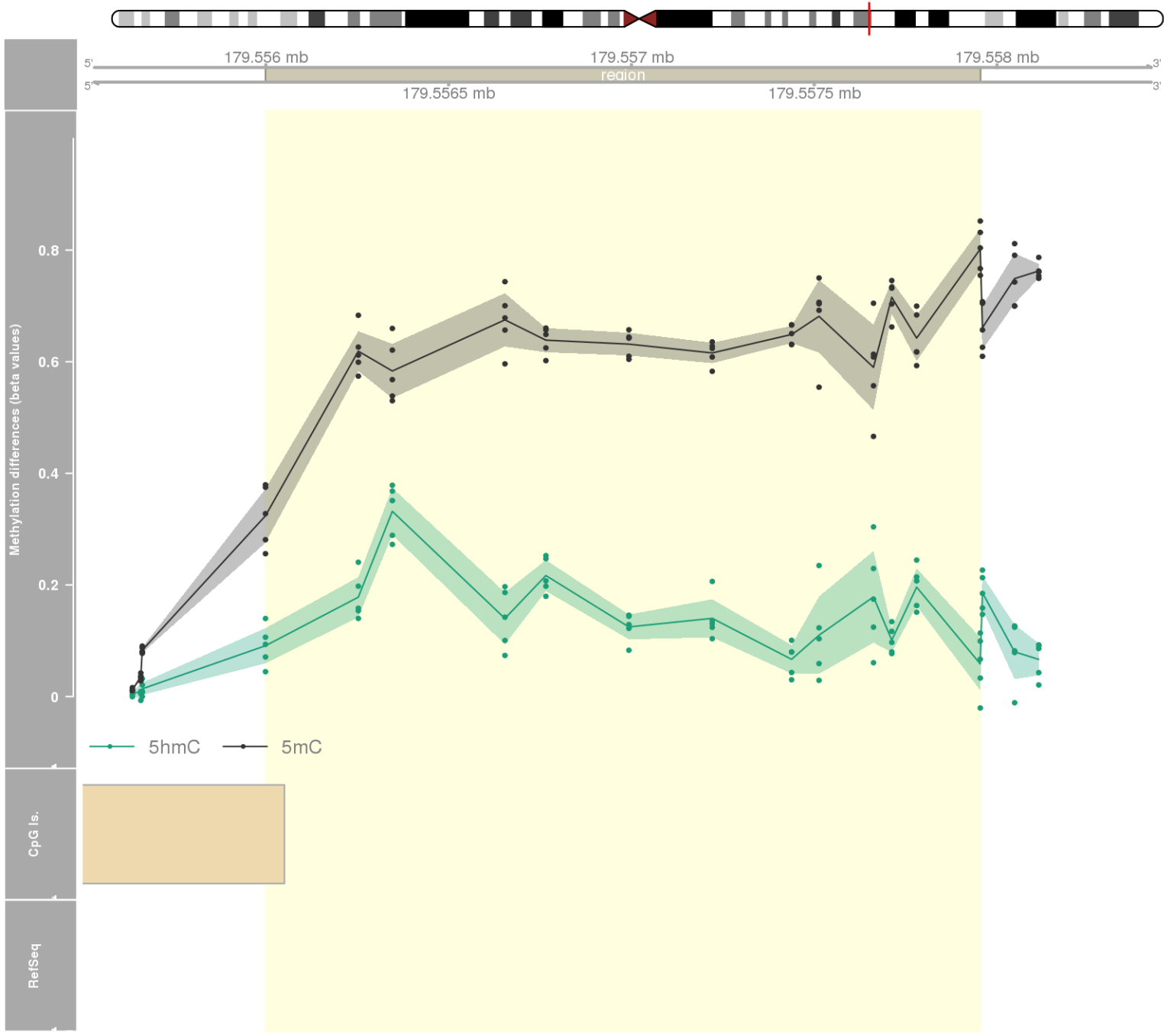
DMR 152 // chr6:43478455-43478829 // 374 pb. (8 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.011 // fwerArea: 1
- genes: POLR1C -



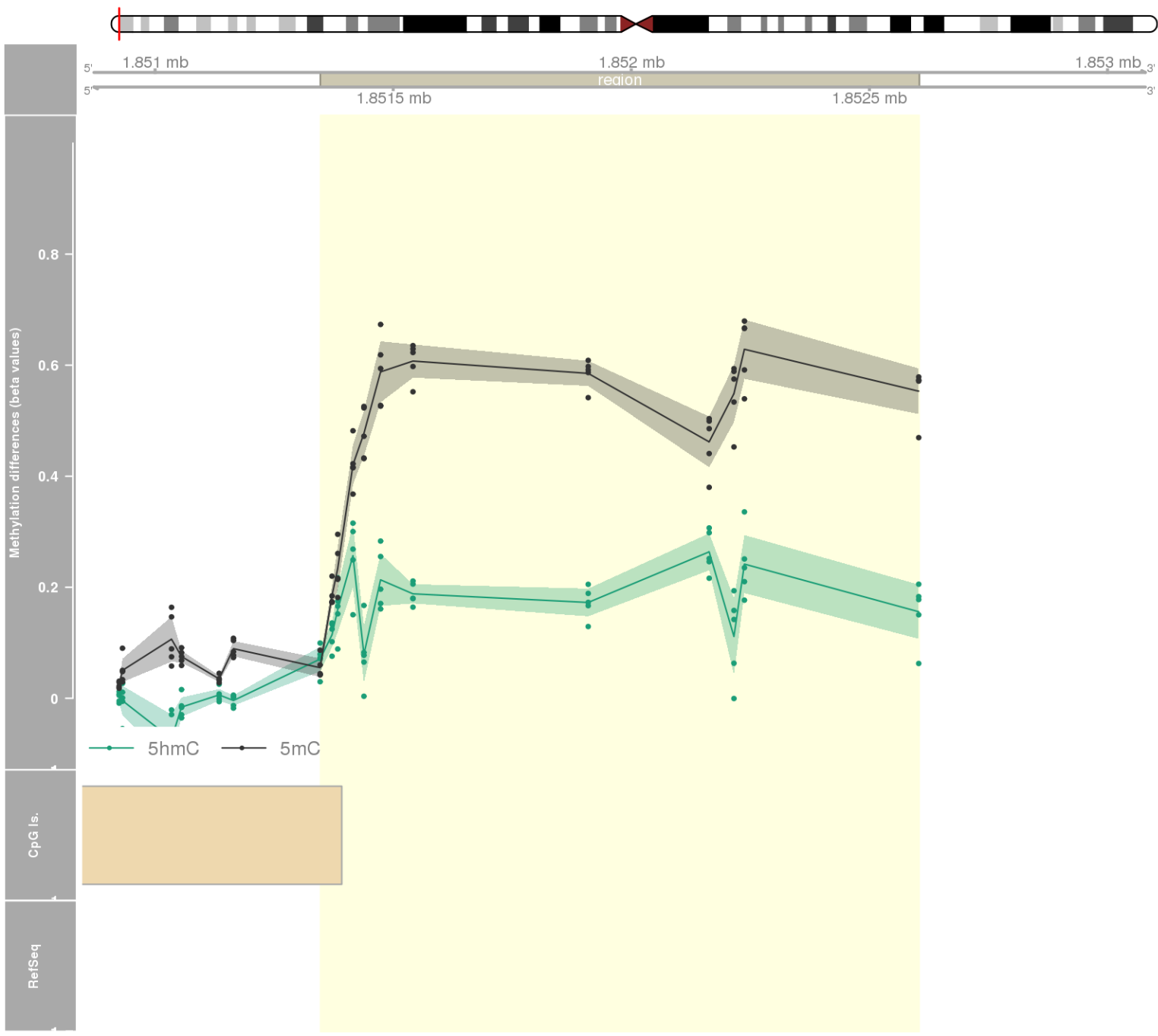
DMR 153 // chr6:32096686-32097257 // 571 pb. (14 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.011 // fwerArea: 1
- genes: FKBPL -



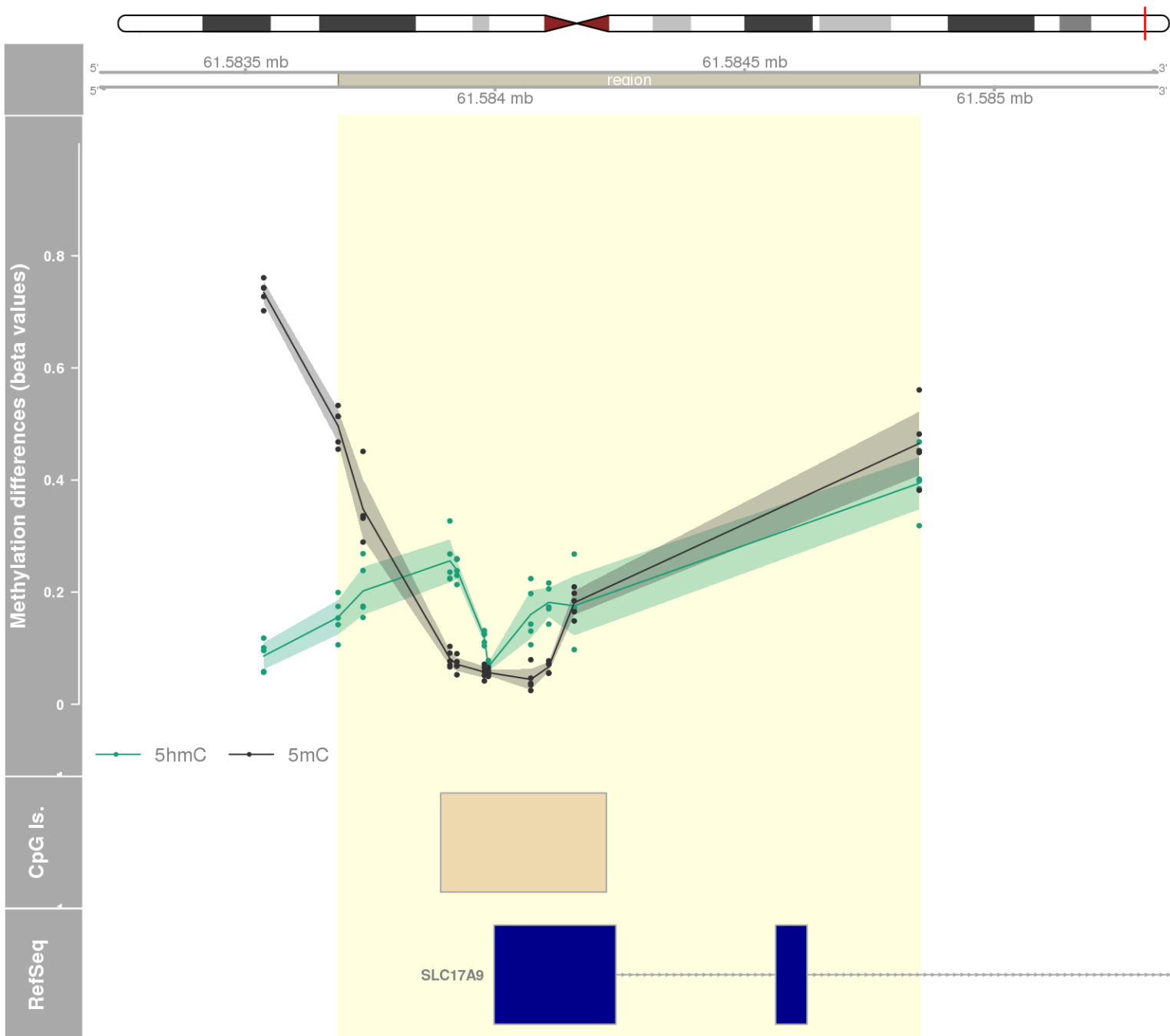
DMR 154 // chr1:179555999-179557954 // 1955 pb. (13 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.011 // fwerArea: 1



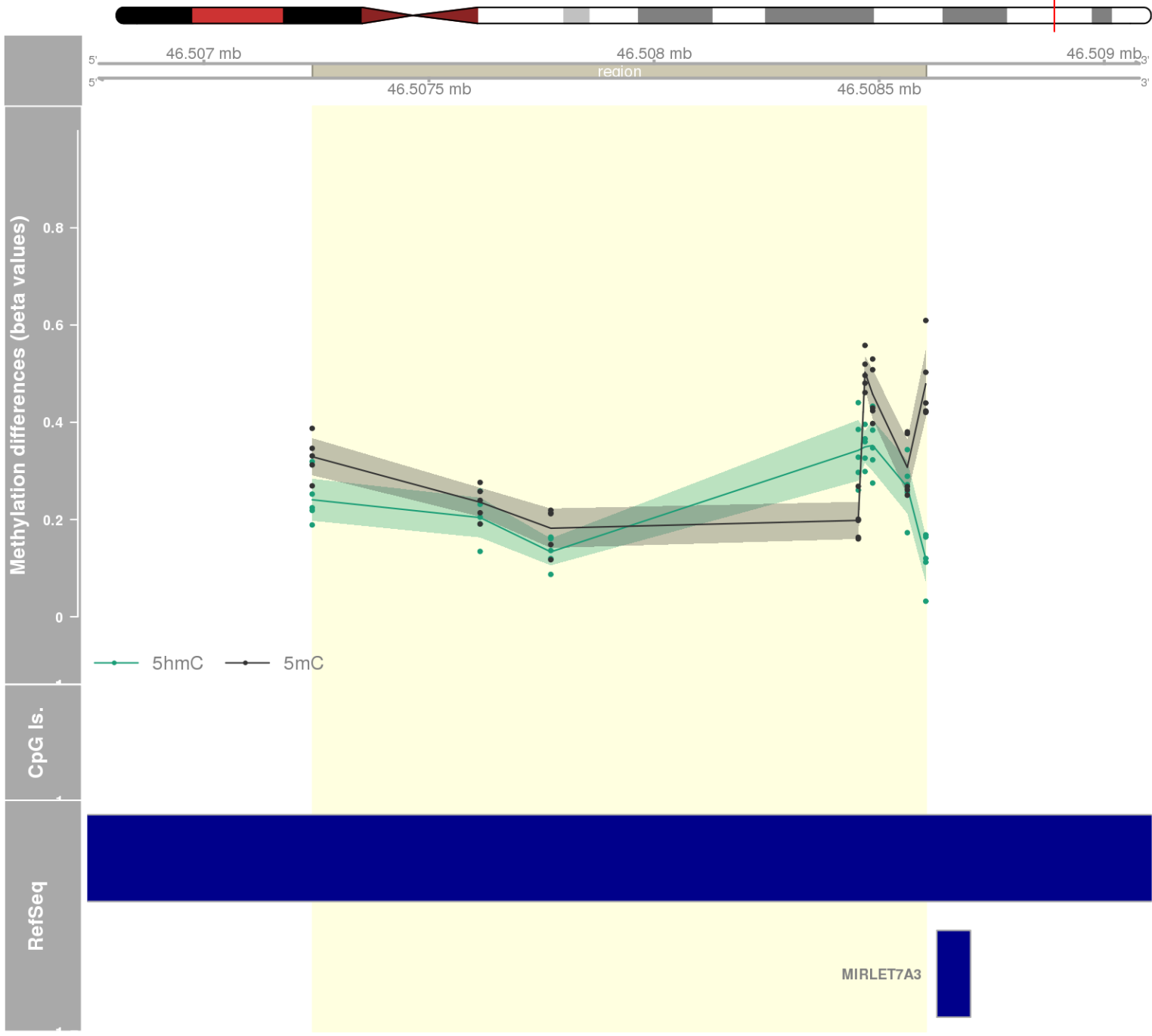
DMR 155 // chr1:1851347-1852604 // 1257 pb. (12 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.011 // fwerArea: 1



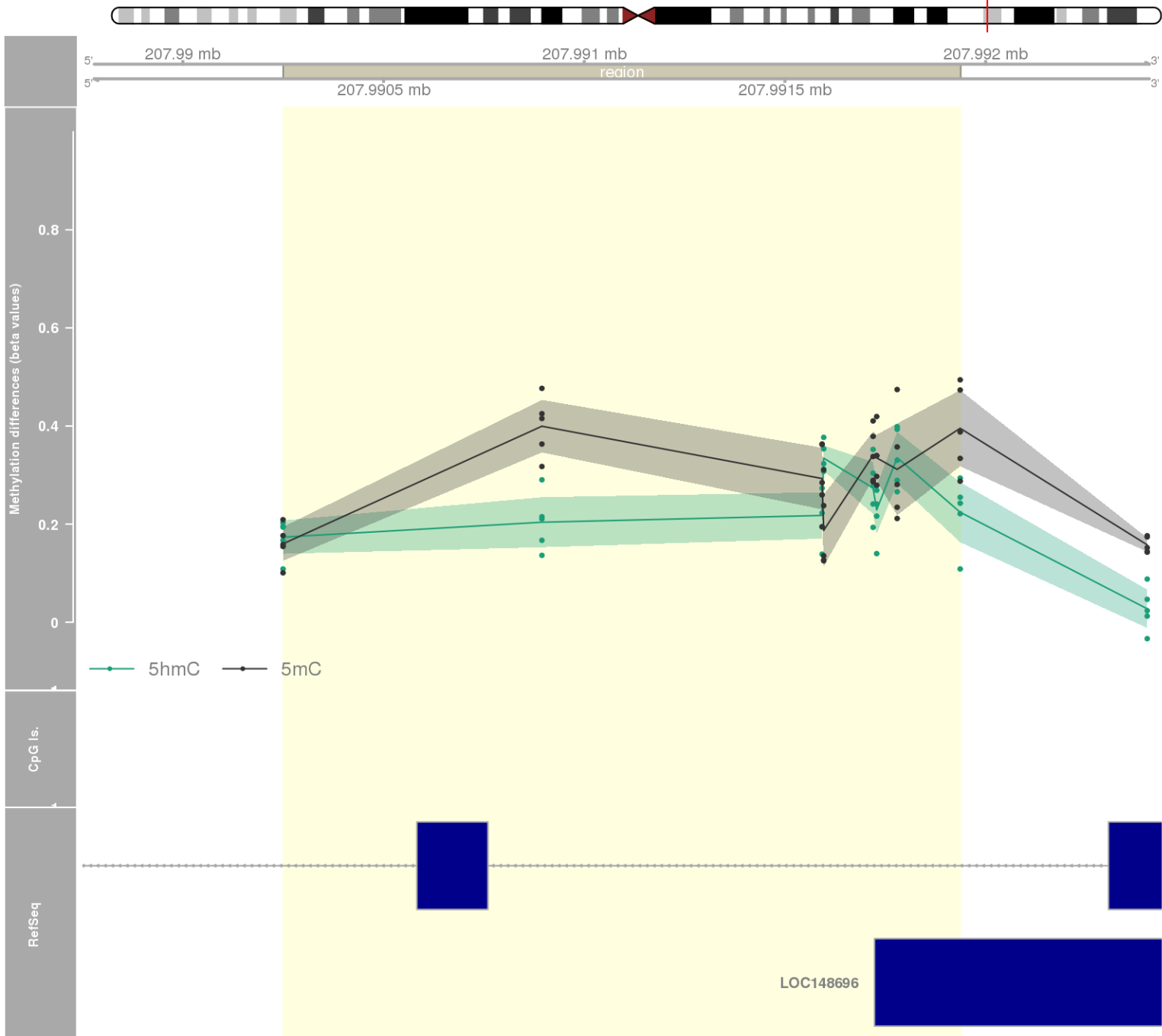
DMR 156 // chr20:61583686-61584850 // 1164 pb. (10 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.011 // fwerArea: 1
- genes: SLC17A9 -



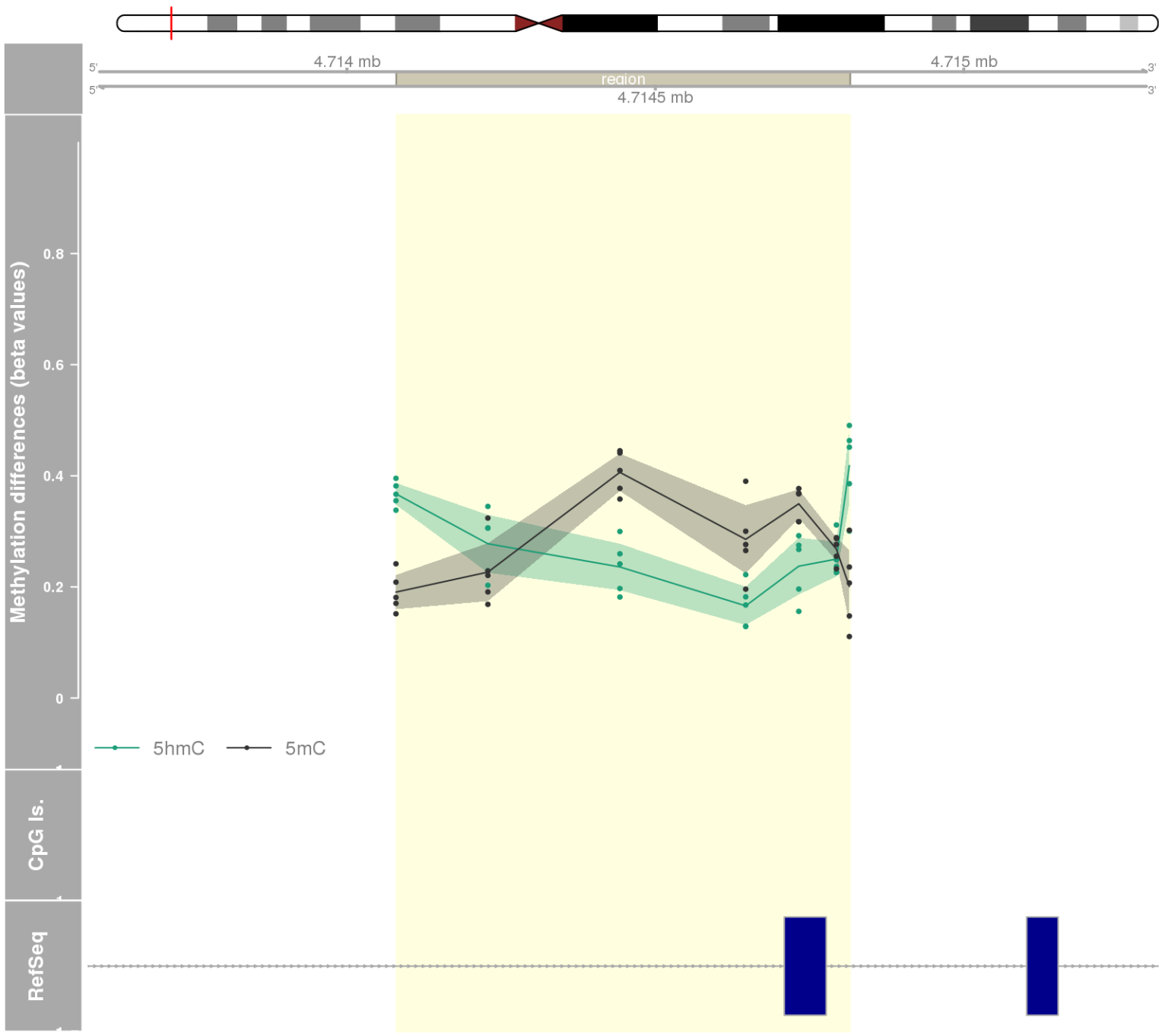
DMR 157 // chr22:46507241-46508604 // 1363 pb. (8 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.011 // fwerArea: 1



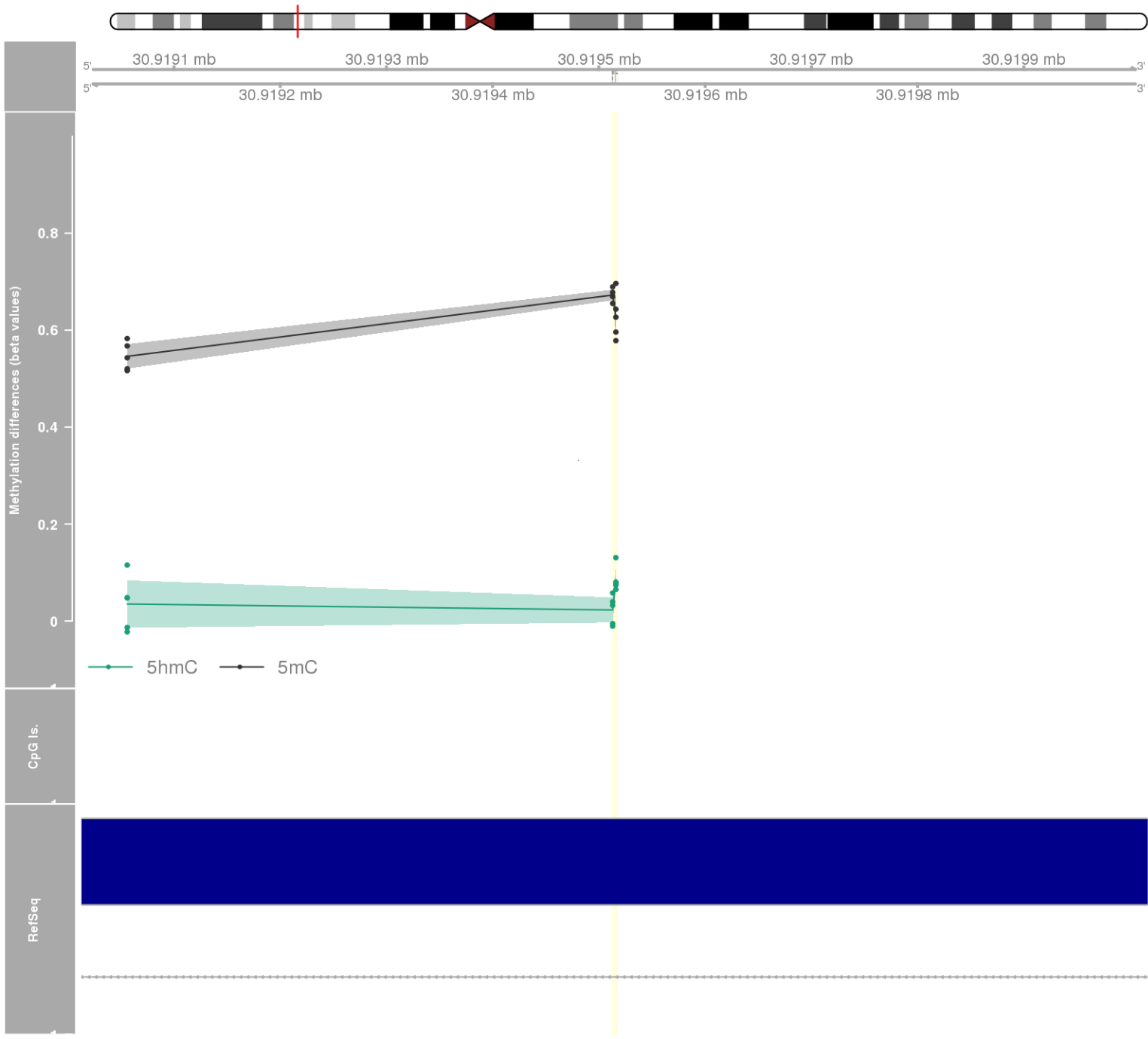
DMR 158 // chr1:207990250-207991937 // 1687 pb. (8 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.011 // fwerArea: 1
- genes: C1orf132 -



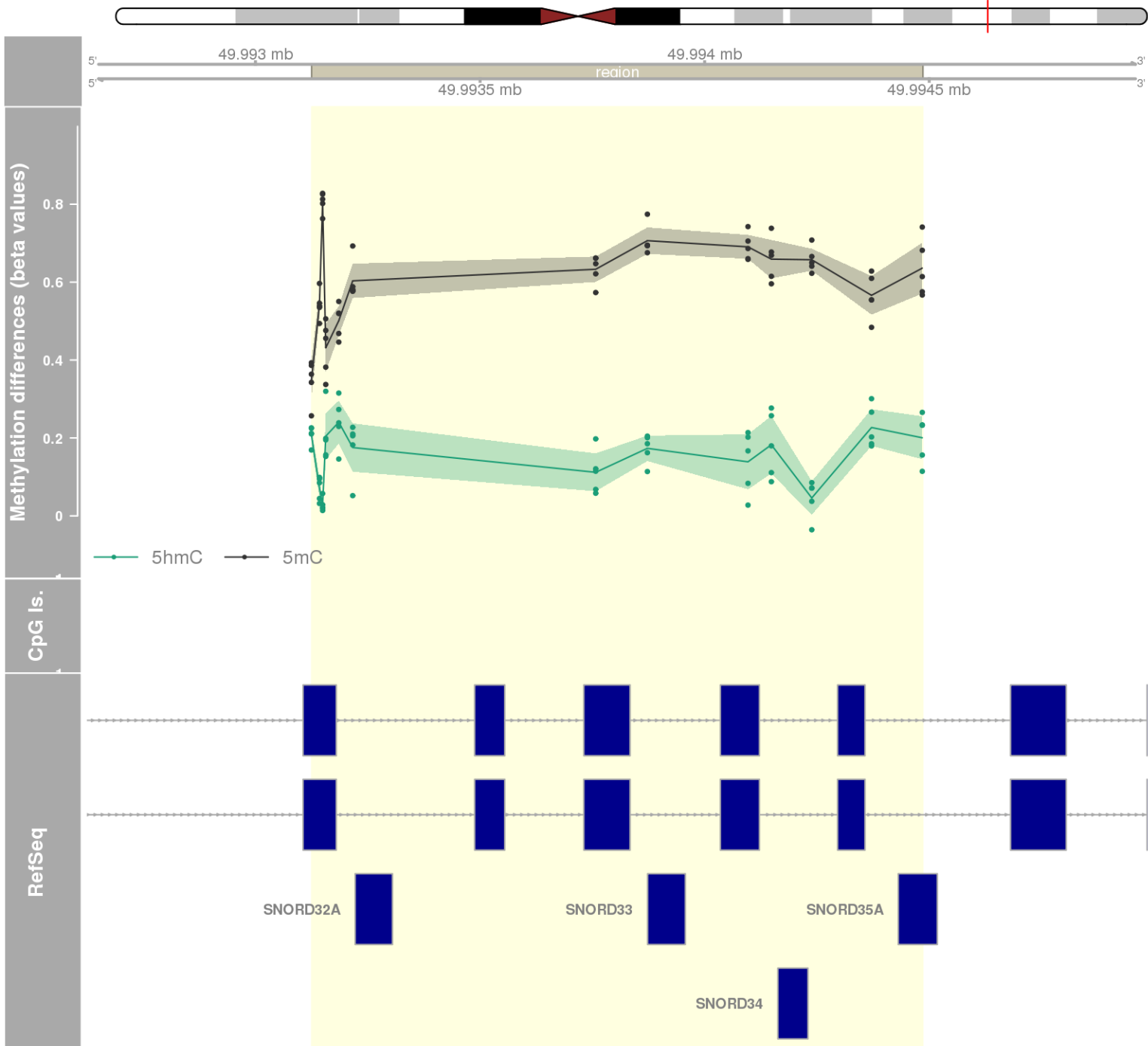
DMR 159 // chr16:4714080-4714815 // 735 pb. (7 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.011 // fwerArea: 1
- genes: MGRN1 -



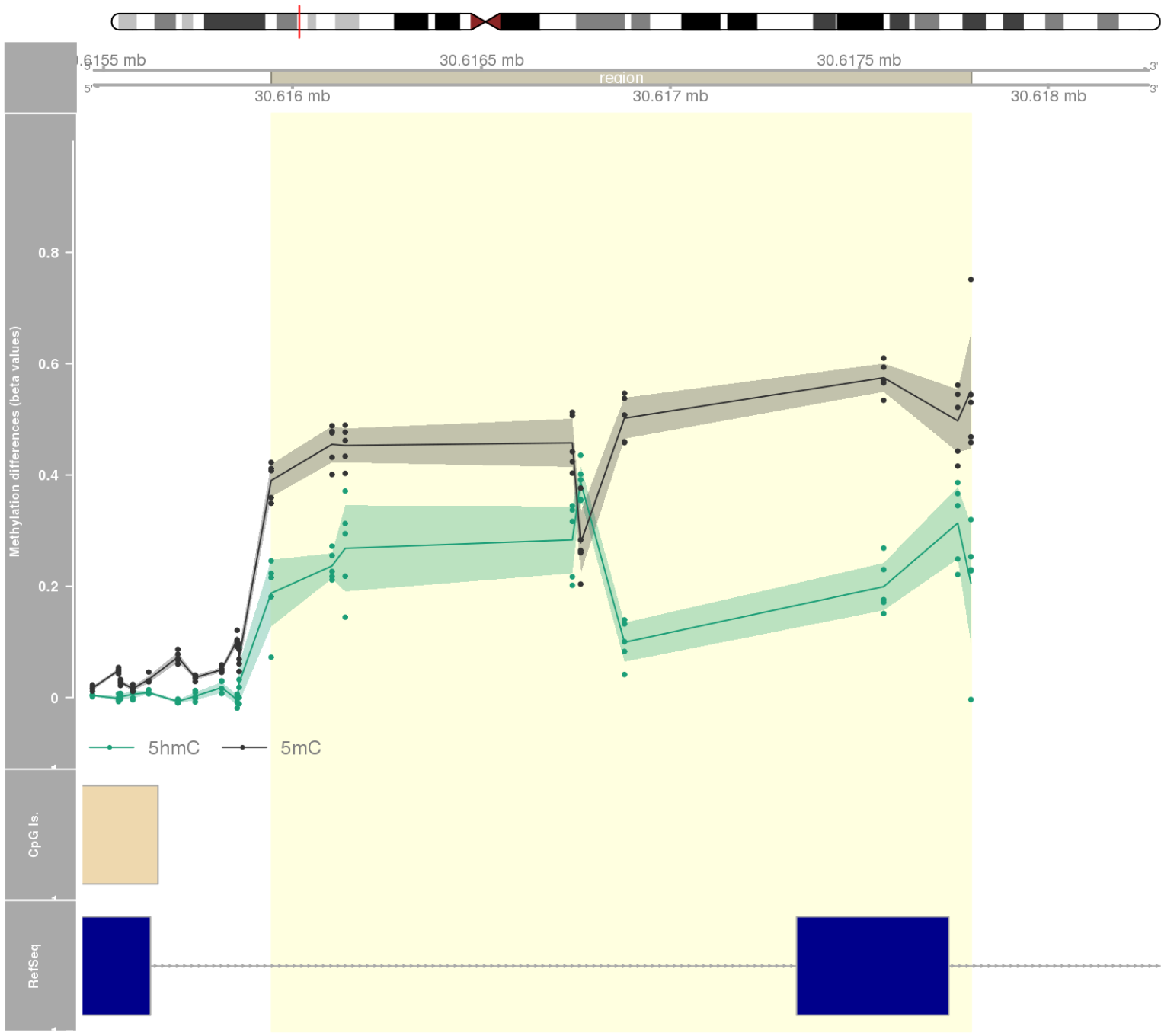
DMR 160 // chr6:30919513-30919516 // 3 pb. (2 probes) // pvalue: 0 // fwer: 0.497 // pvalueArea: 0.011 // fwerArea: 1
- genes: HCG21 / DPCR1 -



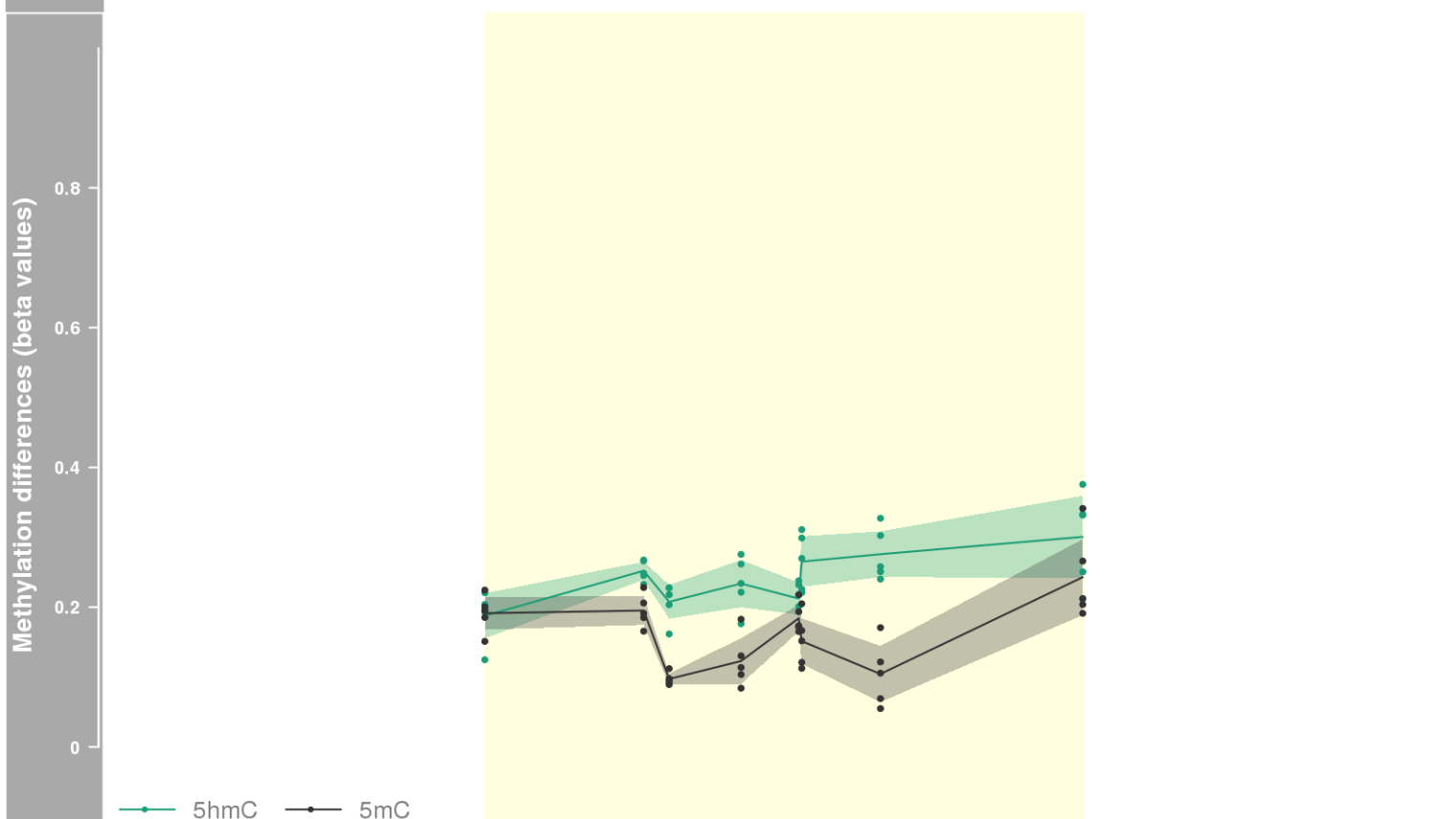
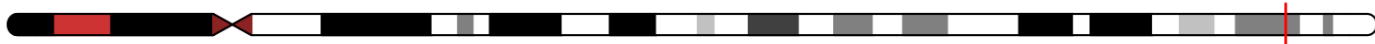
DMR 161 // chr19:49993125-49994485 // 1360 pb. (13 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.011 // fwerArea: 1
- genes: SNORD32A / SNORD33 / SNORD34 / SNORD35A / RPL13A -



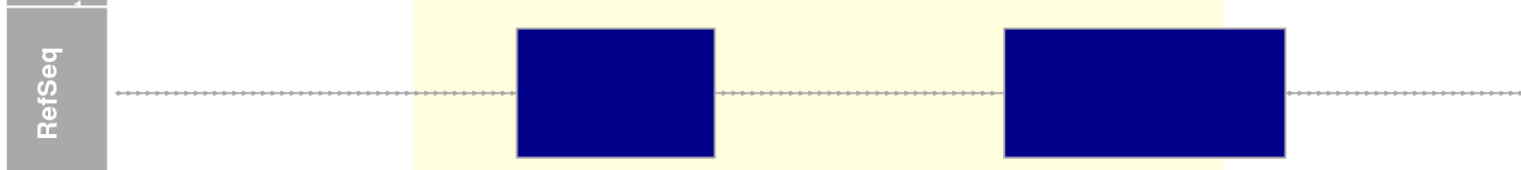
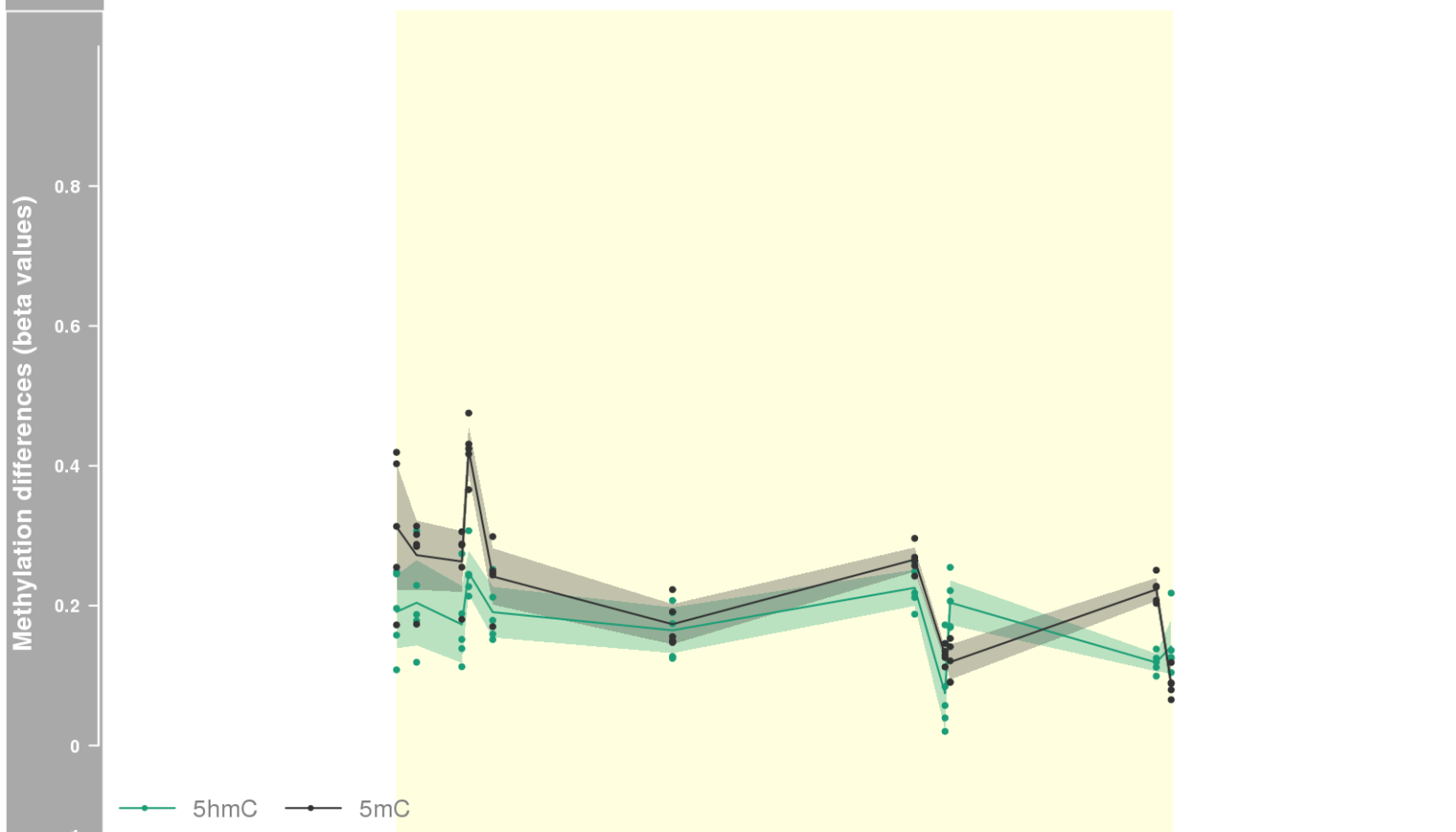
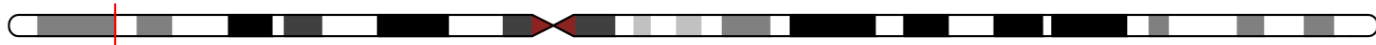
DMR 162 // chr6:30615944-30617796 // 1852 pb. (9 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.012 // fwerArea: 1
- genes: C6orf136 -



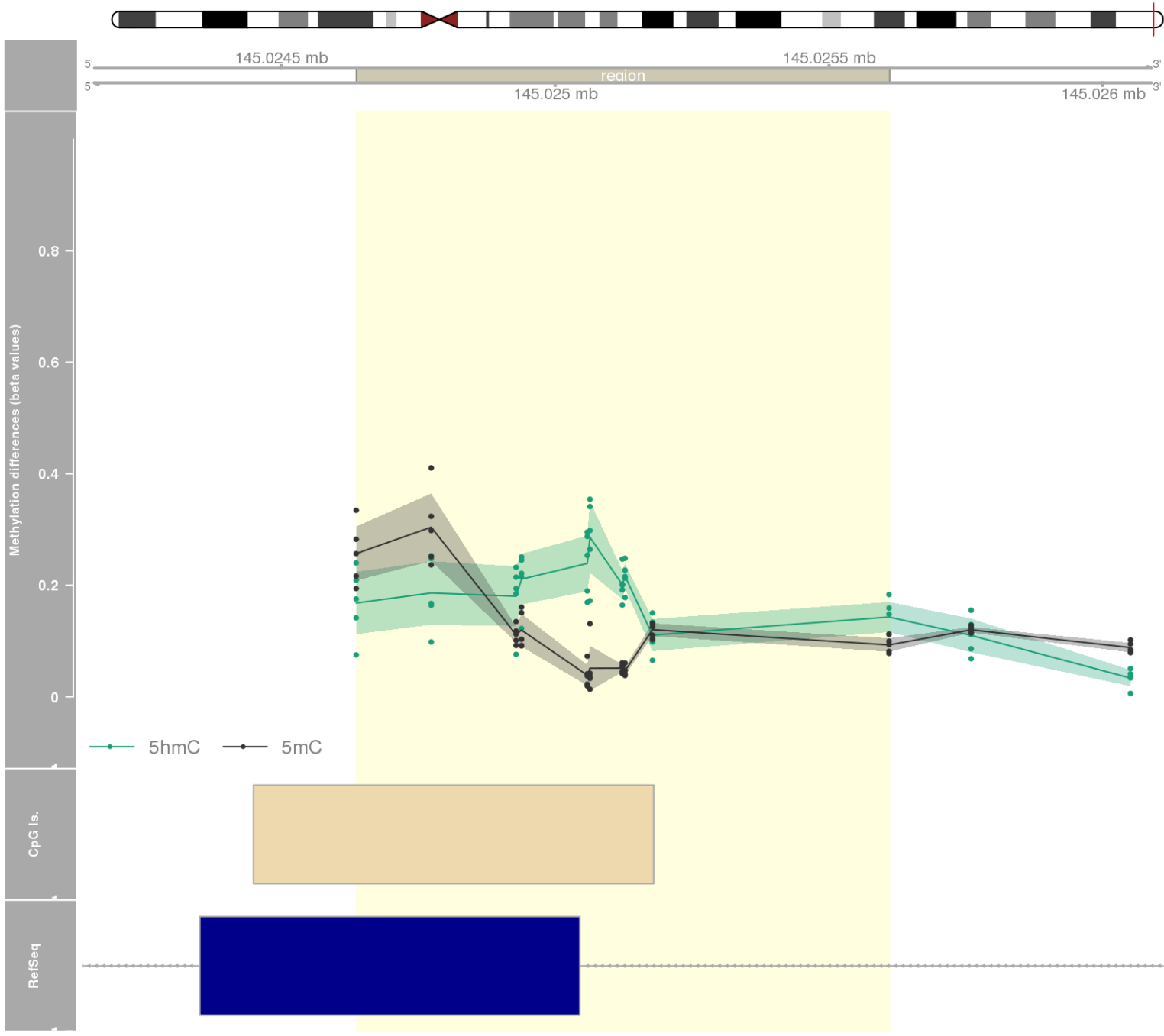
DMR 163 // chr14:100203730-100204528 // 798 pb. (8 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.012 // fwerArea: 1
- genes: EML1 -



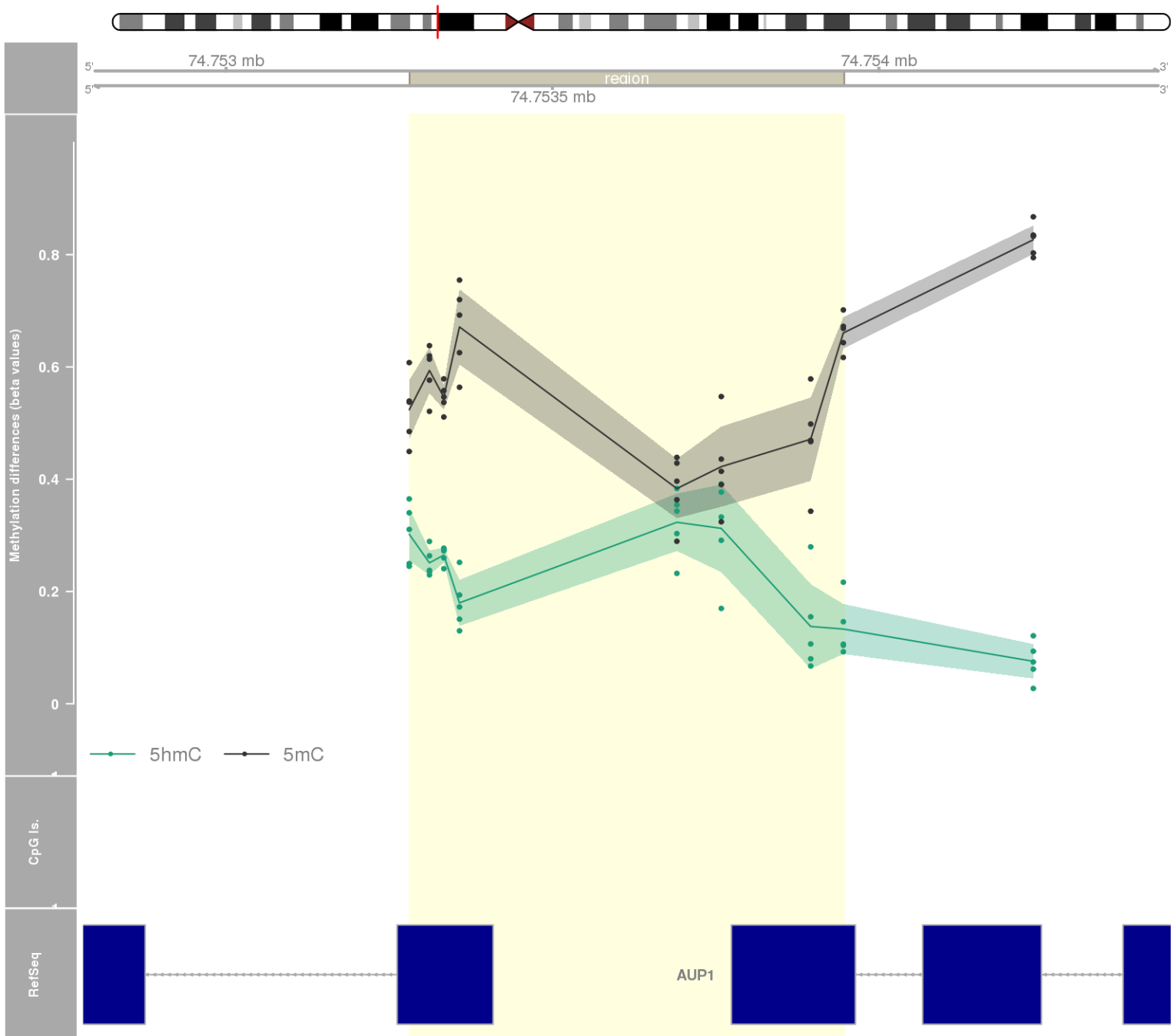
DMR 164 // chr11:10476494-10477847 // 1353 pb. (11 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.012 // fwerArea: 1
- genes: AMPD3 -



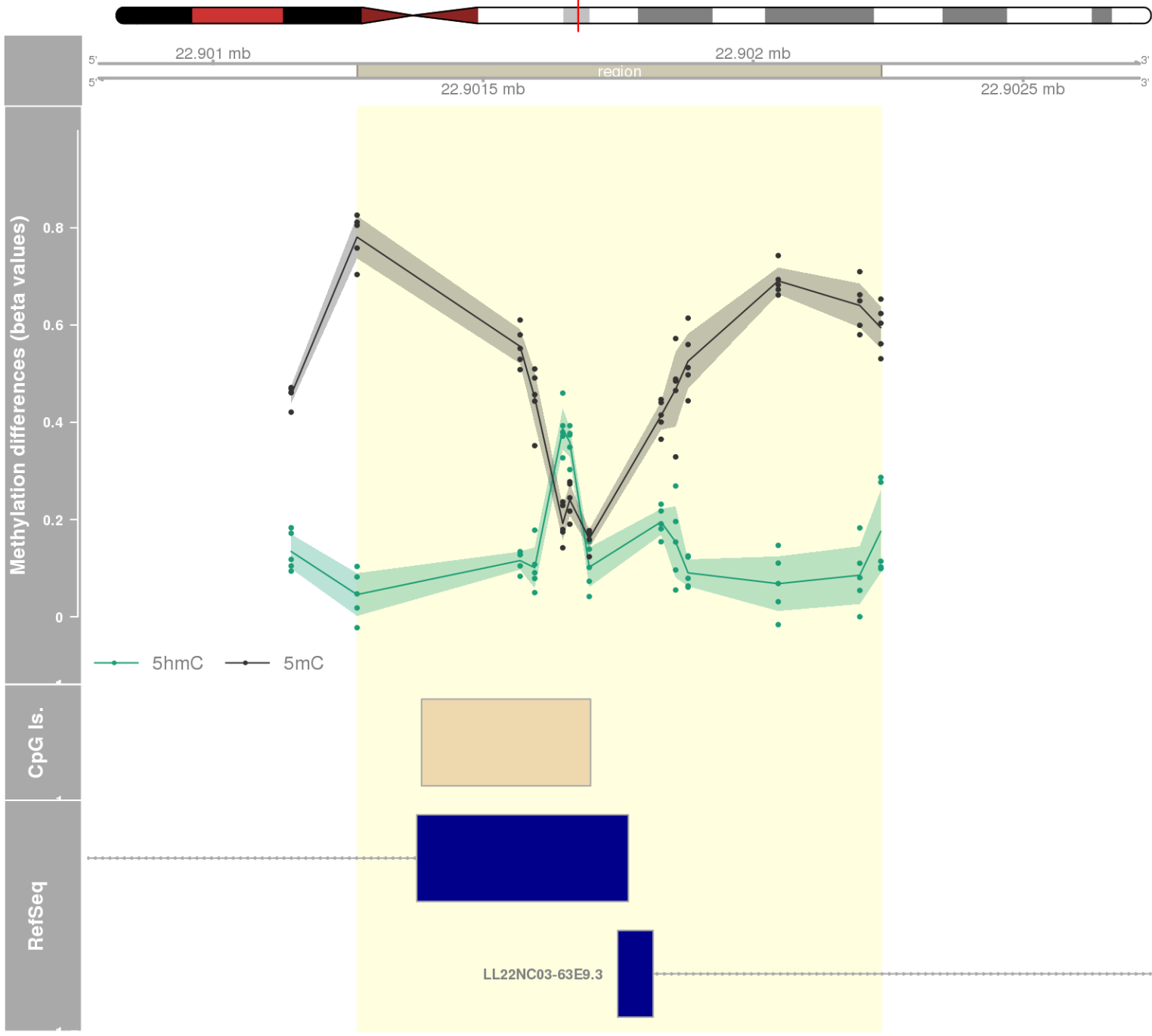
DMR 165 // chr8:145024637-145025610 // 973 pb. (10 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.012 // fwerArea: 1
- genes: PLEC -



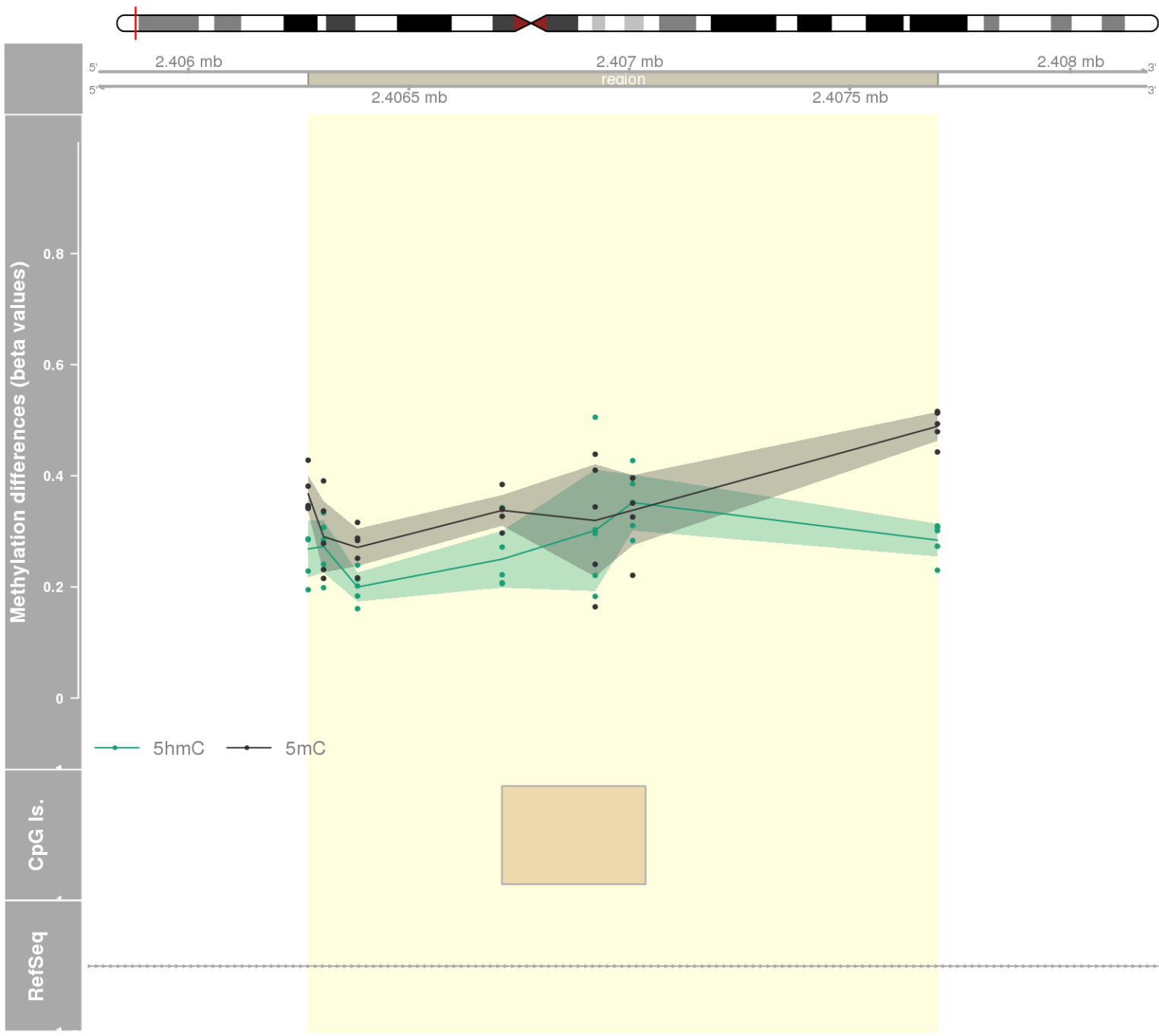
DMR 166 // chr2:74753281-74753946 // 665 pb. (8 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.012 // fwerArea: 1
- genes: DQX1 / AUP1 -



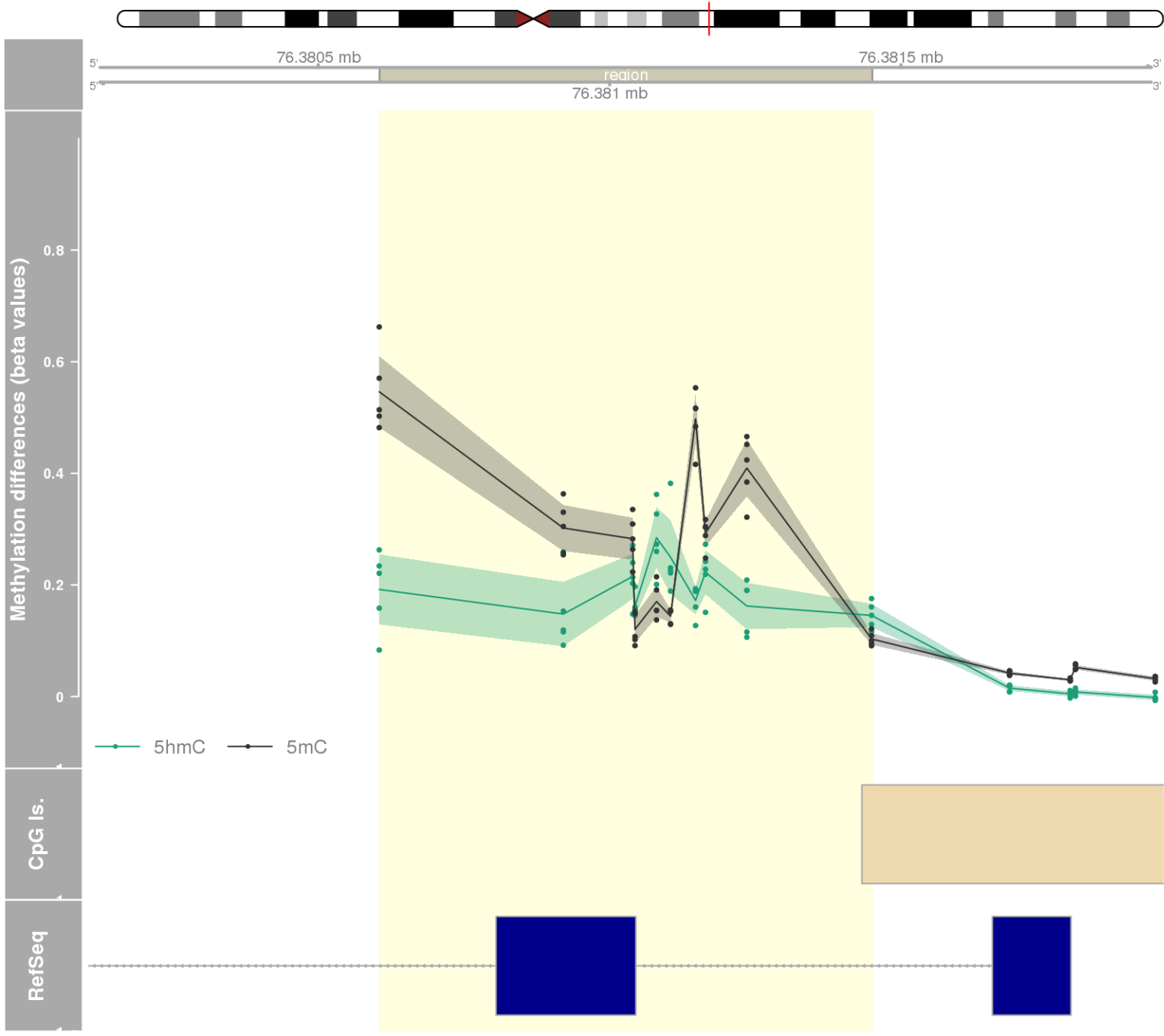
DMR 167 // chr22:22901267-22902237 // 970 pb. (12 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.012 // fwerArea: 1
- genes: PRAME -



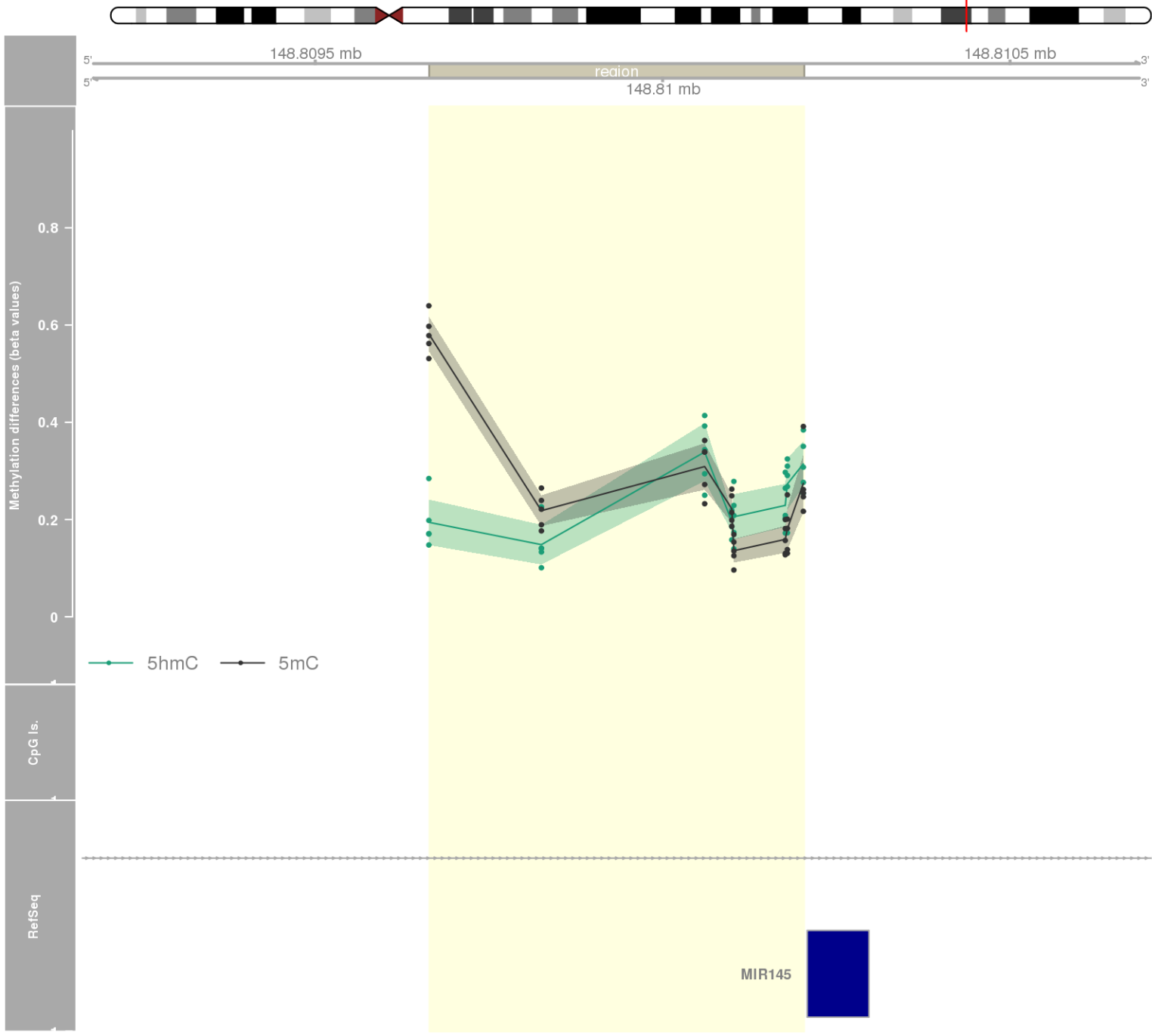
DMR 168 // chr11:2406272-2407699 // 1427 pb. (7 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.012 // fwerArea: 1
- genes: CD81 -



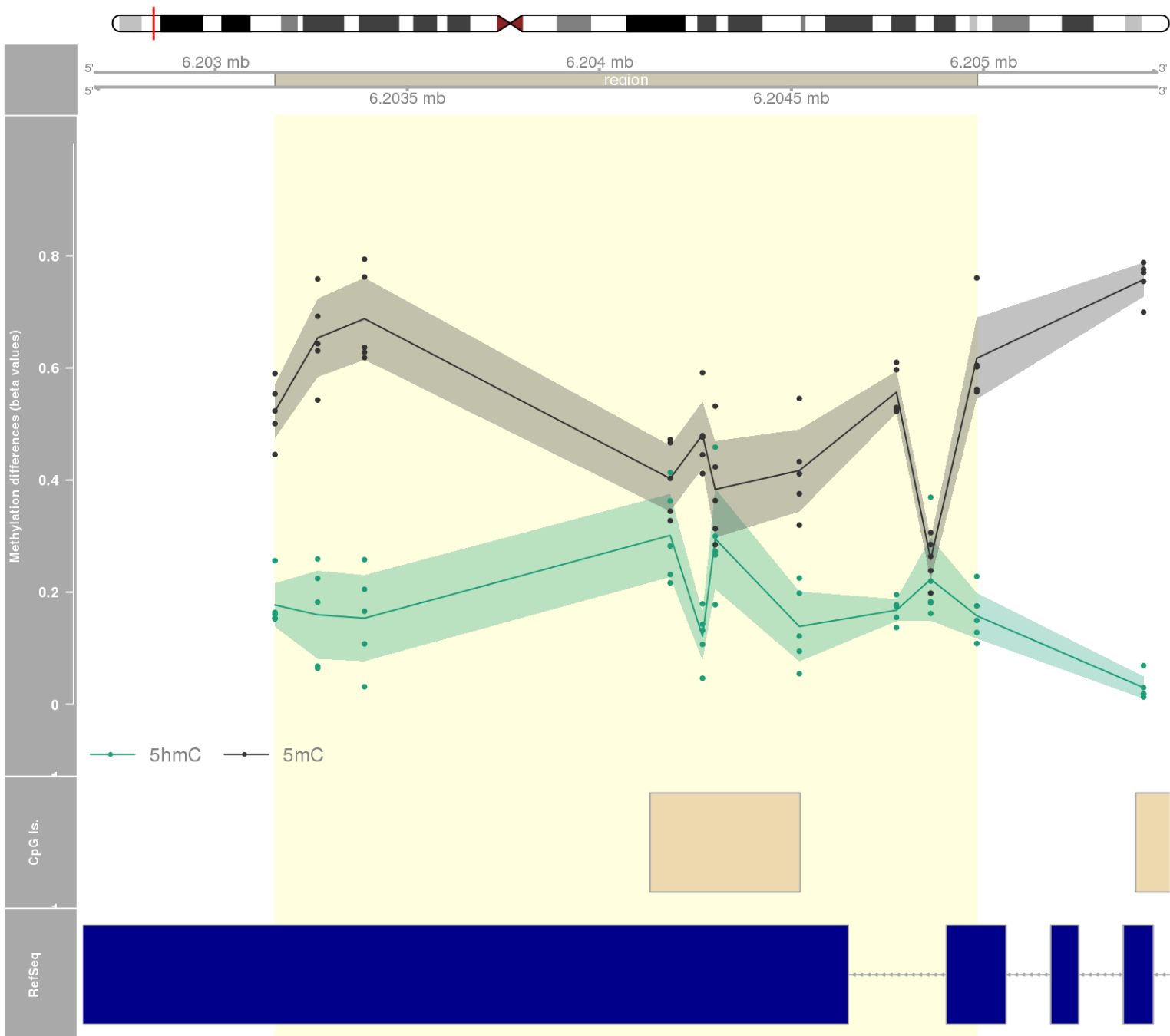
DMR 169 // chr11:76380605-76381450 // 845 pb. (10 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.012 // fwerArea: 1
- genes: LRRC32 -



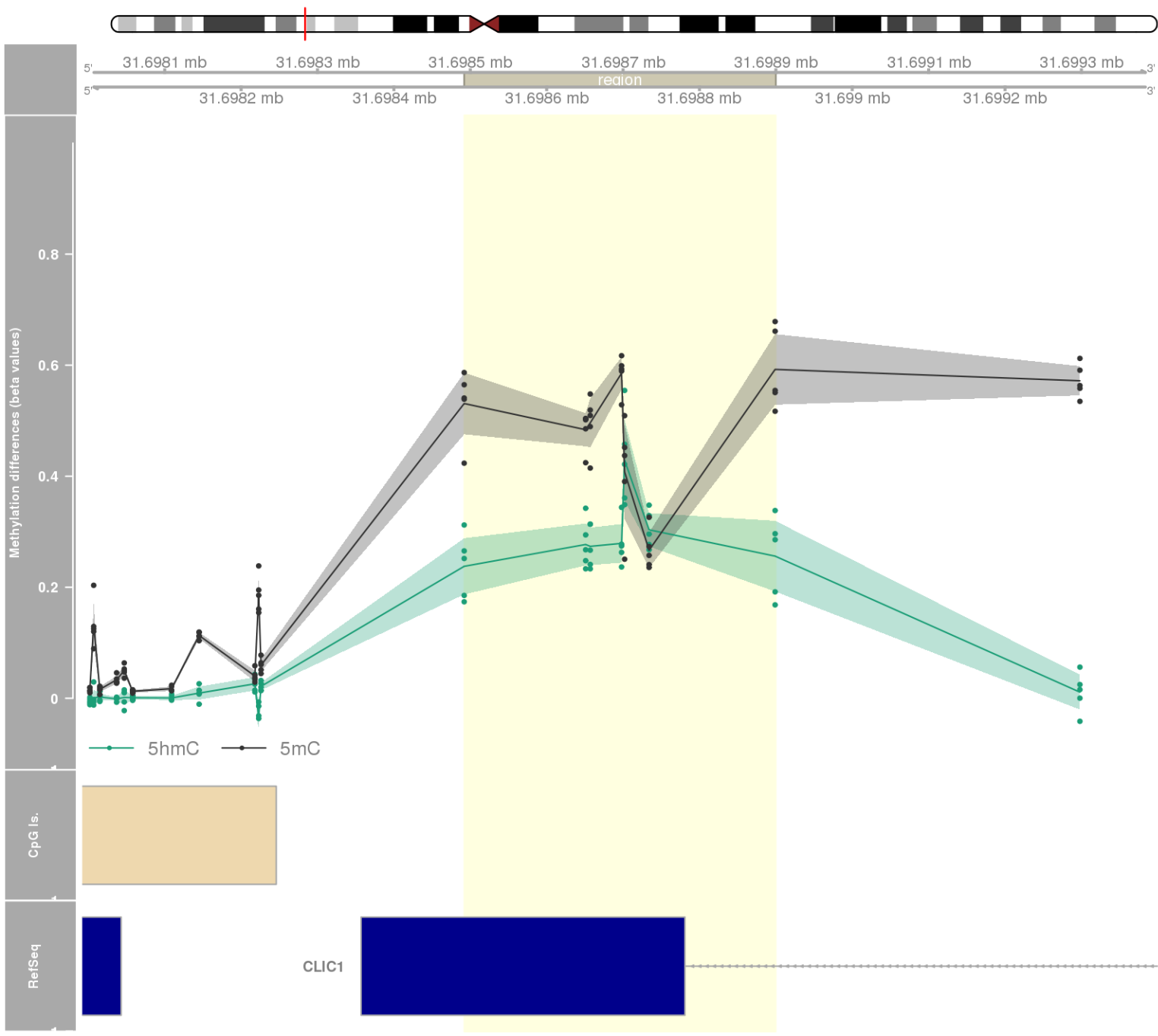
DMR 170 // chr5:148809664-148810203 // 539 pb. (8 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.012 // fwerArea: 1
- genes: MIR145 -



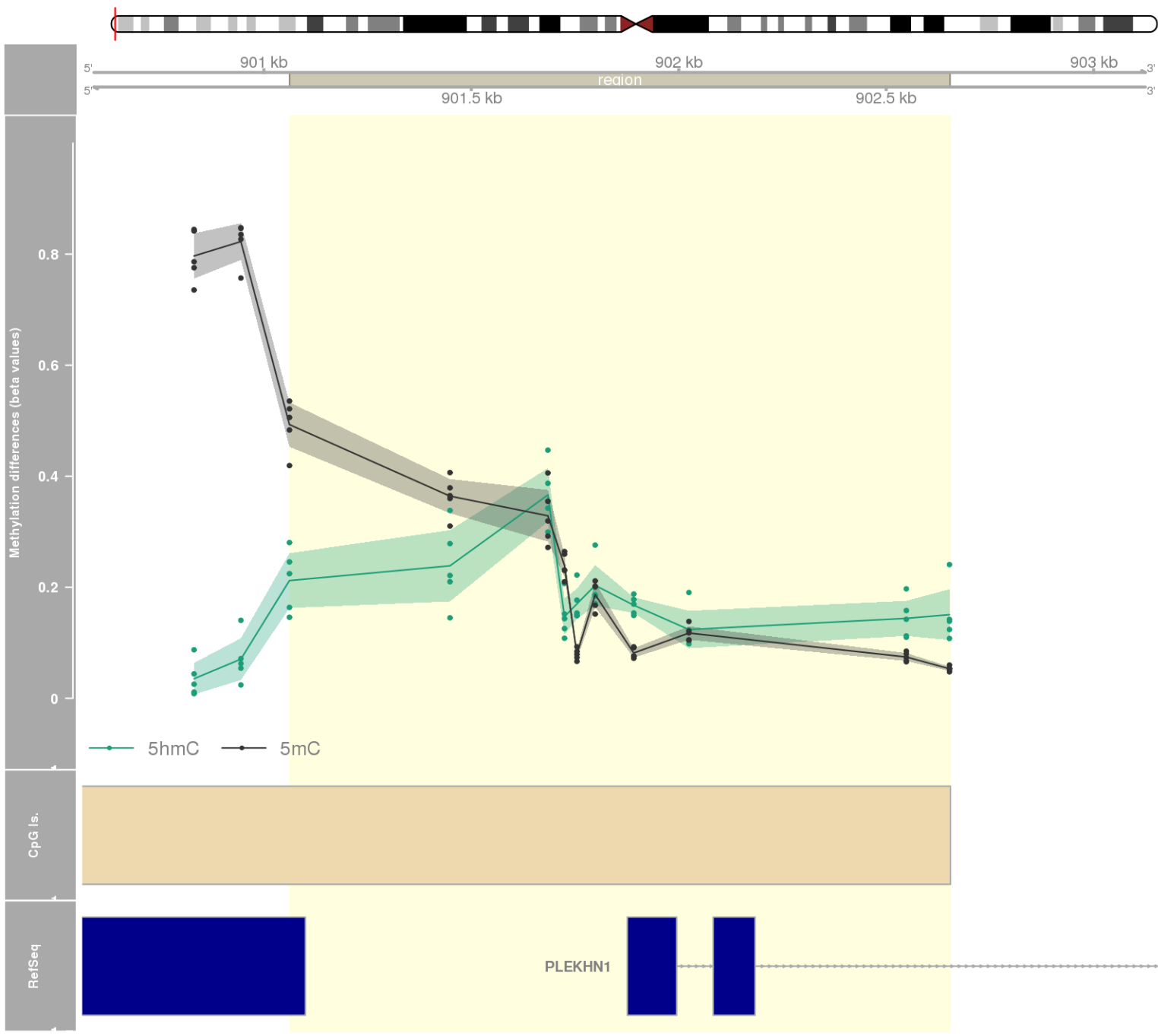
DMR 171 // chr7:6203156-6204983 // 1827 pb. (10 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.012 // fwerArea: 1
- genes: CYTH3 -



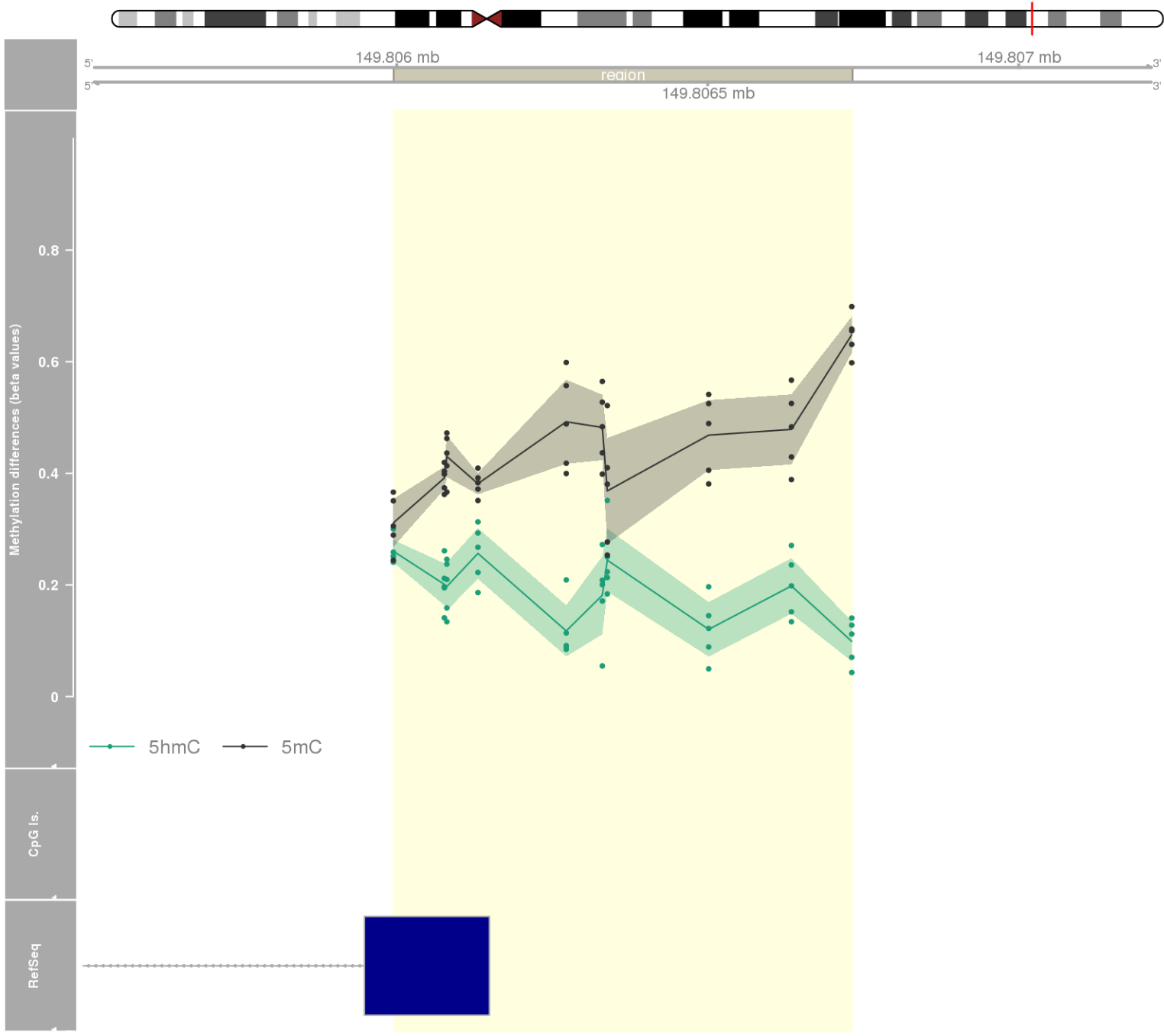
DMR 172 // chr6:31698492-31698899 // 407 pb. (7 probes) // pvalue: 0 // fwer: 0.173 // pvalueArea: 0.012 // fwerArea: 1
- genes: CLIC1 -



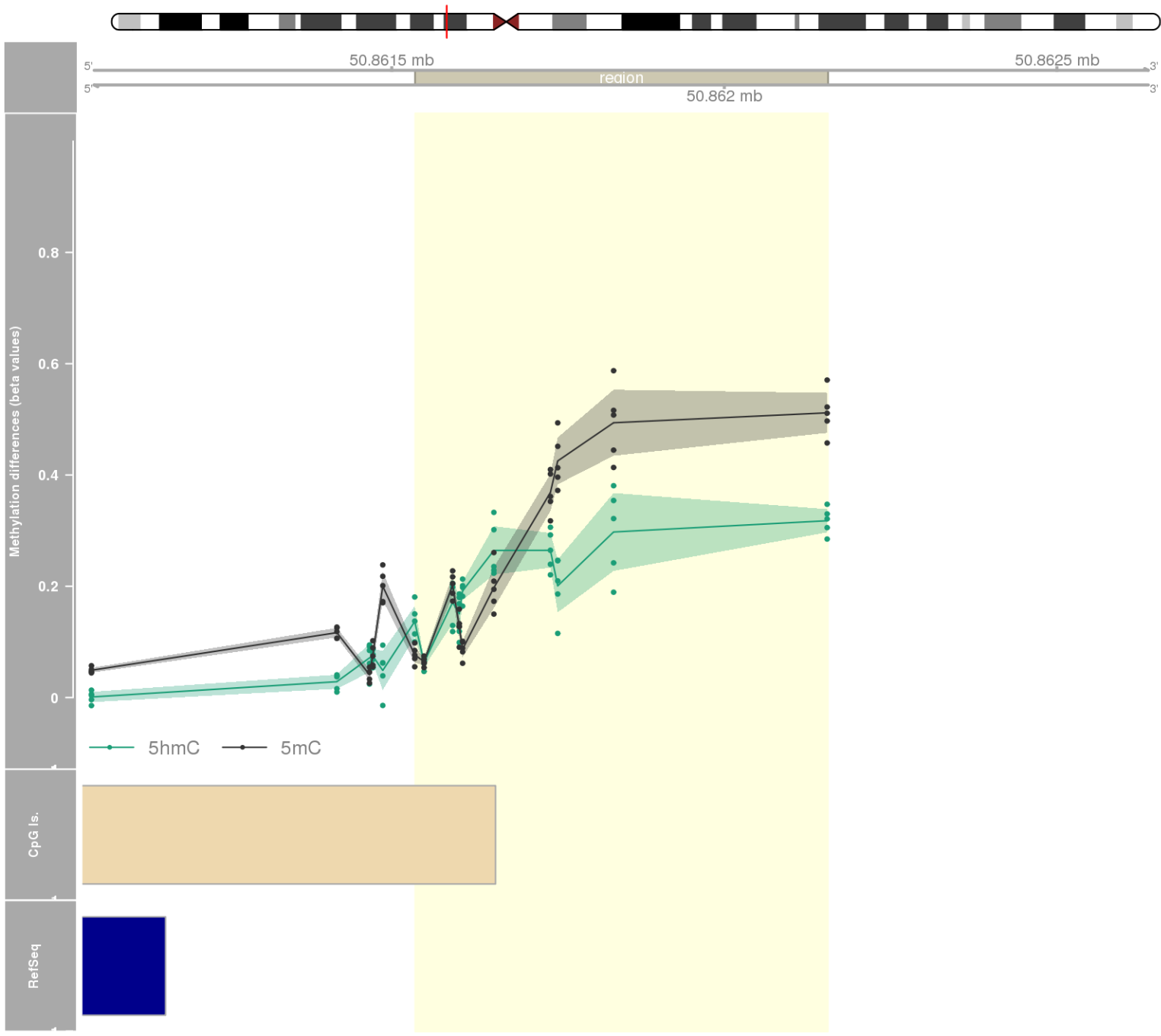
DMR 173 // chr1:901062-902653 // 1591 pb. (10 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.012 // fwerArea: 1
- genes: KLHL17 / PLEKHN1 -



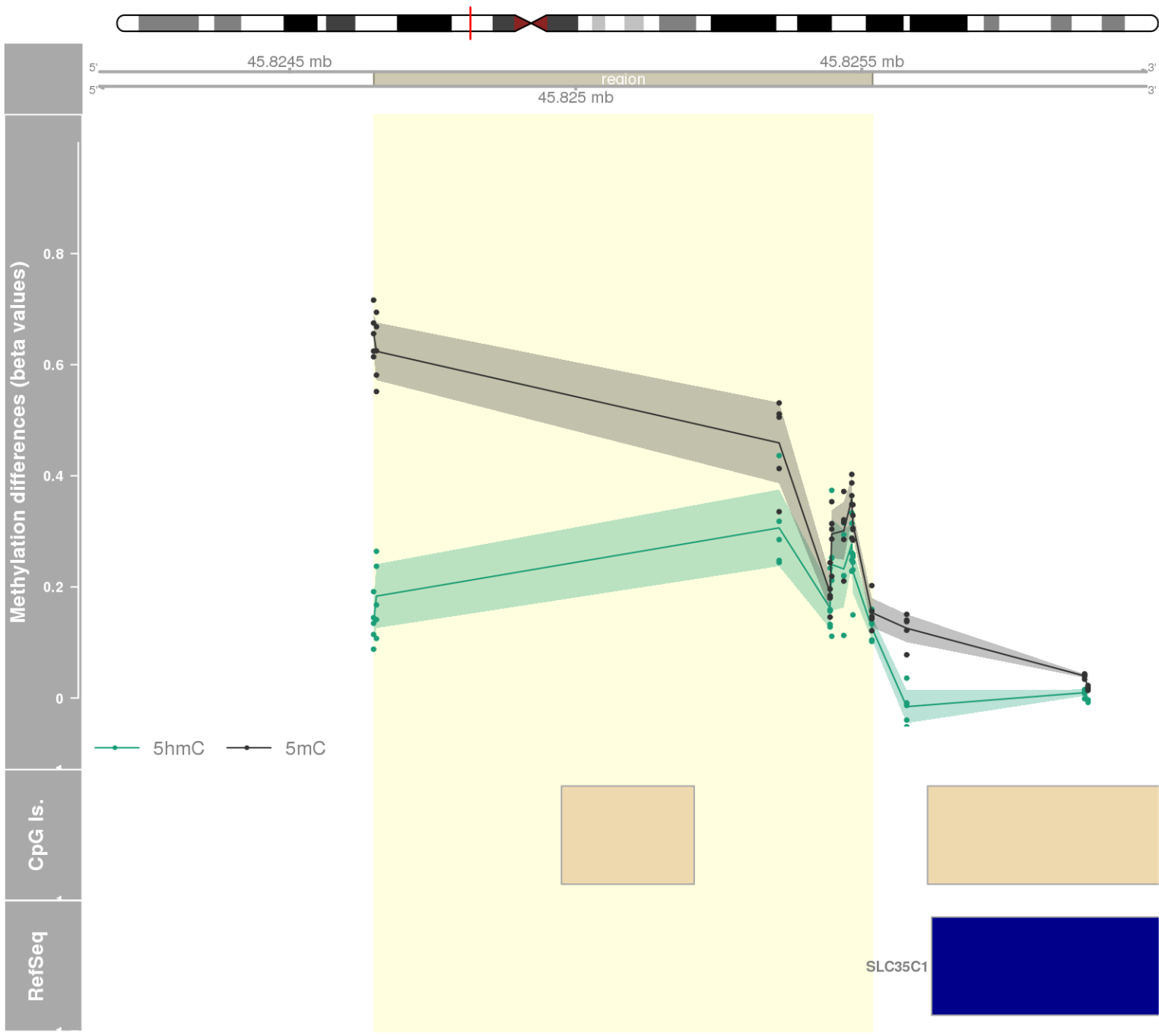
DMR 174 // chr6:149805995-149806732 // 737 pb. (10 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.012 // fwerArea: 1
- genes: ZC3H12D -



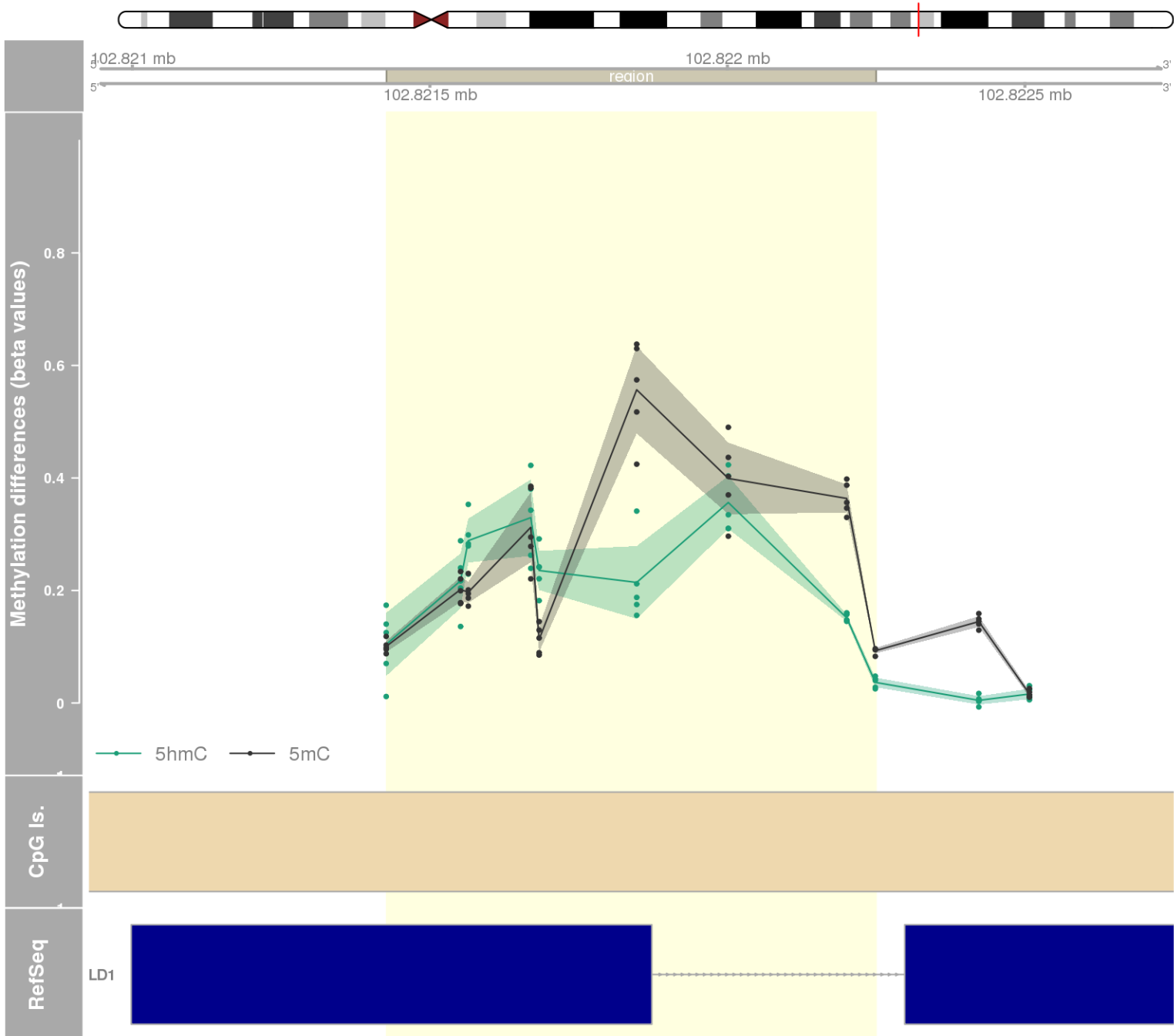
DMR 175 // chr7:50861535-50862155 // 620 pb. (10 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.012 // fwerArea: 1



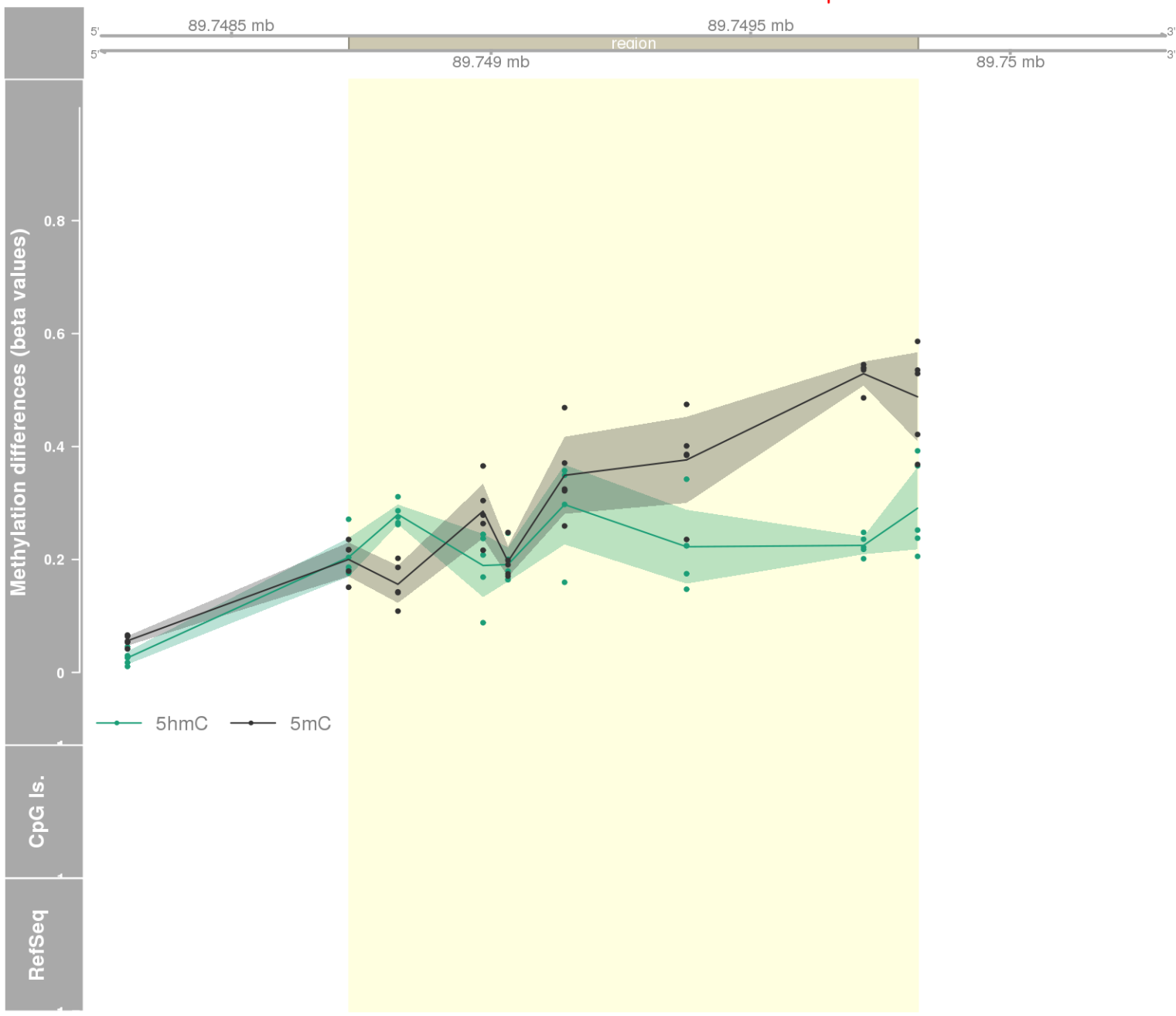
DMR 176 // chr11:45824647-45825518 // 871 pb. (9 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.013 // fwerArea: 1



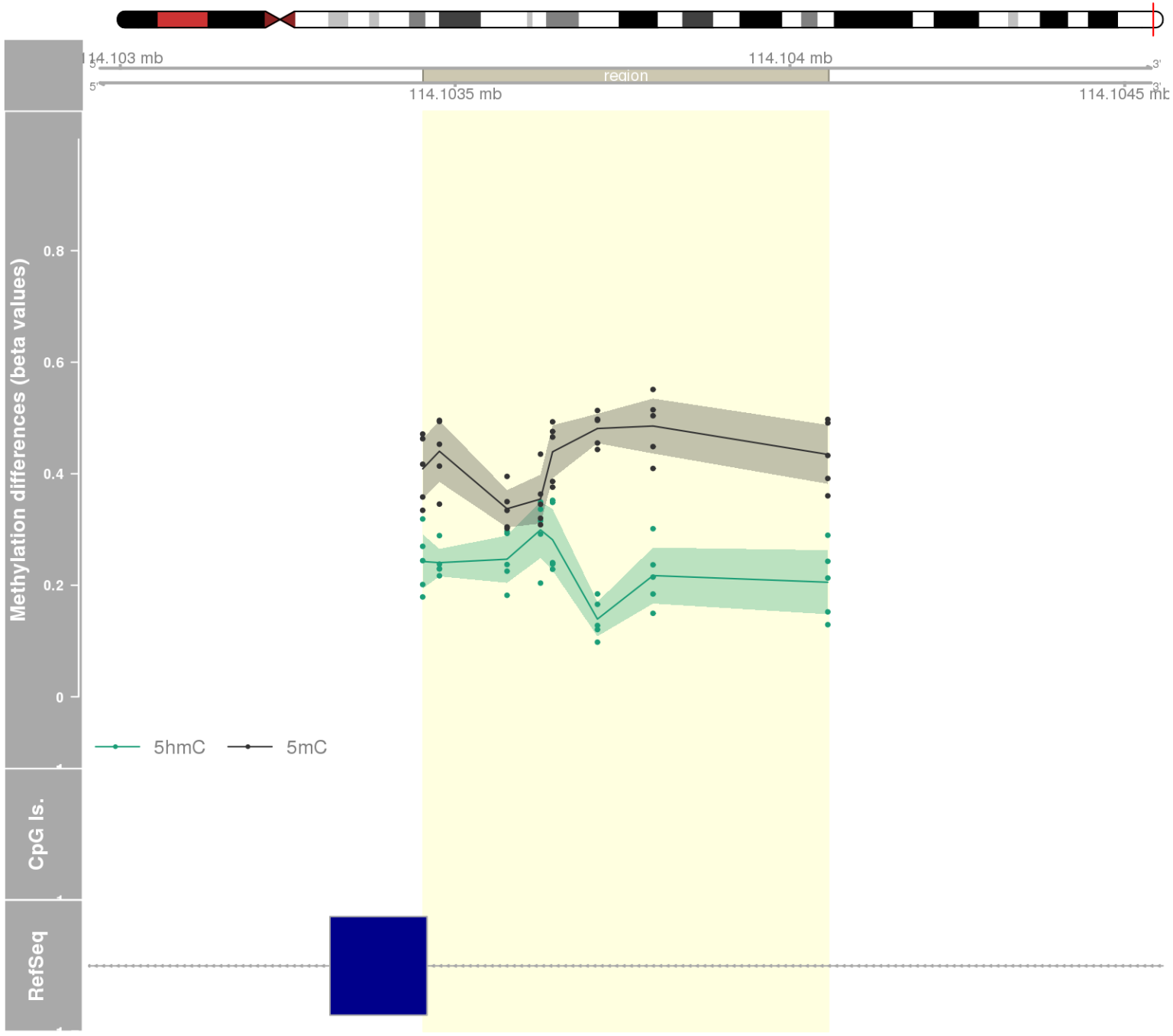
DMR 177 // chr10:102821427-102822249 // 822 pb. (9 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.013 // fwerArea: 1
- genes: KAZALD1 -



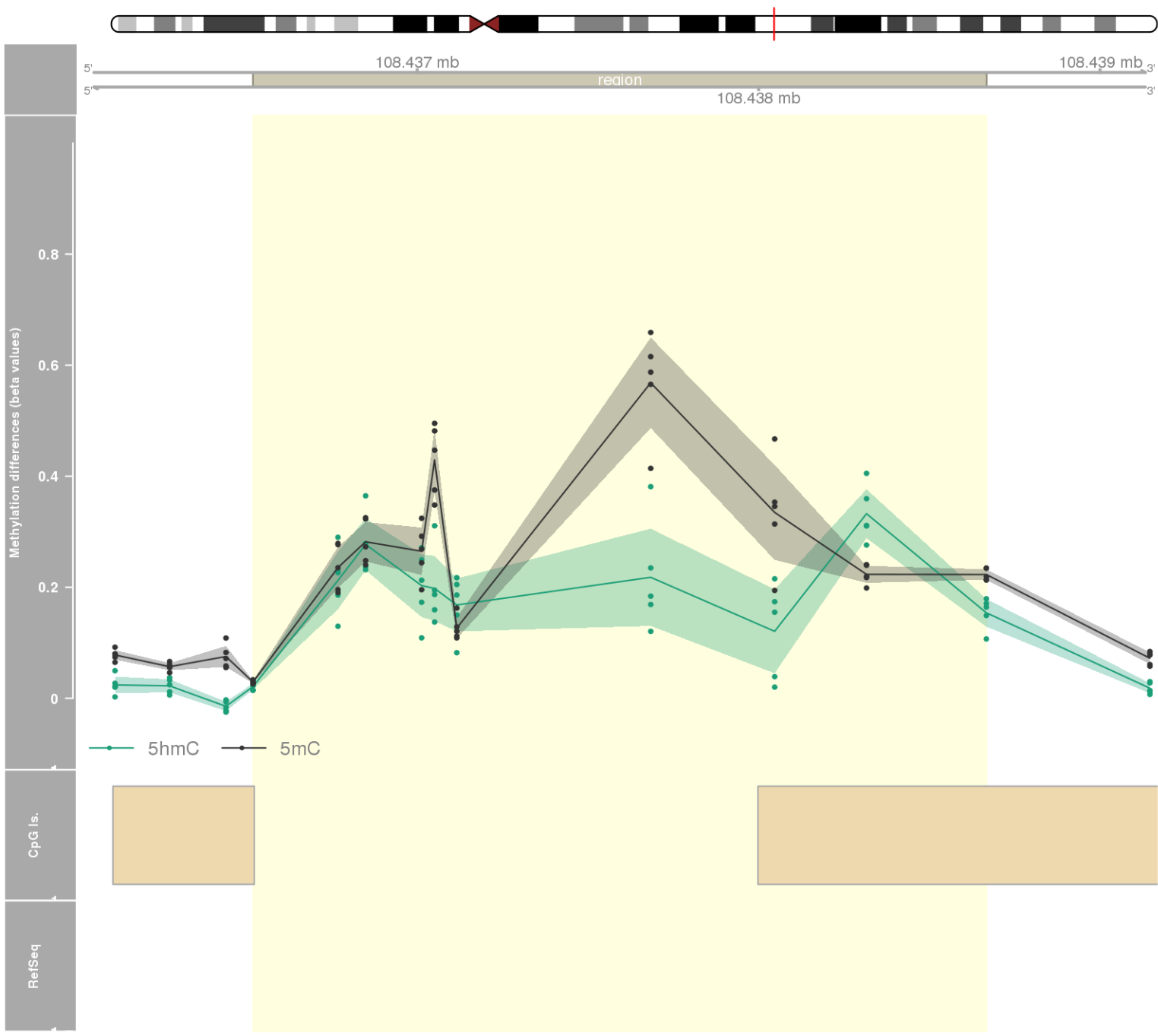
DMR 178 // chr12:89748726-89749822 // 1096 pb. (8 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.013 // fwerArea: 1



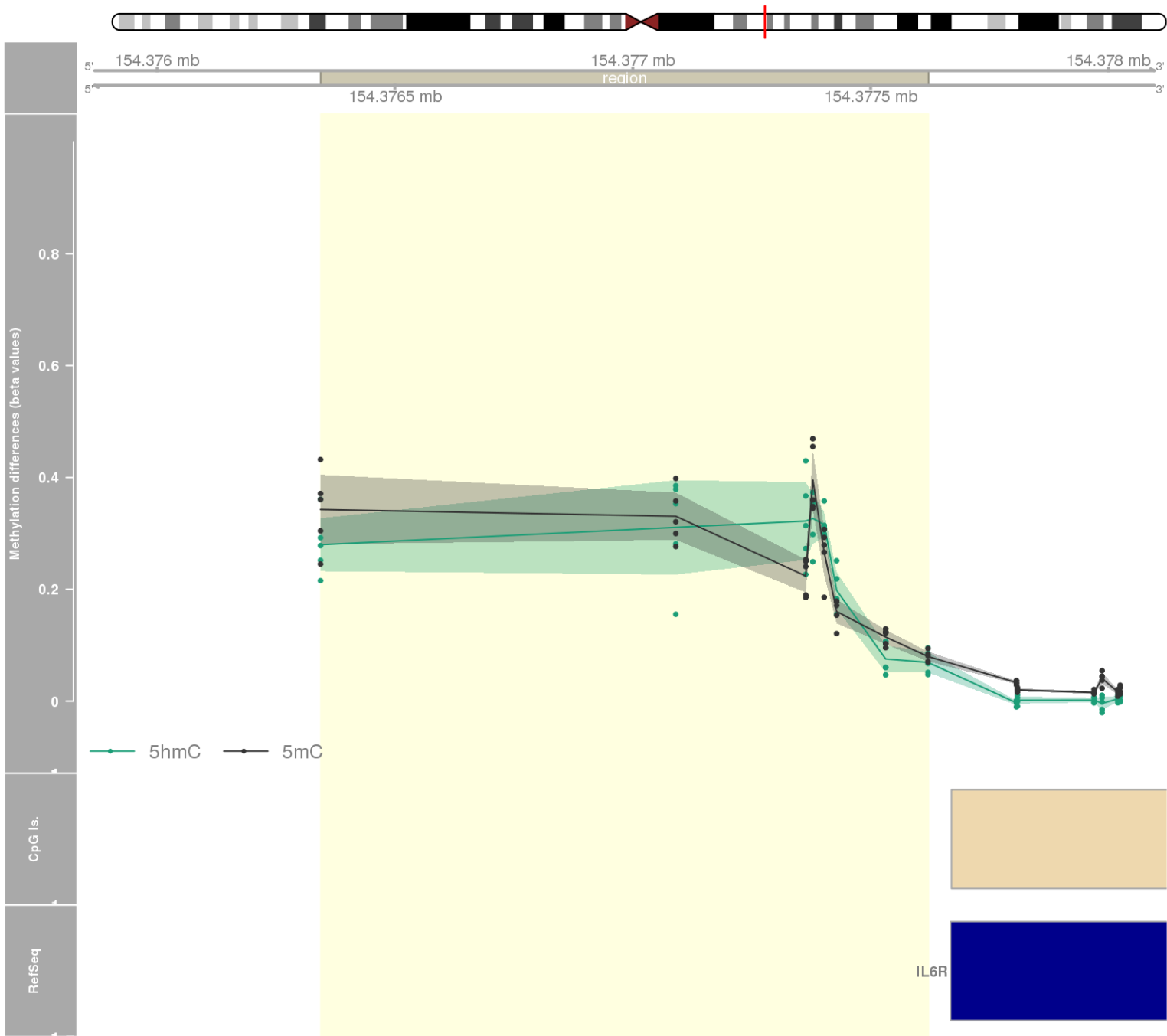
DMR 179 // chr13:114103452-114104057 // 605 pb. (8 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.013 // fwerArea: 1
- genes: ADPRHL1 -



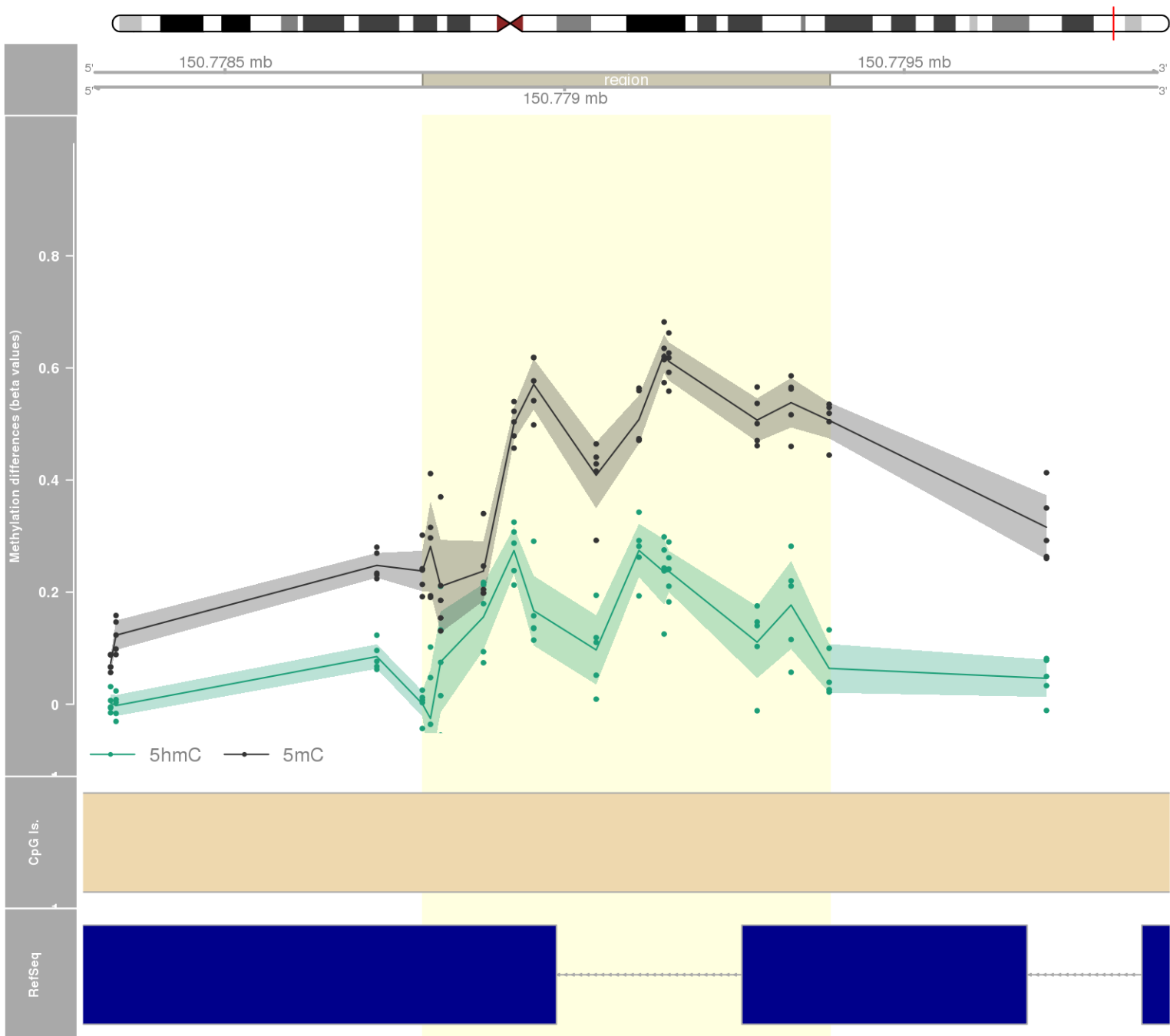
DMR 180 // chr6:108436519-108438667 // 2148 pb. (10 probes) // pvalue: 0.004 // fwer: 0.36 // pvalueArea: 0.013 // fwerArea: 1
- genes: OSTM1 -



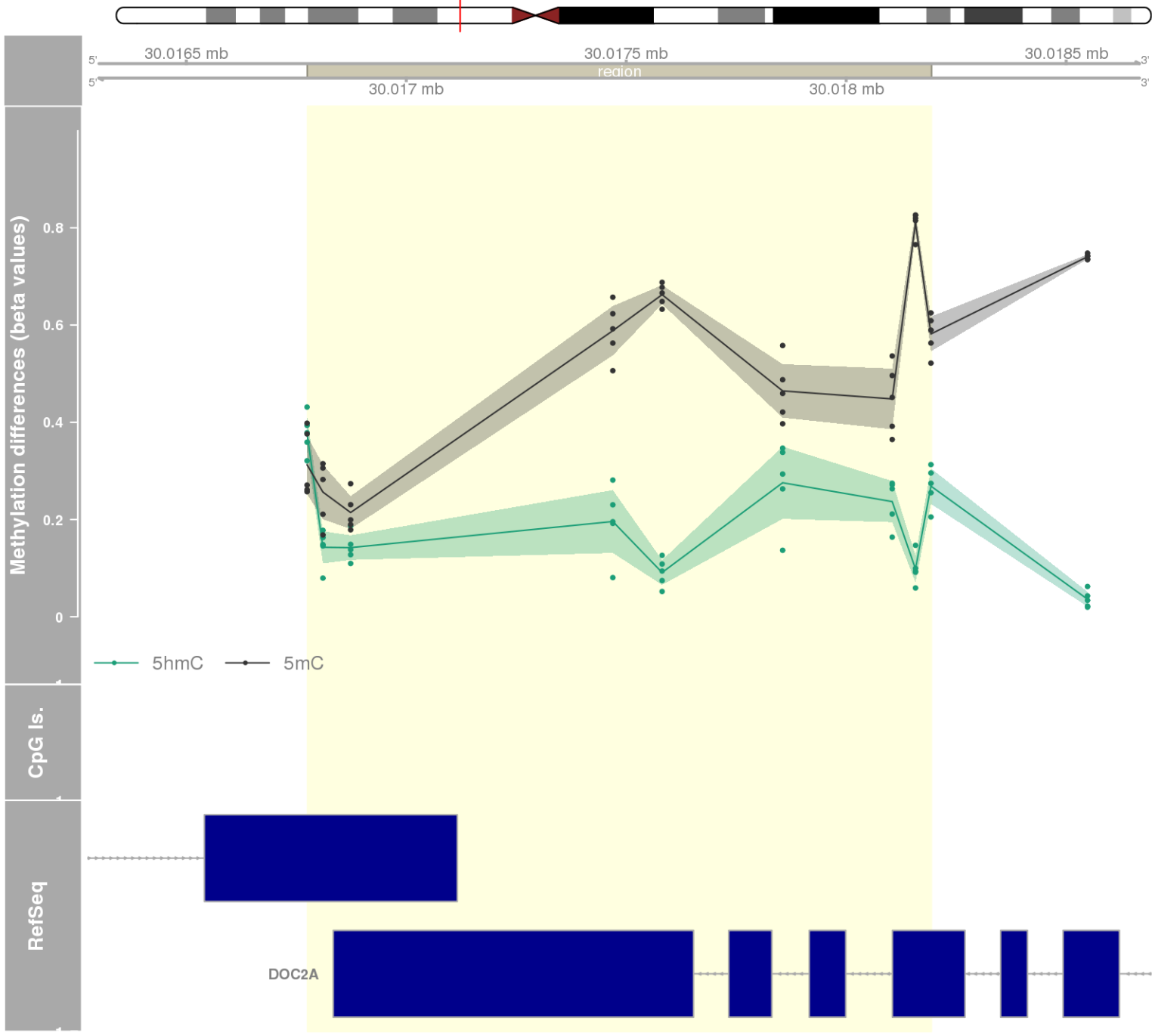
DMR 181 // chr1:154376344-154377621 // 1277 pb. (8 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.013 // fwerArea: 1



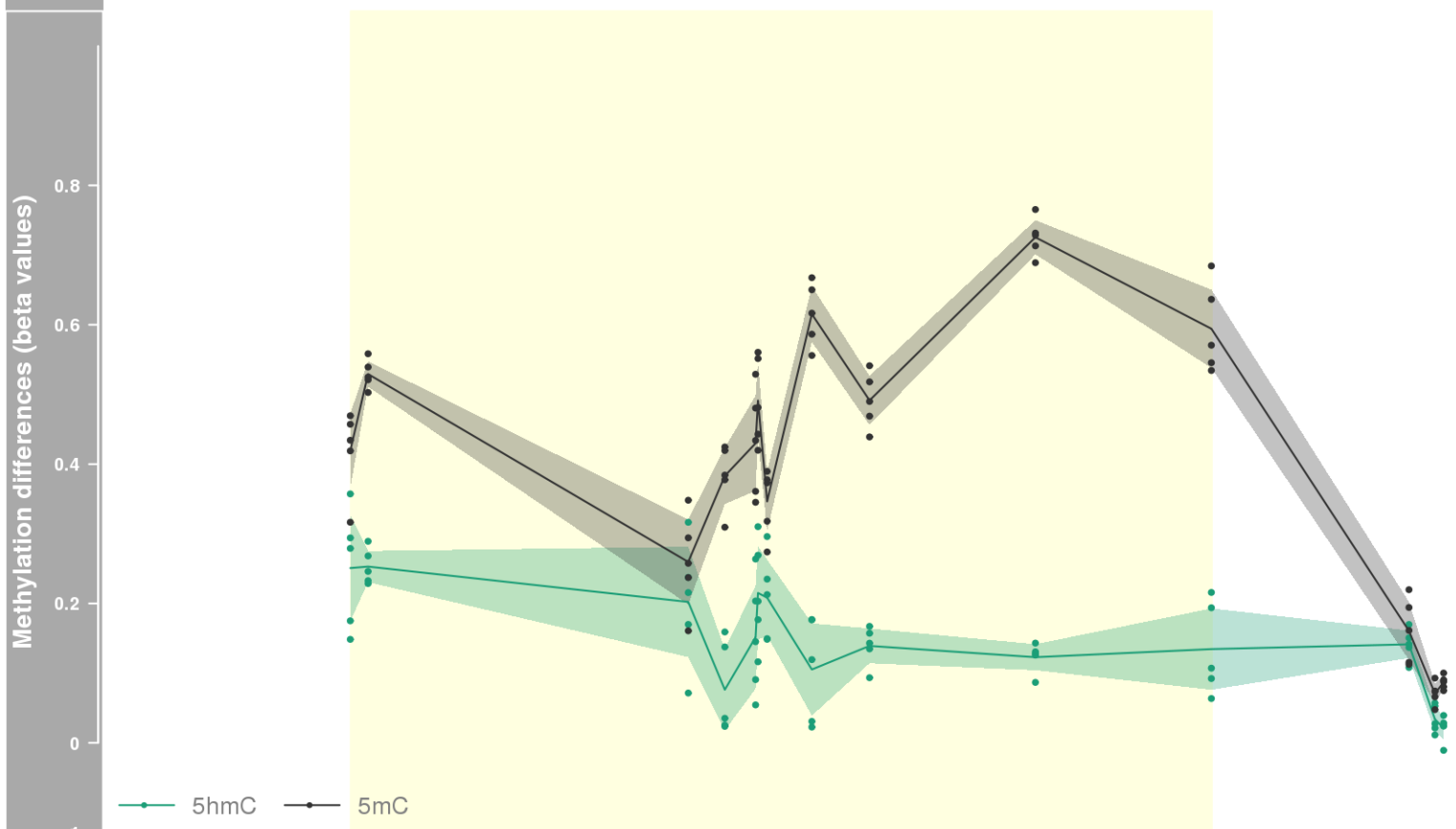
DMR 182 // chr7:150778791-150779390 // 599 pb. (13 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.013 // fwerArea: 1
- genes: TMUB1 -



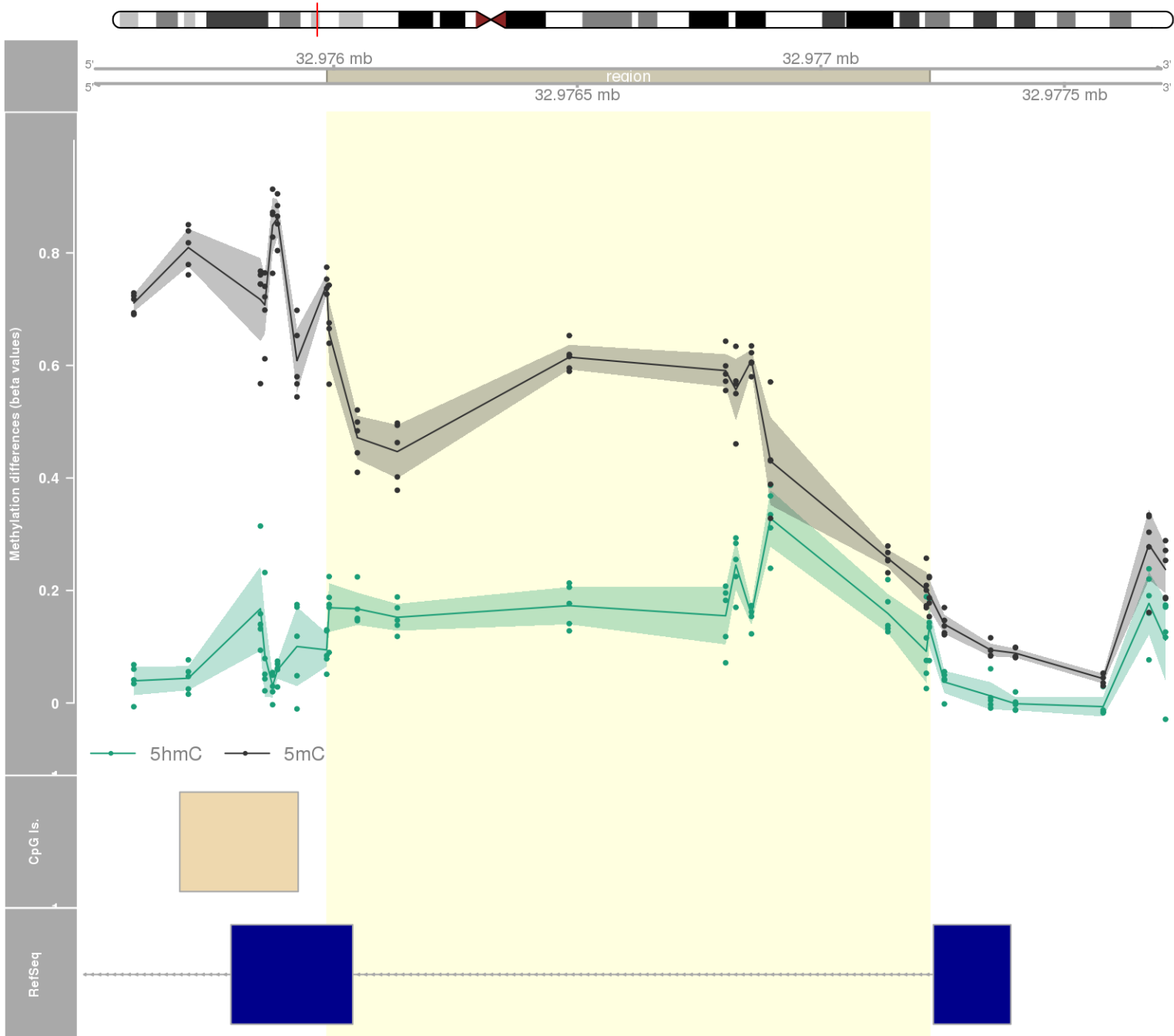
DMR 183 // chr16:30016775-30018193 // 1418 pb. (9 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.013 // fwerArea: 1
- genes: INO80E / DOC2A -



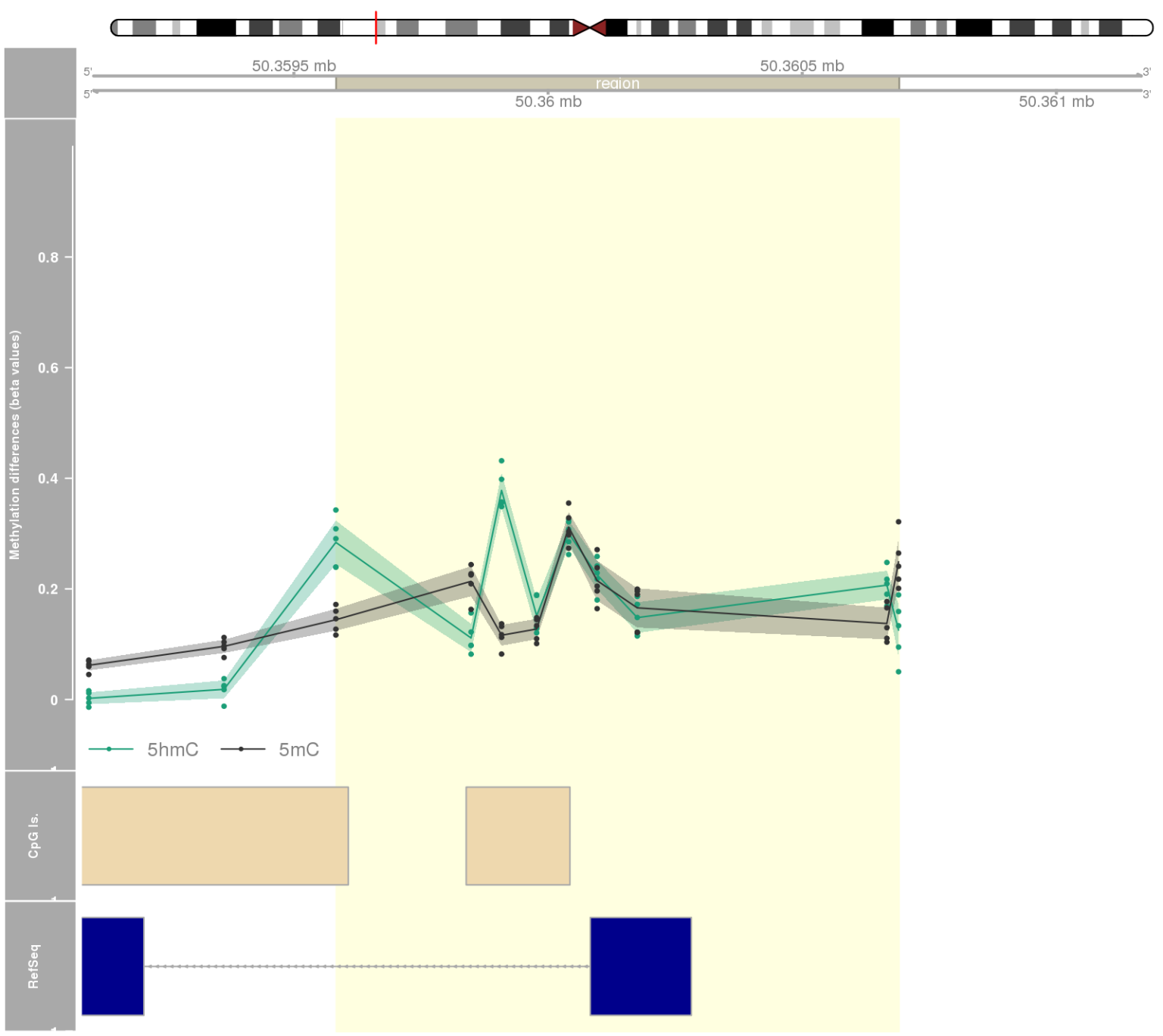
DMR 184 // chr11:73356316-73358107 // 1791 pb. (11 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.013 // fwerArea: 1
- genes: PLEKHB1 -



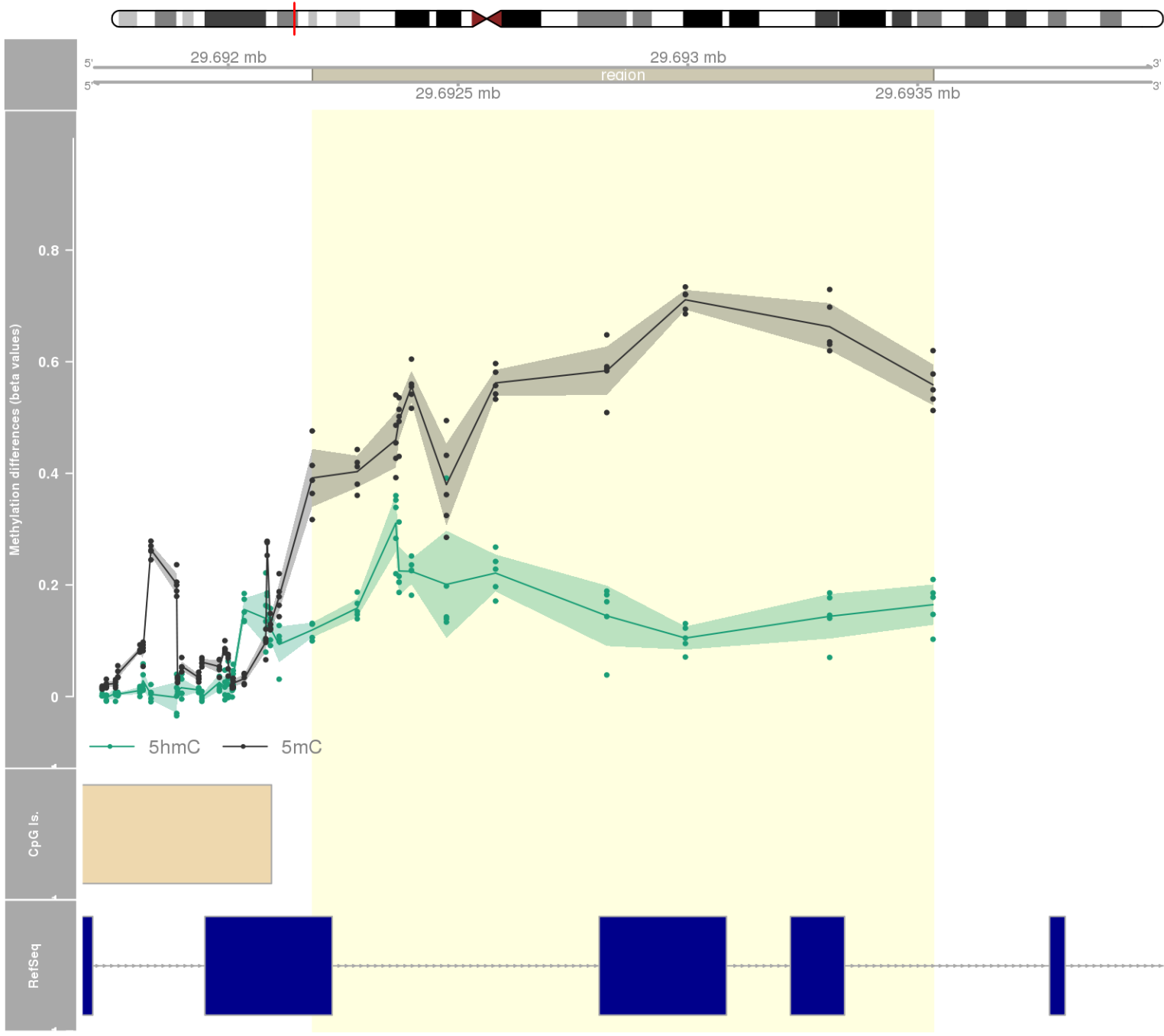
DMR 185 // chr6:32975986-32977223 // 1237 pb. (12 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.013 // fwerArea: 1
- genes: HLA-DOA -



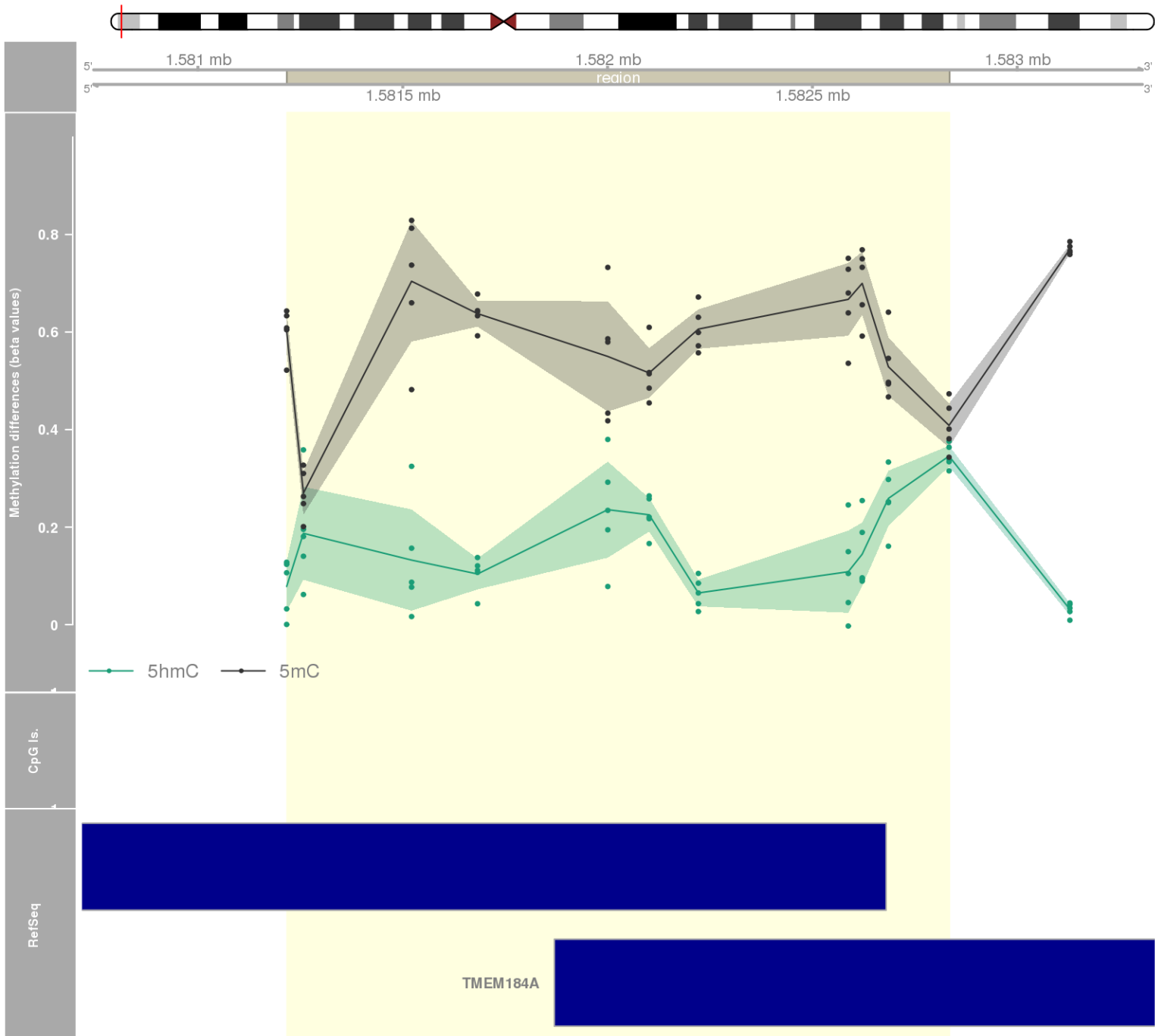
DMR 186 // chr3:50359583-50360690 // 1107 pb. (9 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.013 // fwerArea: 1
- genes: HYAL2 / TUSC2 -



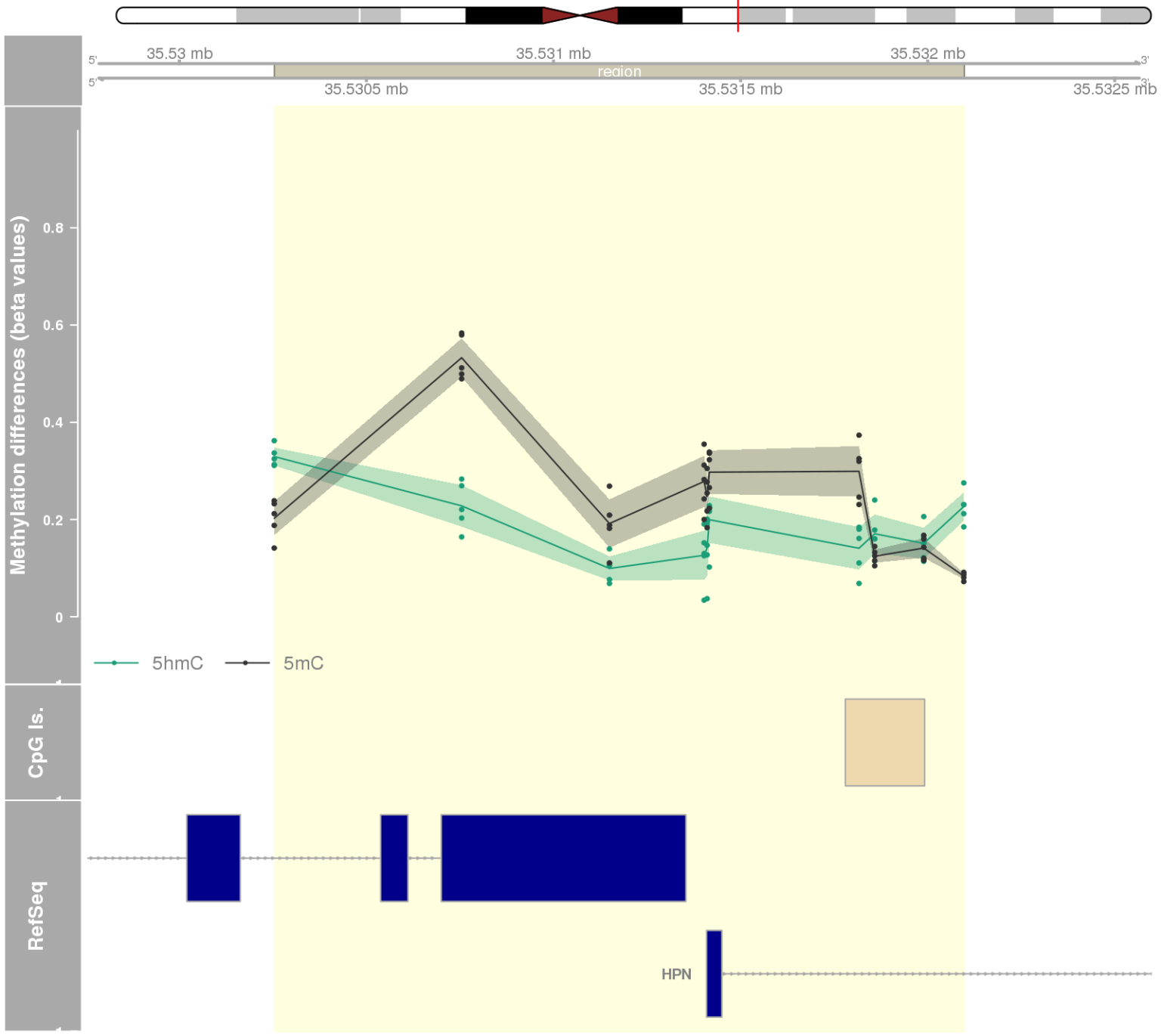
DMR 187 // chr6:29692183-29693534 // 1351 pb. (11 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.013 // fwerArea: 1
- genes: HLA-F -



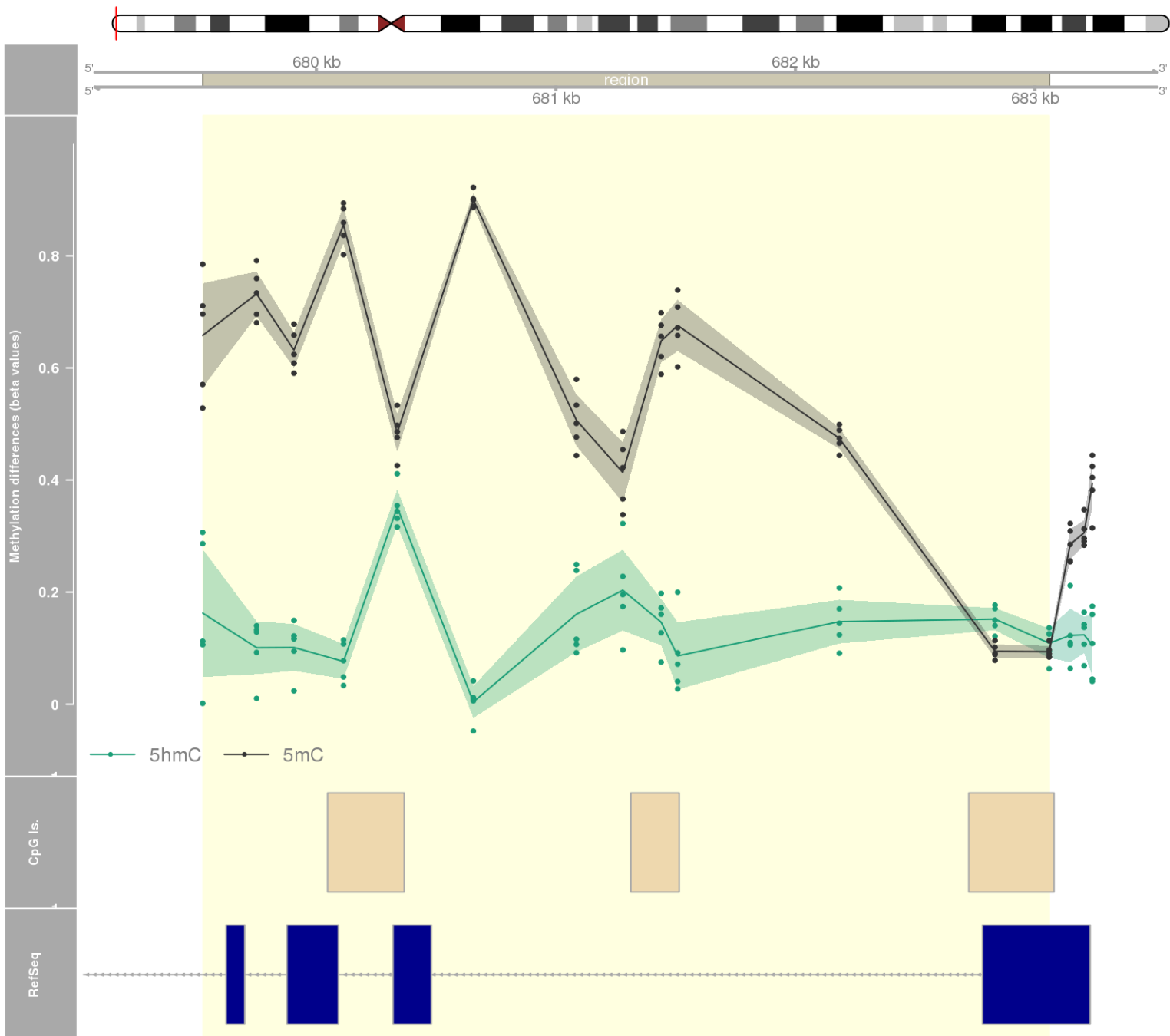
DMR 188 // chr7:1581217-1582834 // 1617 pb. (11 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.013 // fwerArea: 1
- genes: MAFK / TMEM184A -



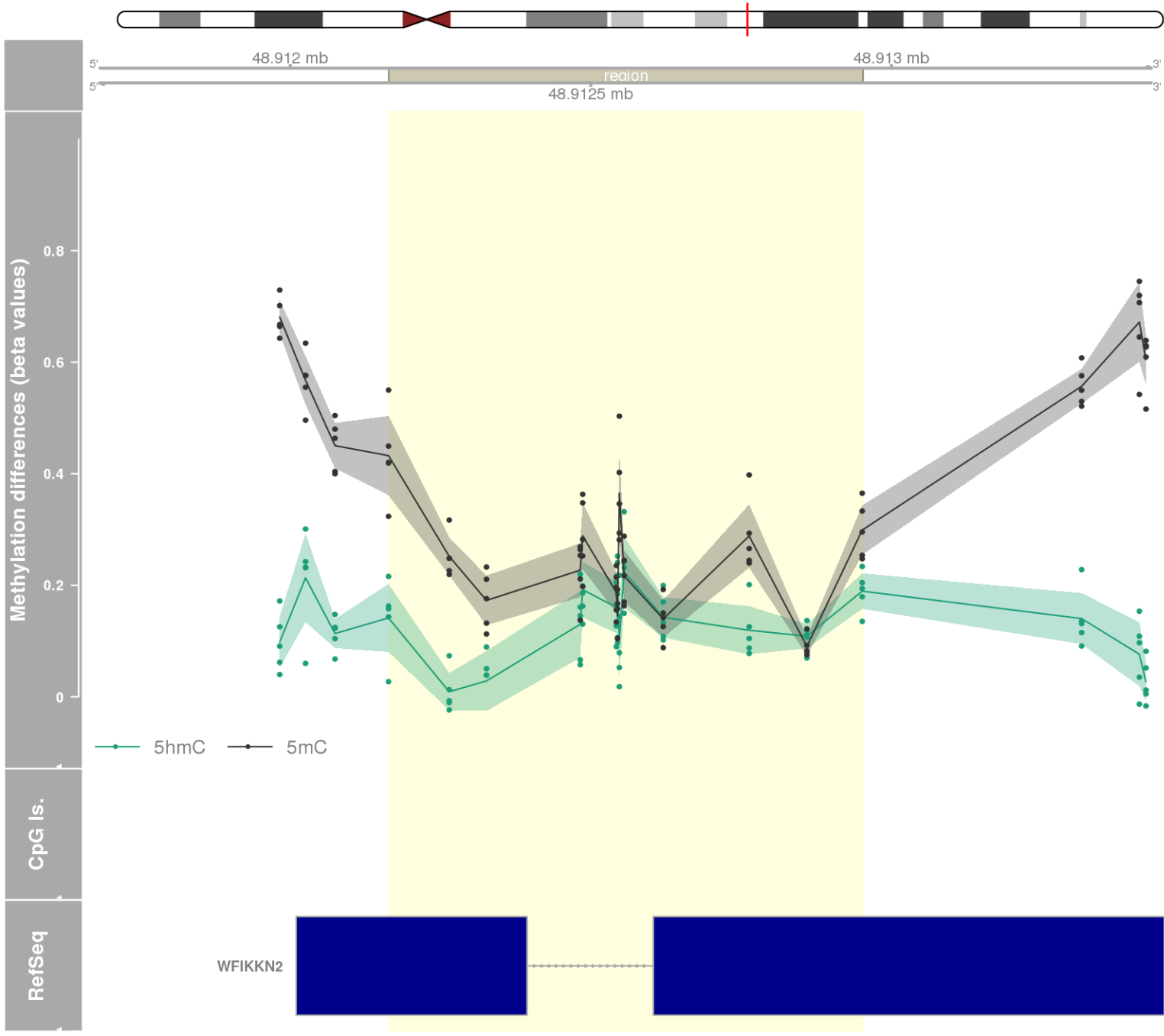
DMR 189 // chr19:35530254-35532097 // 1843 pb. (10 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.013 // fwerArea: 1
- genes: SCN1B / HPN -



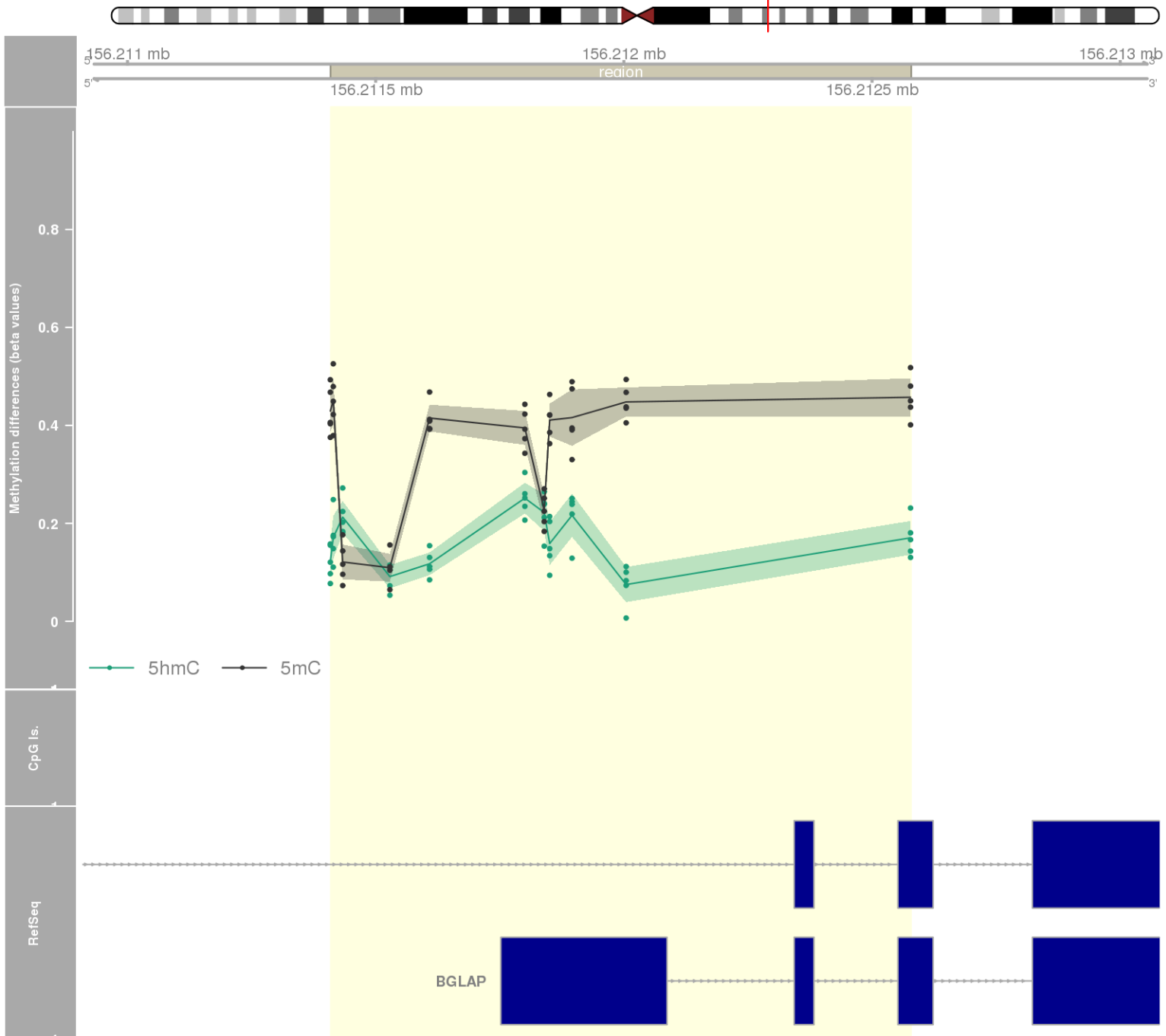
DMR 190 // chr4:679526-683060 // 3534 pb. (13 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.013 // fwerArea: 1
- genes: MFSD7 -



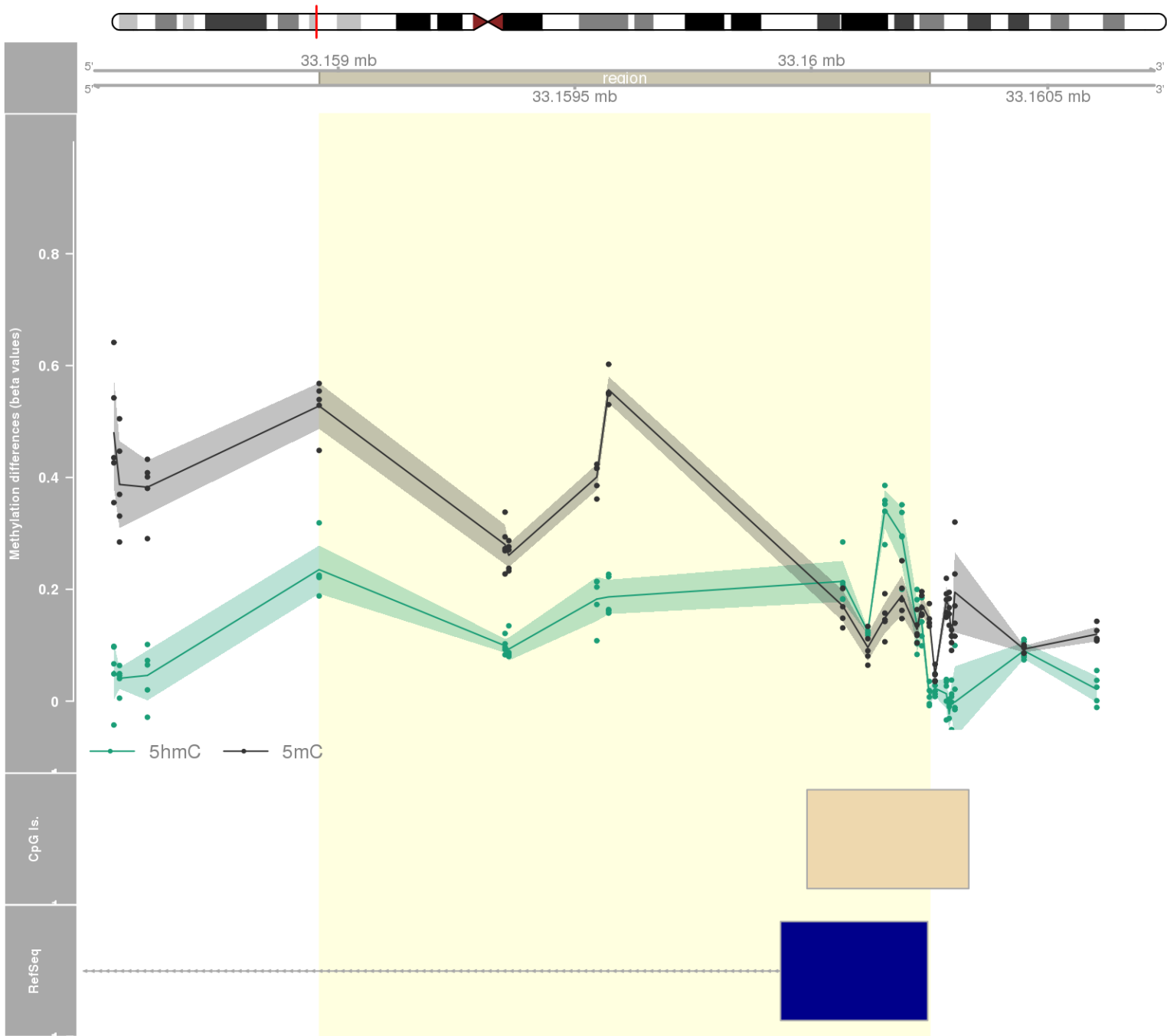
DMR 191 // chr17:48912164-48912952 // 788 pb. (13 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.014 // fwerArea: 1
- genes: WFIKKN2 -



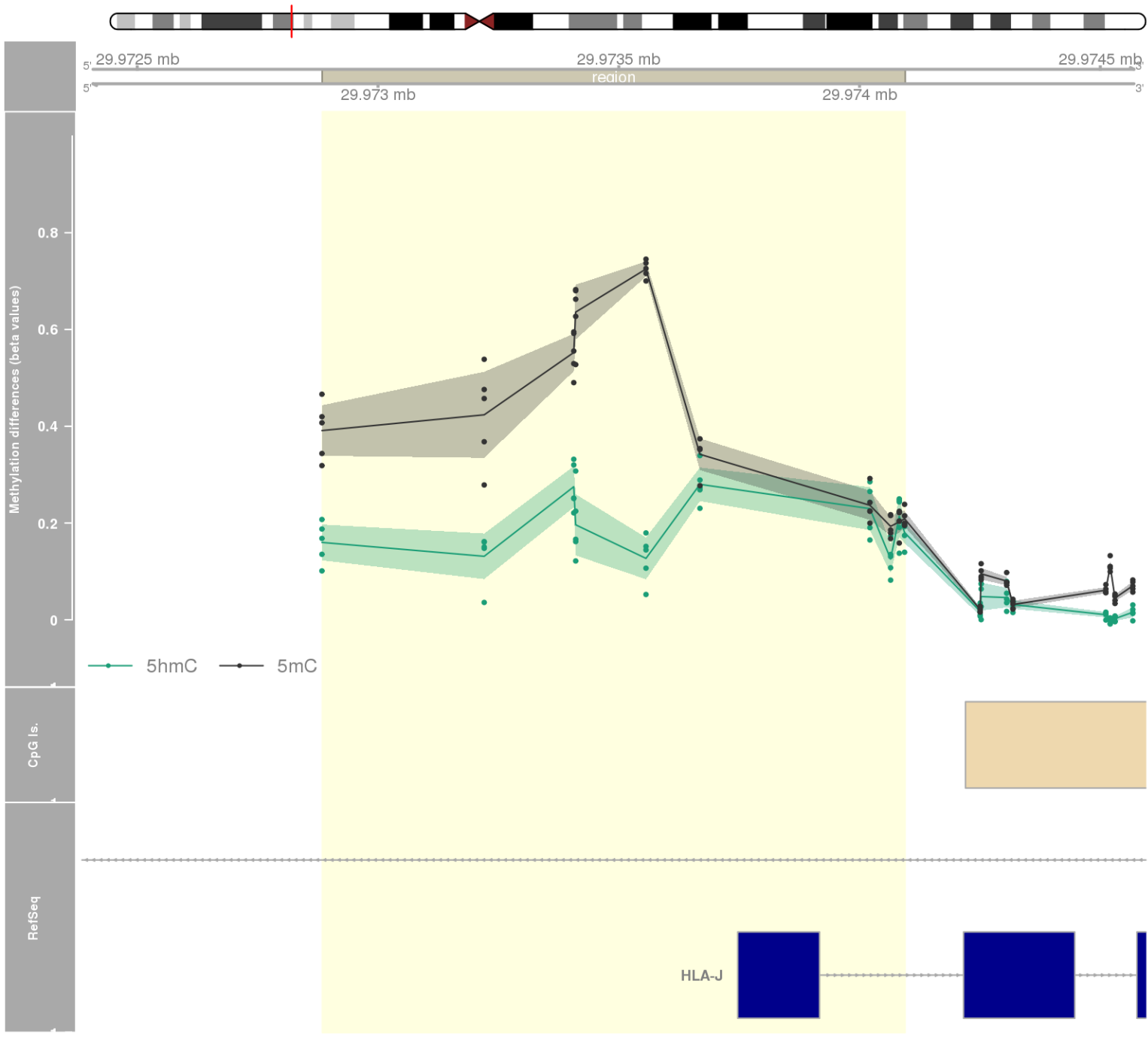
DMR 192 // chr1:156211409-156212578 // 1169 pb. (11 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.014 // fwerArea: 1
- genes: PMF1-BGLAP / PMF1 / BGLAP -



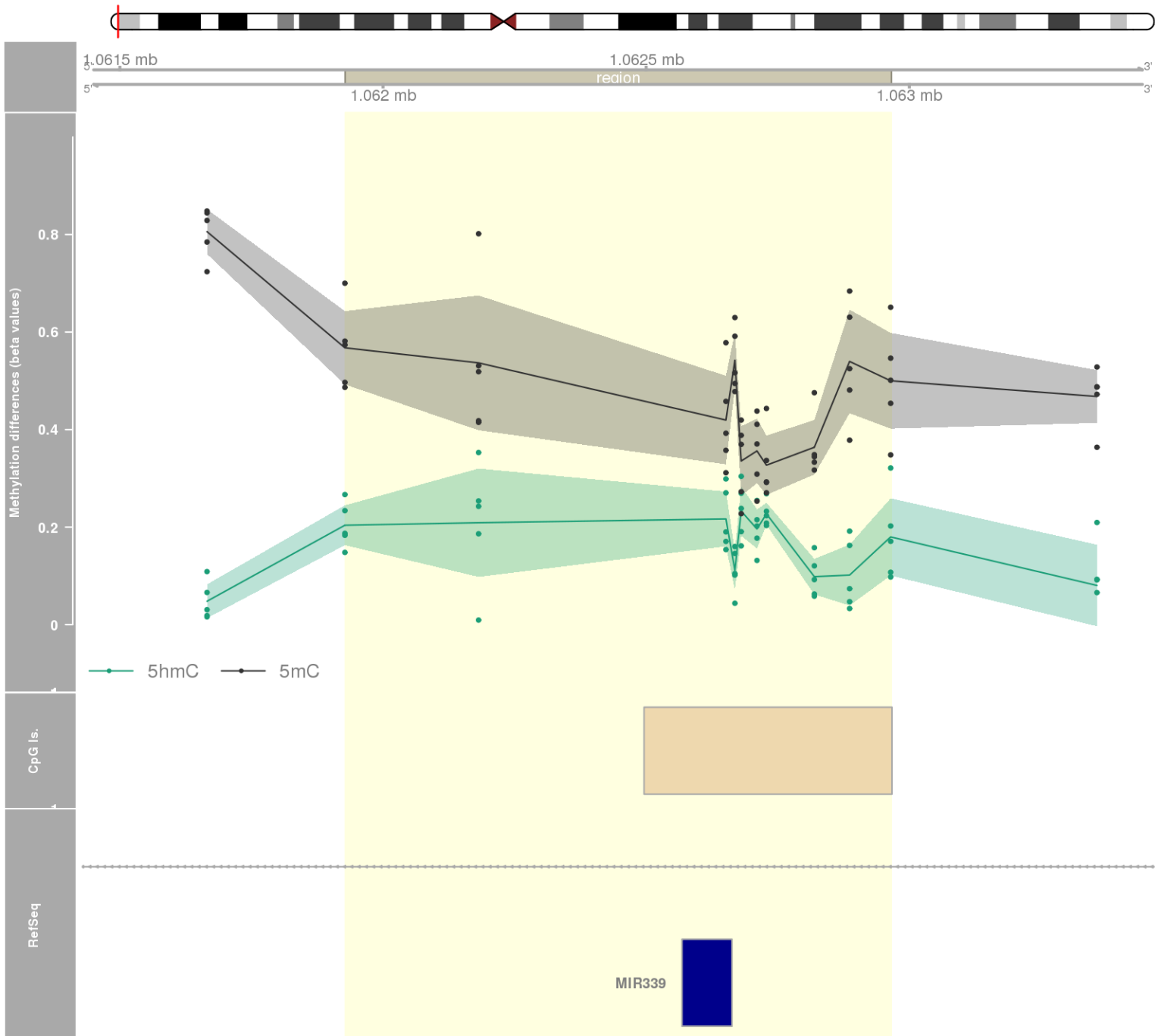
DMR 193 // chr6:33158960-33160250 // 1290 pb. (12 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.014 // fwerArea: 1
- genes: COL11A2 -



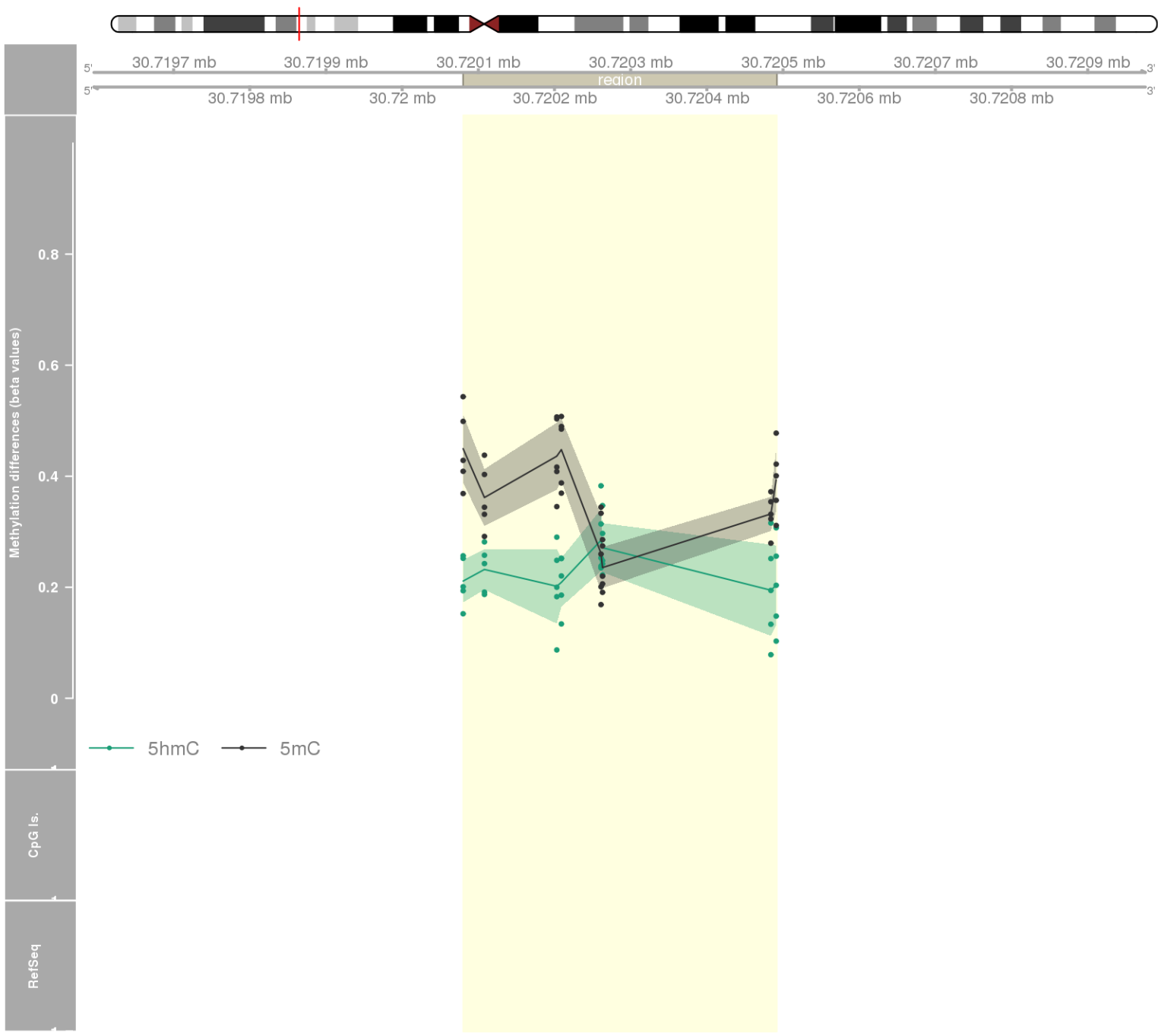
DMR 194 // chr6:29972884-29974094 // 1210 pb. (10 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.014 // fwerArea: 1
- genes: ZNRD1-AS1 / HCG4P3 -



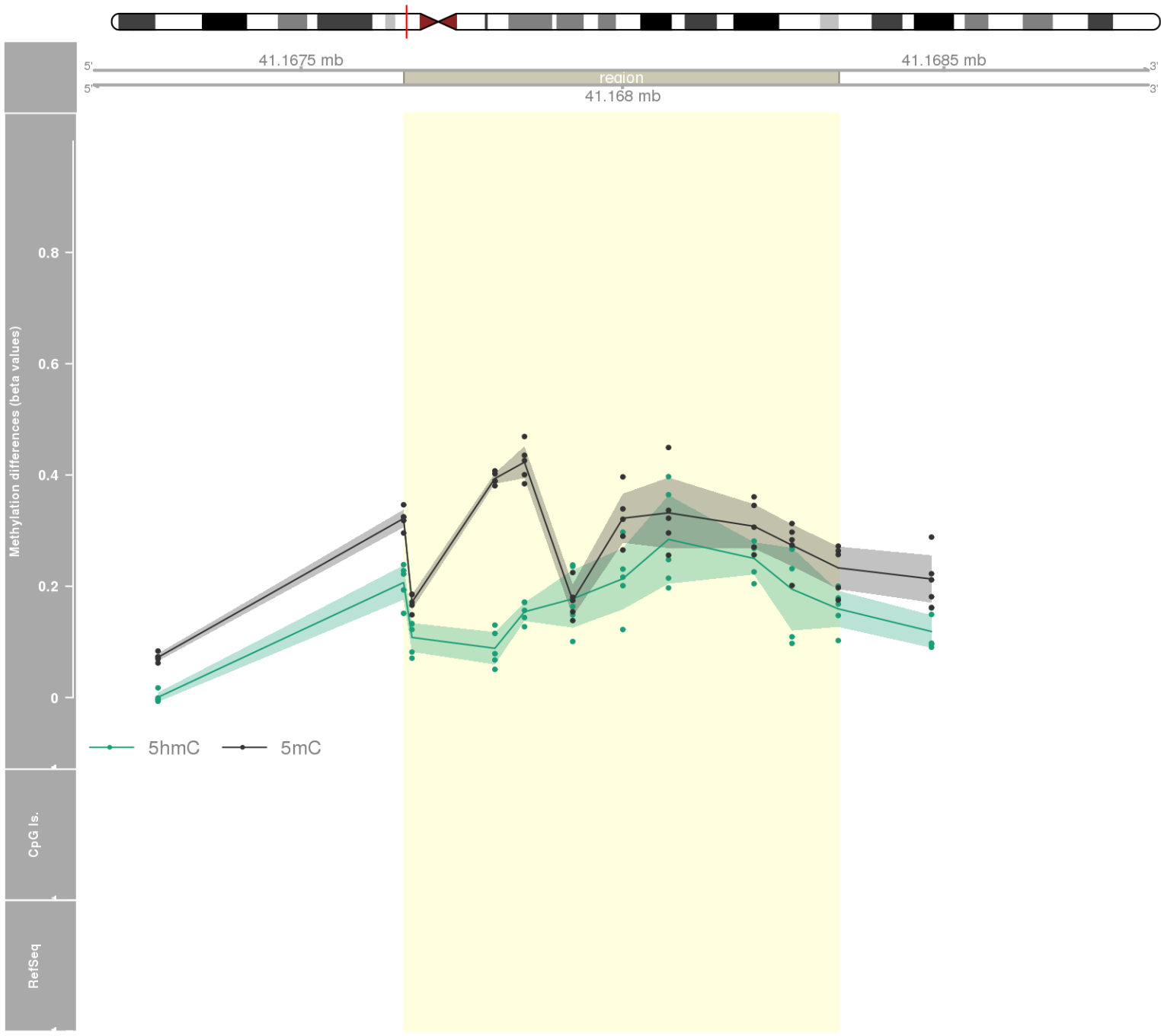
DMR 195 // chr7:1061928-1062965 // 1037 pb. (10 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.014 // fwerArea: 1
- genes: C7orf50 / MIR339 -



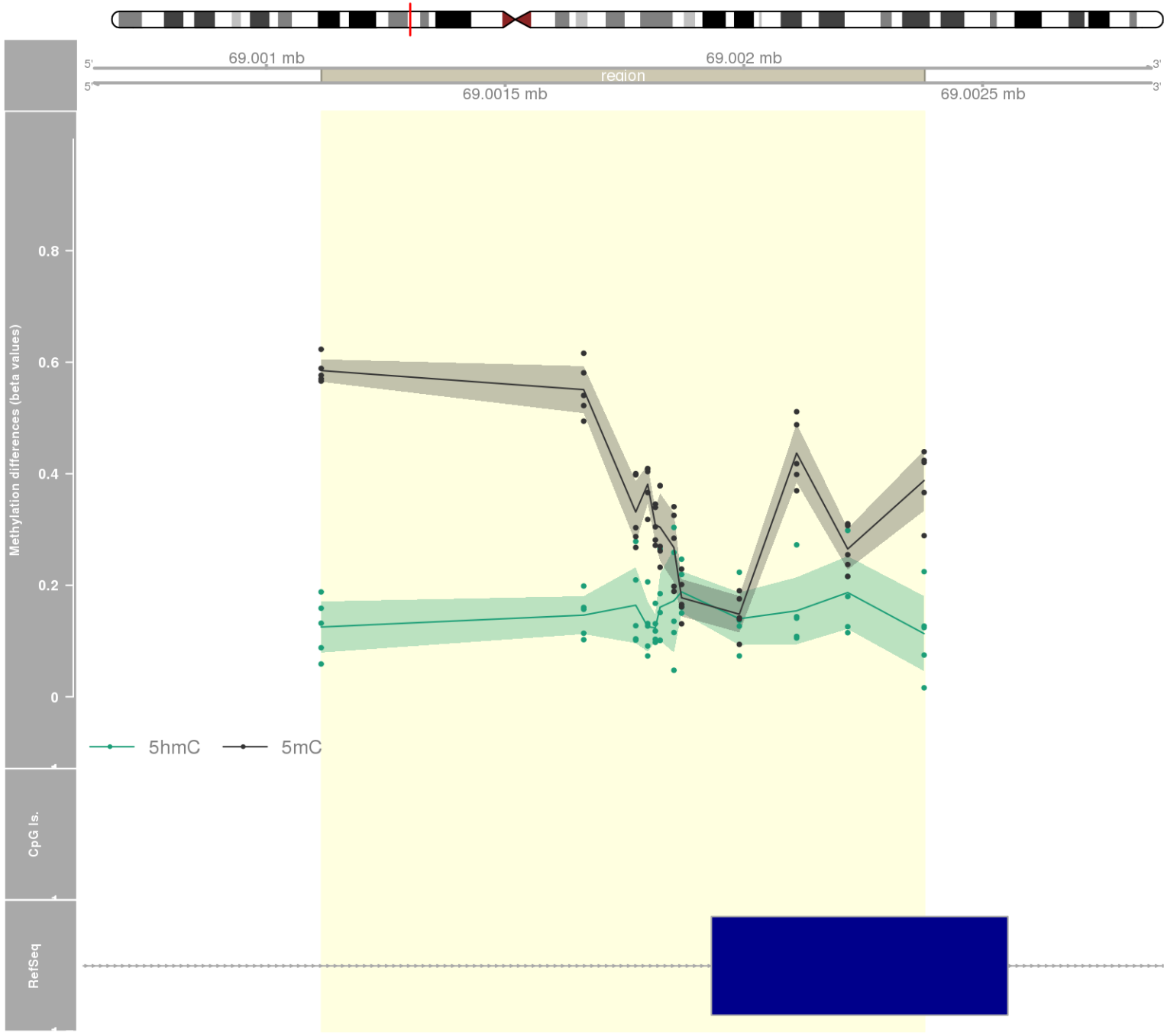
DMR 196 // chr6:30720080-30720491 // 411 pb. (8 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.014 // fwerArea: 1



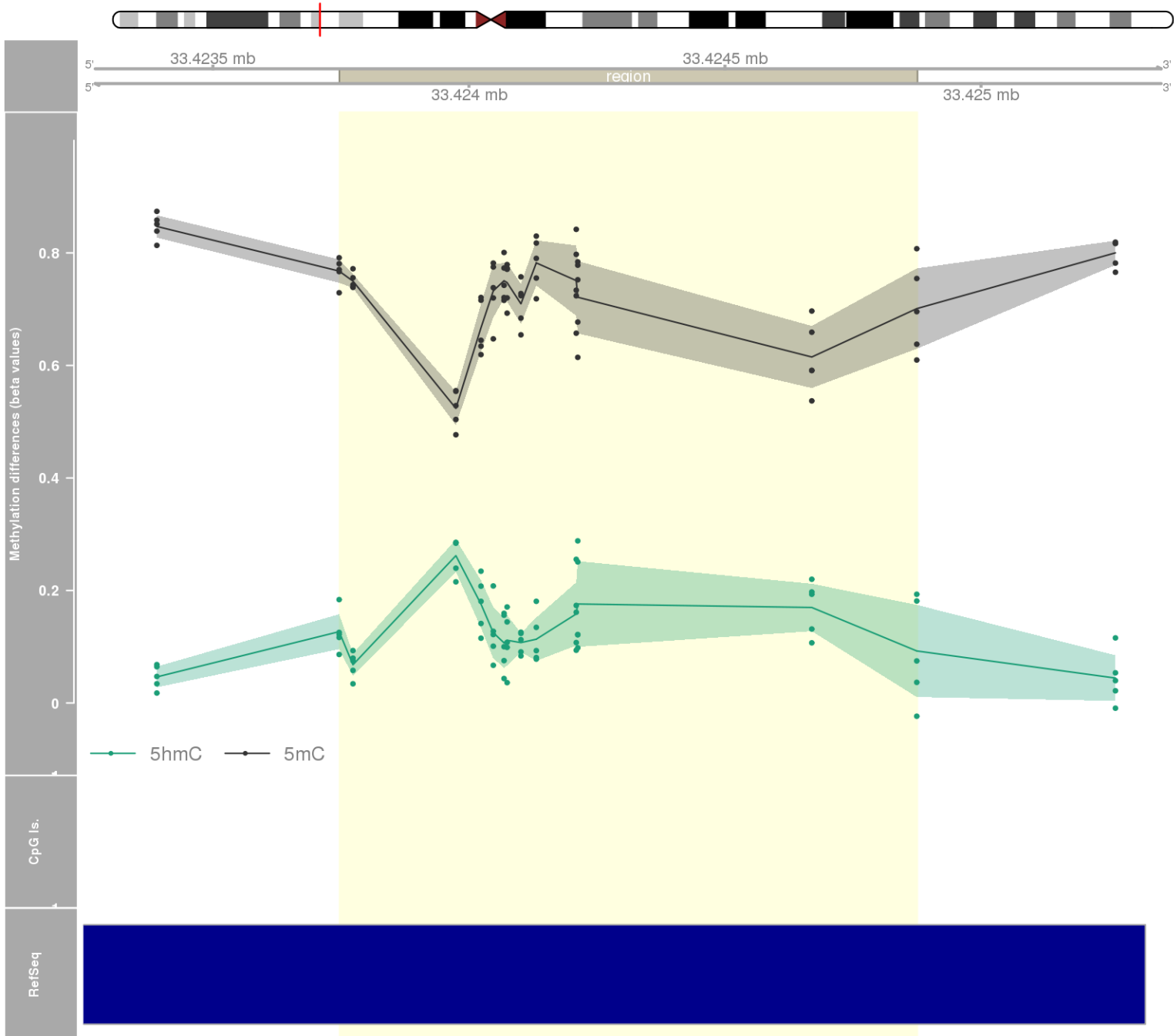
DMR 197 // chr8:41167660-41168336 // 676 pb. (10 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.014 // fwerArea: 1



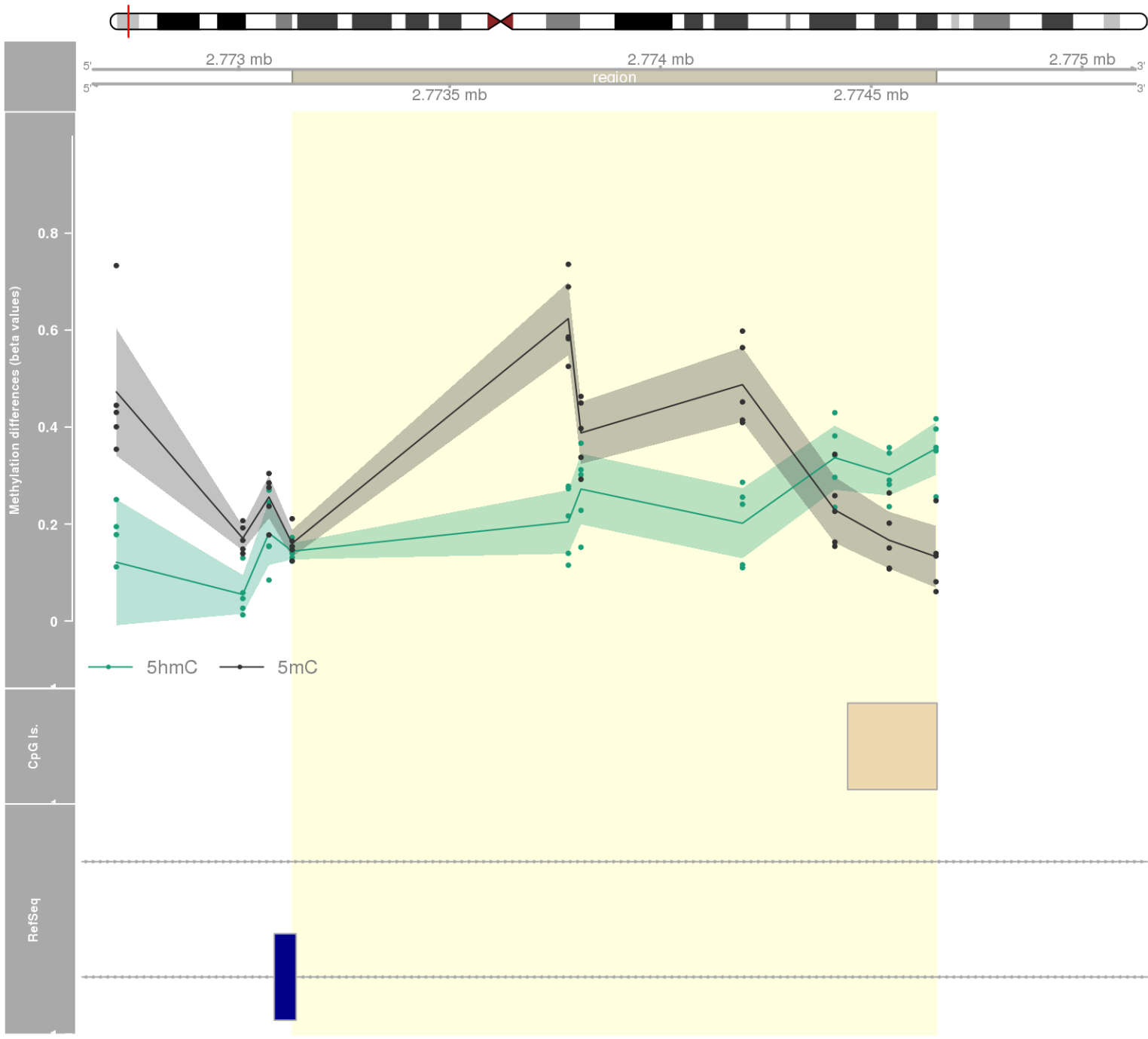
DMR 198 // chr2:69001115-69002378 // 1263 pb. (12 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.014 // fwerArea: 1
- genes: ARHGAP25 -



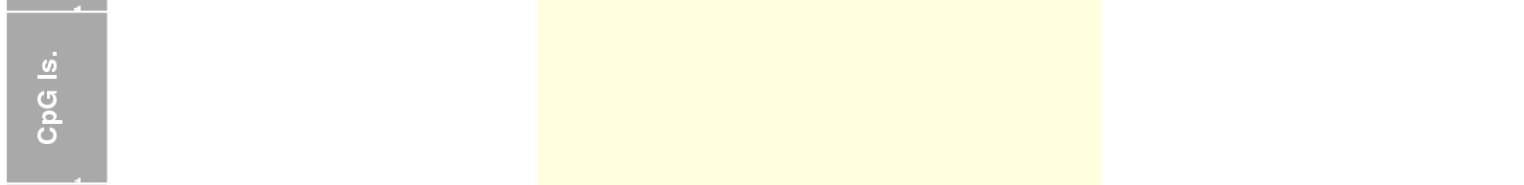
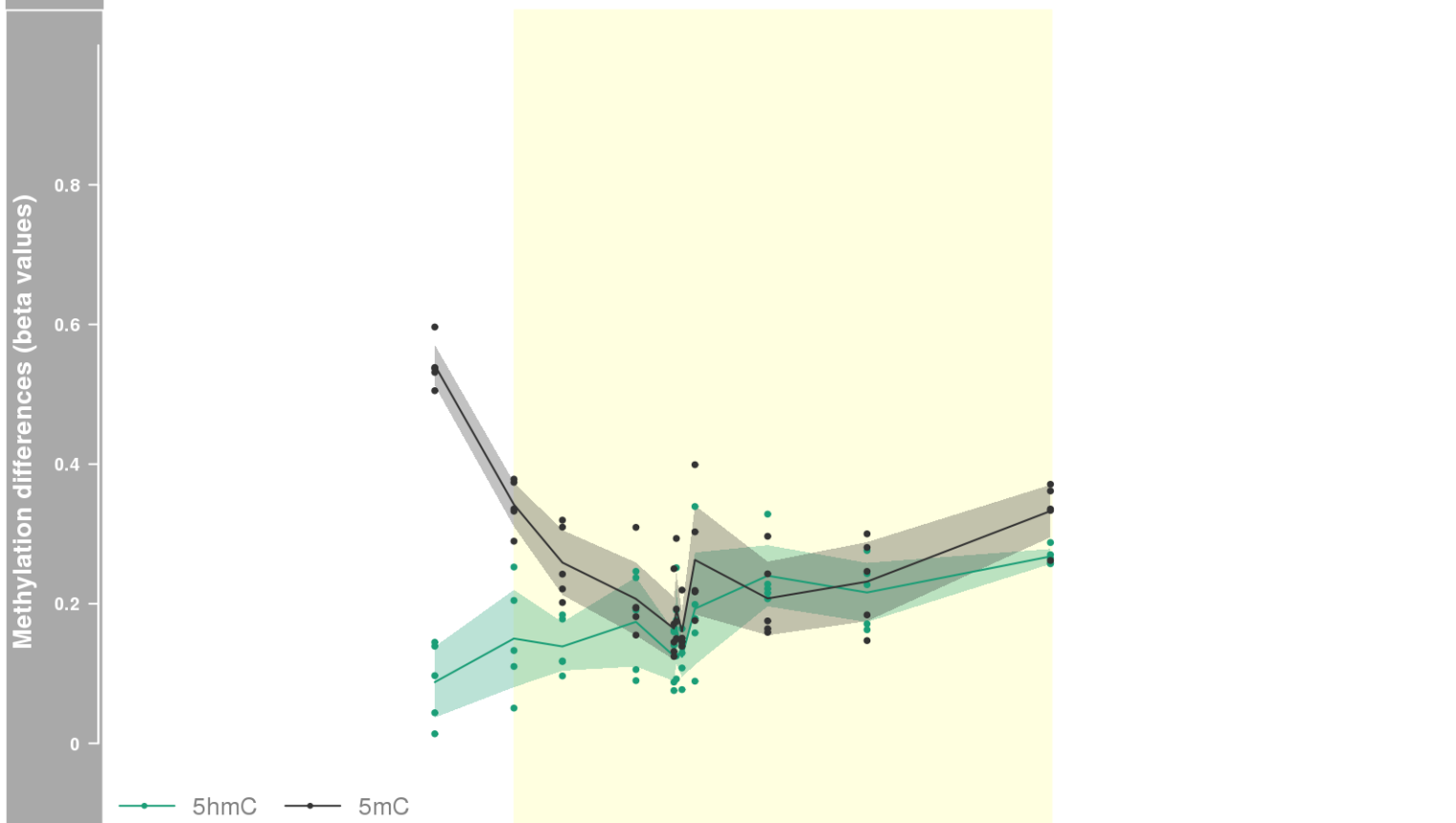
DMR 199 // chr6:33423747-33424875 // 1128 pb. (13 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.014 // fwerArea: 1
- genes: ZBTB9 -



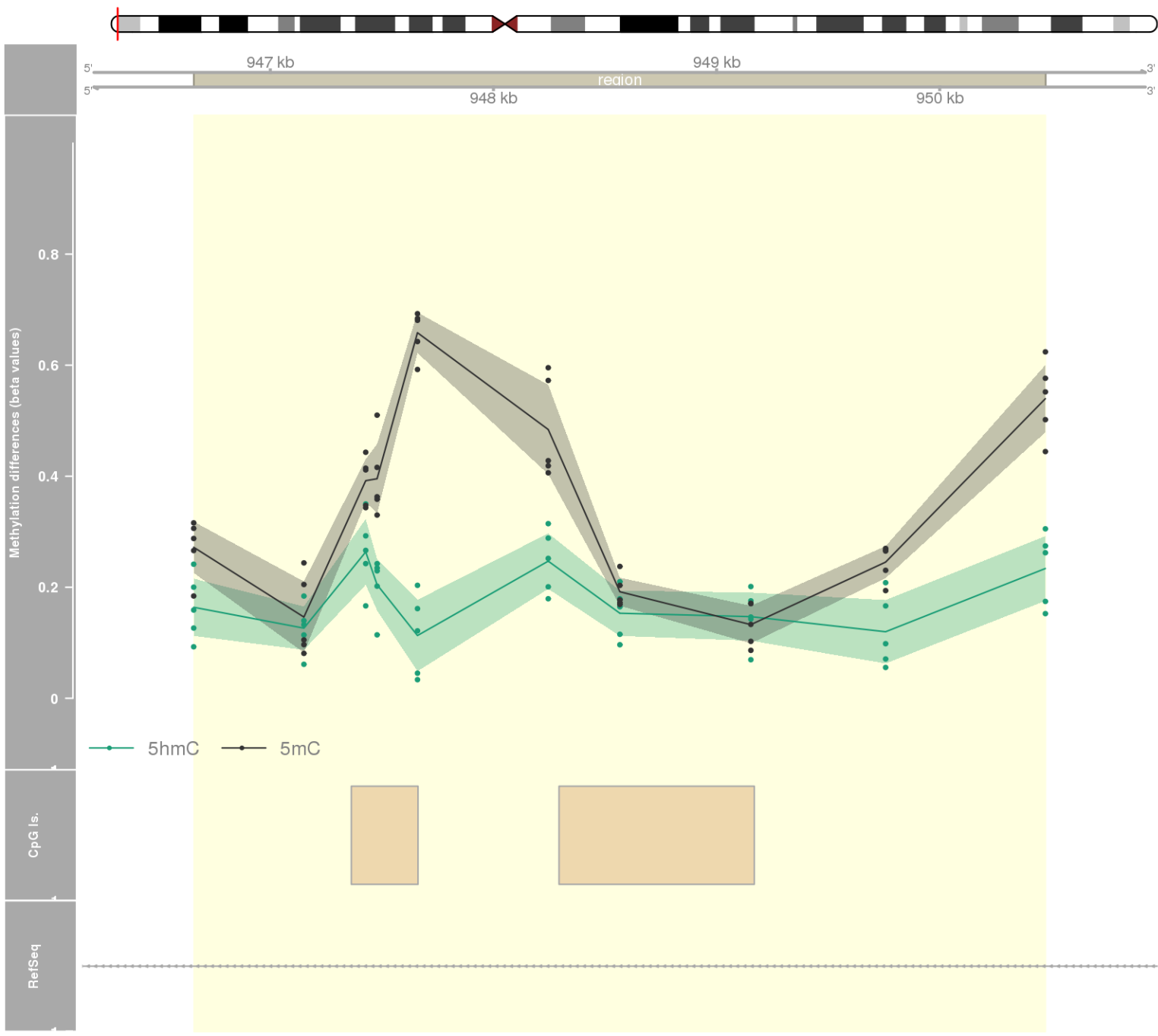
DMR 200 // chr7:2773127-2774654 // 1527 pb. (7 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.014 // fwerArea: 1
- genes: AMZ1 / GNA12 -



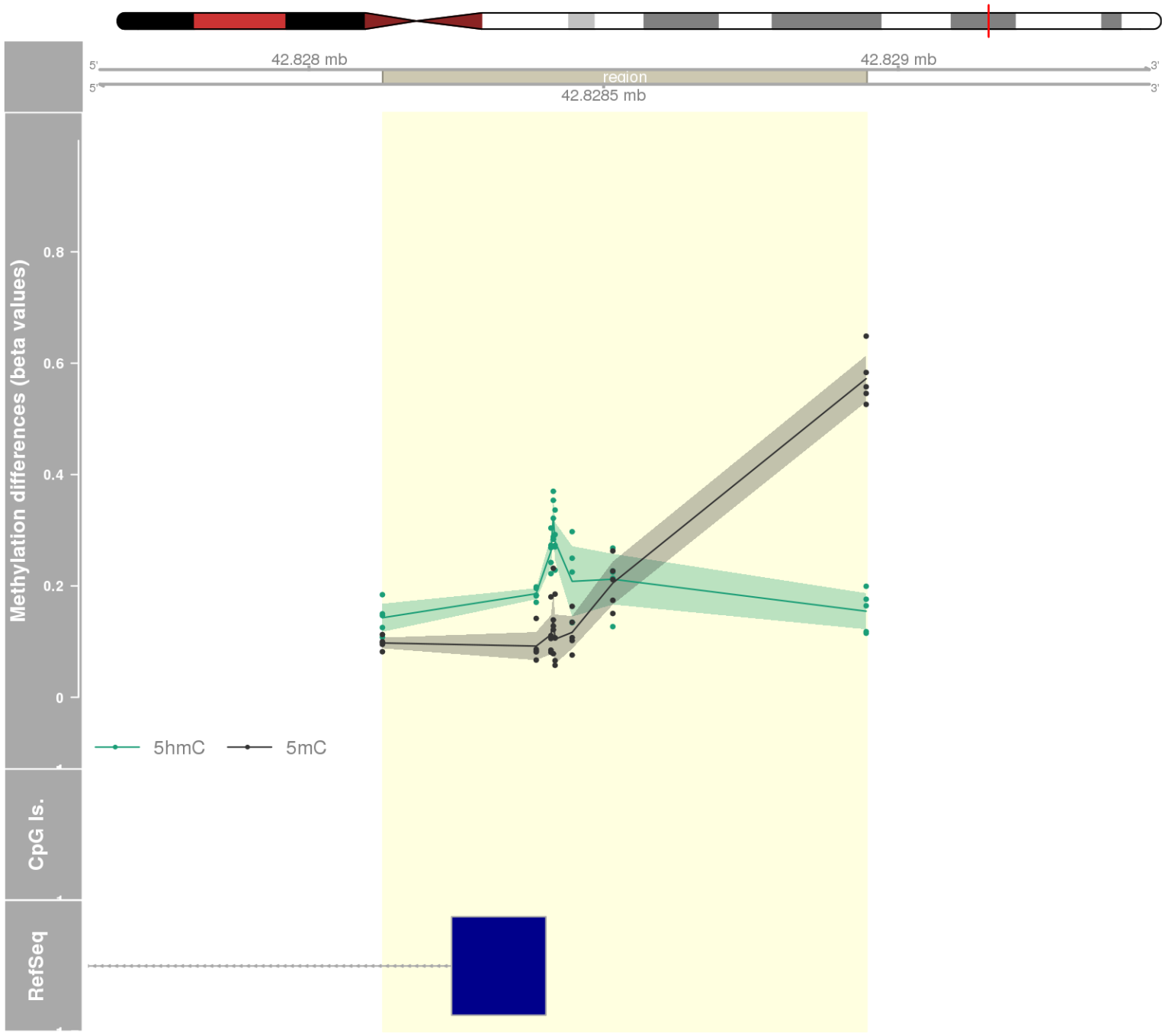
DMR 201 // chr12:116996871-116997535 // 664 pb. (10 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.014 // fwerArea: 1
- genes: MAP1LC3B2 -



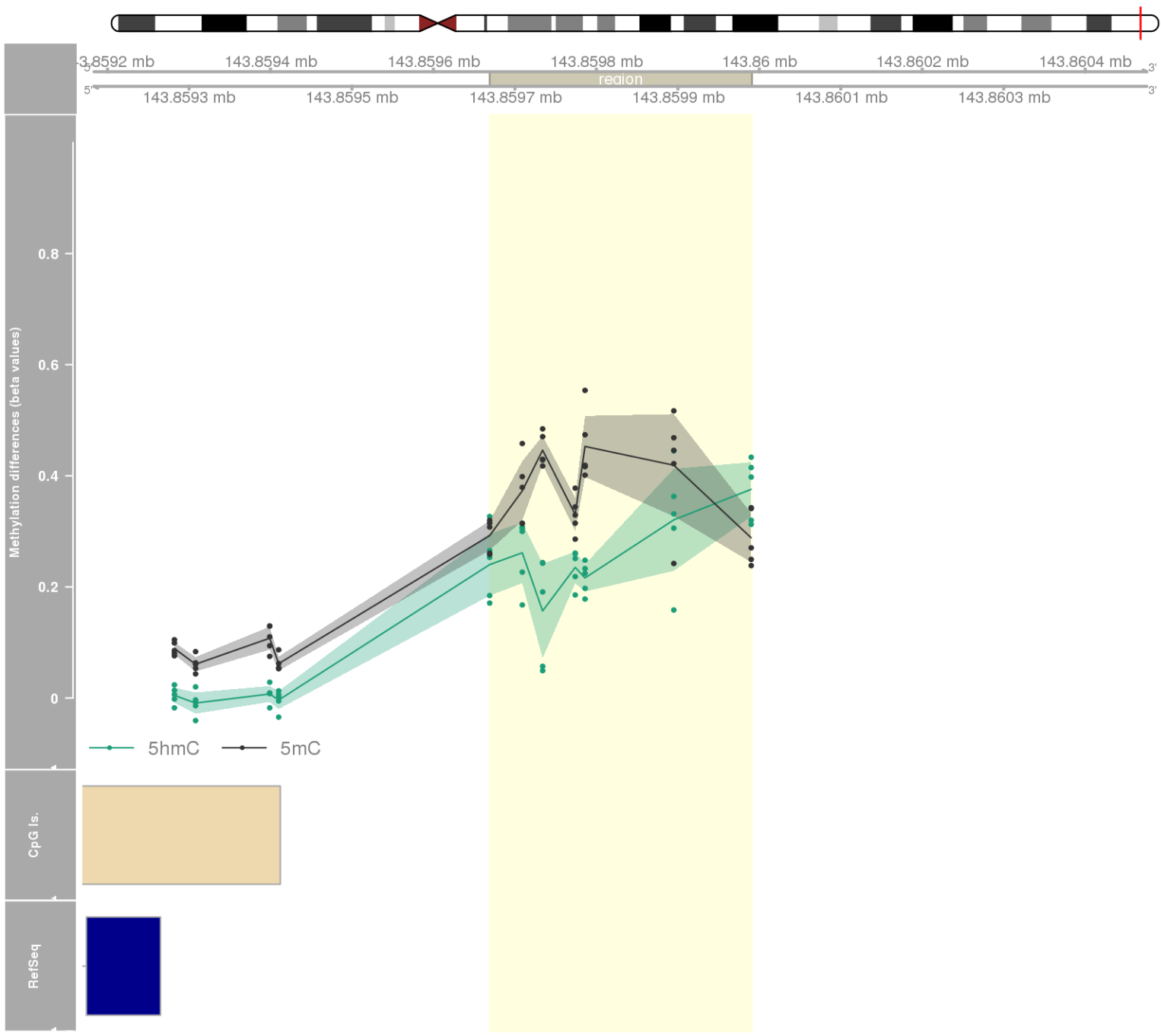
DMR 202 // chr7:946657-950474 // 3817 pb. (10 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.014 // fwerArea: 1
- genes: ADAP1 / COX19 -



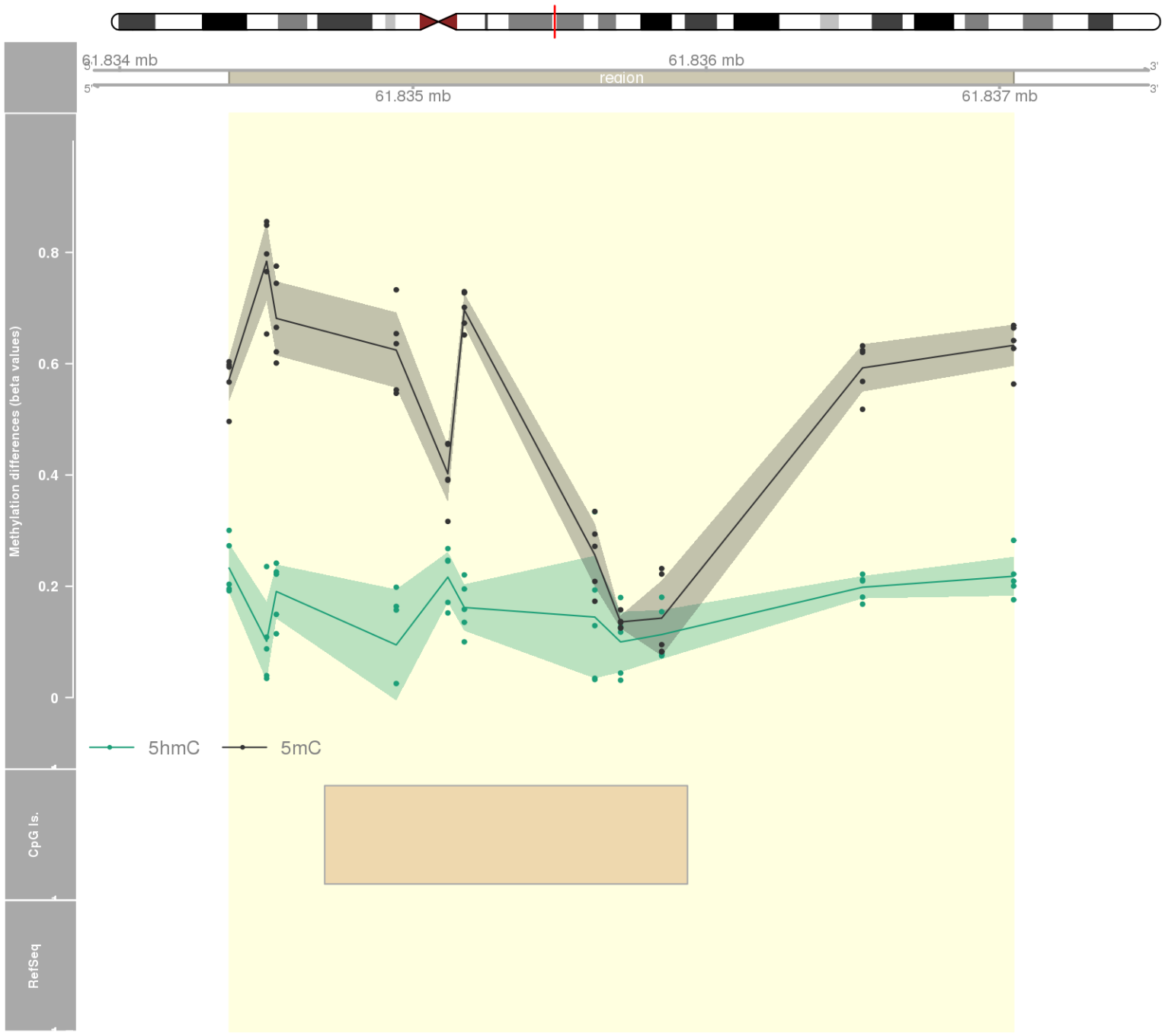
DMR 203 // chr22:42828125-42828946 // 821 pb. (8 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.014 // fwerArea: 1
- genes: NFAM1 -



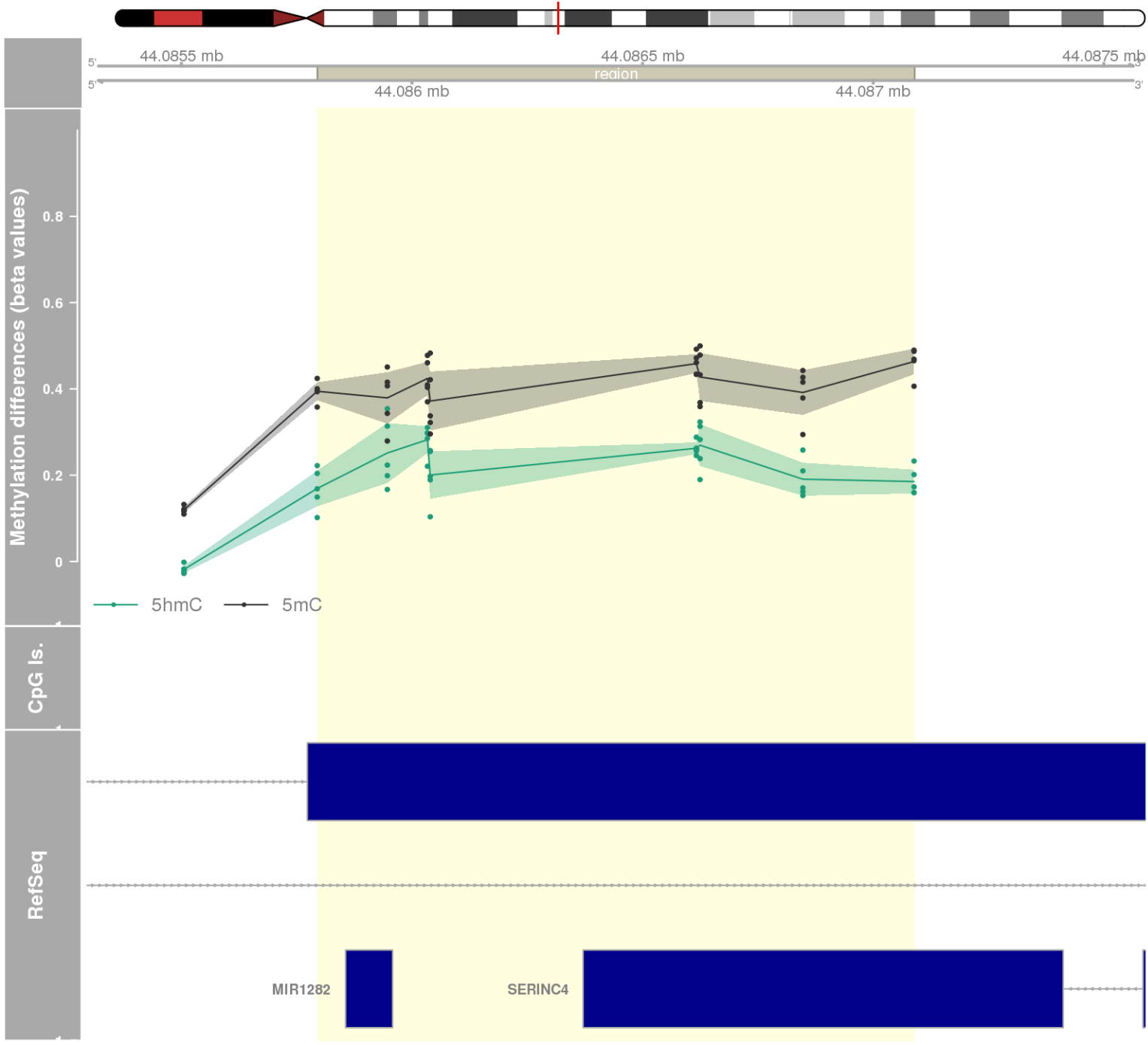
DMR 204 // chr8:143859669-143859990 // 321 pb. (7 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.014 // fwerArea: 1



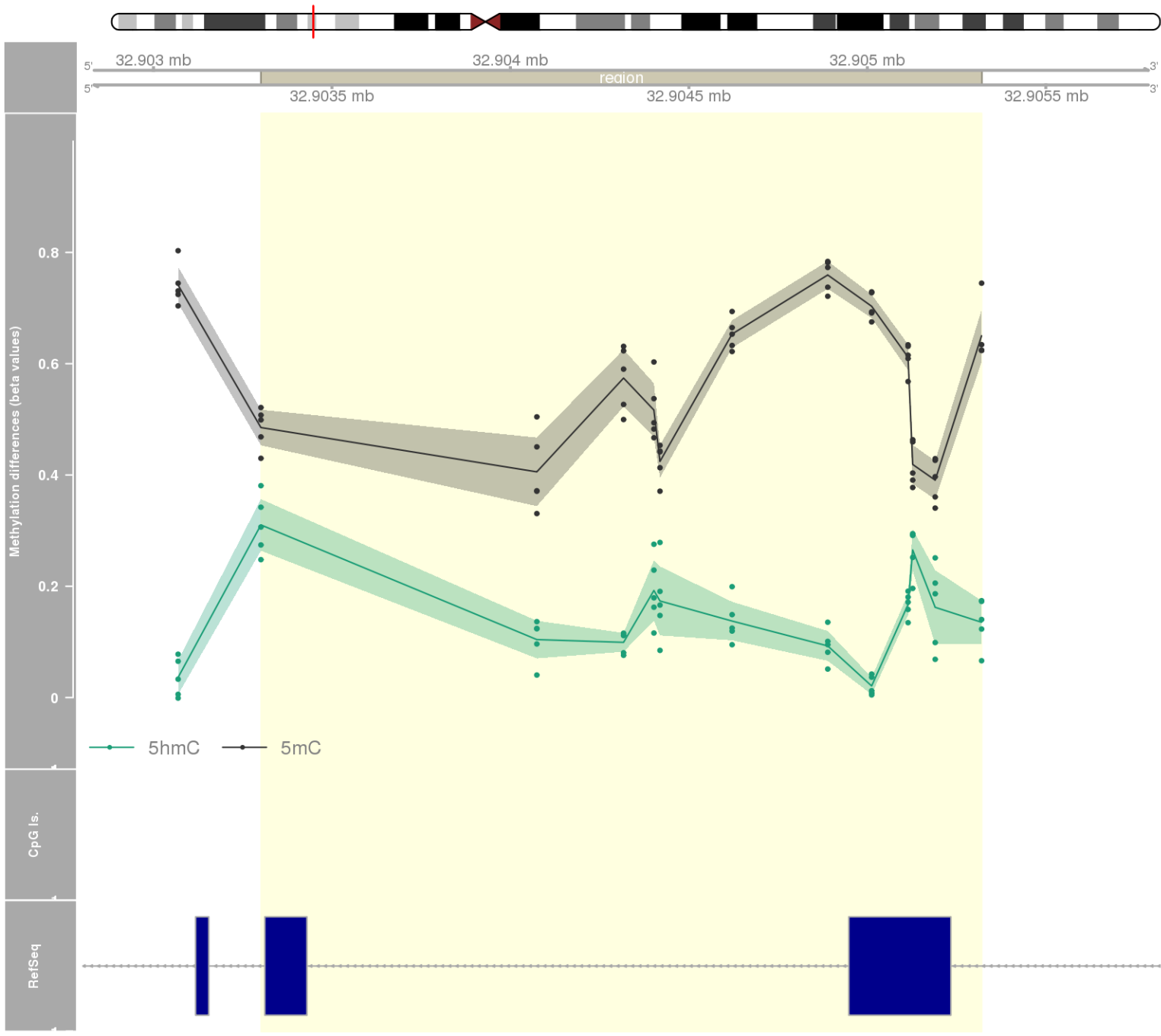
DMR 205 // chr8:61834373-61837048 // 2675 pb. (11 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.014 // fwerArea: 1



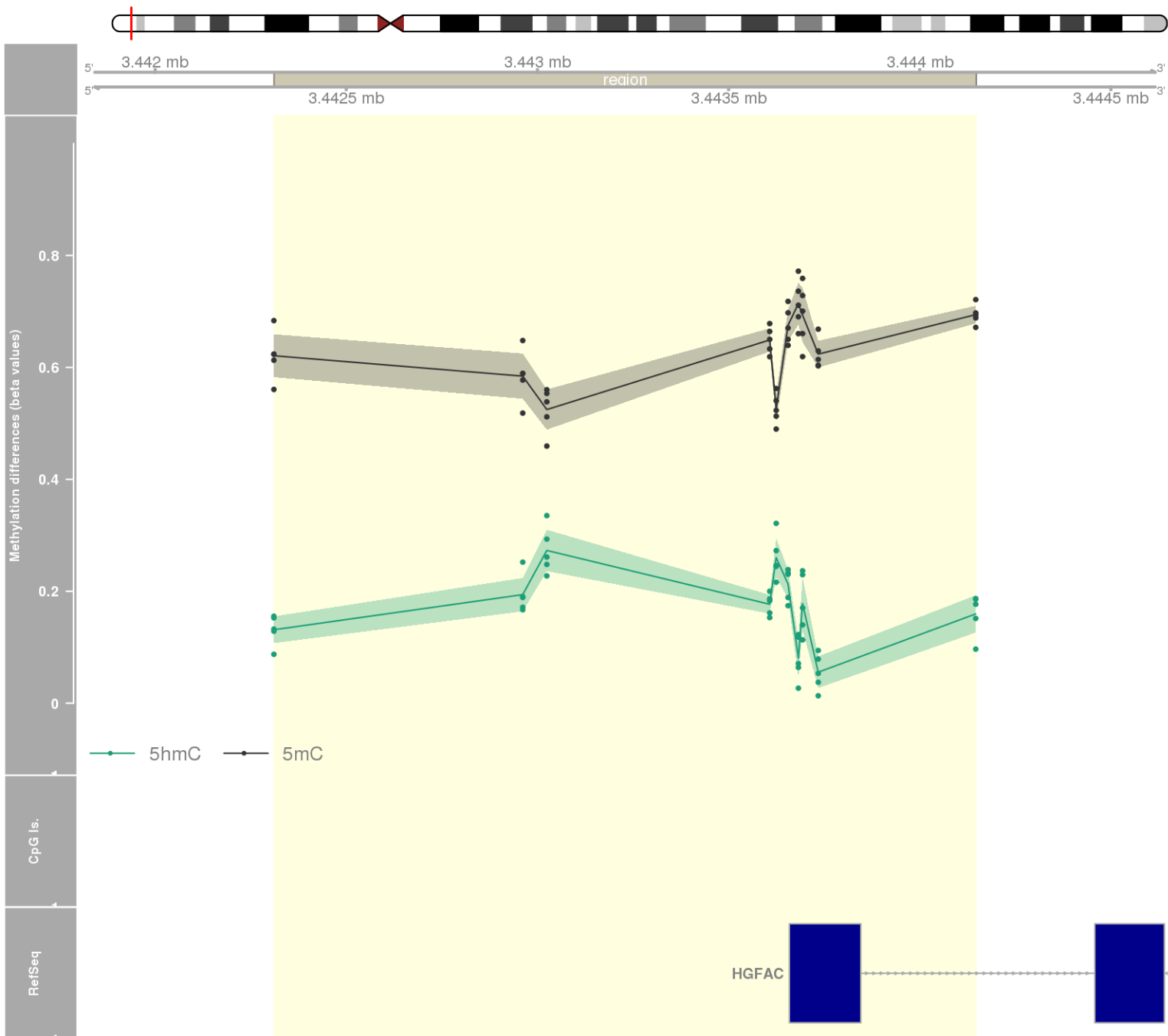
DMR 206 // chr15:44085795-44087089 // 1294 pb. (8 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.015 // fwerArea: 1
- genes: MIR1282 / SERF2 / SERINC4 -



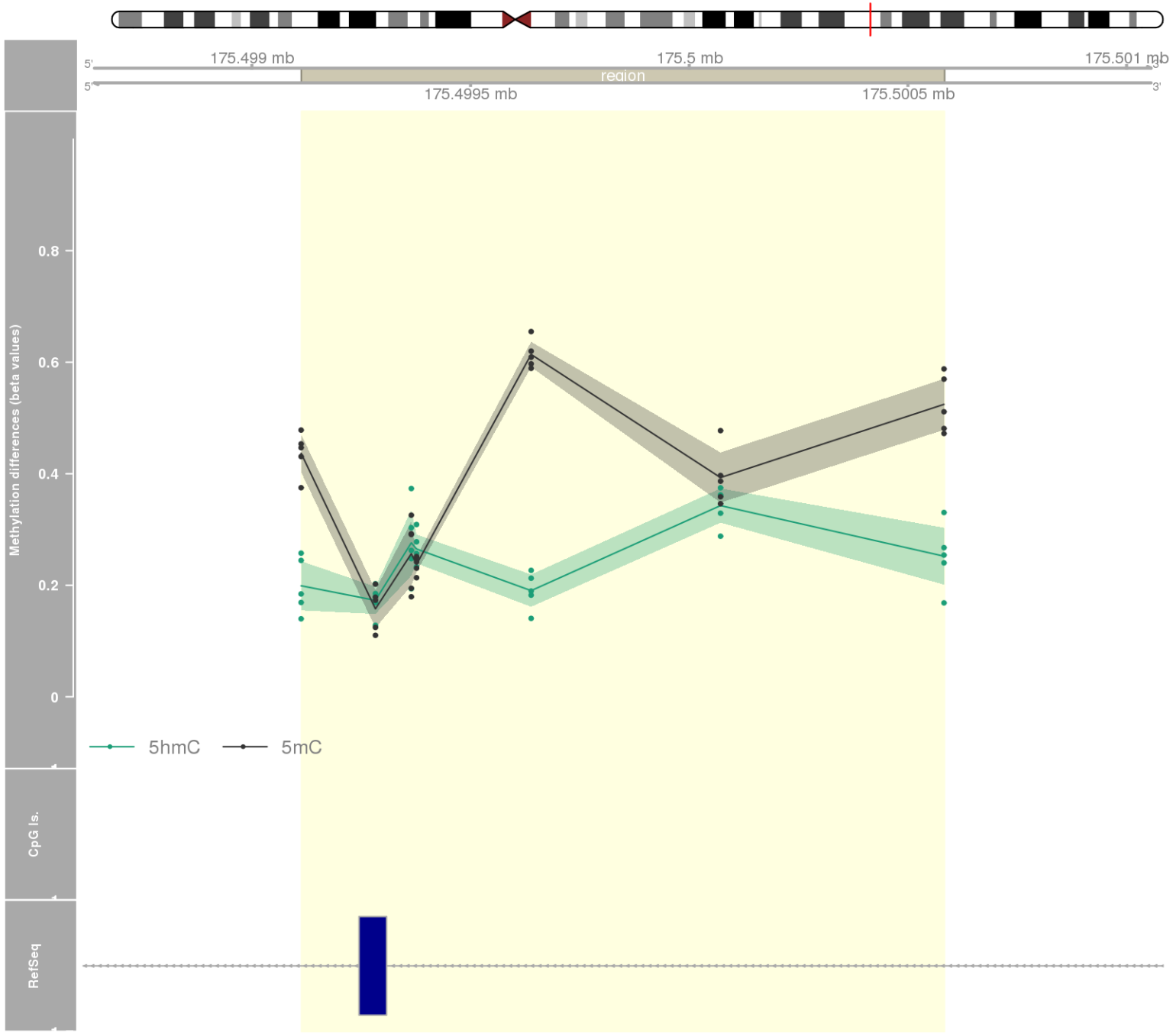
DMR 207 // chr6:32903301-32905320 // 2019 pb. (12 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.015 // fwerArea: 1
- genes: HLA-DMB -



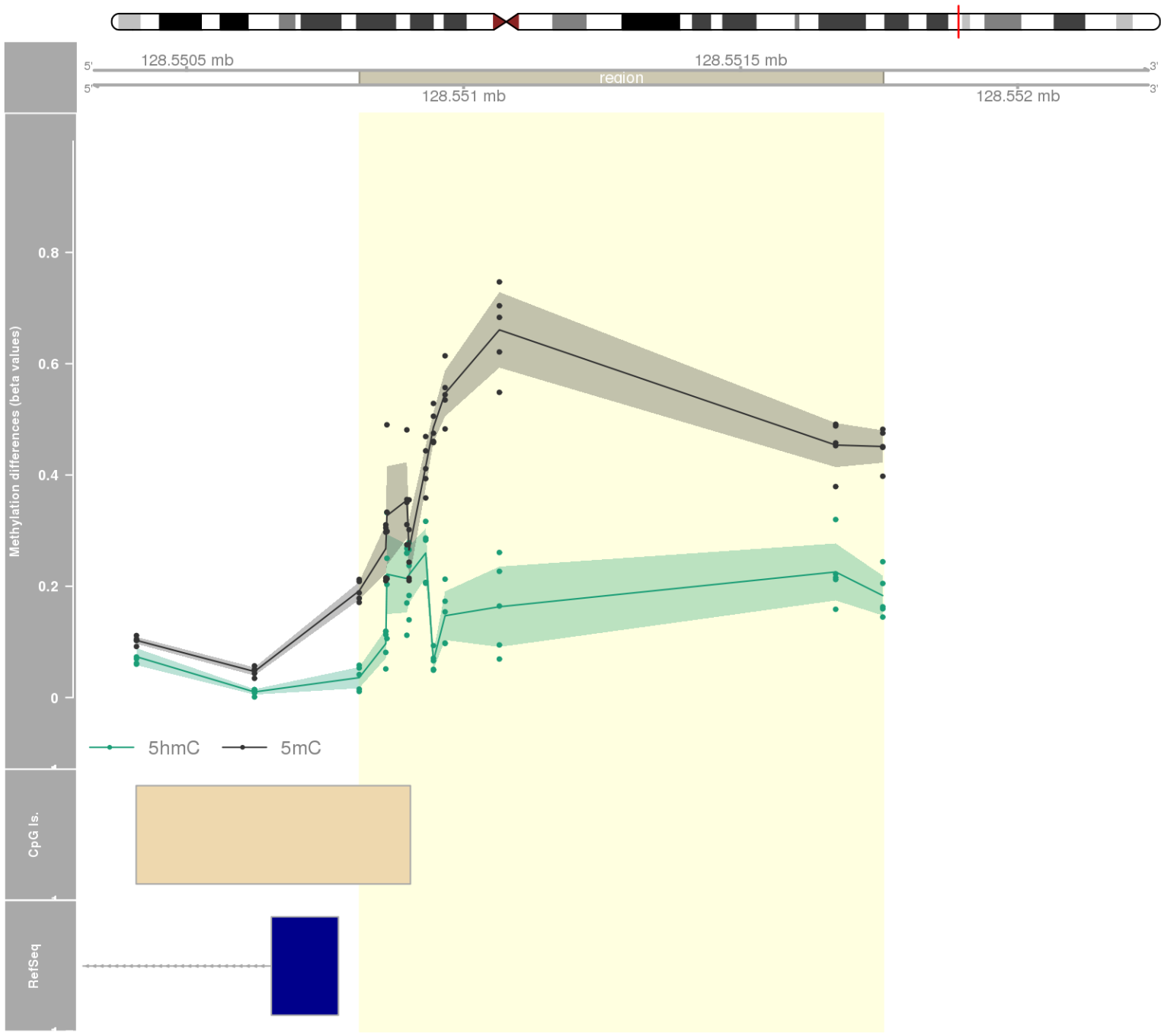
DMR 208 // chr4:3442311-3444147 // 1836 pb. (10 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.015 // fwerArea: 1
- genes: HGFA C -



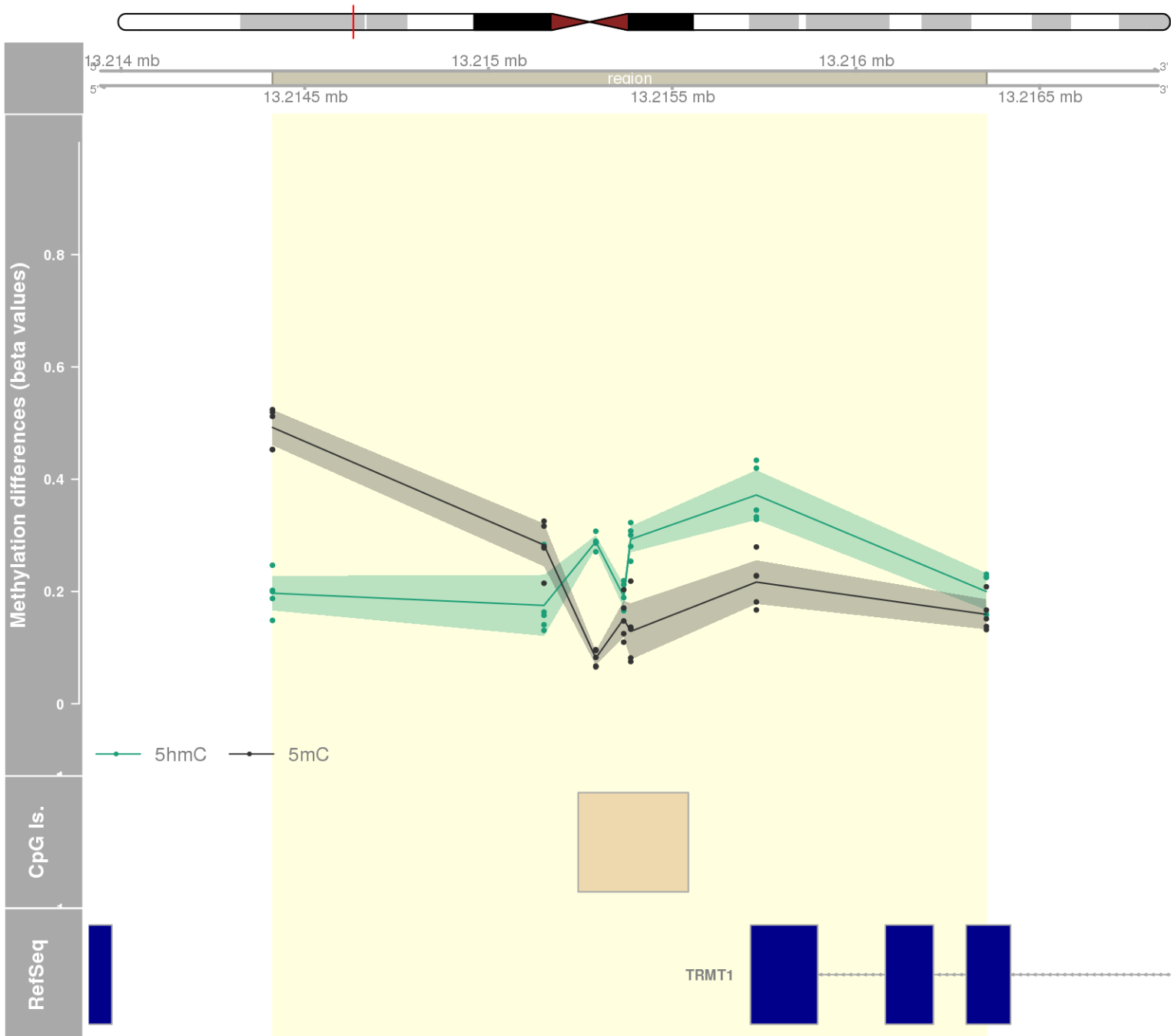
DMR 209 // chr2:175499113-175500583 // 1470 pb. (7 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.015 // fwerArea: 1
- genes: WIPF1 -



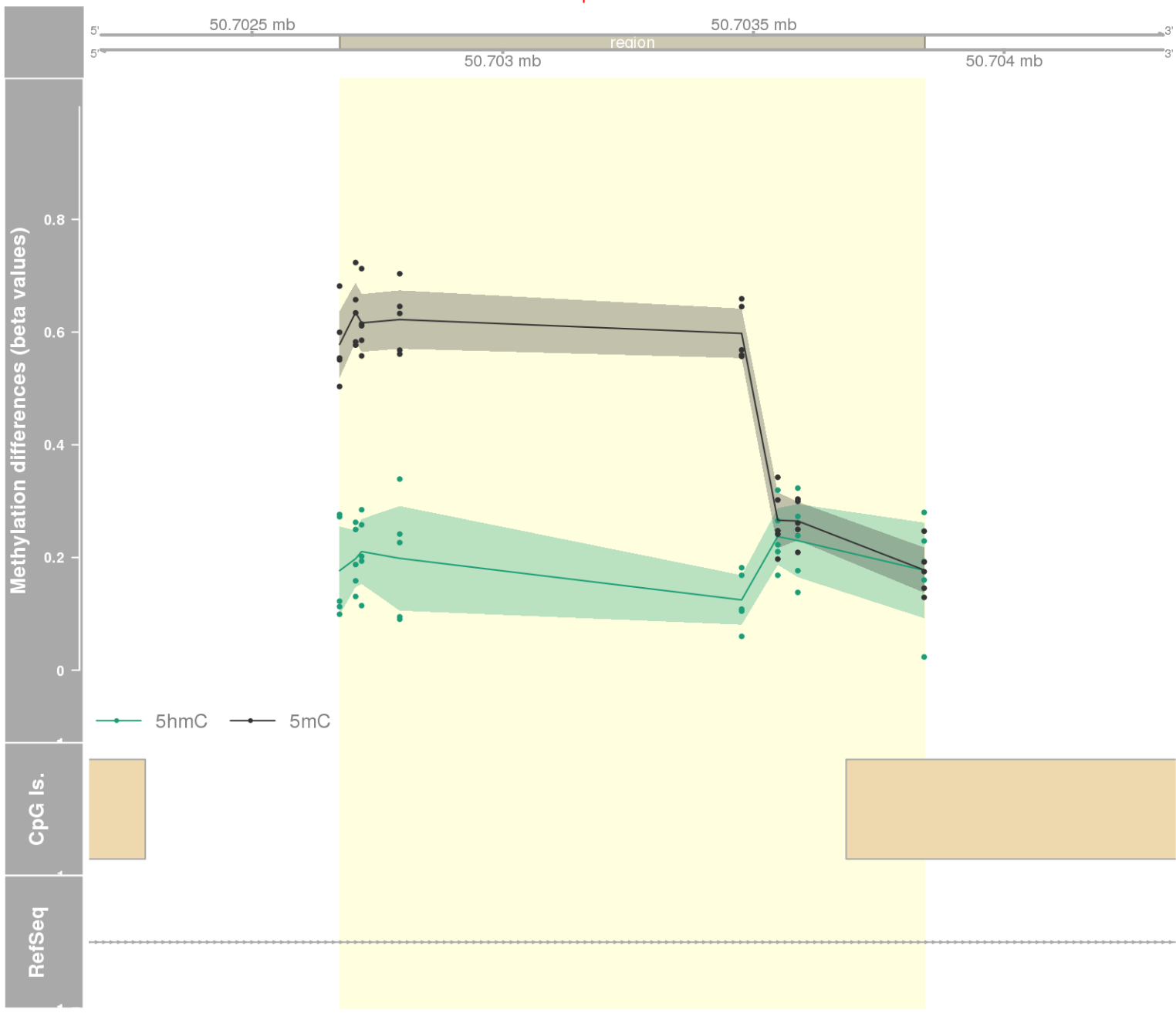
DMR 210 // chr7:128550812-128551757 // 945 pb. (11 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.015 // fwerArea: 1



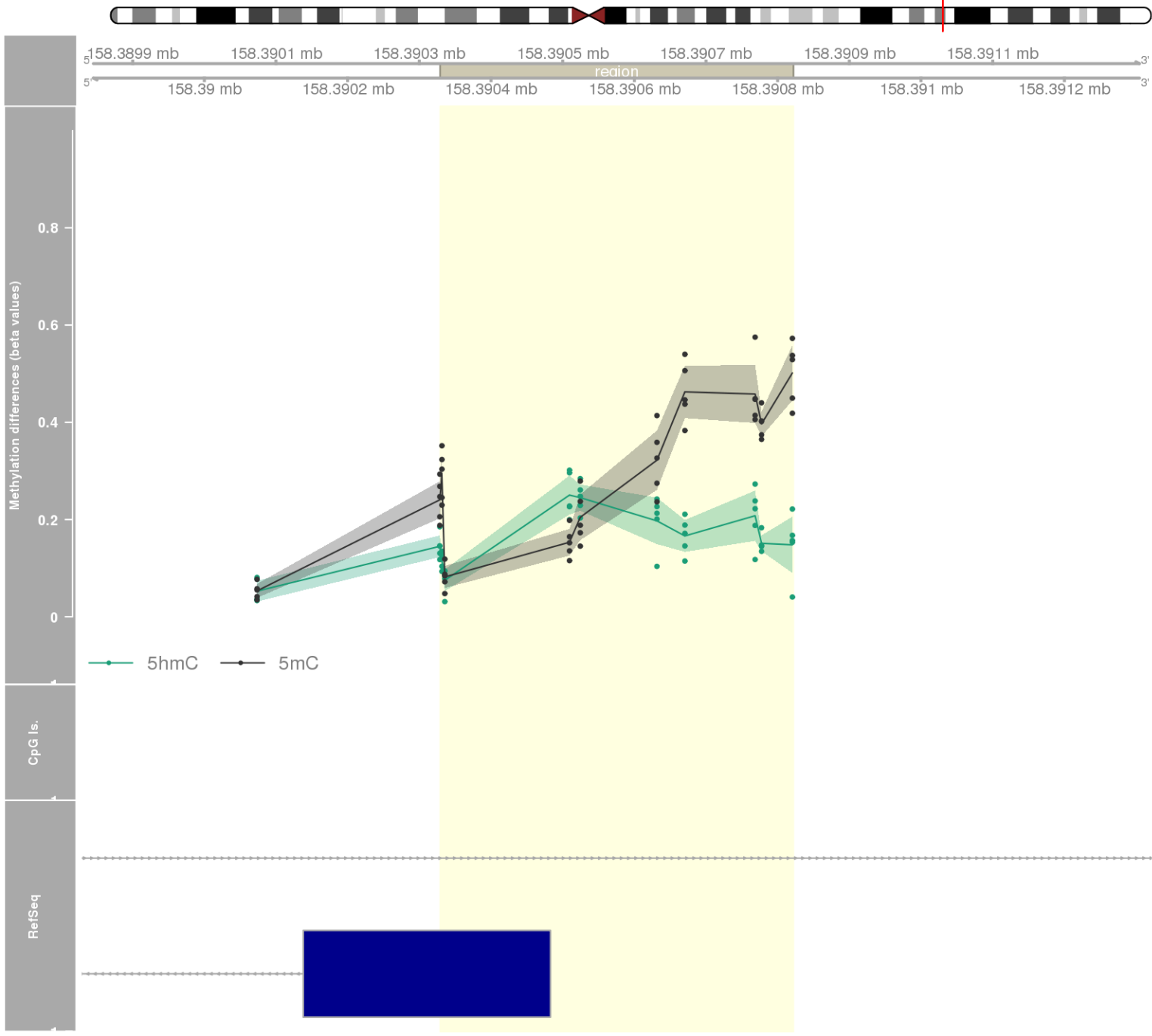
DMR 211 // chr19:13214412-13216355 // 1943 pb. (7 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.015 // fwerArea: 1
- genes: TRMT1 -



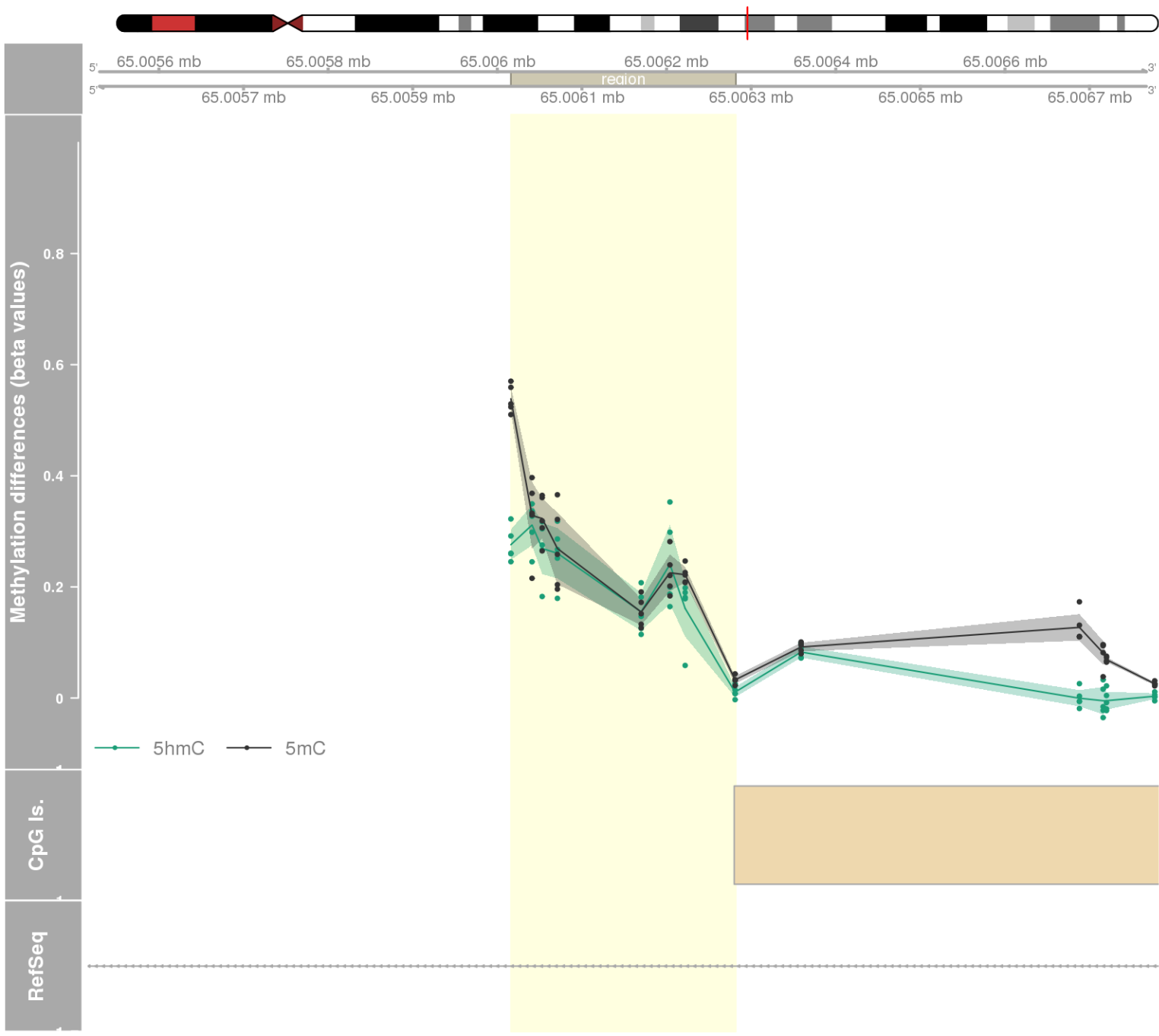
DMR 212 // chr13:50702675-50703841 // 1166 pb. (8 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.015 // fwerArea: 1
- genes: DLEU1 -



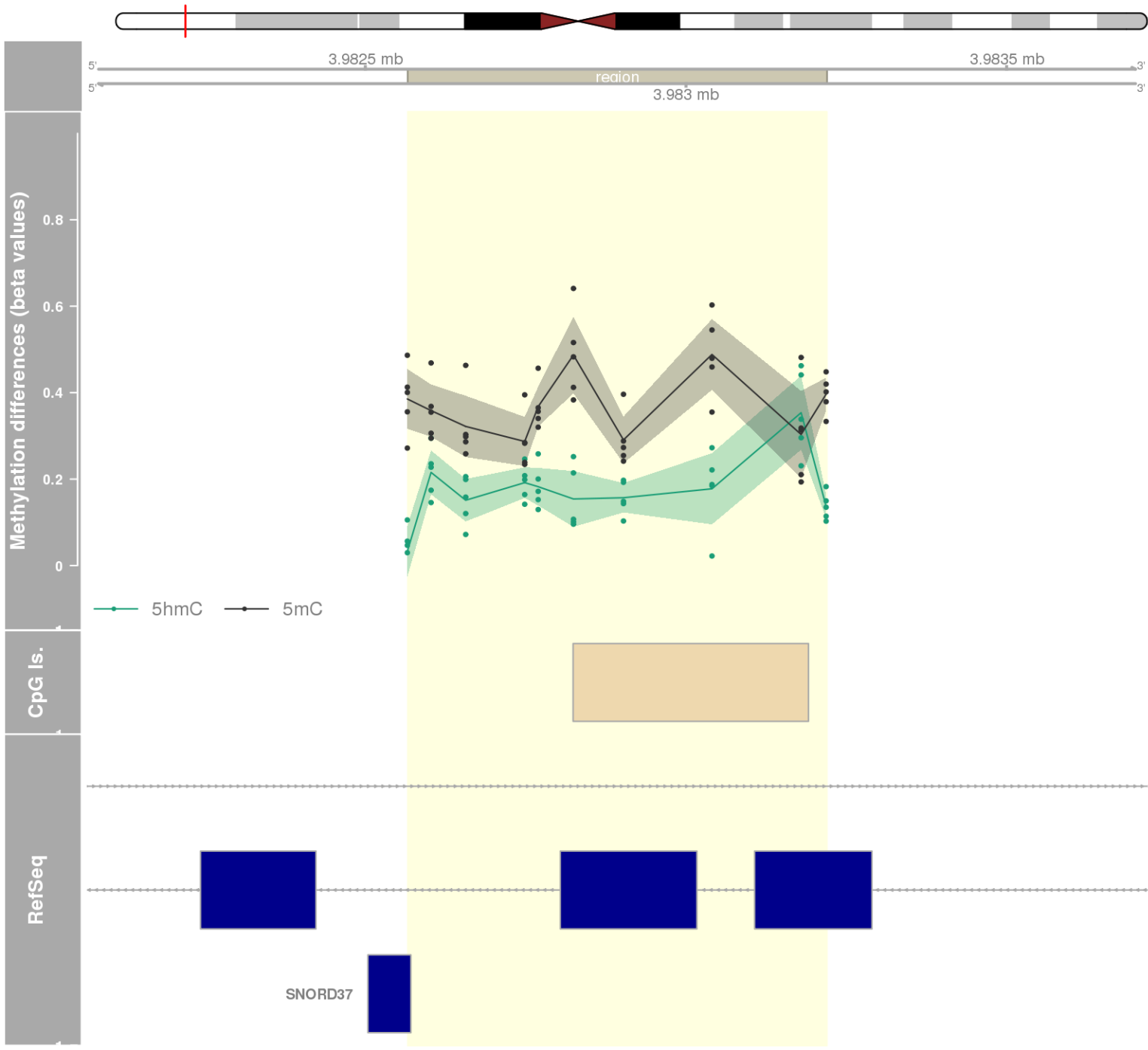
DMR 213 // chr3:158390329-158390821 // 492 pb. (10 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.015 // fwerArea: 1
- genes: GFM1 / LXN -



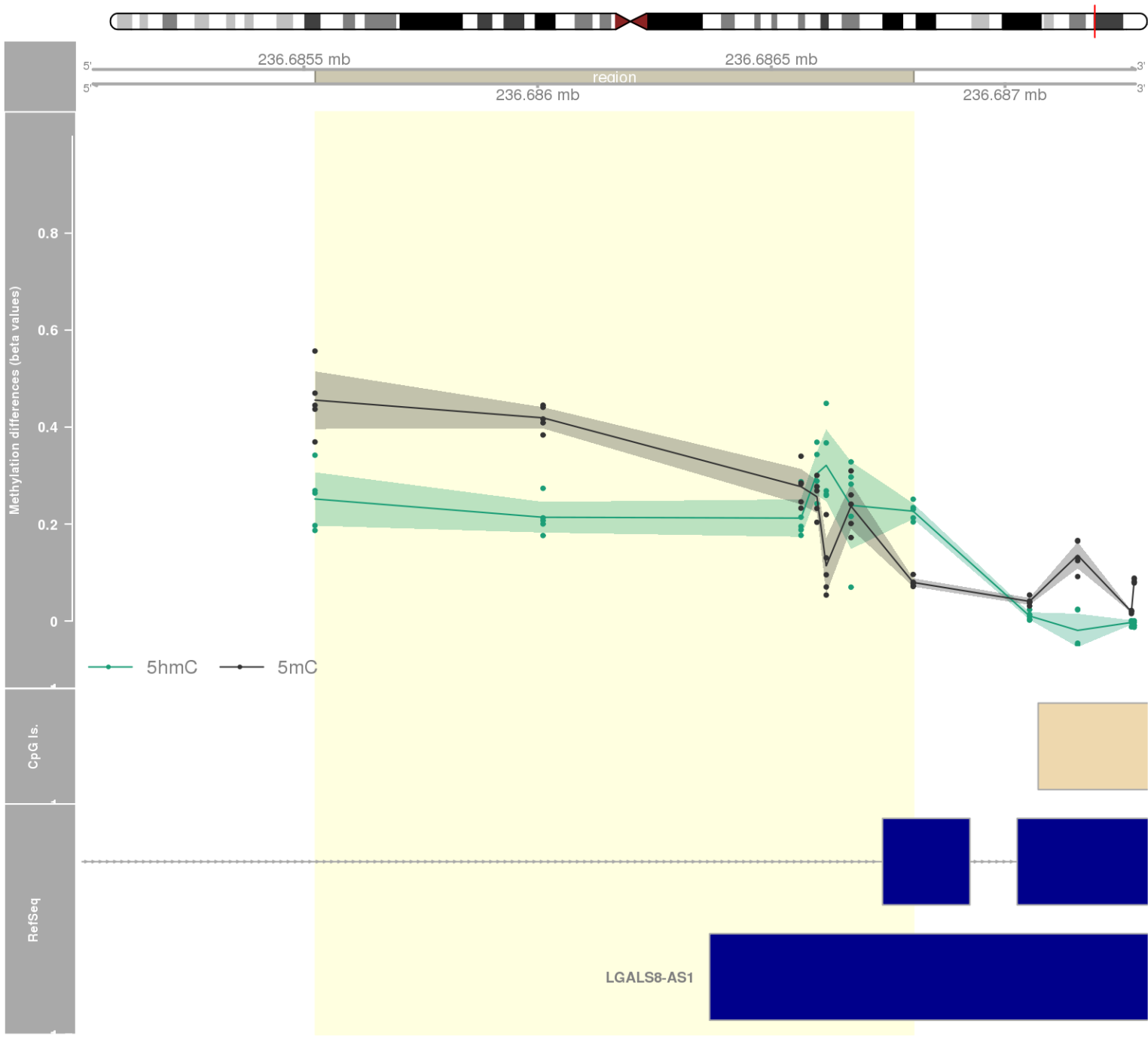
DMR 214 // chr14:65006016-65006281 // 265 pb. (8 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.015 // fwerArea: 1
- genes: HSPA2 -



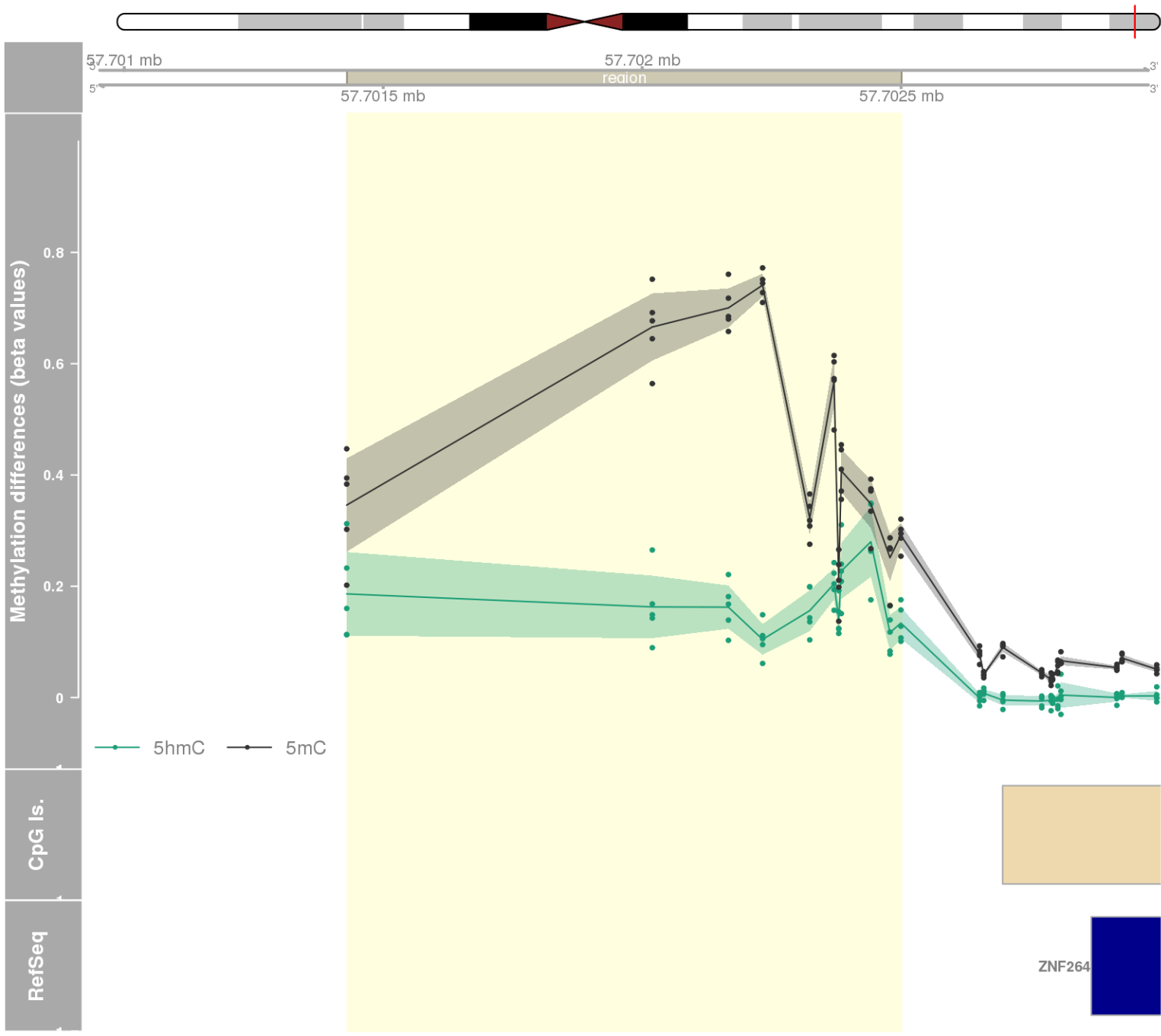
DMR 215 // chr19:3982566-3983219 // 653 pb. (10 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.015 // fwerArea: 1
- genes: SNORD37 / EEF2 -



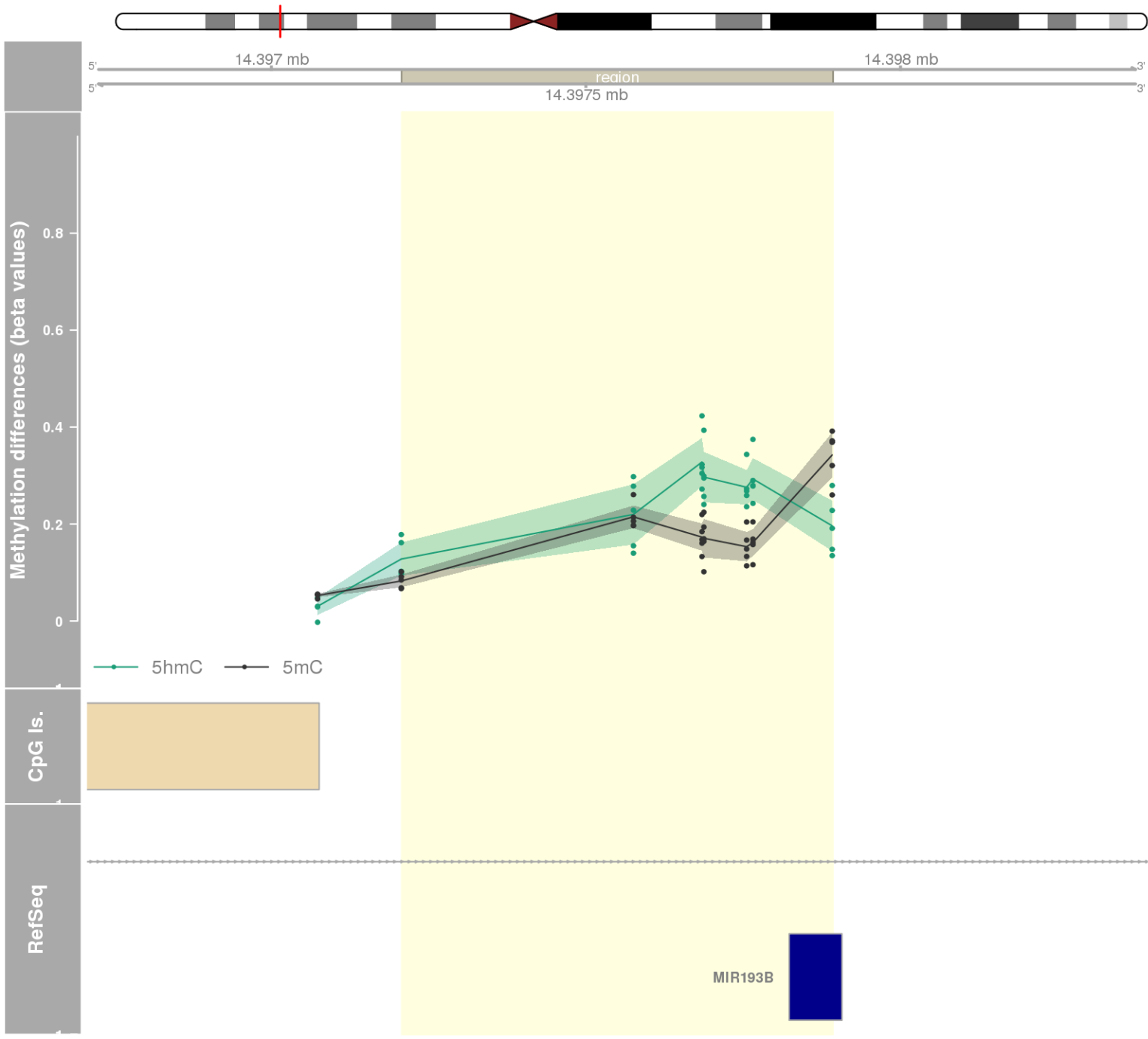
DMR 216 // chr1:236685524-236686804 // 1280 pb. (7 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.015 // fwerArea: 1
- genes: LGALS8-AS1 / LGALS8 -



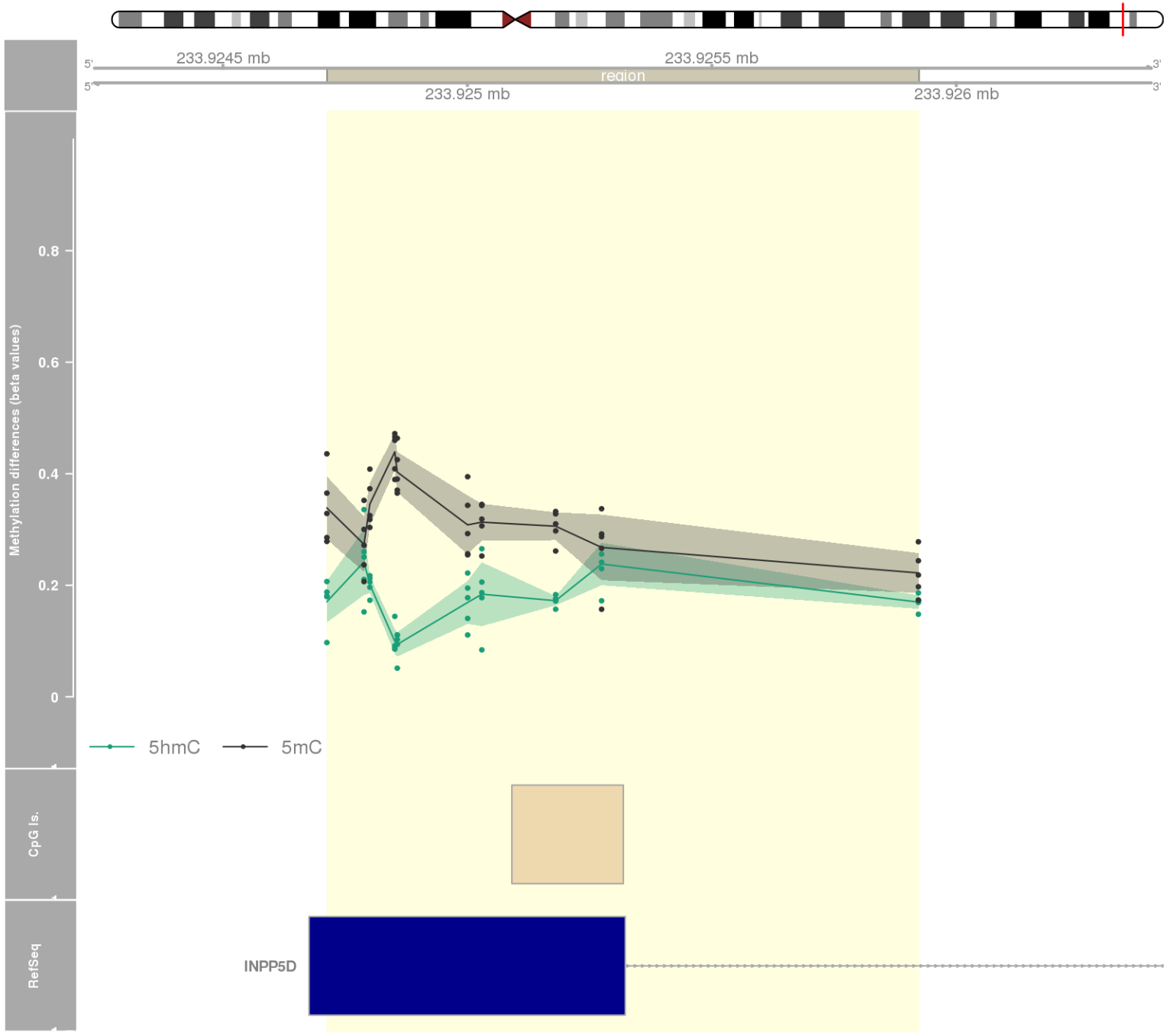
DMR 217 // chr19:57701430-57702500 // 1070 pb. (11 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.015 // fwerArea: 1



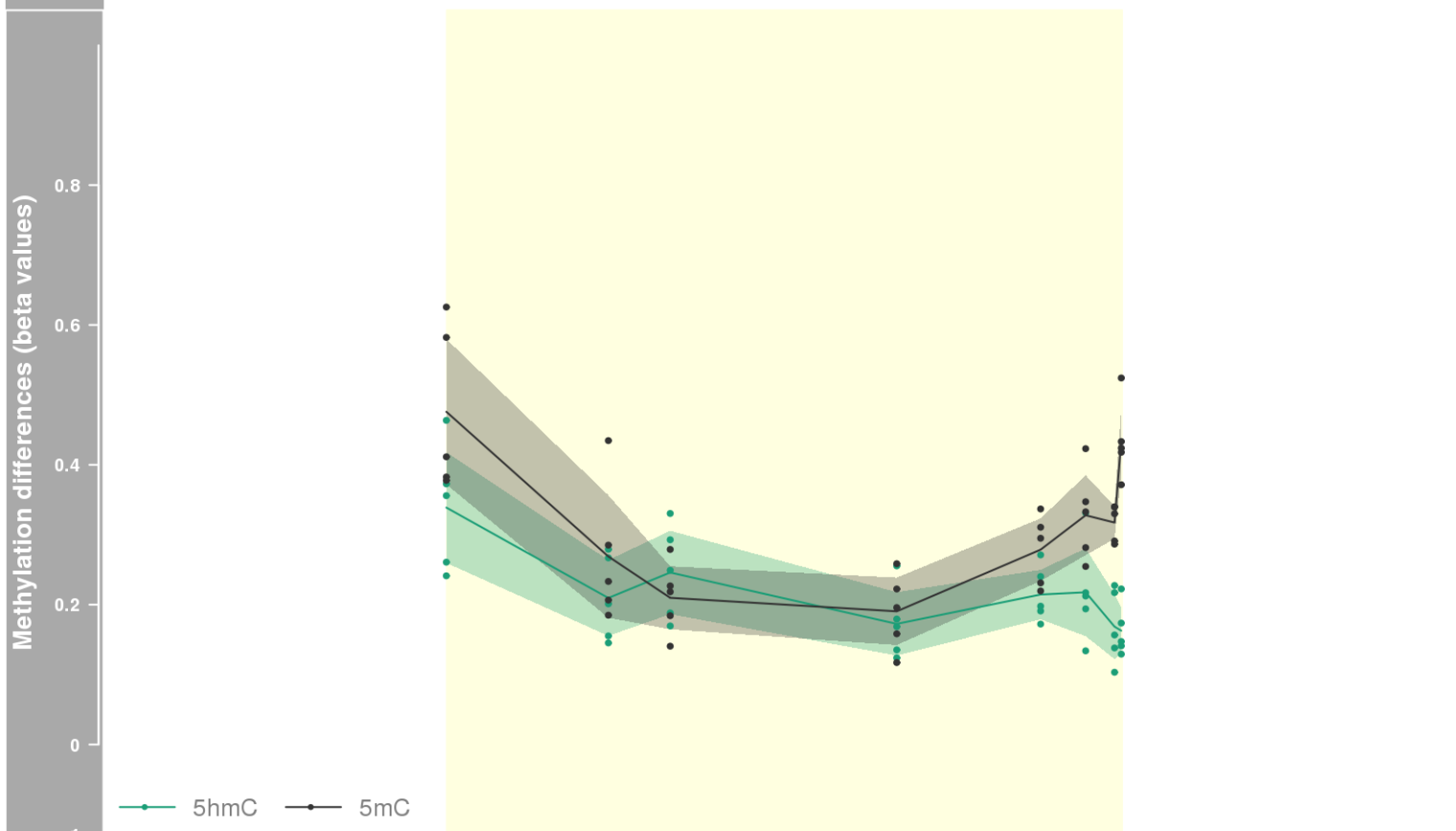
DMR 218 // chr16:14397207-14397892 // 685 pb. (7 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.015 // fwerArea: 1
- genes: MIR193B -



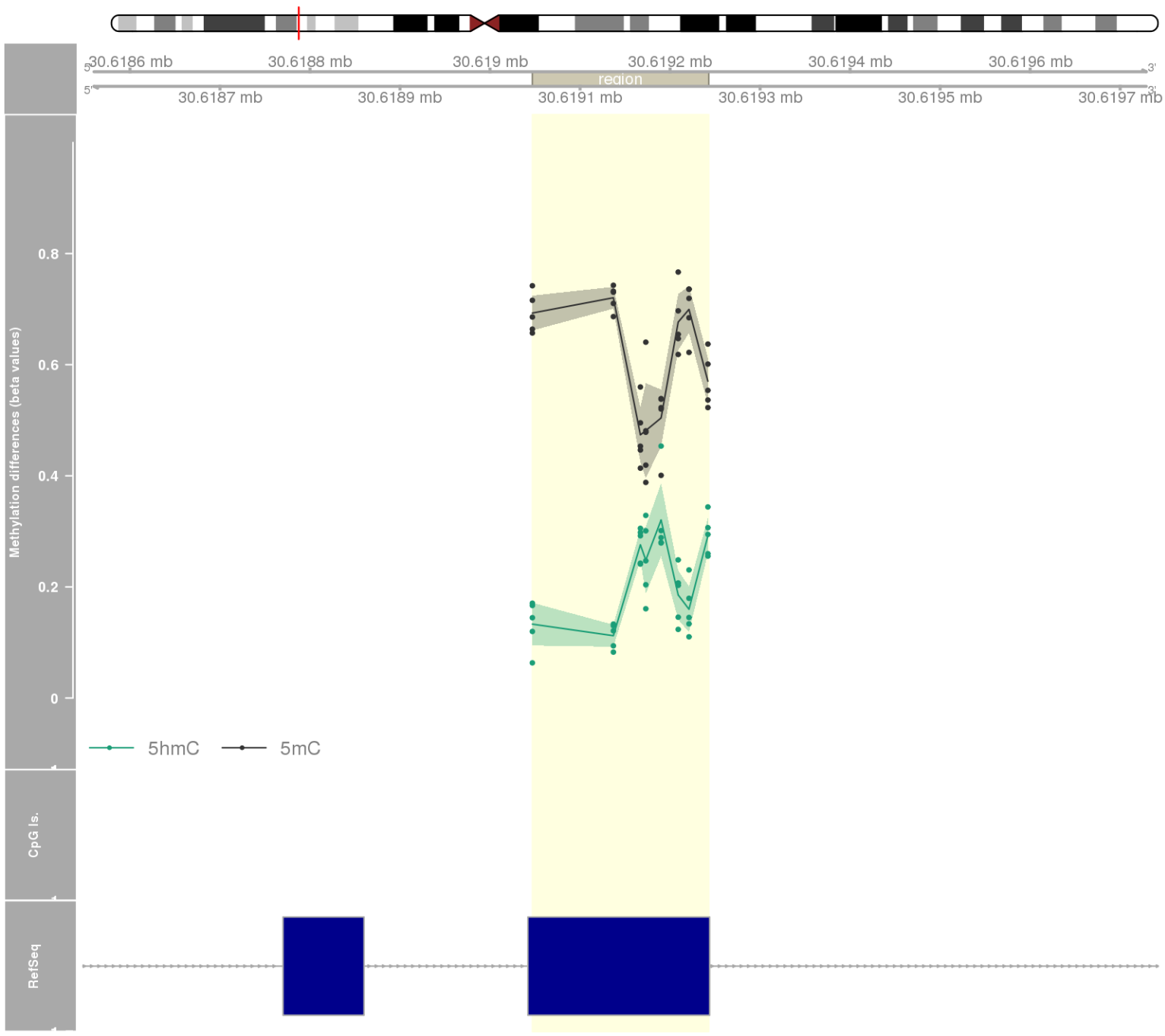
DMR 219 // chr2:233924713-233925923 // 1210 pb. (10 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.015 // fwerArea: 1
- genes: INPP5D -



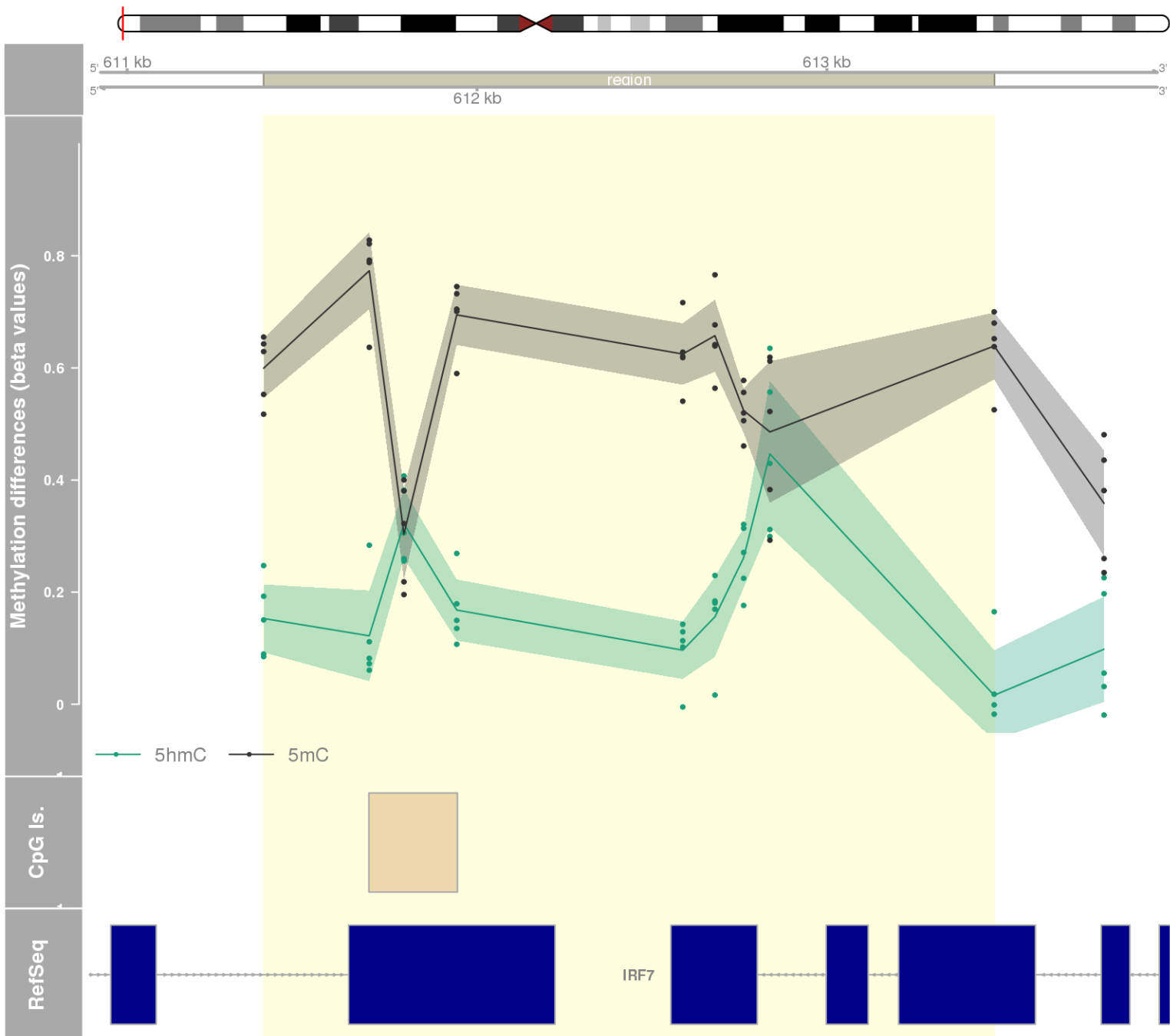
DMR 220 // chr13:111521981-111522985 // 1004 pb. (8 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.015 // fwerArea: 1
- genes: LINC00346 -



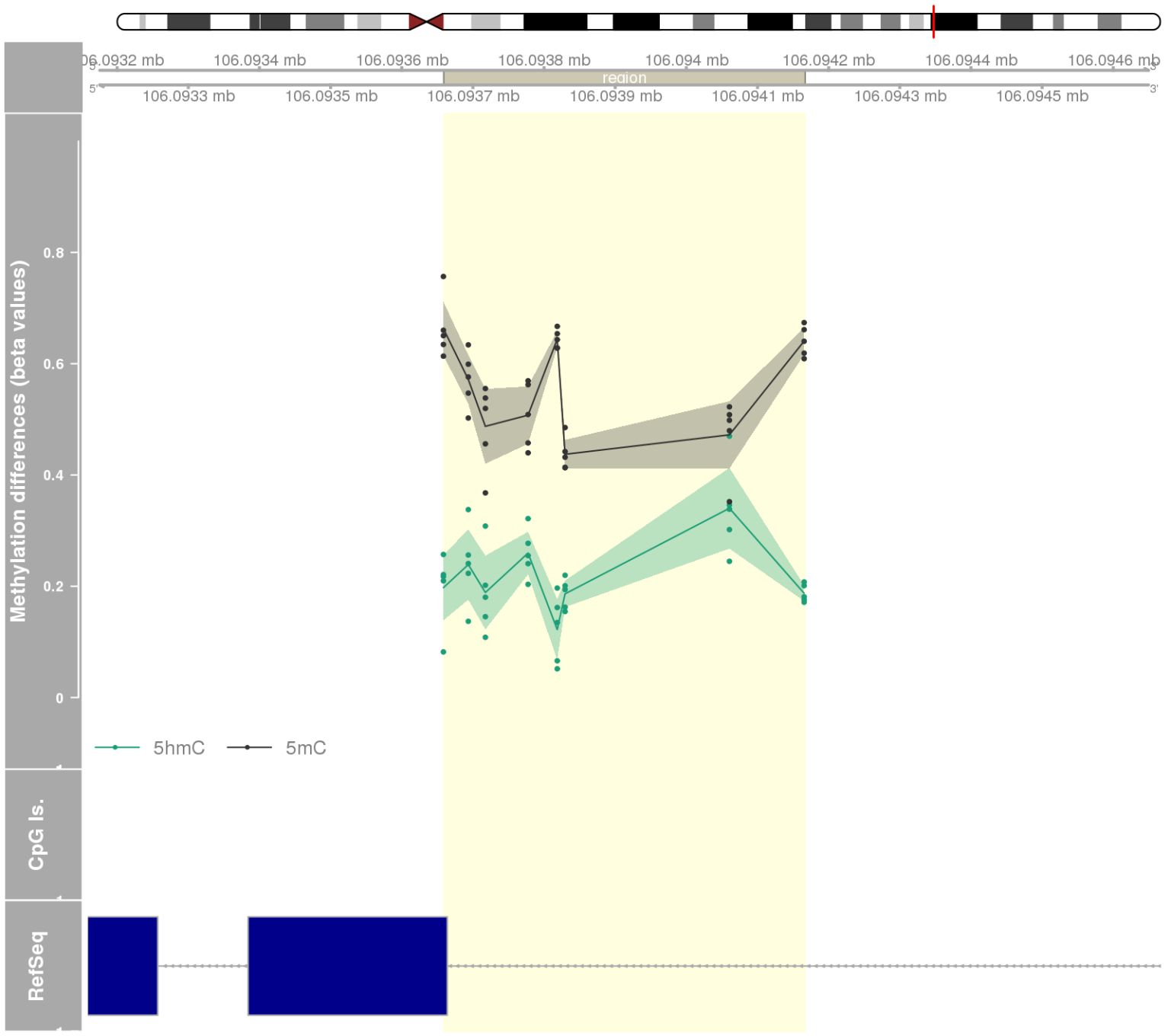
DMR 221 // chr6:30619047-30619242 // 195 pb. (8 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.016 // fwerArea: 1
- genes: C6orf136 -



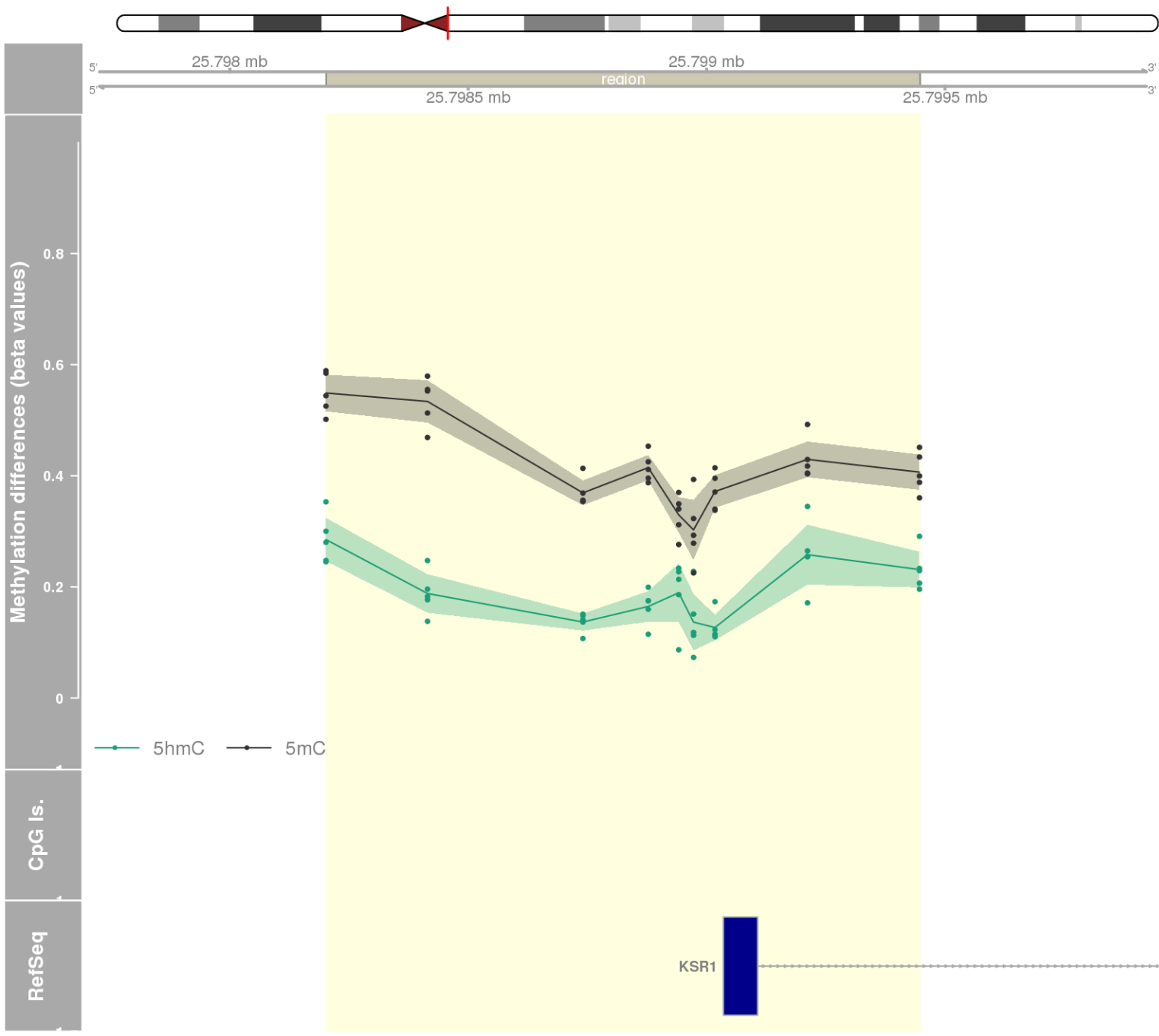
DMR 222 // chr11:611390-613478 // 2088 pb. (9 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.016 // fwerArea: 1
- genes: PHRF1 / IRF7 -



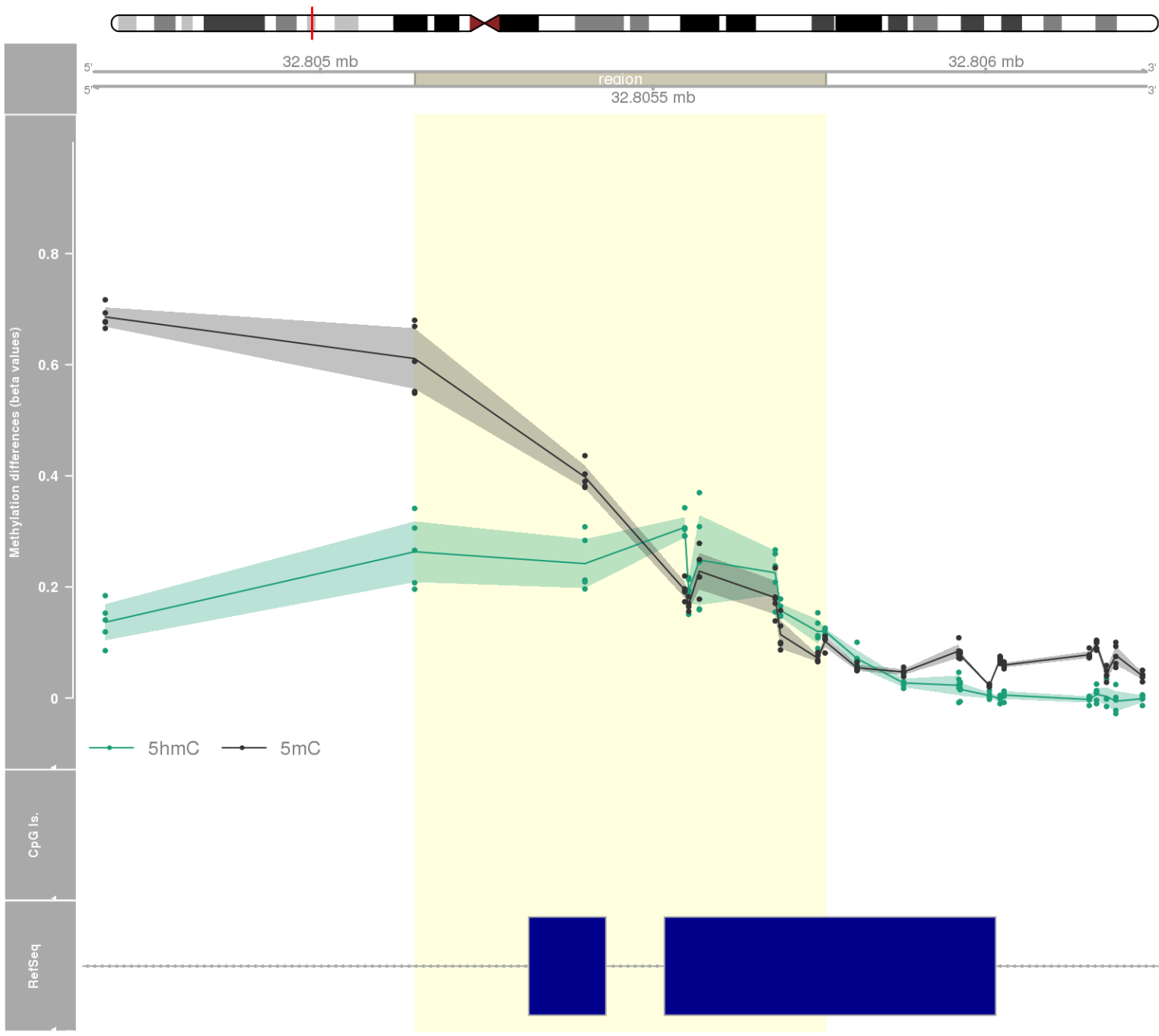
DMR 223 // chr10:106093659-106094166 // 507 pb. (8 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.016 // fwerArea: 1
- genes: ITPRIP -



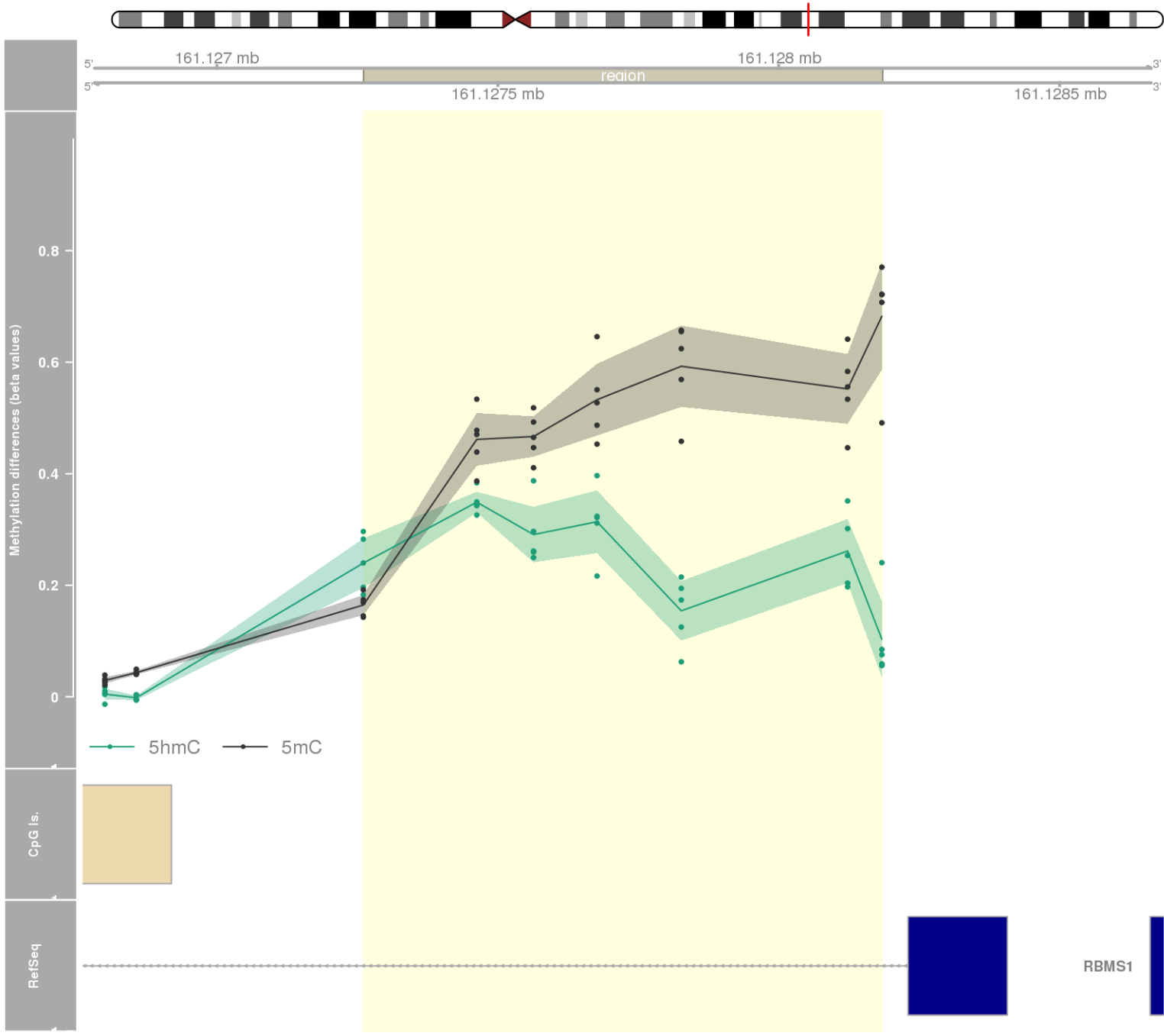
DMR 224 // chr17:25798202-25799447 // 1245 pb. (9 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.016 // fwerArea: 1
- genes: KSR1 -



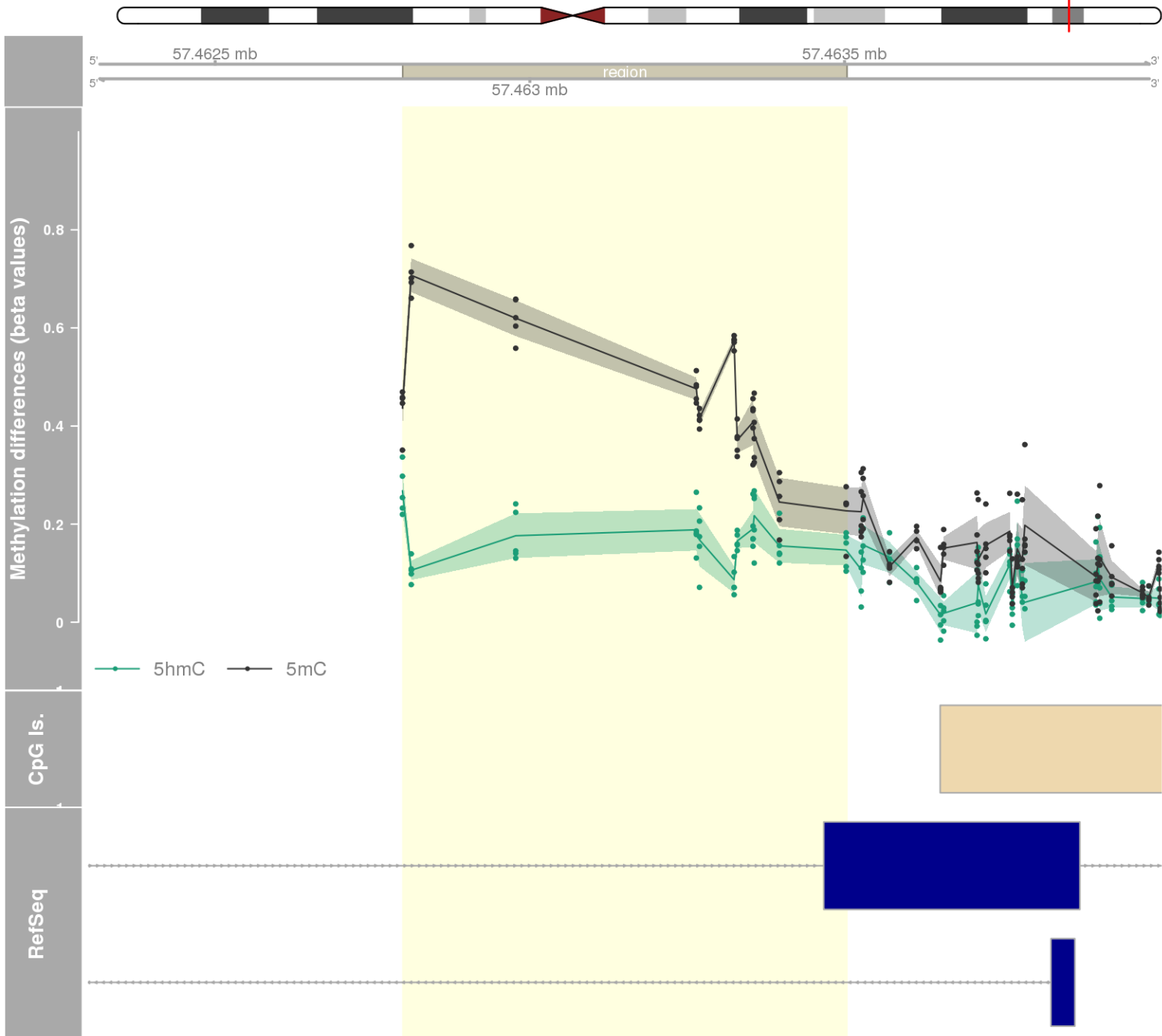
DMR 225 // chr6:32805142-32805759 // 617 pb. (9 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.016 // fwerArea: 1
- genes: TAP2 -



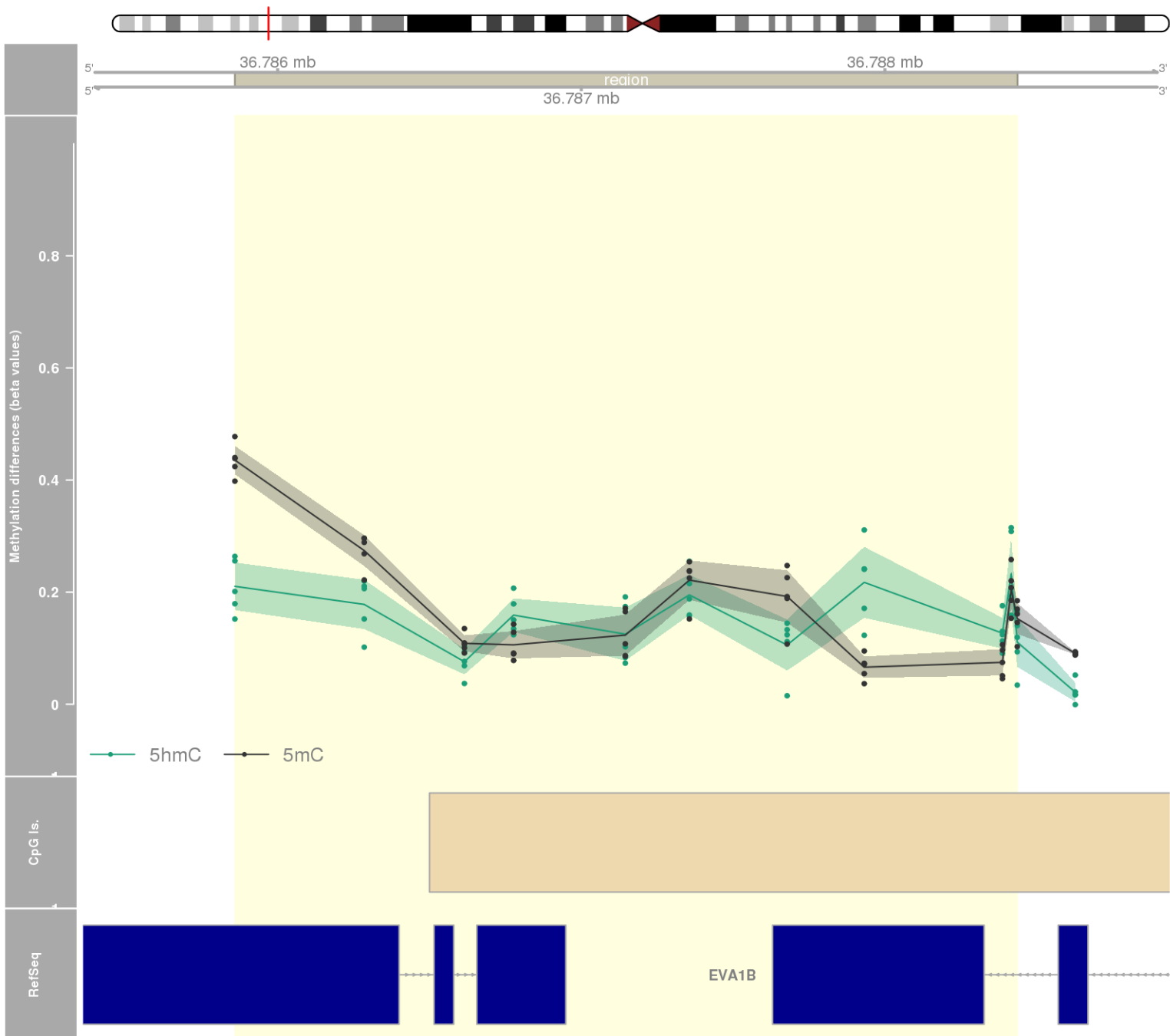
DMR 226 // chr2:161127261-161128184 // 923 pb. (7 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.016 // fwerArea: 1
- genes: ITGB6 -



DMR 227 // chr20:57462798-57463503 // 705 pb. (11 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.016 // fwerArea: 1
- genes: GNAS -



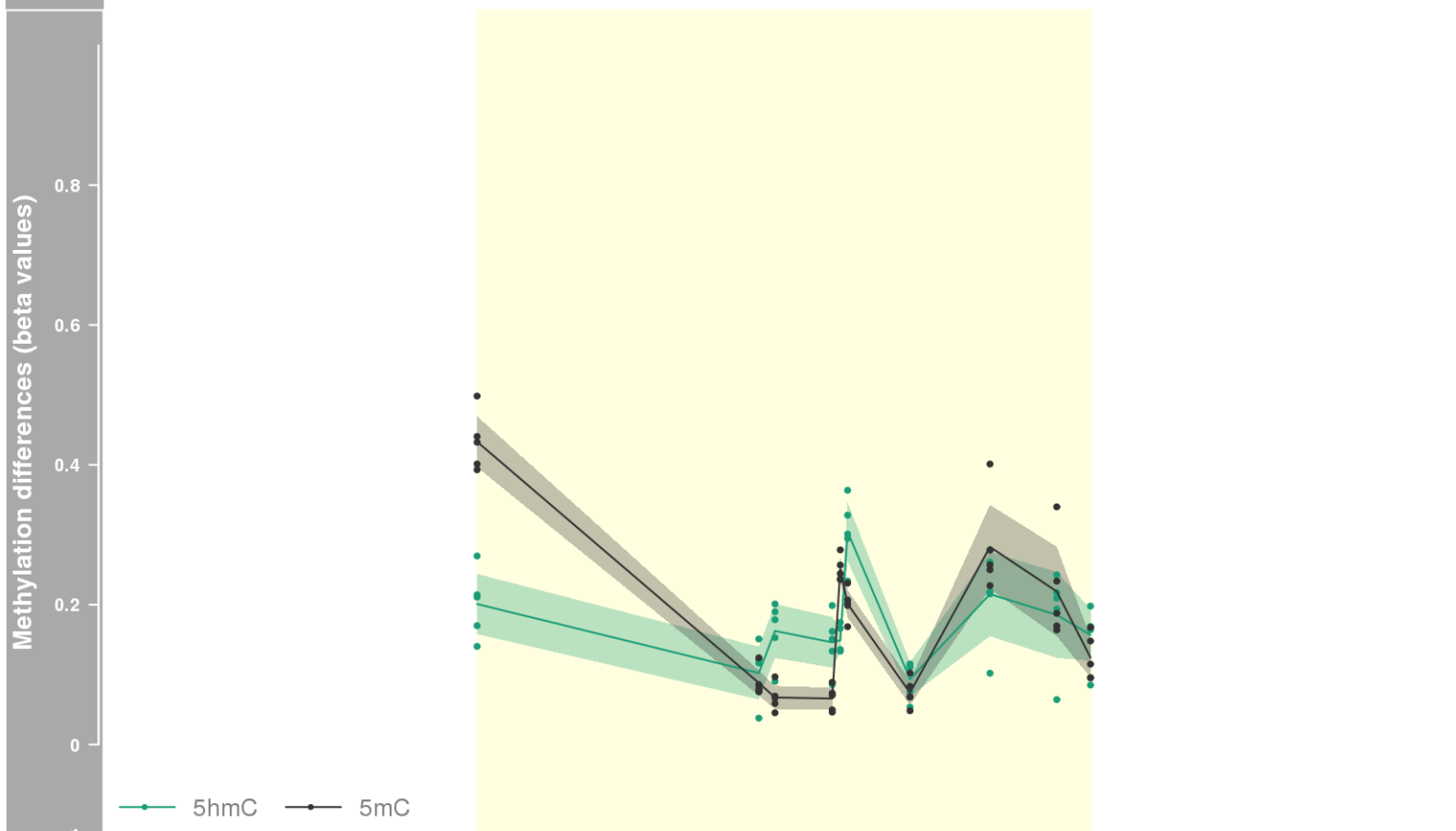
DMR 228 // chr1:36785859-36788436 // 2577 pb. (11 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.016 // fwerArea: 1
- genes: SH3D21 / EVA1B -



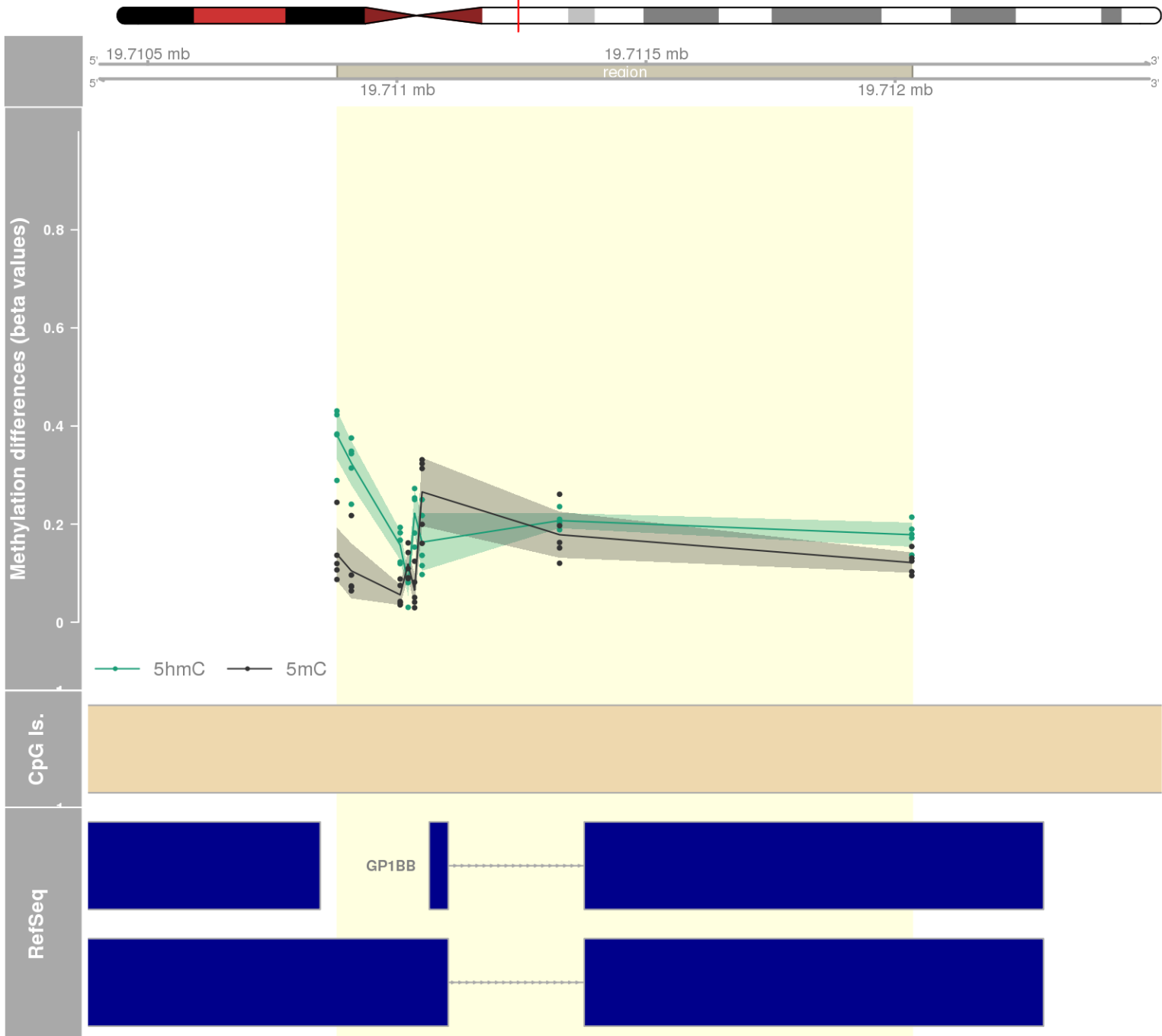
DMR 229 // chr14:24457509-24458345 // 836 pb. (10 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.016 // fwerArea: 1
- genes: DHRS4-AS1 / DHRS4L2 -



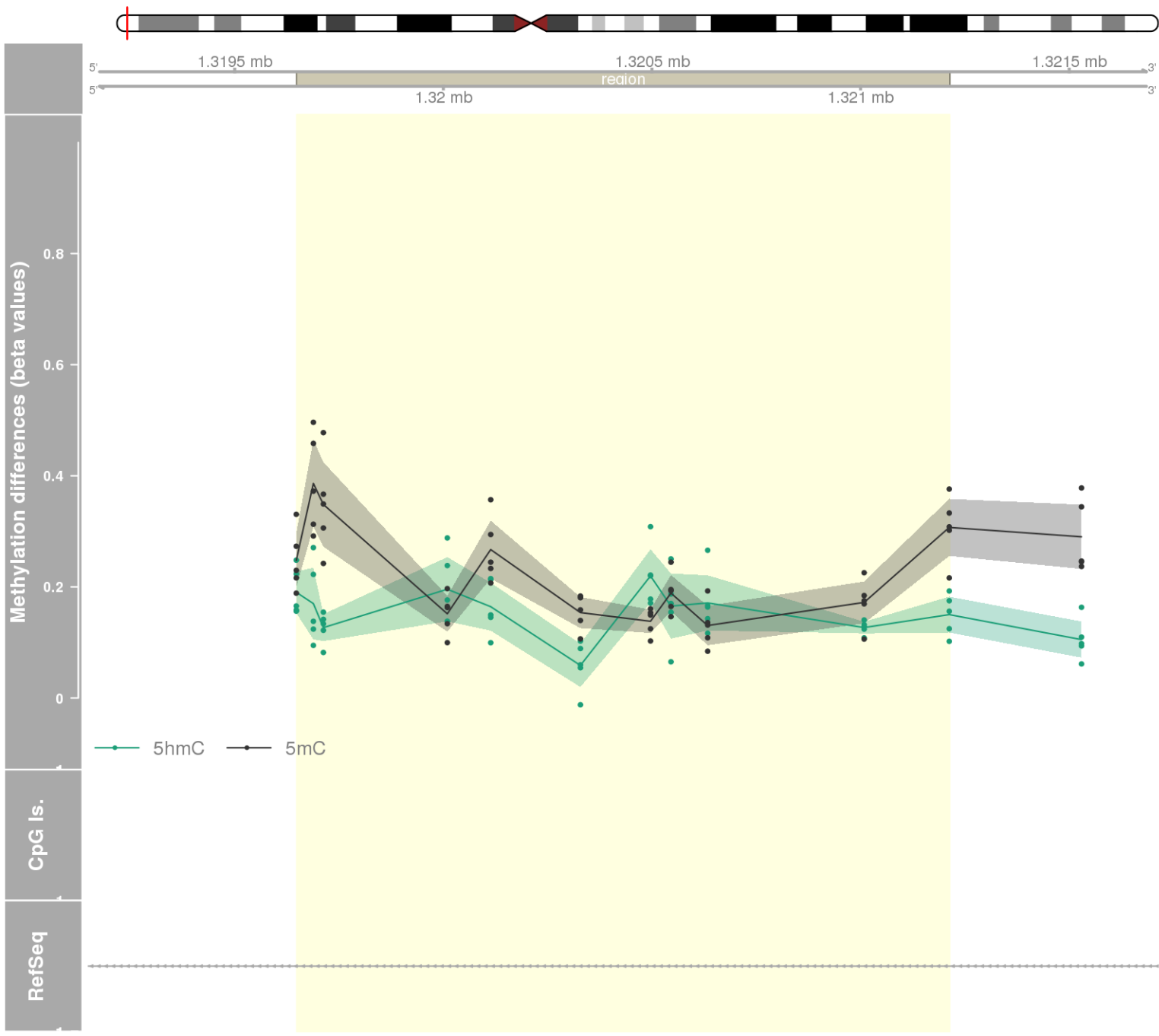
5' 24.4575 mb region 24.4585 mb 3'
5' 24.458 mb 3'



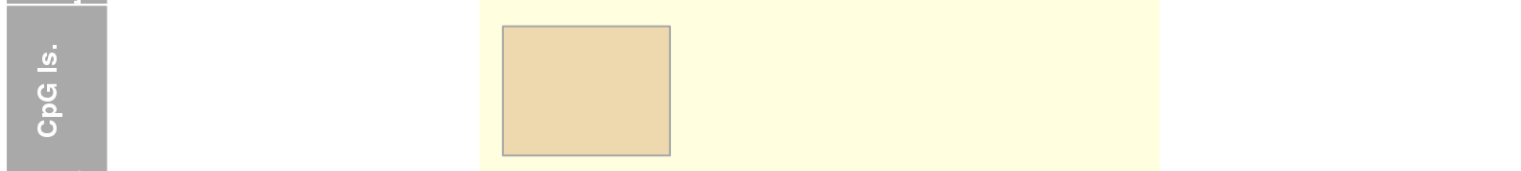
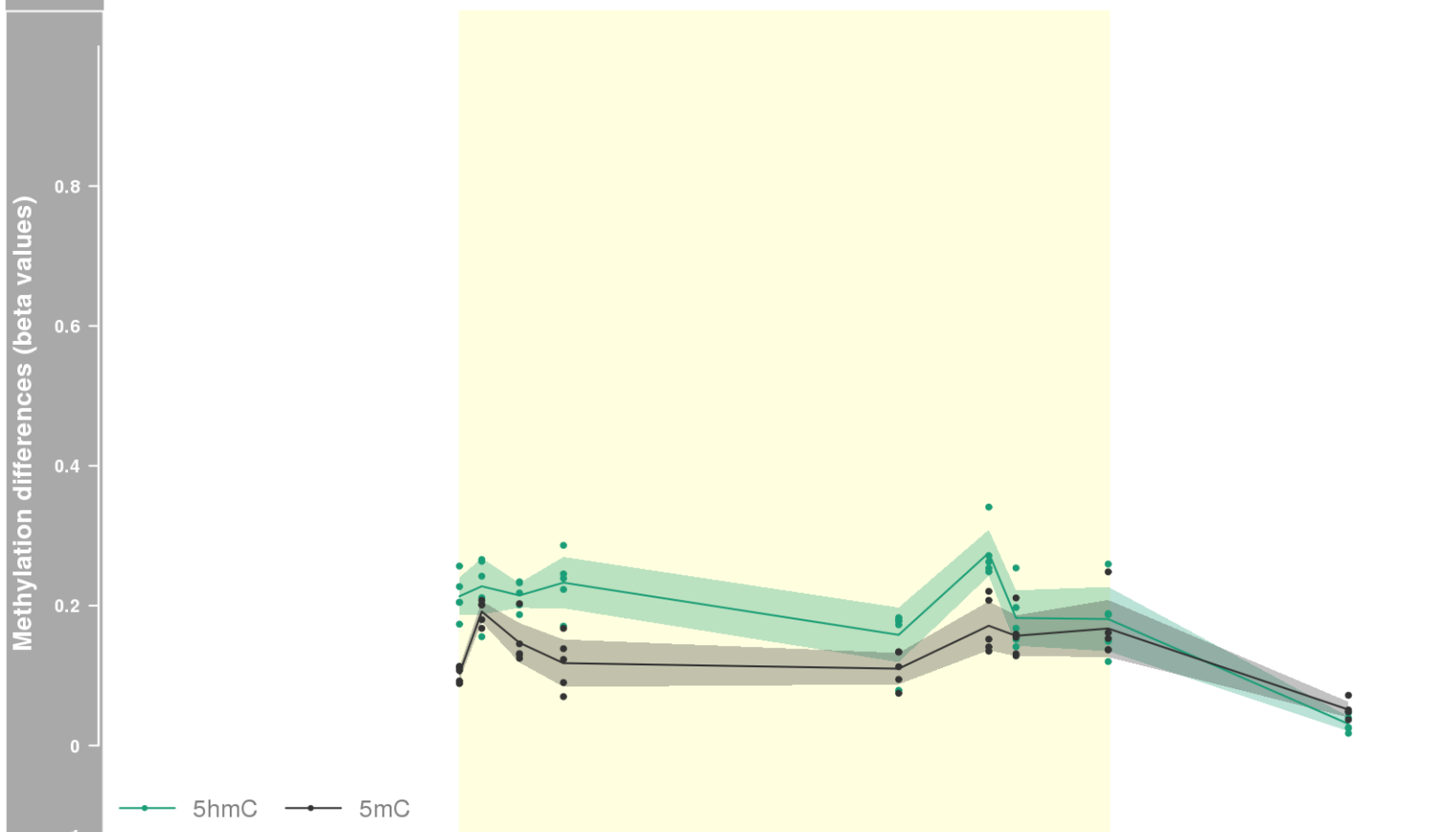
DMR 230 // chr22:19710880-19712034 // 1154 pb. (8 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.016 // fwerArea: 1
- genes: SEPT5 / GP1BB -



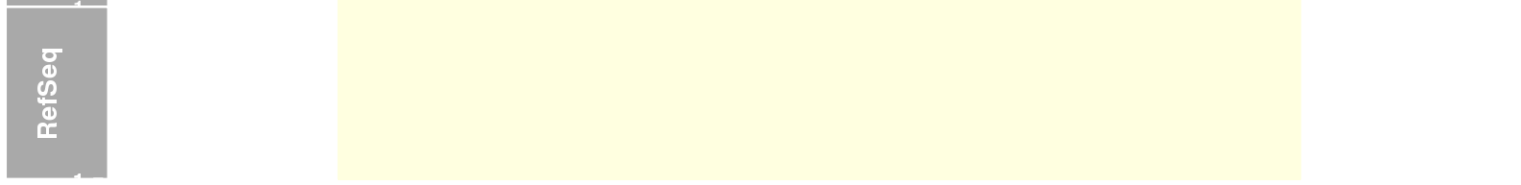
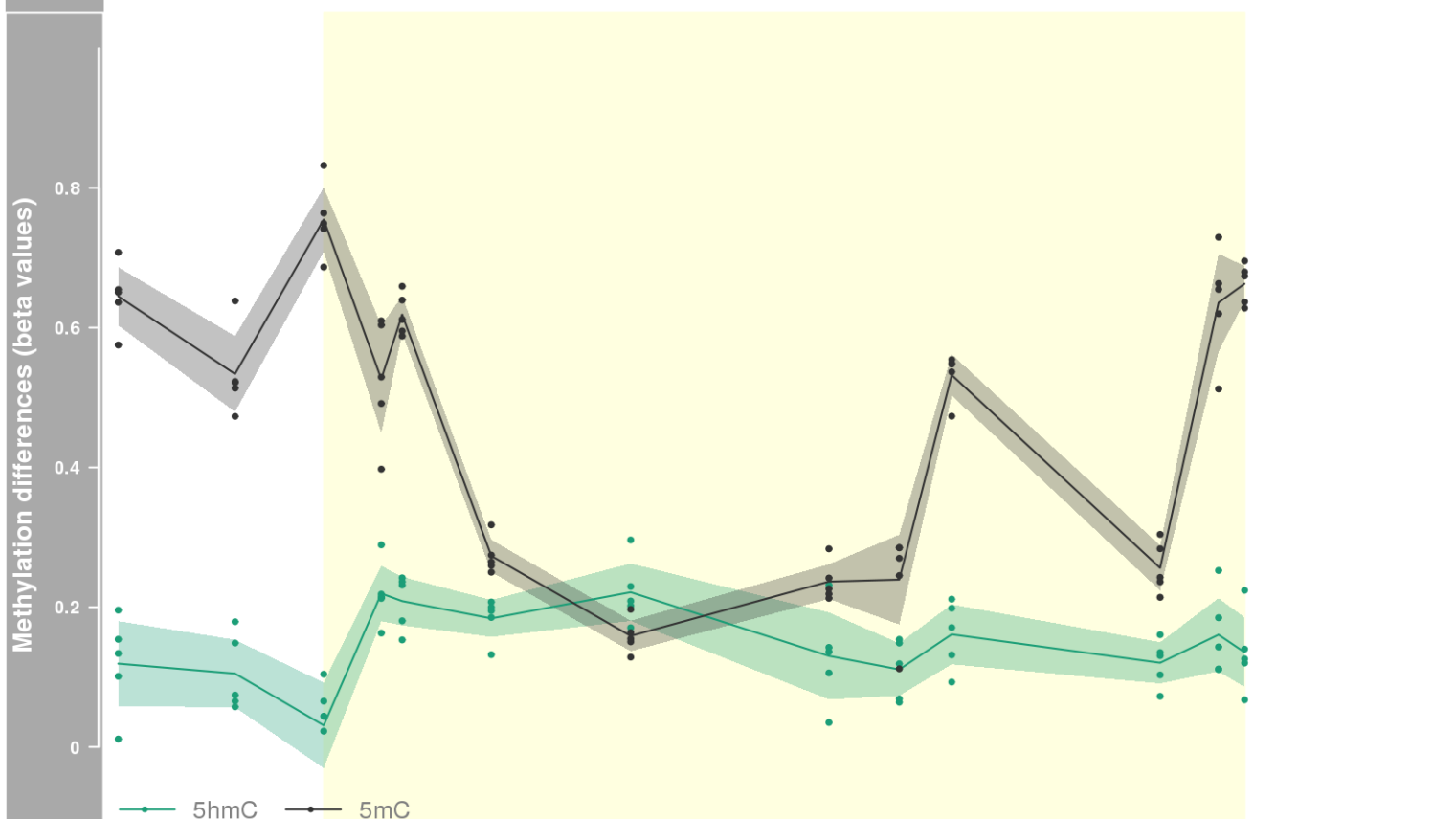
DMR 231 // chr11:1319648-1321213 // 1565 pb. (11 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.016 // fwerArea: 1
- genes: TOLLIP -



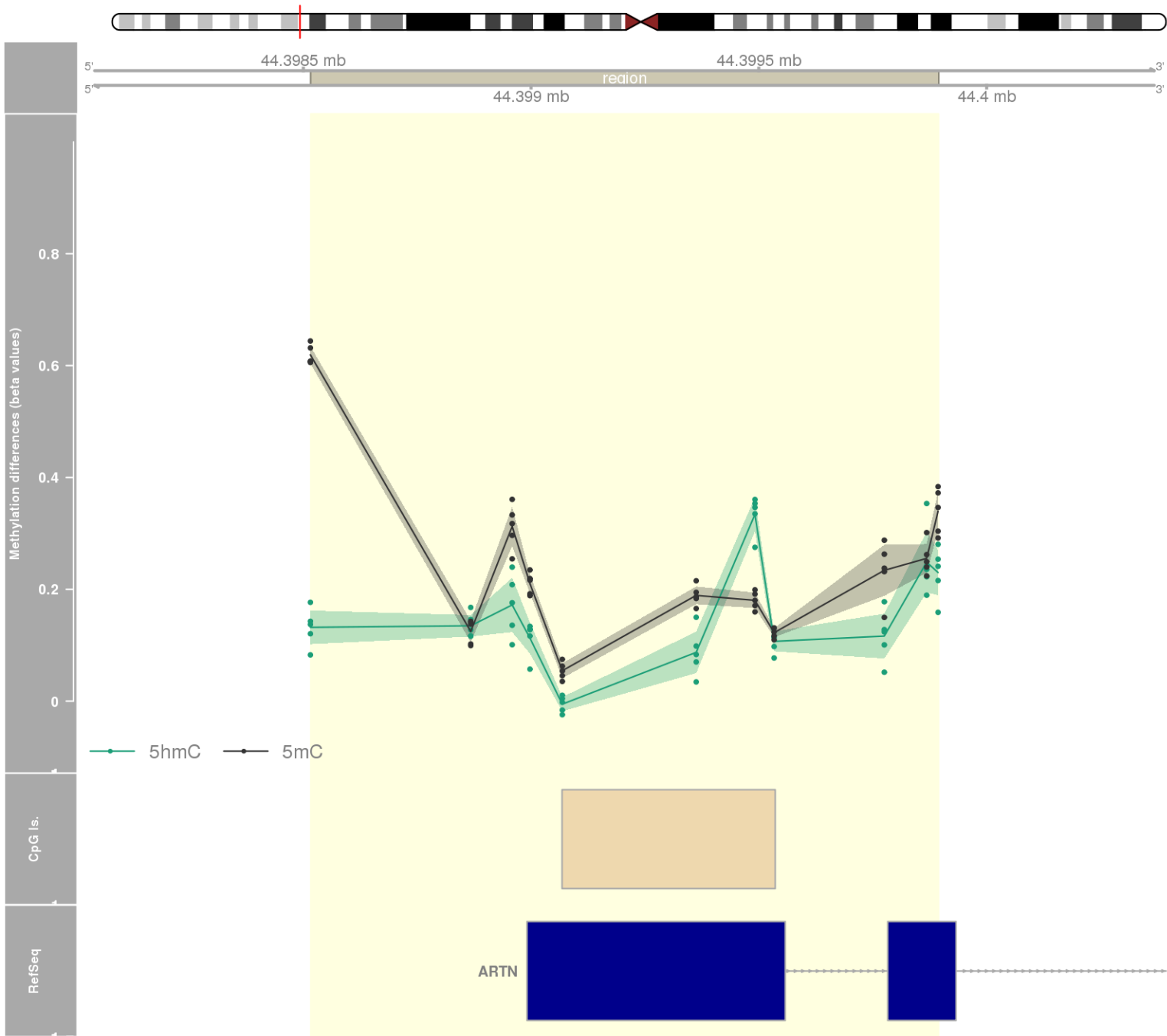
DMR 232 // chr10:3823758-3824687 // 929 pb. (8 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.016 // fwerArea: 1
- genes: KLF6 -



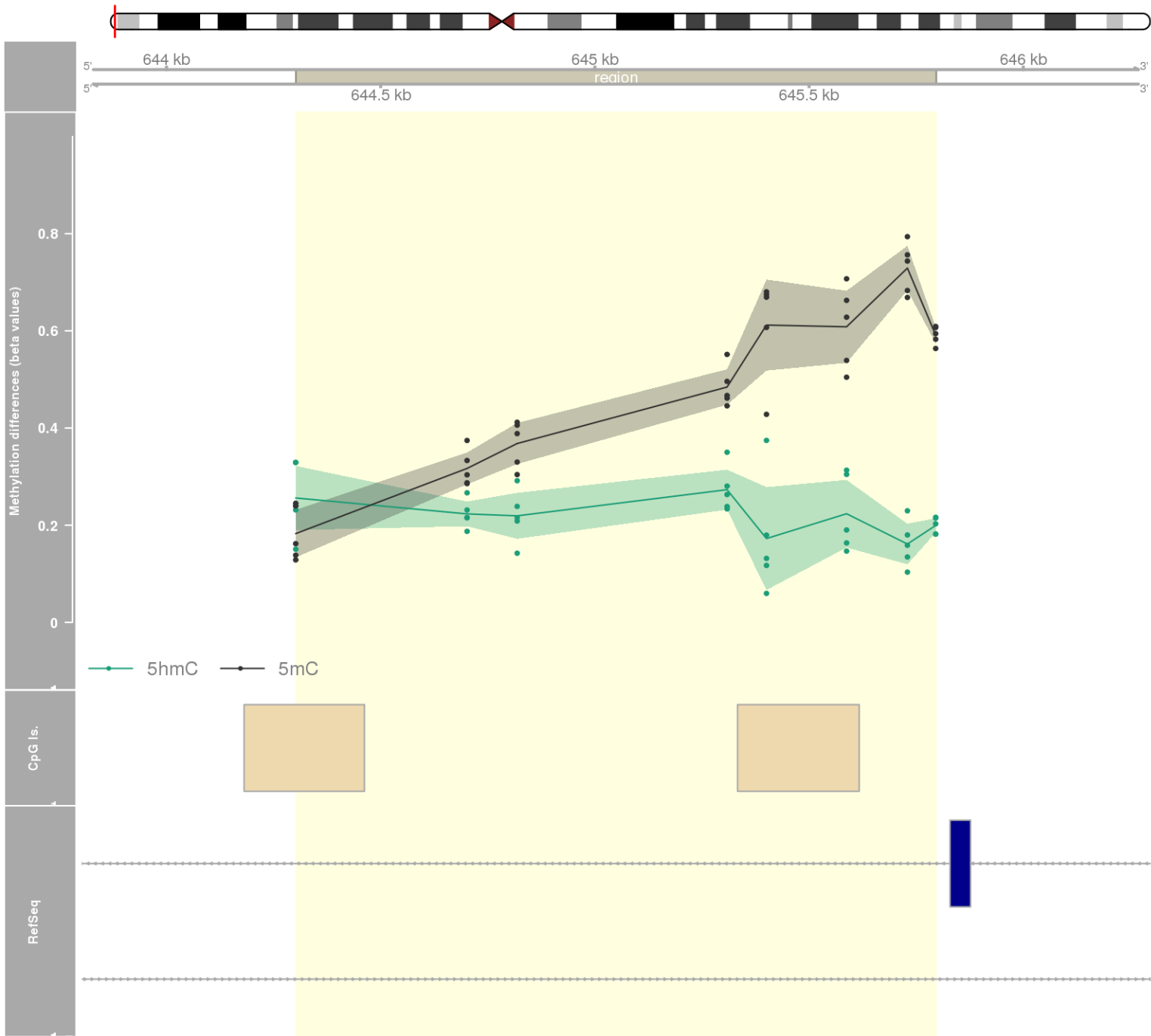
DMR 233 // chr11:1989715-1991875 // 2160 pb. (11 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.016 // fwerArea: 1
- genes: MRPL23 -



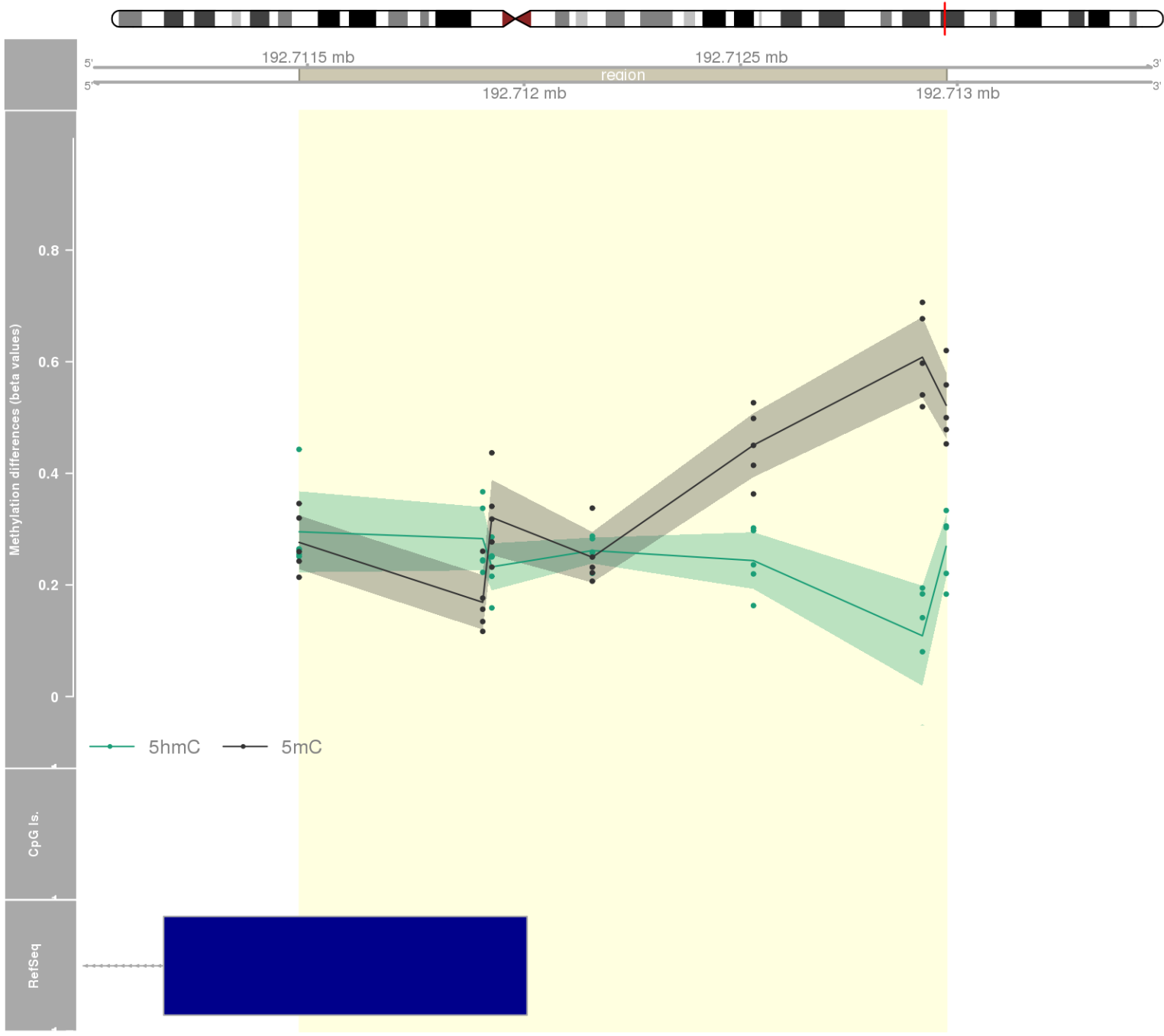
DMR 234 // chr1:44398516-44399894 // 1378 pb. (11 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.016 // fwerArea: 1
- genes: ARTN -



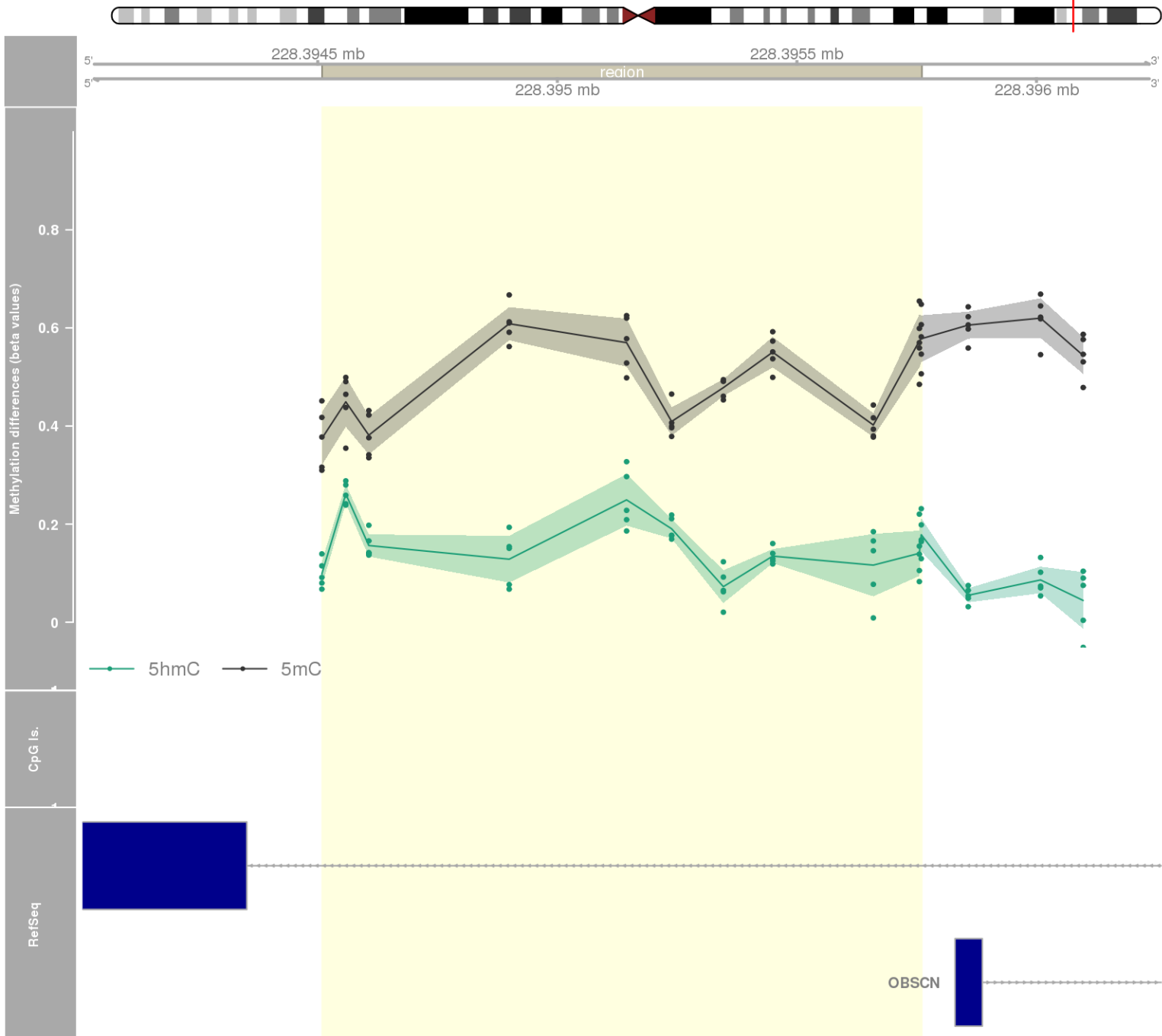
DMR 235 // chr7:644302-645796 // 1494 pb. (8 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.016 // fwerArea: 1
- genes: PRKAR1B -



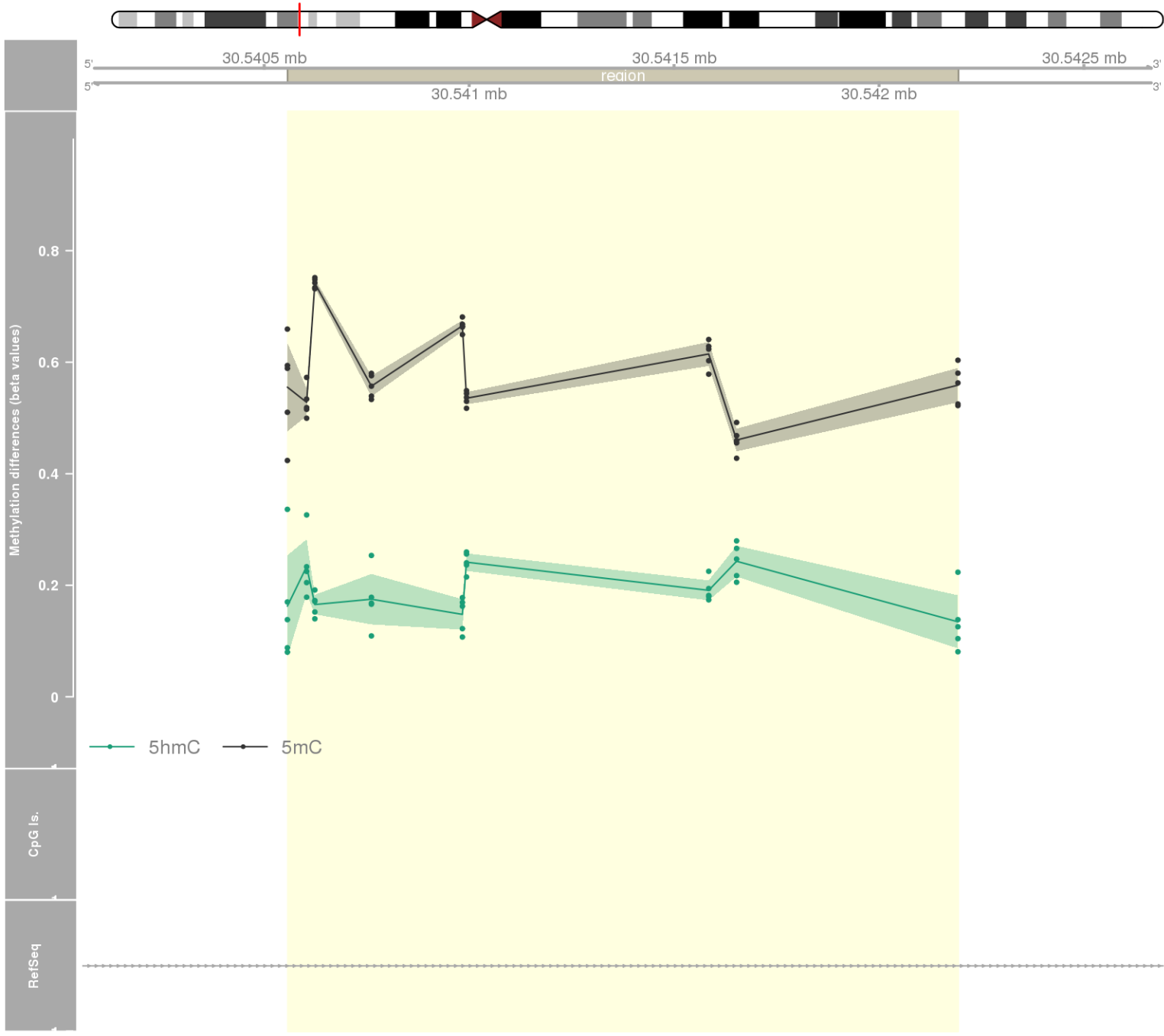
DMR 236 // chr2:192711481-192712975 // 1494 pb. (7 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.017 // fwerArea: 1
- genes: SDPR -



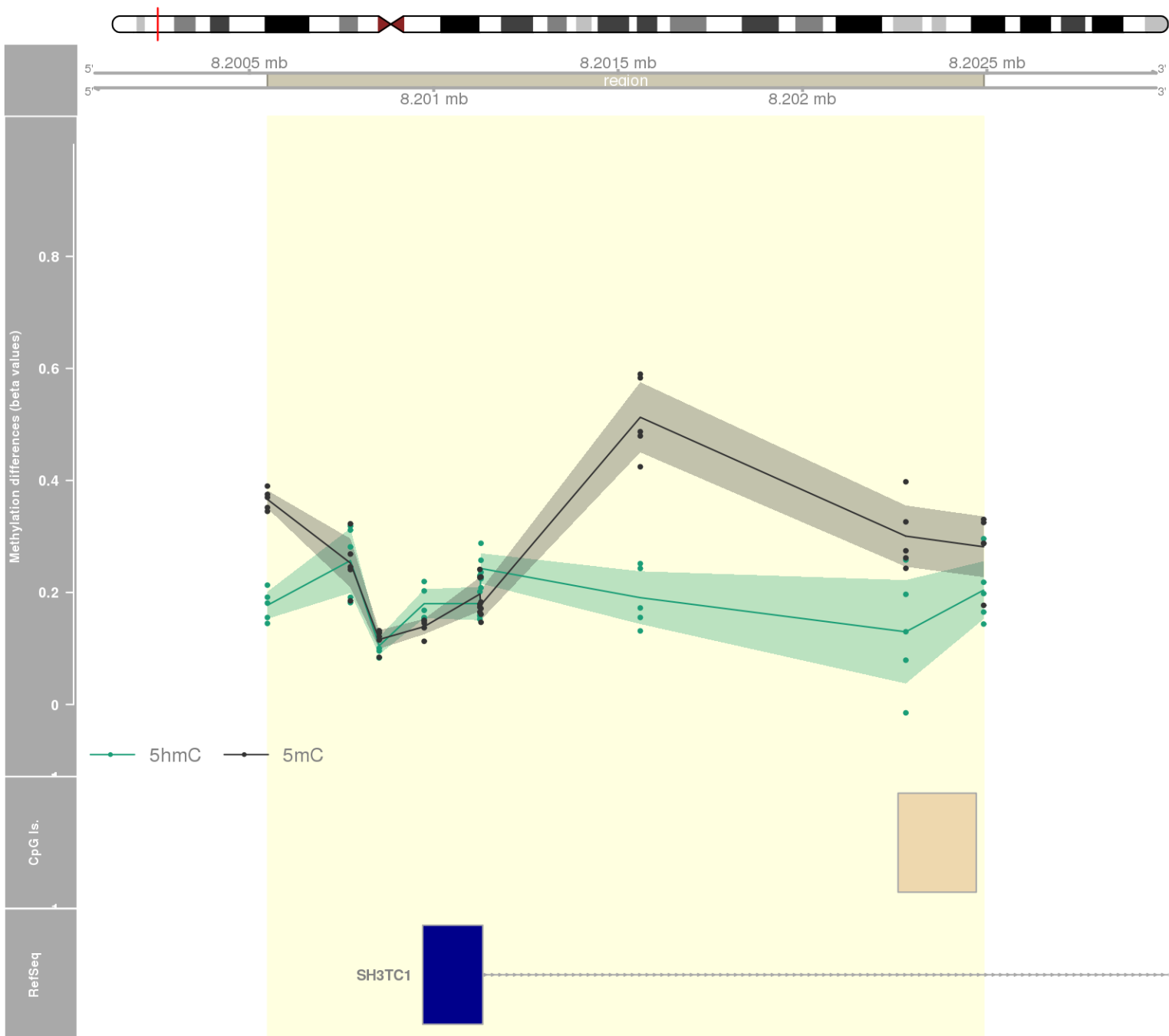
DMR 237 // chr1:228394509-228395760 // 1251 pb. (11 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.017 // fwerArea: 1
- genes: C1orf145 -



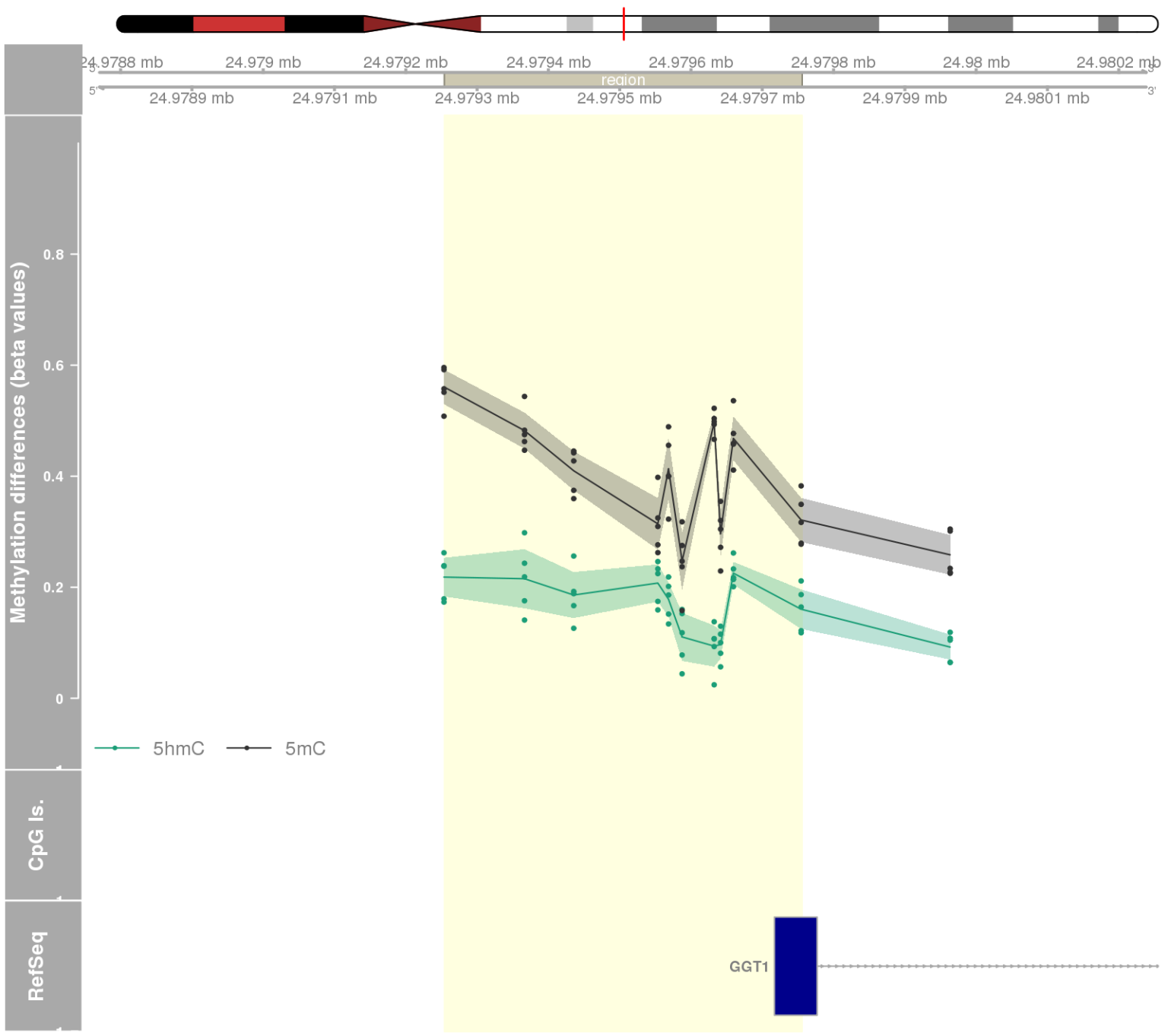
DMR 238 // chr6:30540557-30542193 // 1636 pb. (9 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.017 // fwerArea: 1
- genes: ABCF1 -



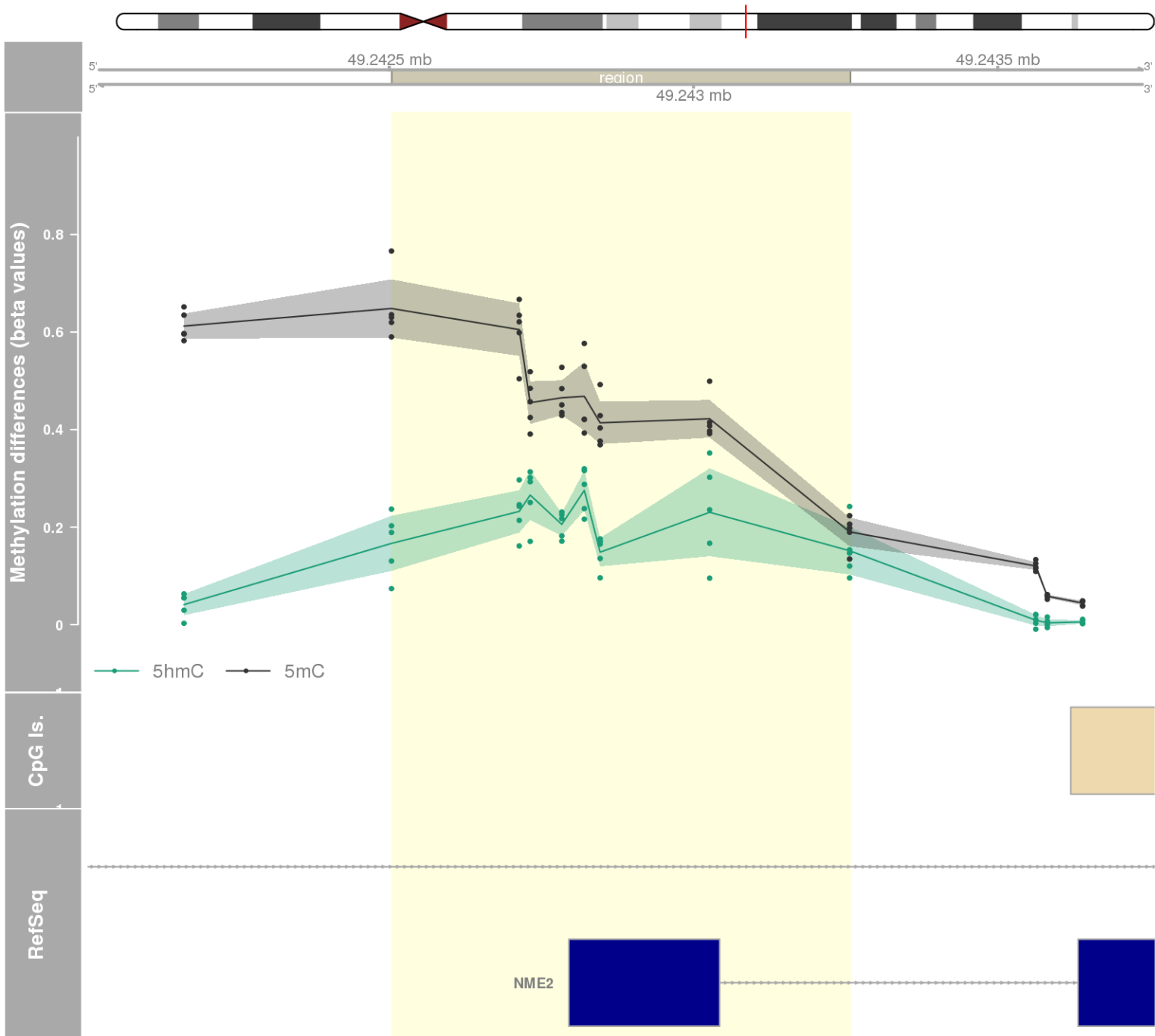
DMR 239 // chr4:8200549-8202491 // 1942 pb. (9 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.017 // fwerArea: 1
- genes: SH3TC1 -



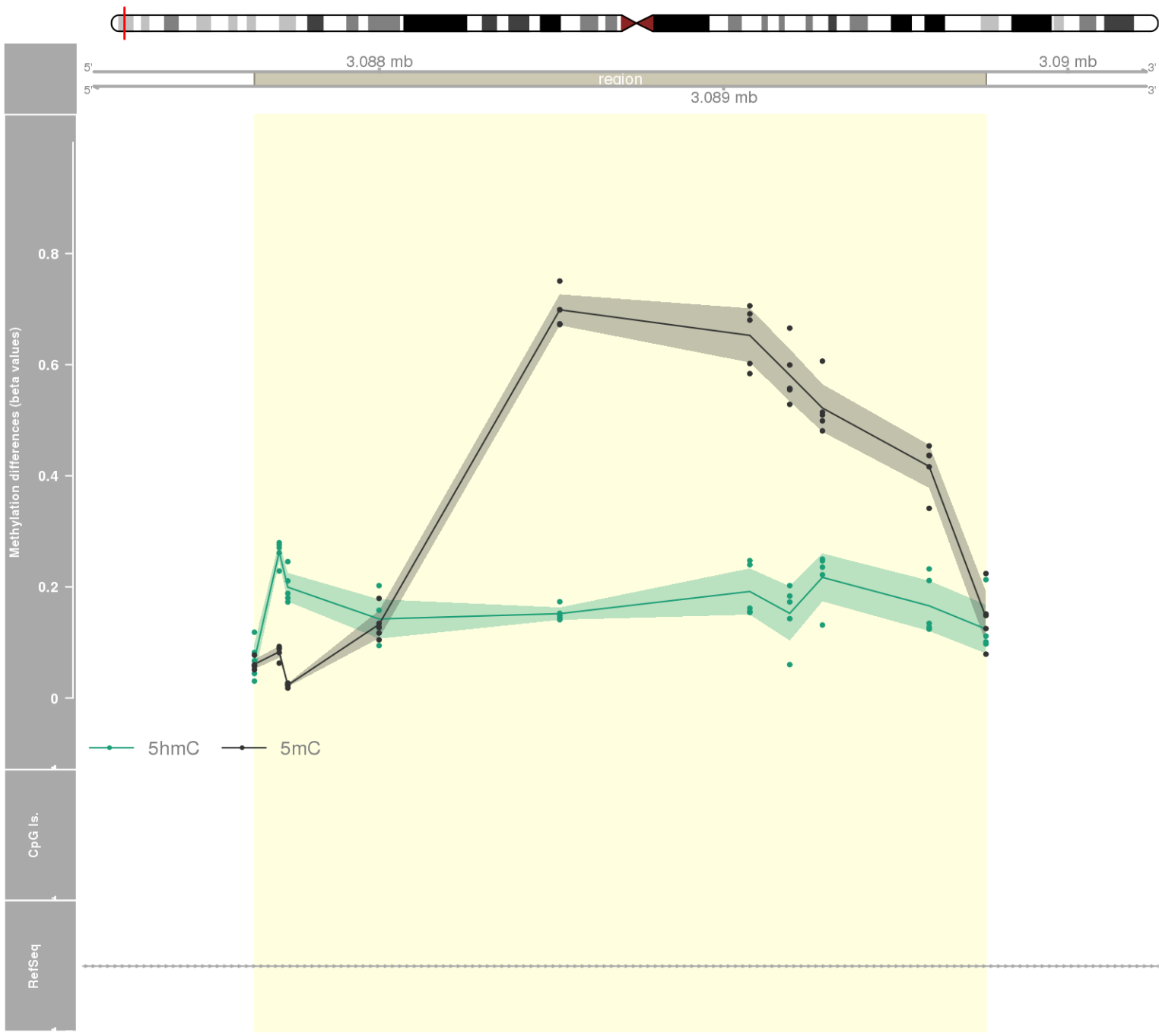
DMR 240 // chr22:24979254-24979755 // 501 pb. (10 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.017 // fwerArea: 1
- genes: SNRPD3 / GGT1 -



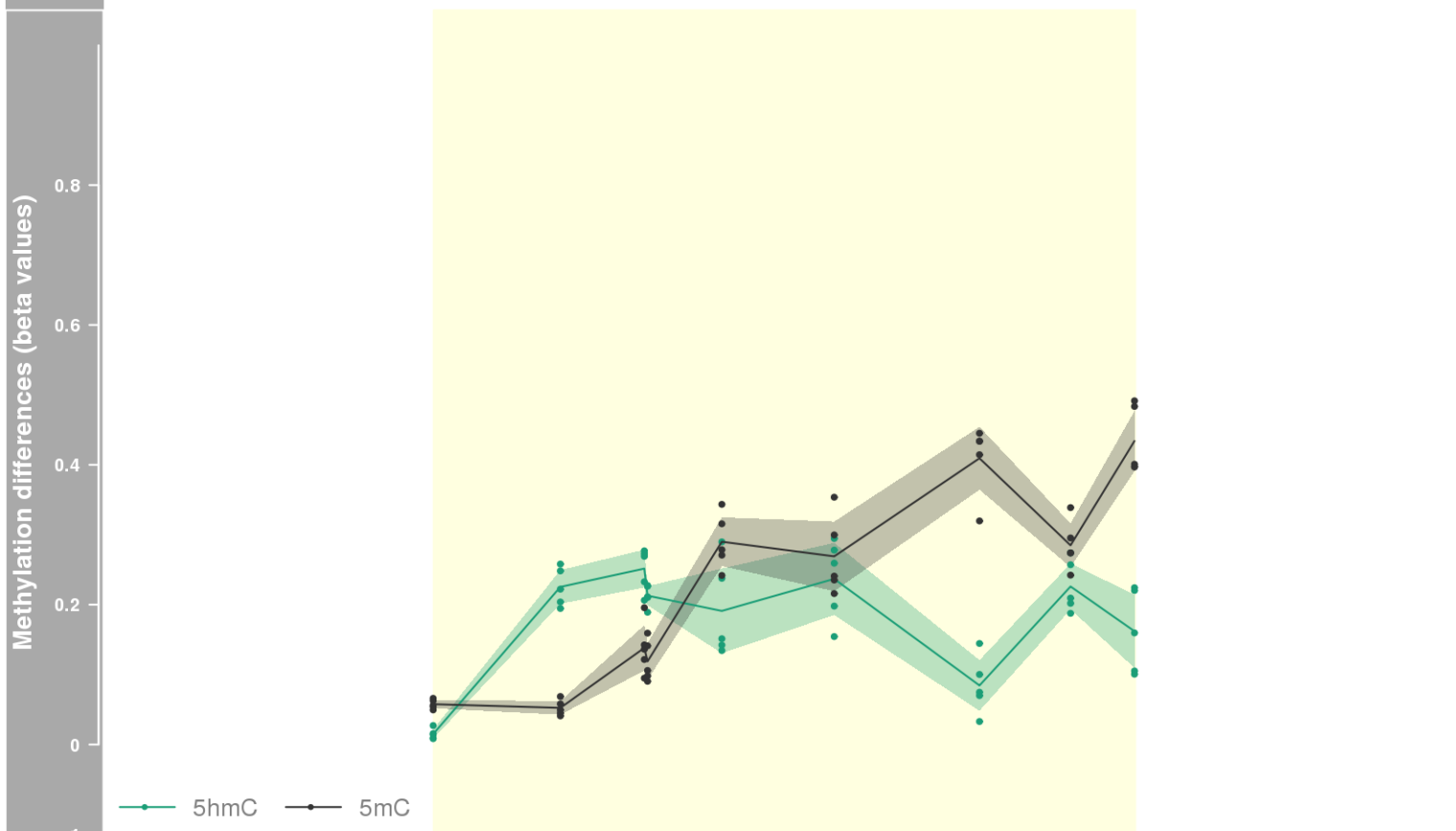
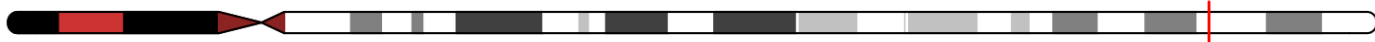
DMR 241 // chr17:49242504-49243257 // 753 pb. (8 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.017 // fwerArea: 1
- genes: NME1-NME2 / NME2 -



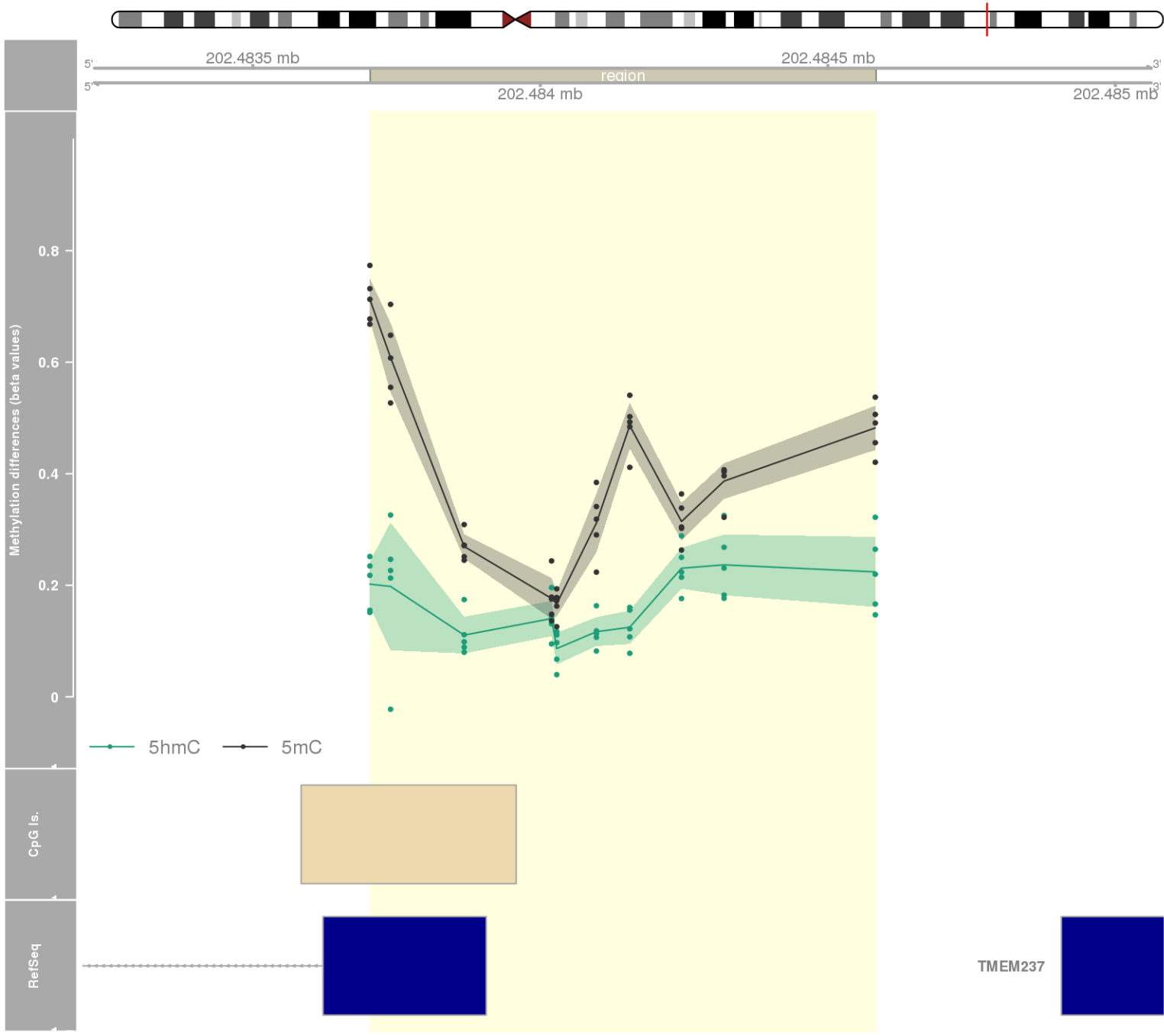
DMR 242 // chr1:3087637-3089762 // 2125 pb. (10 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.017 // fwerArea: 1
- genes: PRDM16 -



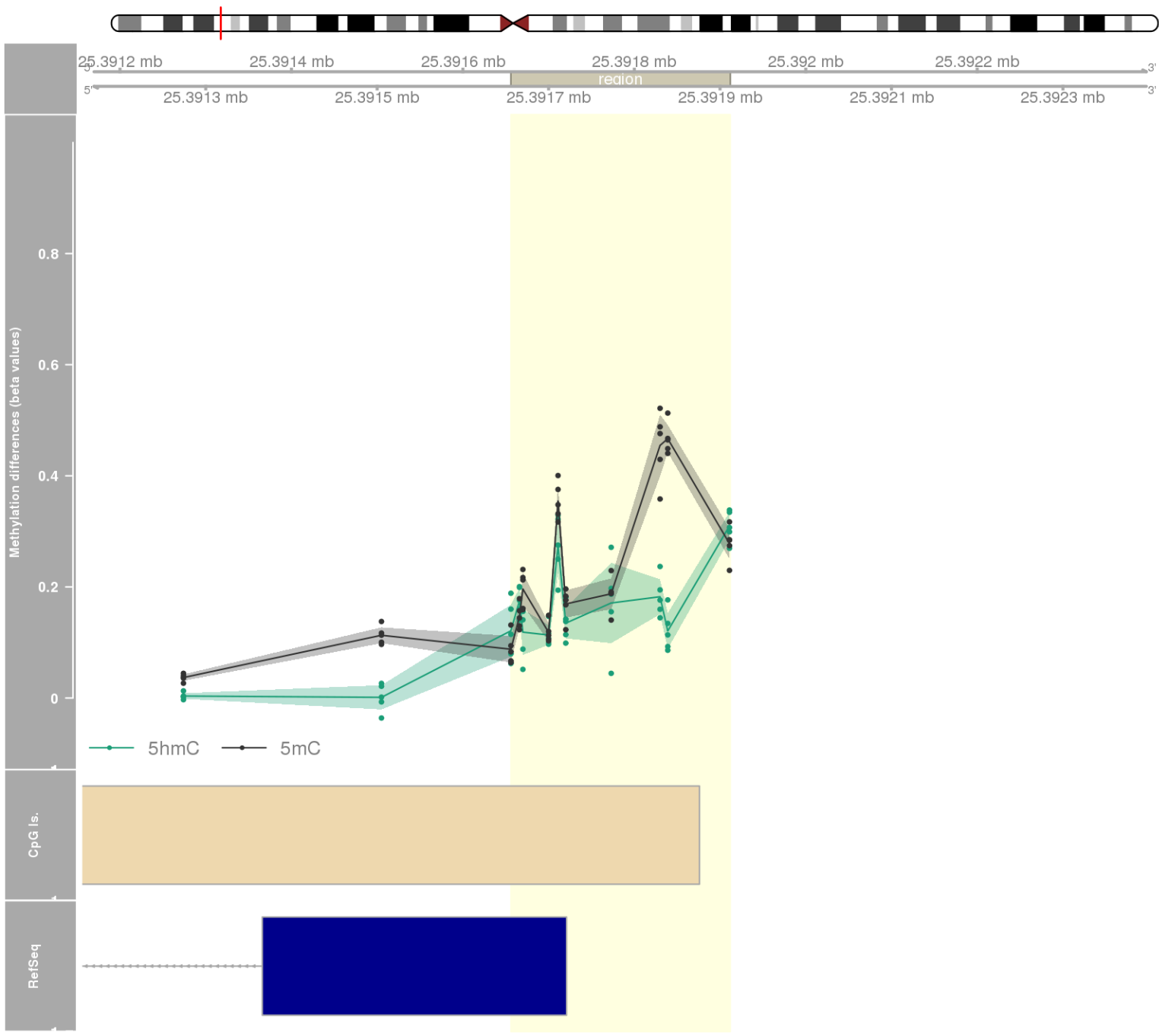
DMR 243 // chr15:89959657-89960743 // 1086 pb. (9 probes) // pvalue: 0.005 // fwer: 0.36 // pvalueArea: 0.017 // fwerArea: 1



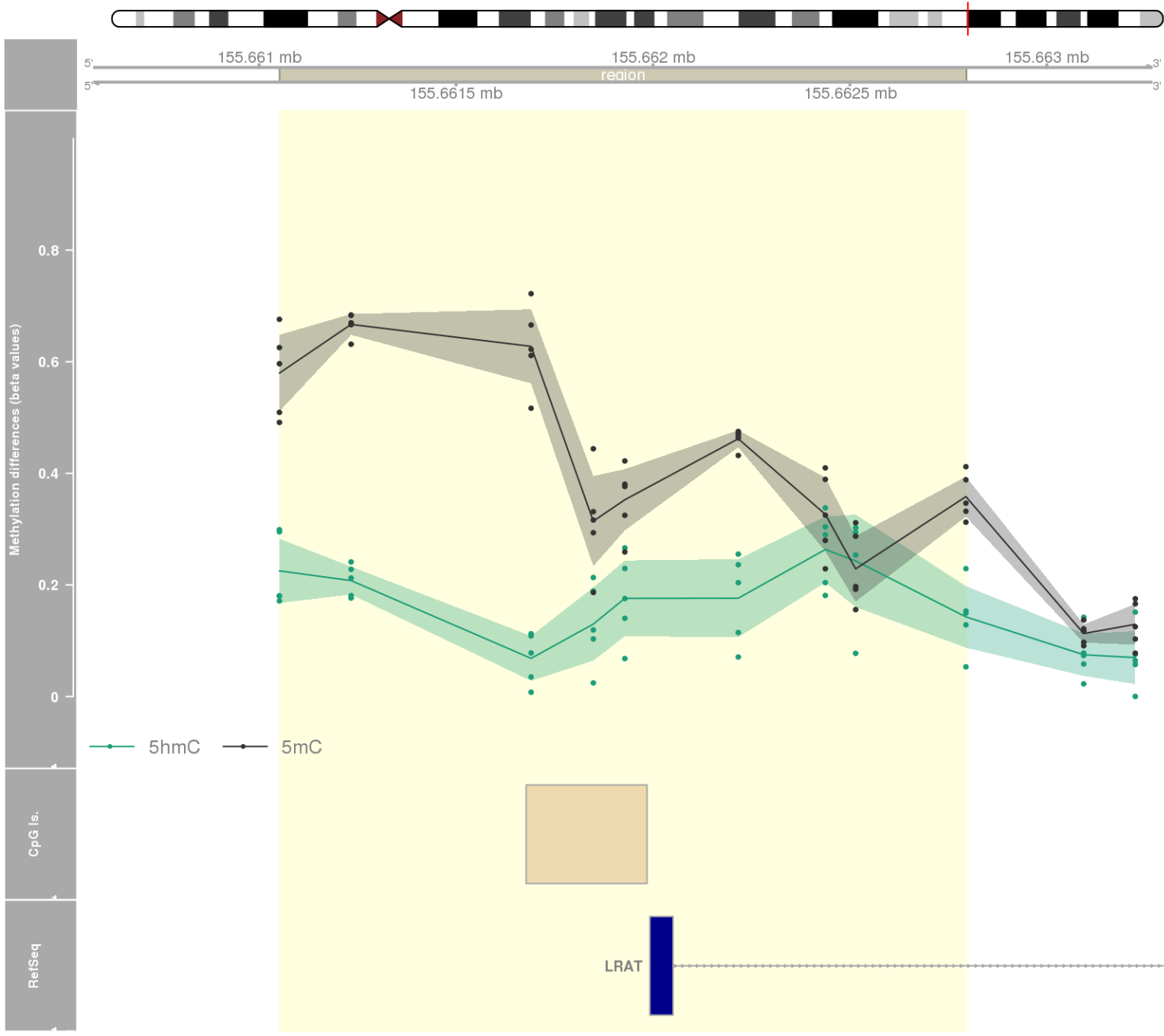
DMR 244 // chr2:202483704-202484583 // 879 pb. (10 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.017 // fwerArea: 1
- genes: ALS2CR11 -



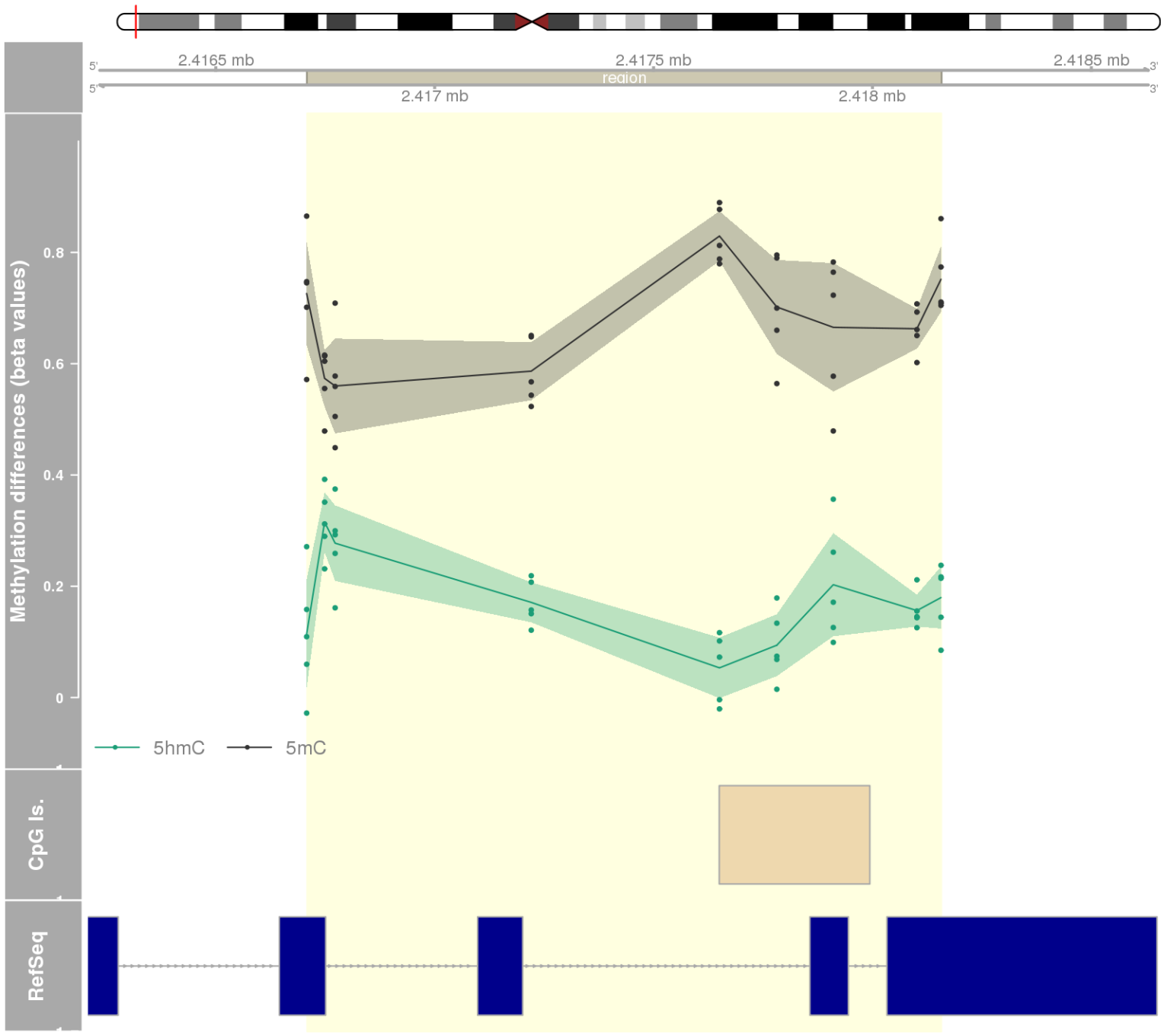
DMR 245 // chr2:25391656-25391911 // 255 pb. (10 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.017 // fwerArea: 1
- genes: POMC -



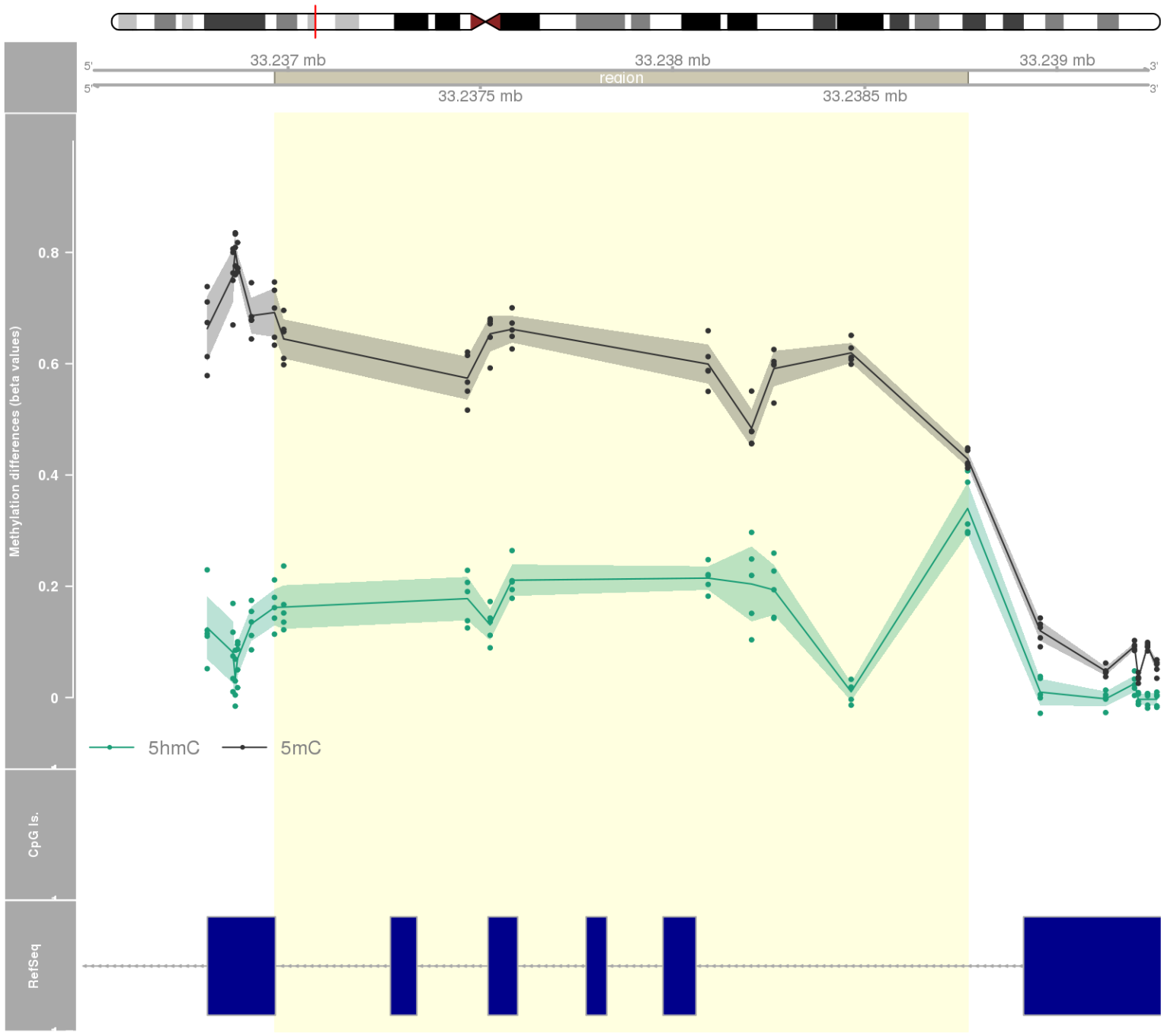
DMR 246 // chr4:155661052-155662795 // 1743 pb. (9 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.017 // fwerArea: 1
- genes: LRAT -



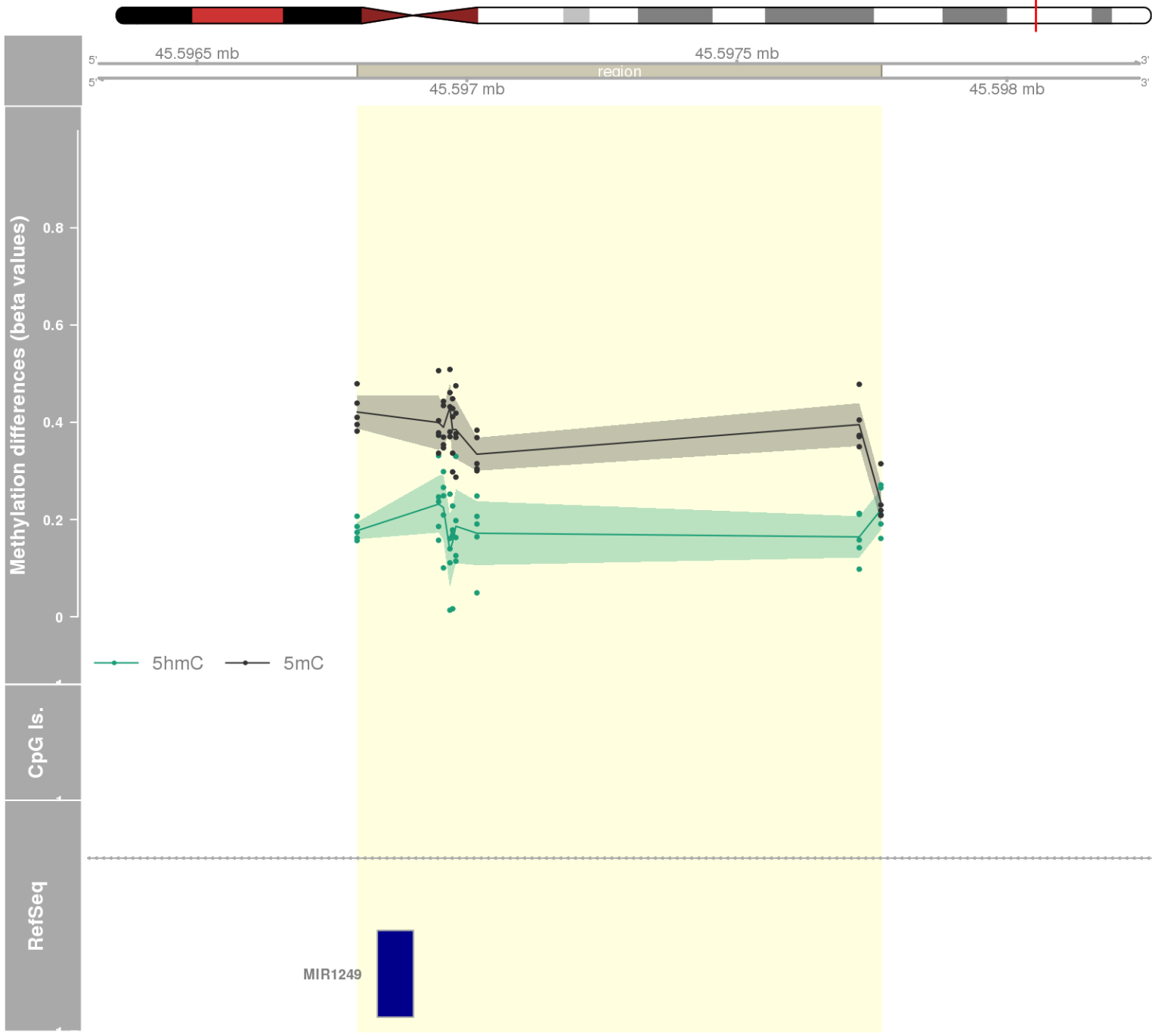
DMR 247 // chr11:2416708-2418157 // 1449 pb. (9 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.017 // fwerArea: 1
- genes: CD81 -



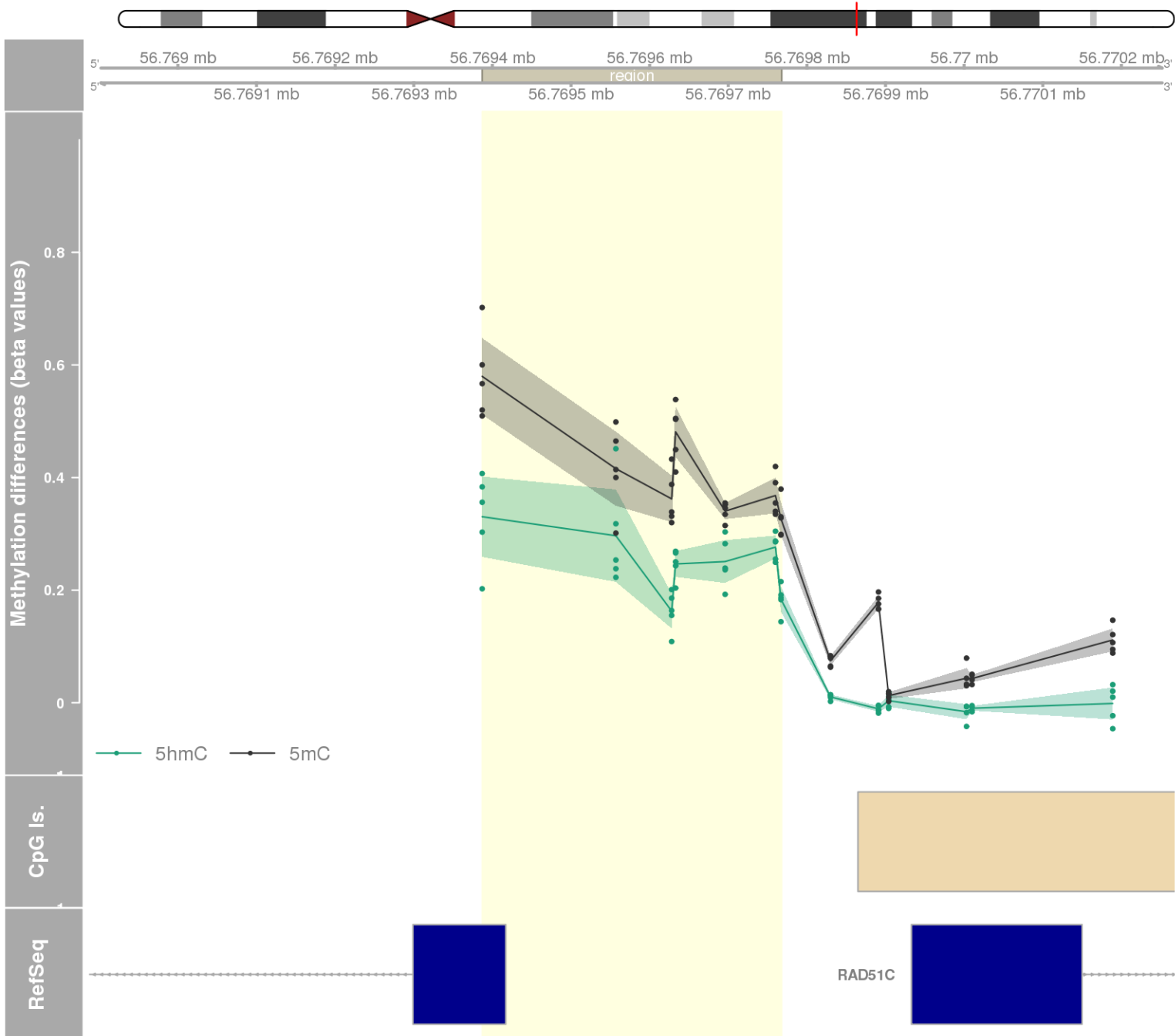
DMR 248 // chr6:33236965-33238768 // 1803 pb. (10 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.017 // fwerArea: 1
- genes: VPS52 -



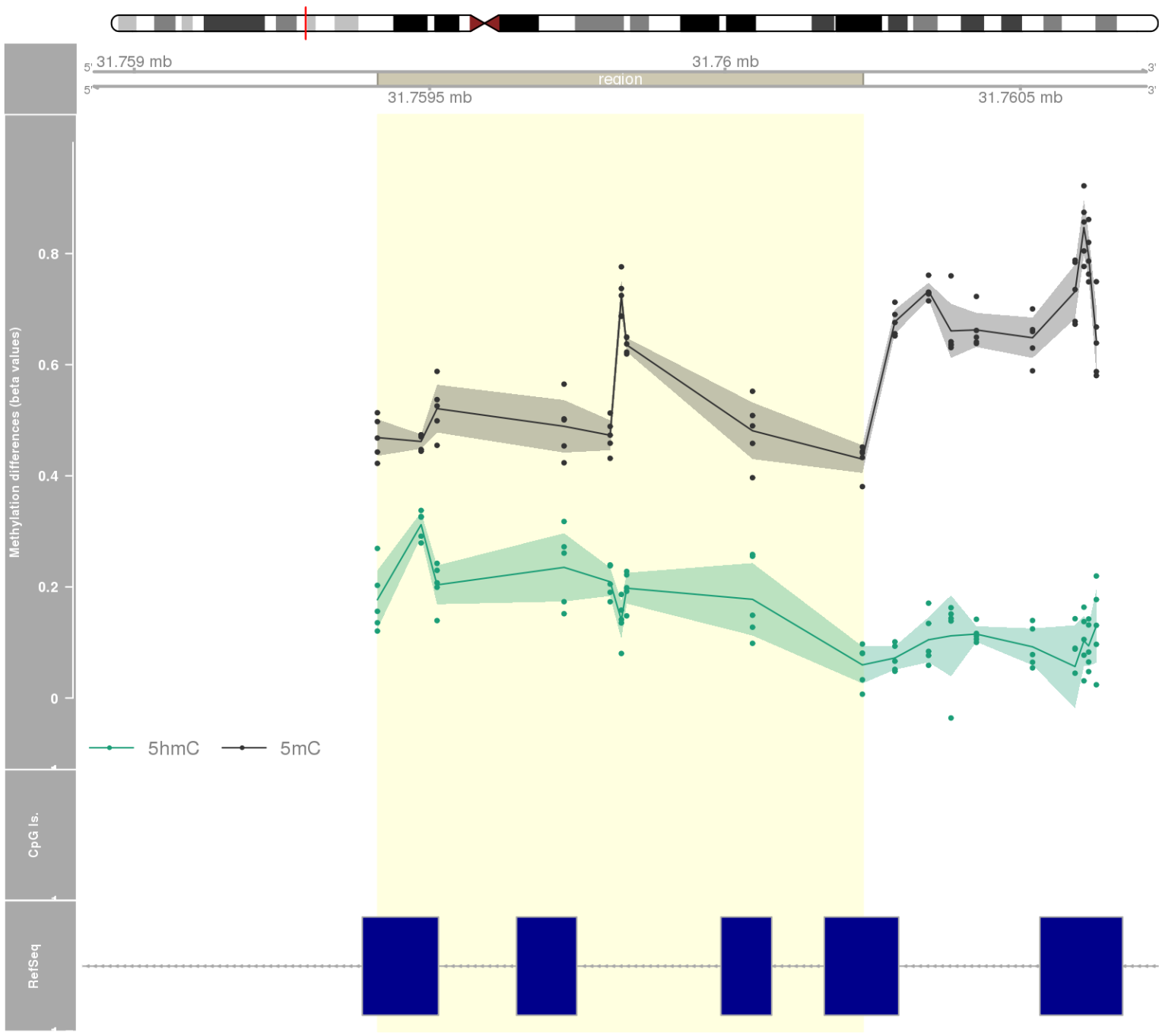
DMR 249 // chr22:45596797-45597767 // 970 pb. (9 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.017 // fwerArea: 1
- genes: MIR1249 / KIAA0930 -



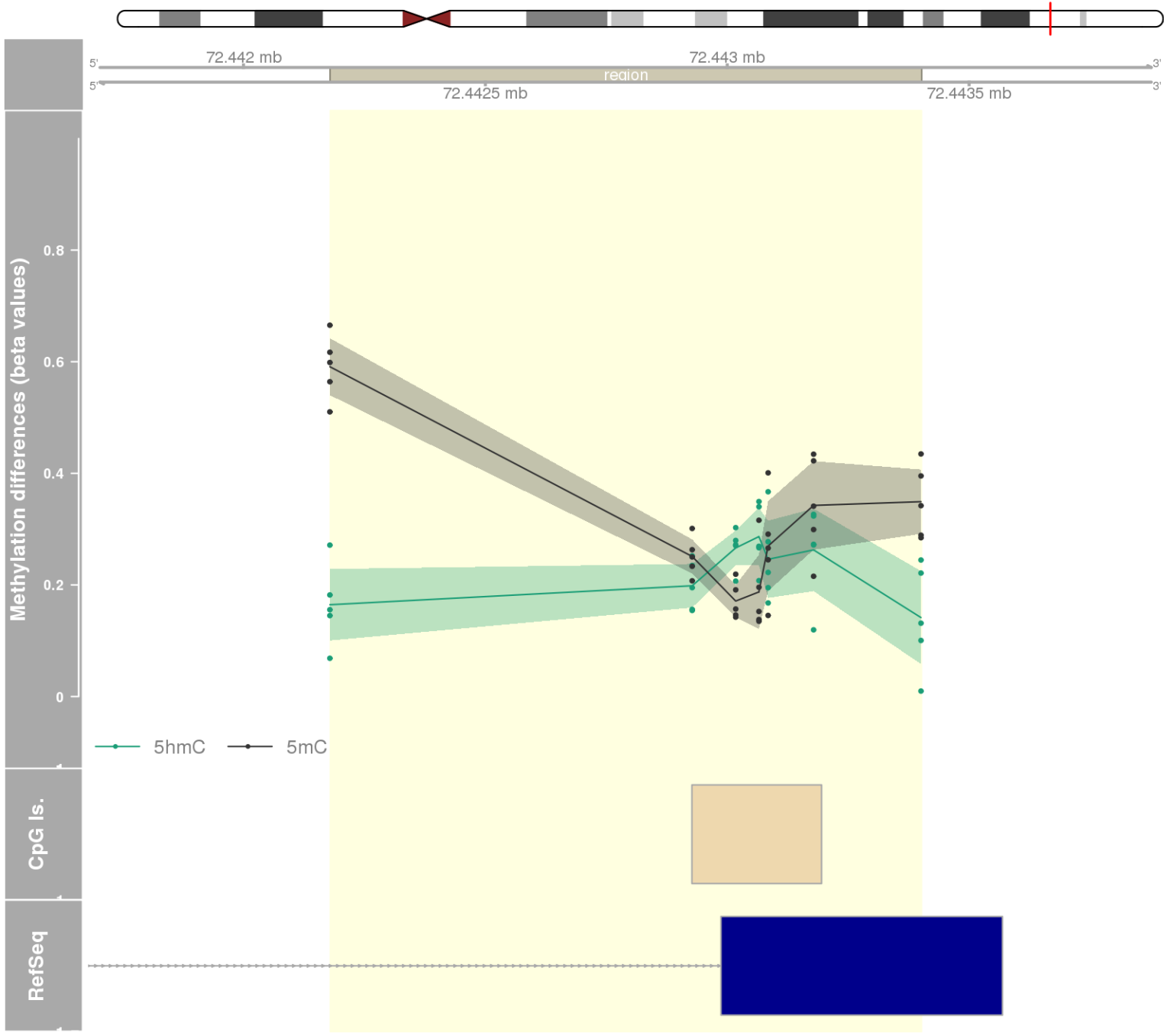
DMR 250 // chr17:56769387-56769767 // 380 pb. (7 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.018 // fwerArea: 1
- genes: TEX14 -



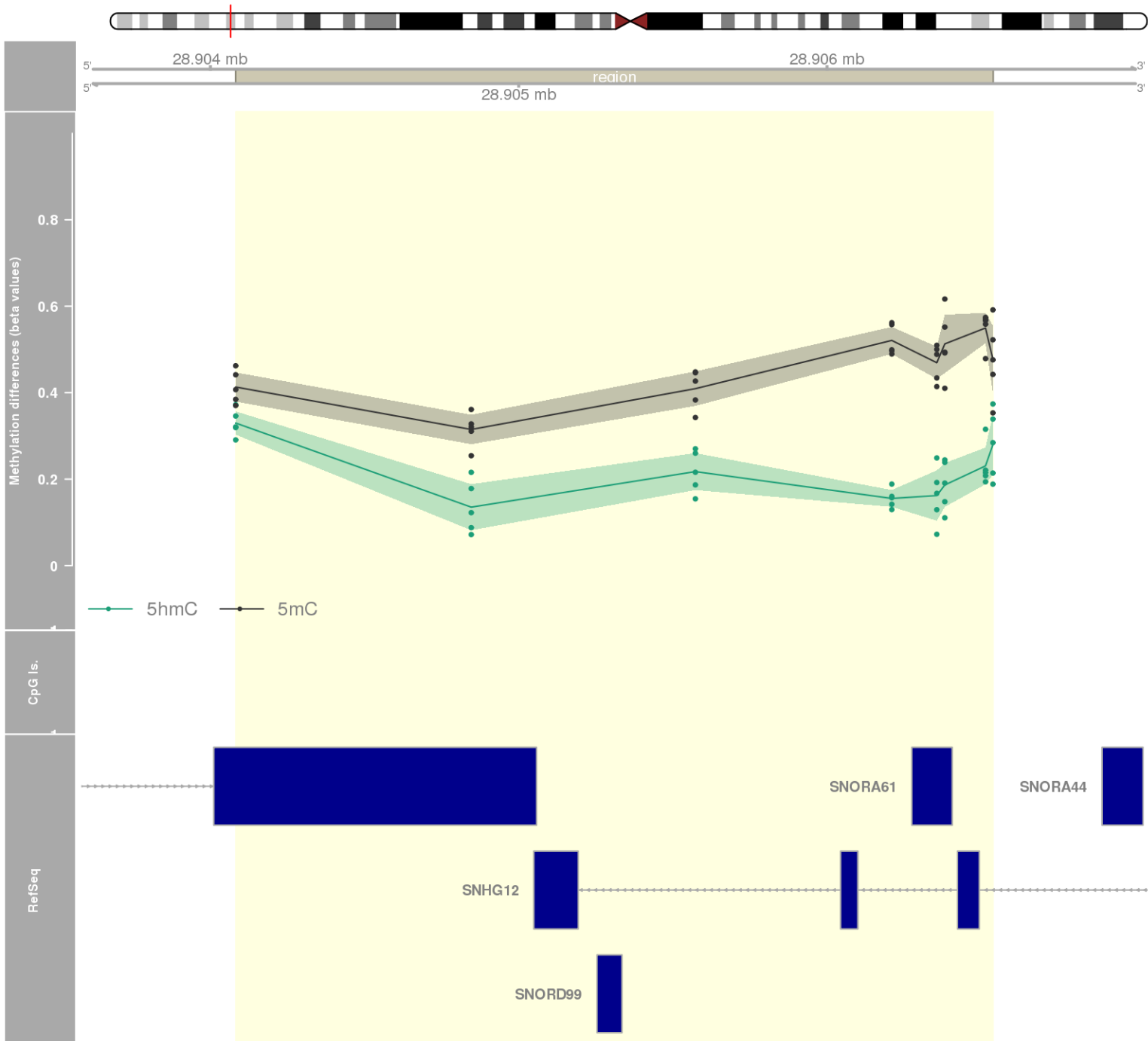
DMR 251 // chr6:31759412-31760233 // 821 pb. (9 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.018 // fwerArea: 1
- genes: VARS -



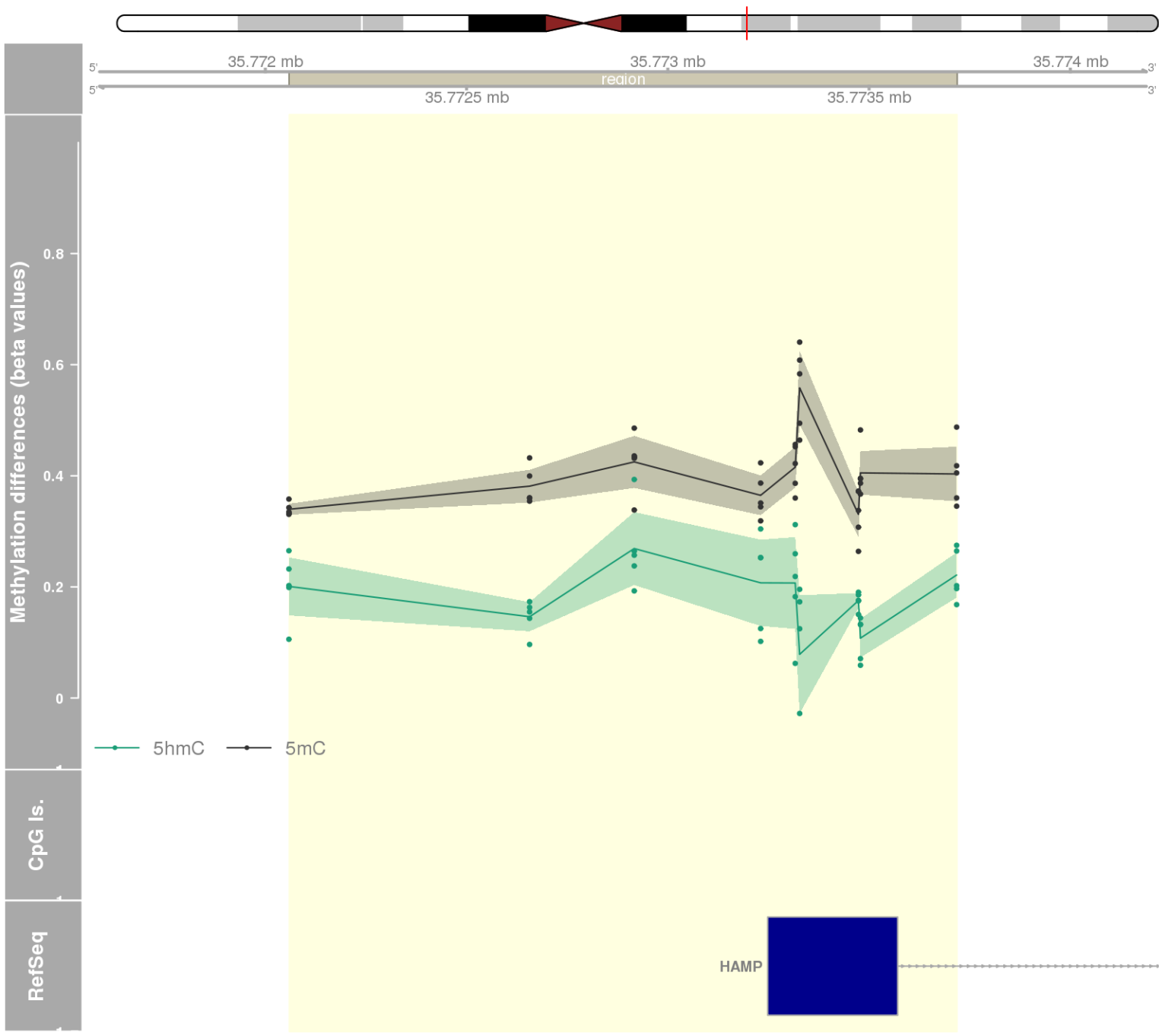
DMR 252 // chr17:72442179-72443401 // 1222 pb. (7 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.018 // fwerArea: 1
- genes: GPRC5C -



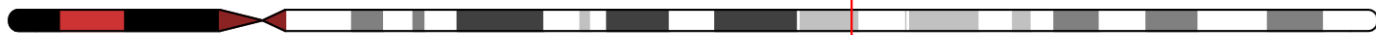
DMR 253 // chr1:28904082-28906538 // 2456 pb. (8 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.018 // fwerArea: 1
- genes: SNHG12 / SNORD99 / TRNAU1AP -



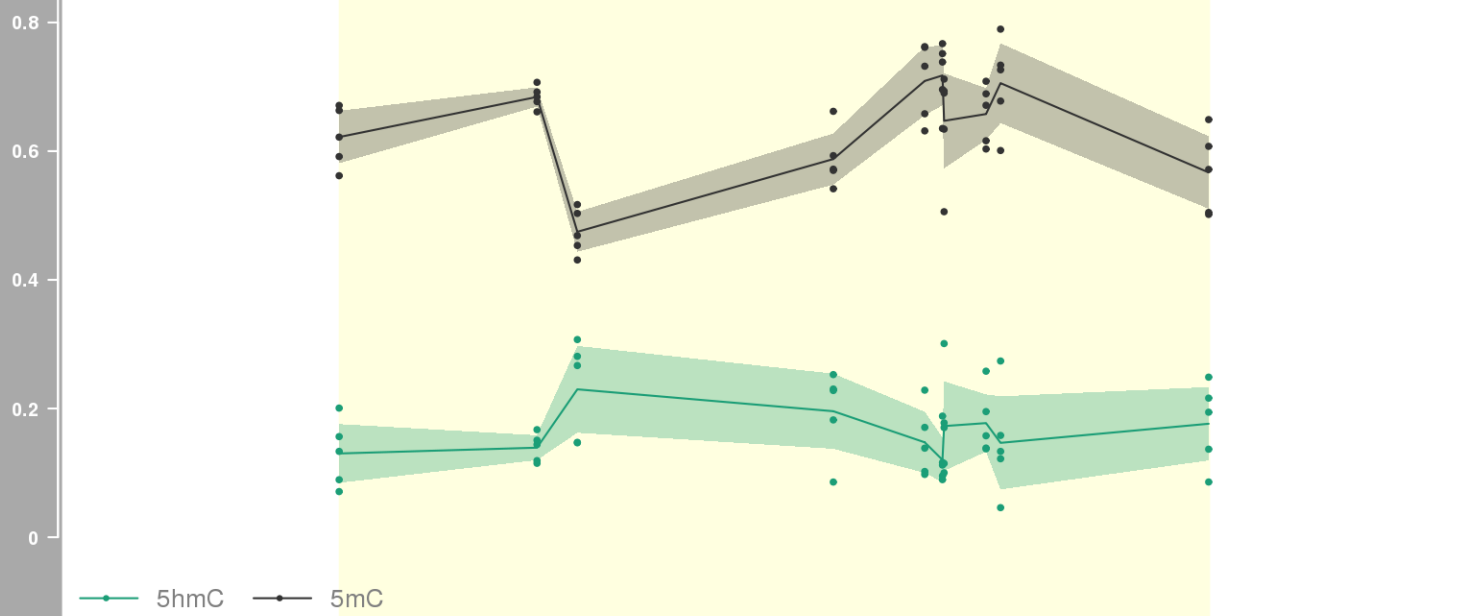
DMR 254 // chr19:35772059-35773718 // 1659 pb. (9 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.018 // fwerArea: 1
- genes: HAMP -



DMR 255 // chr15:63114897-63116514 // 1617 pb. (10 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.018 // fwerArea: 1
- genes: MIR190A / TLN2 -



Methylation differences (beta values)



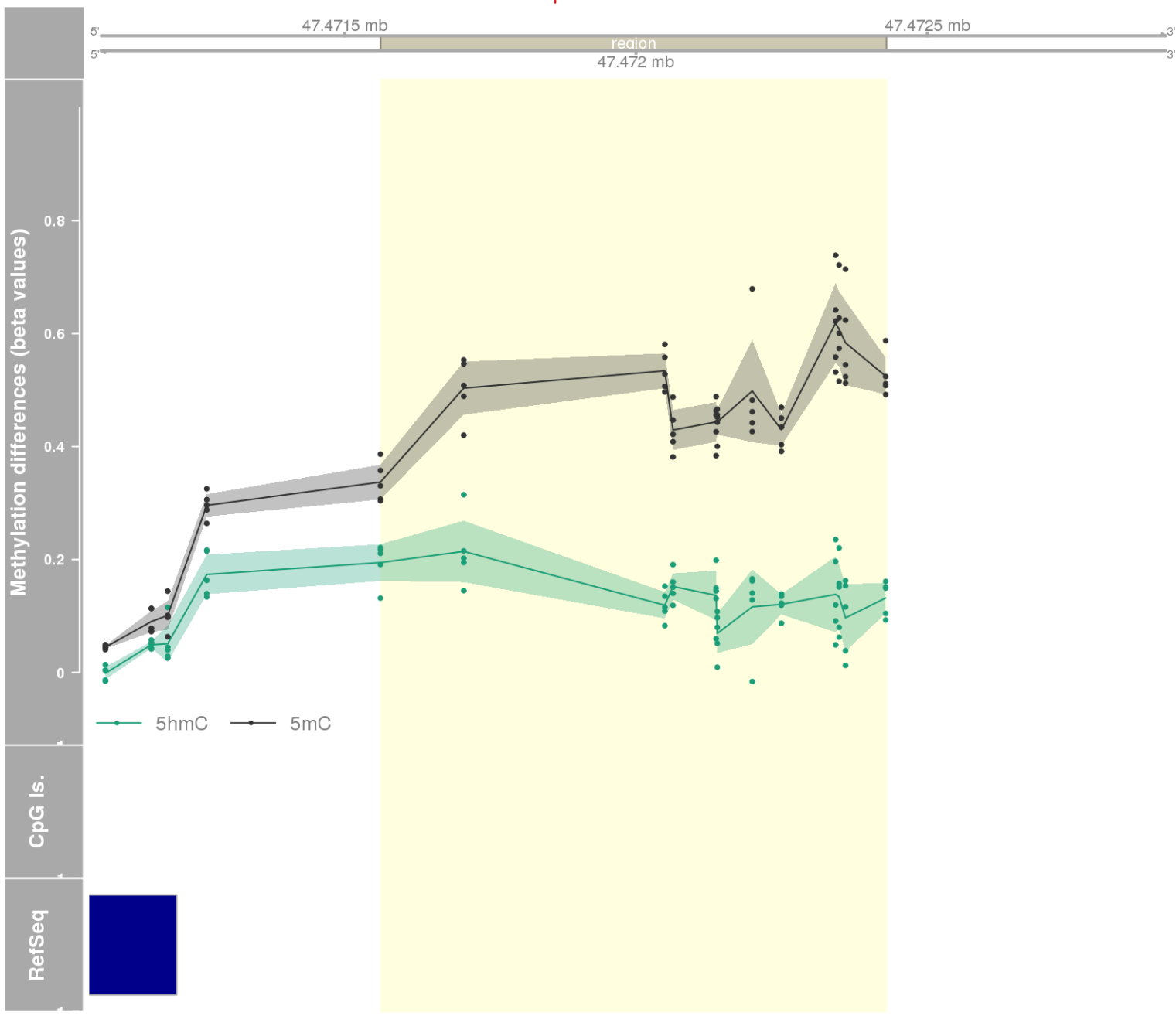
CpG Is.

RefSeq

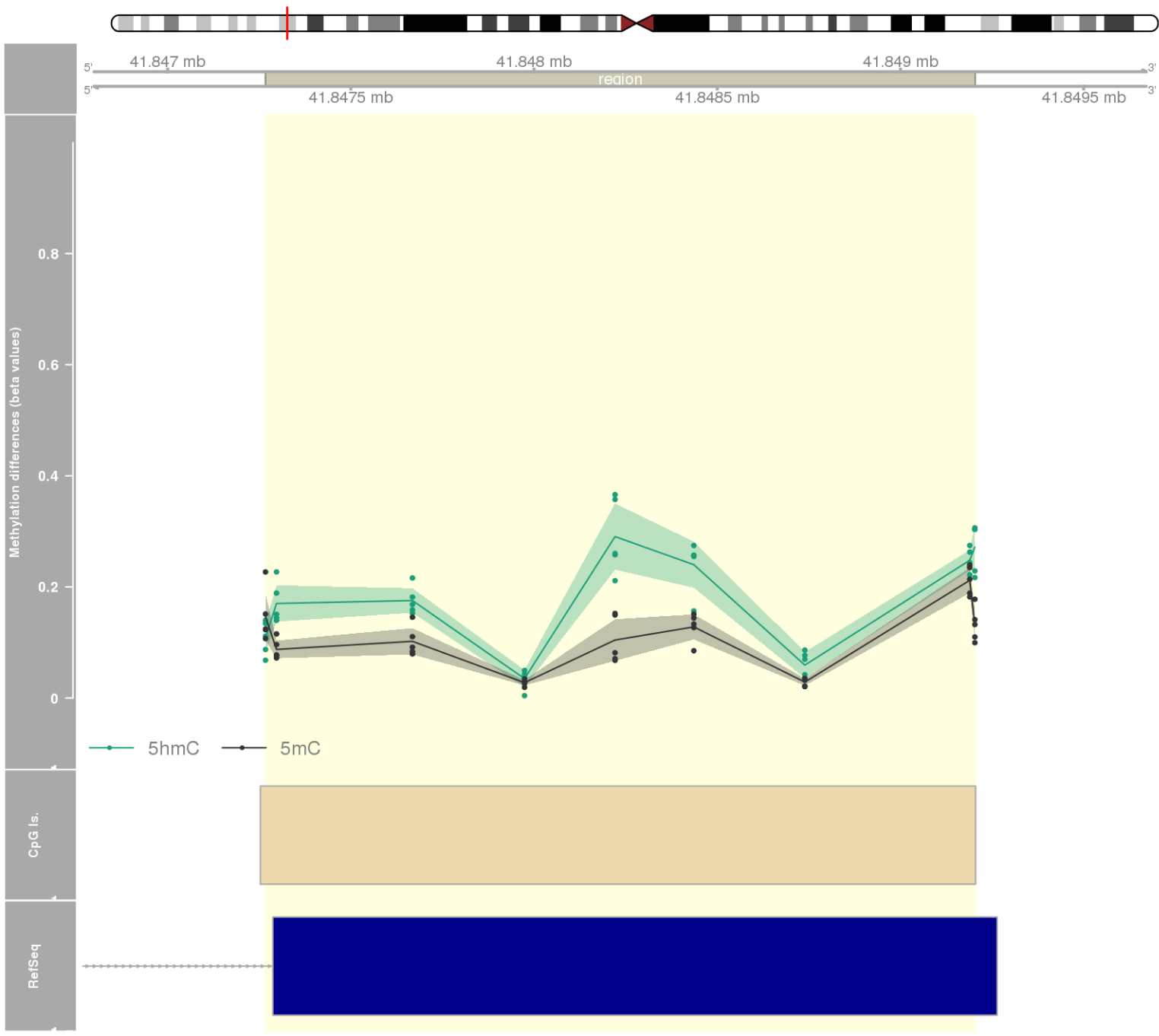
MIR190A



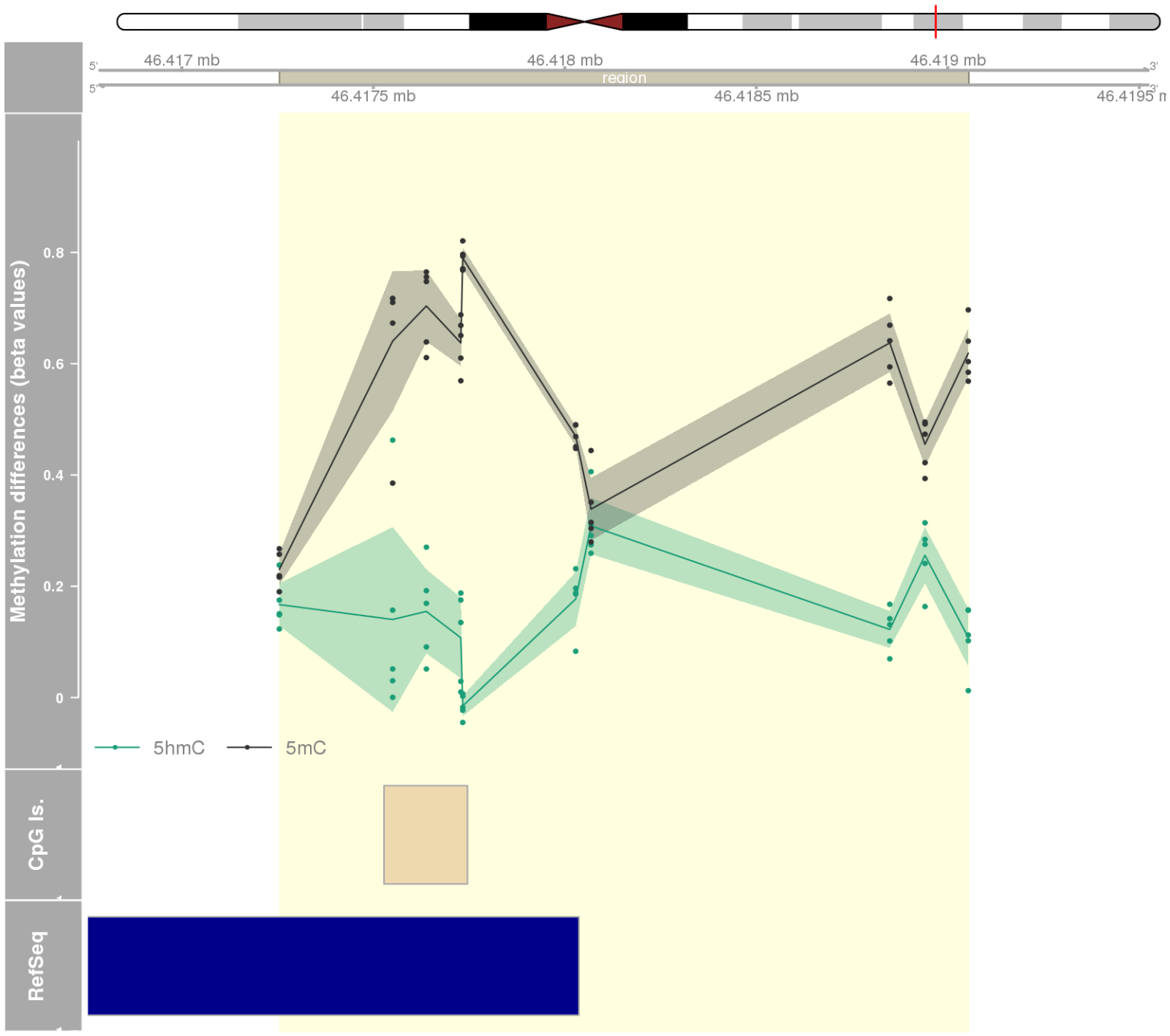
DMR 256 // chr13:47471562-47472429 // 867 pb. (12 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.018 // fwerArea: 1



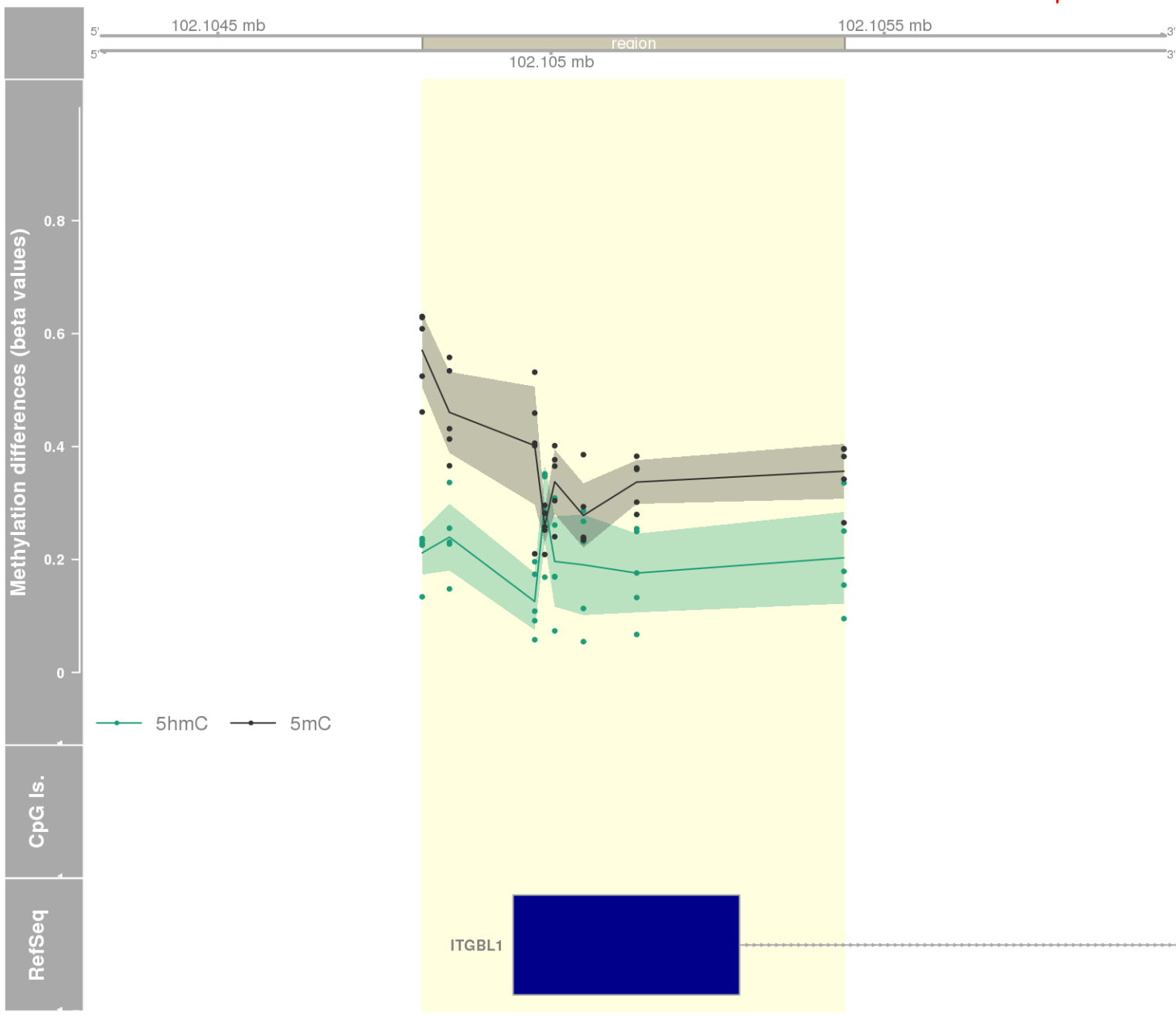
DMR 257 // chr1:41847267-41849203 // 1936 pb. (9 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.018 // fwerArea: 1
- genes: FOXO6 -



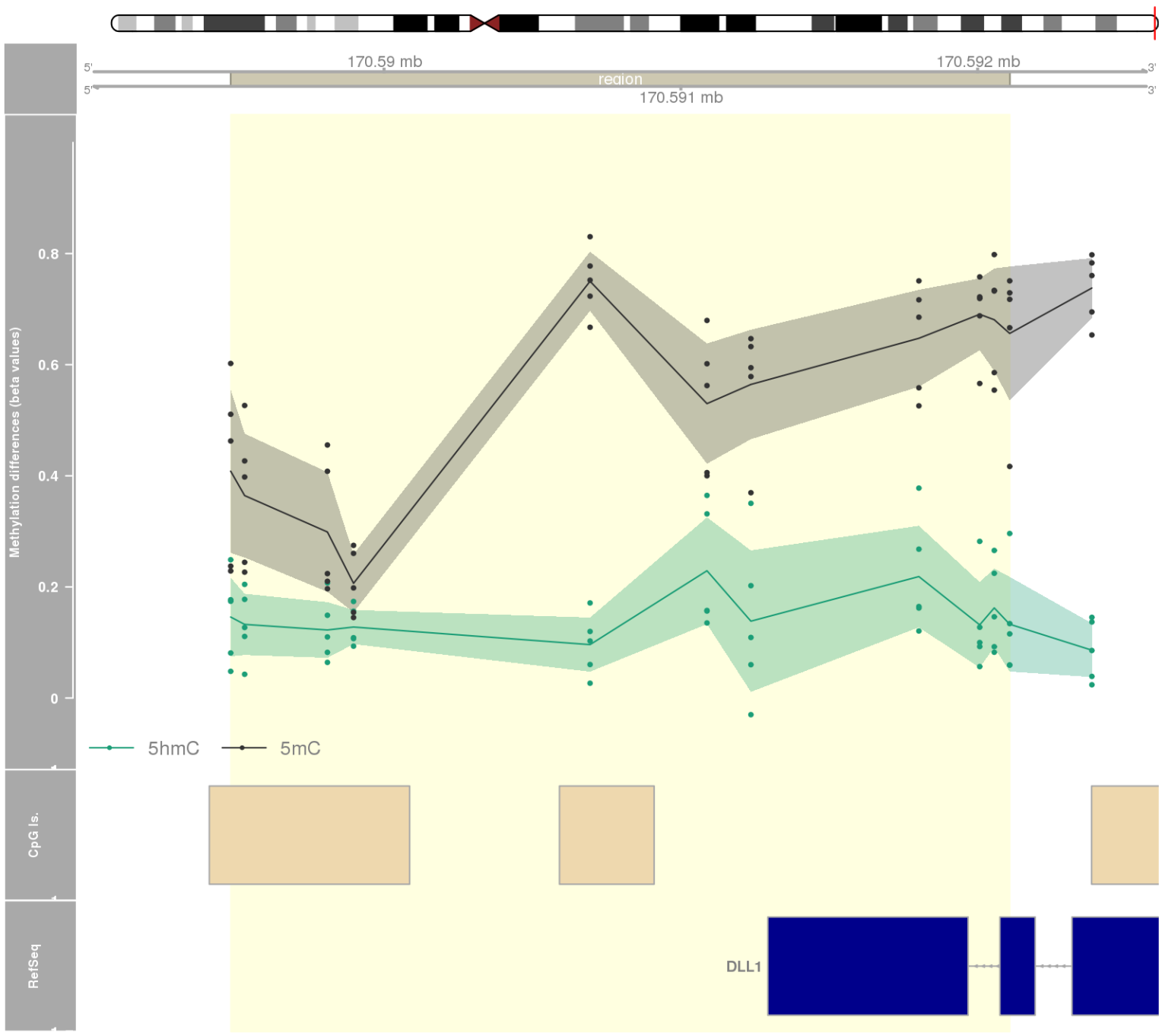
DMR 258 // chr19:46417255-46419054 // 1799 pb. (10 probes) // pvalue: 0.006 // fwer: 0.36 // pvalueArea: 0.018 // fwerArea: 1
- genes: NANOS2 -



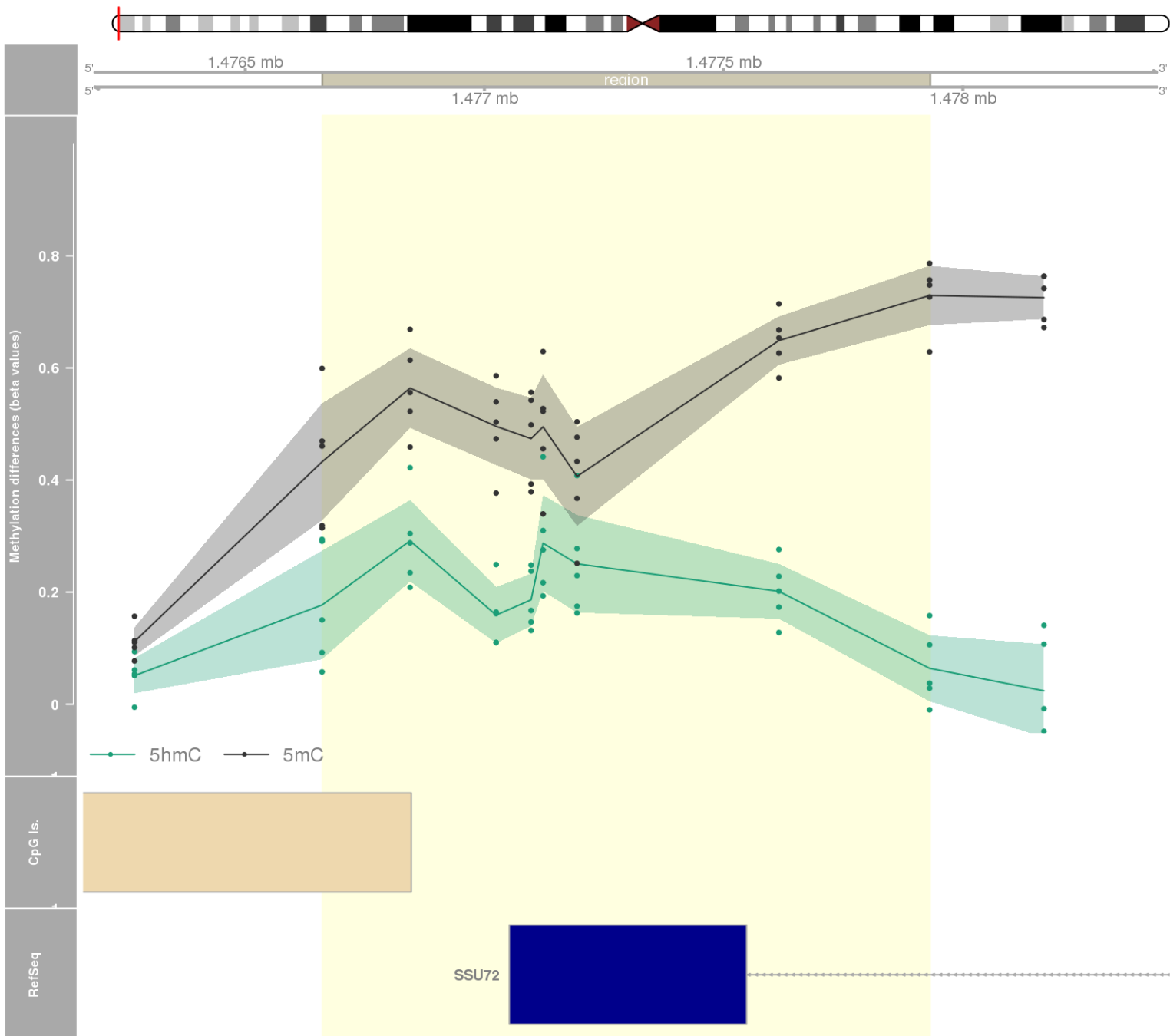
DMR 259 // chr13:102104807-102105440 // 633 pb. (8 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.018 // fwerArea: 1
- genes: ITGBL1 -



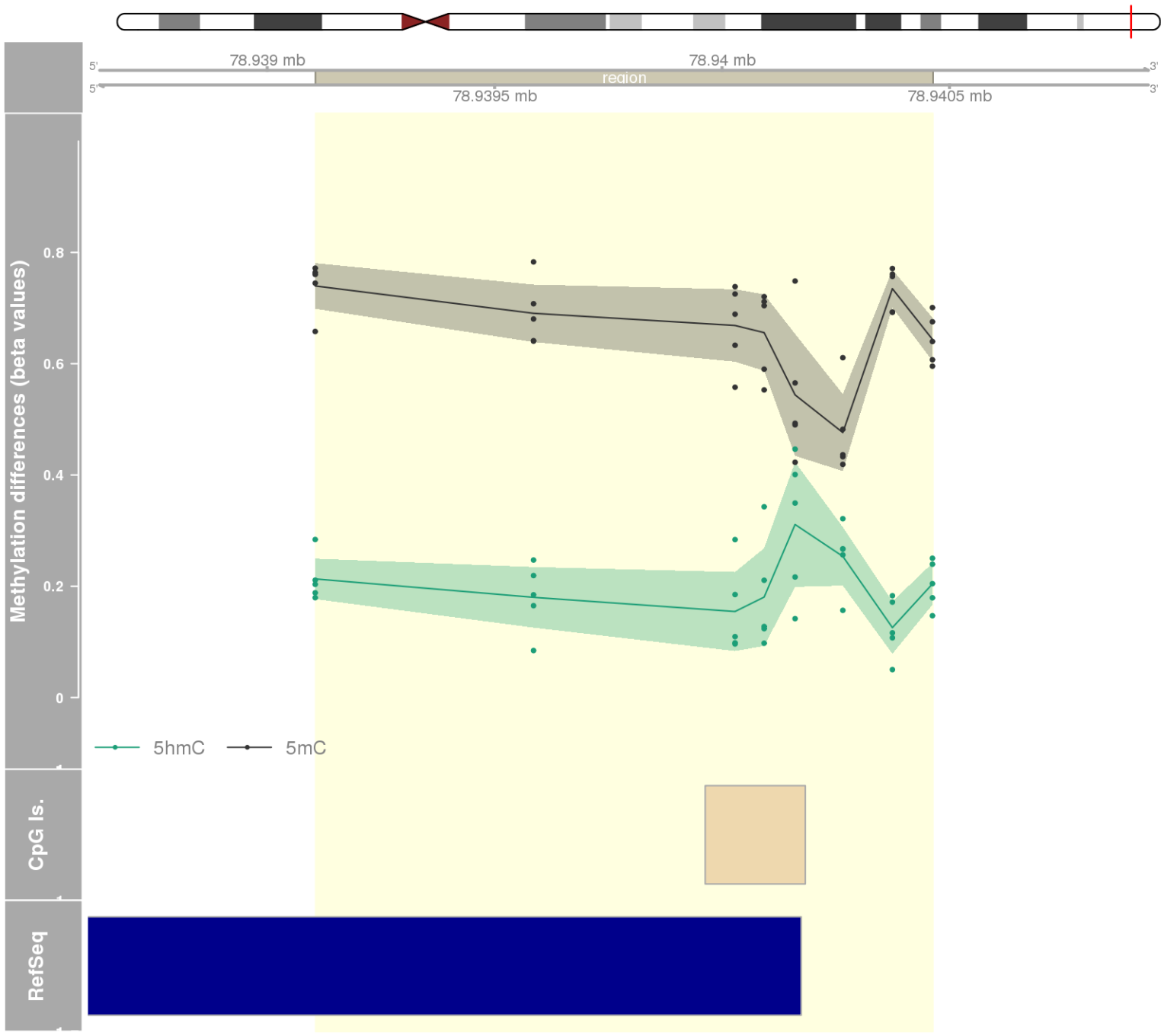
DMR 260 // chr6:170589483-170592108 // 2625 pb. (11 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.018 // fwerArea: 1
- genes: DLL1 -



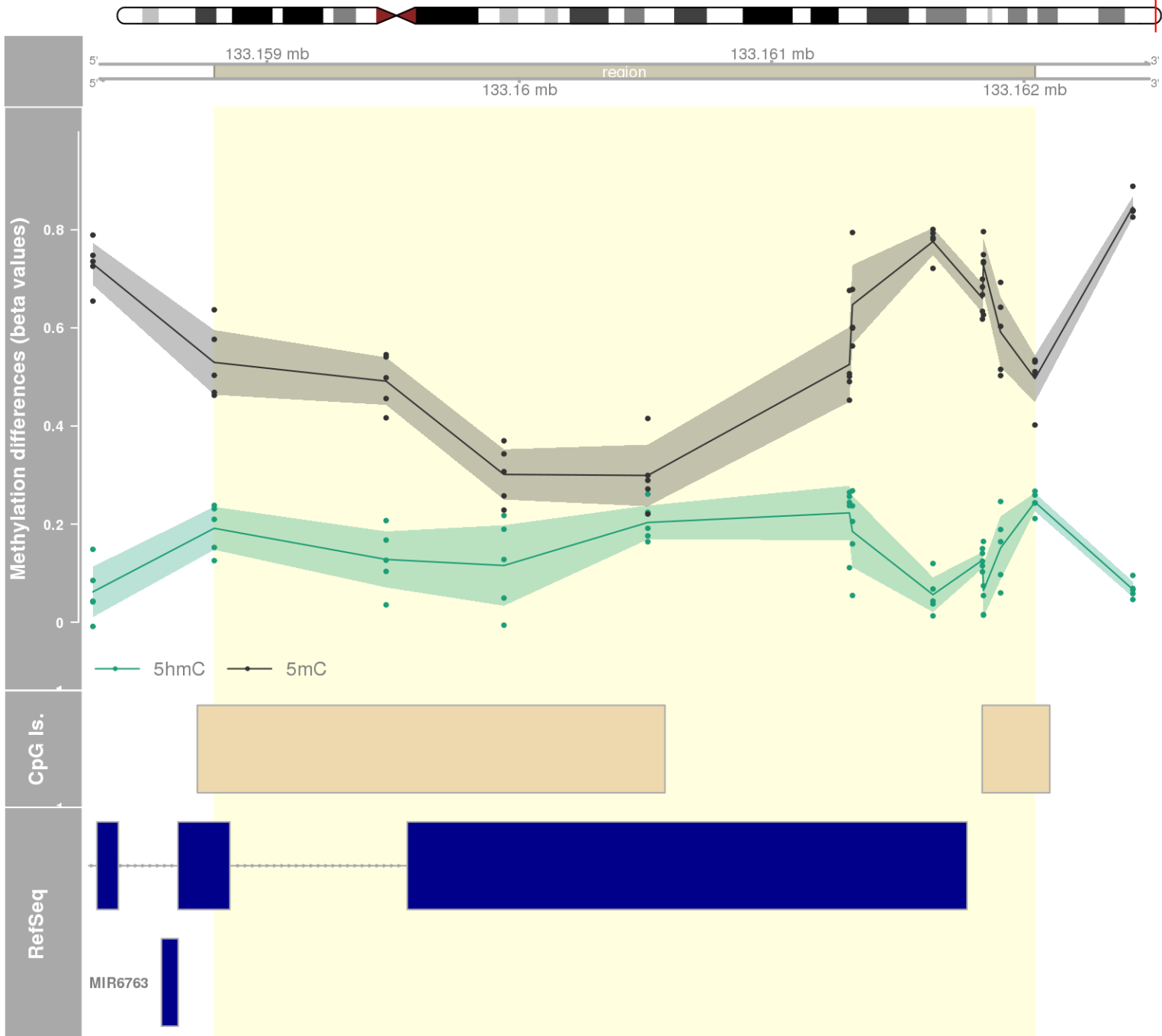
DMR 261 // chr1:1476661-1477931 // 1270 pb. (8 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.018 // fwerArea: 1
- genes: SSU72 -



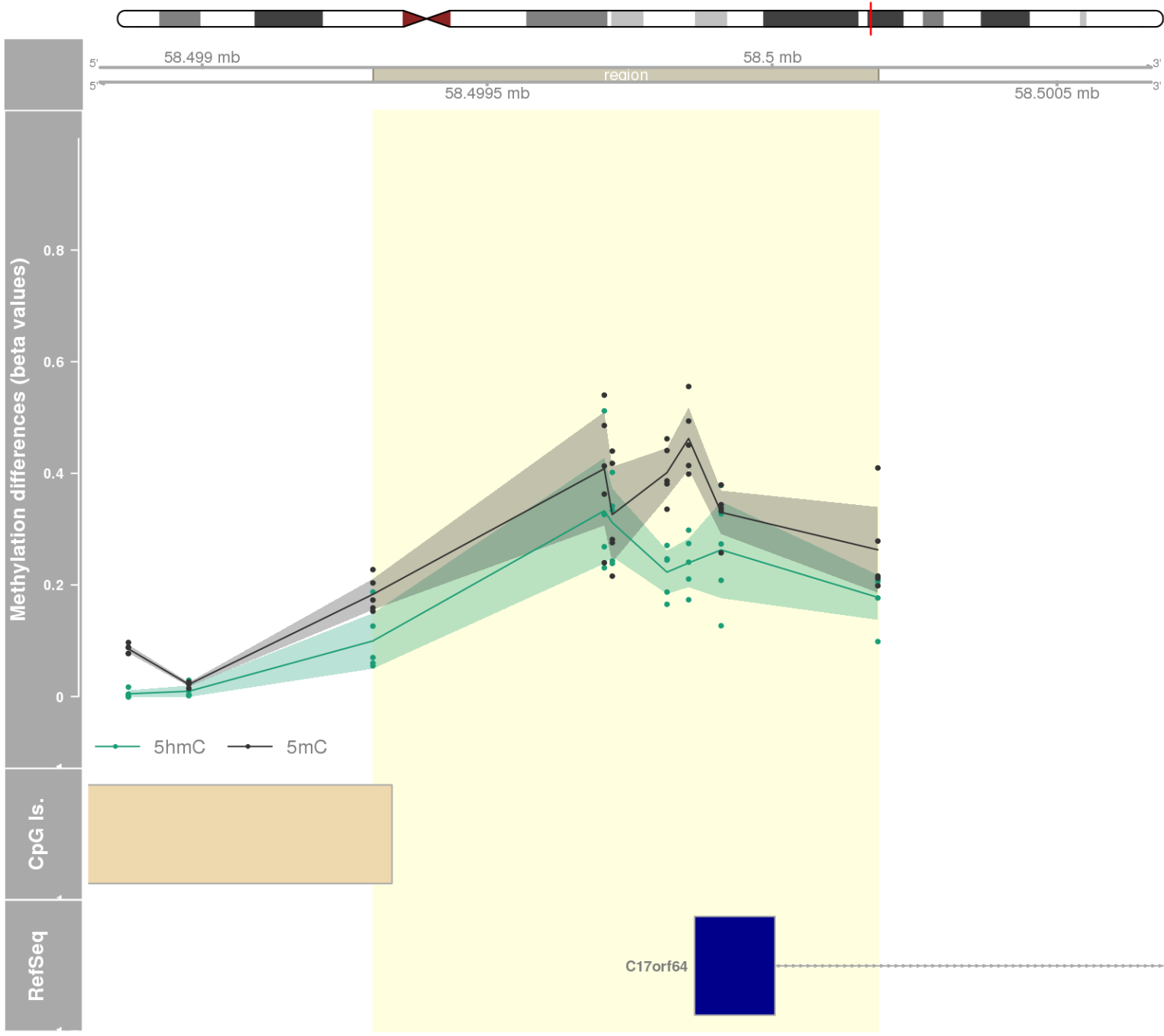
DMR 262 // chr17:78939106-78940463 // 1357 pb. (8 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.018 // fwerArea: 1
- genes: RPTOR -



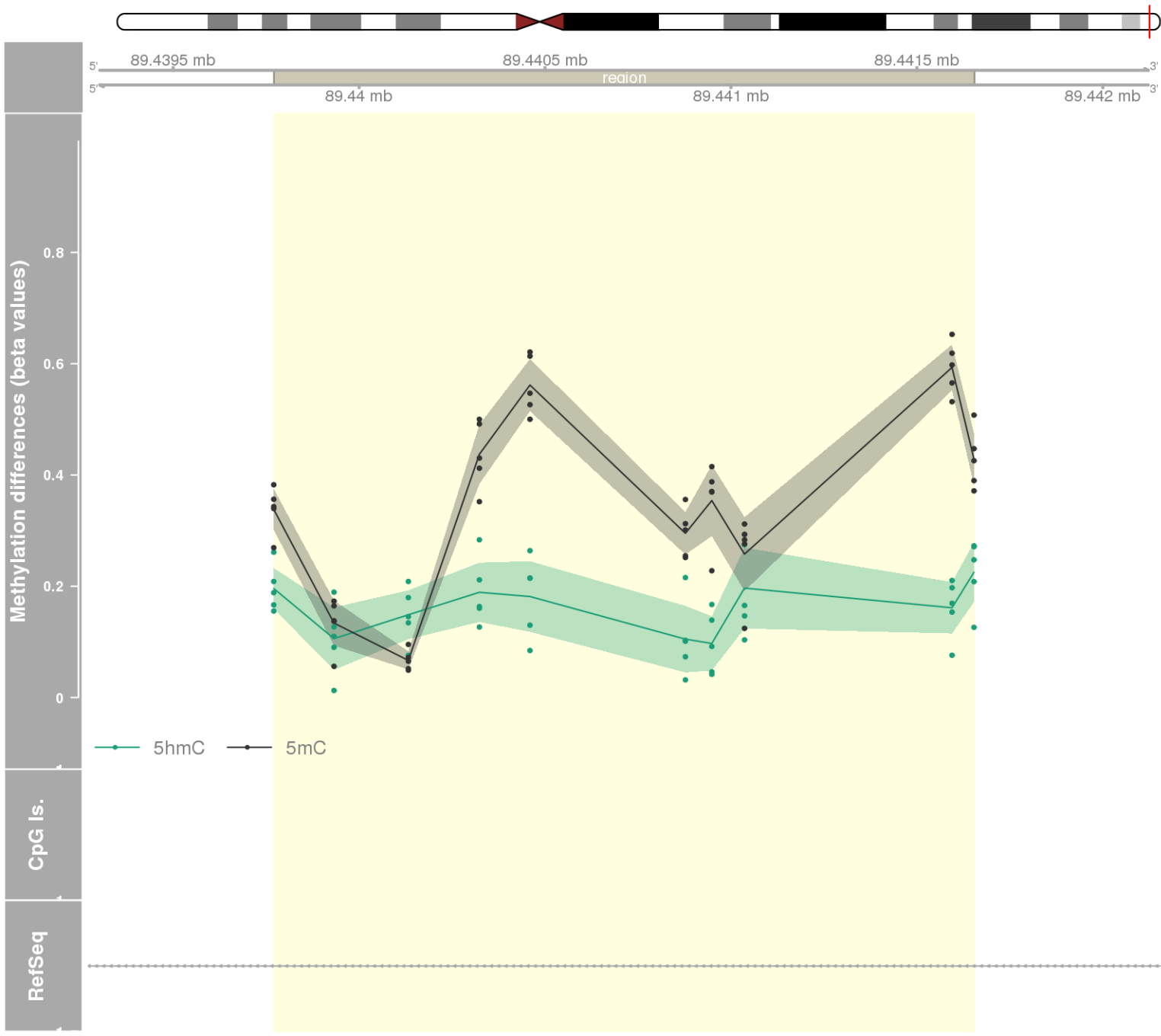
DMR 263 // chr12:133158791-133162043 // 3252 pb. (11 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.019 // fwerArea: 1
- genes: FBRSL1 -



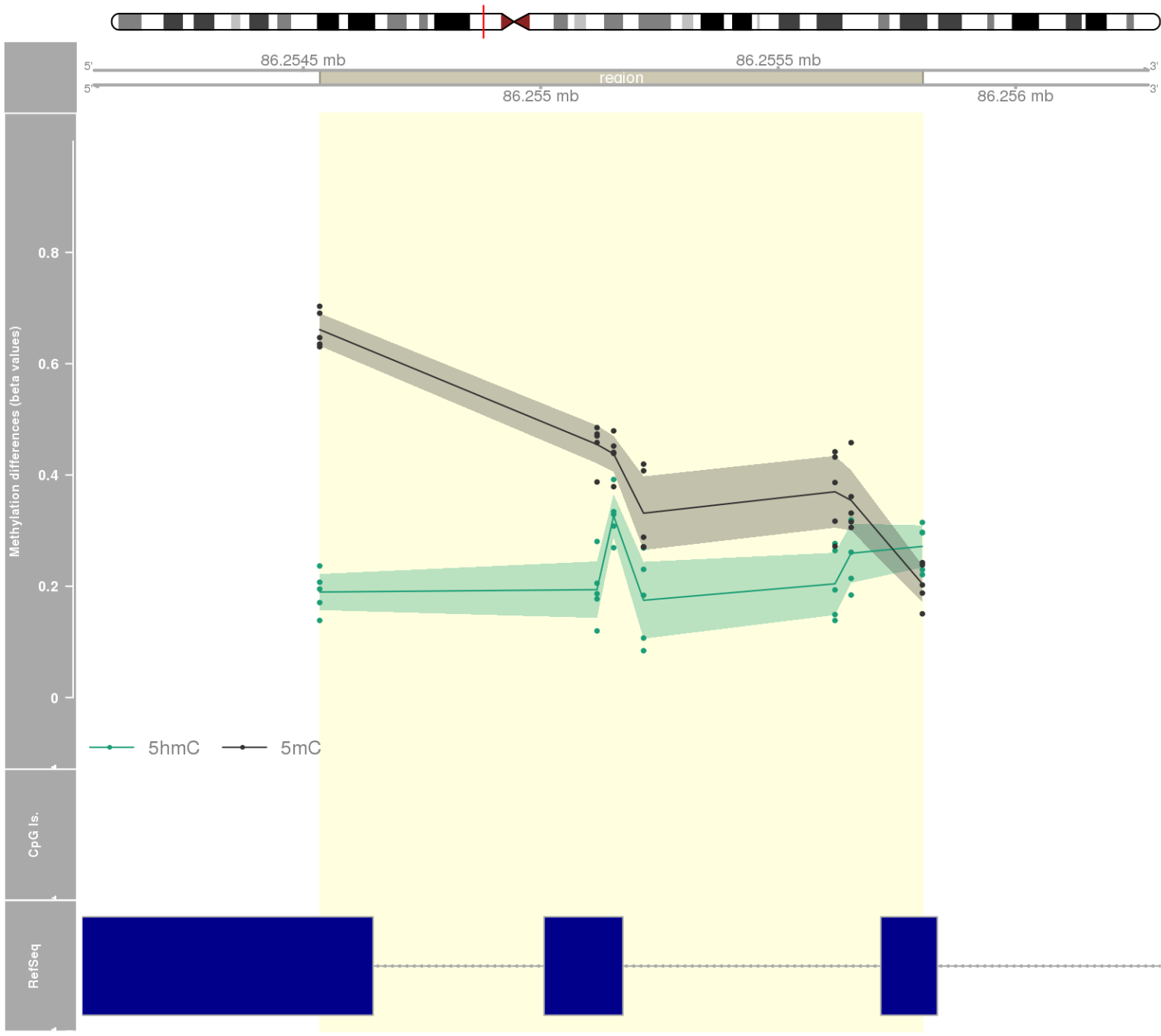
DMR 264 // chr17:58499300-58500186 // 886 pb. (7 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.019 // fwerArea: 1
- genes: USP32 / C17orf64 -



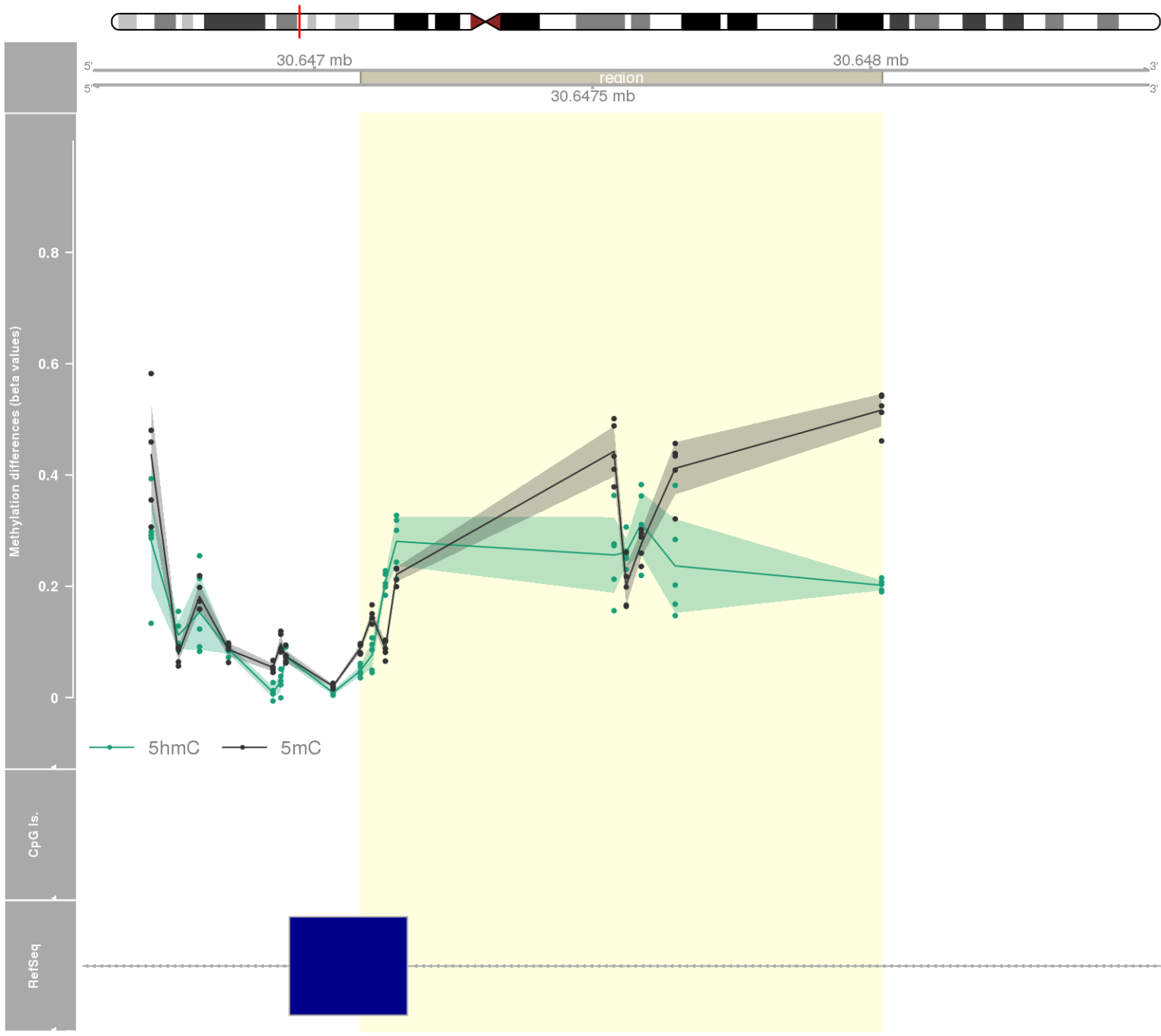
DMR 265 // chr16:89439771-89441654 // 1883 pb. (10 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.019 // fwerArea: 1
- genes: ANKRD11 -



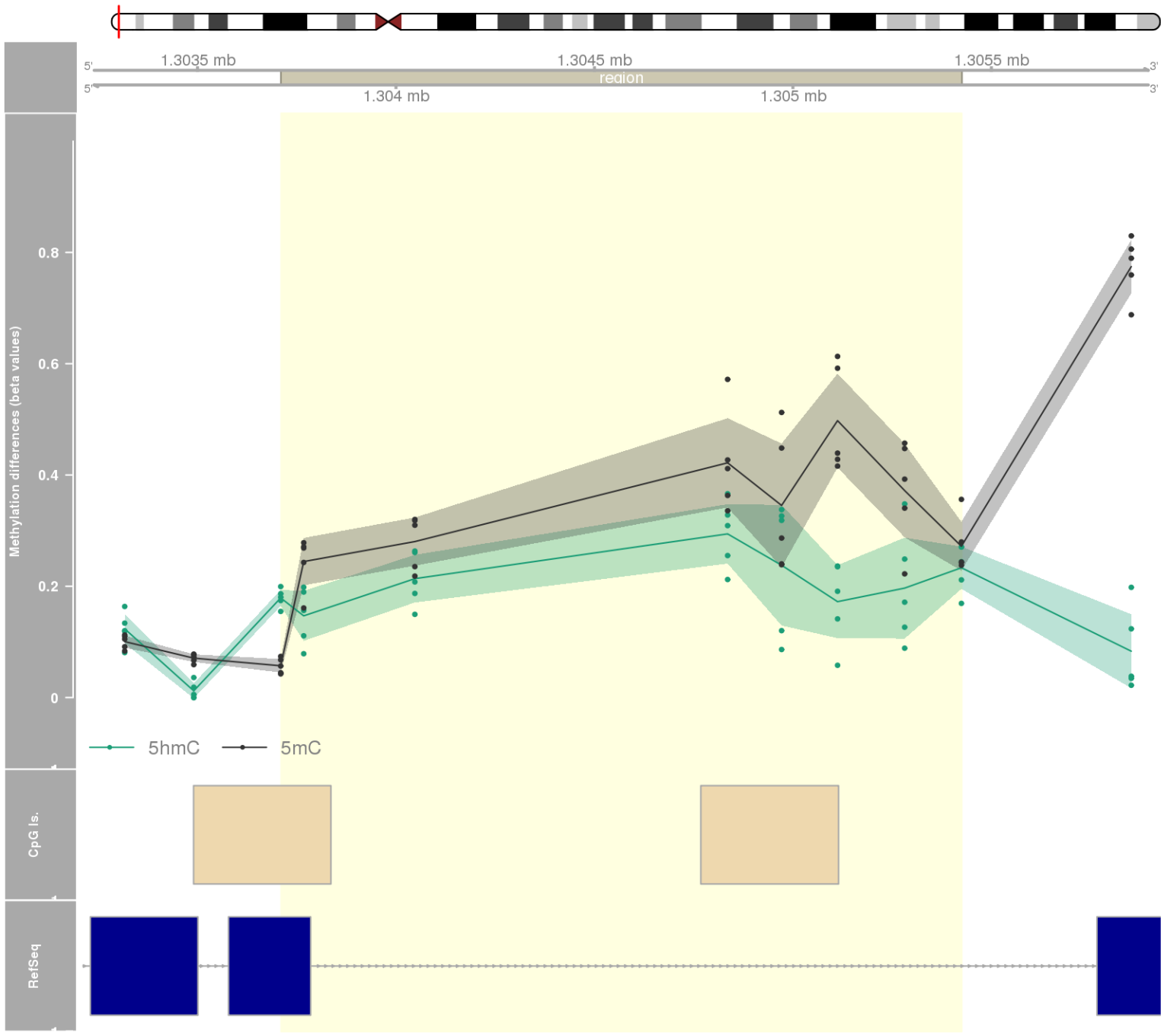
DMR 266 // chr2:86254535-86255804 // 1269 pb. (7 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.019 // fwerArea: 1
- genes: POLR1A -



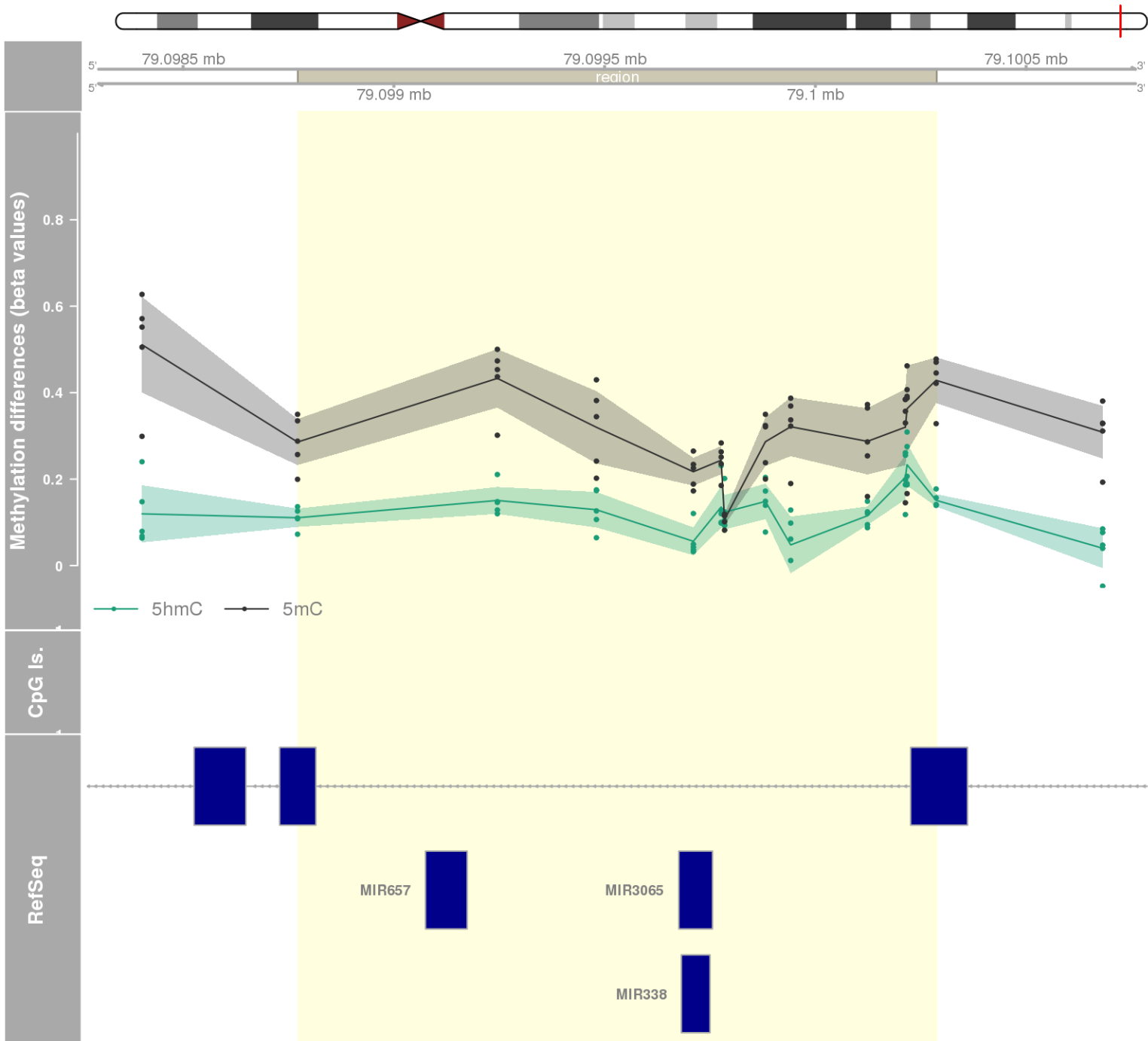
DMR 267 // chr6:30647083-30648020 // 937 pb. (9 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.019 // fwerArea: 1
- genes: PPP1R18 -



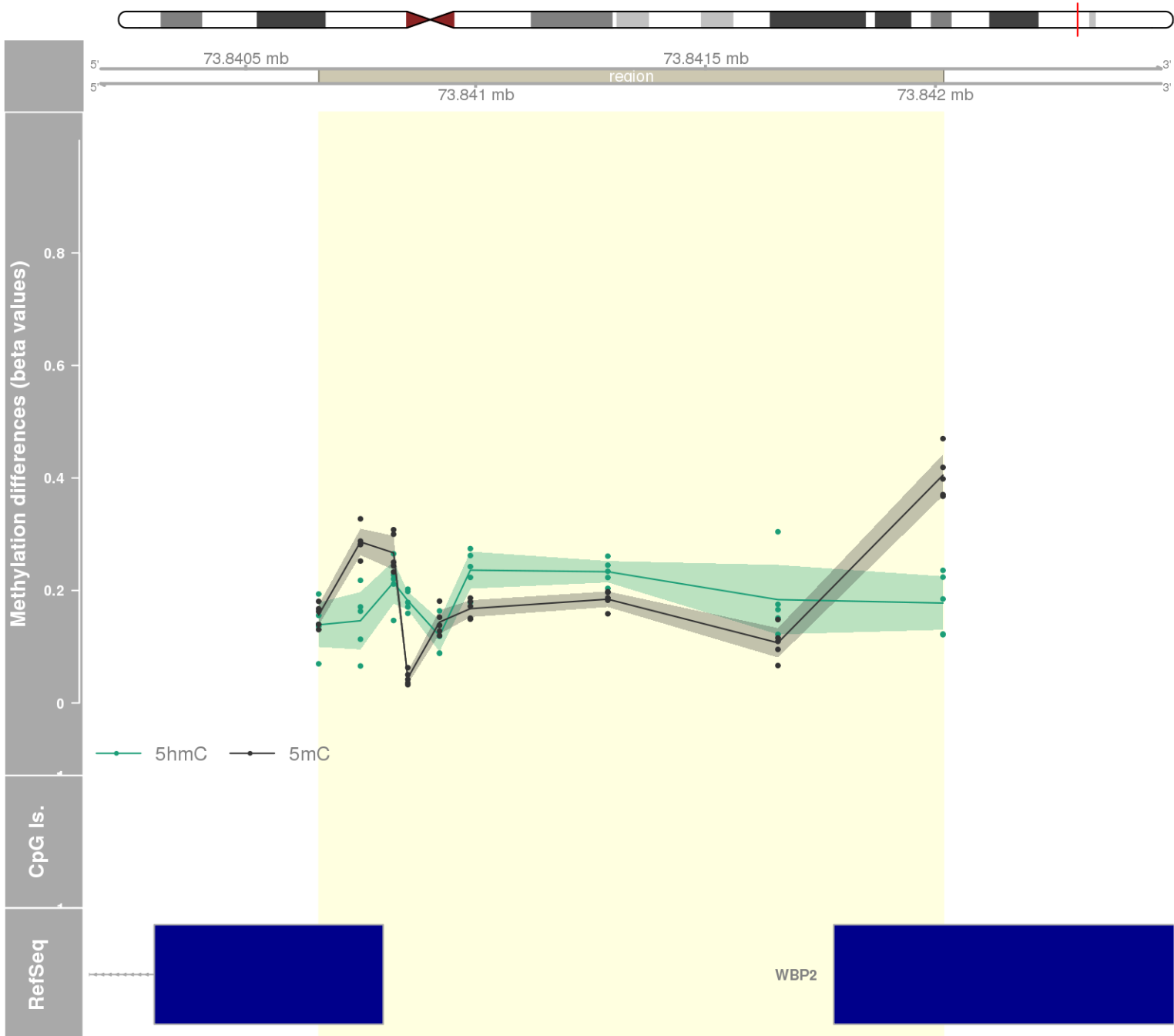
DMR 268 // chr4:1303710-1305425 // 1715 pb. (8 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.019 // fwerArea: 1
- genes: MAEA -



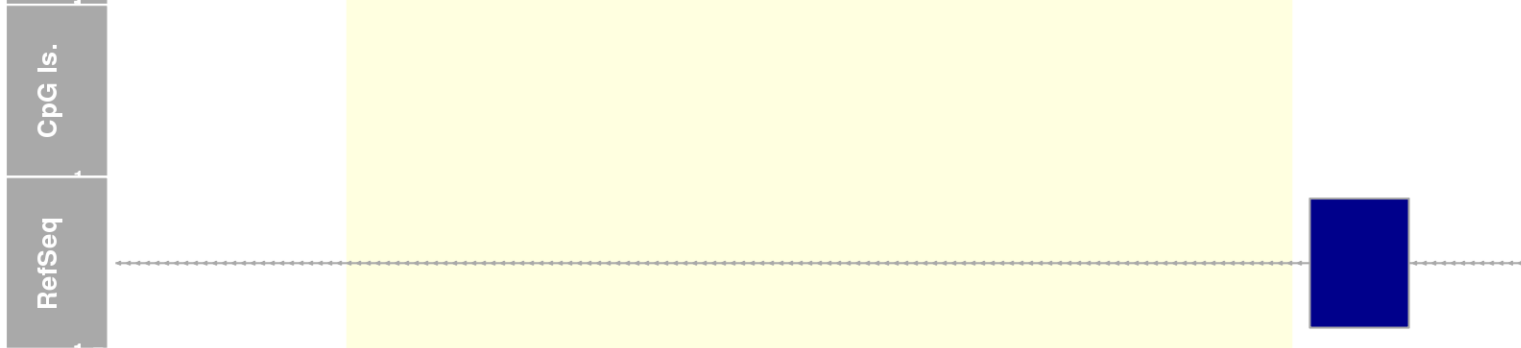
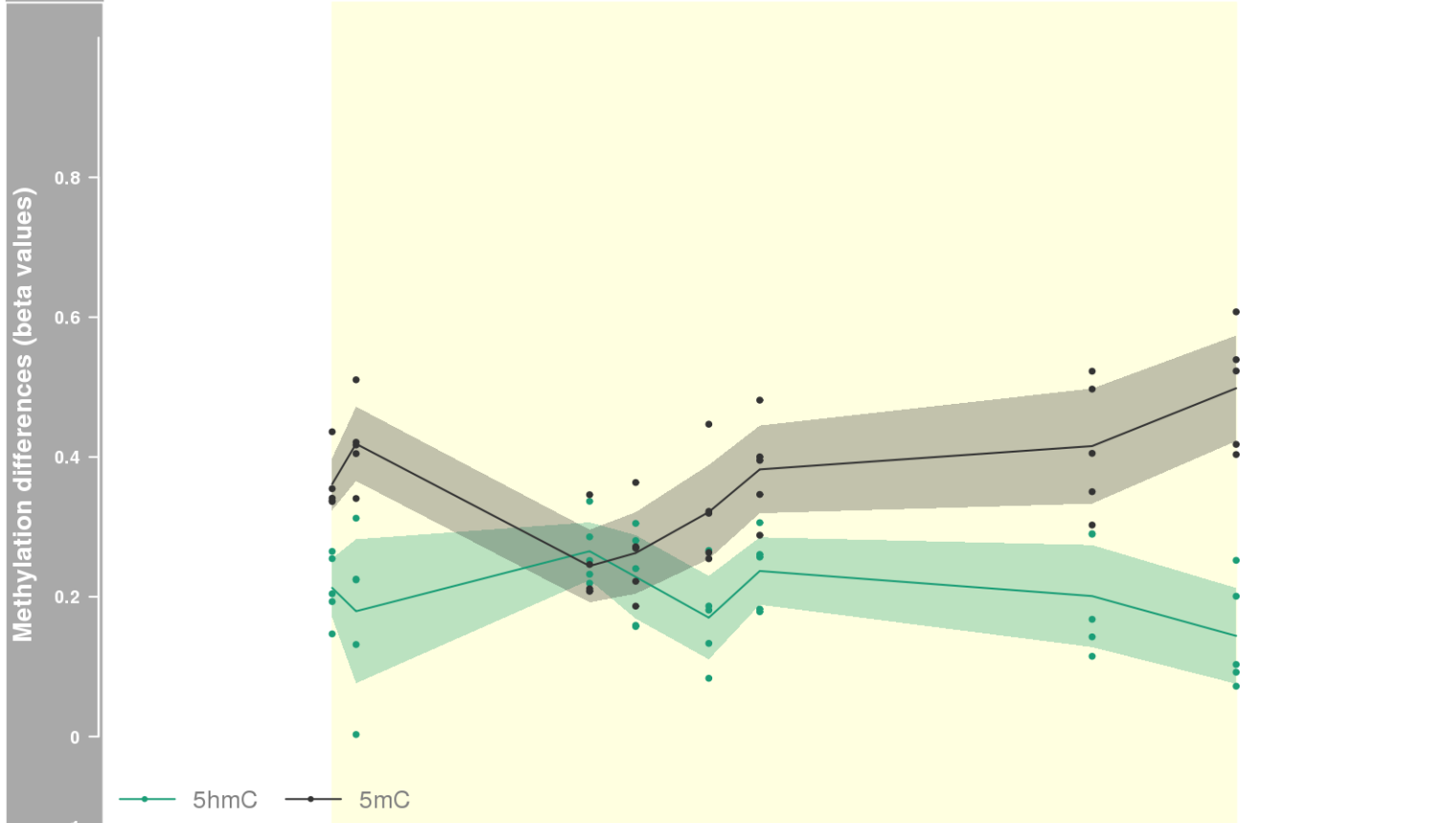
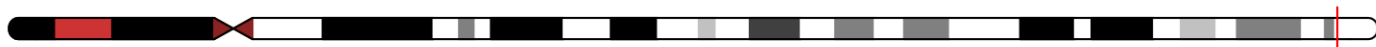
DMR 269 // chr17:79098772-79100287 // 1515 pb. (12 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.019 // fwerArea: 1
- genes: AATK / MIR657 / MIR3065 / MIR338 -



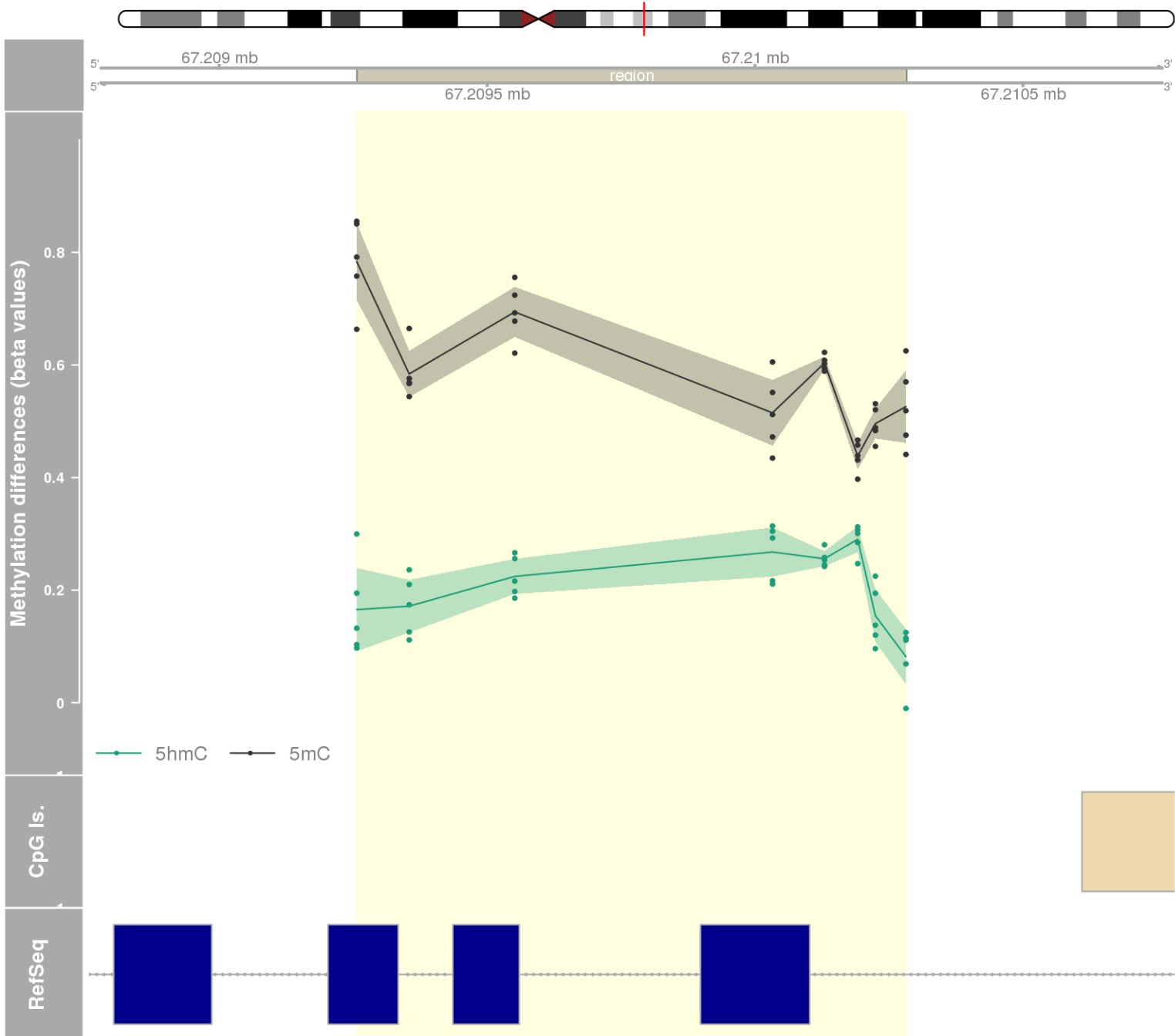
DMR 270 // chr17:73840659-73842017 // 1358 pb. (9 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.019 // fwerArea: 1
- genes: UNC13D / WBP2 -



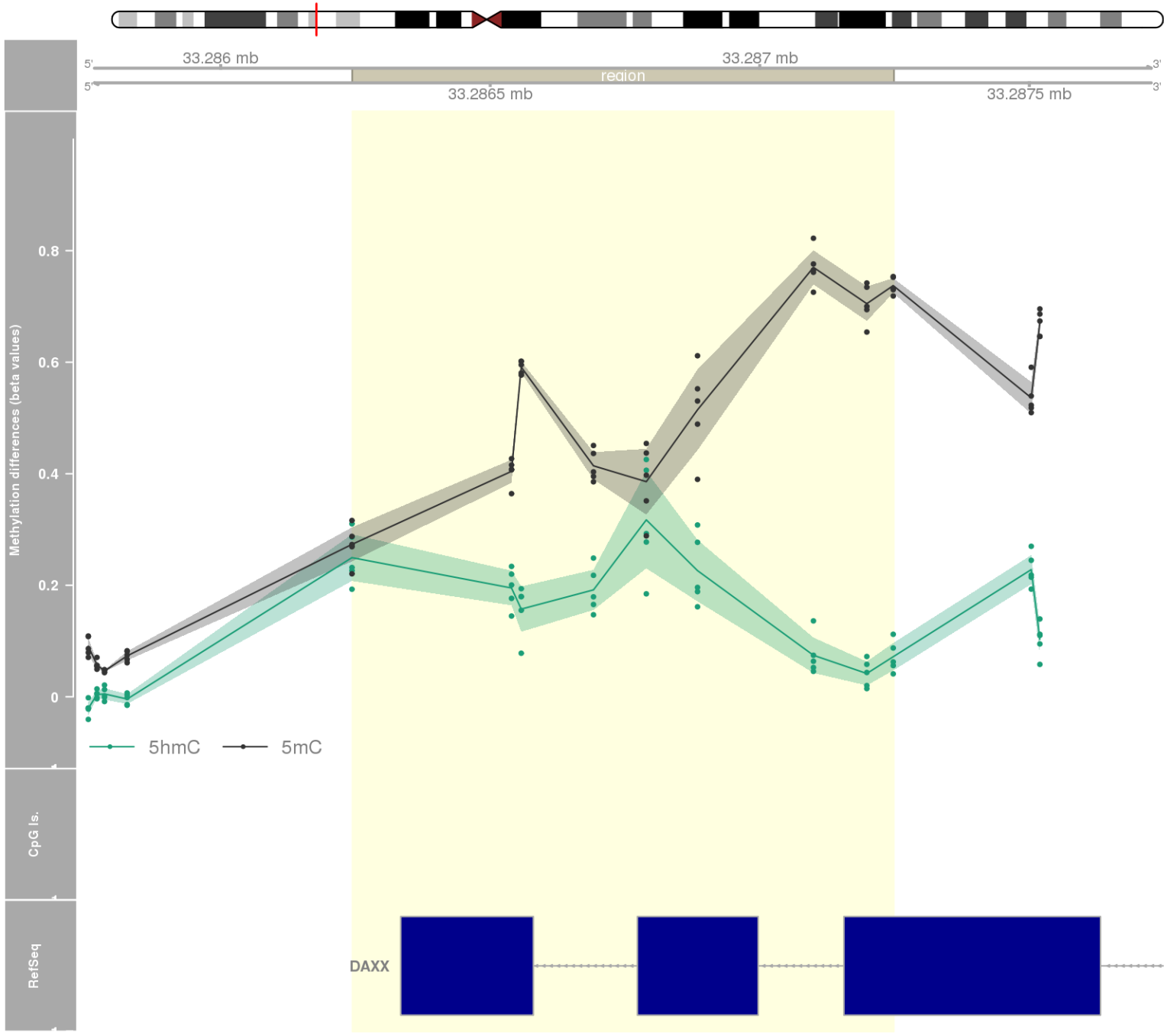
DMR 271 // chr14:104171259-104173299 // 2040 pb. (8 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.019 // fwerArea: 1
- genes: XRCC3 -



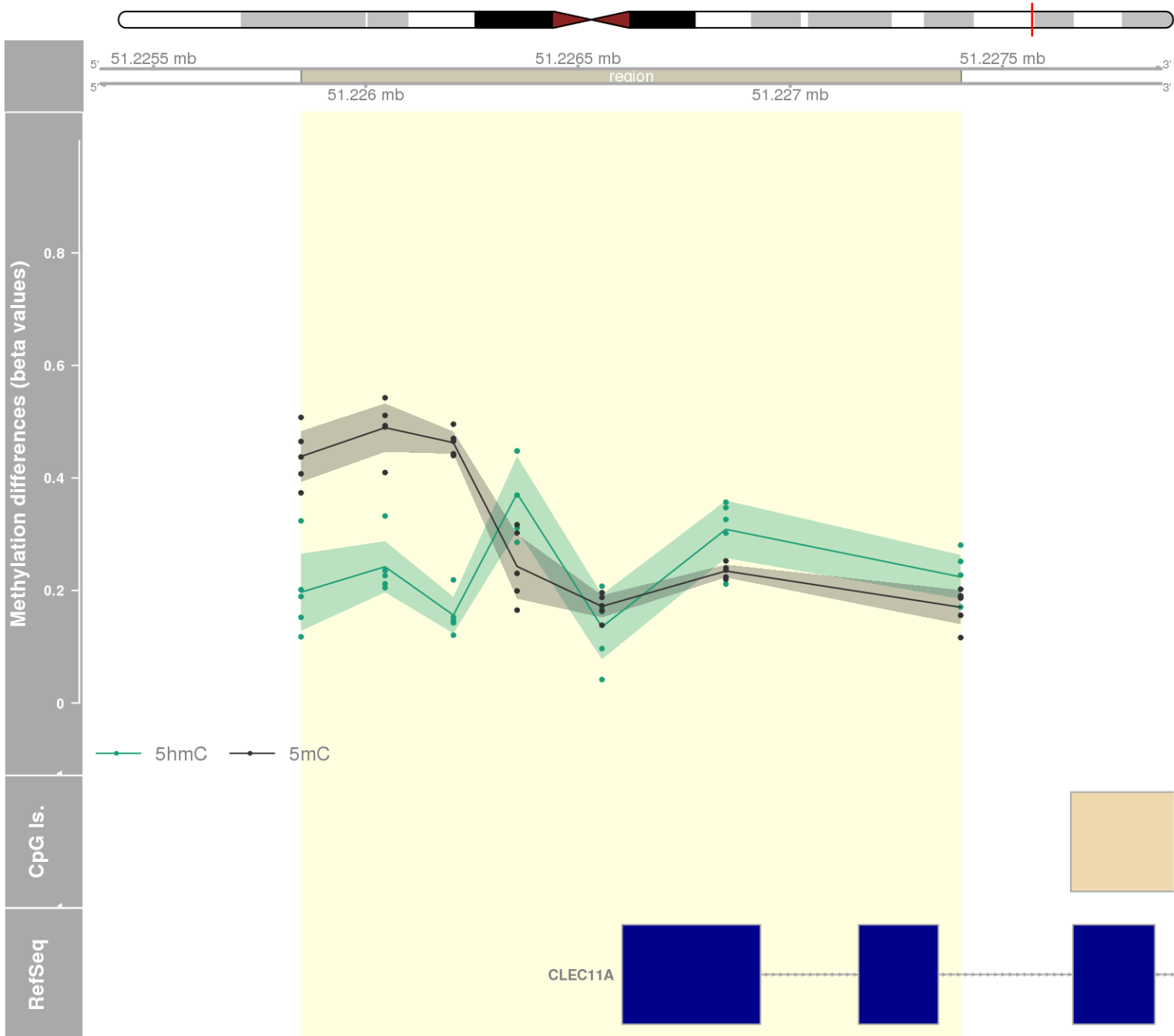
DMR 272 // chr11:67209257-67210282 // 1025 pb. (8 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.019 // fwerArea: 1
- genes: CORO1B -



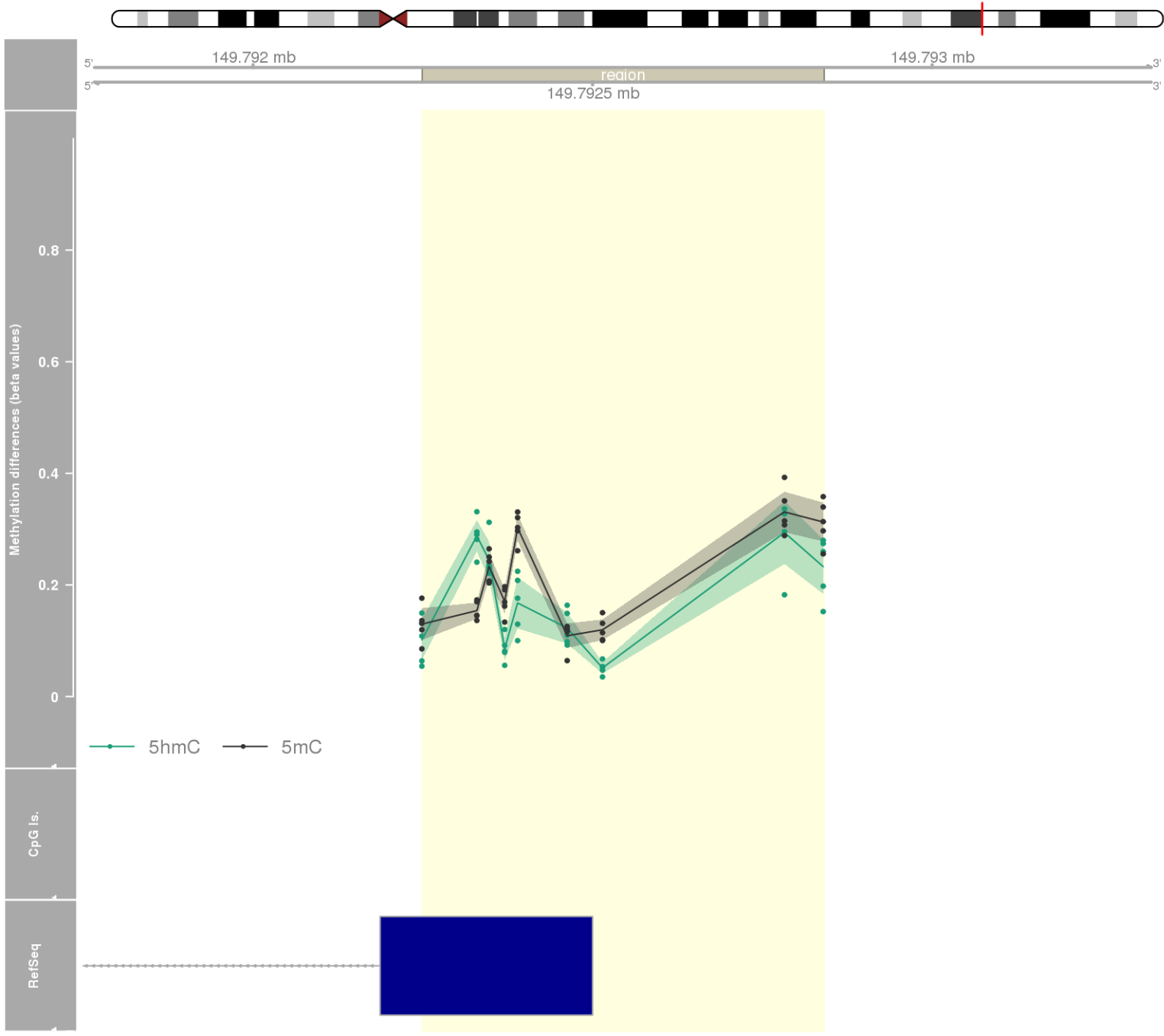
DMR 273 // chr6:33286244-33287248 // 1004 pb. (9 probes) // pvalue: 0.007 // fwer: 0.36 // pvalueArea: 0.019 // fwerArea: 1
- genes: DAXX -



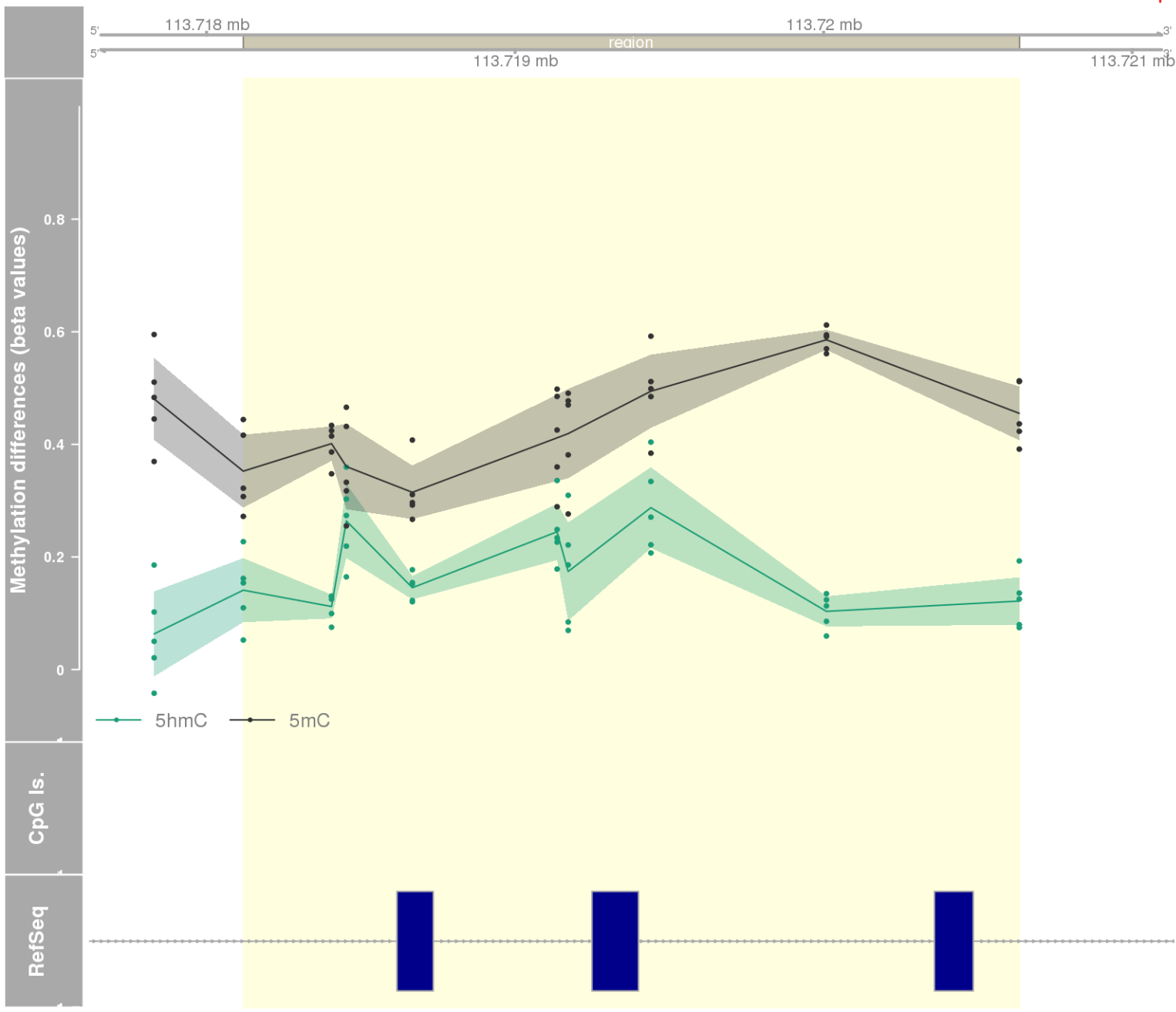
DMR 274 // chr19:51225848-51227402 // 1554 pb. (7 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.019 // fwerArea: 1
- genes: CLEC11A -



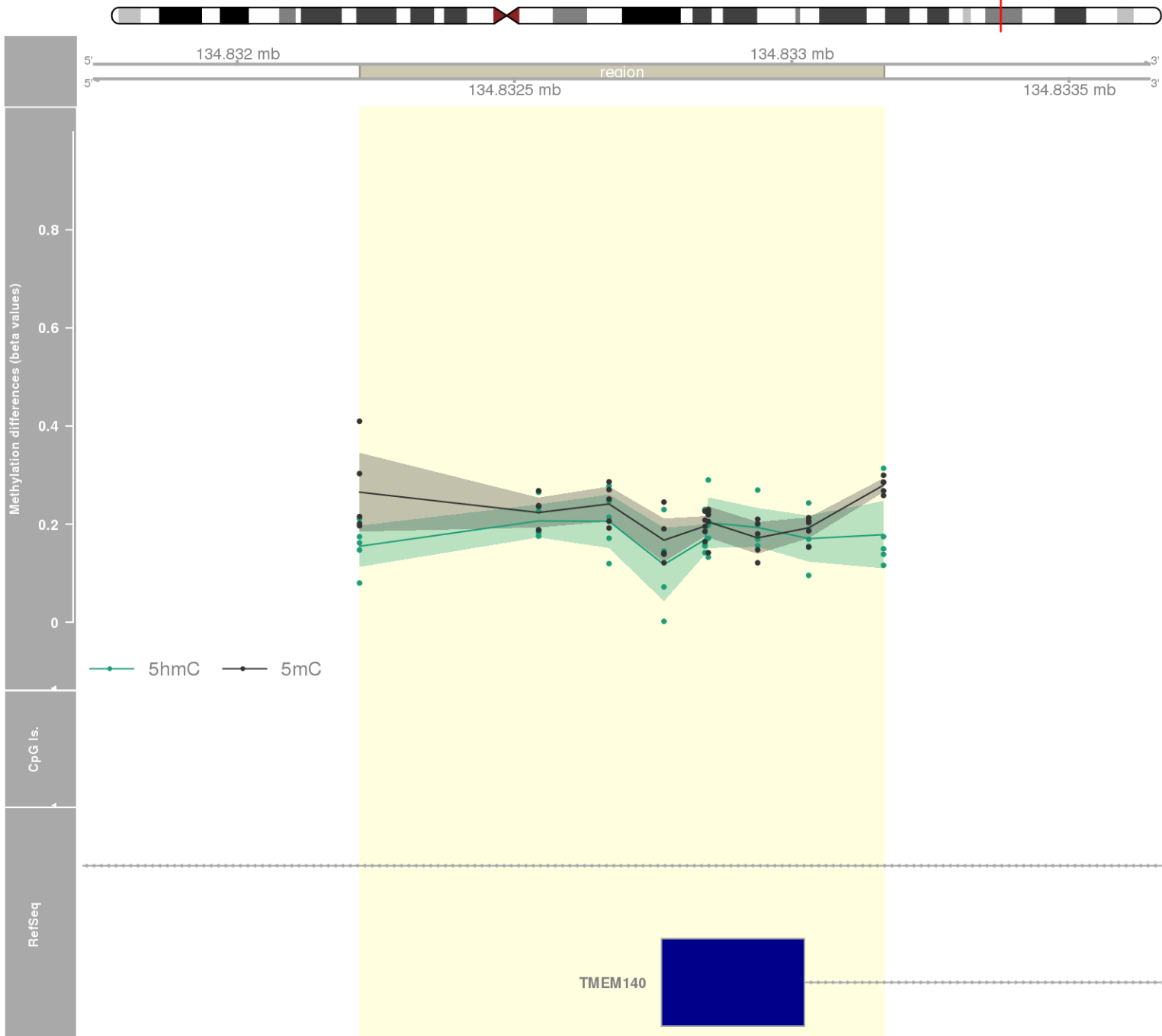
DMR 275 // chr5:149792249-149792840 // 591 pb. (9 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.019 // fwerArea: 1
- genes: CD74 -



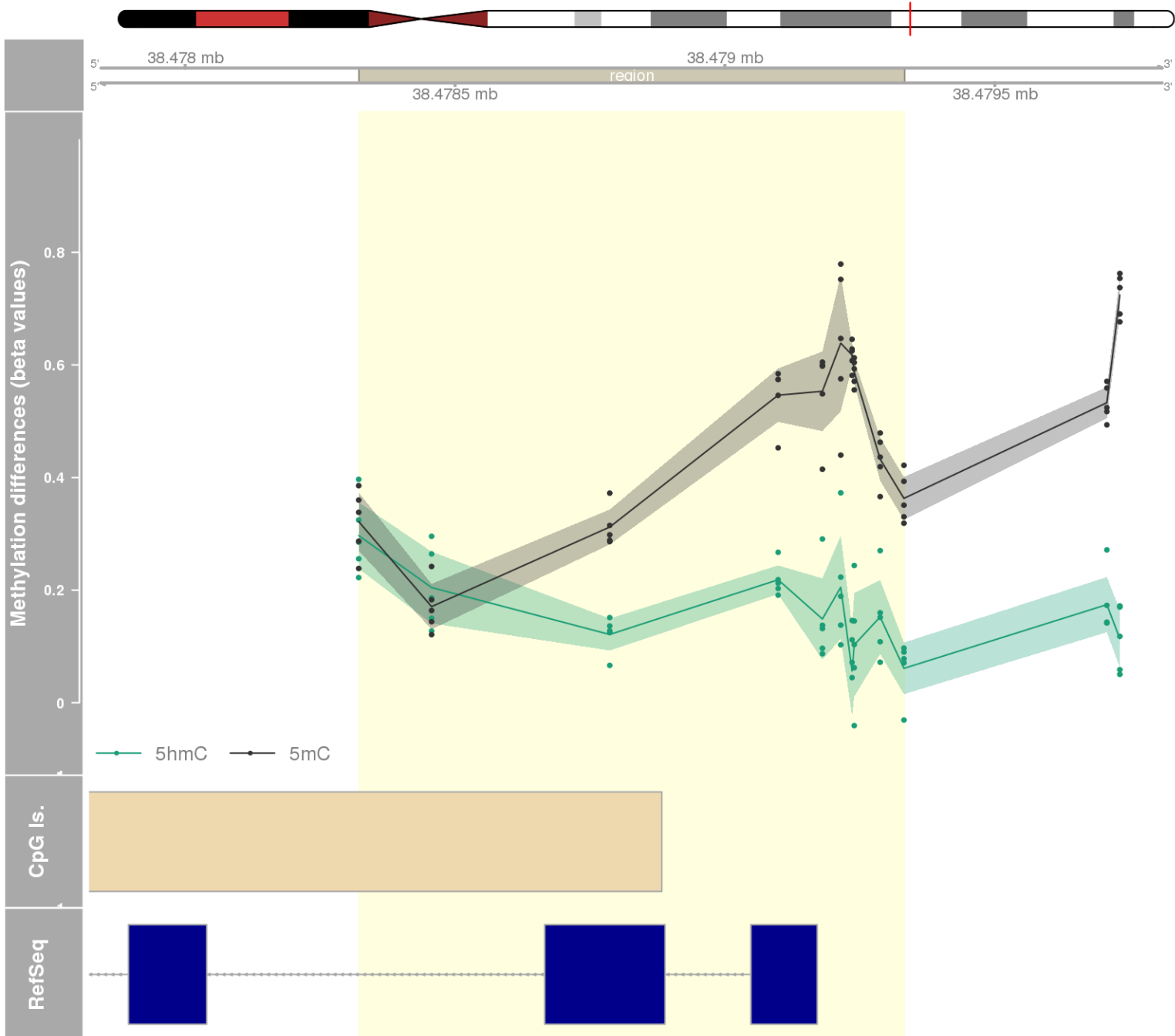
DMR 276 // chr13:113718119-113720634 // 2515 pb. (9 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.019 // fwerArea: 1
- genes: MCF2L -



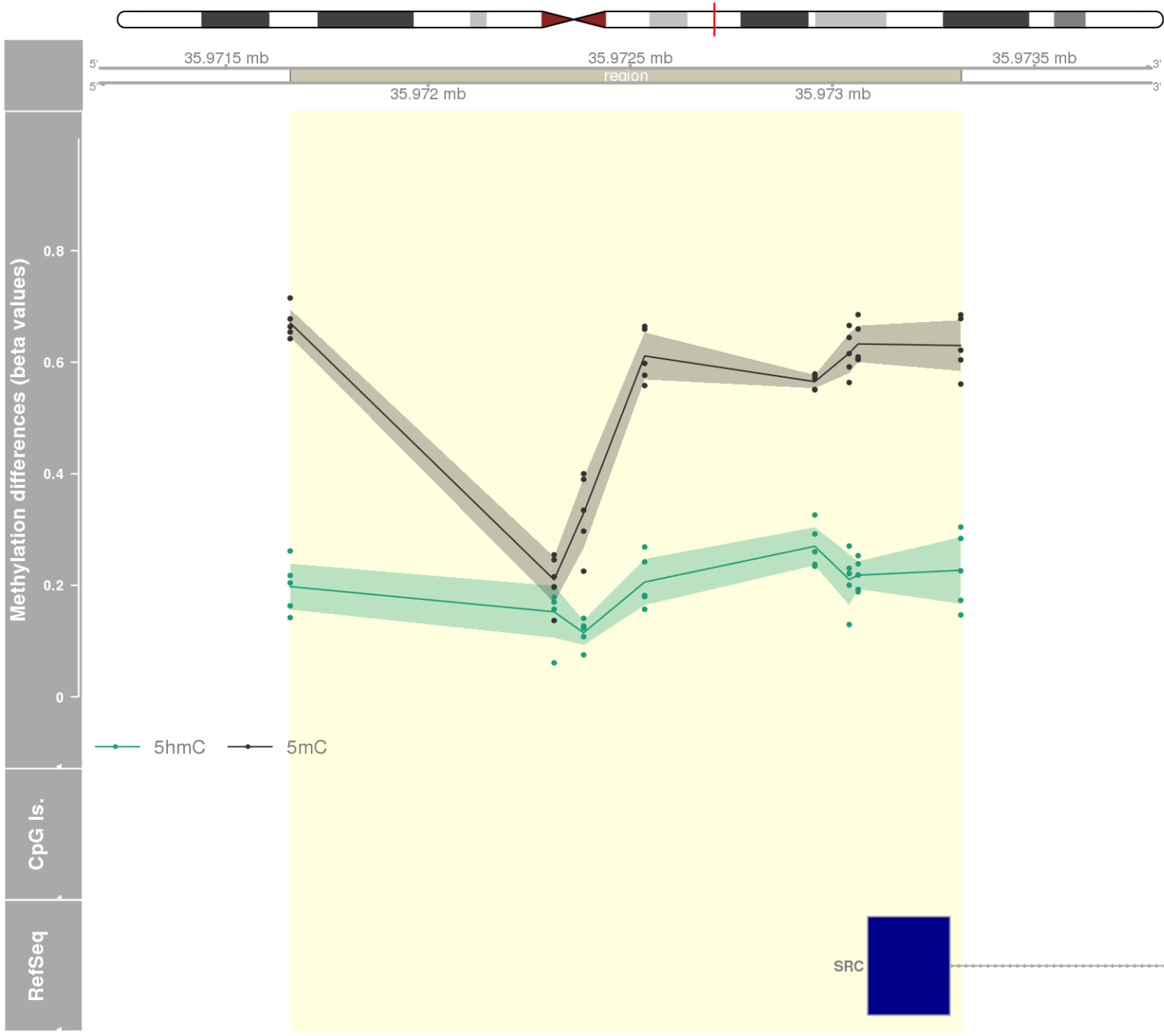
DMR 277 // chr7:134832221-134833166 // 945 pb. (9 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.019 // fwerArea: 1
- genes: AGLB3 / C7orf49 / TMEM140 -



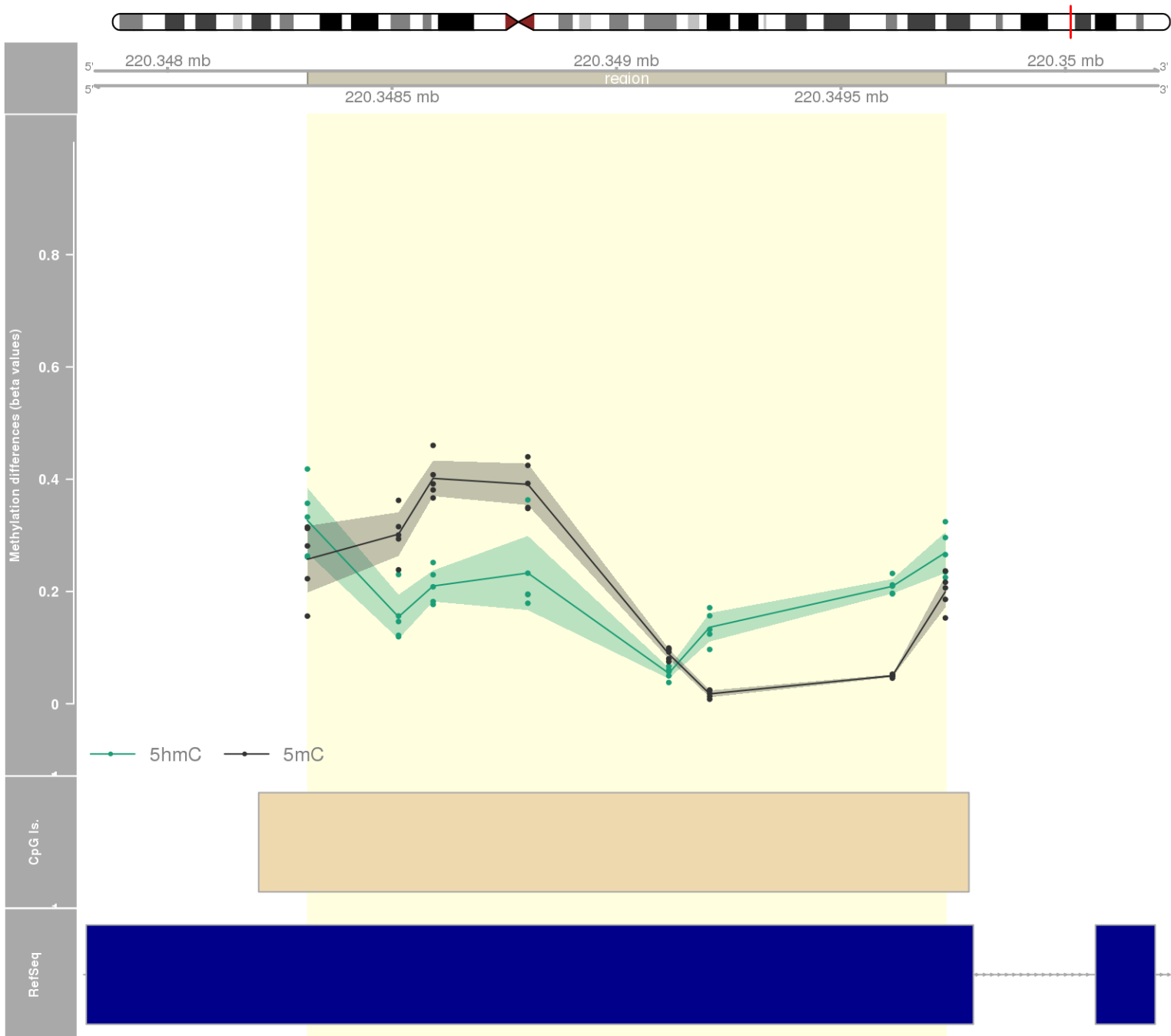
DMR 278 // chr22:38478322-38479332 // 1010 pb. (10 probes) // pvalue: 0.01 // fwer: 0.36 // pvalueArea: 0.02 // fwerArea: 1
- genes: SLC16A8 -



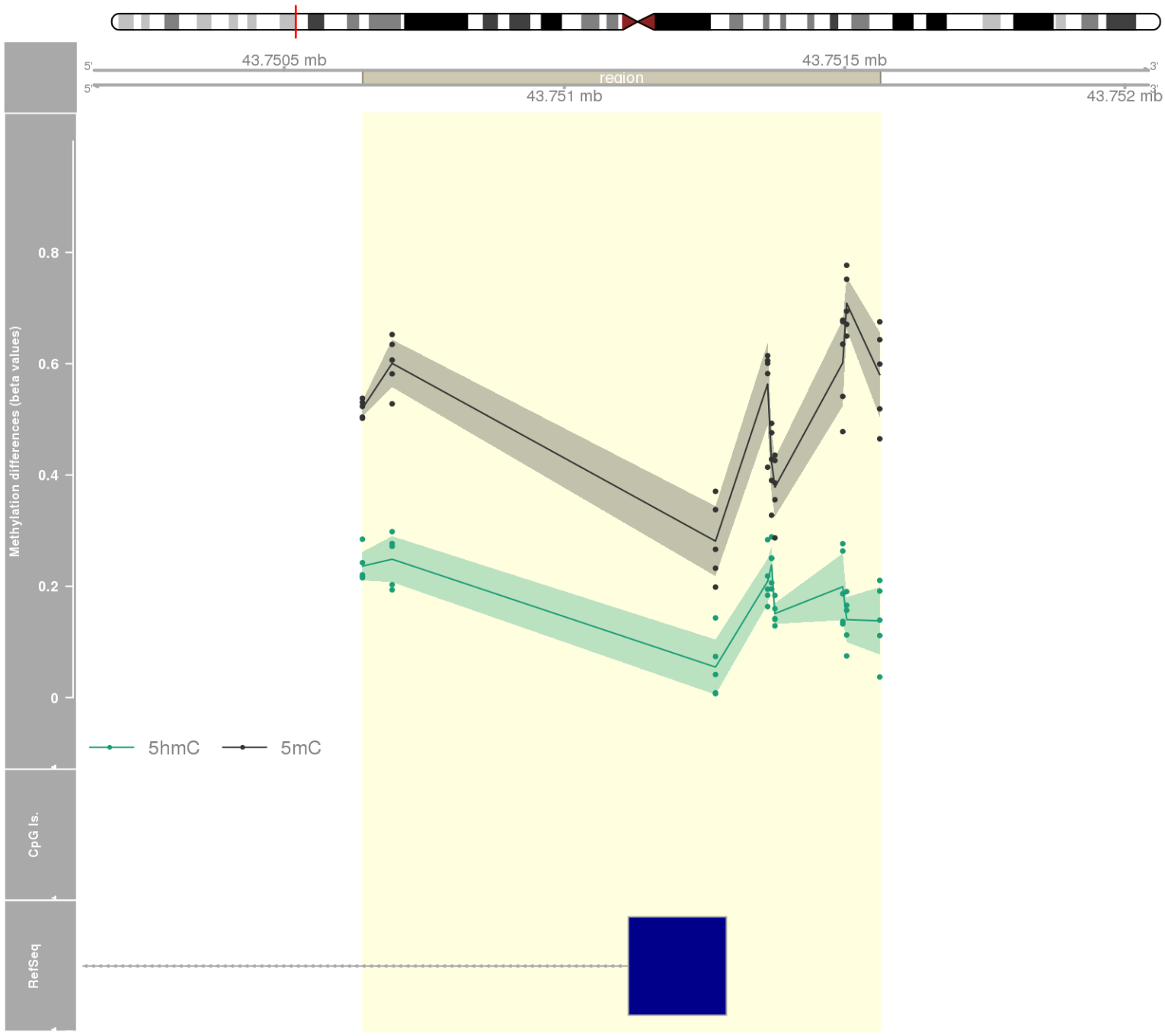
DMR 279 // chr20:35971659-35973318 // 1659 pb. (8 probes) // pvalue: 0.006 // fwer: 0.173 // pvalueArea: 0.02 // fwerArea: 1
- genes: SRC -



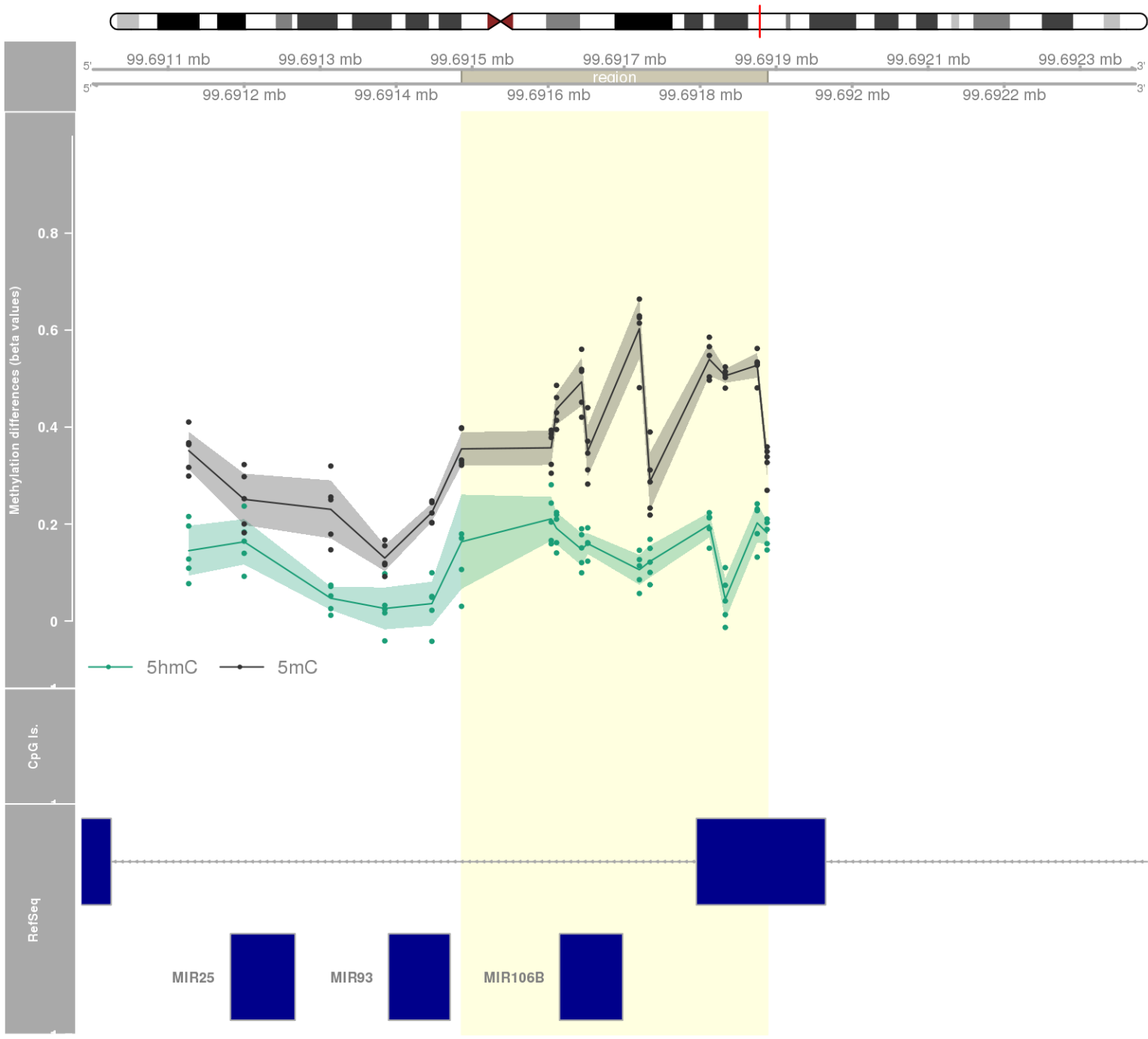
DMR 280 // chr2:220348312-220349733 // 1421 pb. (8 probes) // pvalue: 0.006 // fwer: 0.173 // pvalueArea: 0.02 // fwerArea: 1
- genes: SPEG -



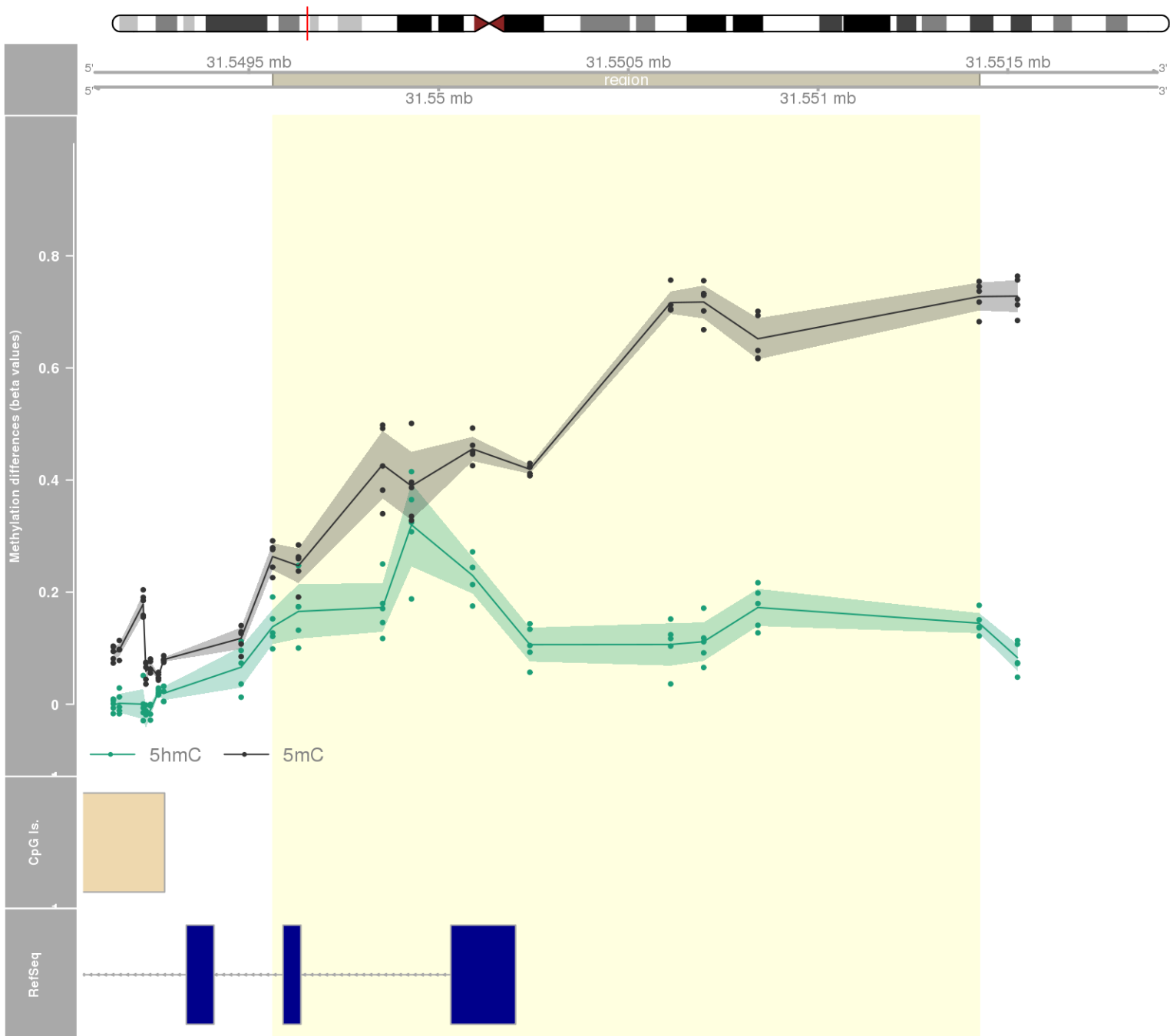
DMR 281 // chr1:43750640-43751563 // 923 pb. (9 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.02 // fwerArea: 1
- genes: C1orf210 -



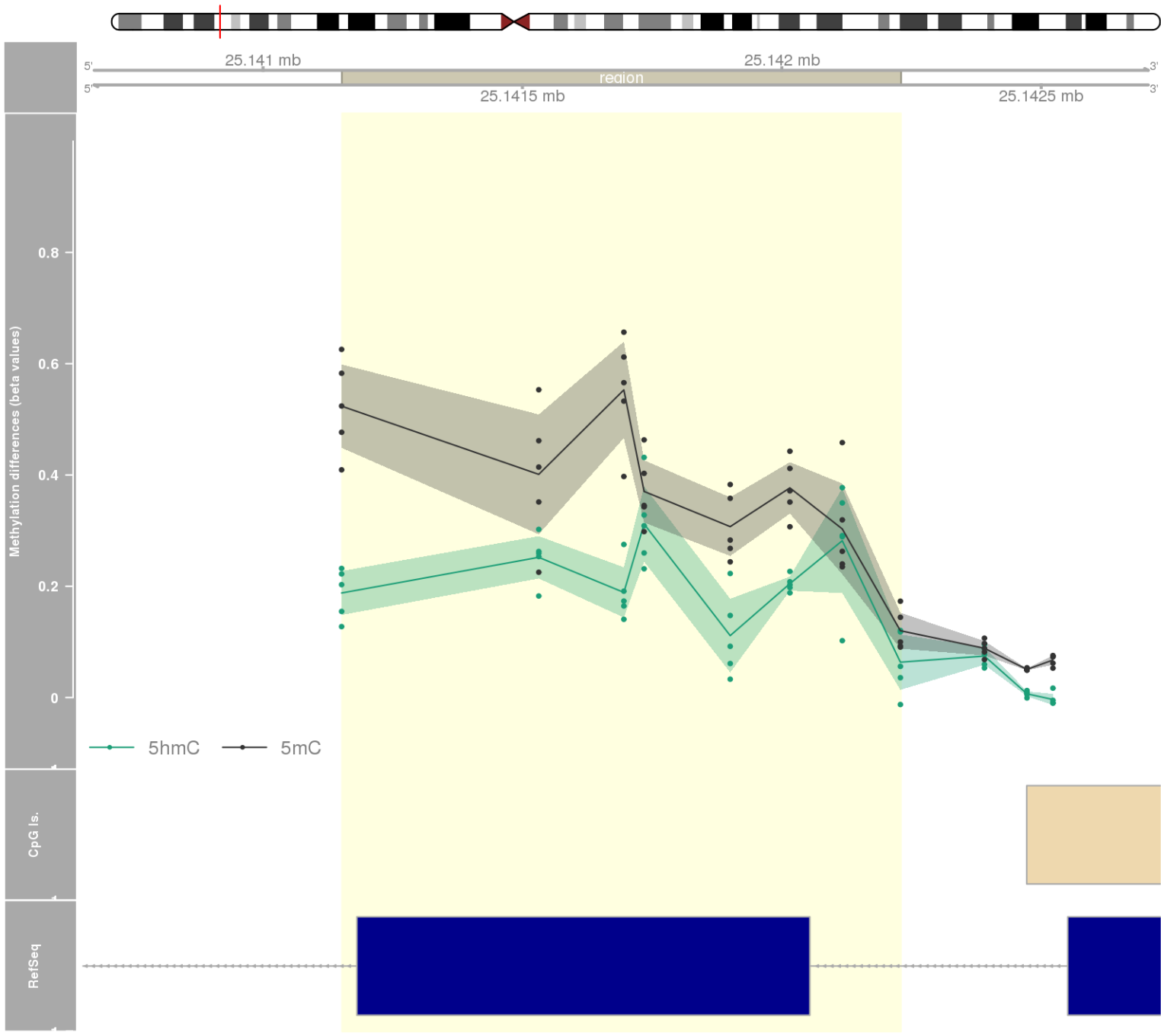
DMR 282 // chr7:99691486-99691888 // 402 pb. (11 probes) // pvalue: 0.01 // fwer: 0.36 // pvalueArea: 0.02 // fwerArea: 1
- genes: MIR106B / MCM7 -



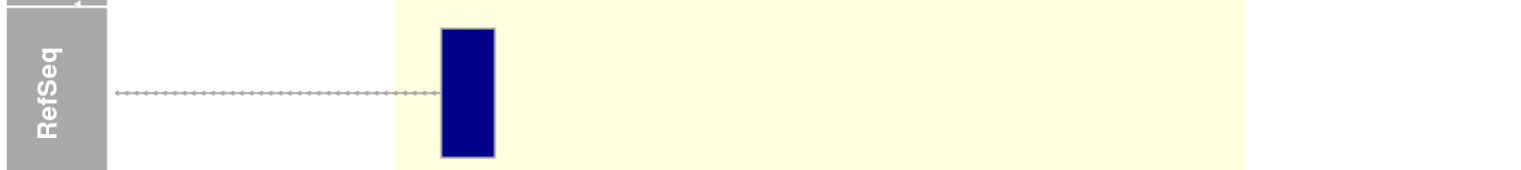
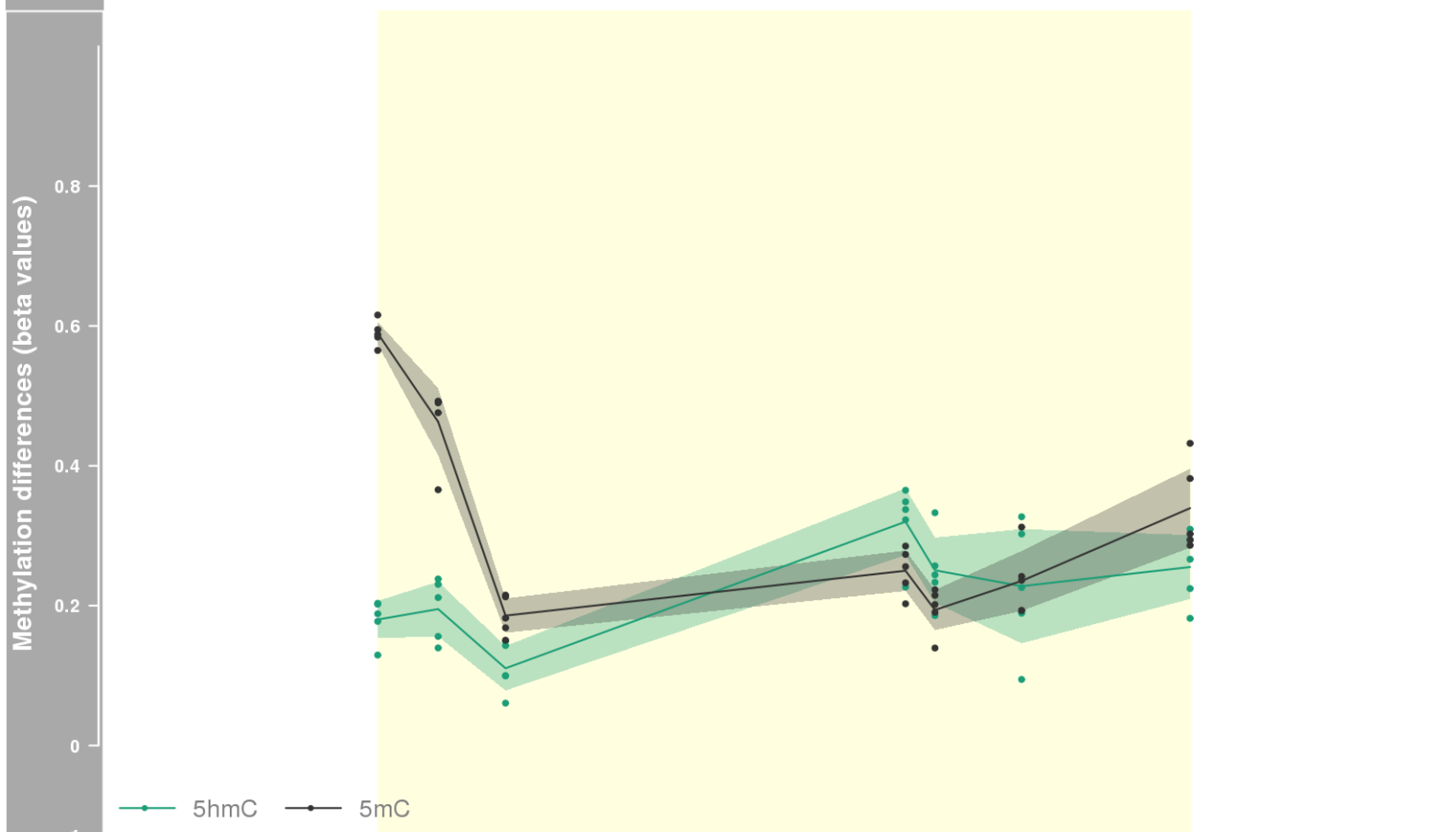
DMR 283 // chr6:31549563-31551425 // 1862 pb. (10 probes) // pvalue: 0.01 // fwer: 0.36 // pvalueArea: 0.02 // fwerArea: 1
- genes: LTB -



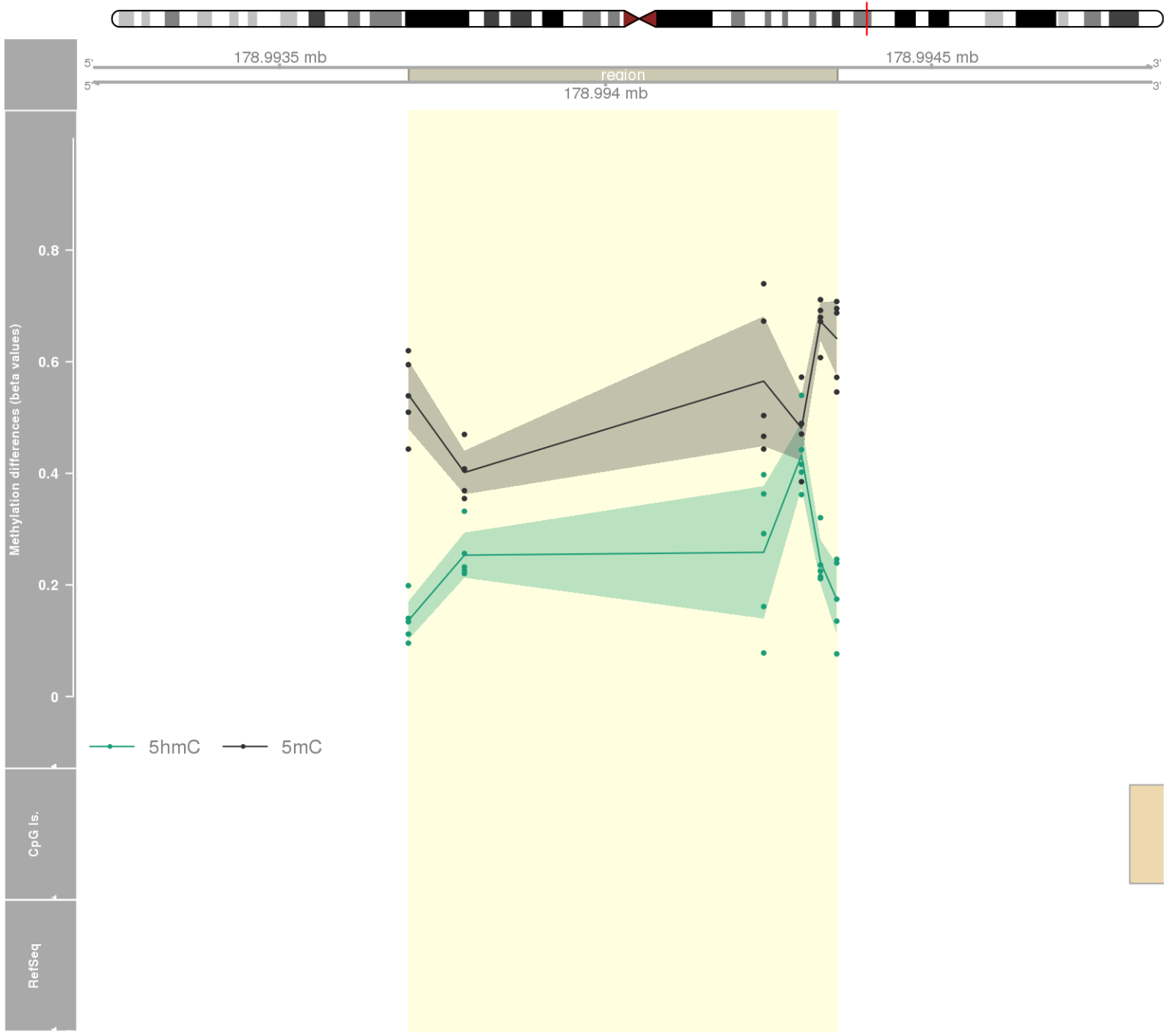
DMR 284 // chr2:25141152-25142229 // 1077 pb. (8 probes) // pvalue: 0.006 // fwer: 0.173 // pvalueArea: 0.02 // fwerArea: 1
- genes: ADCY3 -



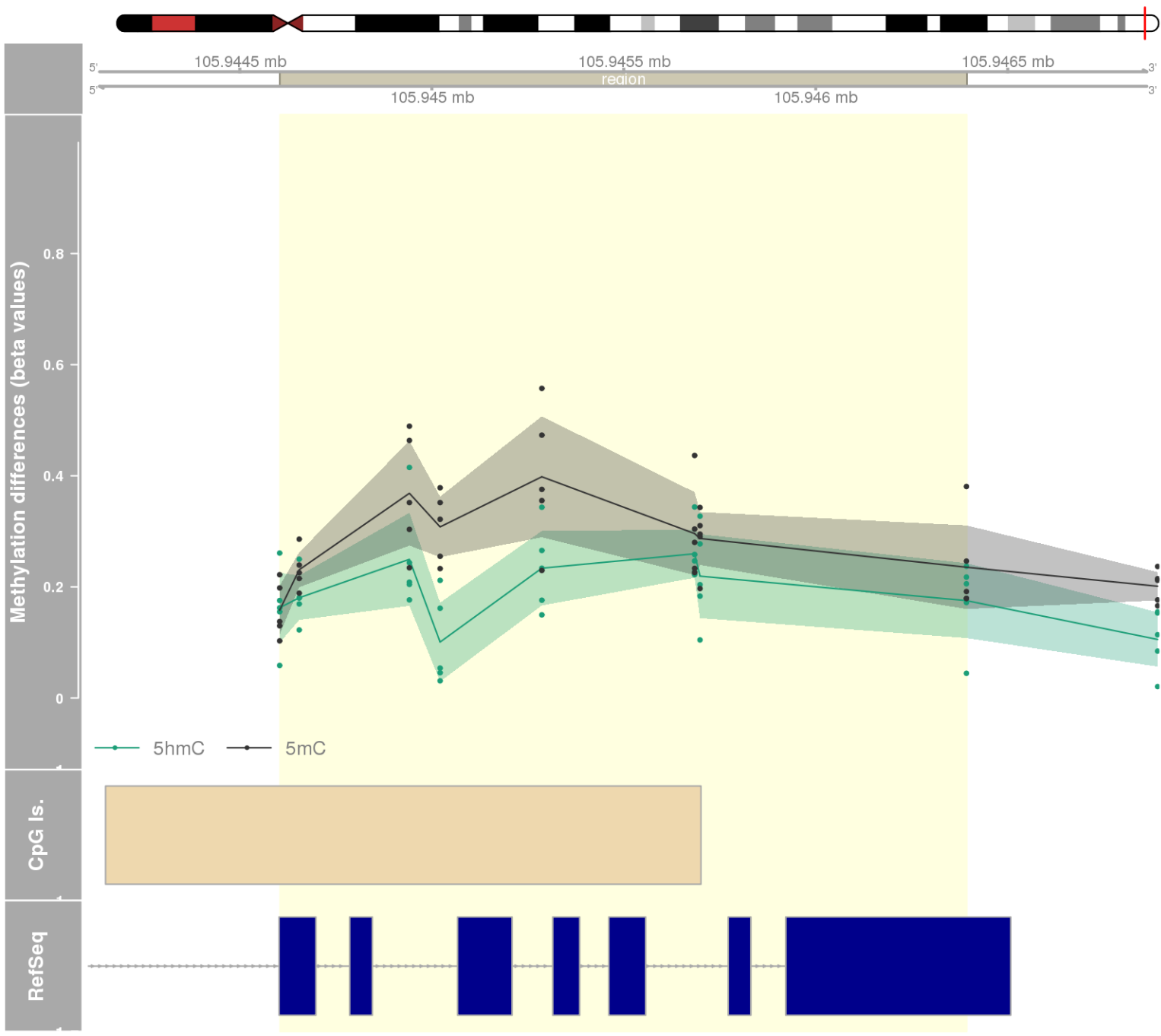
DMR 285 // chr12:58209674-58211193 // 1519 pb. (7 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.02 // fwerArea: 1
- genes: AVIL -



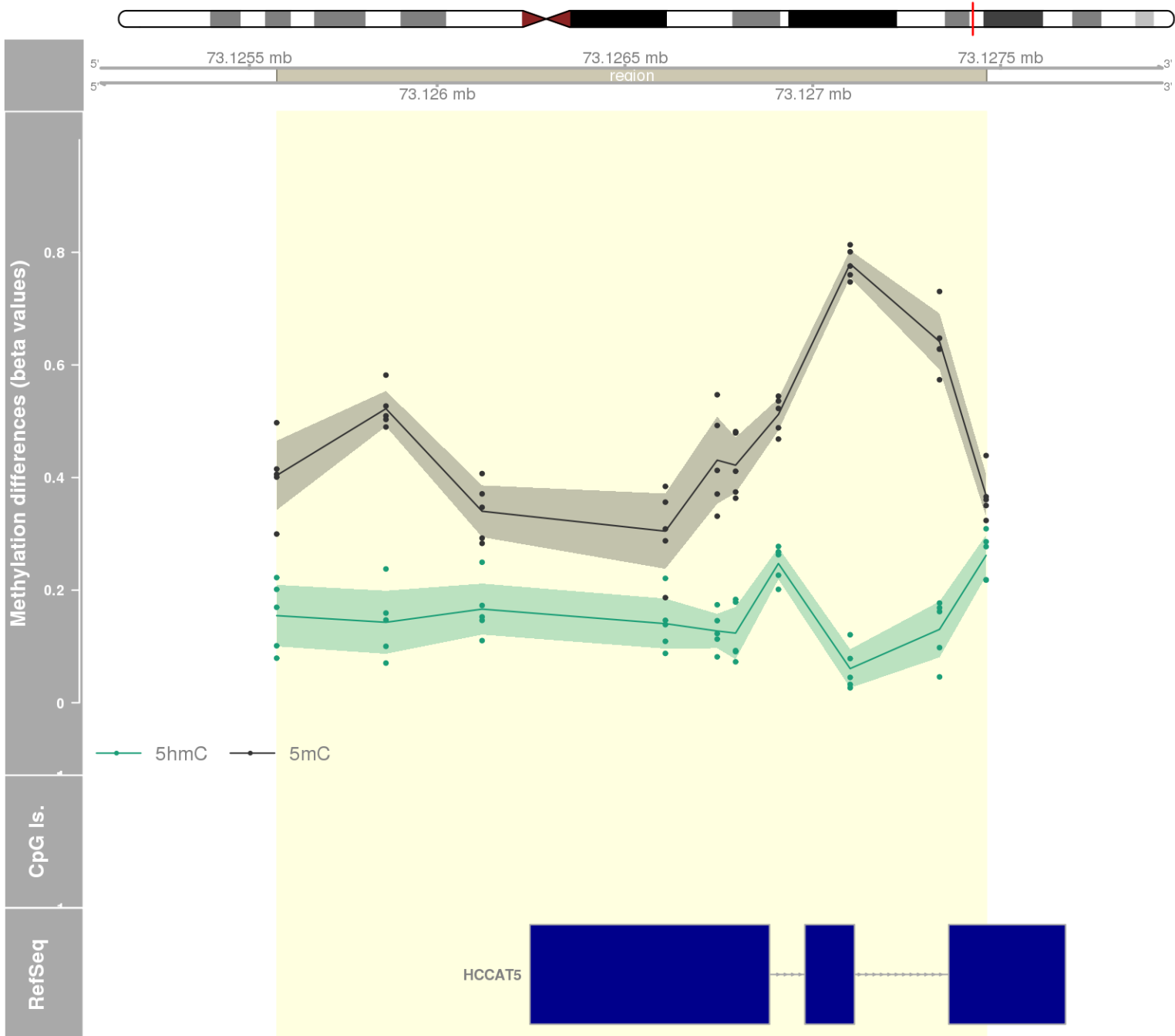
DMR 286 // chr1:178993698-178994355 // 657 pb. (6 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.02 // fwerArea: 1



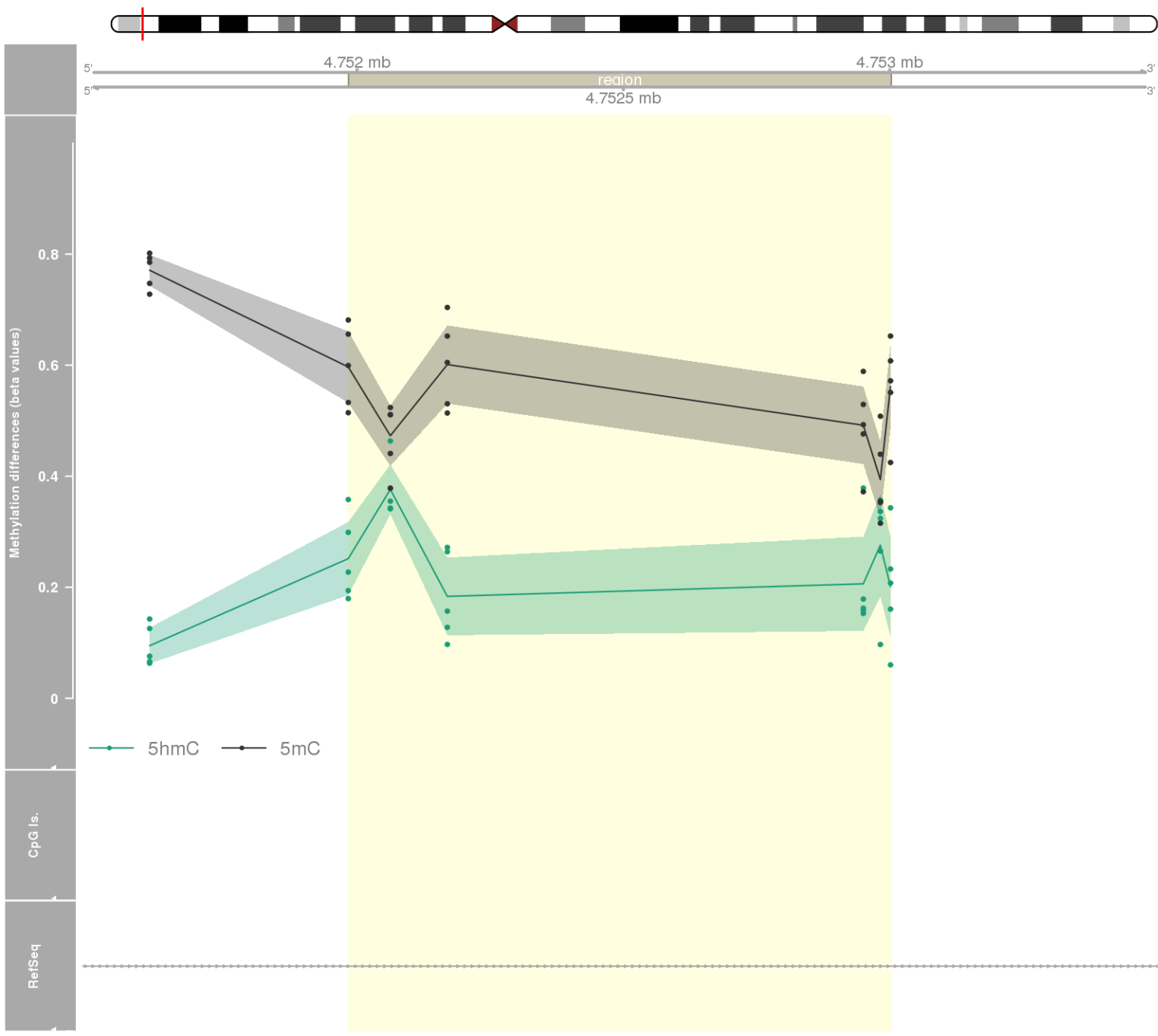
DMR 287 // chr14:105944604-105946393 // 1789 pb. (8 probes) // pvalue: 0.006 // fwer: 0.173 // pvalueArea: 0.02 // fwerArea: 1
- genes: CRIP2 -



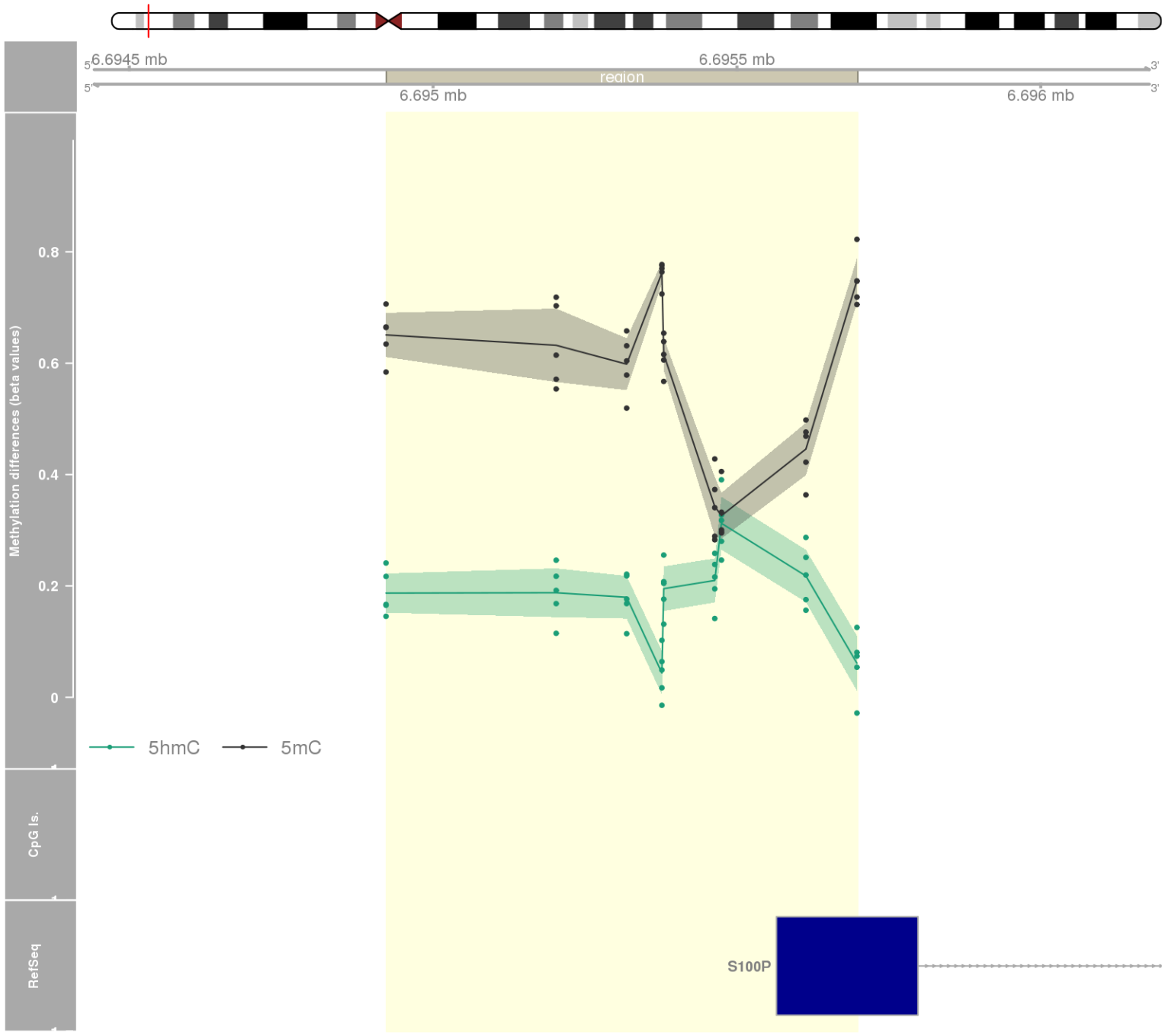
DMR 288 // chr16:73125573-73127462 // 1889 pb. (10 probes) // pvalue: 0.01 // fwer: 0.36 // pvalueArea: 0.02 // fwerArea: 1
- genes: HCCAT5 -



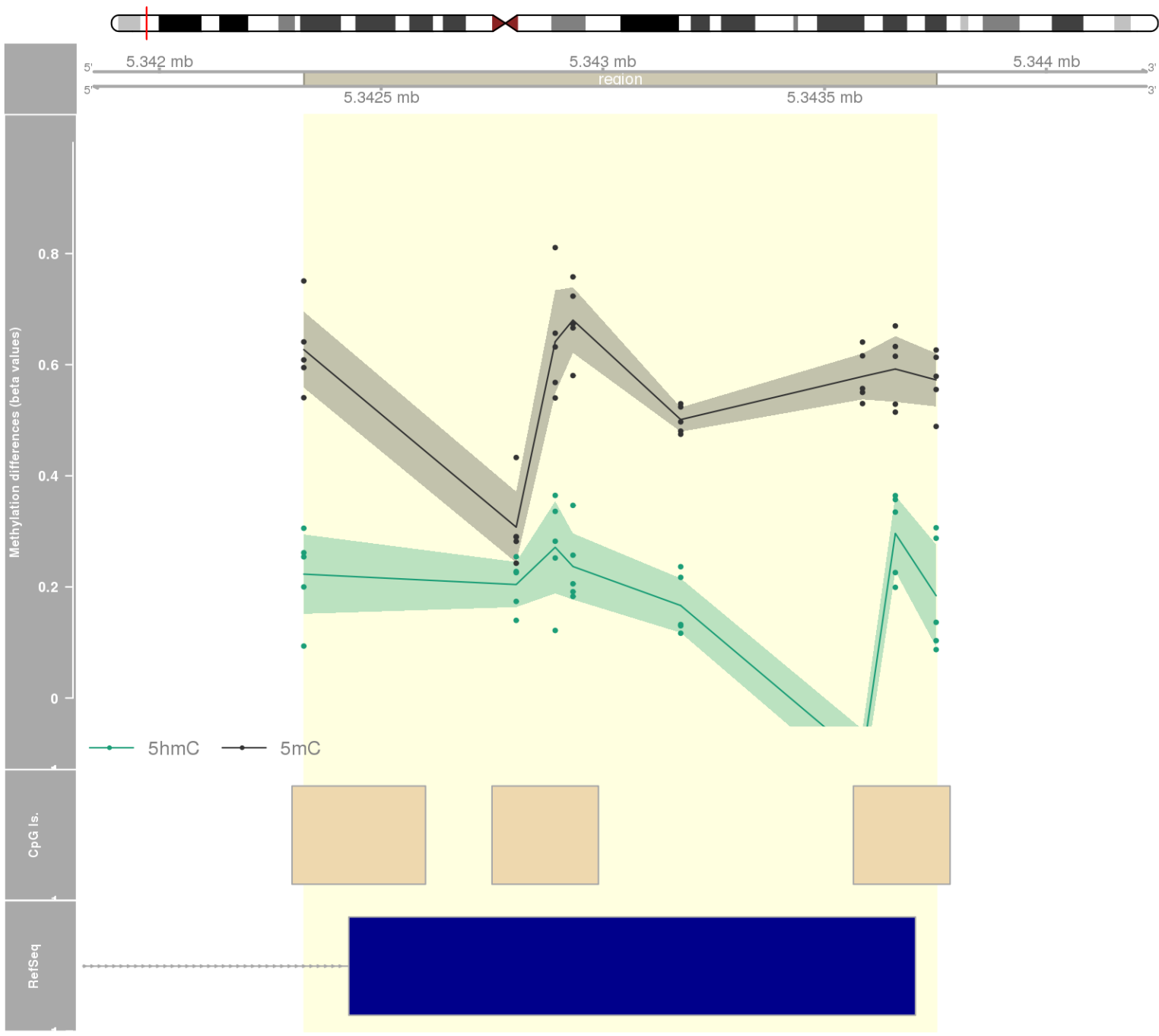
DMR 289 // chr7:4751984-4753002 // 1018 pb. (6 probes) // pvalue: 0.001 // fwer: 0.173 // pvalueArea: 0.02 // fwerArea: 1
- genes: FO XK1 -



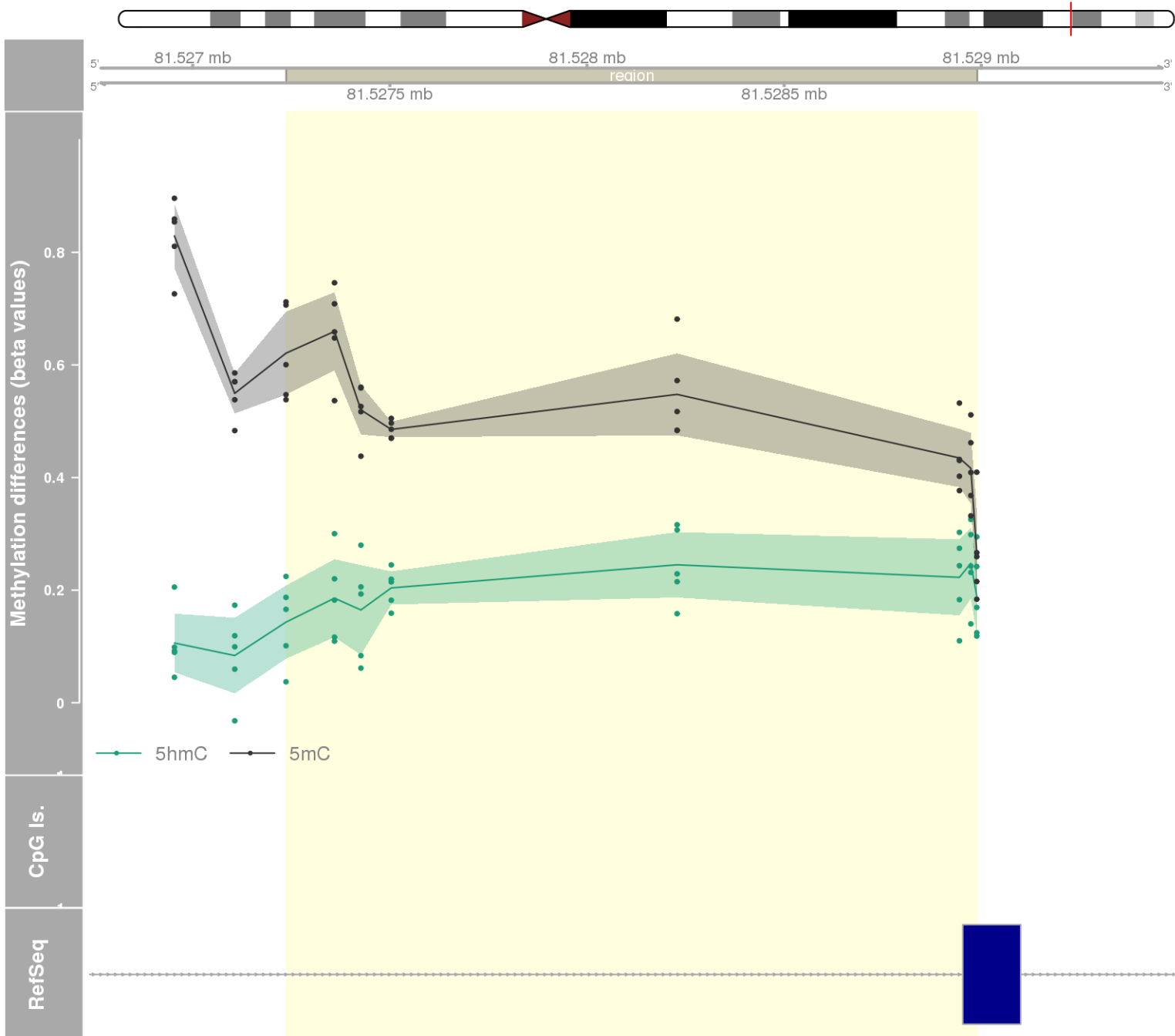
DMR 290 // chr4:6694923-6695698 // 775 pb. (9 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.02 // fwerArea: 1
- genes: S100P -



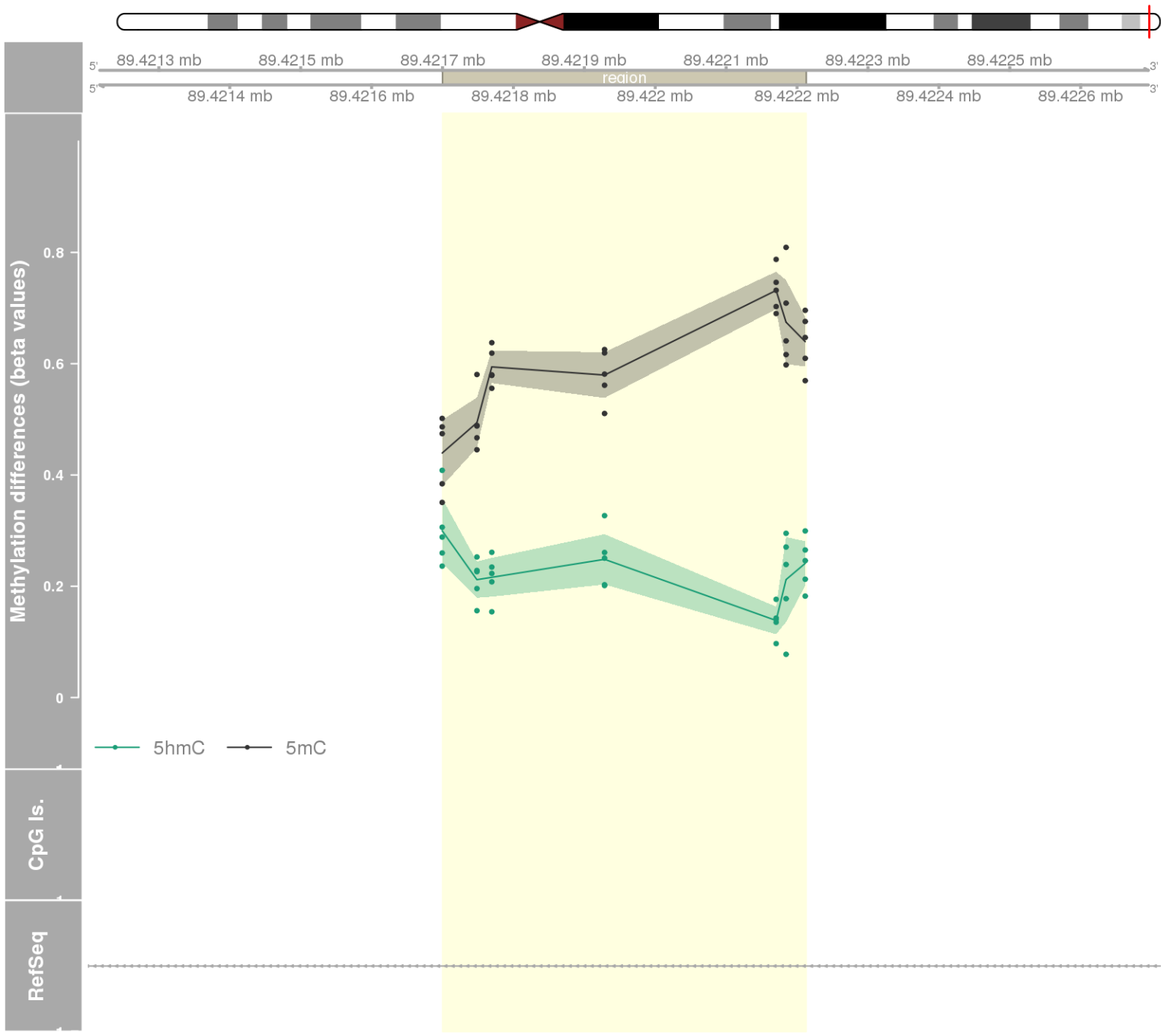
DMR 291 // chr7:5342326-5343752 // 1426 pb. (8 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.02 // fwerArea: 1
- genes: SLC29A4 -



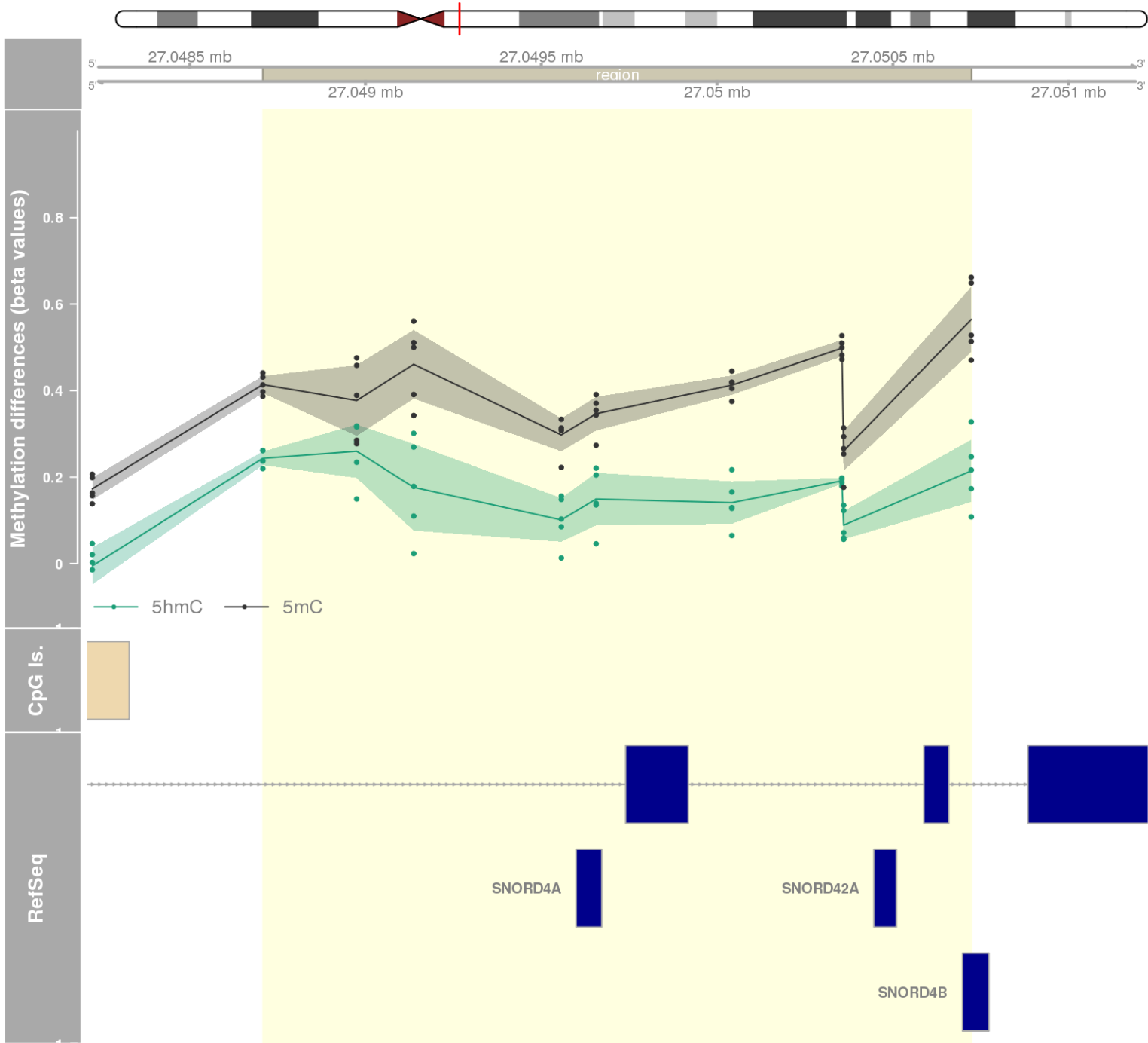
DMR 292 // chr16:81527237-81528989 // 1752 pb. (8 probes) // pvalue: 0.006 // fwer: 0.173 // pvalueArea: 0.02 // fwerArea: 1
- genes: CMIP -



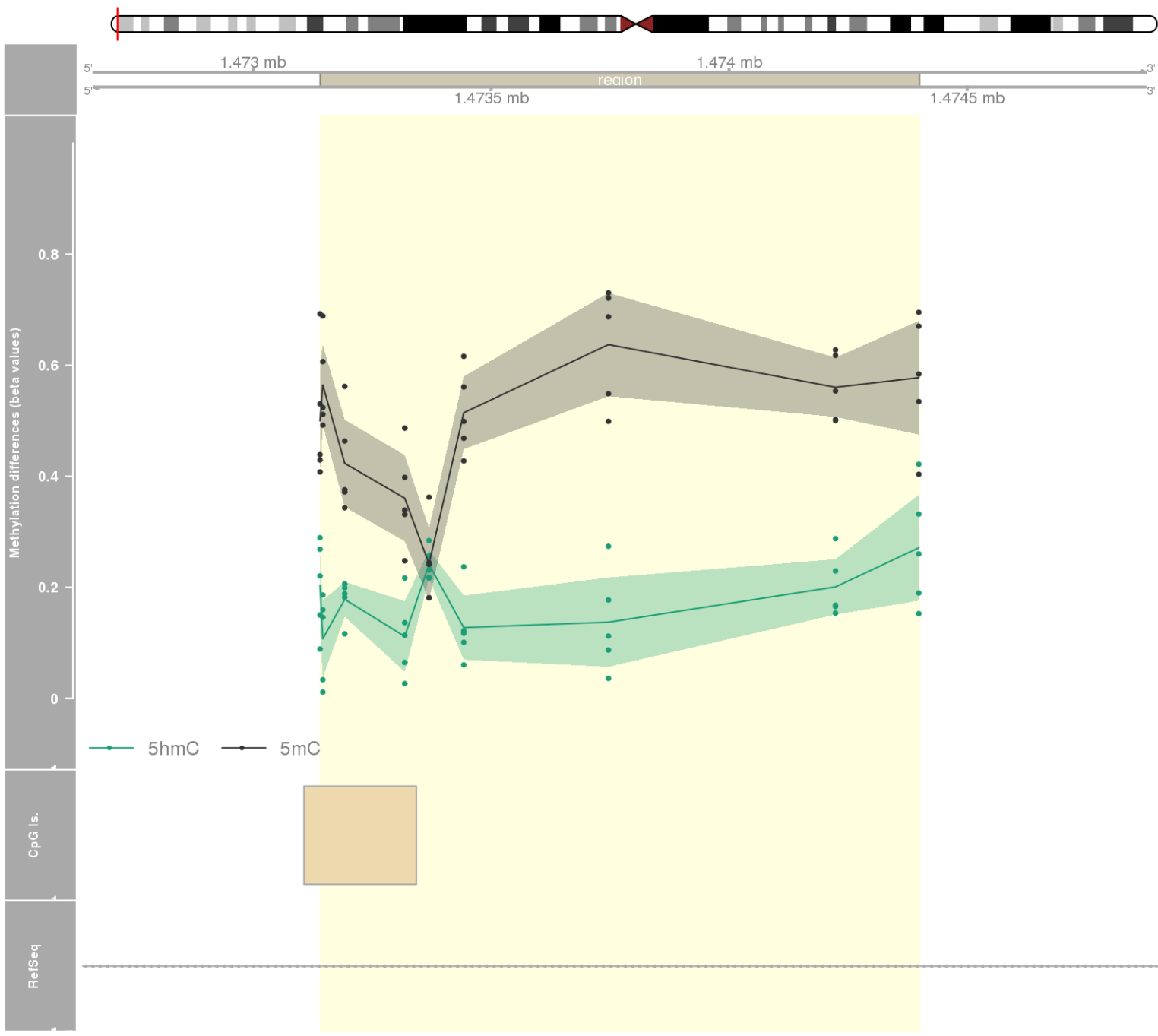
DMR 293 // chr16:89421700-89422212 // 512 pb. (7 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.021 // fwerArea: 1
- genes: ANKRD11 -



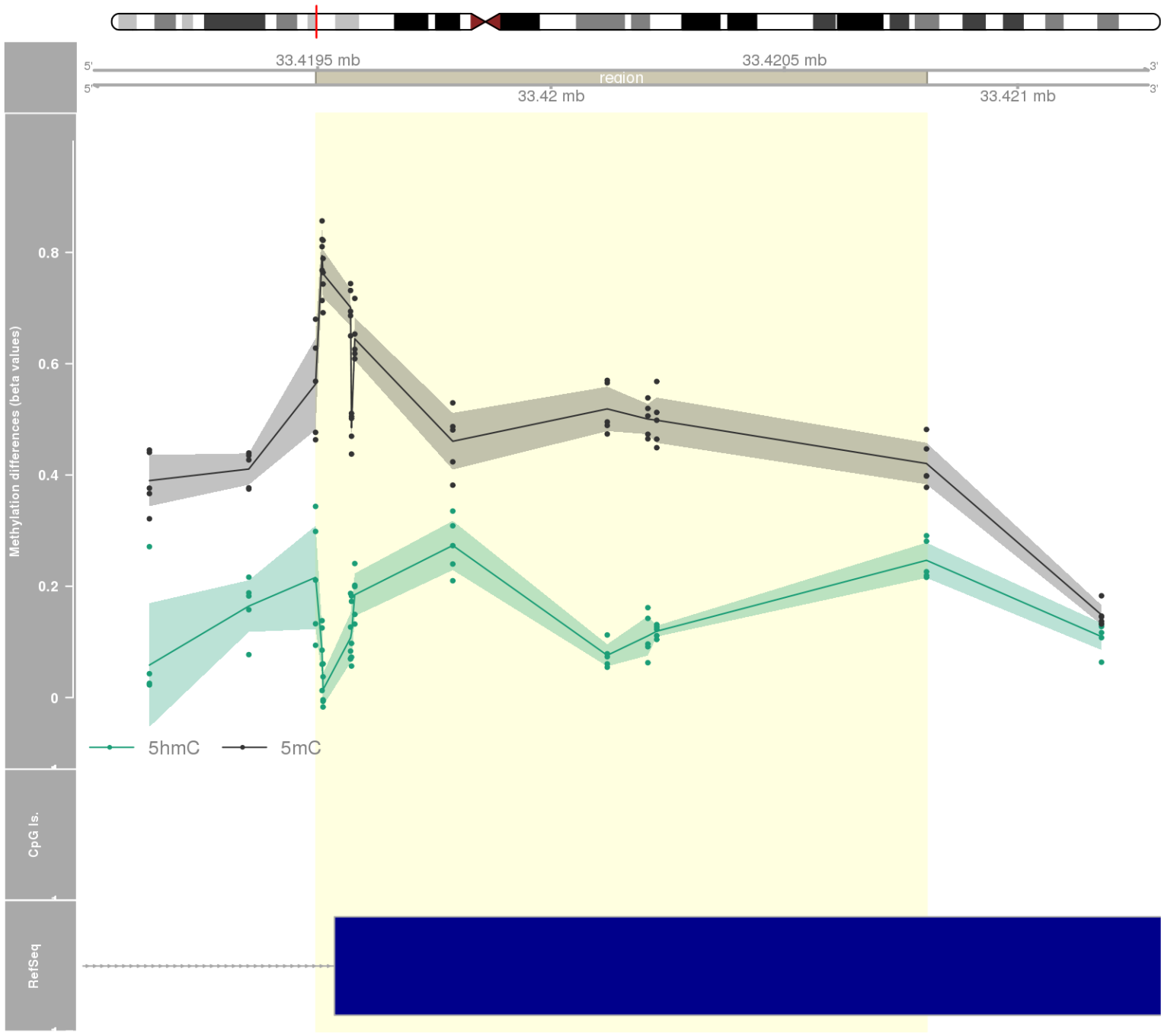
DMR 294 // chr17:27048708-27050723 // 2015 pb. (9 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.021 // fwerArea: 1
- genes: RPL23A / SNORD4A / SNORD42A / SNORD4B -



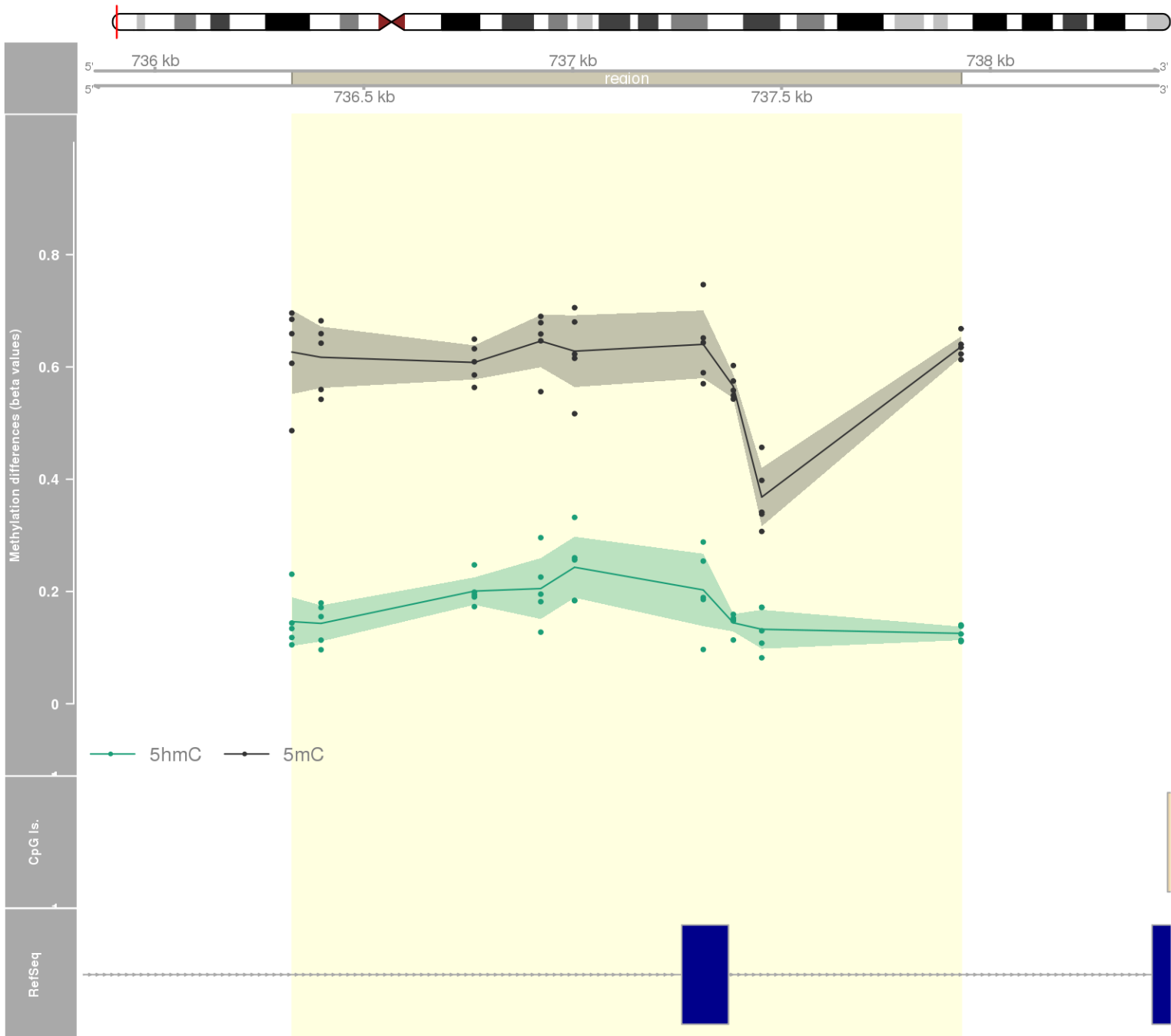
DMR 295 // chr1:1473141-1474399 // 1258 pb. (9 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.021 // fwerArea: 1
- genes: TMEM240 -



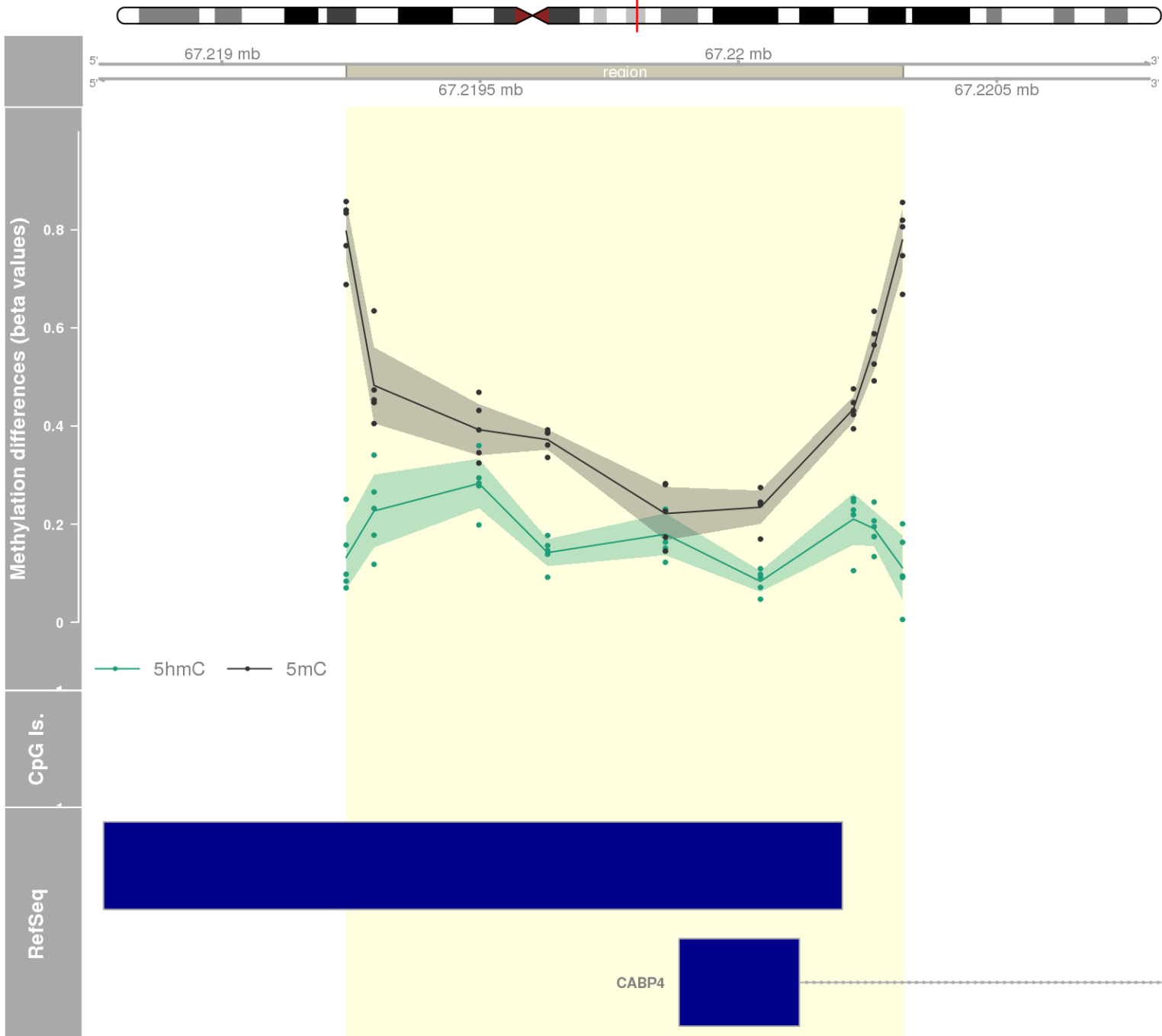
DMR 296 // chr6:33419496-33420805 // 1309 pb. (11 probes) // pvalue: 0.01 // fwer: 0.36 // pvalueArea: 0.021 // fwerArea: 1
- genes: SYNGAP1 -



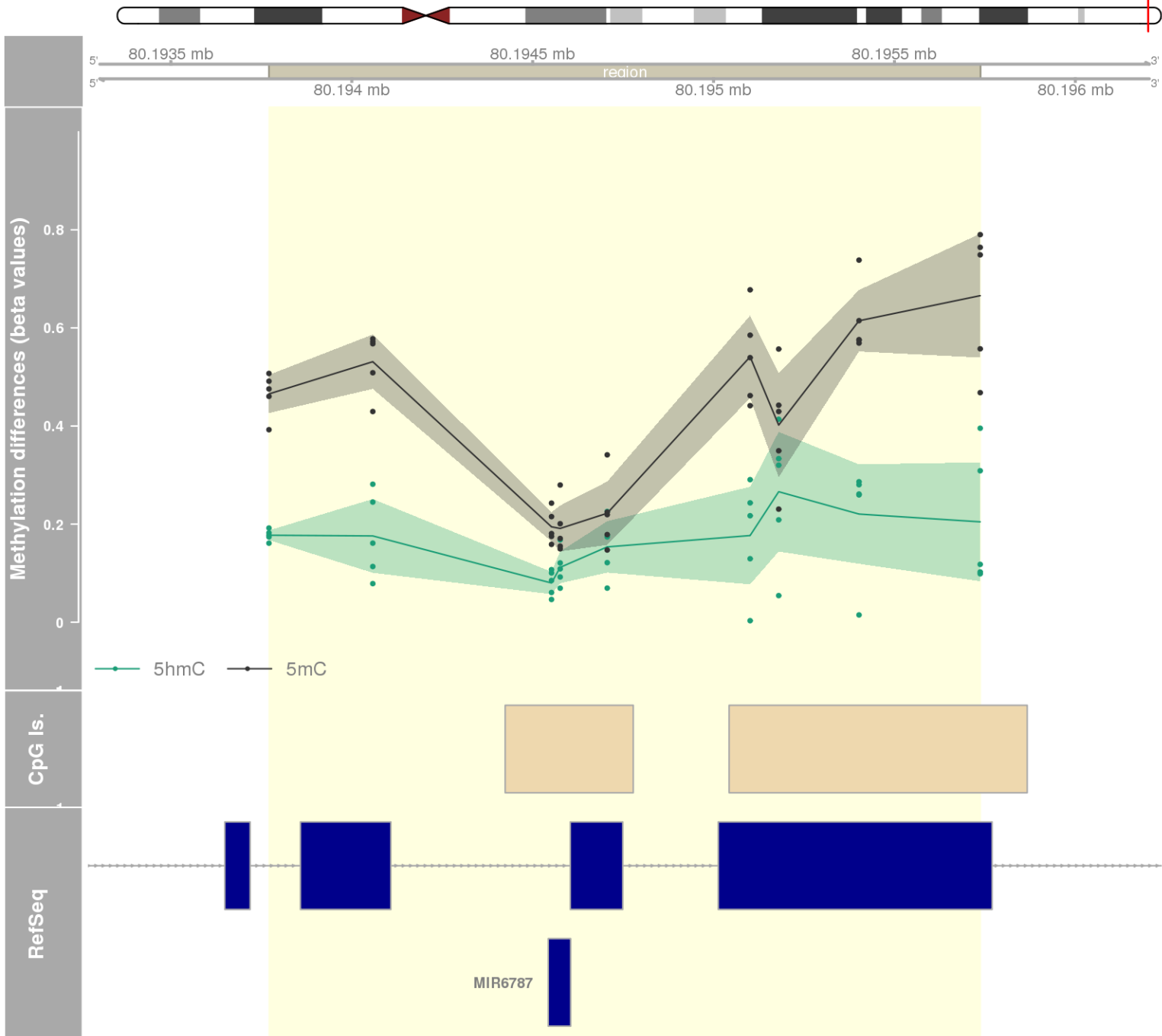
DMR 297 // chr4:736328-737930 // 1602 pb. (9 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.021 // fwerArea: 1
- genes: PCGF3 -



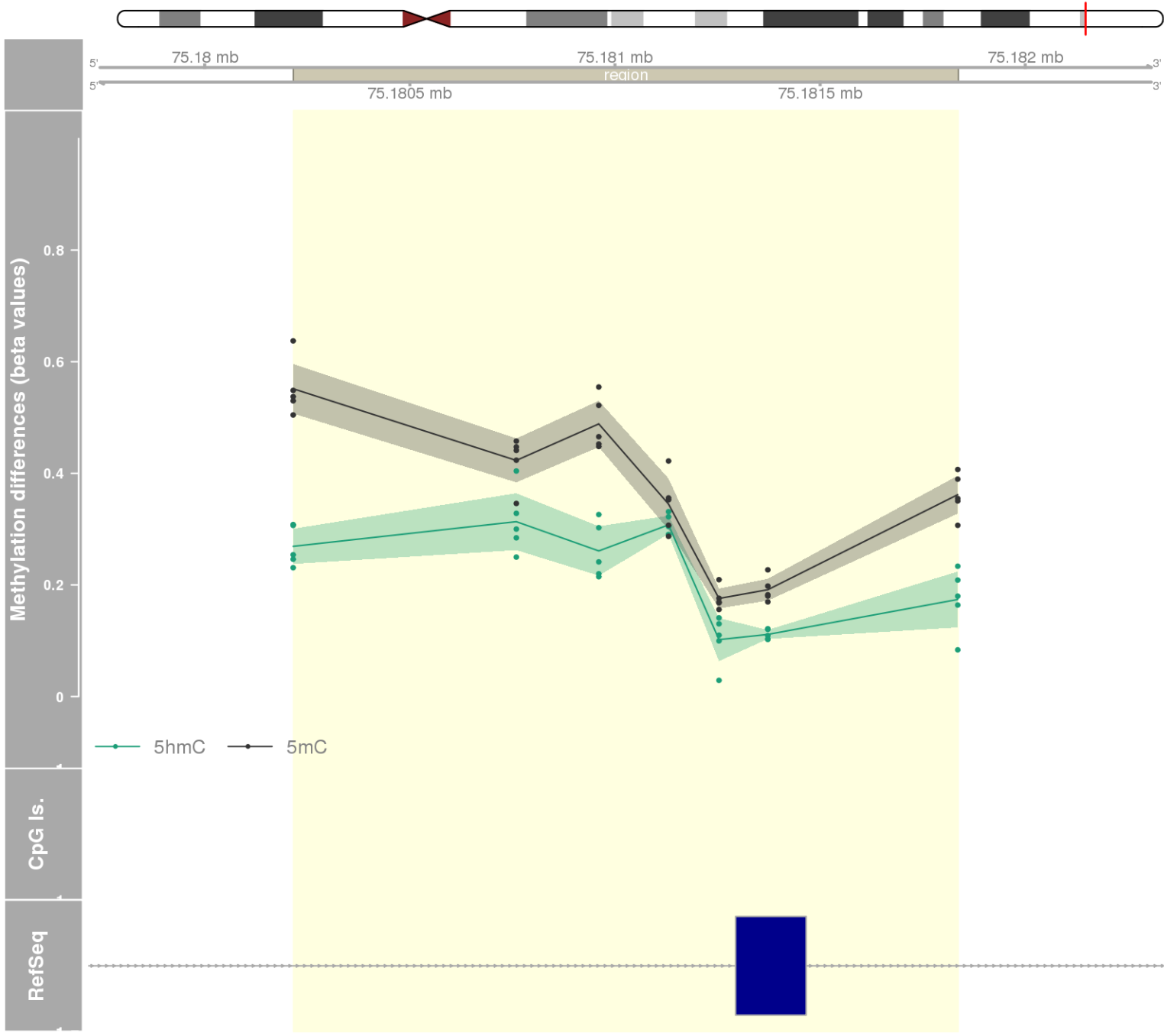
DMR 298 // chr11:67219241-67220318 // 1077 pb. (9 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.021 // fwerArea: 1
- genes: GPR152 / CABP4 -



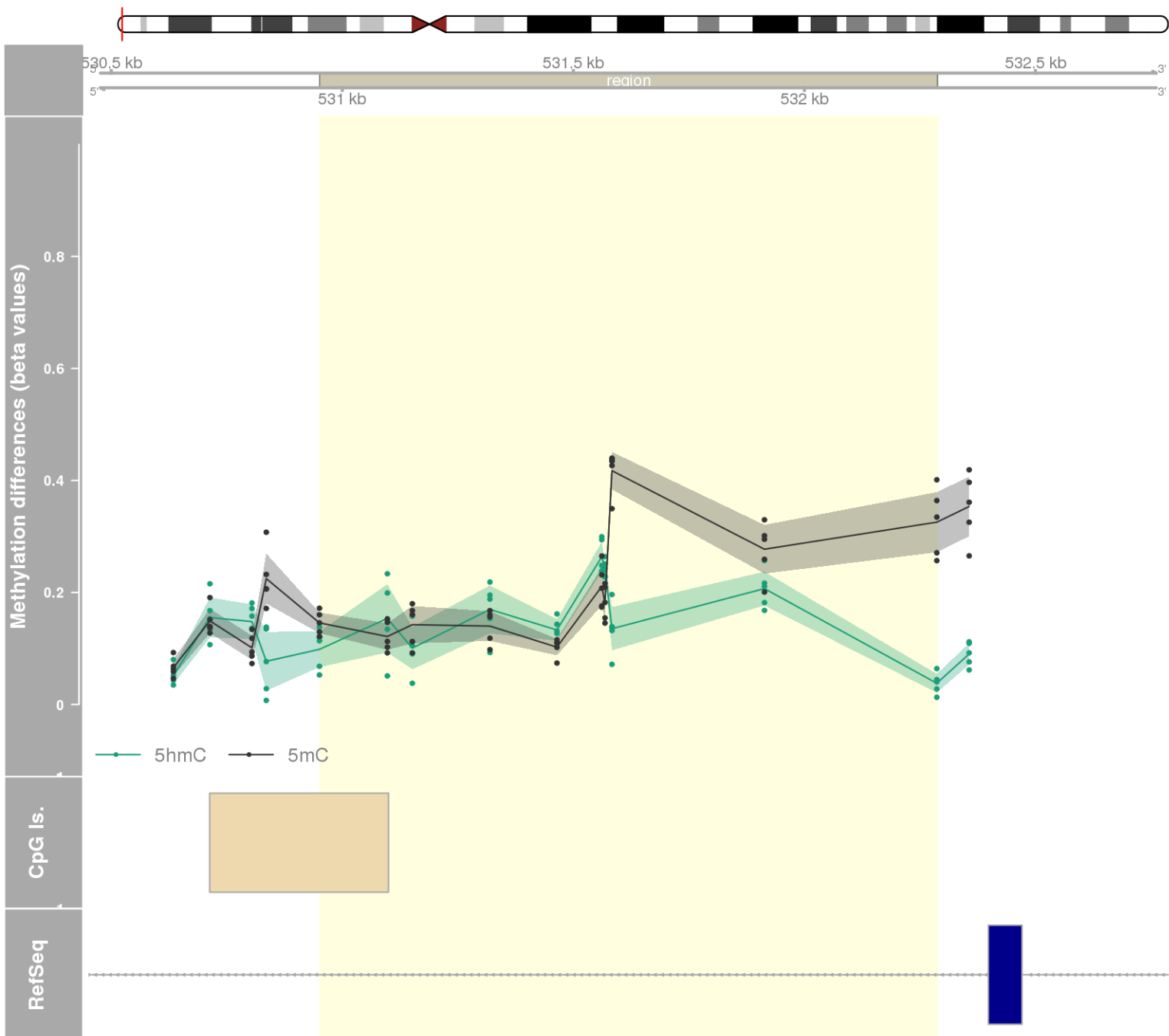
DMR 299 // chr17:80193771-80195737 // 1966 pb. (9 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.021 // fwerArea: 1
- genes: SLC16A3 -



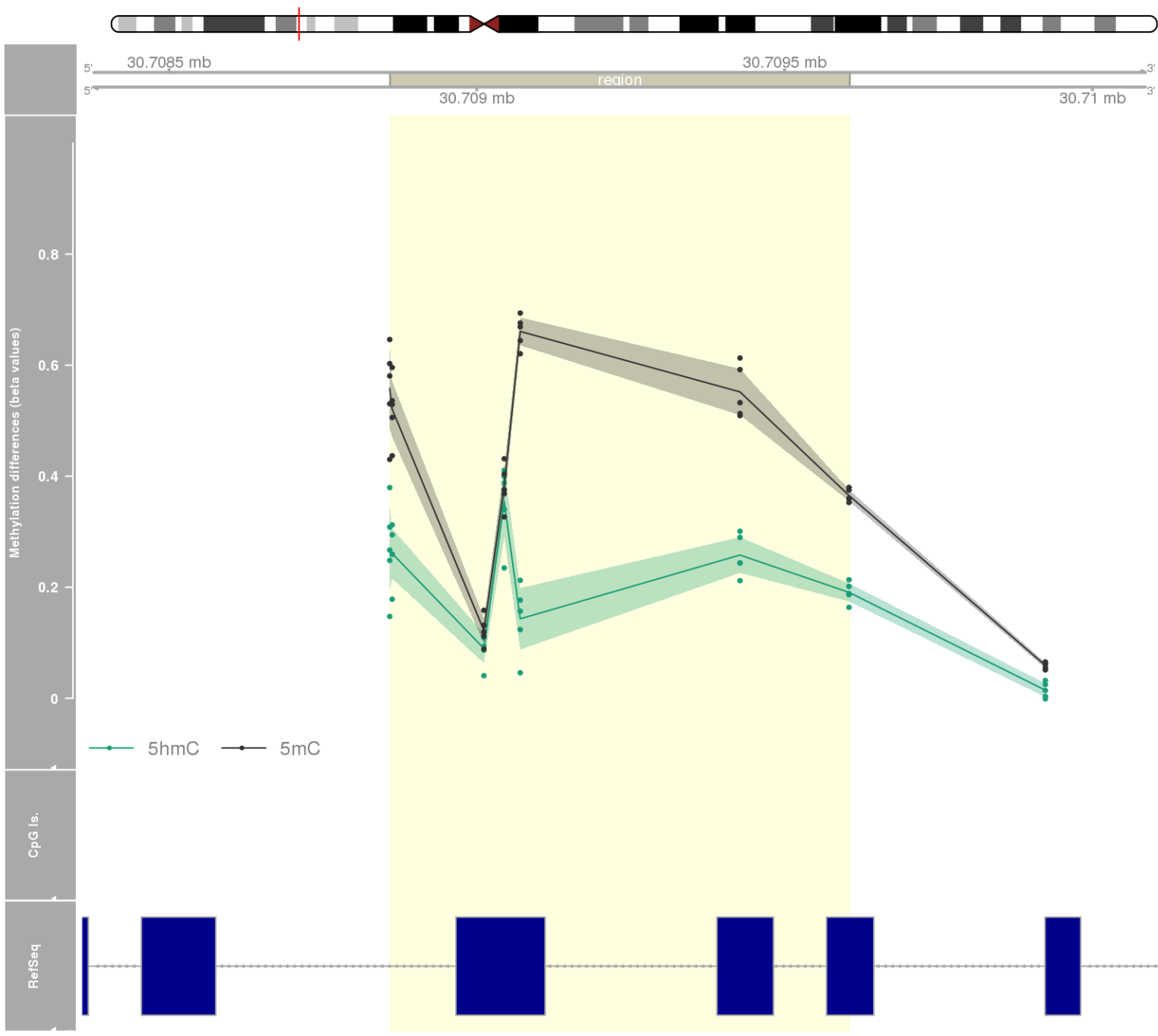
DMR 300 // chr17:75180216-75181836 // 1620 pb. (7 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.021 // fwerArea: 1
- genes: SEC14L1 -



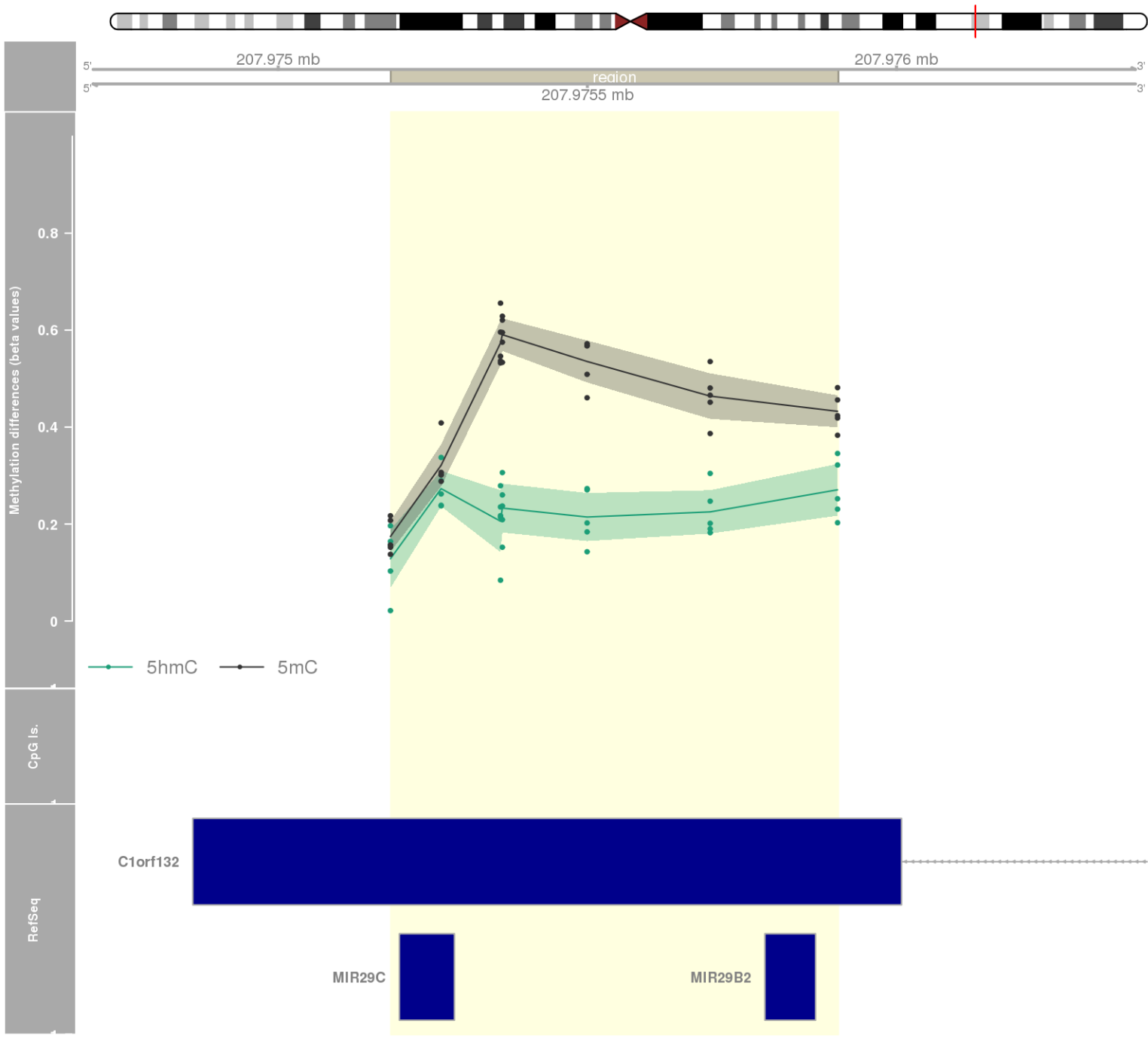
DMR 301 // chr10:530951-532287 // 1336 pb. (10 probes) // pvalue: 0.011 // fwer: 0.36 // pvalueArea: 0.021 // fwerArea: 1
- genes: DIP2C -



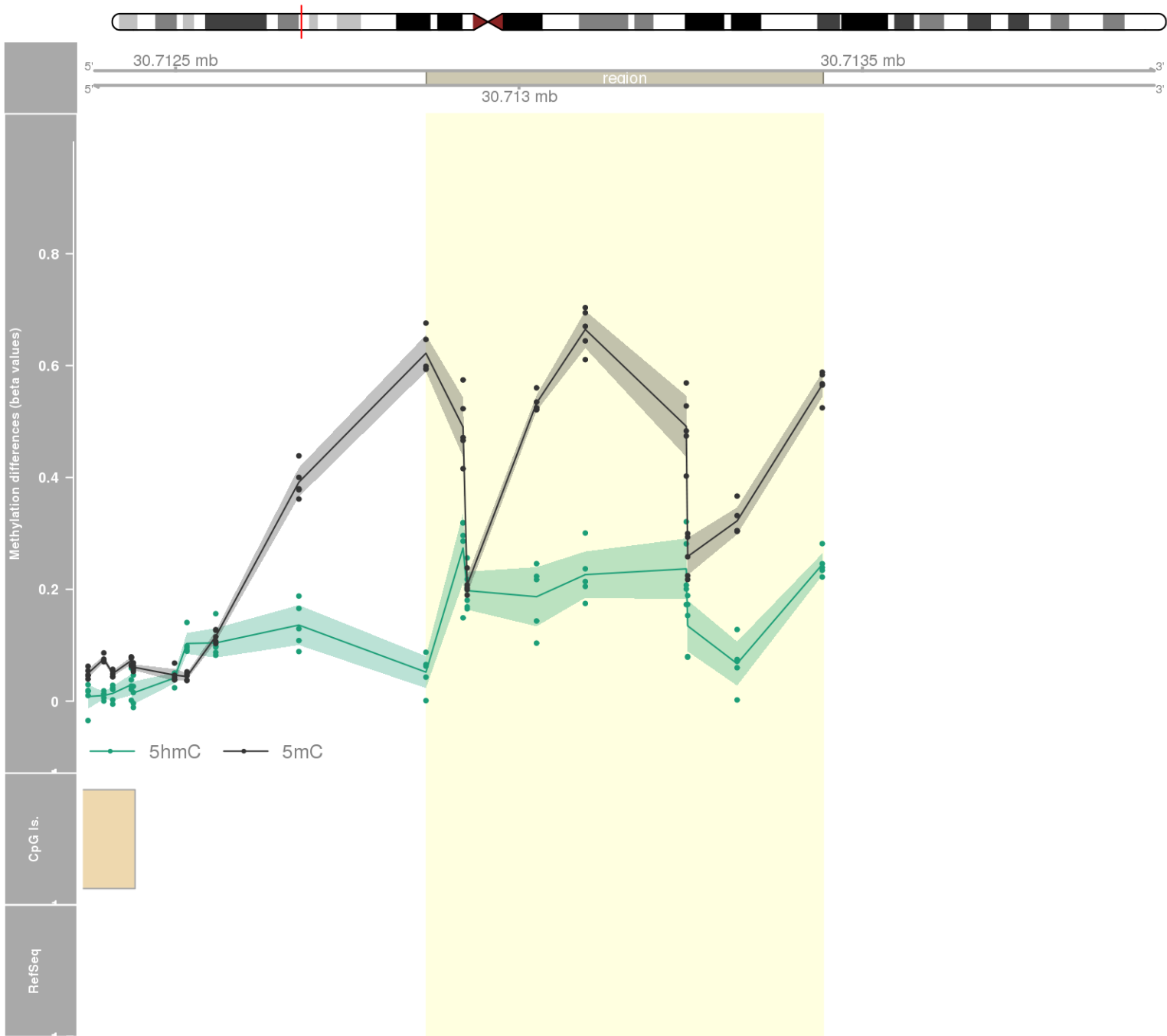
DMR 302 // chr6:30708859-30709605 // 746 pb. (7 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.021 // fwerArea: 1
- genes: FLOT1 -



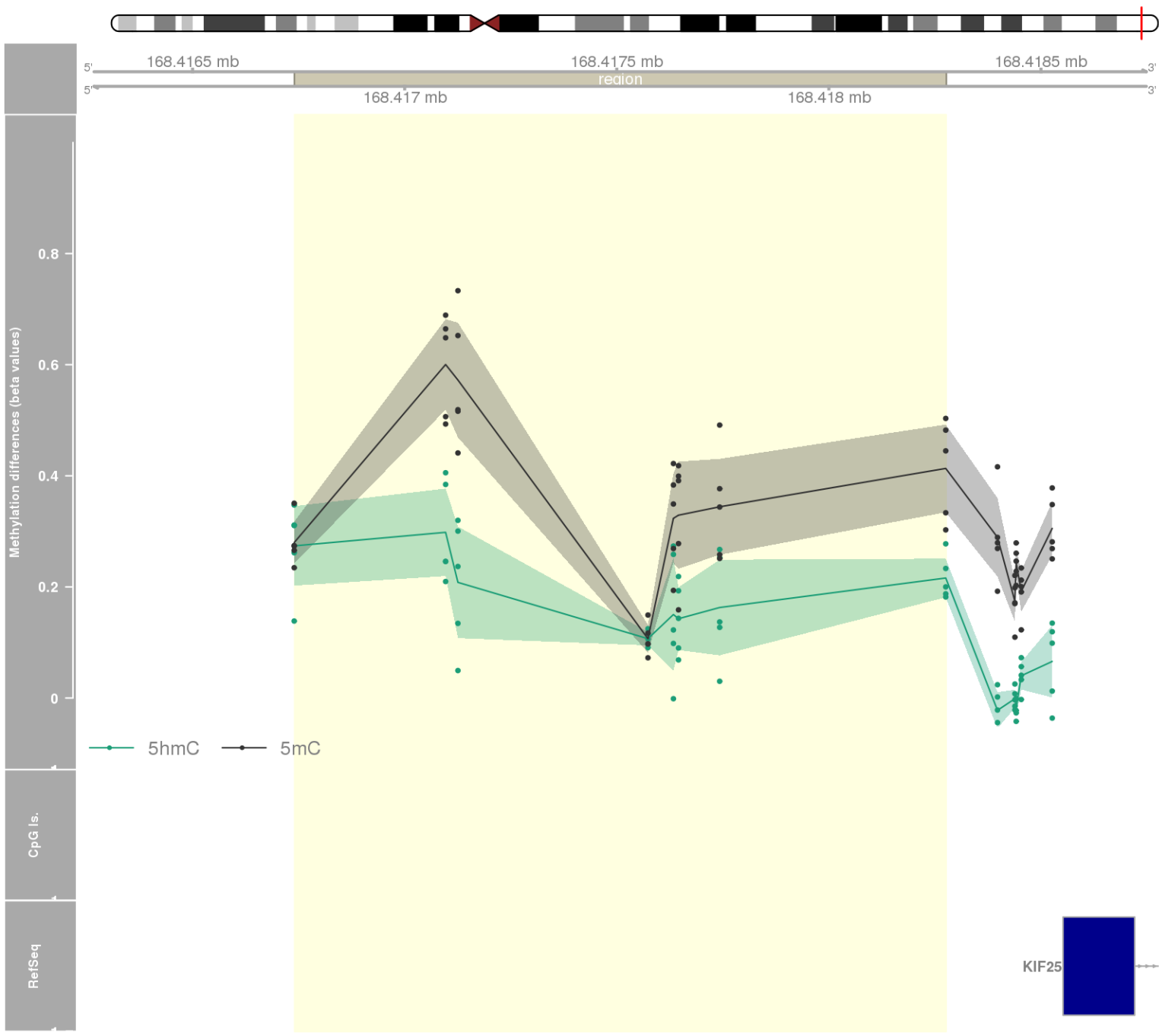
DMR 303 // chr1:207975182-207975905 // 723 pb. (7 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.021 // fwerArea: 1
- genes: C1orf132 -



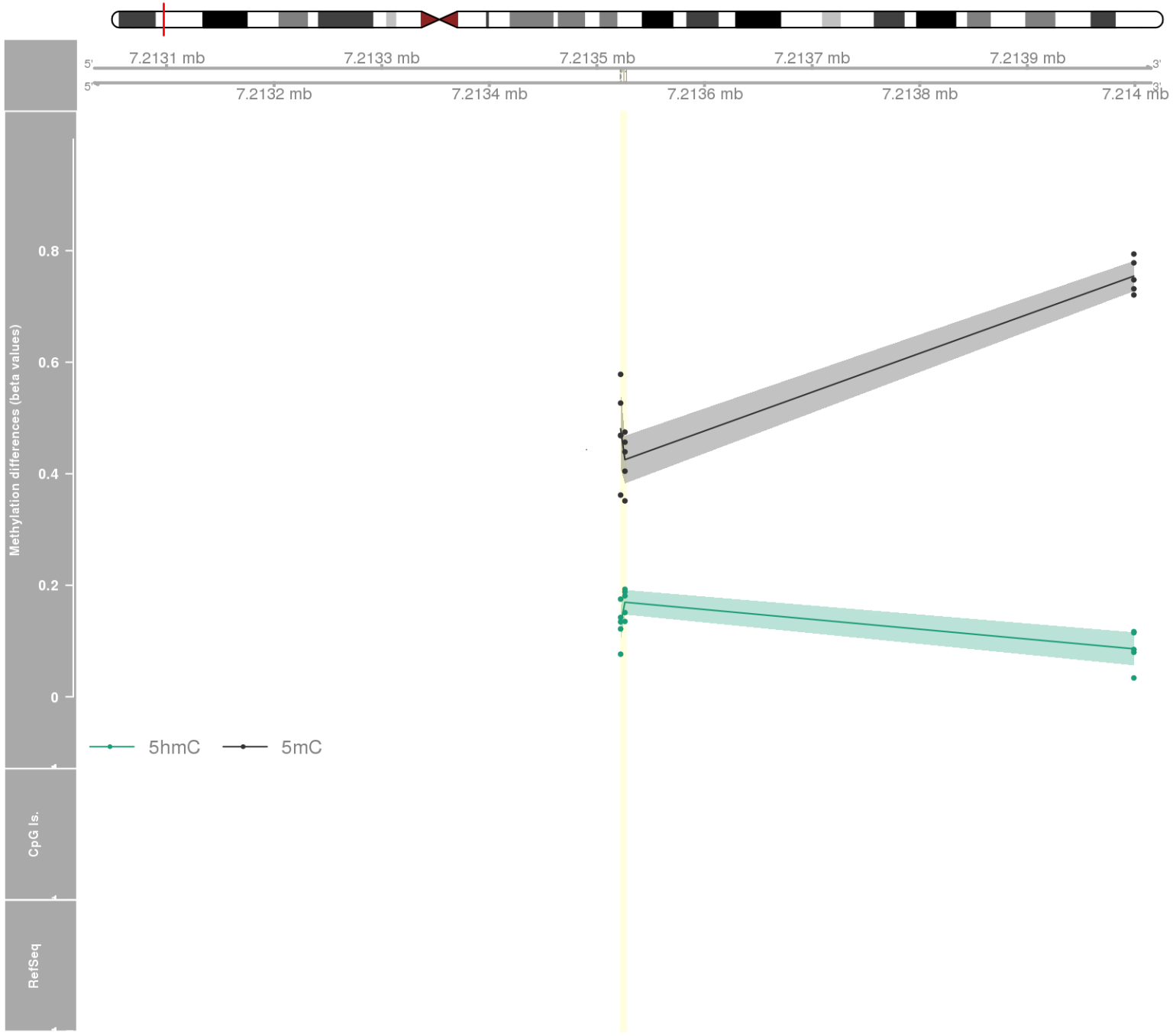
DMR 304 // chr6:30712865-30713442 // 577 pb. (9 probes) // pvalue: 0.01 // fwer: 0.36 // pvalueArea: 0.021 // fwerArea: 1



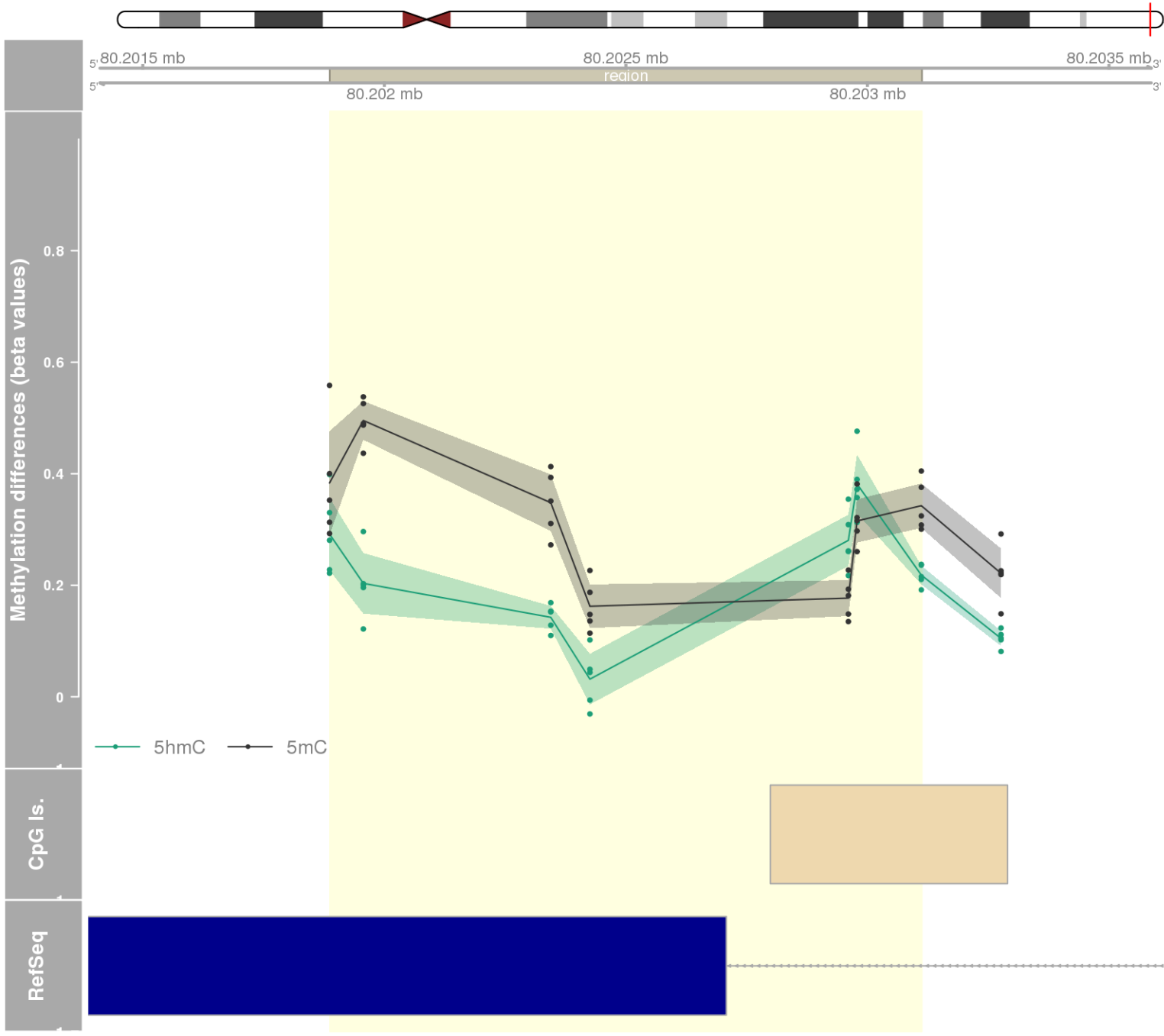
DMR 305 // chr6:168416740-168418276 // 1536 pb. (8 probes) // pvalue: 0.007 // fwer: 0.173 // pvalueArea: 0.021 // fwerArea: 1
- genes: KIF25 -



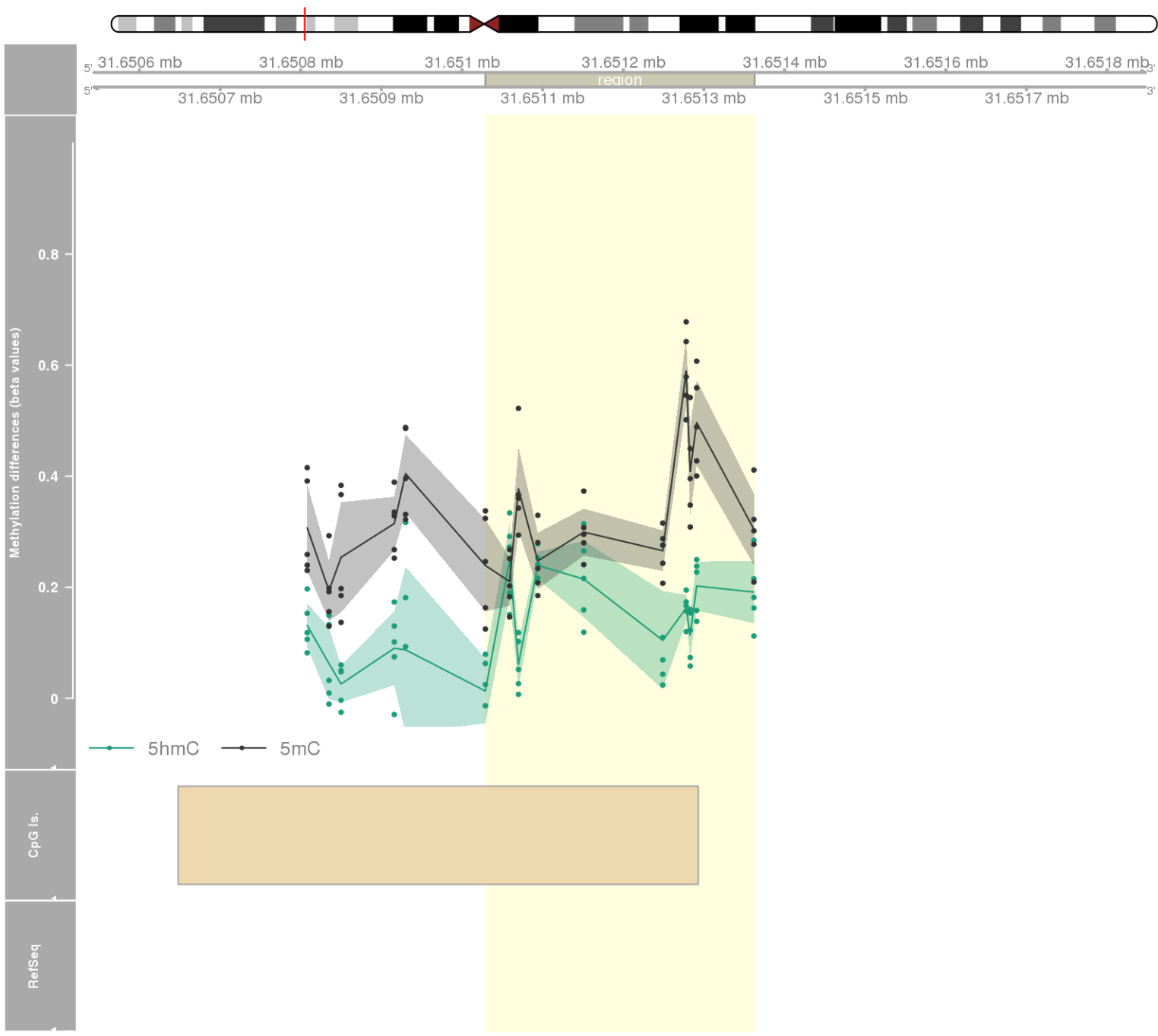
DMR 306 // chr8:7213522-7213526 // 4 pb. (2 probes) // pvalue: 0.001 // fwer: 0.659 // pvalueArea: 0.021 // fwerArea: 1
- genes: ZNF705G -



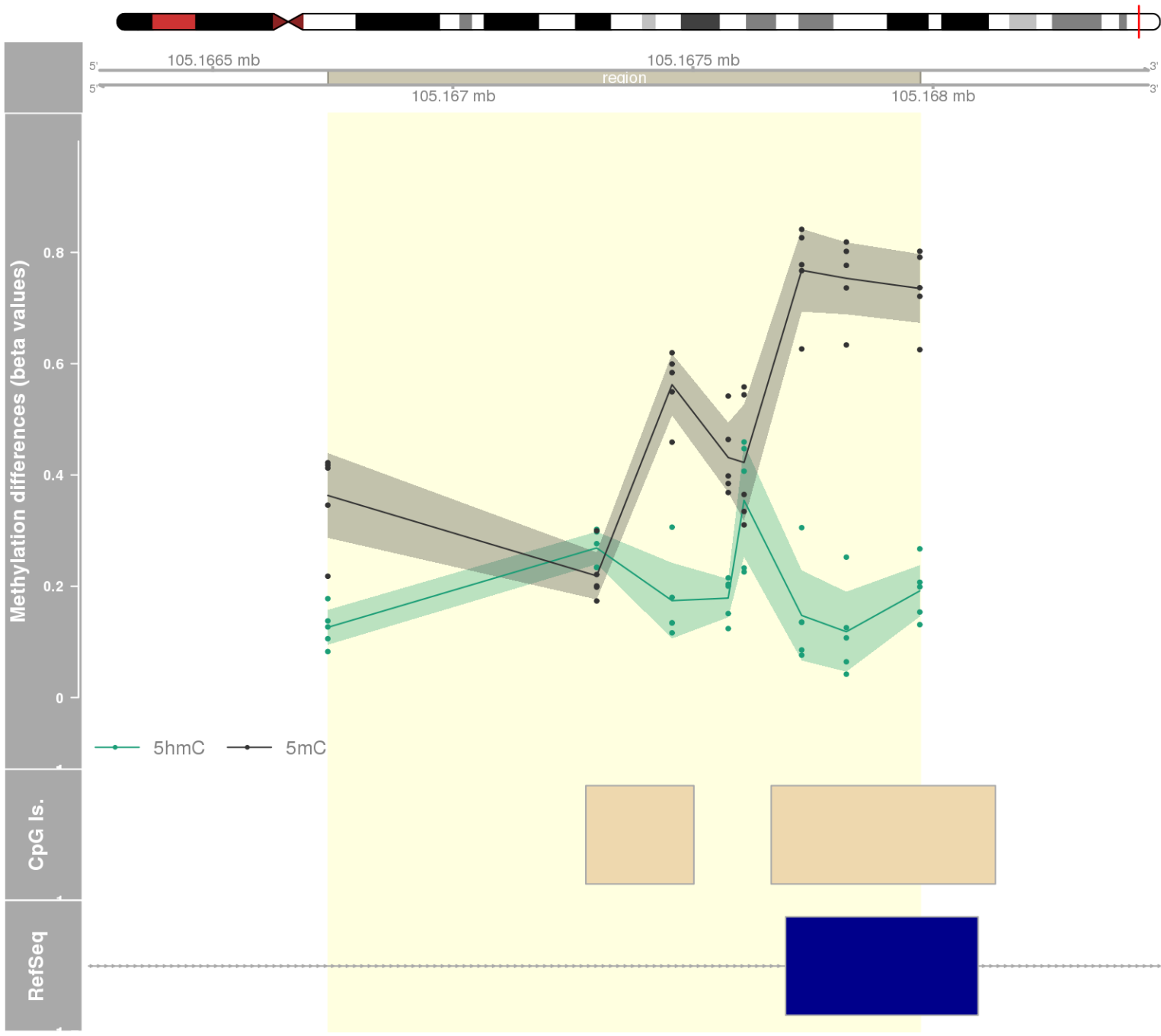
DMR 307 // chr17:80201887-80203112 // 1225 pb. (7 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.022 // fwerArea: 1
- genes: SLC16A3 / CSNK1D -



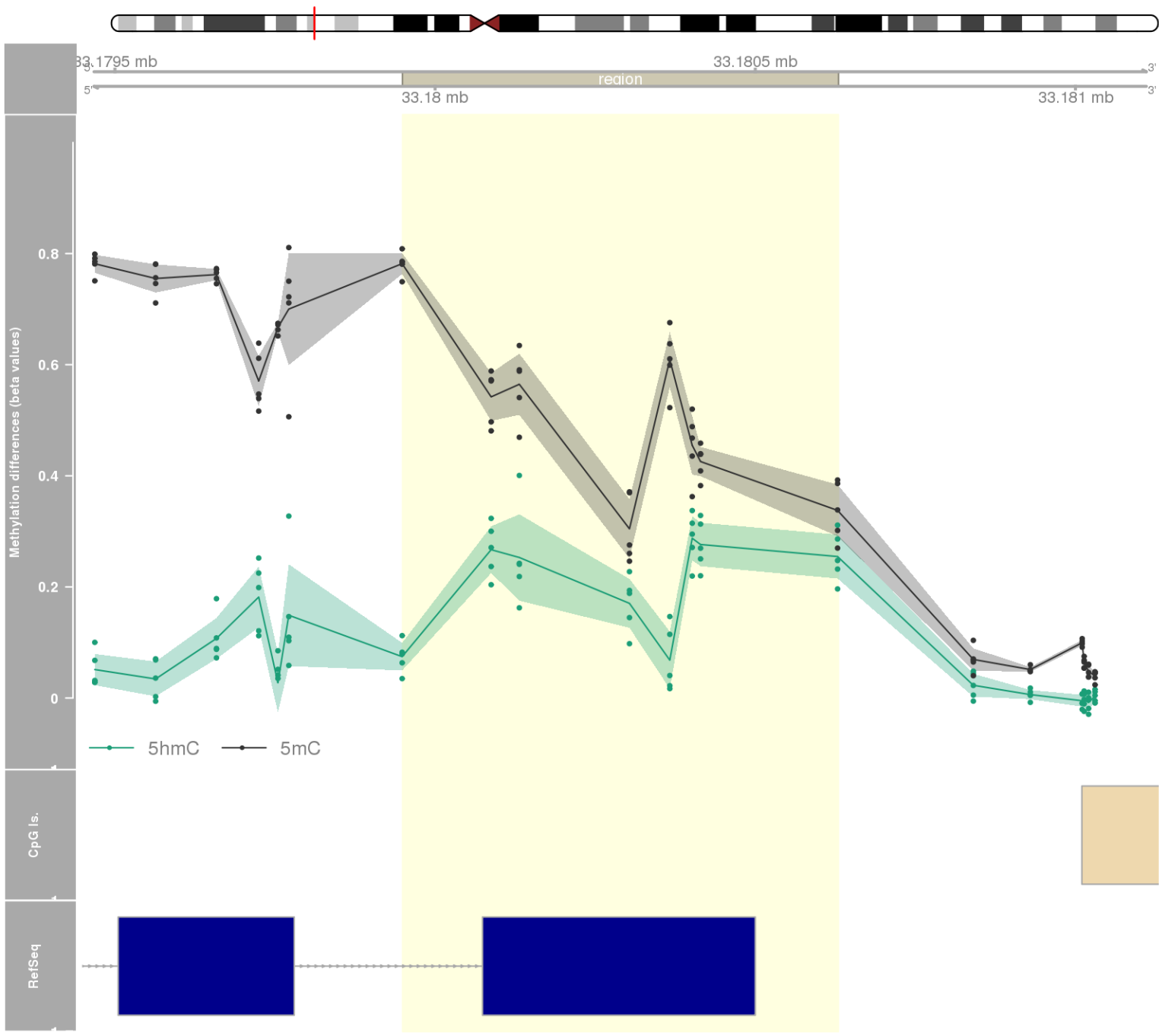
DMR 308 // chr6:31651029-31651362 // 333 pb. (10 probes) // pvalue: 0.011 // fwer: 0.36 // pvalueArea: 0.022 // fwerArea: 1
- genes: LY6G5C -



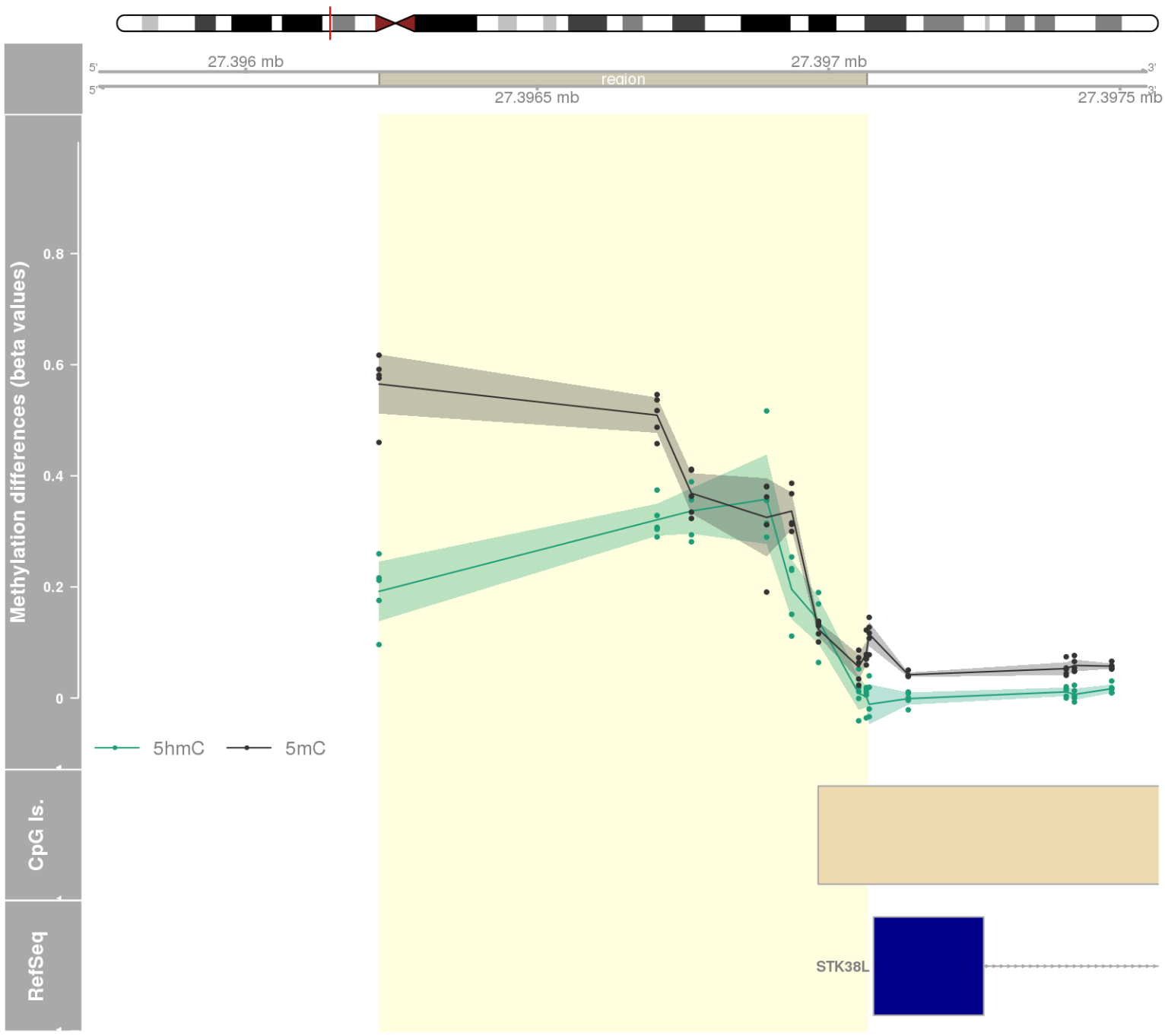
DMR 309 // chr14:105166740-105167973 // 1233 pb. (8 probes) // pvalue: 0.007 // fwer: 0.173 // pvalueArea: 0.022 // fwerArea: 1
- genes: INF2 -



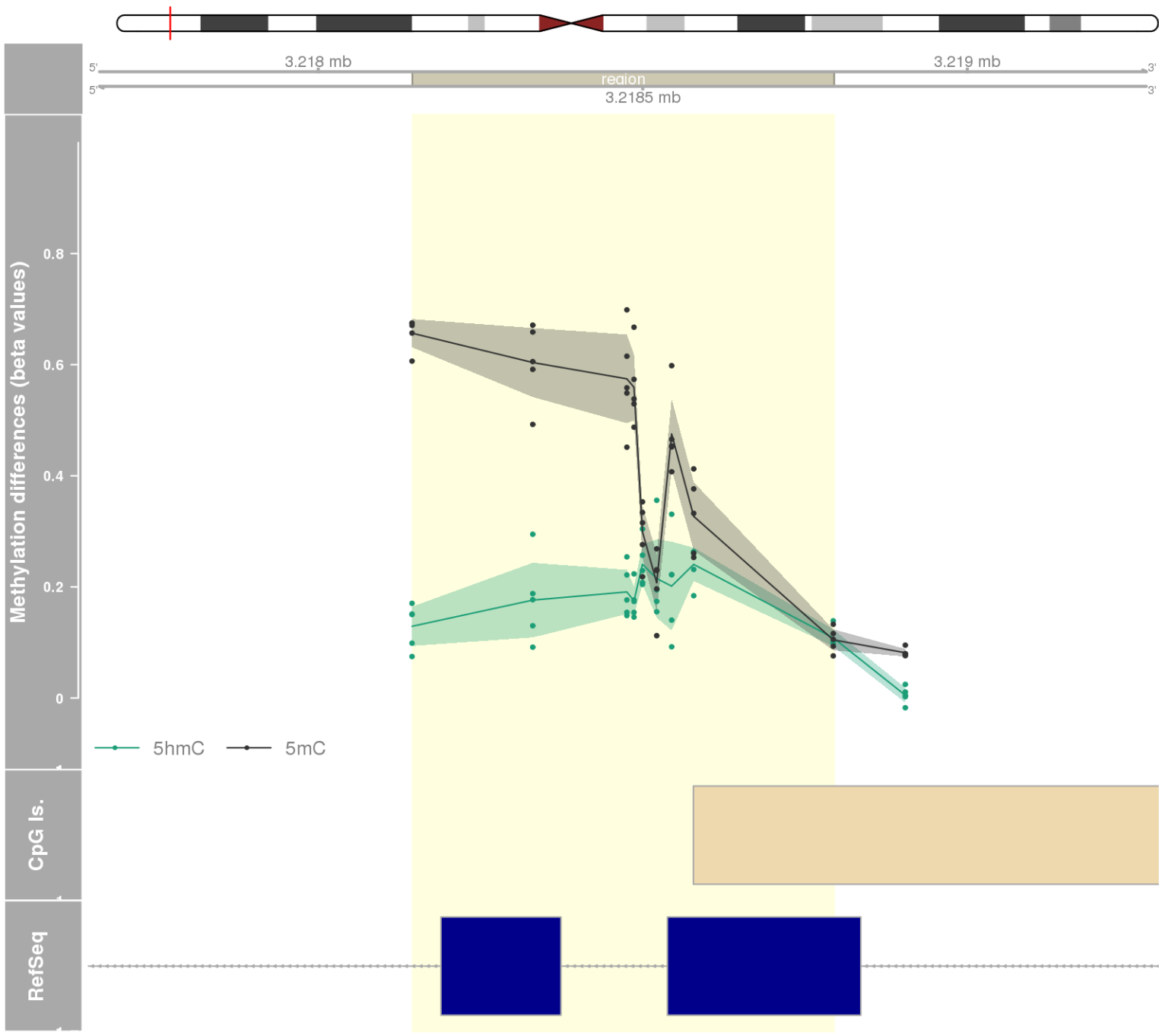
DMR 310 // chr6:33179949-33180629 // 680 pb. (8 probes) // pvalue: 0.007 // fwer: 0.173 // pvalueArea: 0.022 // fwerArea: 1
- genes: RING1 -



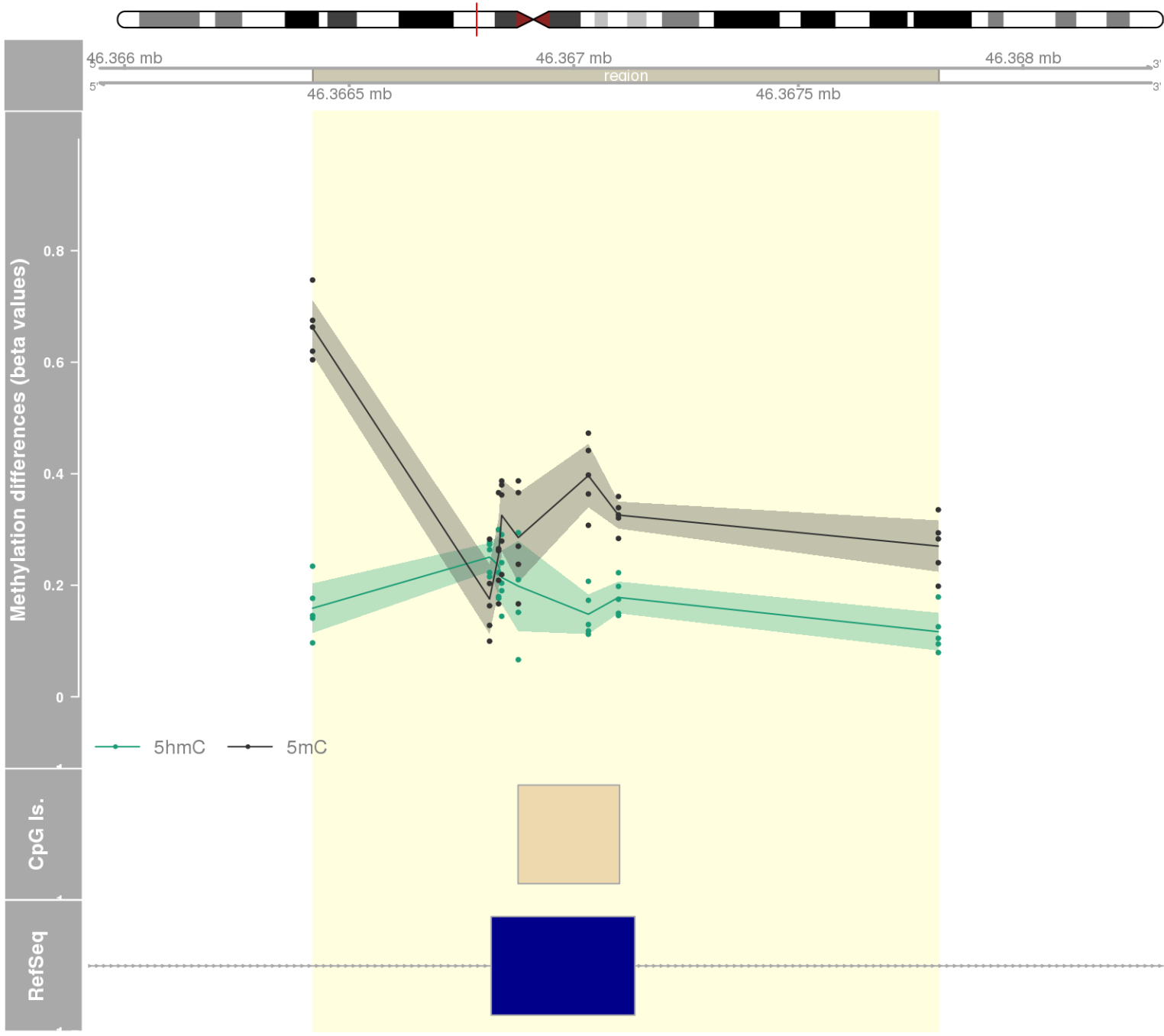
DMR 311 // chr12:27396229-27397065 // 836 pb. (8 probes) // pvalue: 0.007 // fwer: 0.173 // pvalueArea: 0.022 // fwerArea: 1
- genes: STK38L -



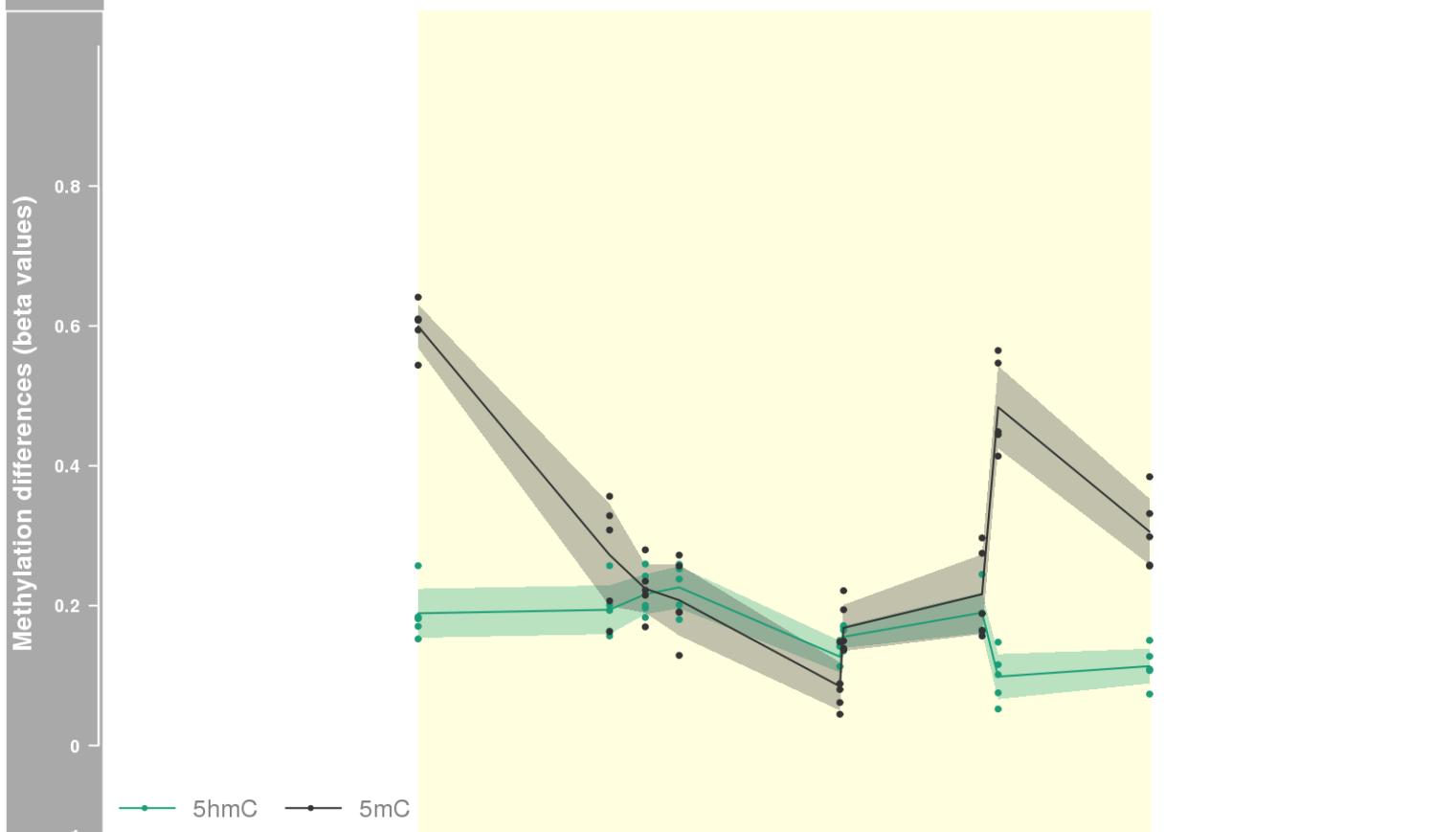
DMR 312 // chr20:3218145-3218794 // 649 pb. (9 probes) // pvalue: 0.01 // fwer: 0.36 // pvalueArea: 0.022 // fwerArea: 1
- genes: SLC4A11 -



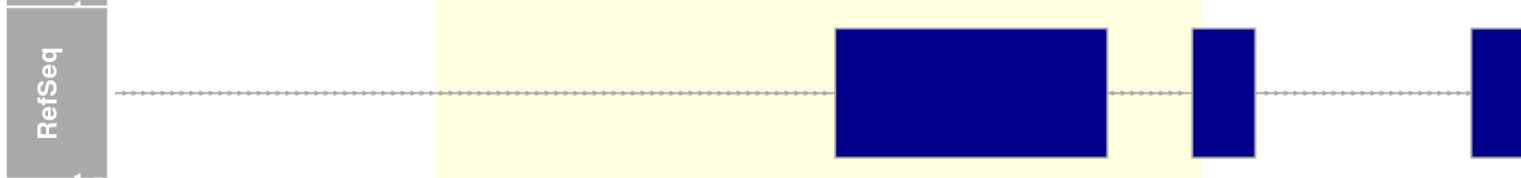
DMR 313 // chr11:46366419-46367812 // 1393 pb. (8 probes) // pvalue: 0.007 // fwer: 0.173 // pvalueArea: 0.022 // fwerArea: 1
- genes: DGKZ -



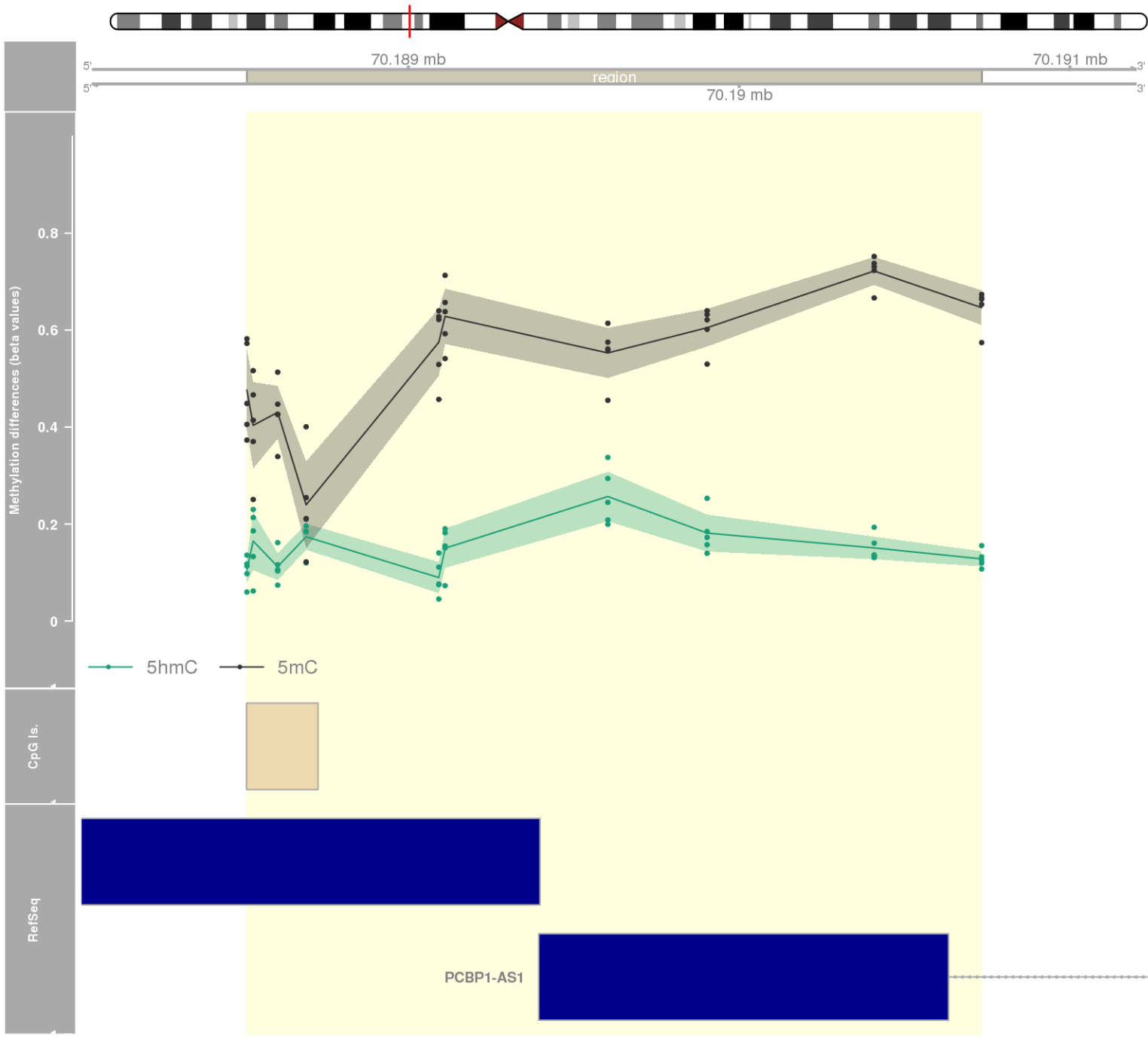
DMR 314 // chr12:6492579-6493767 // 1188 pb. (9 probes) // pvalue: 0.01 // fwer: 0.36 // pvalueArea: 0.022 // fwerArea: 1
- genes: LTBR -



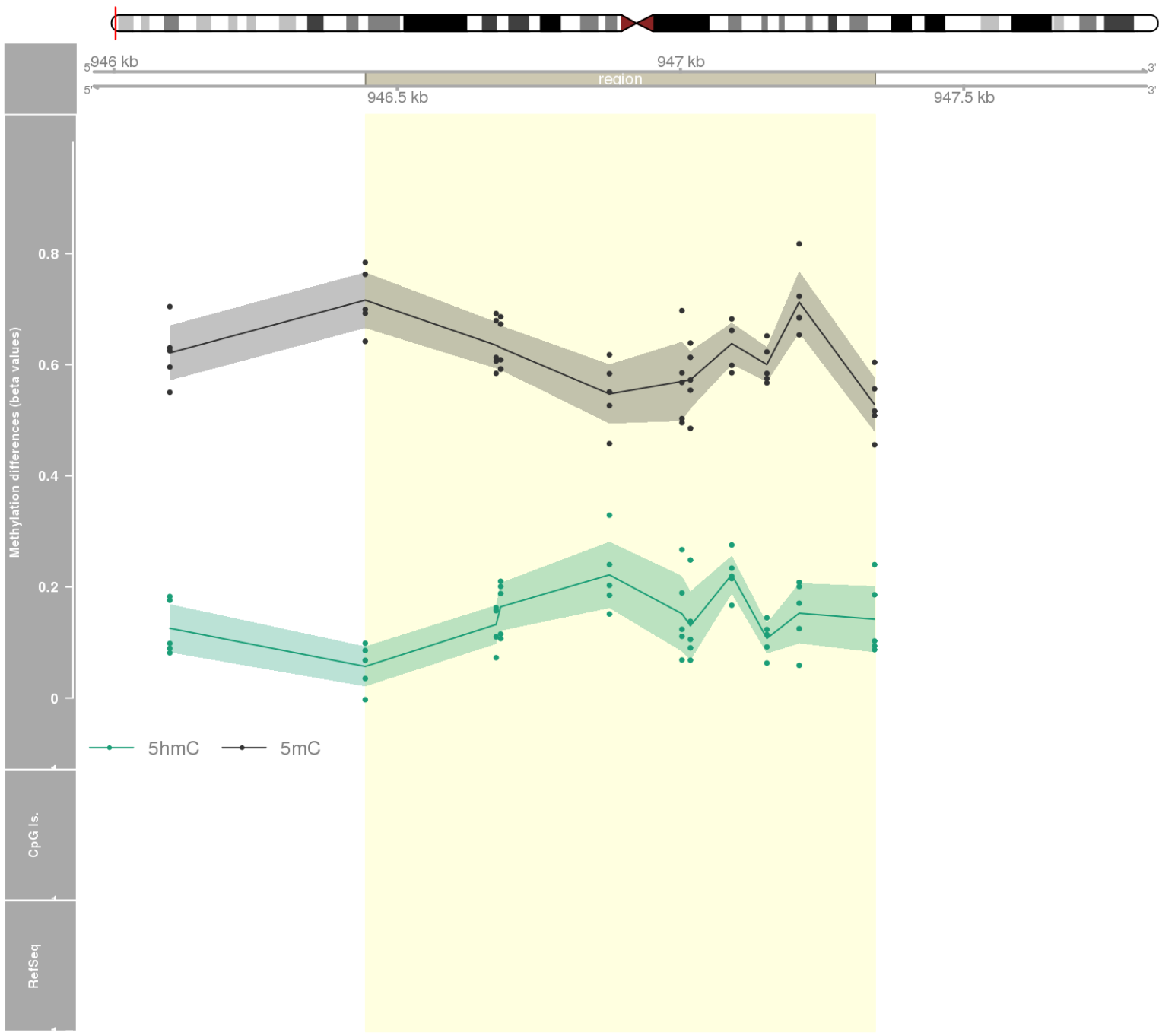
—●— 5hmC —●— 5mC



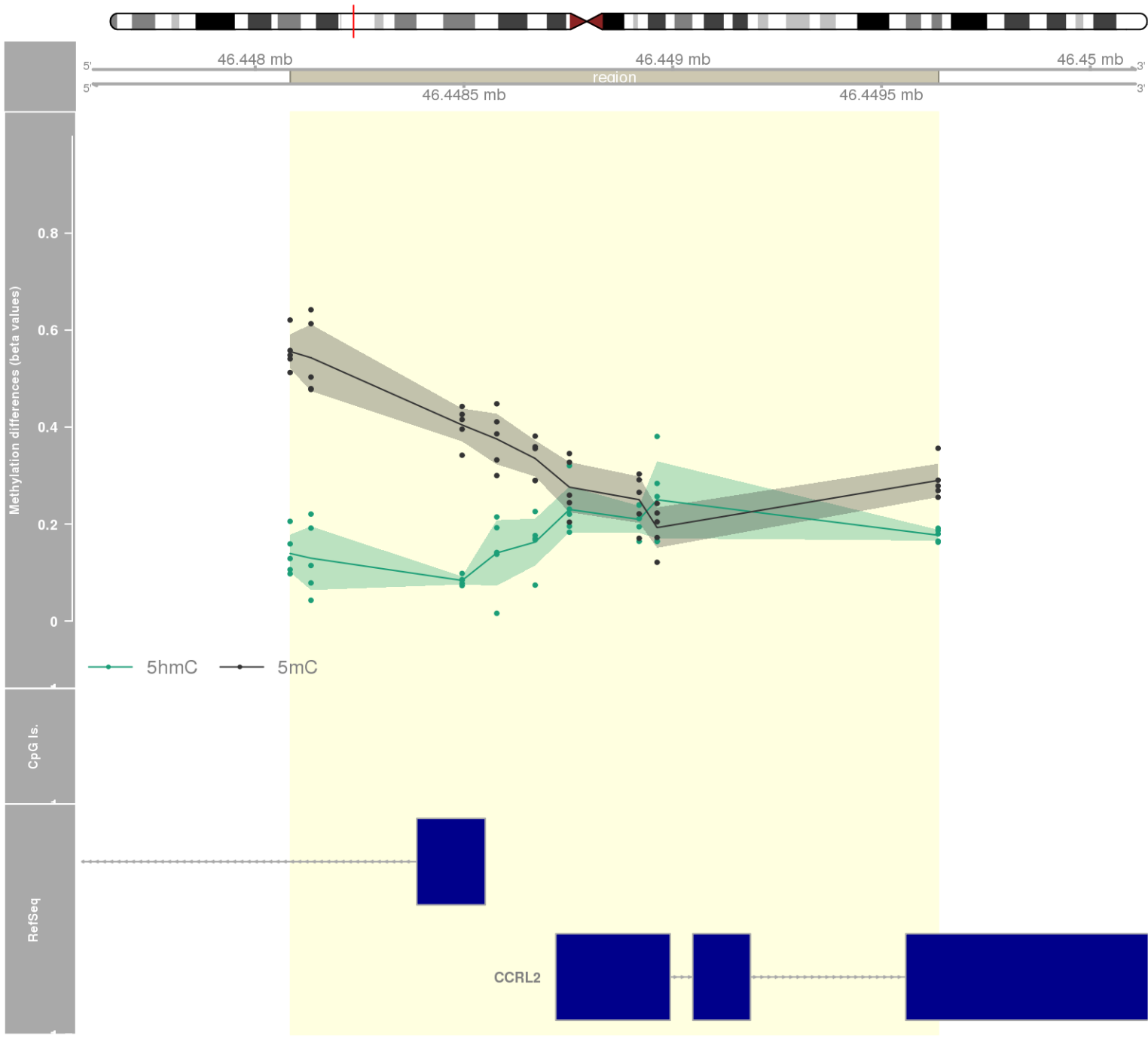
DMR 315 // chr2:70188512-70190733 // 2221 pb. (10 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.022 // fwerArea: 1
- genes: PCBP1-AS1 / ASPRV1 -



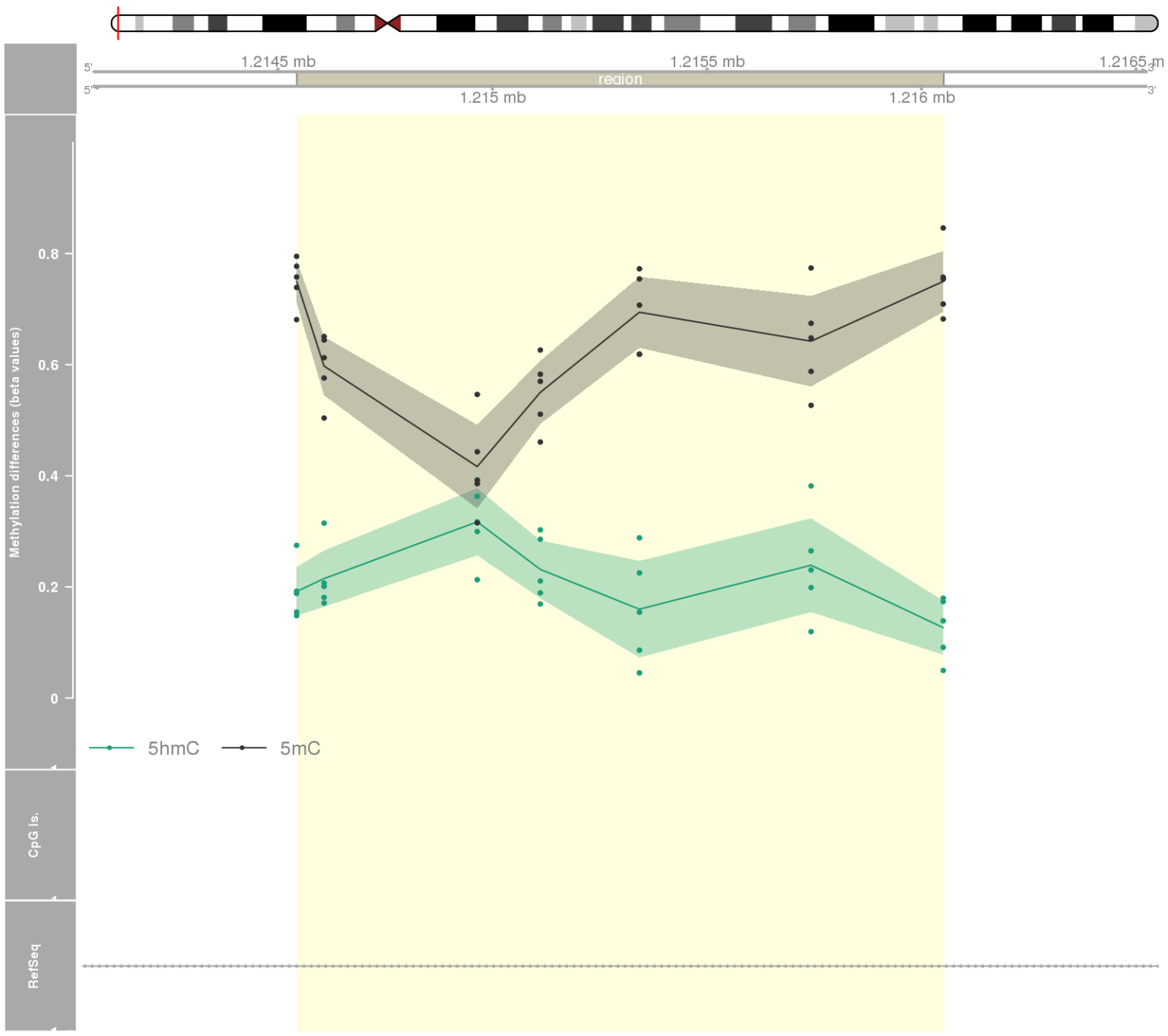
DMR 316 // chr1:946444-947343 // 899 pb. (10 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.022 // fwerArea: 1



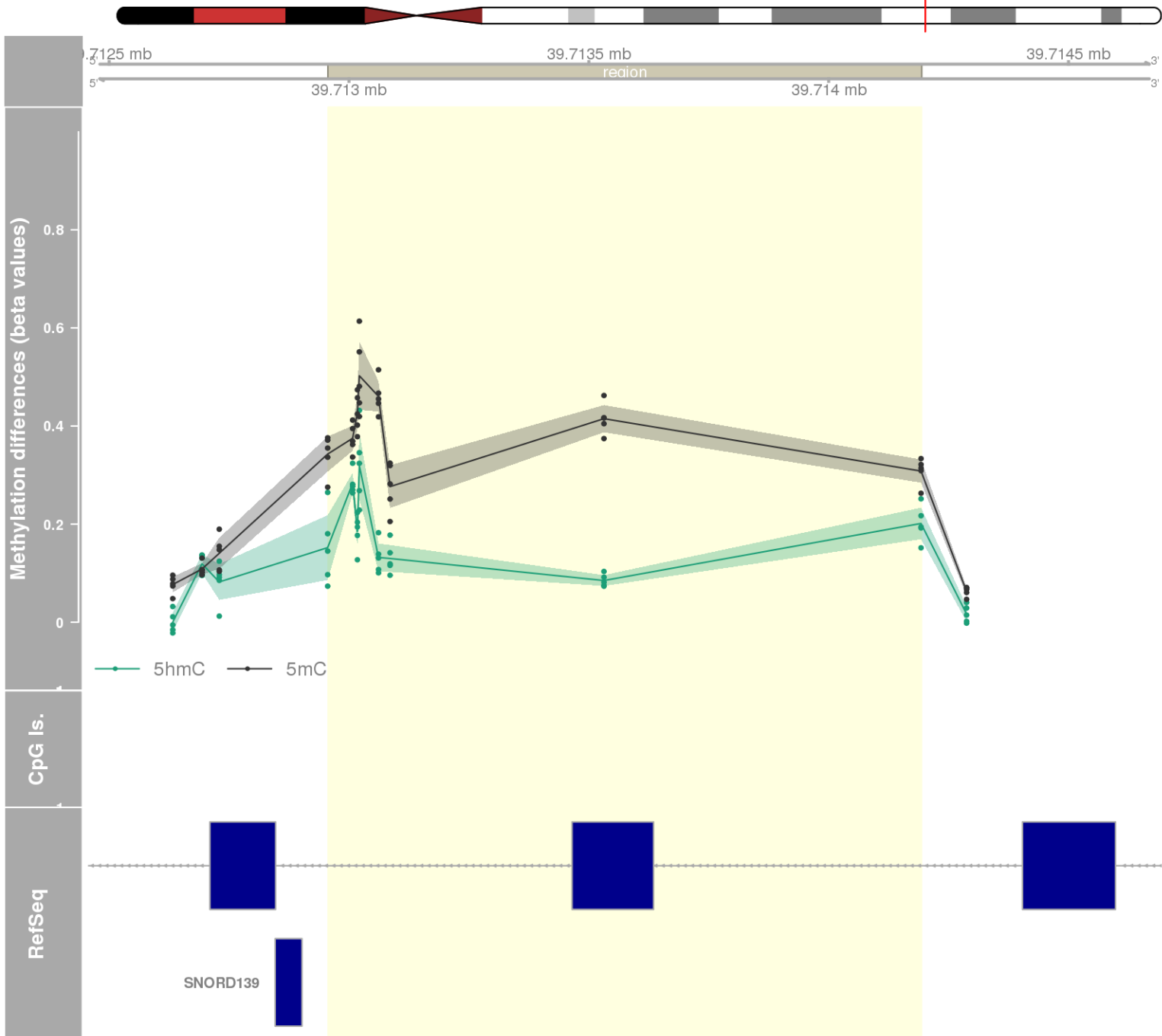
DMR 317 // chr3:46448084-46449636 // 1552 pb. (9 probes) // pvalue: 0.011 // fwer: 0.36 // pvalueArea: 0.022 // fwerArea: 1
- genes: CCRL2 -



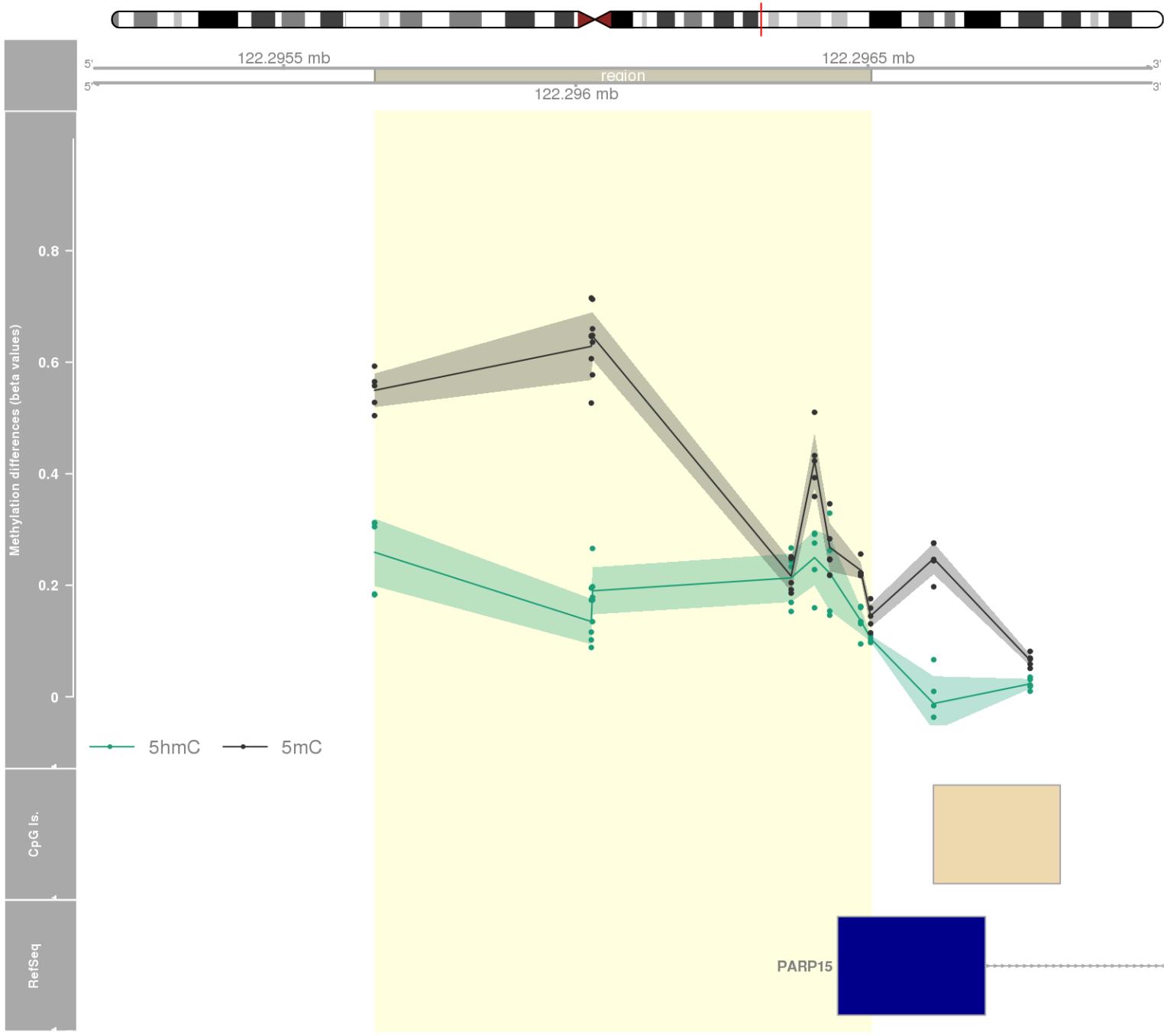
DMR 318 // chr4:1214544-1216051 // 1507 pb. (7 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.022 // fwerArea: 1
- genes: CTBP1 -



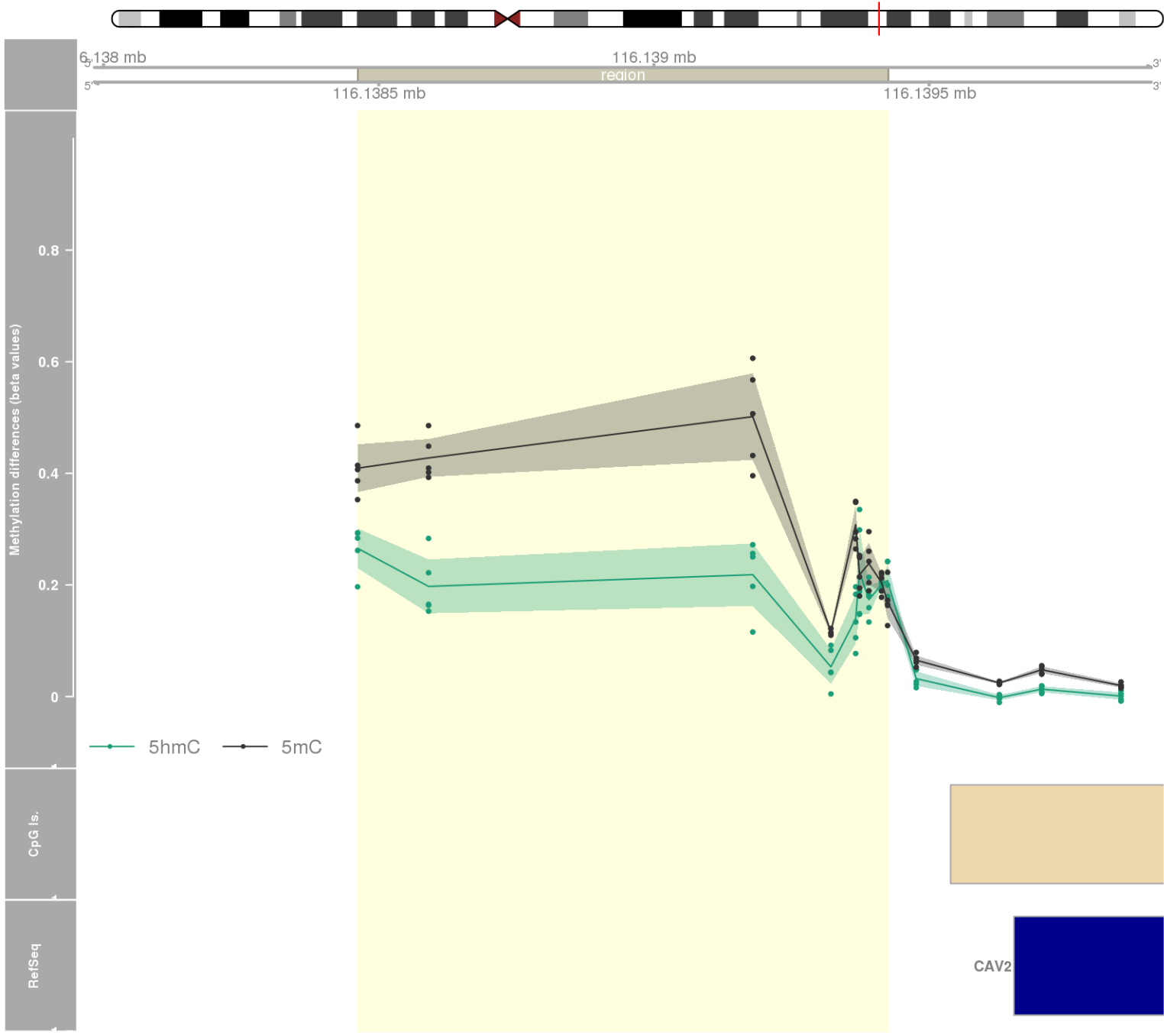
DMR 319 // chr22:39712956-39714193 // 1237 pb. (8 probes) // pvalue: 0.008 // fwer: 0.173 // pvalueArea: 0.022 // fwerArea: 1
- genes: RPL3 -



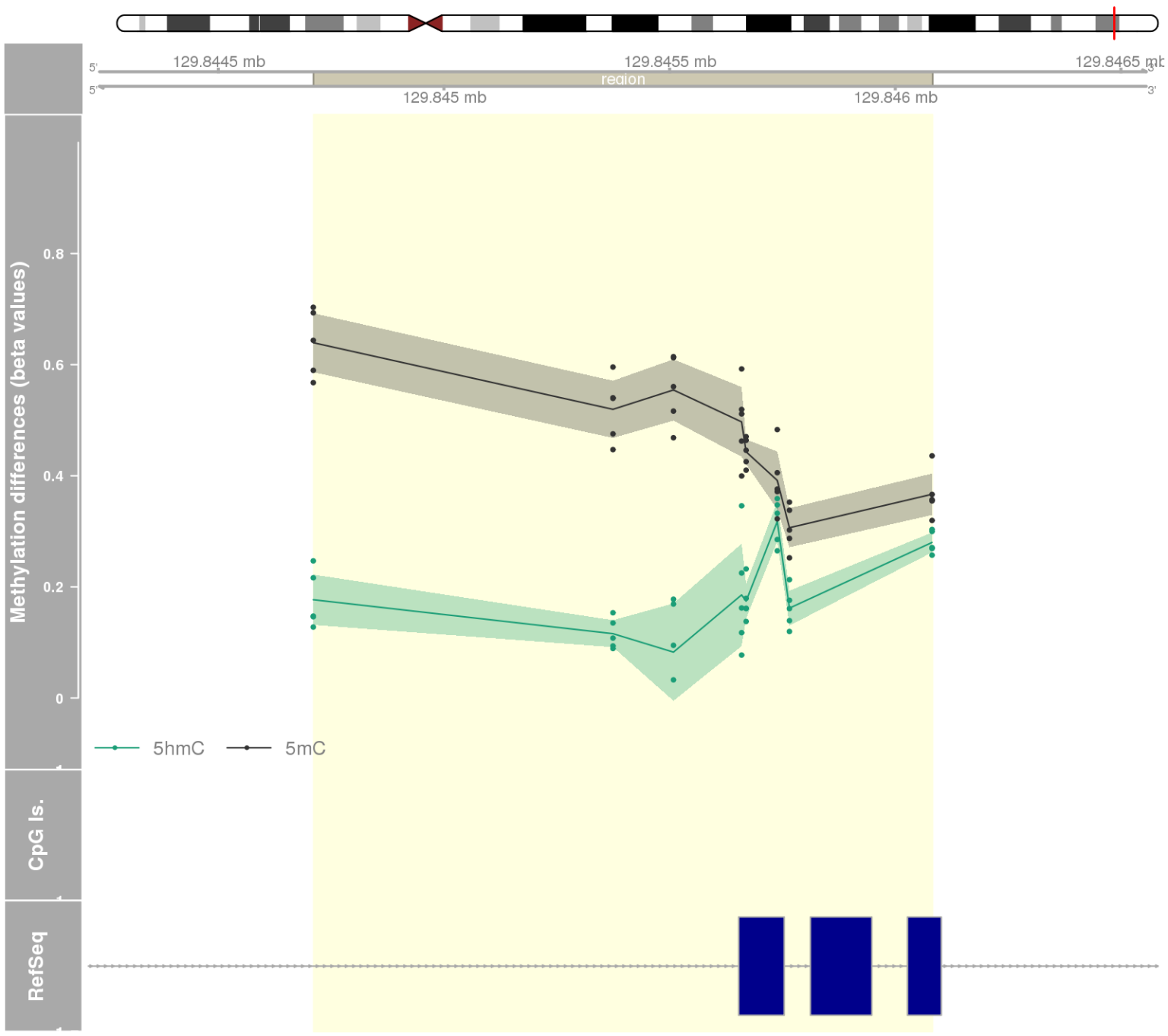
DMR 320 // chr3:122295656-122296505 // 849 pb. (8 probes) // pvalue: 0.008 // fwer: 0.173 // pvalueArea: 0.022 // fwerArea: 1
- genes: PARP15 -



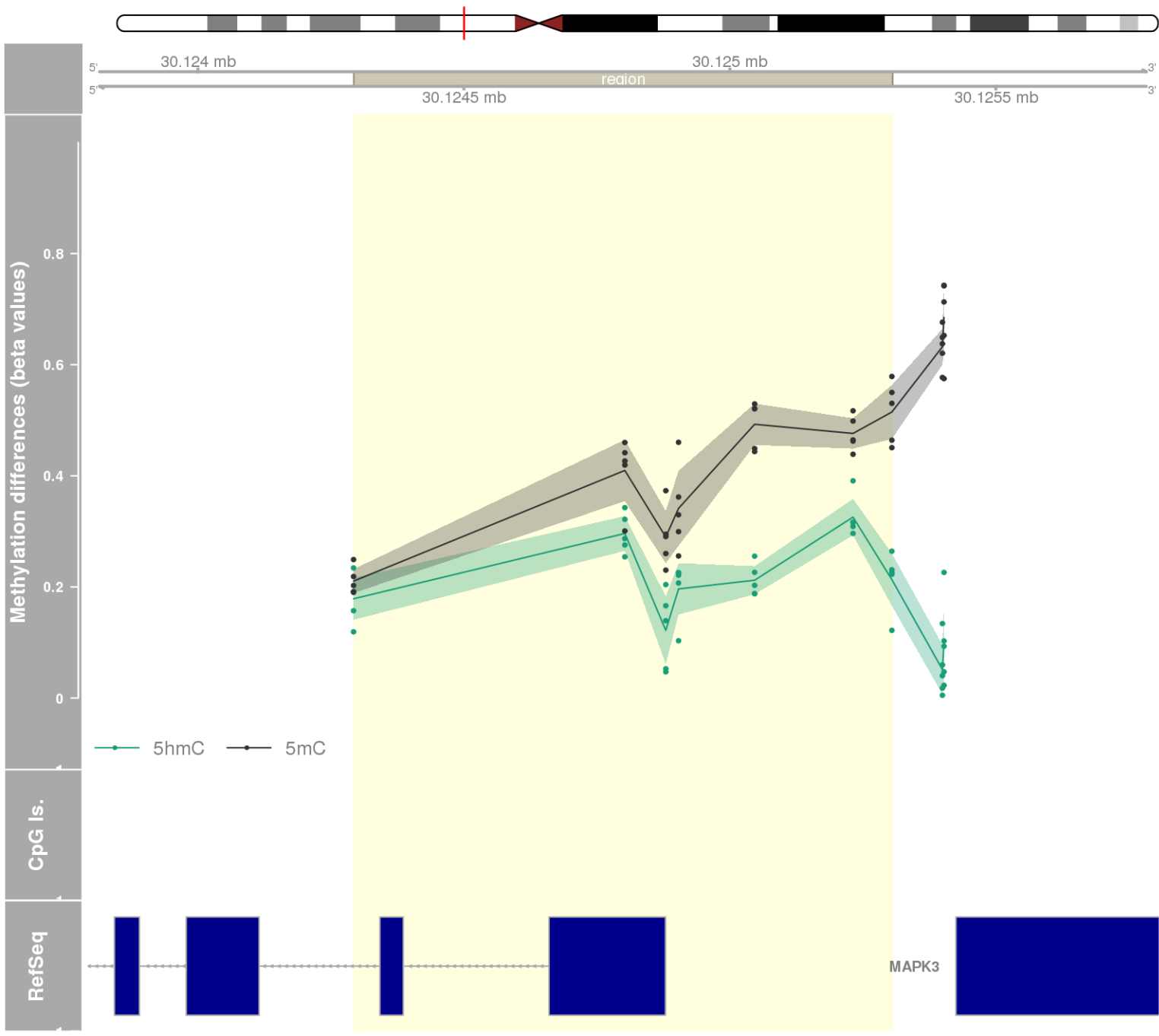
DMR 321 // chr7:116138462-116139425 // 963 pb. (9 probes) // pvalue: 0.011 // fwer: 0.36 // pvalueArea: 0.022 // fwerArea: 1
- genes: CAV2 -



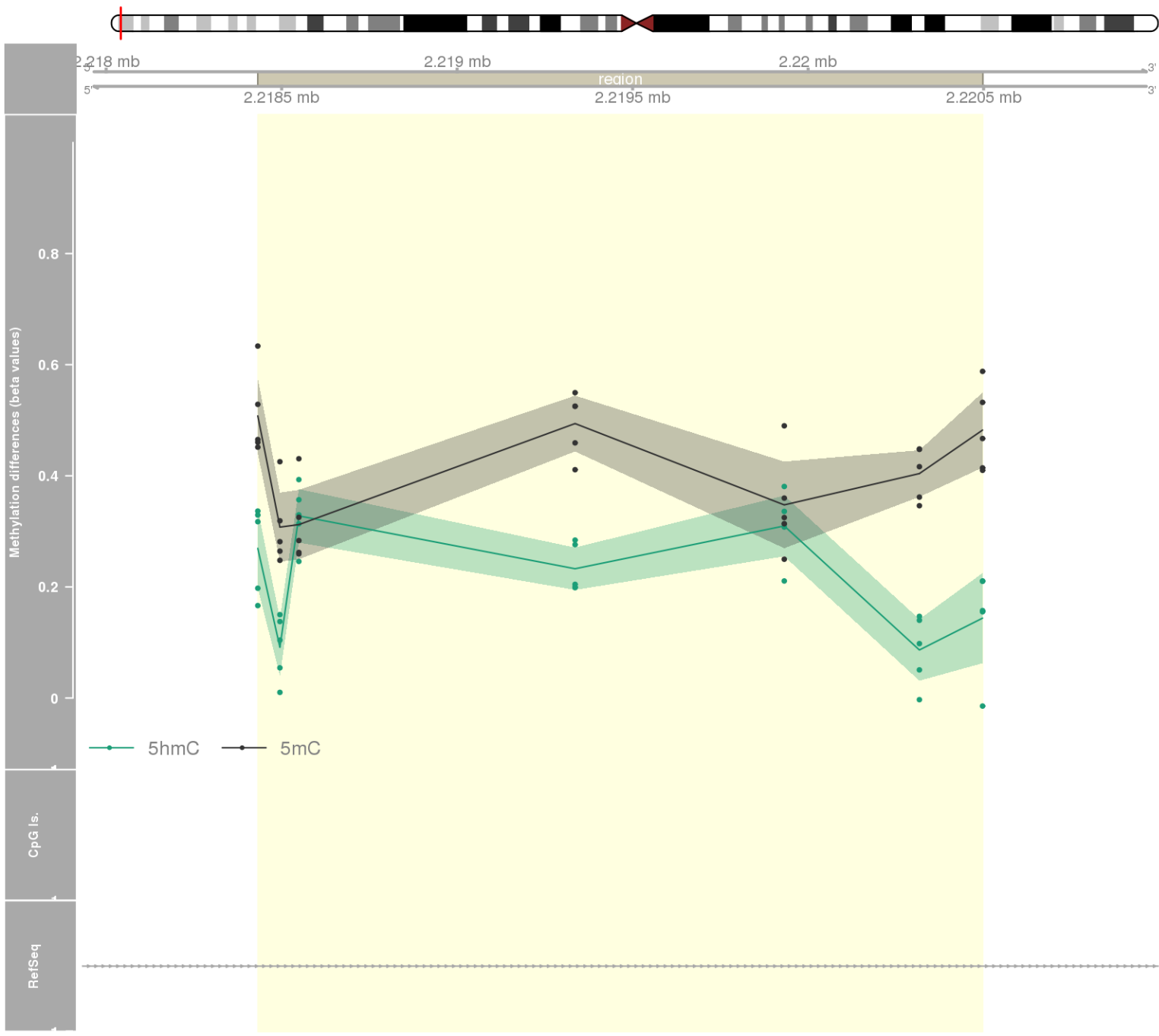
DMR 322 // chr10:129844711-129846082 // 1371 pb. (8 probes) // pvalue: 0.008 // fwer: 0.173 // pvalueArea: 0.023 // fwerArea: 1
- genes: PTPRE -



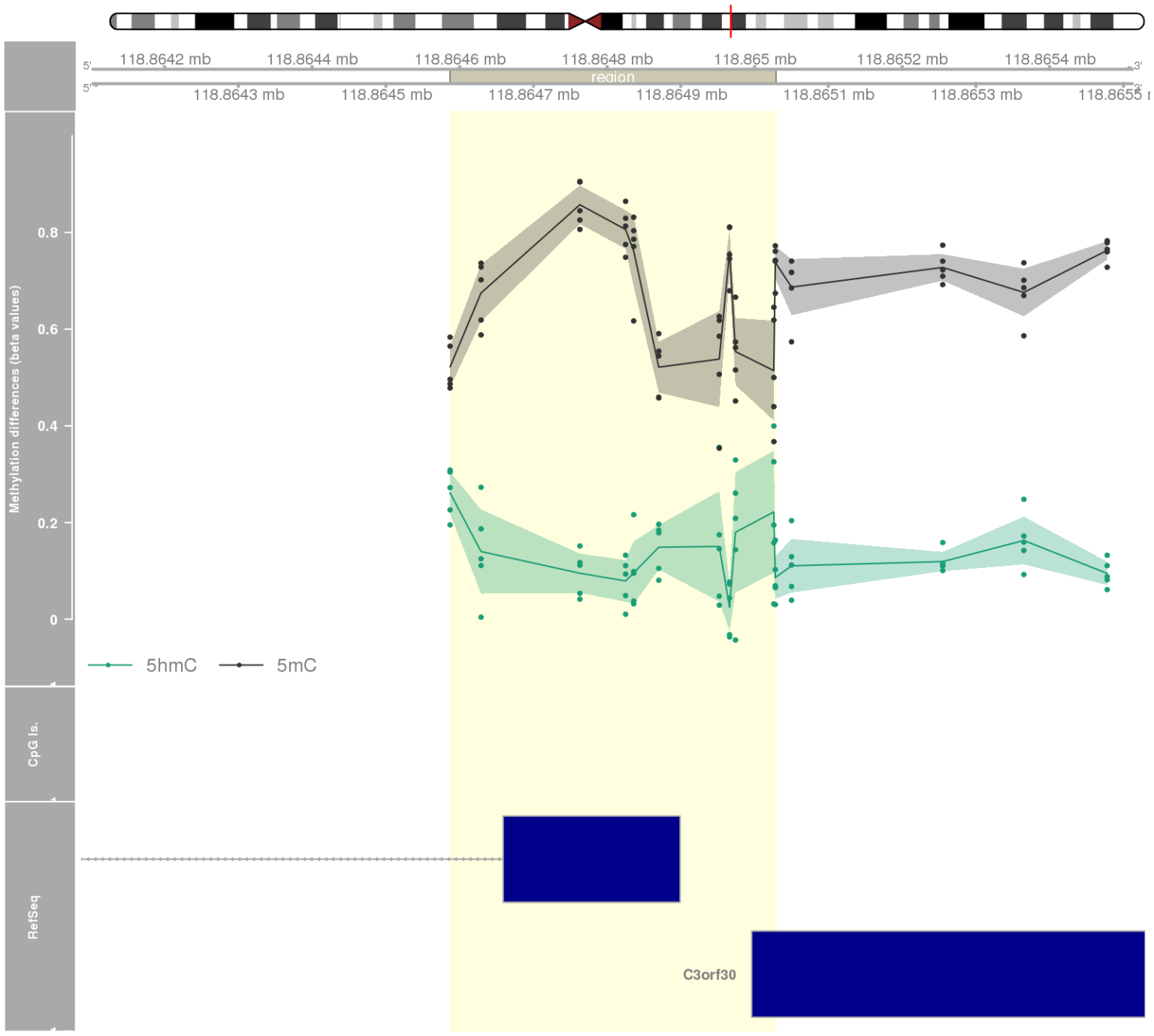
DMR 323 // chr16:30124293-30125305 // 1012 pb. (7 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.023 // fwerArea: 1
- genes: GDPD3 -



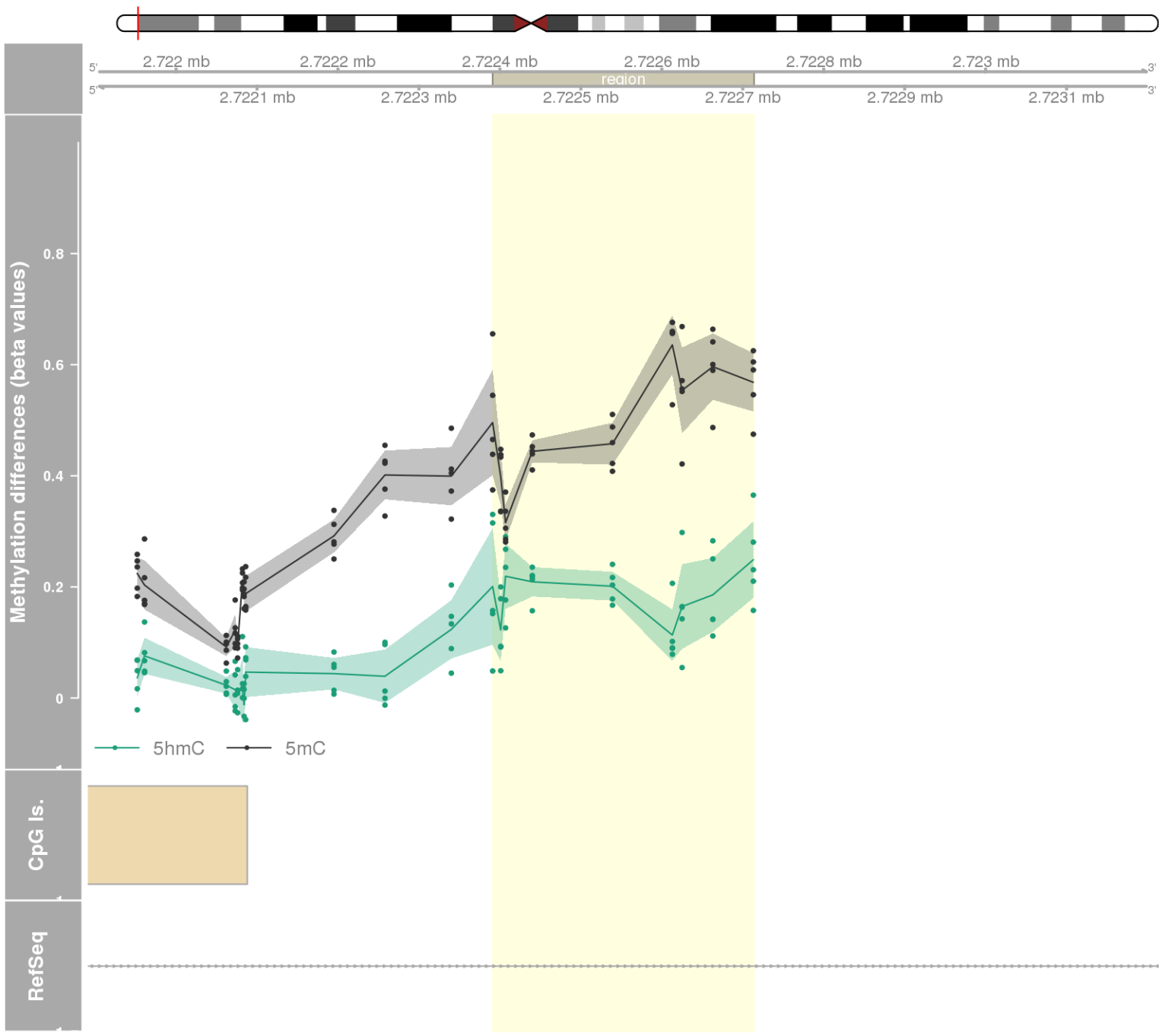
DMR 324 // chr1:2218432-2220497 // 2065 pb. (7 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.023 // fwerArea: 1
- genes: SKI -



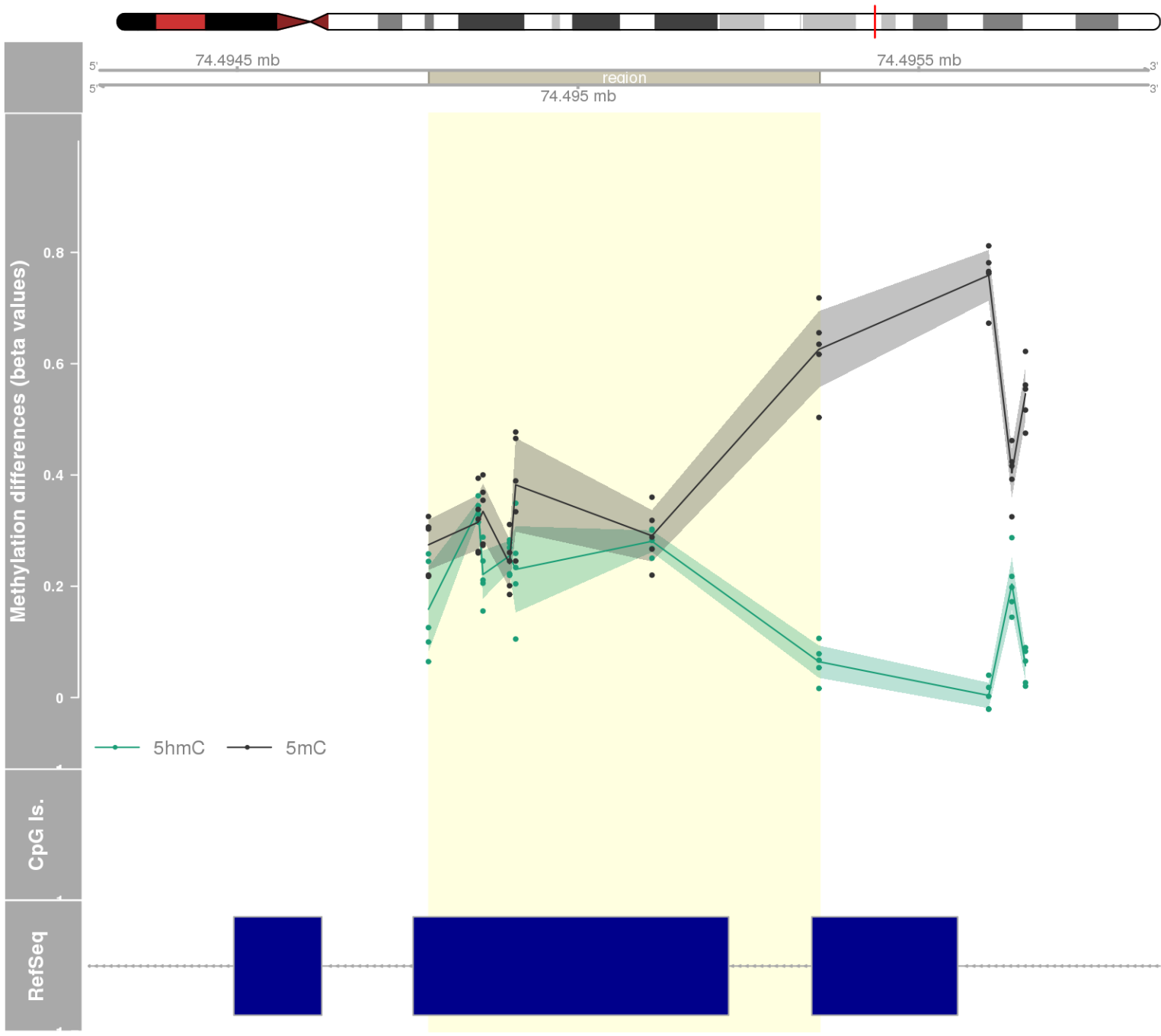
DMR 325 // chr3:118864587-118865028 // 441 pb. (11 probes) // pvalue: 0.011 // fwer: 0.36 // pvalueArea: 0.023 // fwerArea: 1
- genes: IGSF11 / C3orf30 -



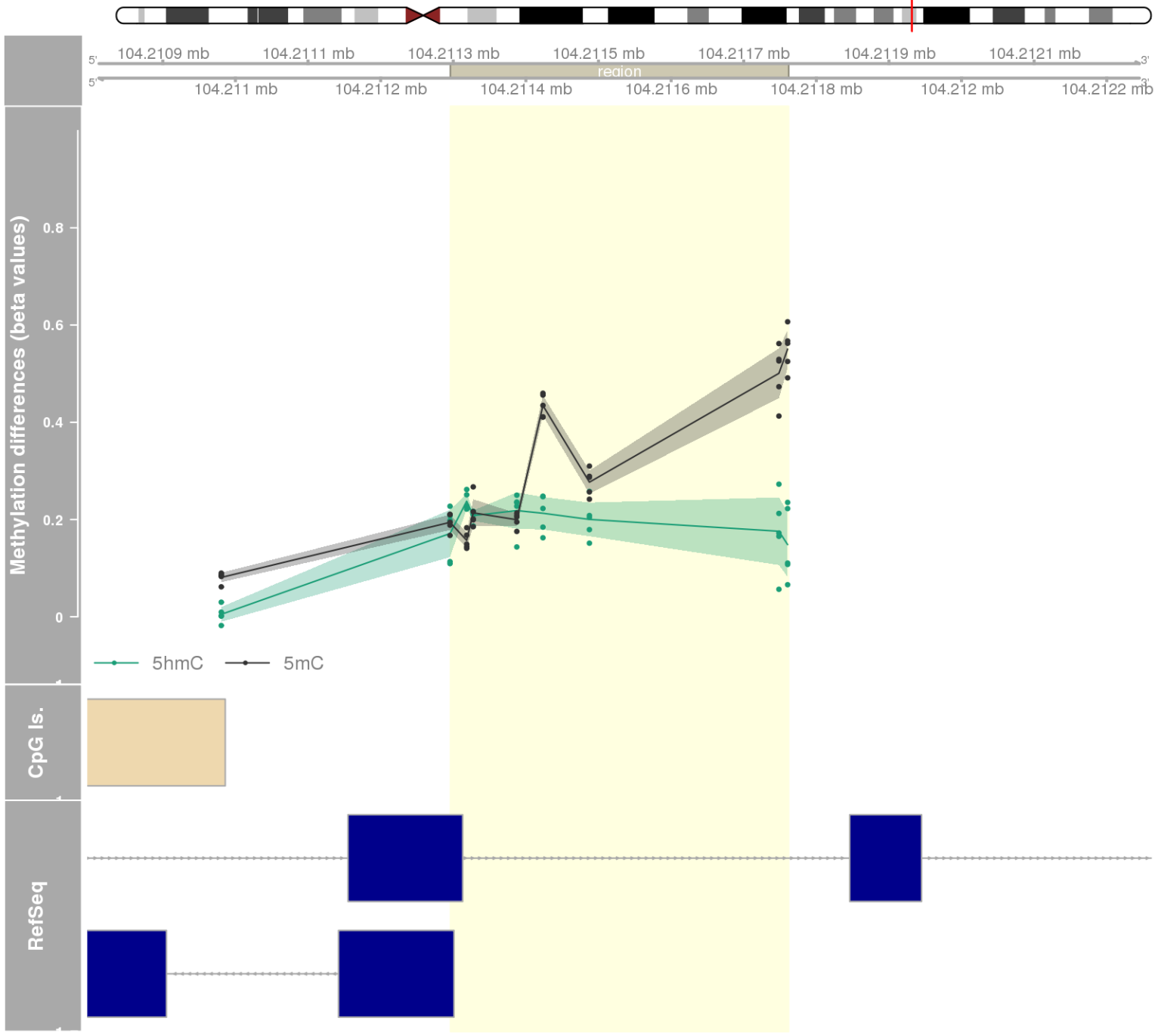
DMR 326 // chr11:2722391-2722713 // 322 pb. (9 probes) // pvalue: 0.011 // fwer: 0.36 // pvalueArea: 0.023 // fwerArea: 1
- genes: KCNQ1 -



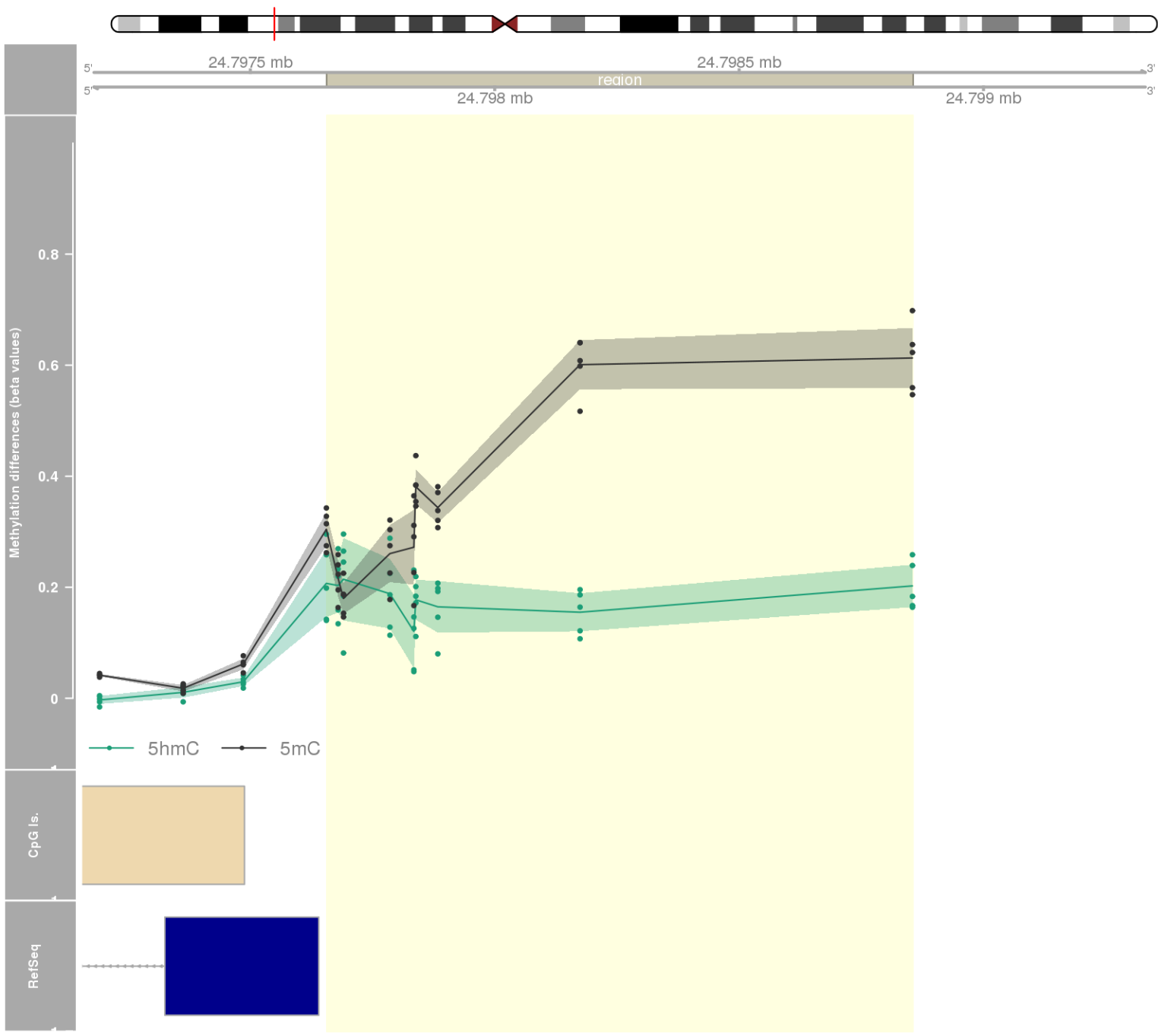
DMR 327 // chr15:74494781-74495354 // 573 pb. (7 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.023 // fwerArea: 1
- genes: STRA6 -



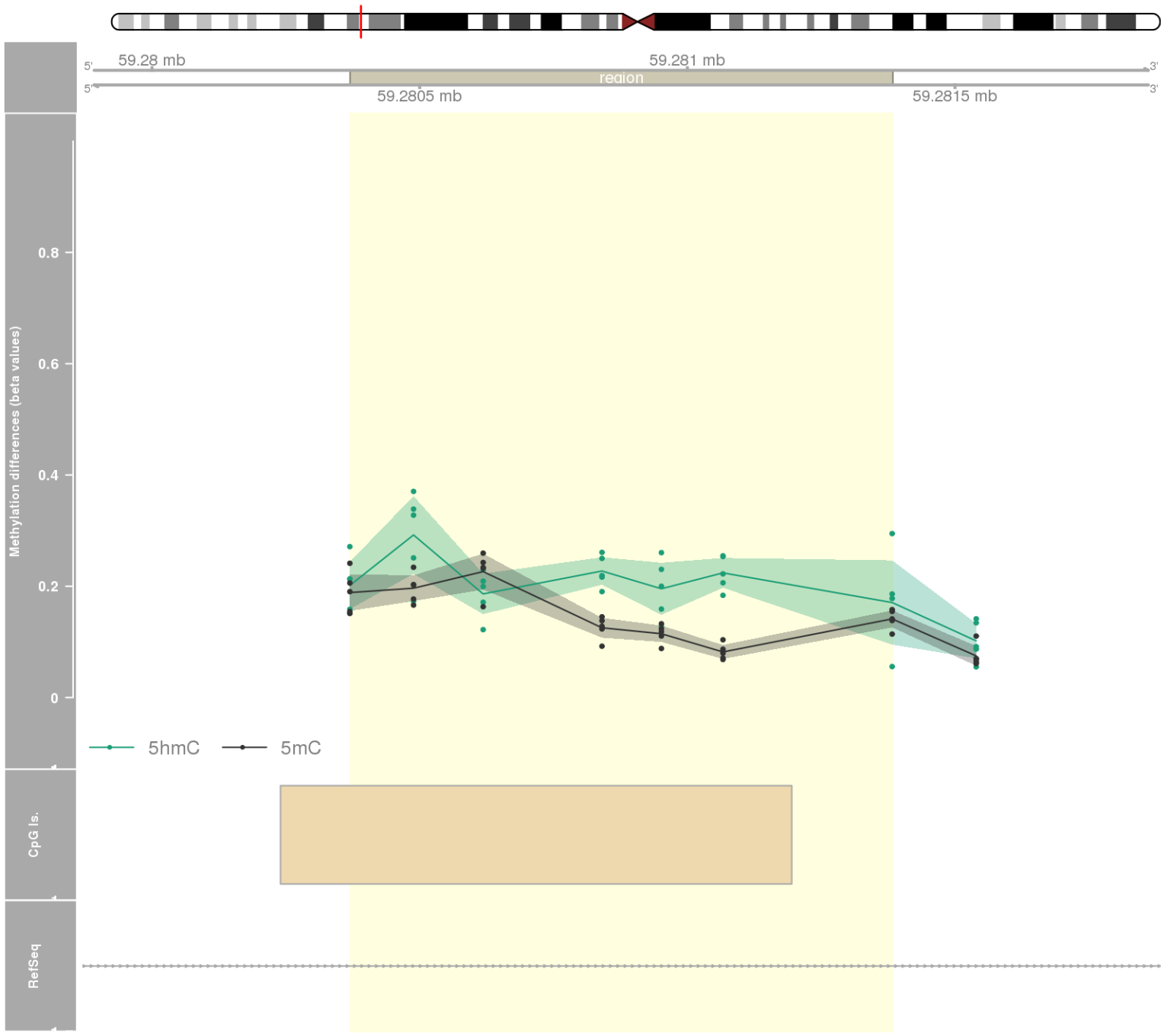
DMR 328 // chr10:104211296-104211761 // 465 pb. (8 probes) // pvalue: 0.008 // fwer: 0.36 // pvalueArea: 0.023 // fwerArea: 1
- genes: C10orf95 -



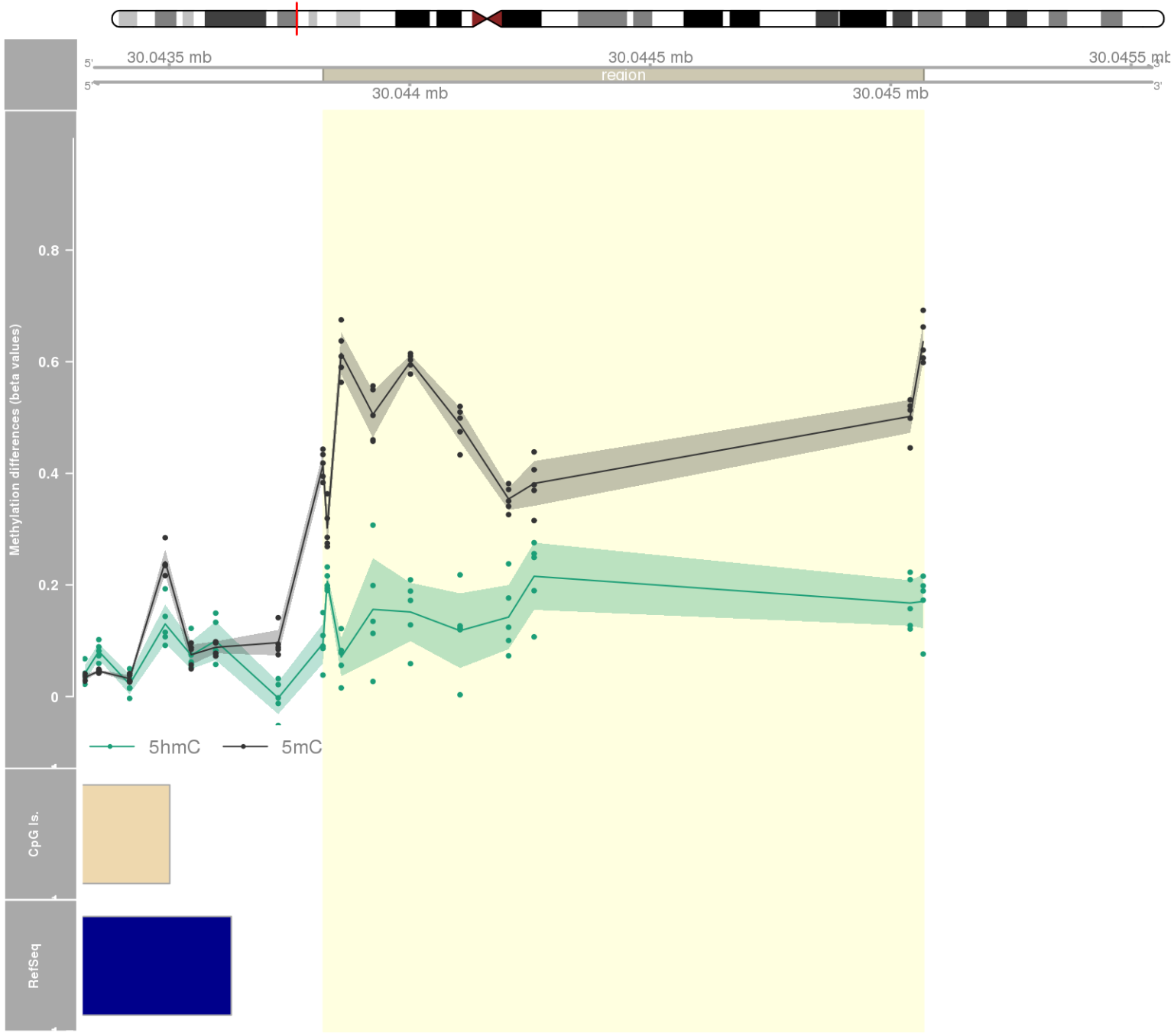
DMR 329 // chr7:24797656-24798855 // 1199 pb. (9 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.023 // fwerArea: 1
- genes: DFNA5 -



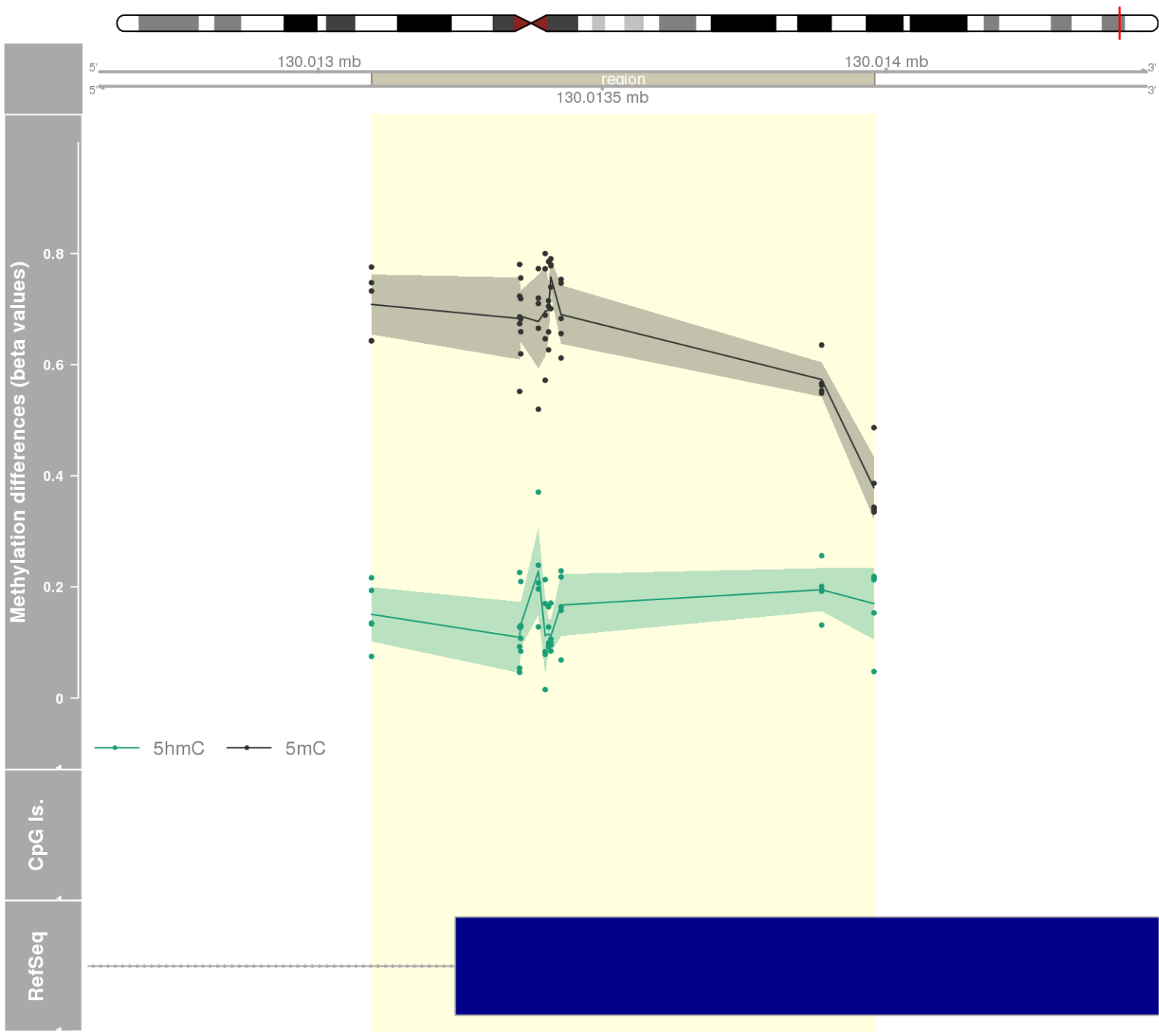
DMR 330 // chr1:59280370-59281383 // 1013 pb. (7 probes) // pvalue: 0.006 // fwer: 0.173 // pvalueArea: 0.023 // fwerArea: 1
- genes: LINC01135 -



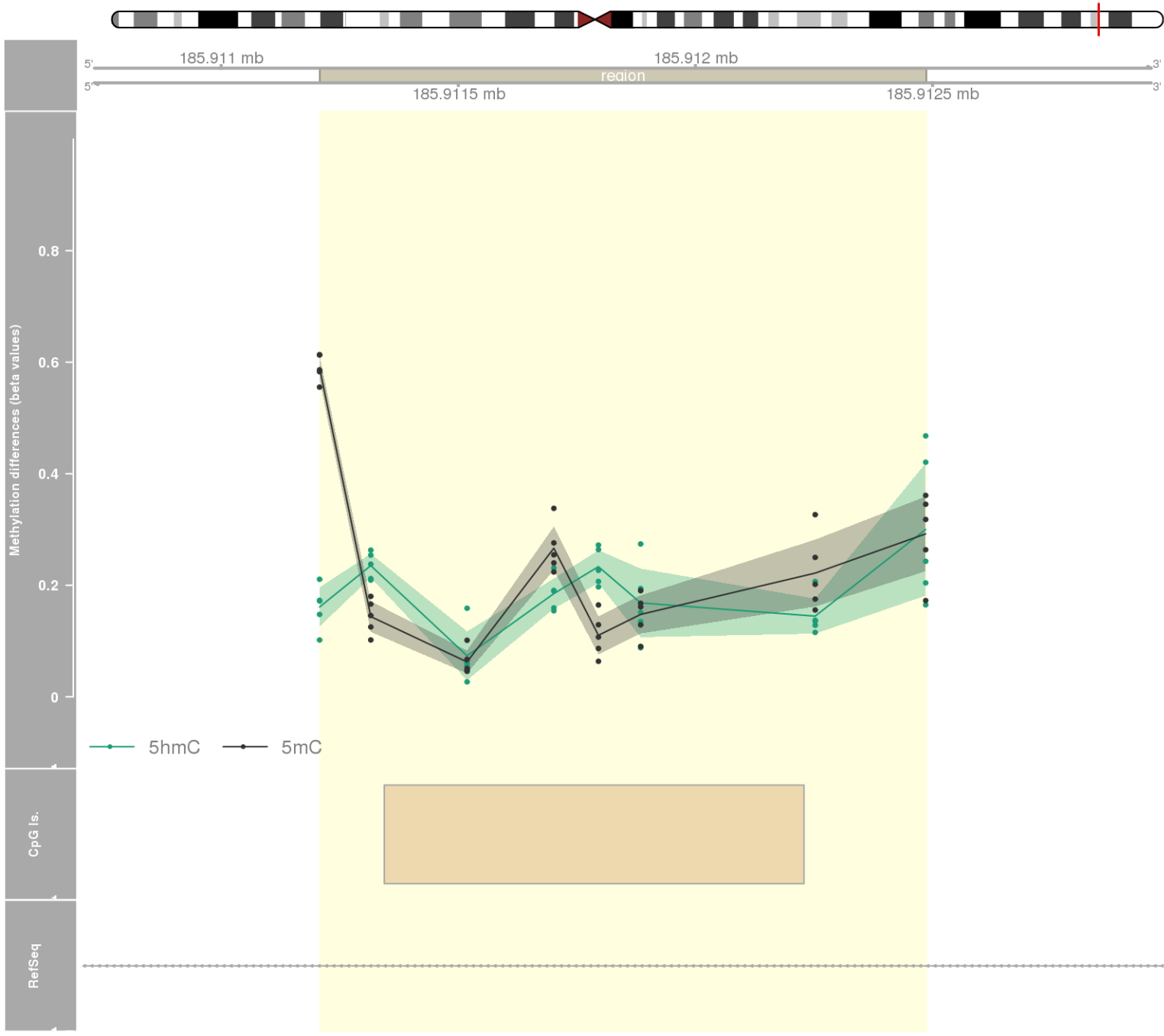
DMR 331 // chr6:30043820-30045068 // 1248 pb. (10 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.023 // fwerArea: 1



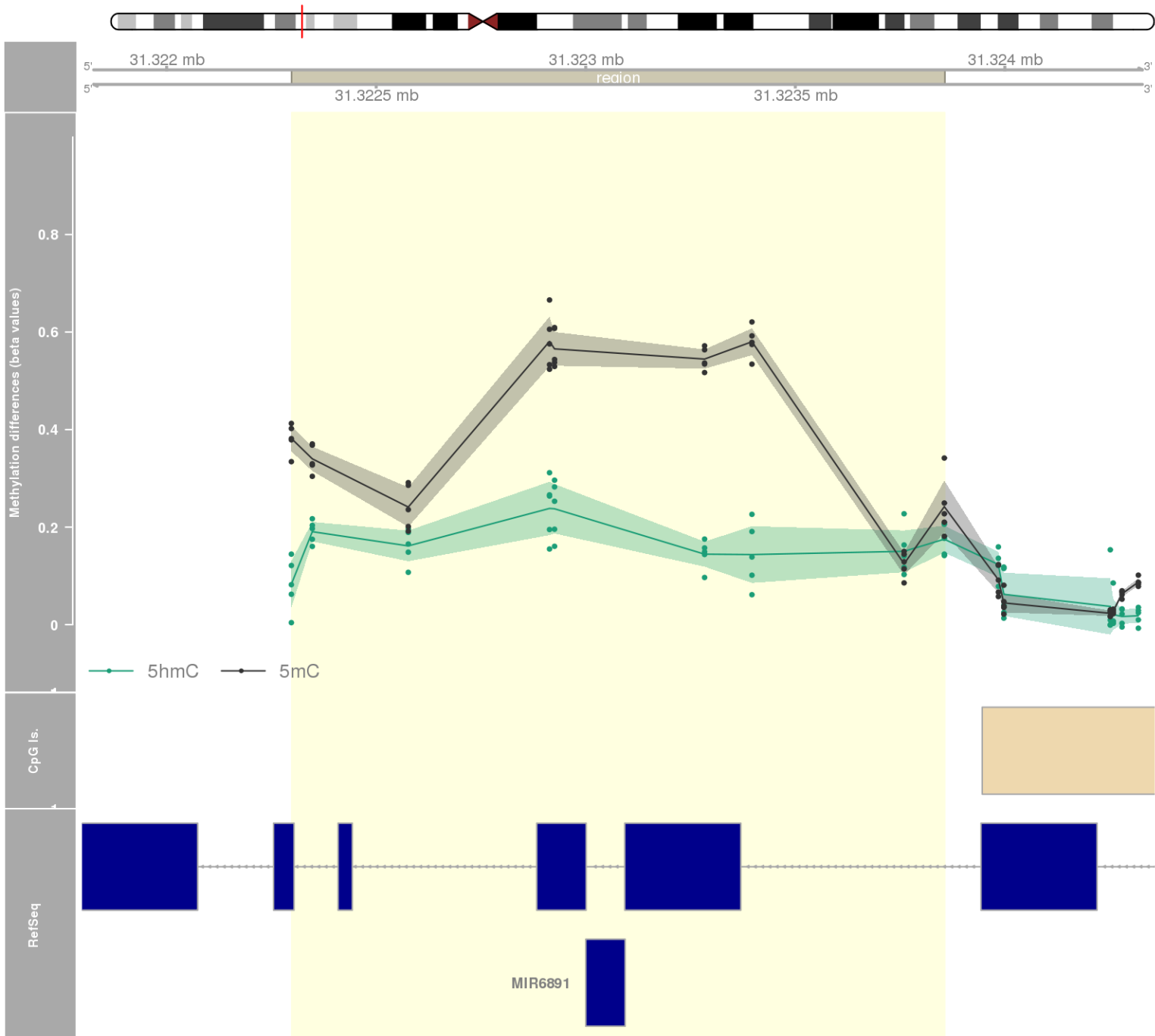
DMR 332 // chr11:130013094-130013979 // 885 pb. (10 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.023 // fwerArea: 1
- genes: APLP2 -



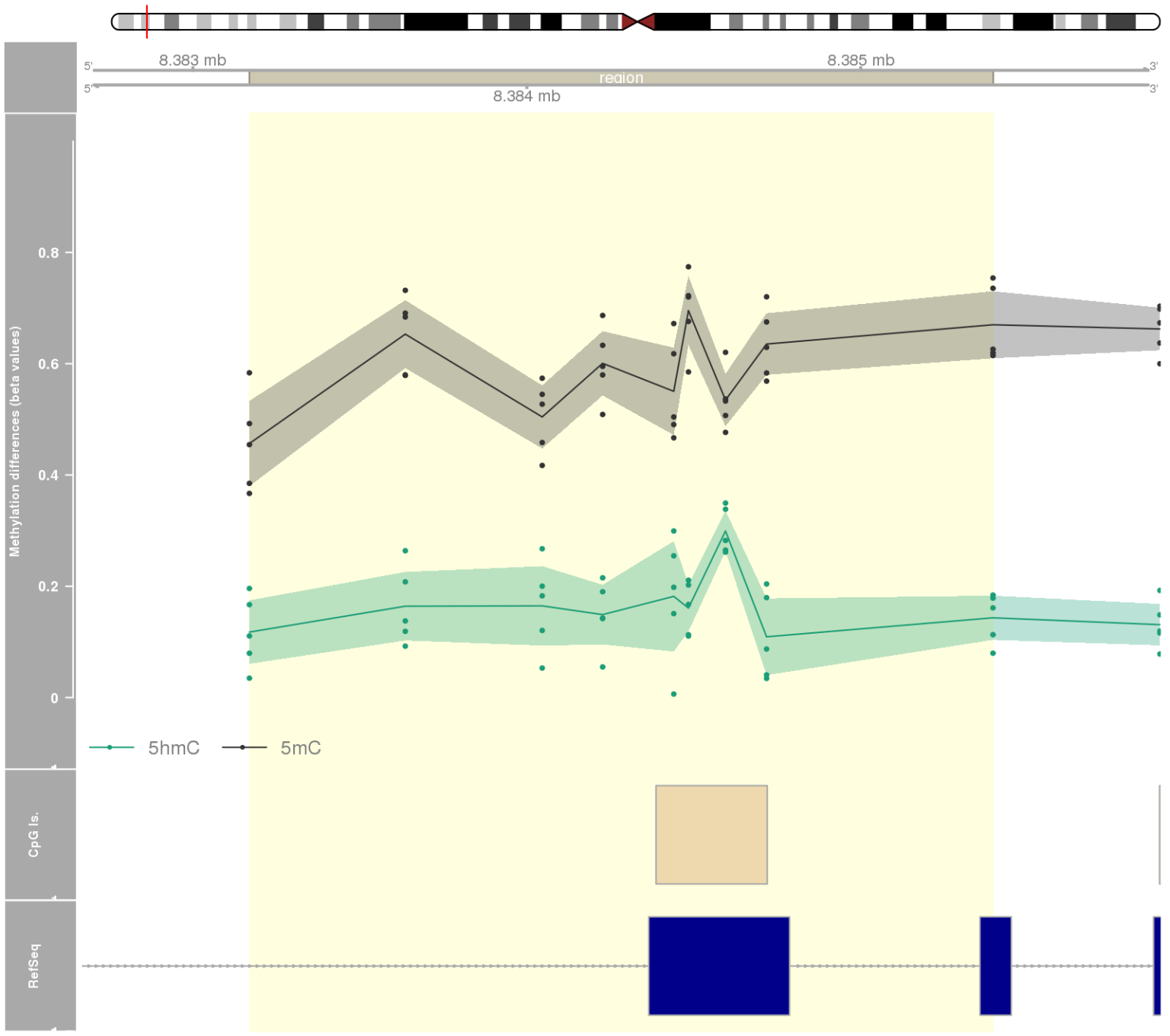
DMR 333 // chr3:185911208-185912486 // 1278 pb. (8 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.023 // fwerArea: 1
- genes: DGKG -



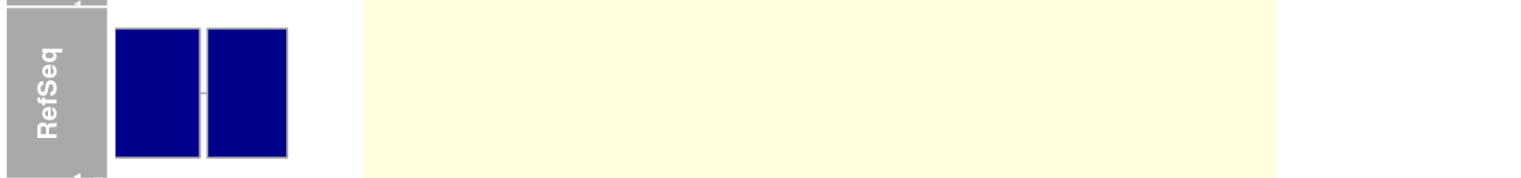
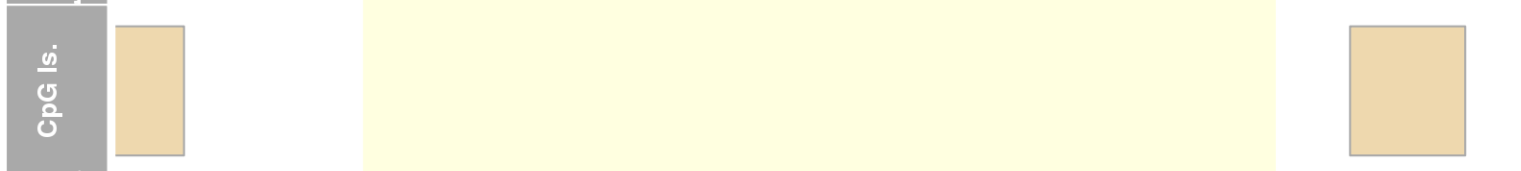
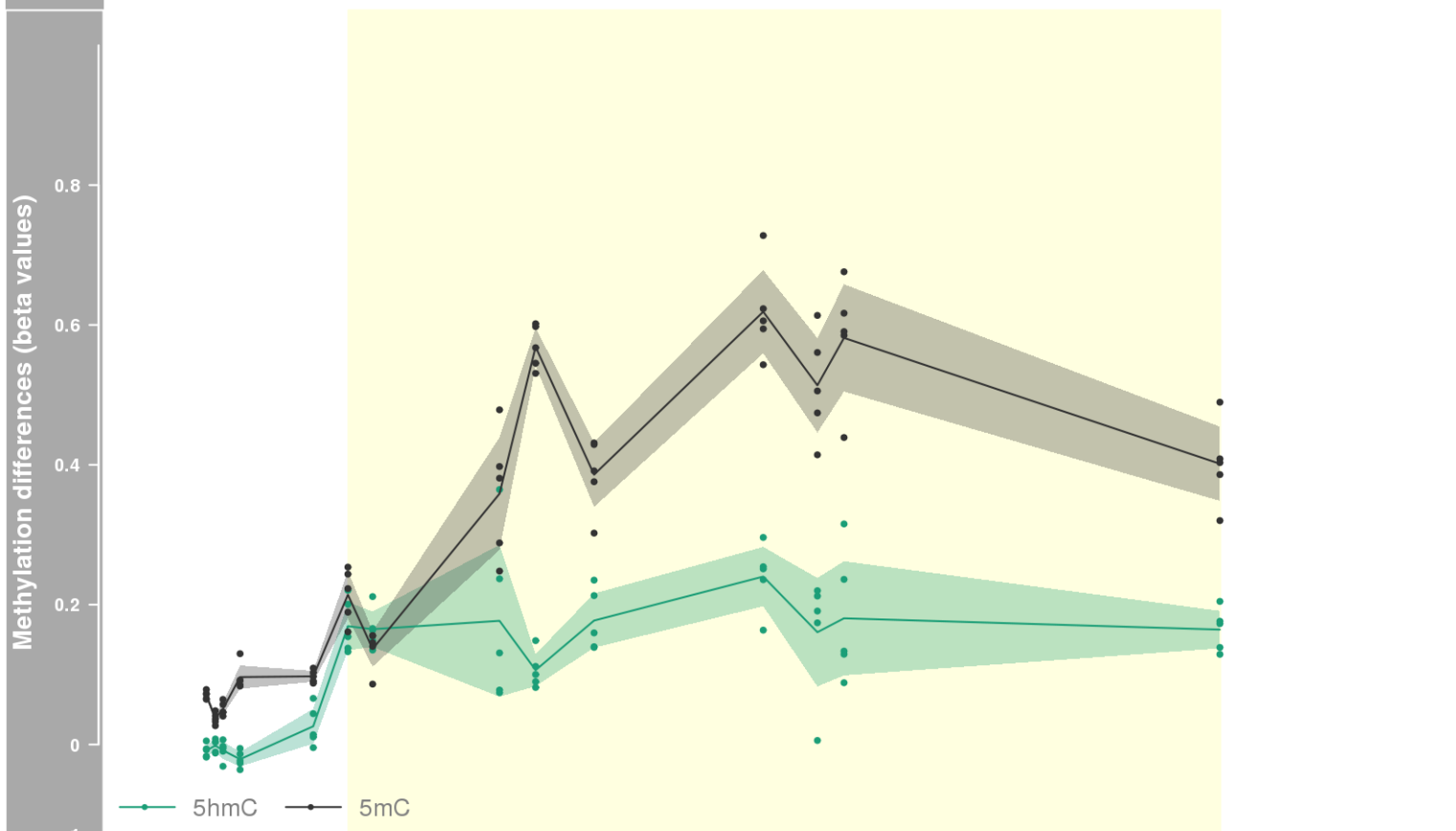
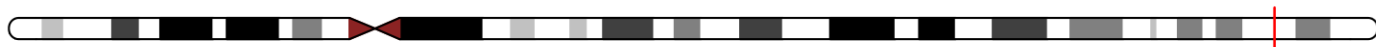
DMR 334 // chr6:31322298-31323856 // 1558 pb. (9 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.023 // fwerArea: 1
- genes: HLA-B -



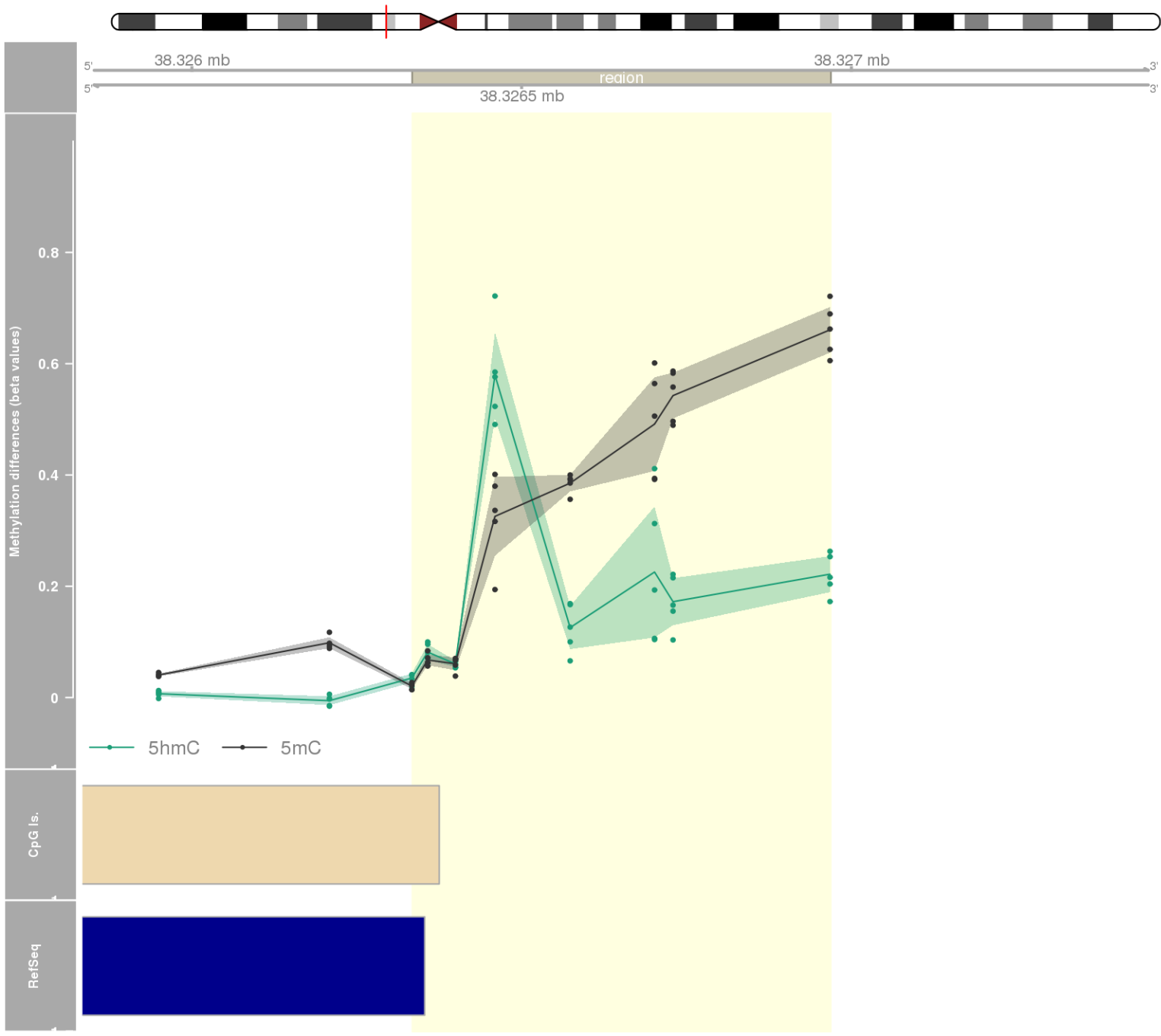
DMR 335 // chr1:8383169-8385396 // 2227 pb. (9 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.023 // fwerArea: 1
- genes: SLC45A1 -



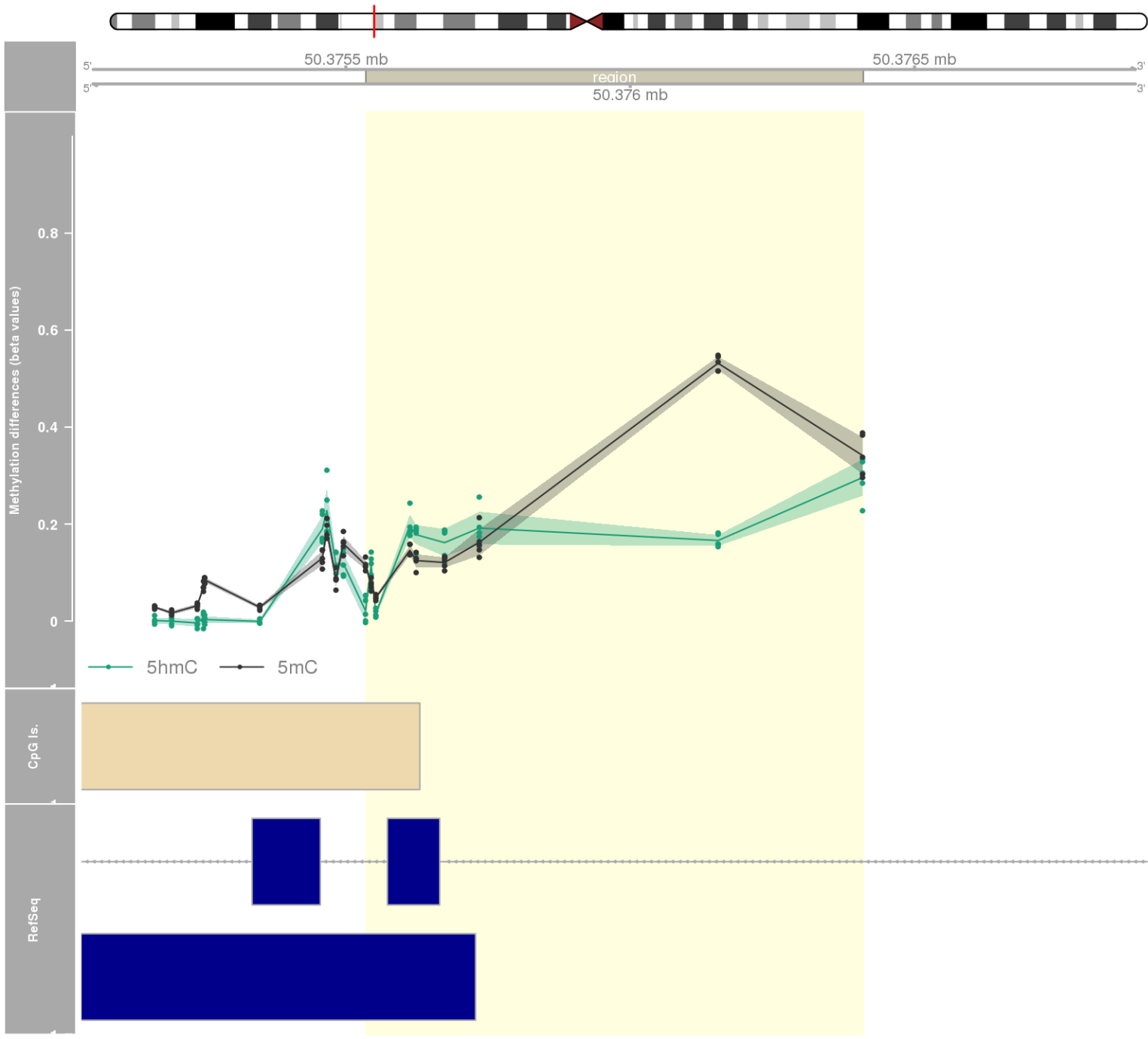
DMR 336 // chr12:123757018-123758853 // 1835 pb. (9 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.023 // fwerArea: 1



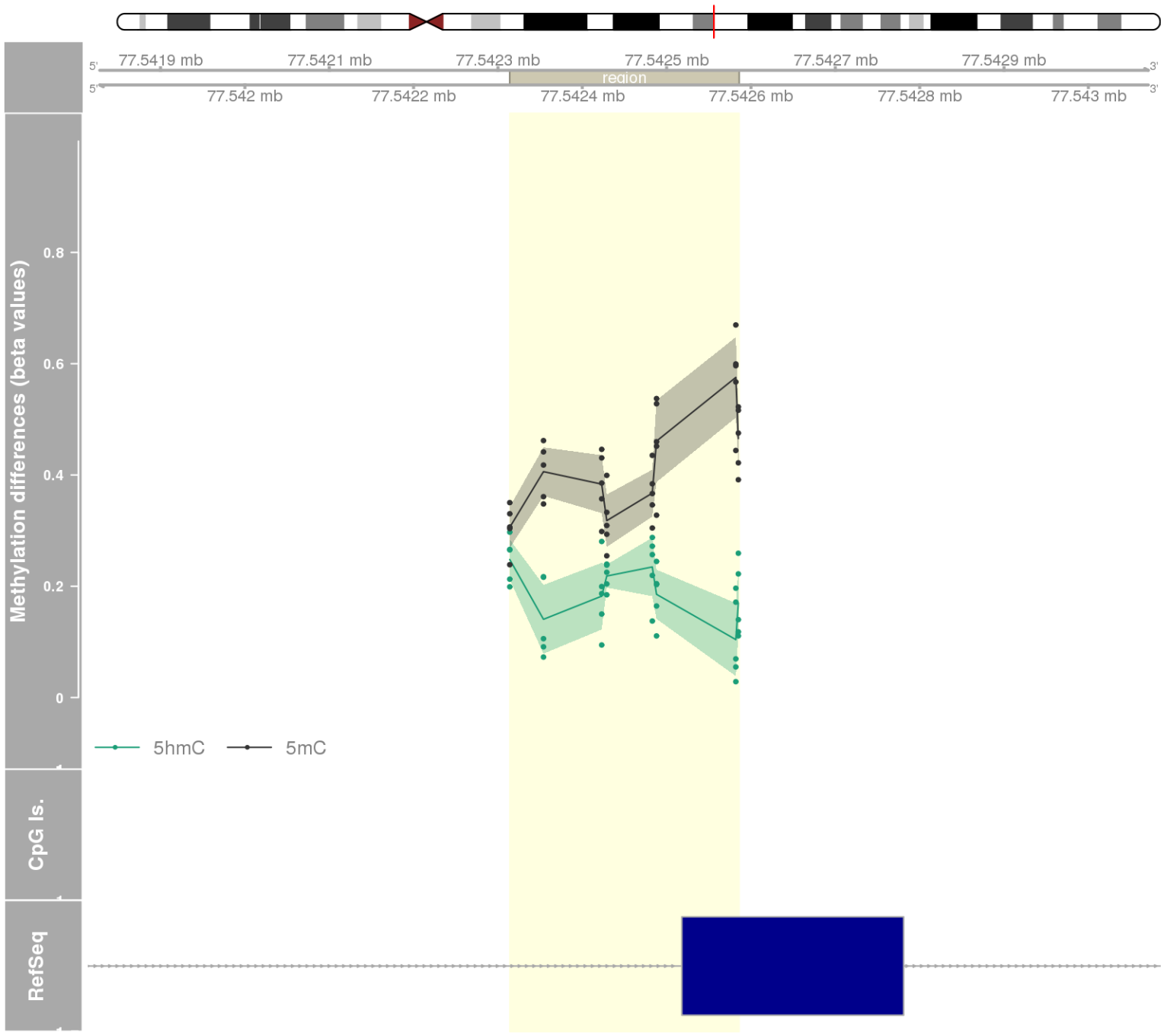
DMR 337 // chr8:38326334-38326968 // 634 pb. (8 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.024 // fwerArea: 1
- genes: FGFR1 -



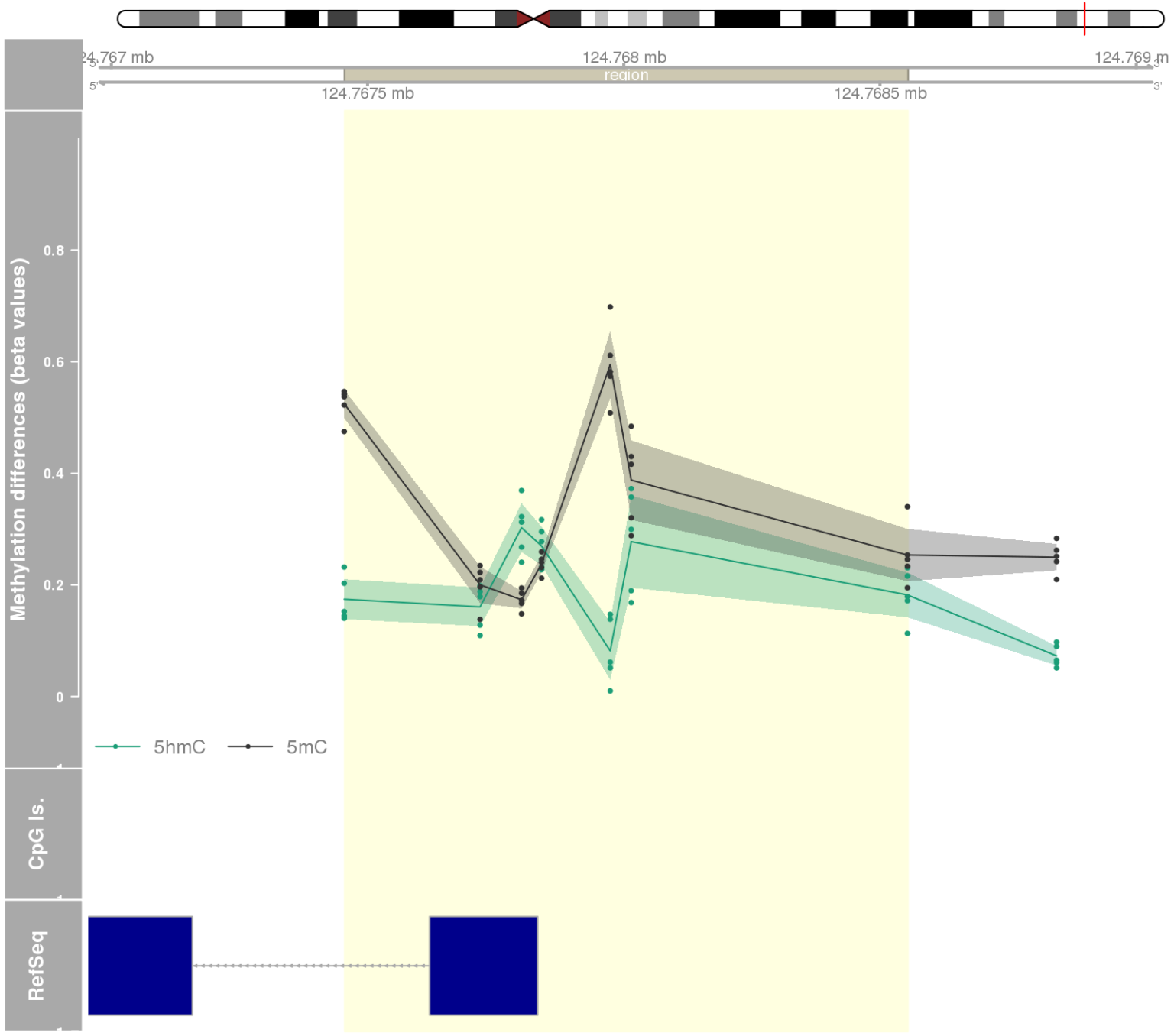
DMR 338 // chr3:50375535-50376409 // 874 pb. (9 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.024 // fwerArea: 1
- genes: RASSF1 -



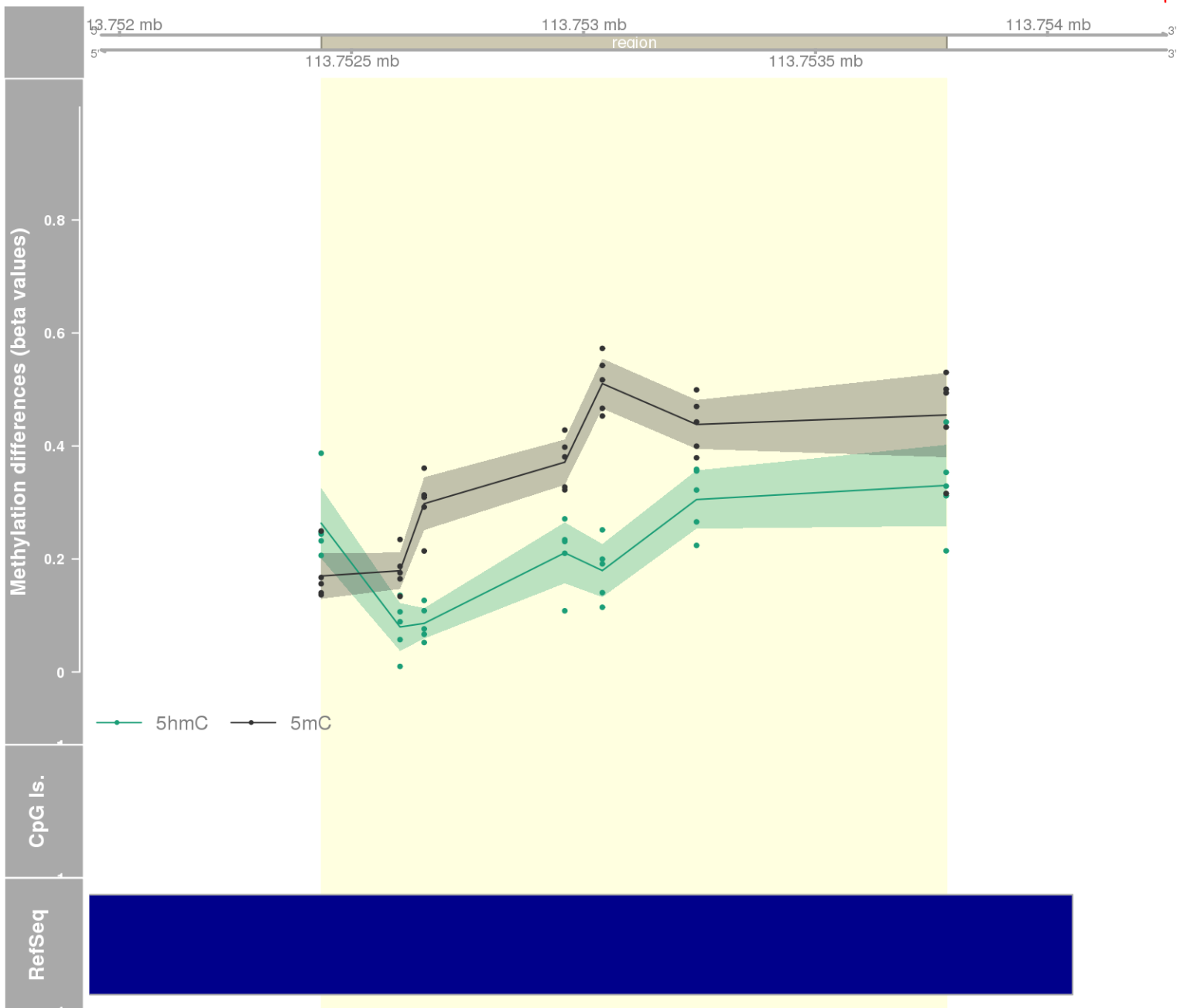
DMR 339 // chr10:77542314-77542585 // 271 pb. (8 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.024 // fwerArea: 1
- genes: C10orf11 -



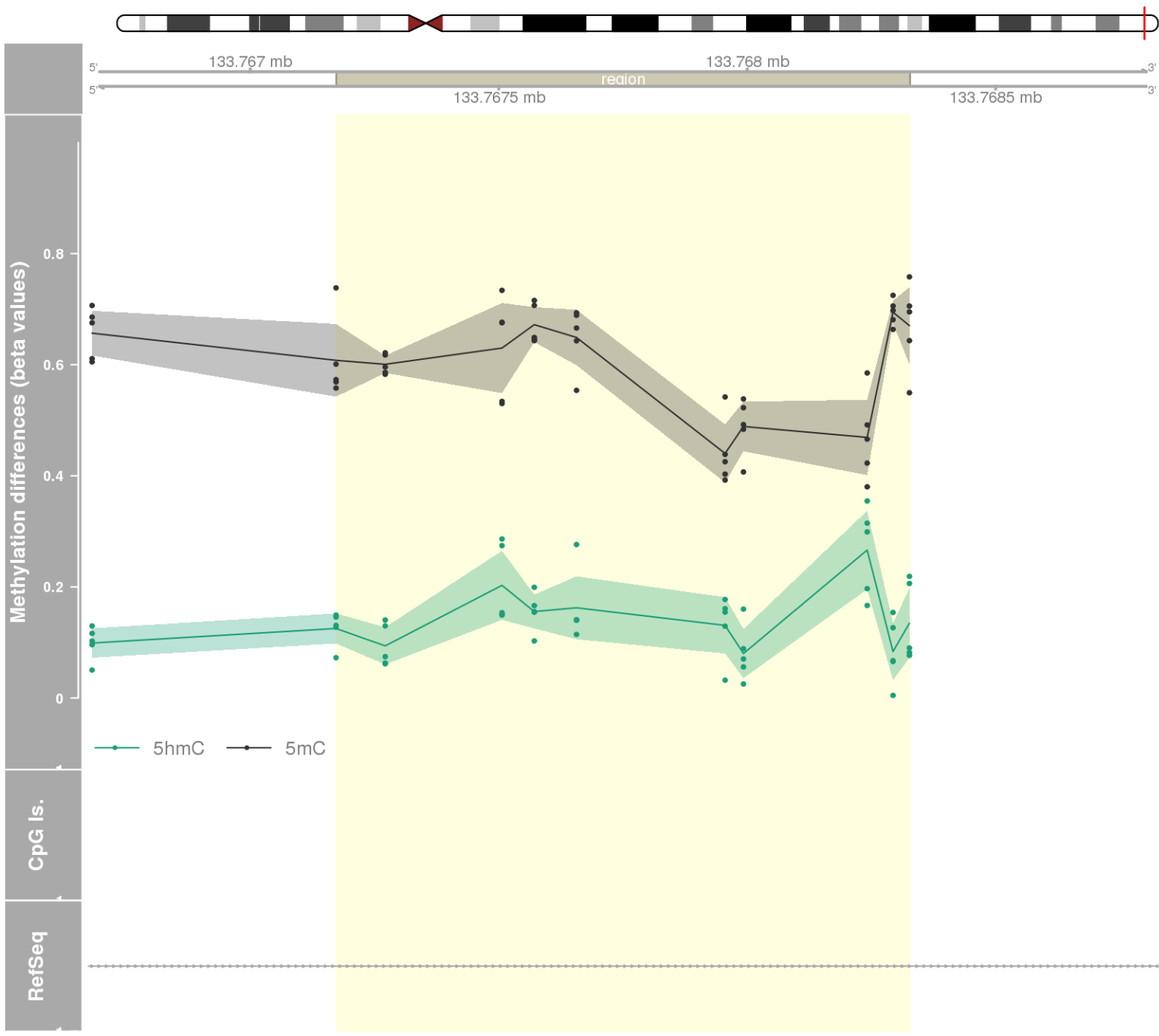
DMR 340 // chr11:124767455-124768554 // 1099 pb. (7 probes) // pvalue: 0.006 // fwer: 0.173 // pvalueArea: 0.024 // fwerArea: 1
- genes: ROBO4 -



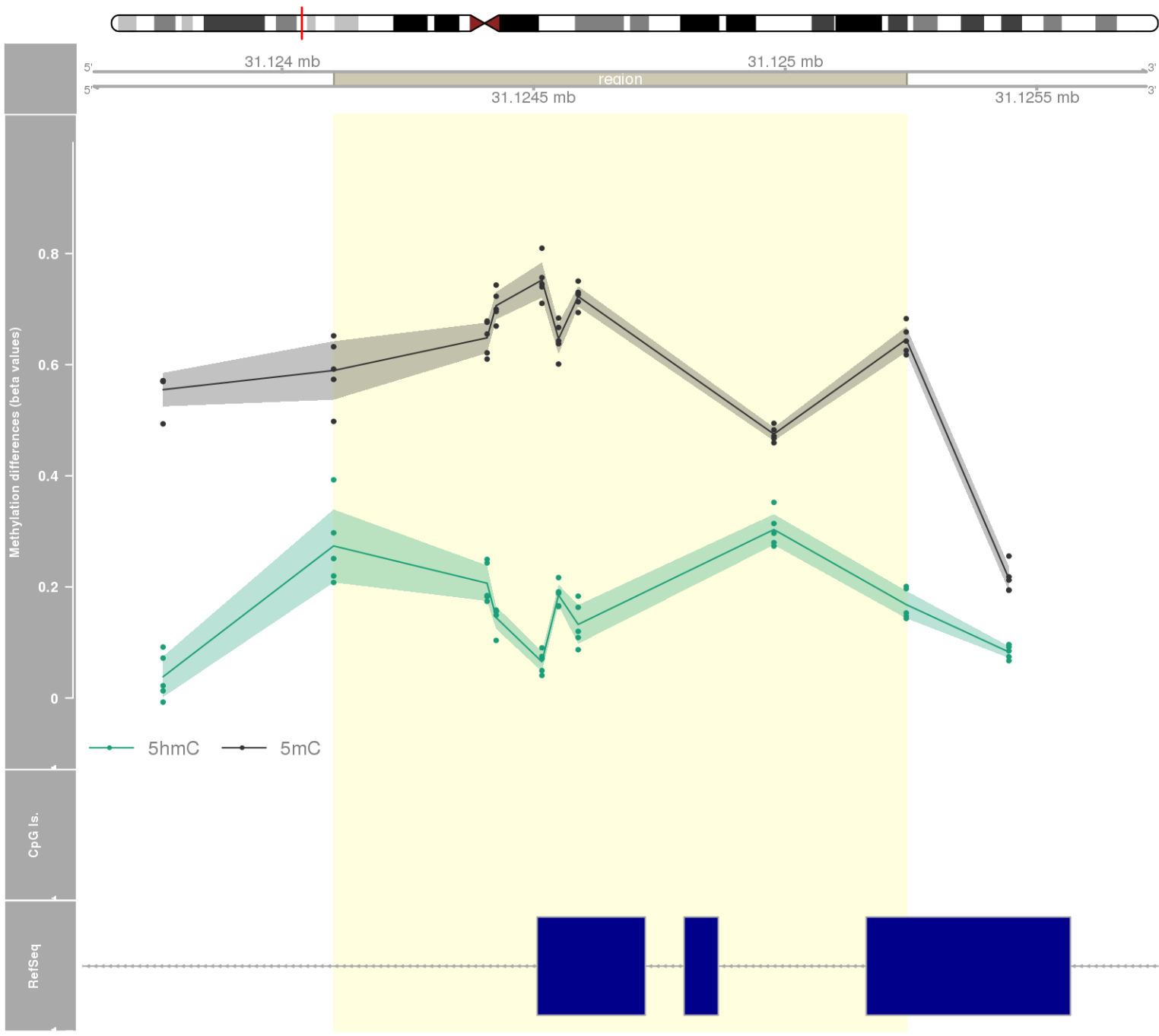
DMR 341 // chr13:113752435-113753782 // 1347 pb. (7 probes) // pvalue: 0.006 // fwer: 0.173 // pvalueArea: 0.024 // fwerArea: 1
- genes: MCF2L -



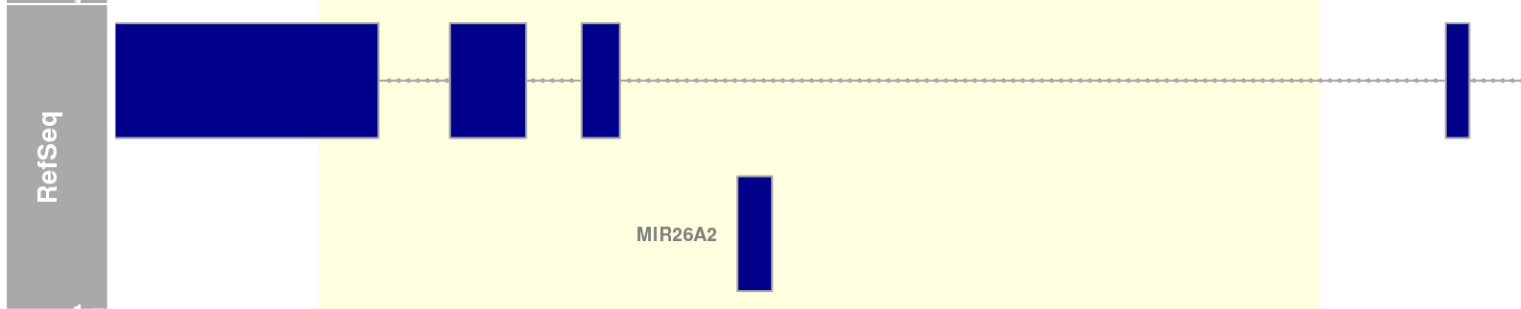
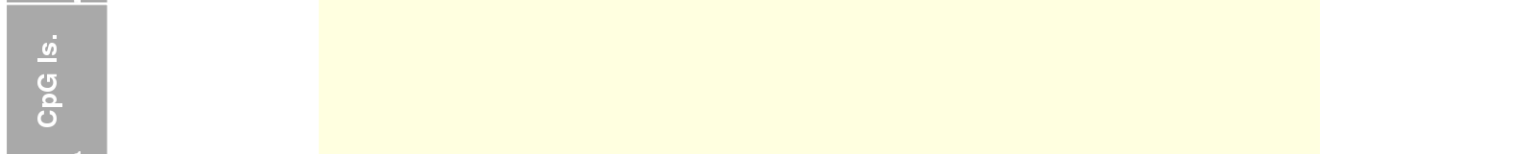
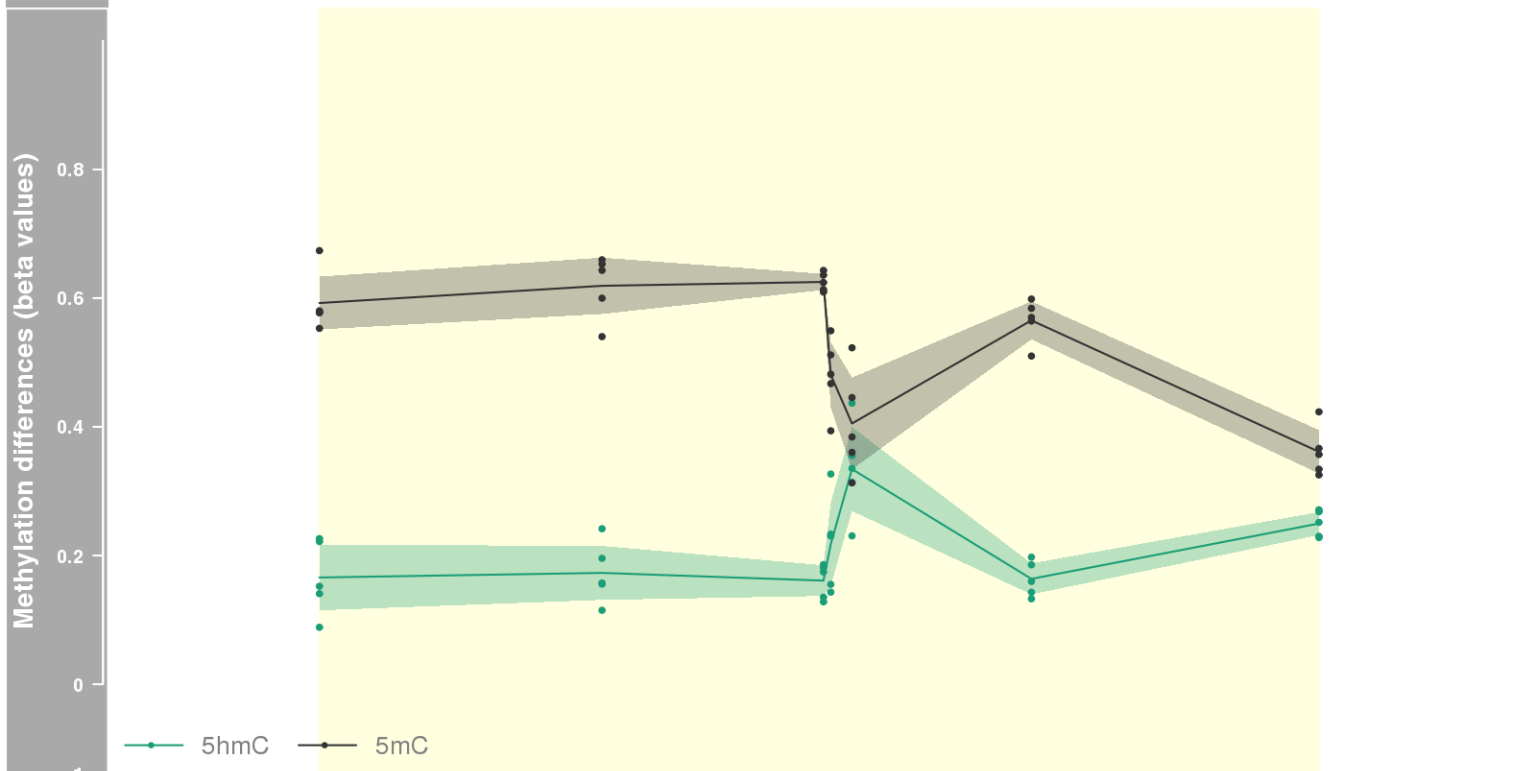
DMR 342 // chr10:133767173-133768327 // 1154 pb. (10 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.024 // fwerArea: 1
- genes: PPP2R2D -



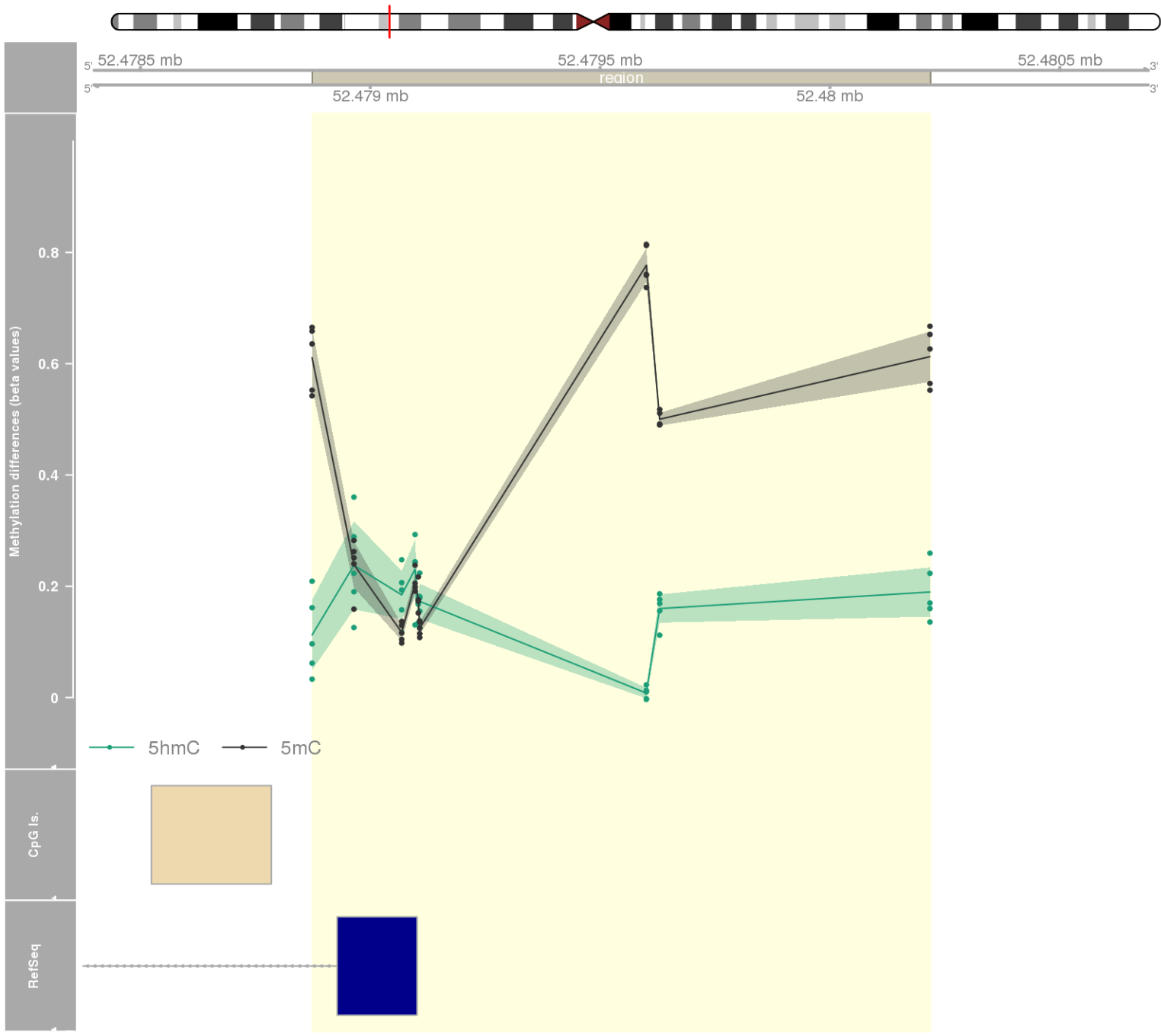
DMR 343 // chr6:31124103-31125241 // 1138 pb. (8 probes) // pvalue: 0.009 // fwer: 0.36 // pvalueArea: 0.024 // fwerArea: 1
- genes: CCHCR1 -



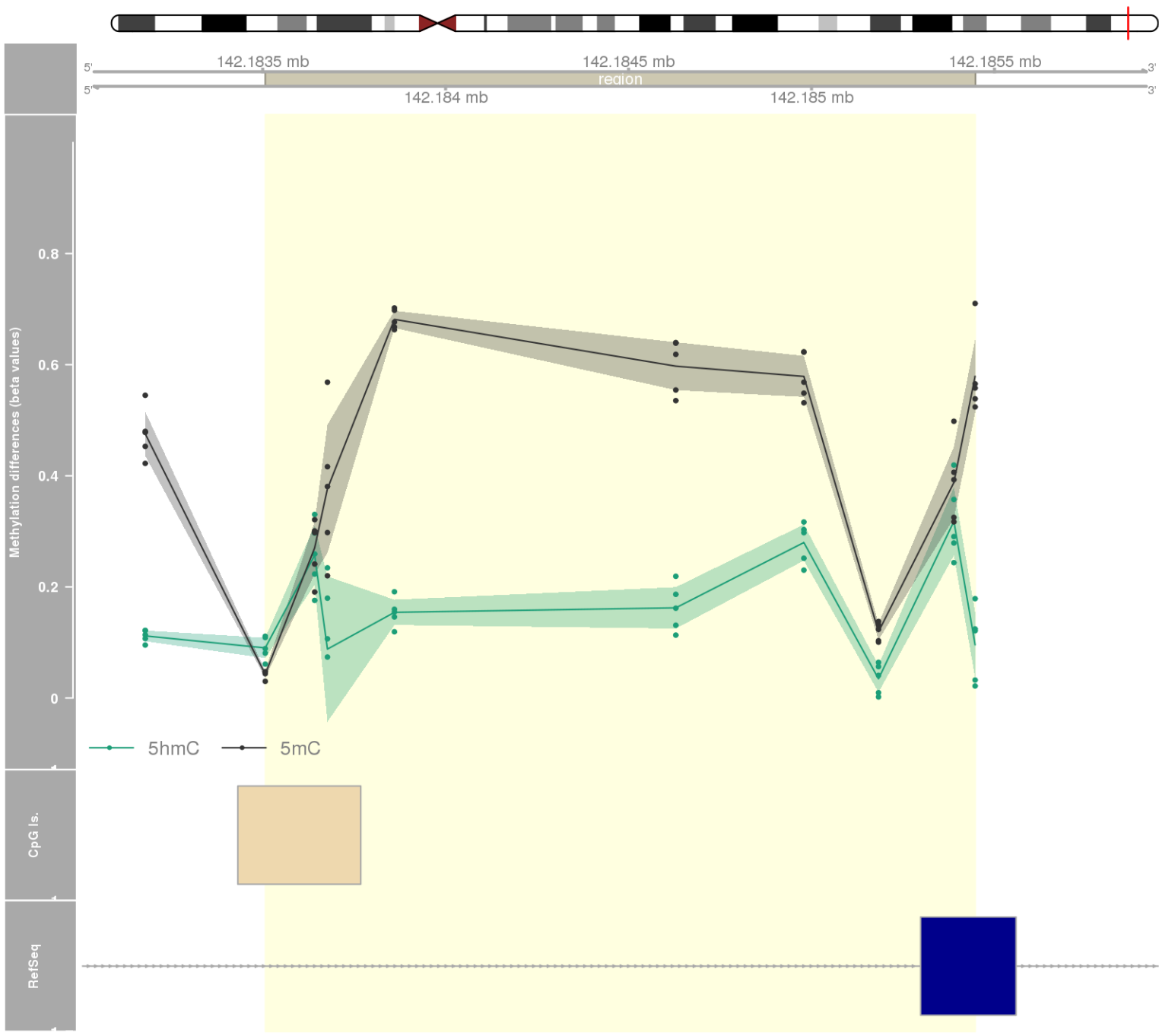
DMR 344 // chr12:58217366-58219817 // 2451 pb. (7 probes) // pvalue: 0.006 // fwer: 0.173 // pvalueArea: 0.024 // fwerArea: 1
- genes: MIR26A2 / CTDSP2 -



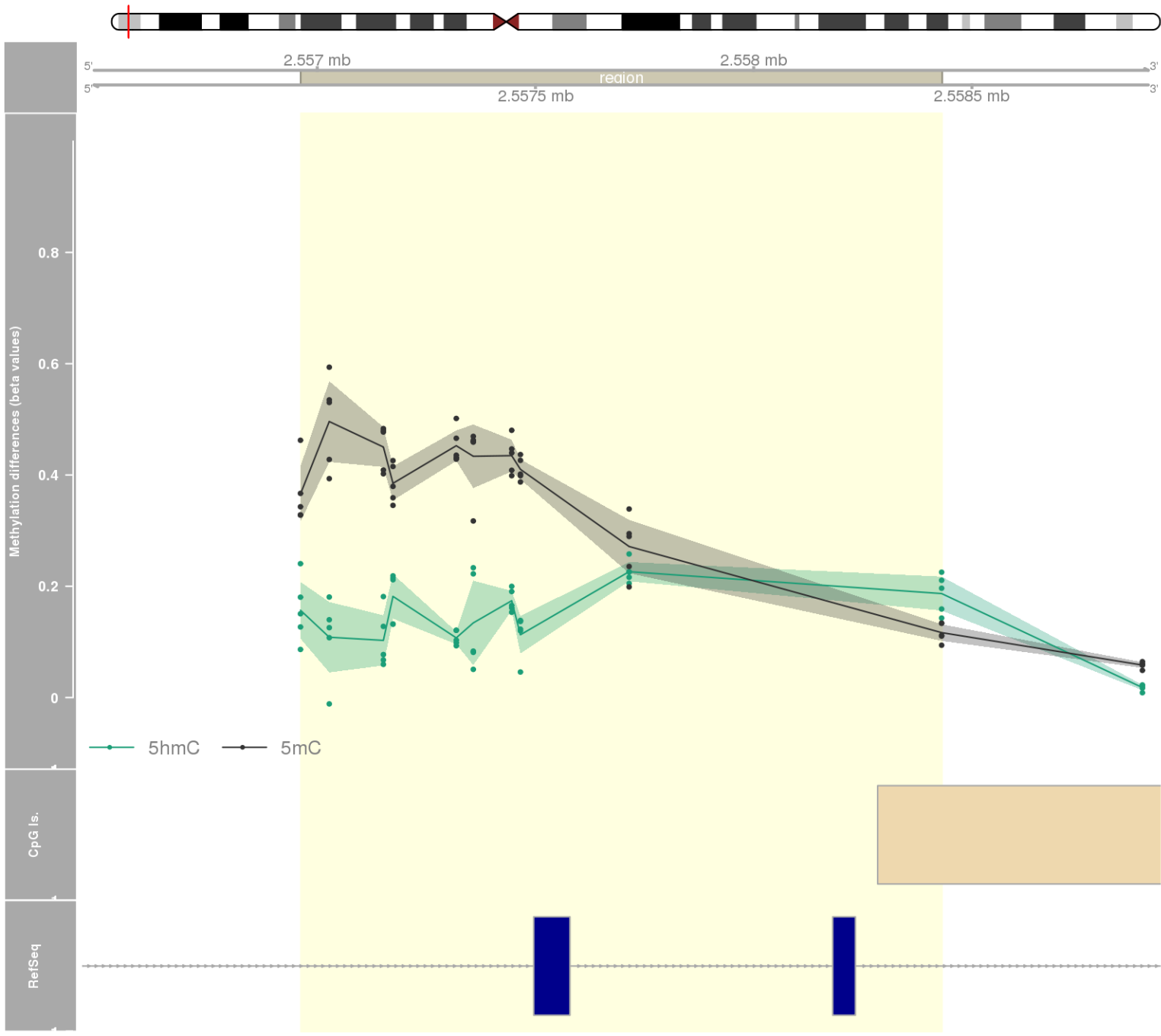
DMR 345 // chr3:52478874-52480218 // 1344 pb. (9 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.024 // fwerArea: 1
- genes: SEMA3G -



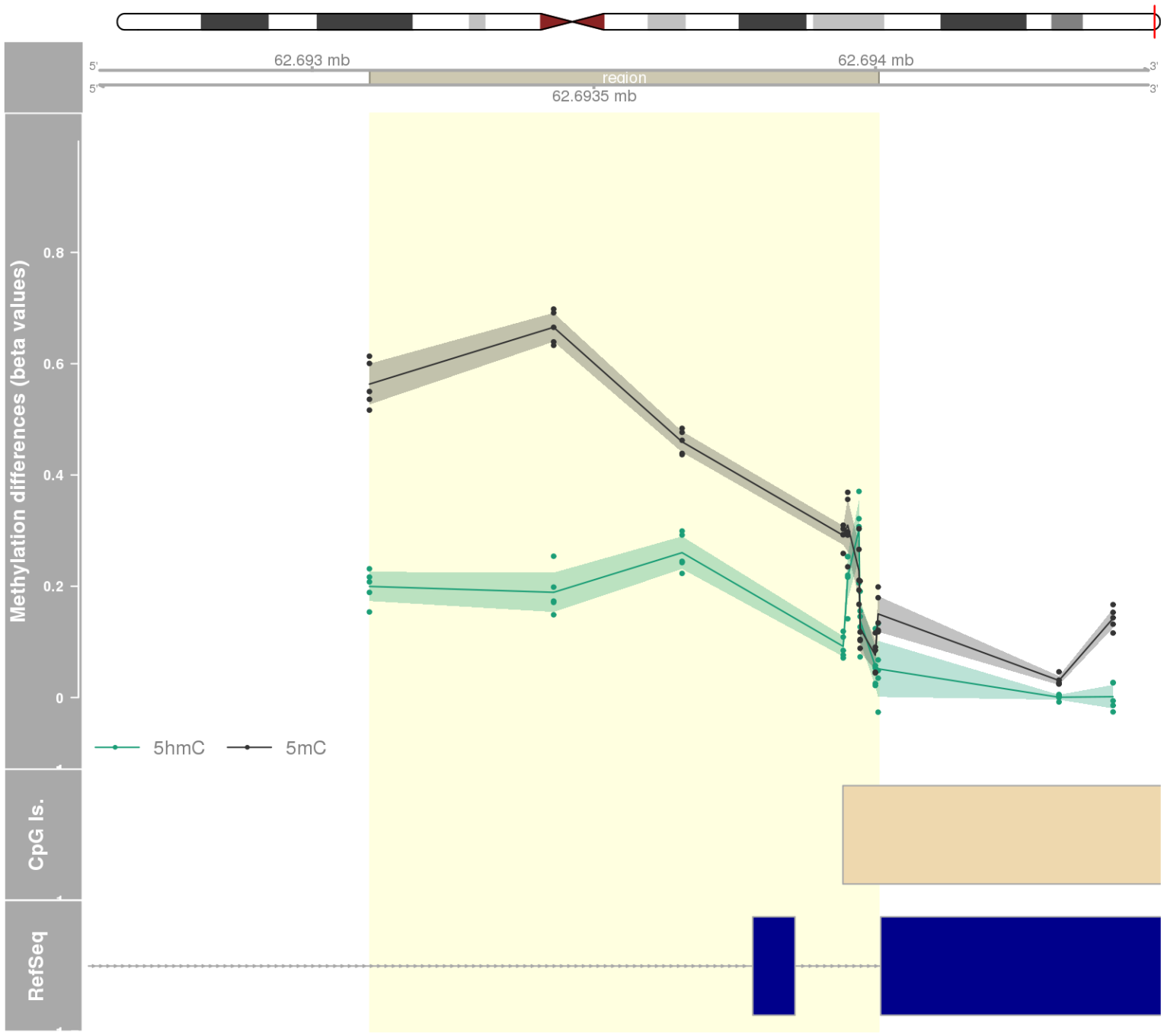
DMR 346 // chr8:142183507-142185447 // 1940 pb. (9 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.024 // fwerArea: 1
- genes: DENND3 -



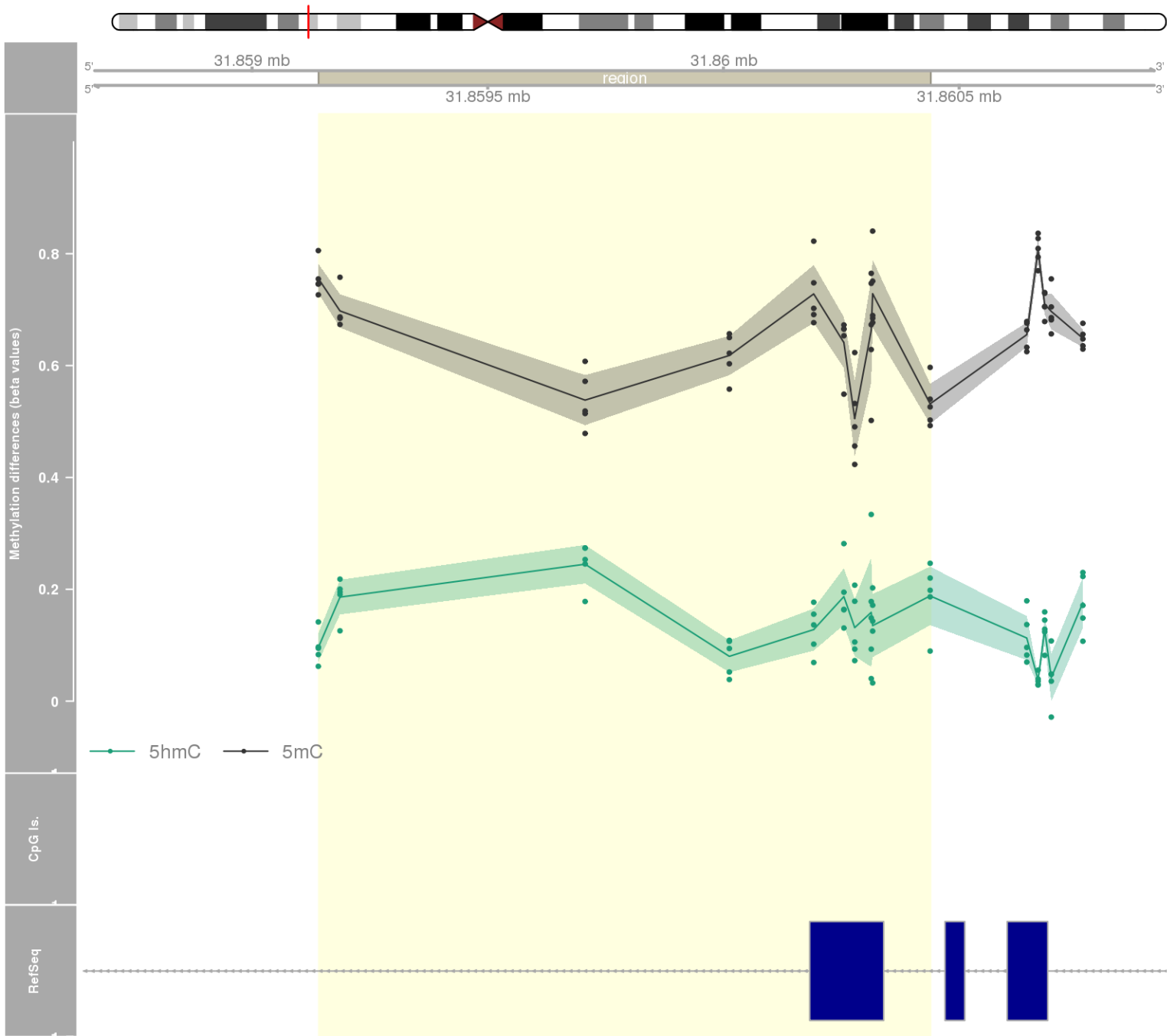
DMR 347 // chr7:2556962-2558431 // 1469 pb. (10 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.024 // fwerArea: 1
- genes: LFNG -



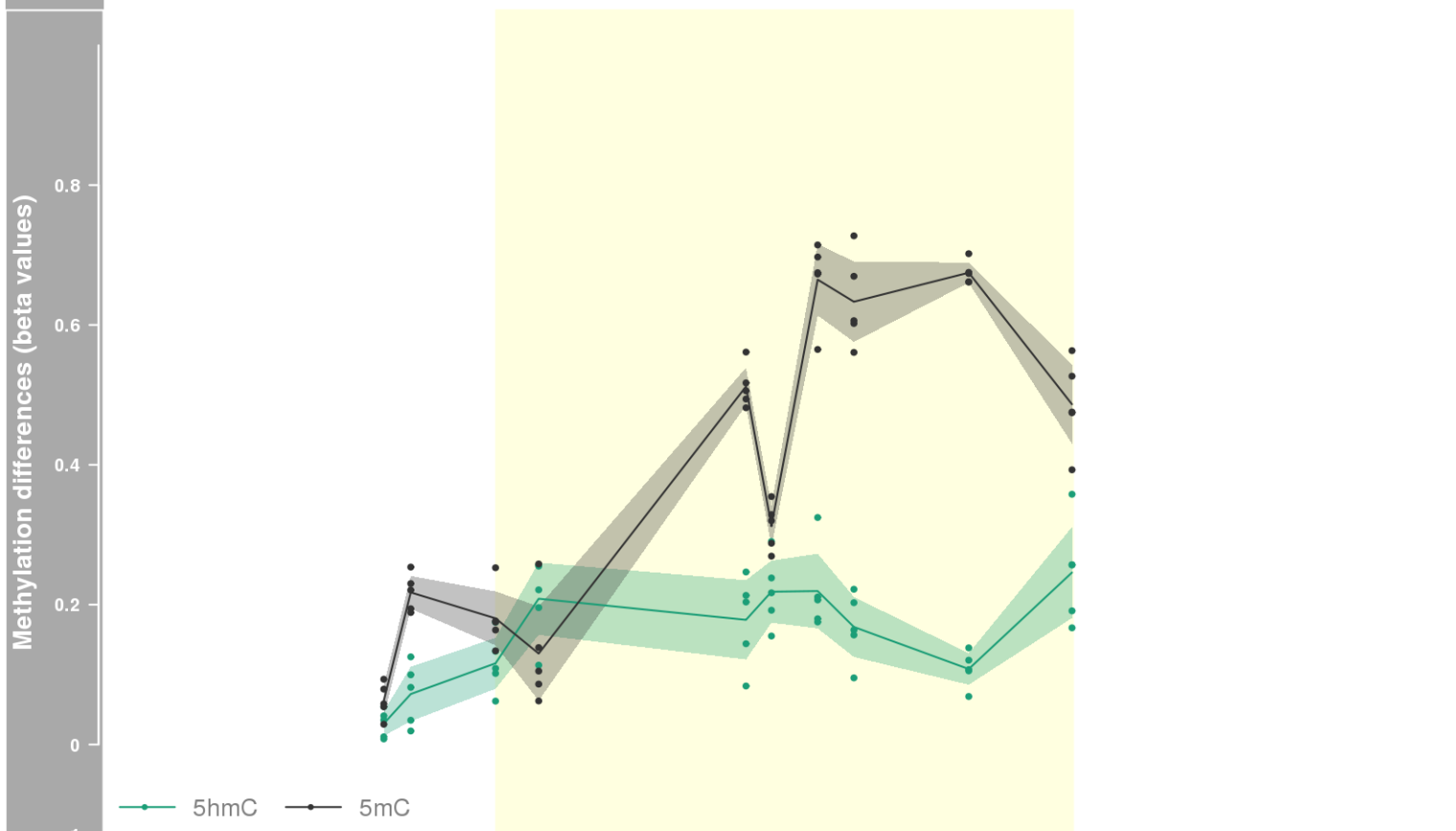
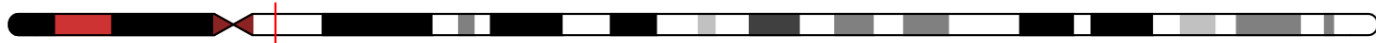
DMR 348 // chr20:62693102-62694005 // 903 pb. (9 probes) // pvalue: 0.012 // fwer: 0.36 // pvalueArea: 0.024 // fwerArea: 1
- genes: TCEA2 -



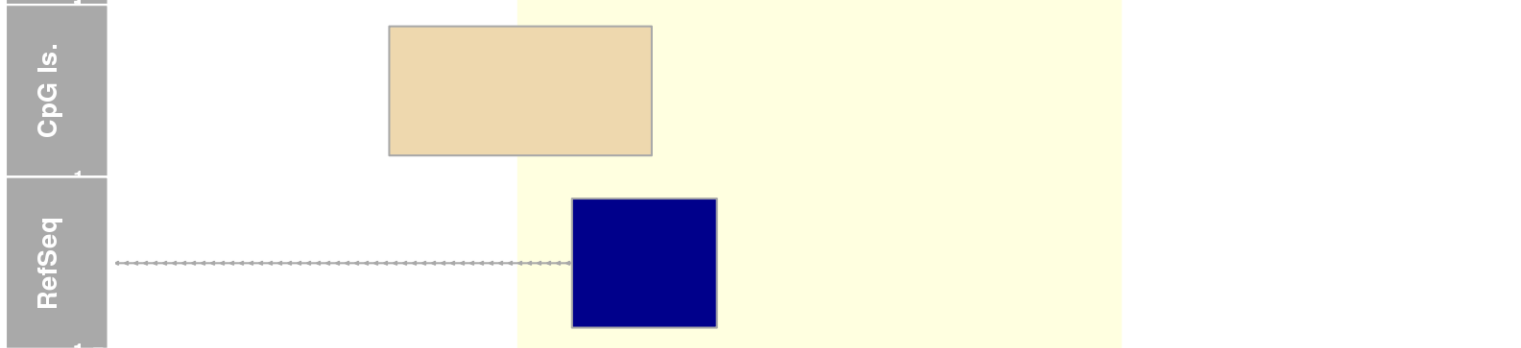
DMR 349 // chr6:31859141-31860439 // 1298 pb. (10 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.024 // fwerArea: 1
- genes: EHMT2 -



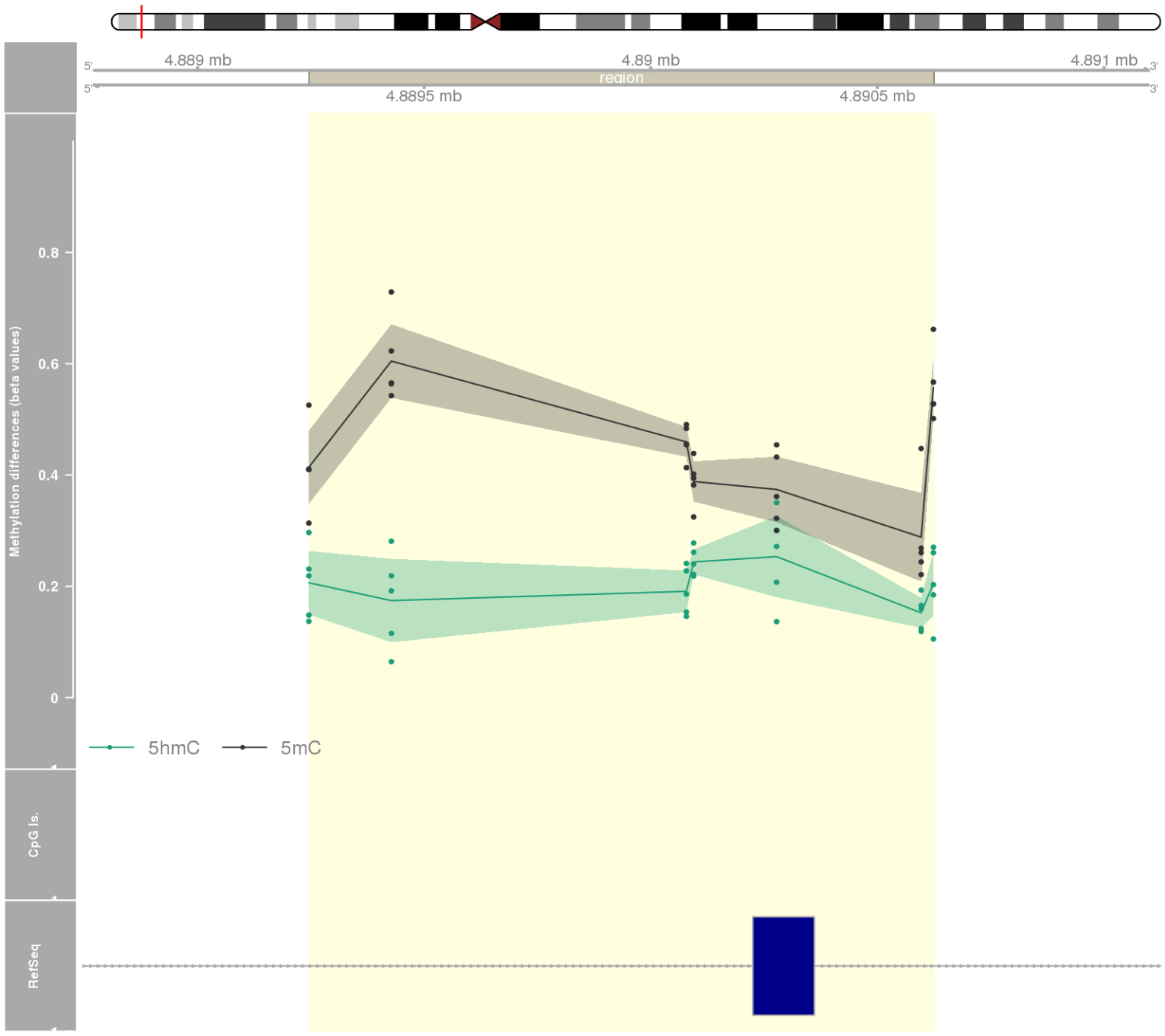
DMR 350 // chr14:20903555-20904303 // 748 pb. (8 probes) // pvalue: 0.01 // fwer: 0.522 // pvalueArea: 0.024 // fwerArea: 1
- genes: KLHL33 -



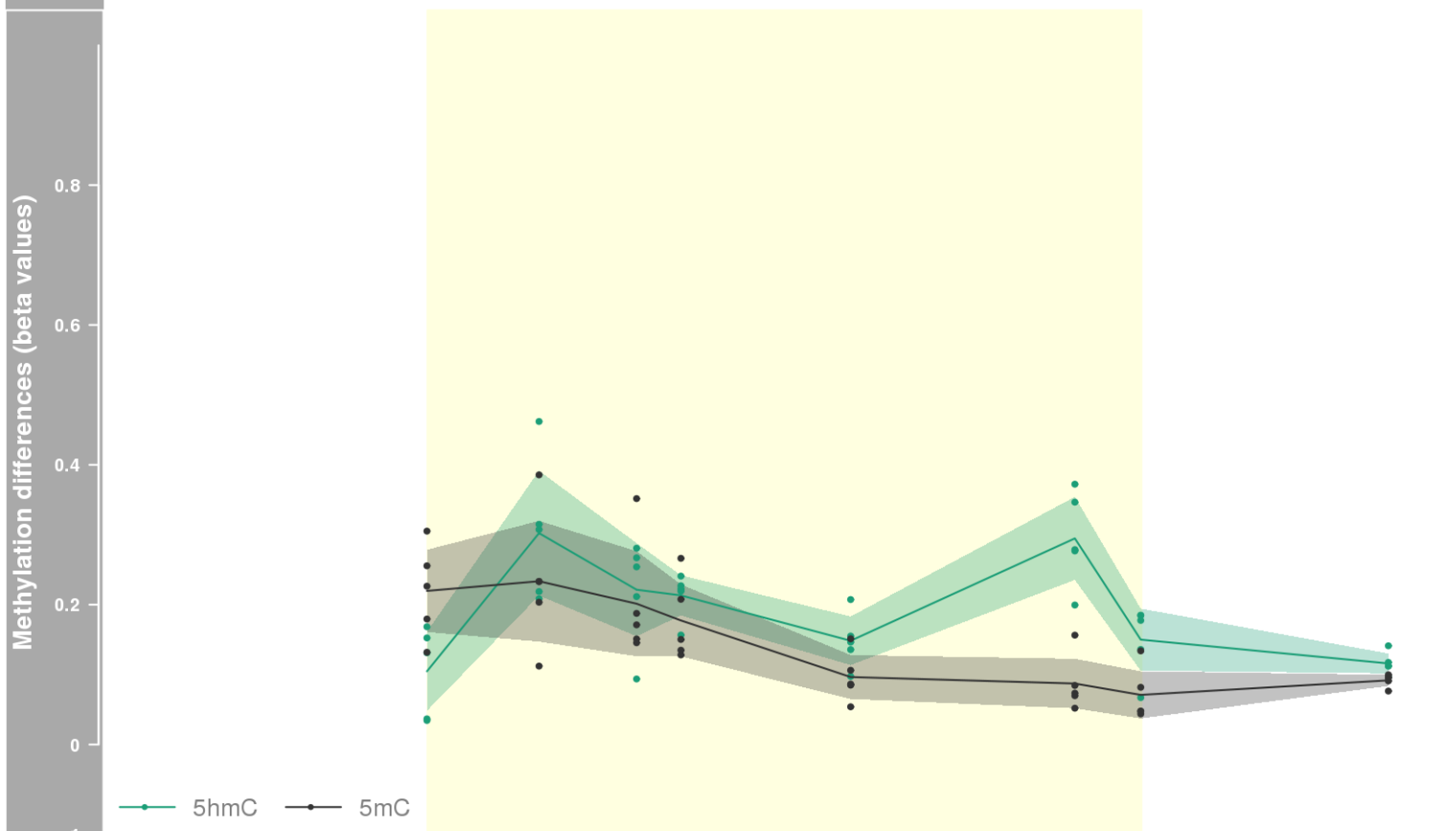
5hmC 5mC



DMR 351 // chr6:4889246-4890624 // 1378 pb. (7 probes) // pvalue: 0.006 // fwer: 0.173 // pvalueArea: 0.025 // fwerArea: 1
- genes: CDYL -



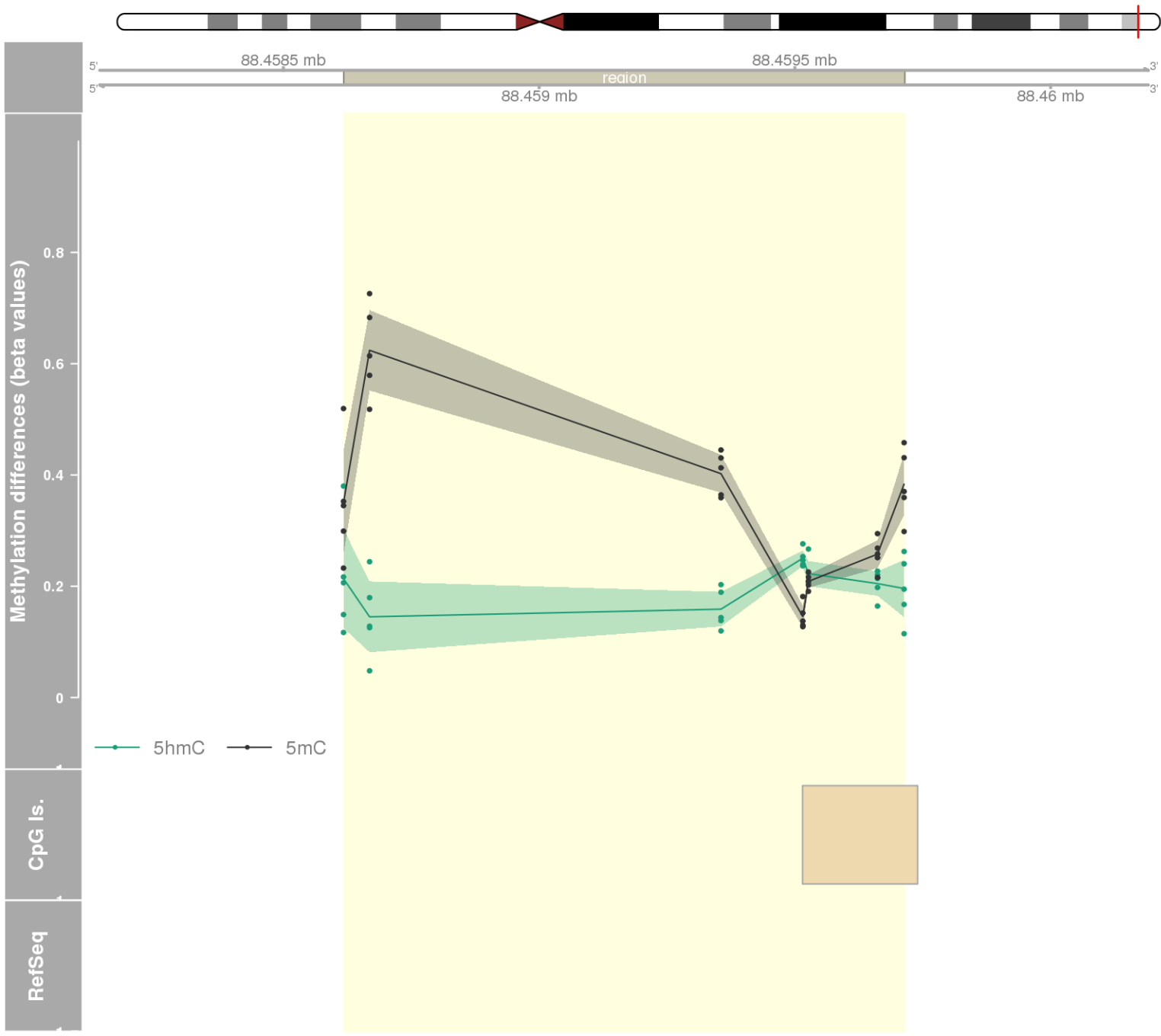
DMR 352 // chr17:1961109-1962236 // 1127 pb. (7 probes) // pvalue: 0.007 // fwer: 0.173 // pvalueArea: 0.025 // fwerArea: 1
- genes: HIC1 -



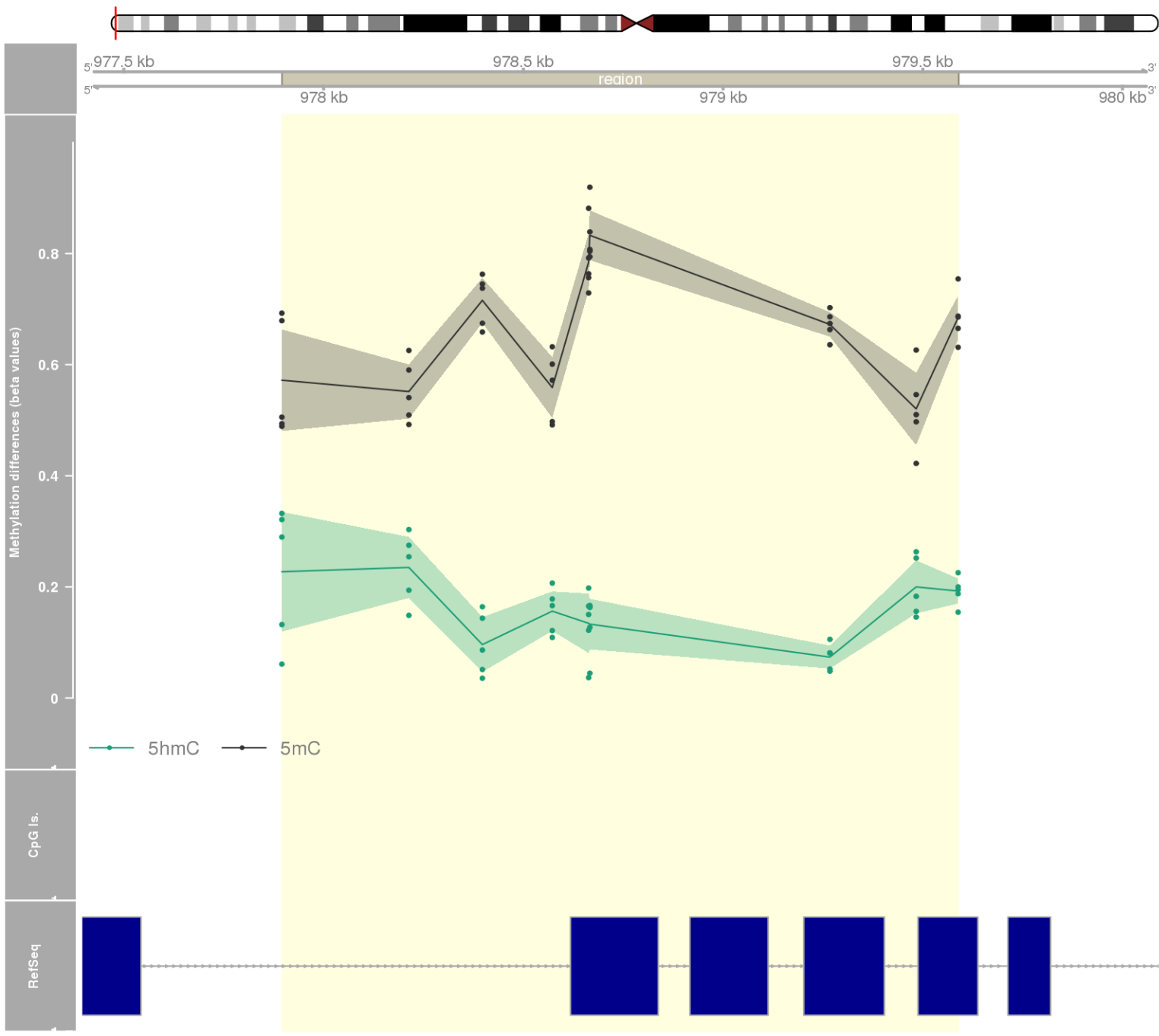
—●— 5hmC —●— 5mC



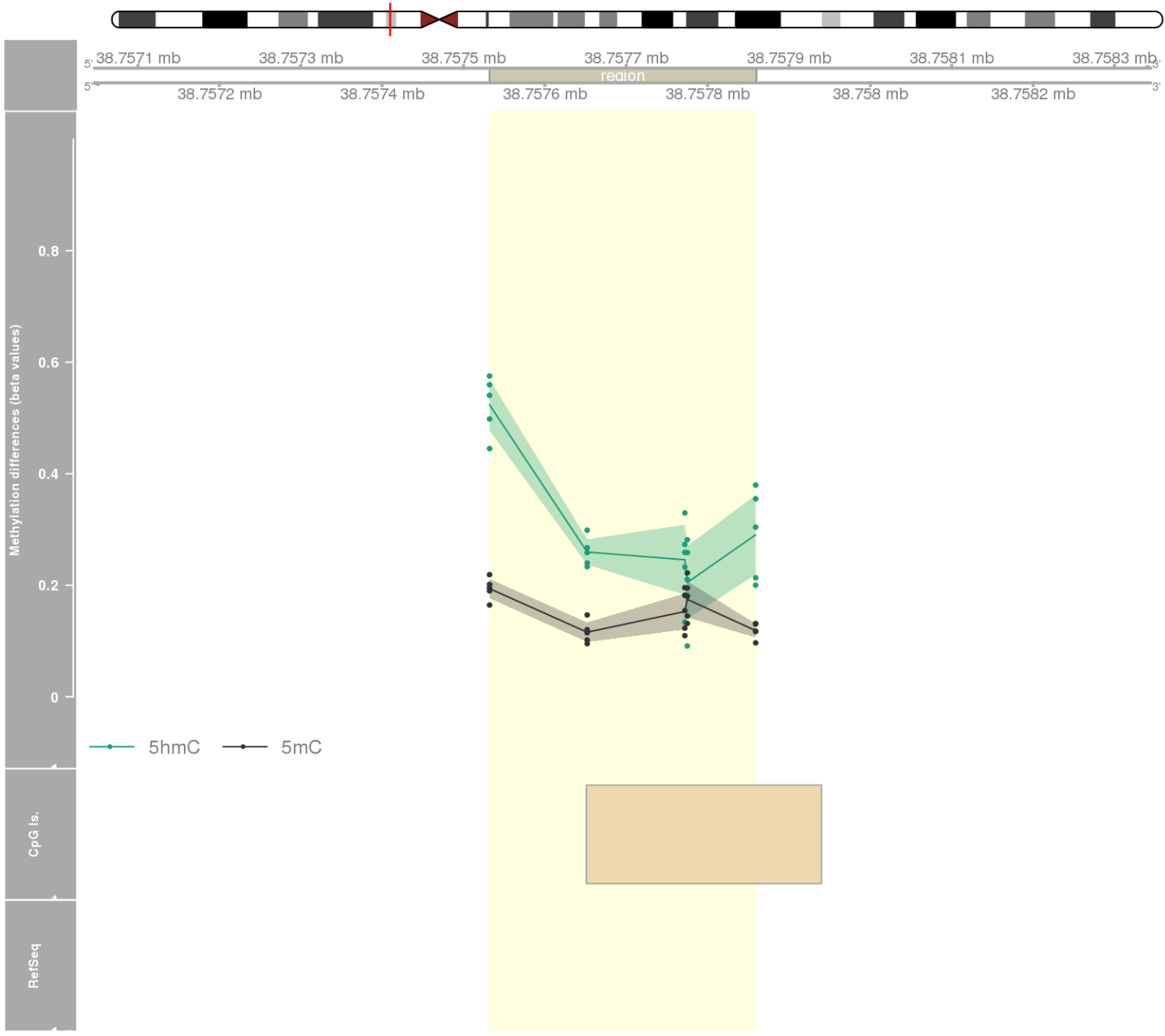
DMR 353 // chr16:88458618-88459714 // 1096 pb. (7 probes) // pvalue: 0.007 // fwer: 0.173 // pvalueArea: 0.025 // fwerArea: 1



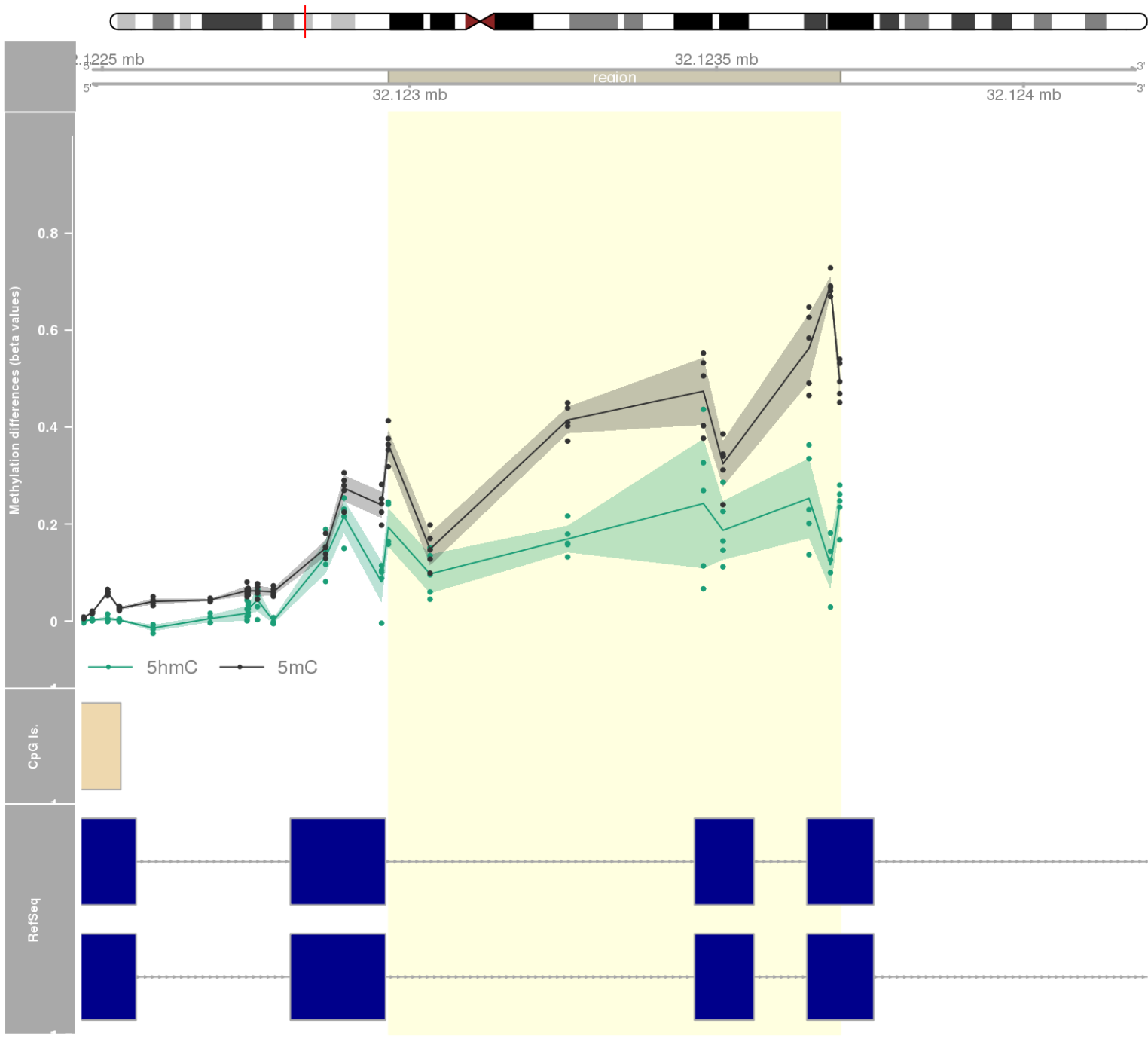
DMR 354 // chr1:977896-979589 // 1693 pb. (9 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.025 // fwerArea: 1
- genes: AGRN -



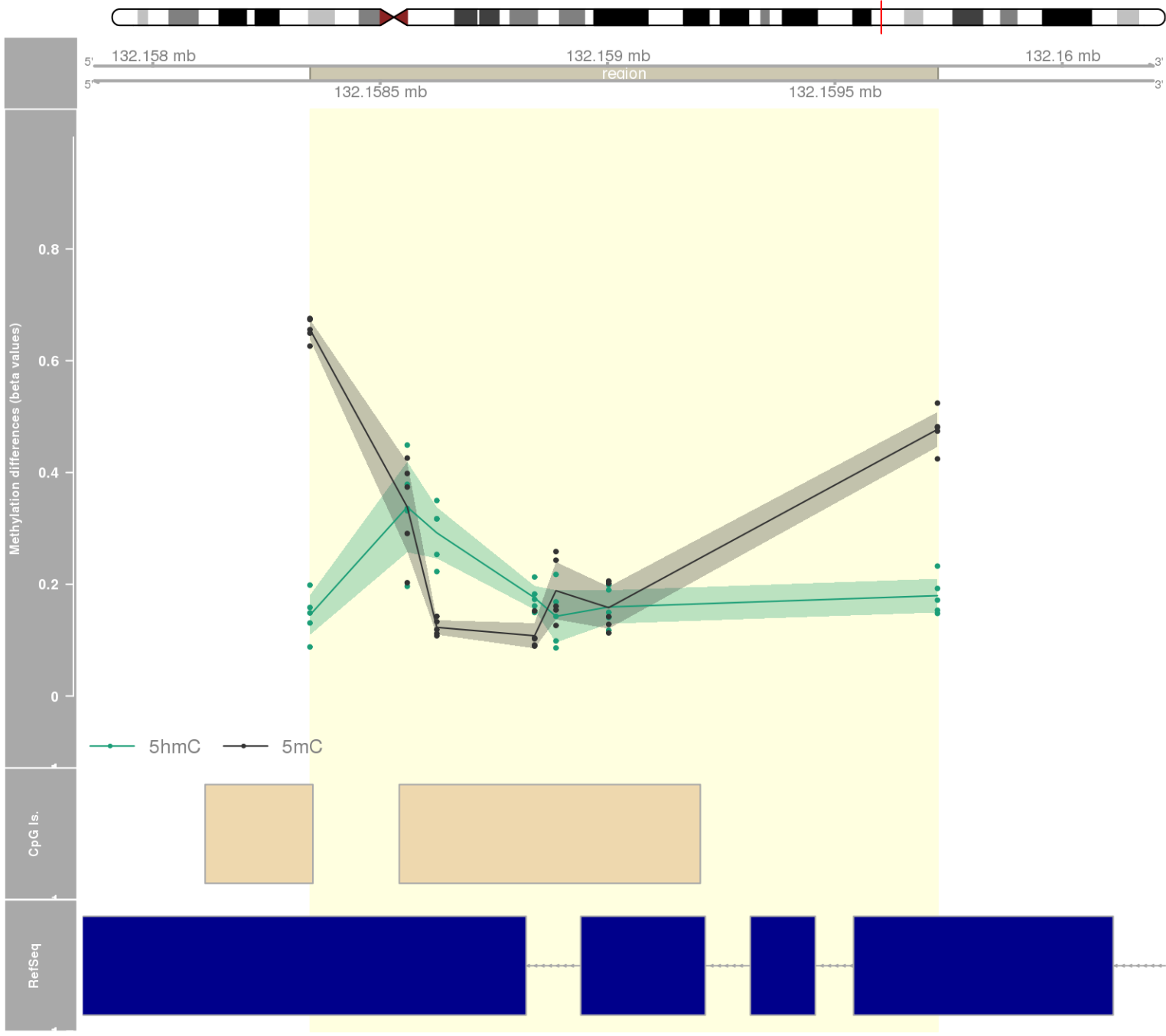
DMR 355 // chr8:38757532-38757859 // 327 pb. (5 probes) // pvalue: 0 // fwer: 0.496 // pvalueArea: 0.025 // fwerArea: 1



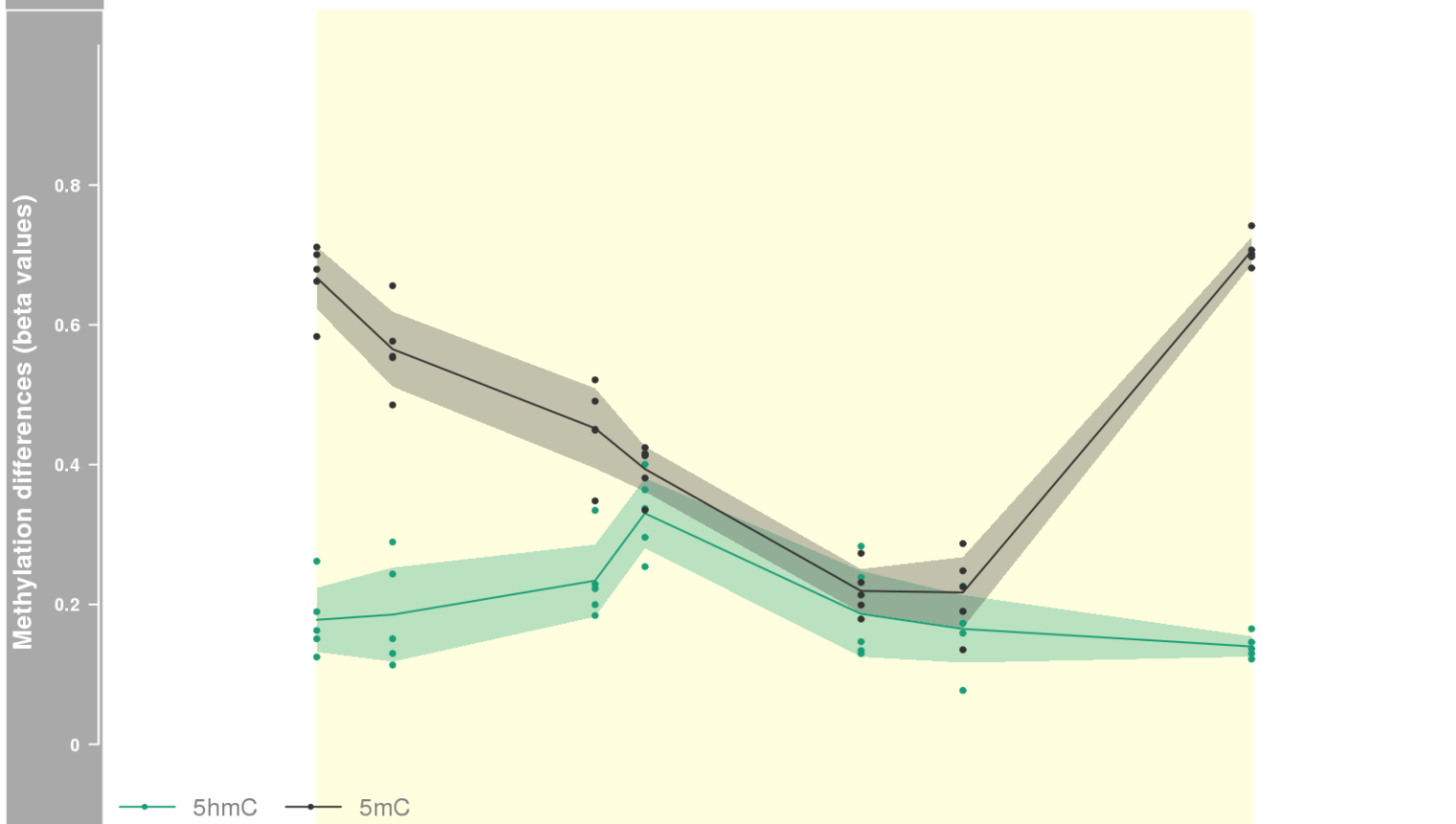
DMR 356 // chr6:32122966-32123701 // 735 pb. (8 probes) // pvalue: 0.01 // fwer: 0.522 // pvalueArea: 0.025 // fwerArea: 1
- genes: PPT2 / PPT2-EGFL8 -



DMR 357 // chr5:132158346-132159726 // 1380 pb. (7 probes) // pvalue: 0.007 // fwer: 0.173 // pvalueArea: 0.025 // fwerArea: 1
- genes: SHROOM1 -



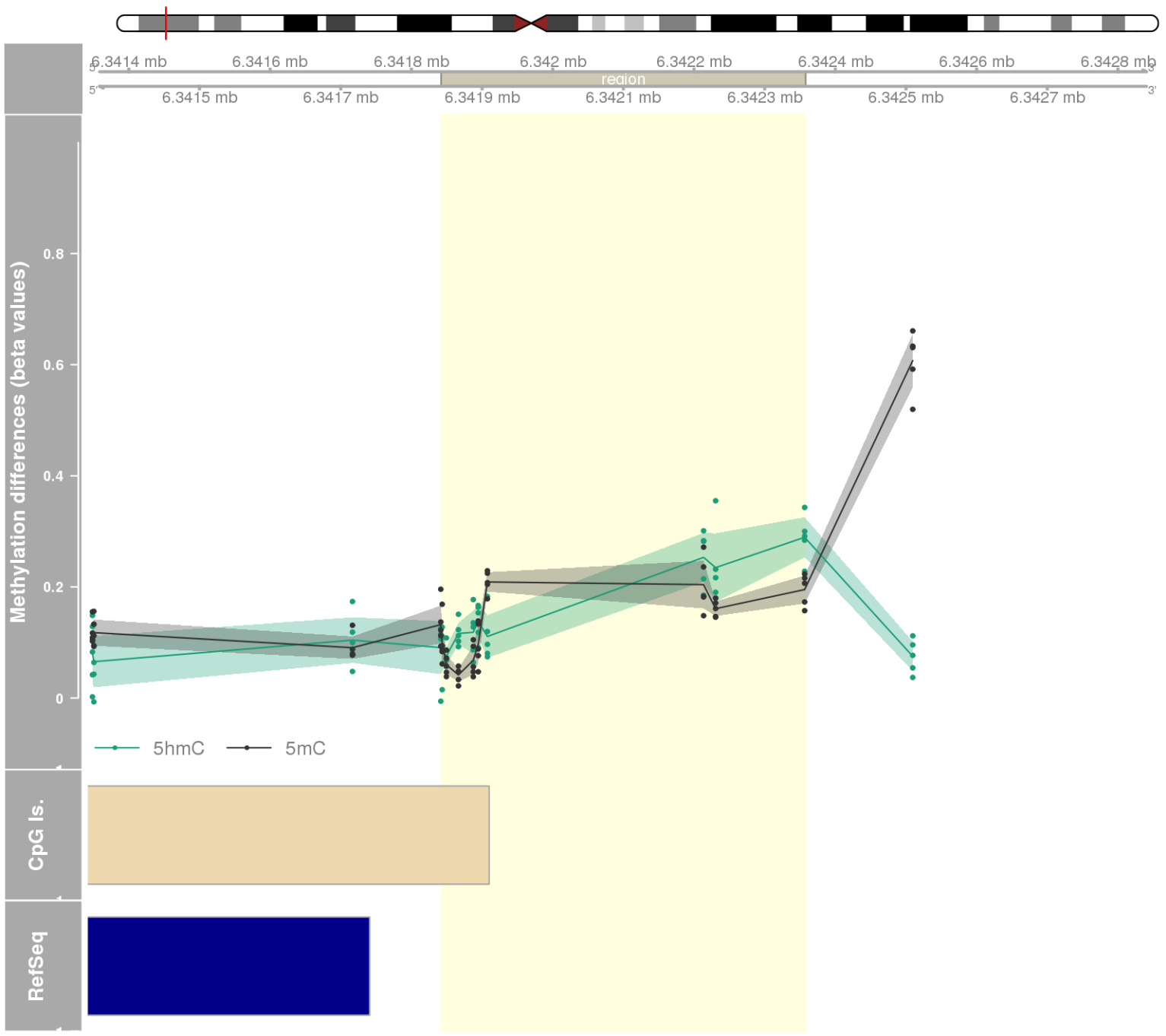
DMR 358 // chr17:3906674-3908941 // 2267 pb. (7 probes) // pvalue: 0.007 // fwer: 0.173 // pvalueArea: 0.025 // fwerArea: 1
- genes: ZZEF1 -



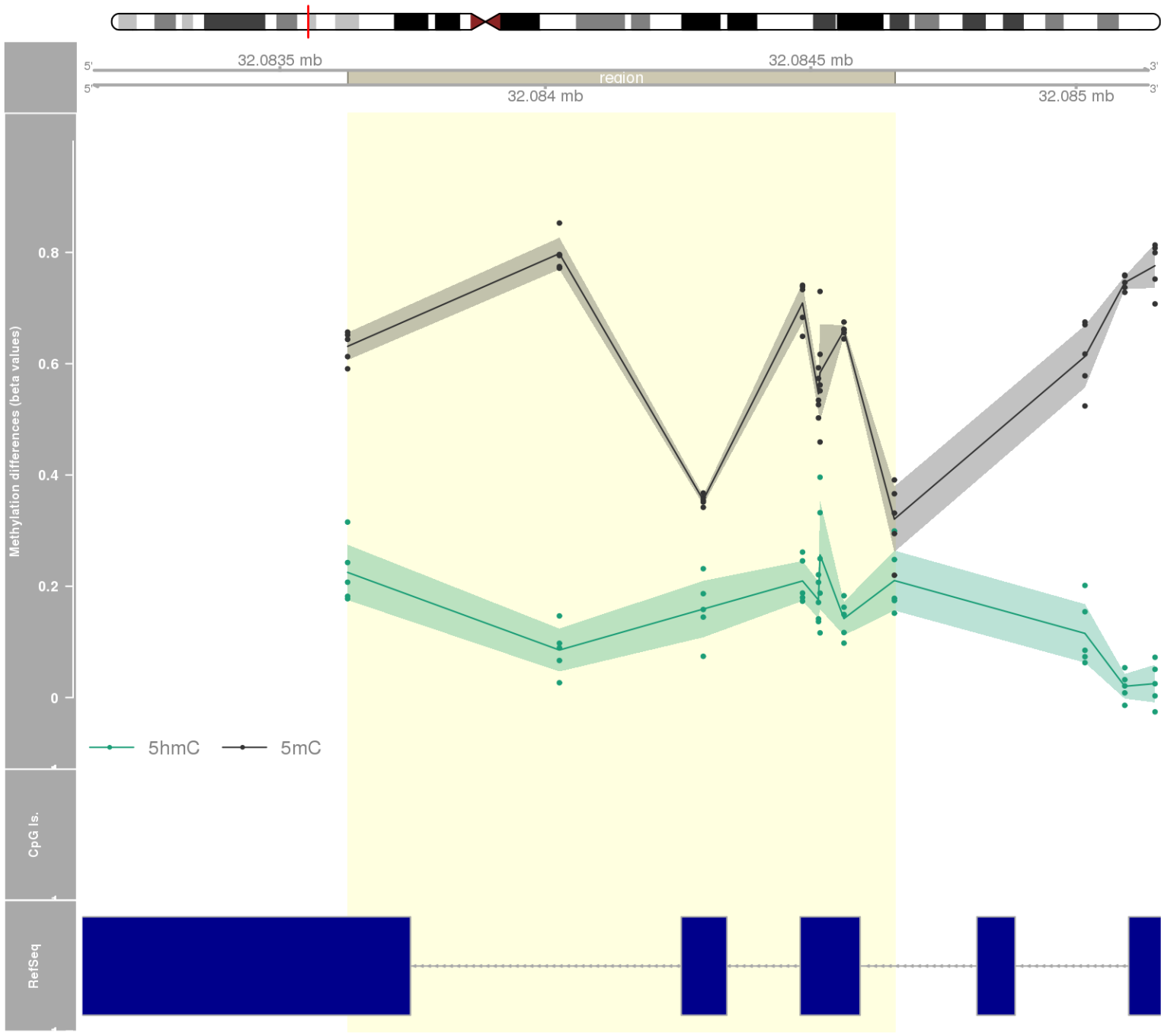
—●— 5hmC —●— 5mC



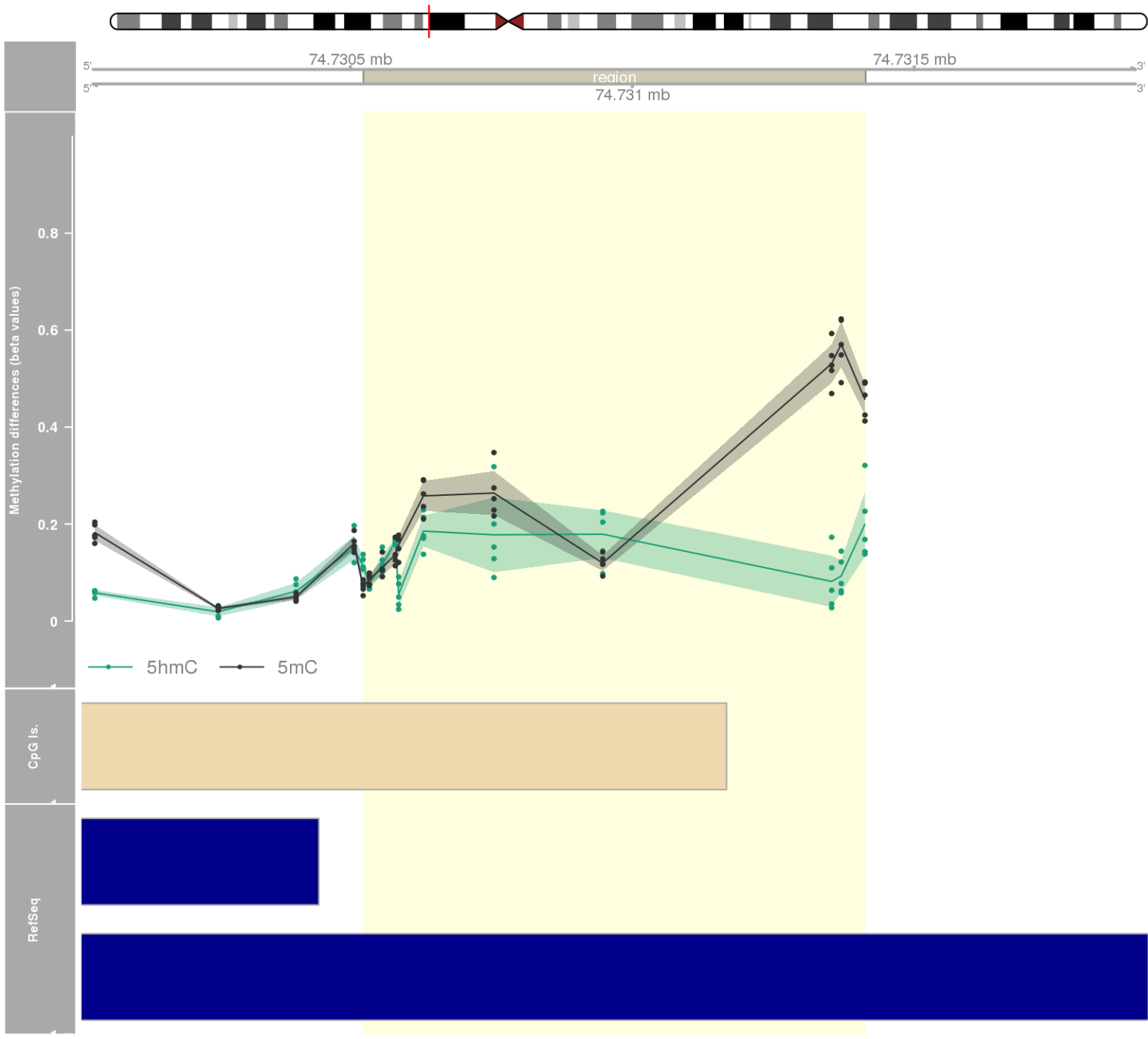
DMR 359 // chr11:6341842-6342357 // 515 pb. (10 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.025 // fwerArea: 1
- genes: PRKCDBP -



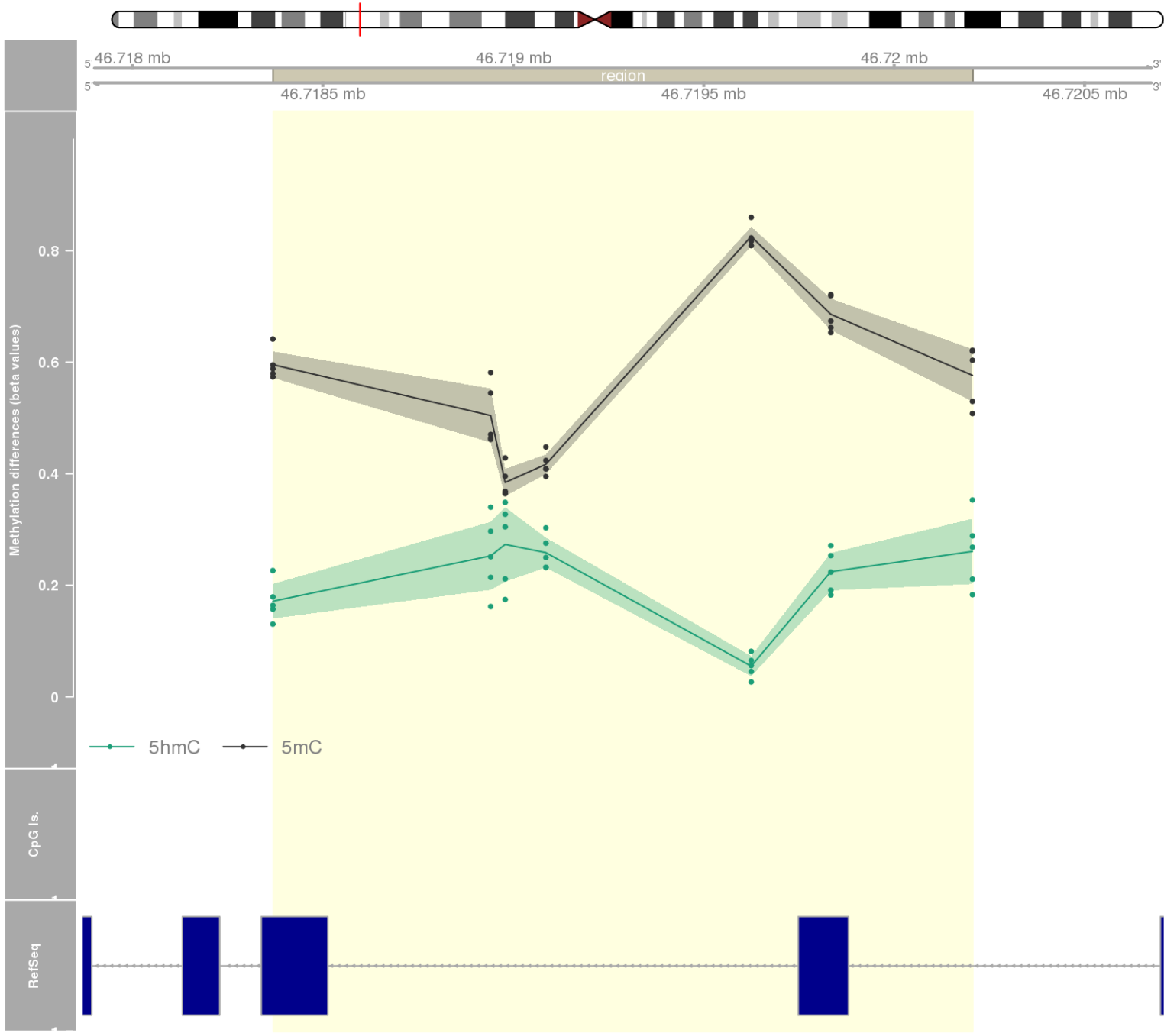
DMR 360 // chr6:32083628-32084658 // 1030 pb. (8 probes) // pvalue: 0.011 // fwer: 0.522 // pvalueArea: 0.025 // fwerArea: 1
- genes: ATF6B -



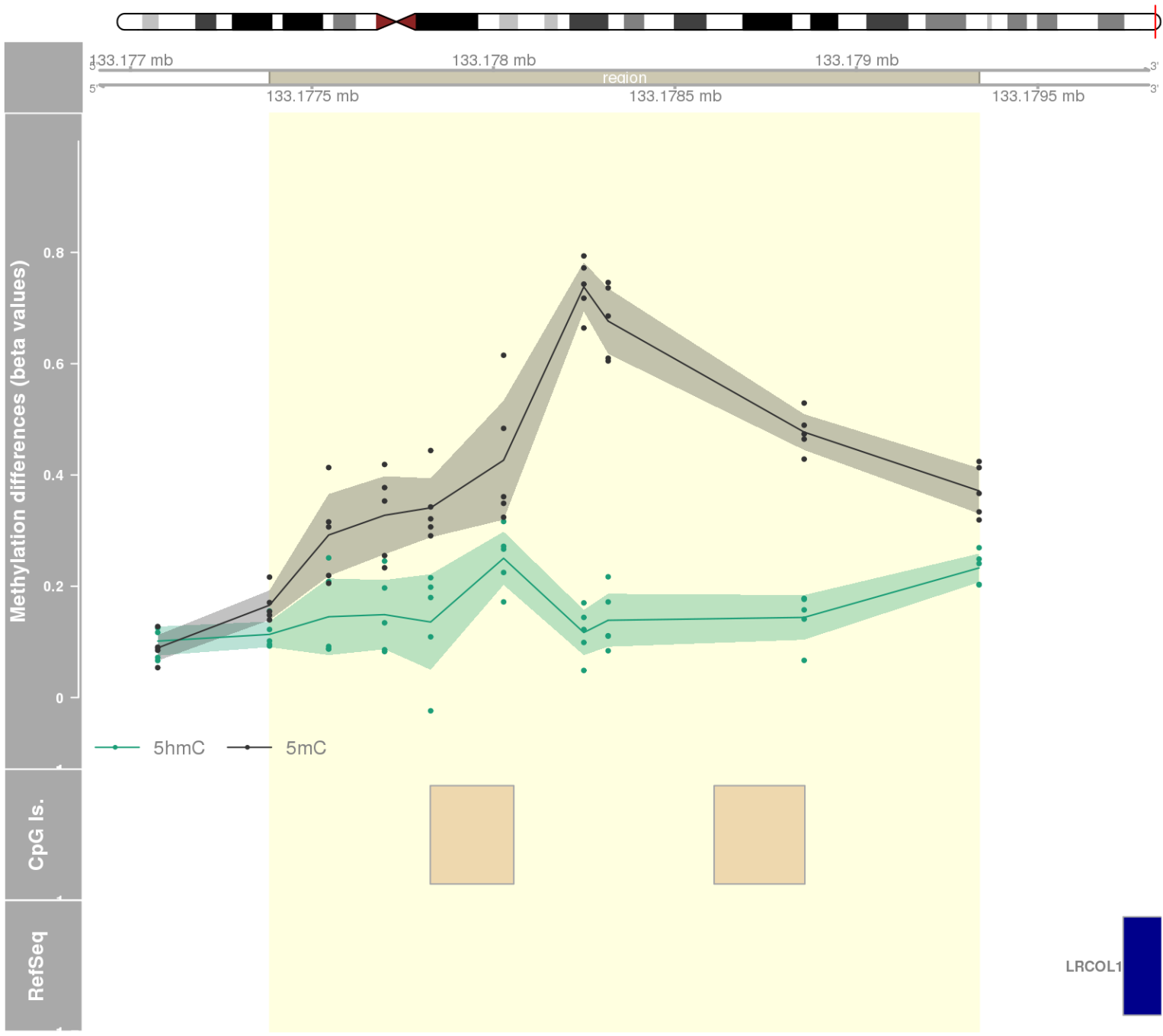
DMR 361 // chr2:74730523-74731413 // 890 pb. (11 probes) // pvalue: 0.011 // fwer: 0.36 // pvalueArea: 0.025 // fwerArea: 1
- genes: LBX2-AS1 -



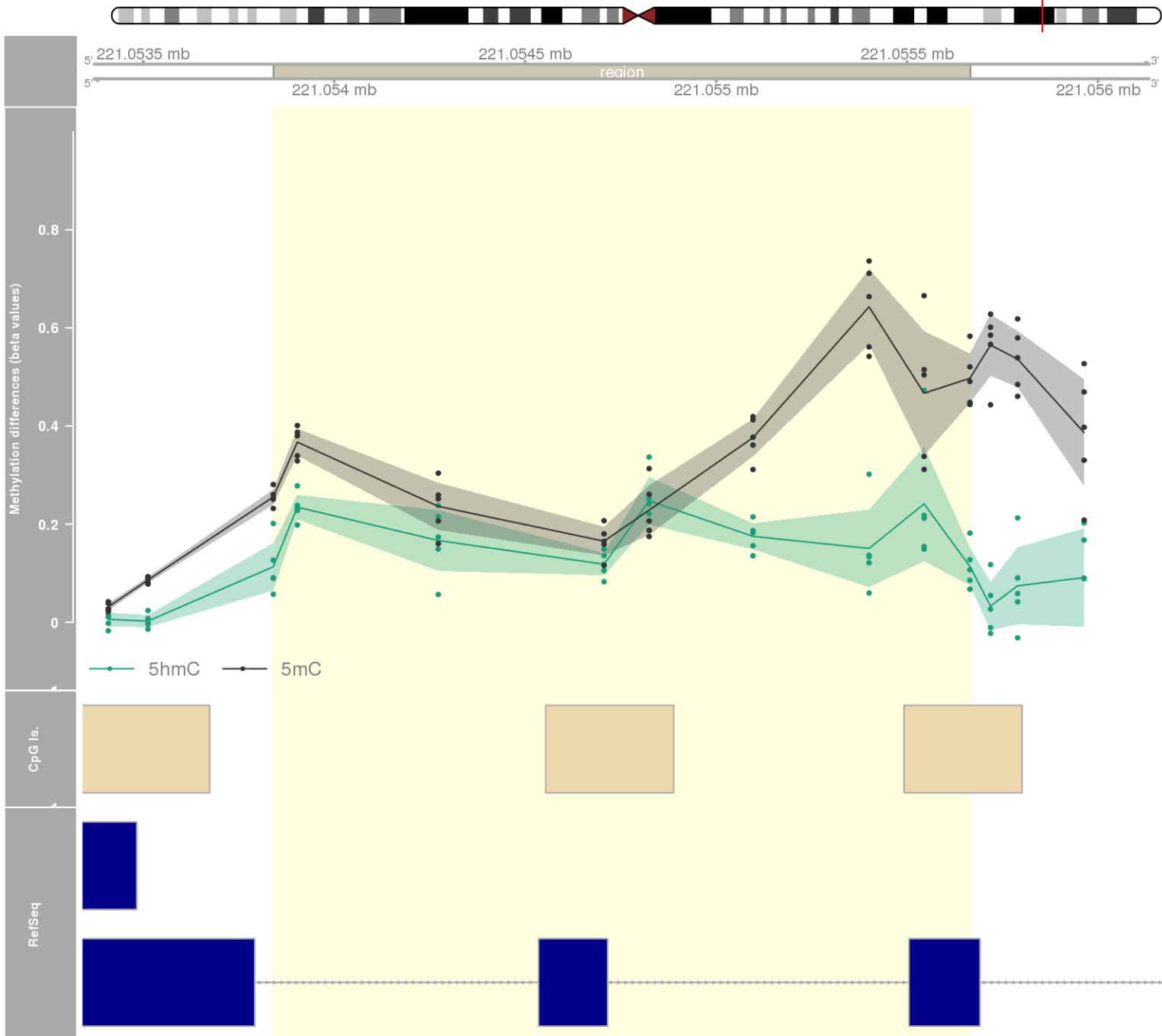
DMR 362 // chr3:46718369-46720206 // 1837 pb. (7 probes) // pvalue: 0.007 // fwer: 0.173 // pvalueArea: 0.025 // fwerArea: 1
- genes: ALS2CL -



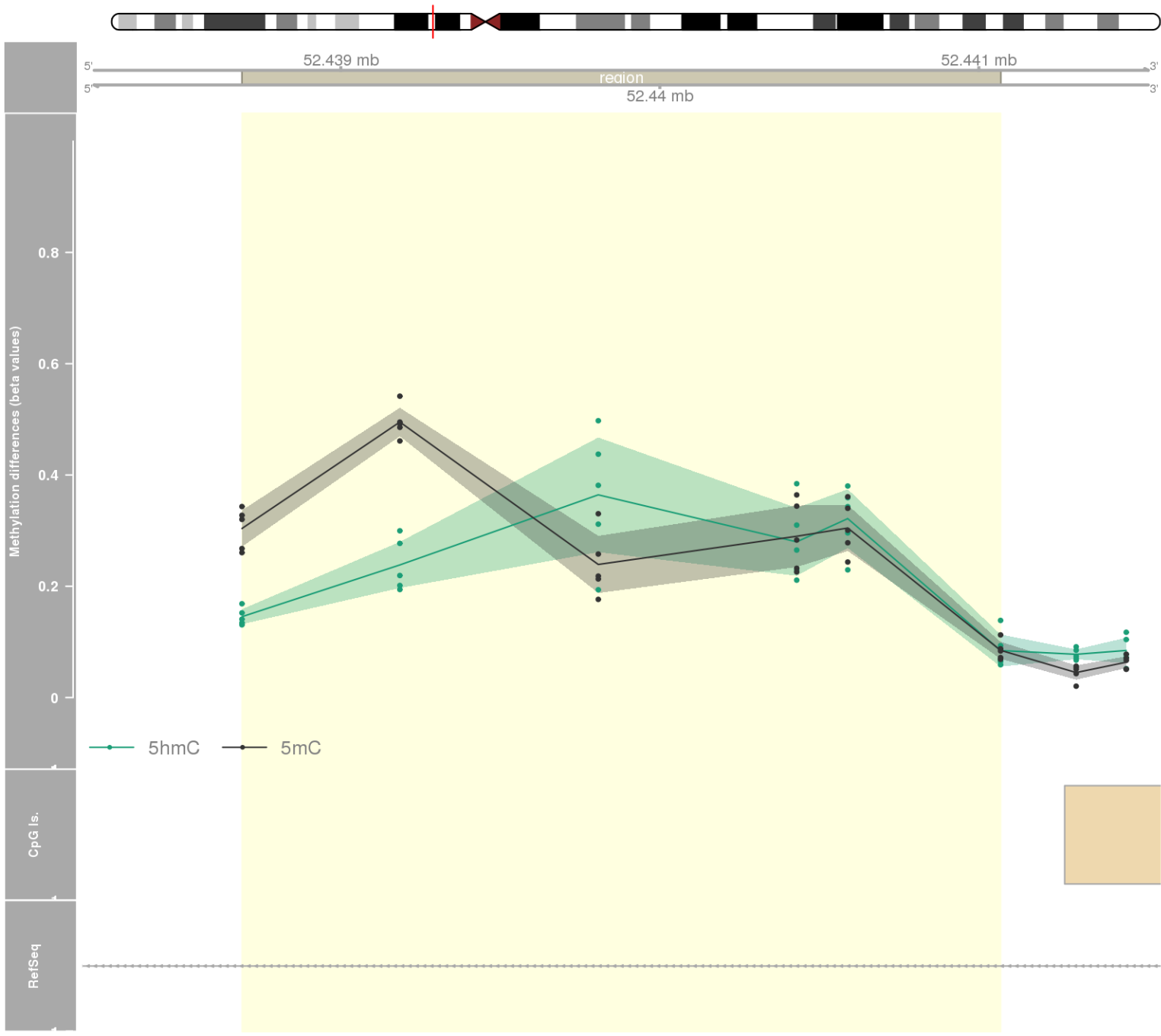
DMR 363 // chr12:133177383-133179338 // 1955 pb. (9 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.025 // fwerArea: 1



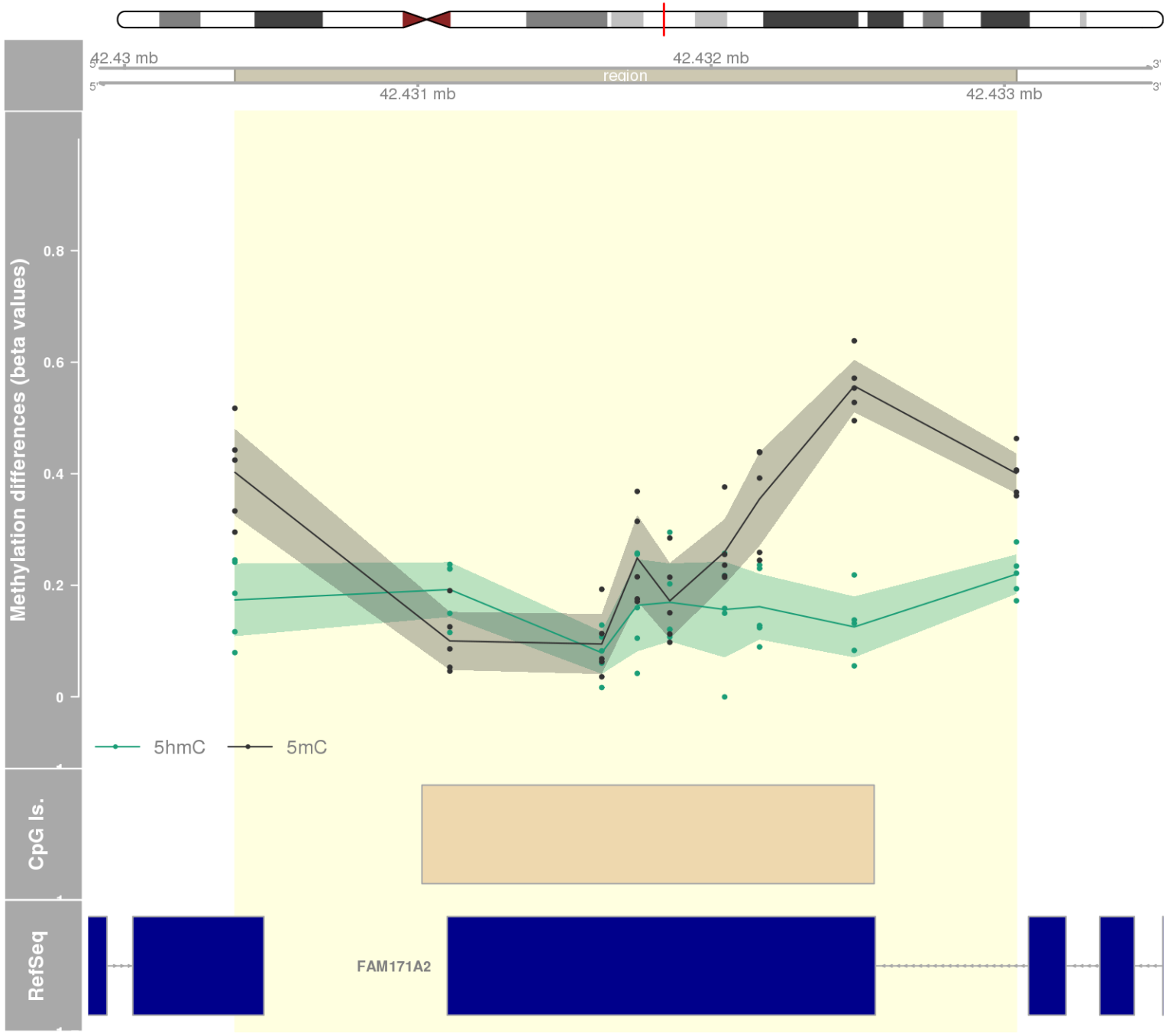
DMR 364 // chr1:221053841-221055665 // 1824 pb. (9 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.025 // fwerArea: 1
- genes: HLX -



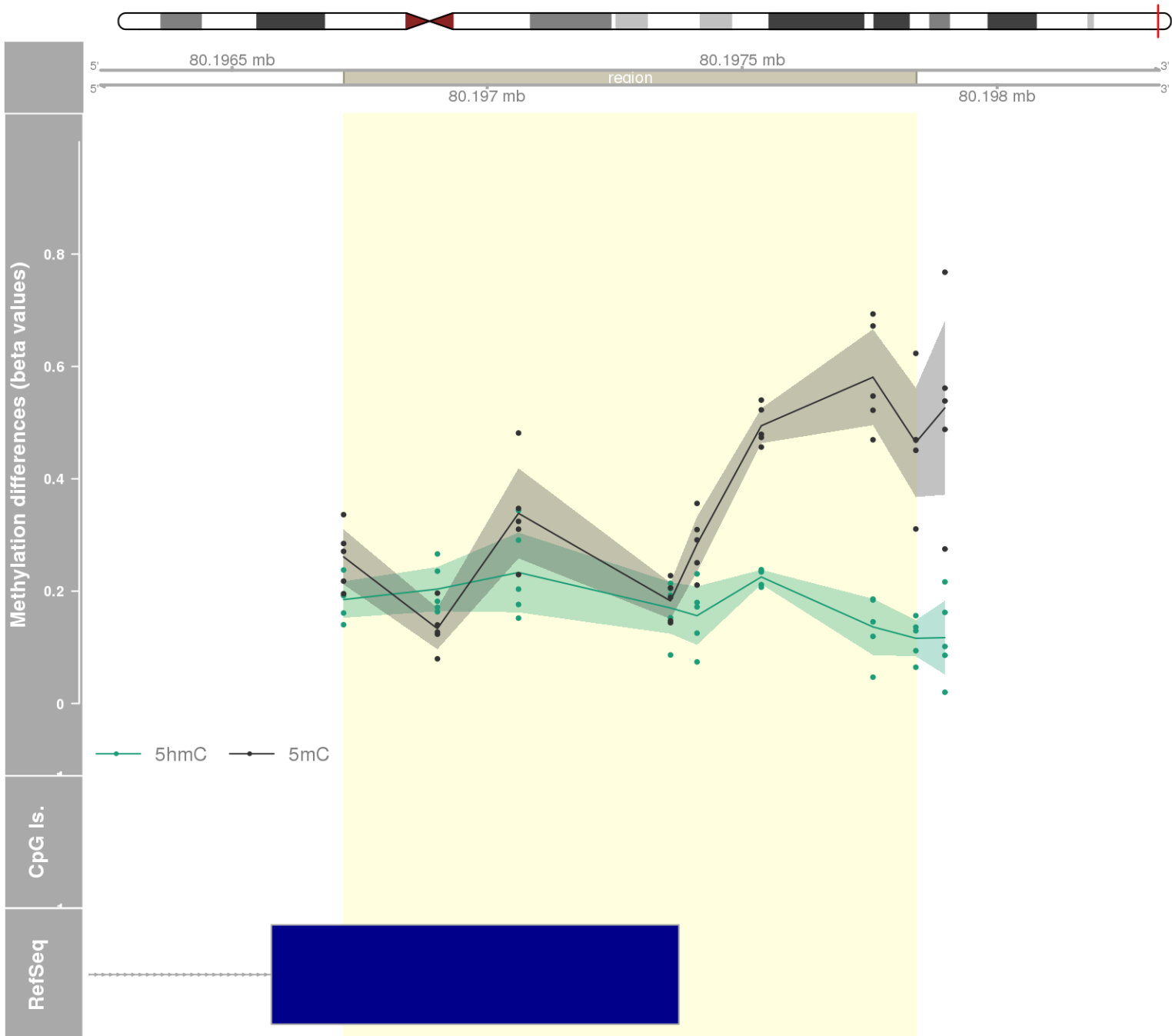
DMR 365 // chr6:52438690-52441068 // 2378 pb. (6 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.025 // fwerArea: 1
- genes: TRAM2 -



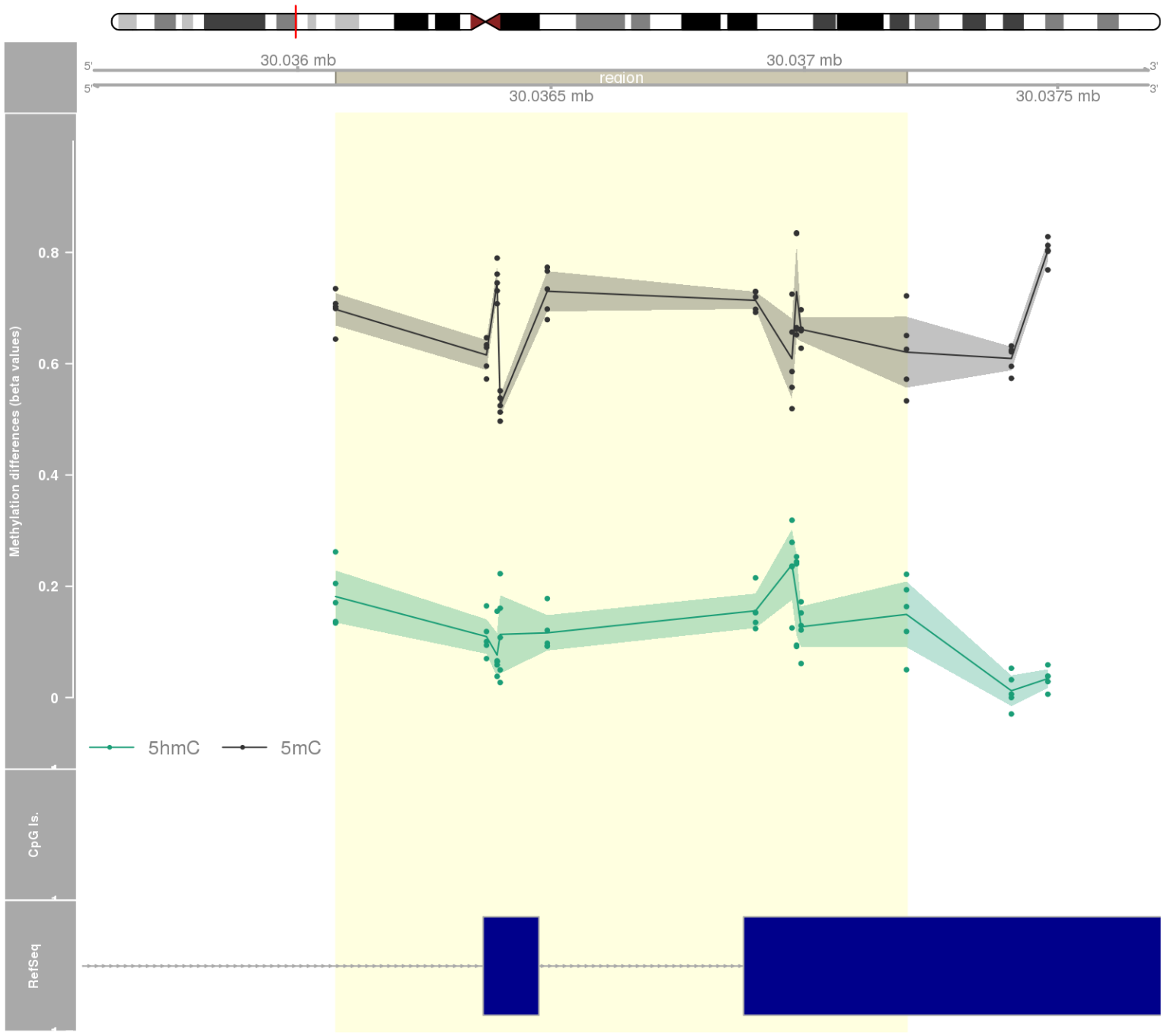
DMR 366 // chr17:42430376-42433041 // 2665 pb. (9 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.026 // fwerArea: 1
- genes: GRN / FAM171A2 -



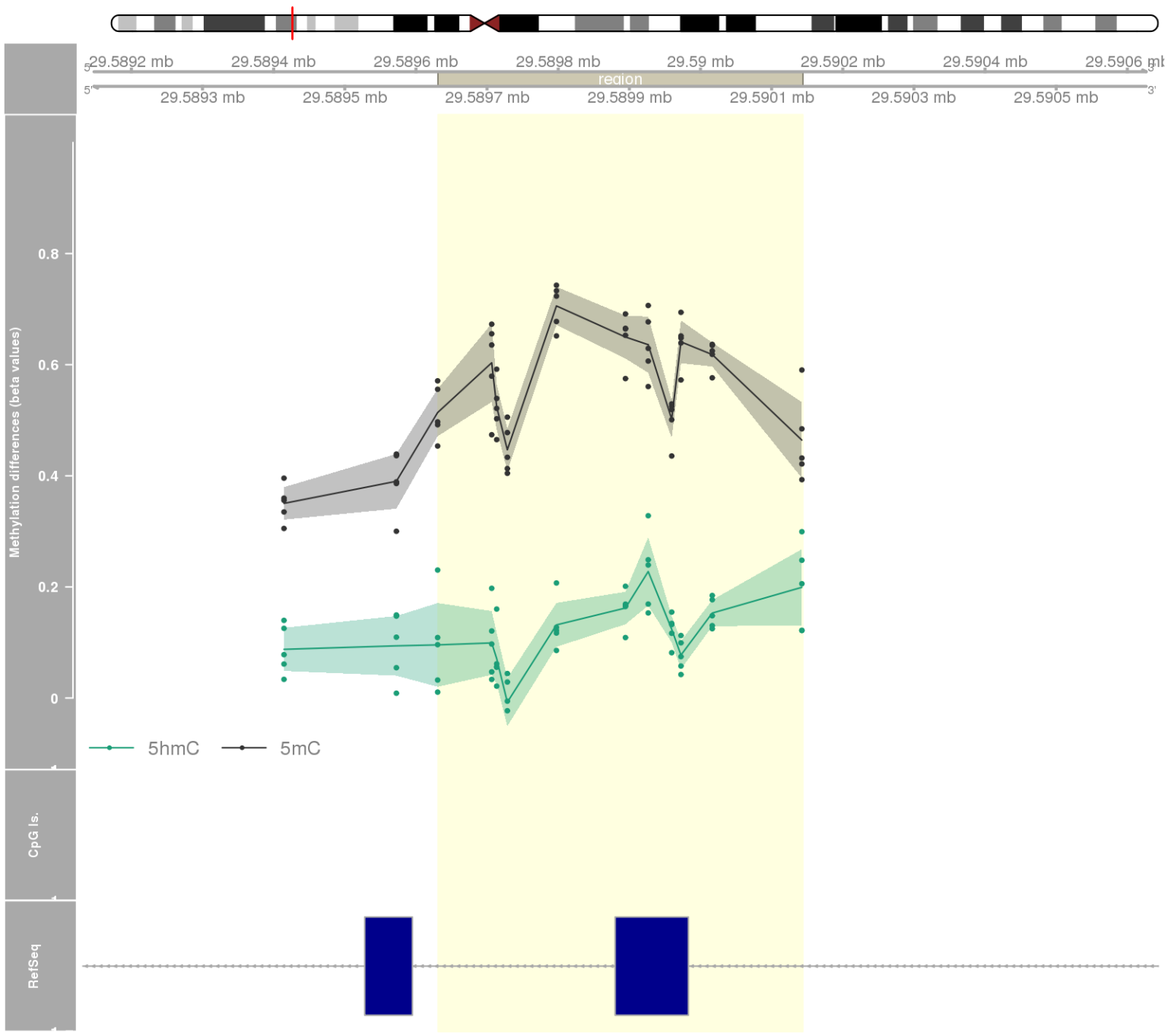
DMR 367 // chr17:80196719-80197841 // 1122 pb. (8 probes) // pvalue: 0.011 // fwer: 0.522 // pvalueArea: 0.026 // fwerArea: 1
- genes: SLC16A3 / CSNK1D -



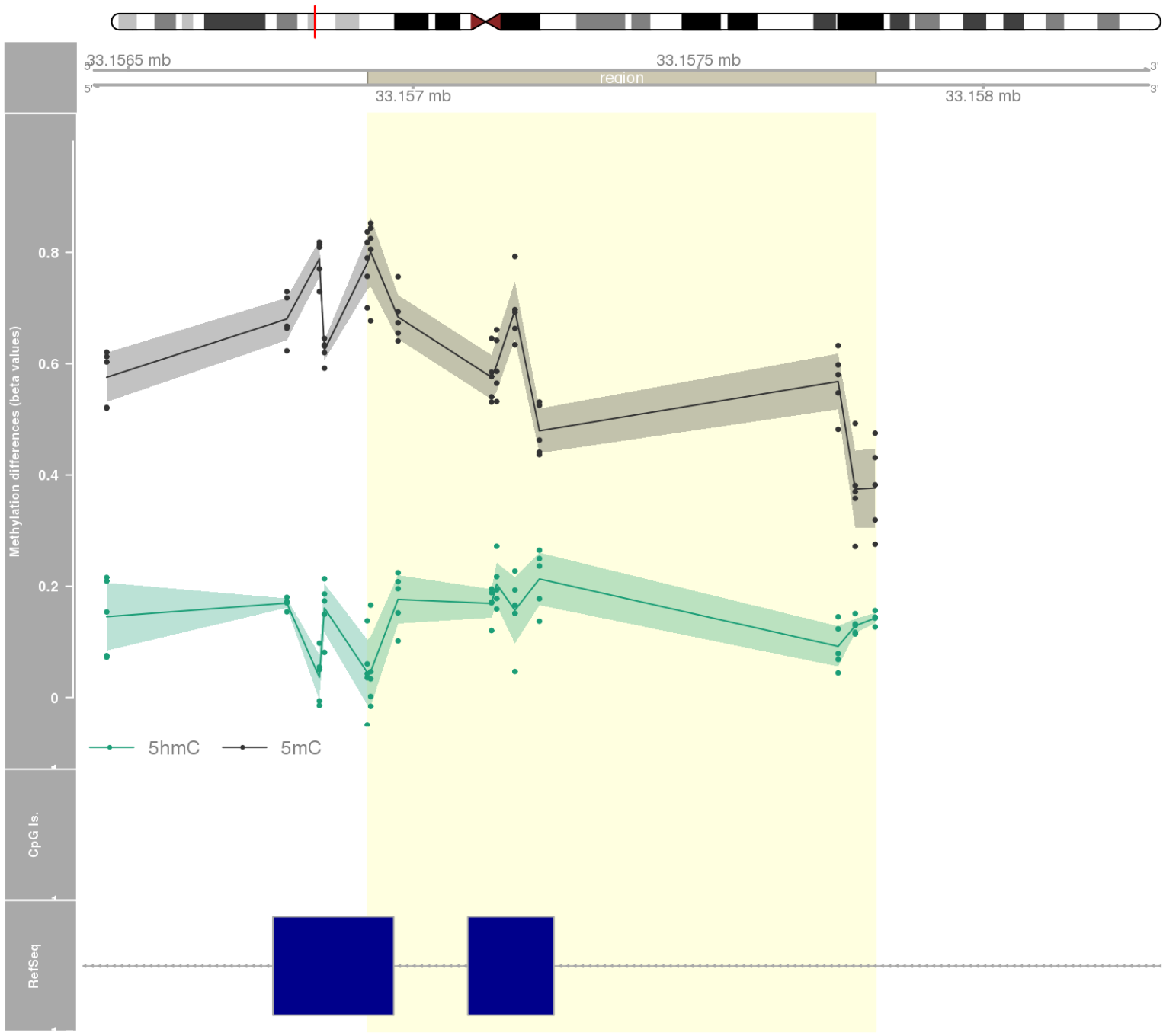
DMR 368 // chr6:30036075-30037202 // 1127 pb. (10 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.026 // fwerArea: 1
- genes: PPP1R11 -



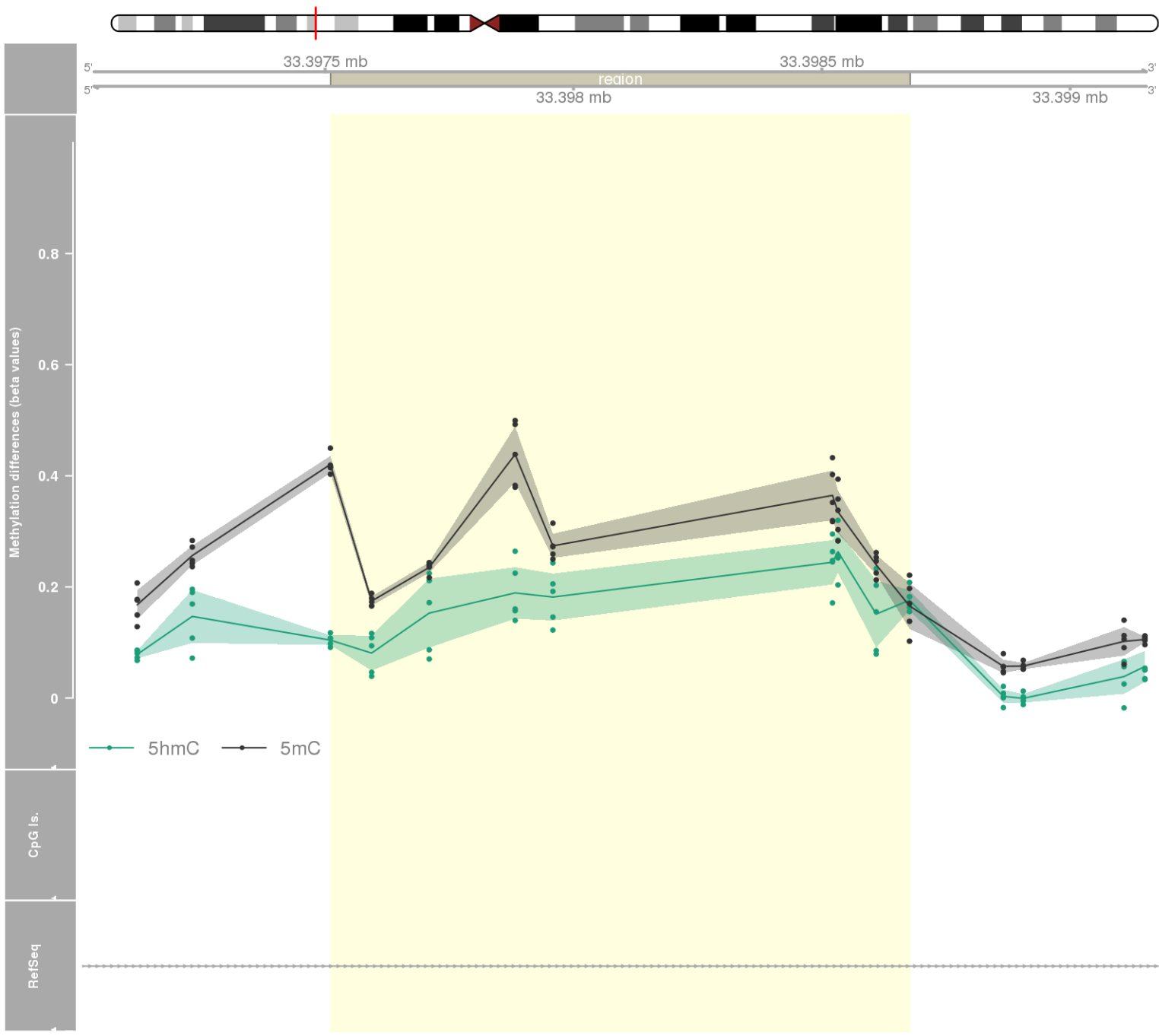
DMR 369 // chr6:29589631-29590143 // 512 pb. (11 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.026 // fwerArea: 1
- genes: GABBR1 -



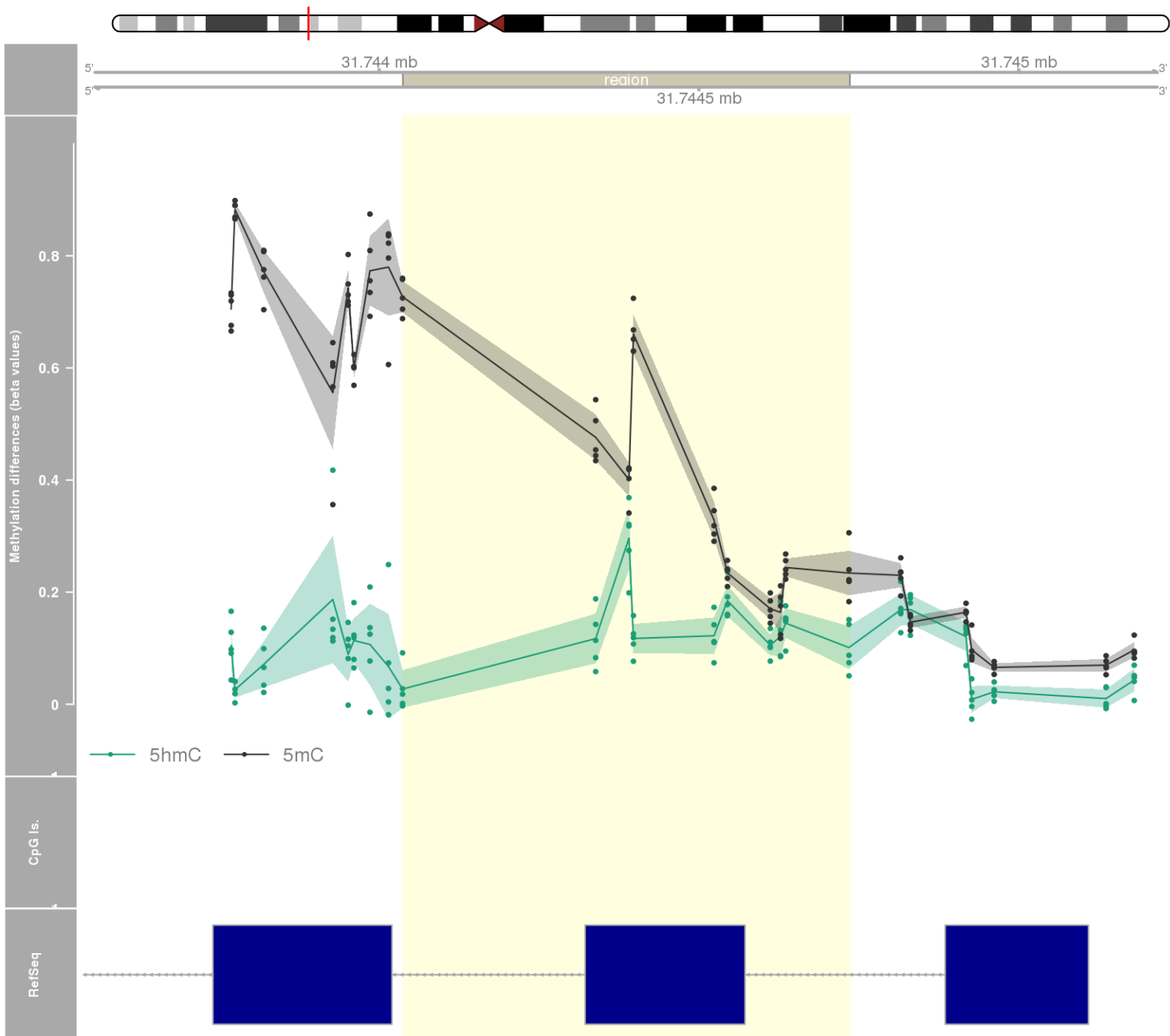
DMR 370 // chr6:33156920-33157811 // 891 pb. (10 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.026 // fwerArea: 1
- genes: COL11A2 -



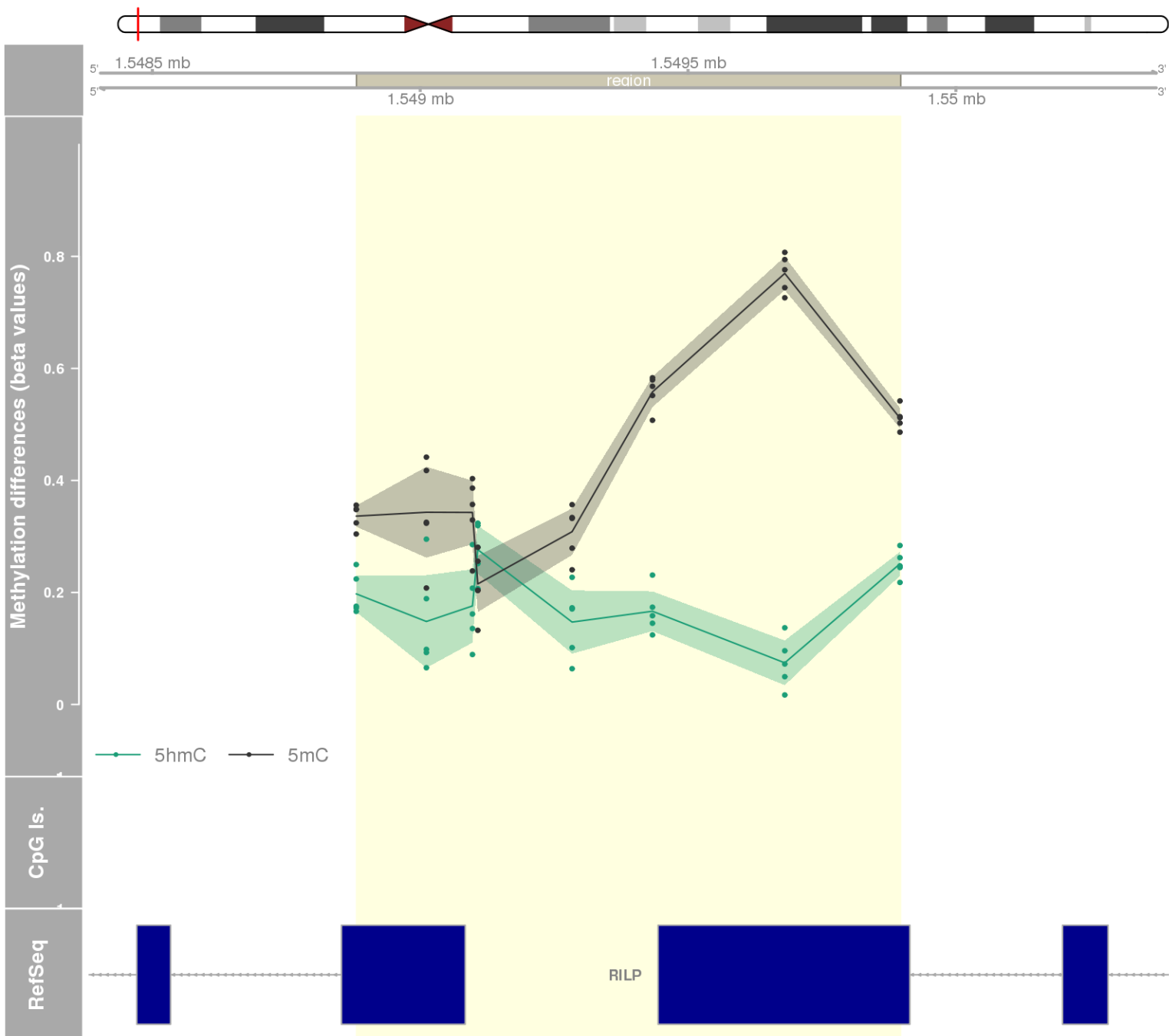
DMR 371 // chr6:33397511-33398677 // 1166 pb. (9 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.026 // fwerArea: 1
- genes: SYNGAP1 -



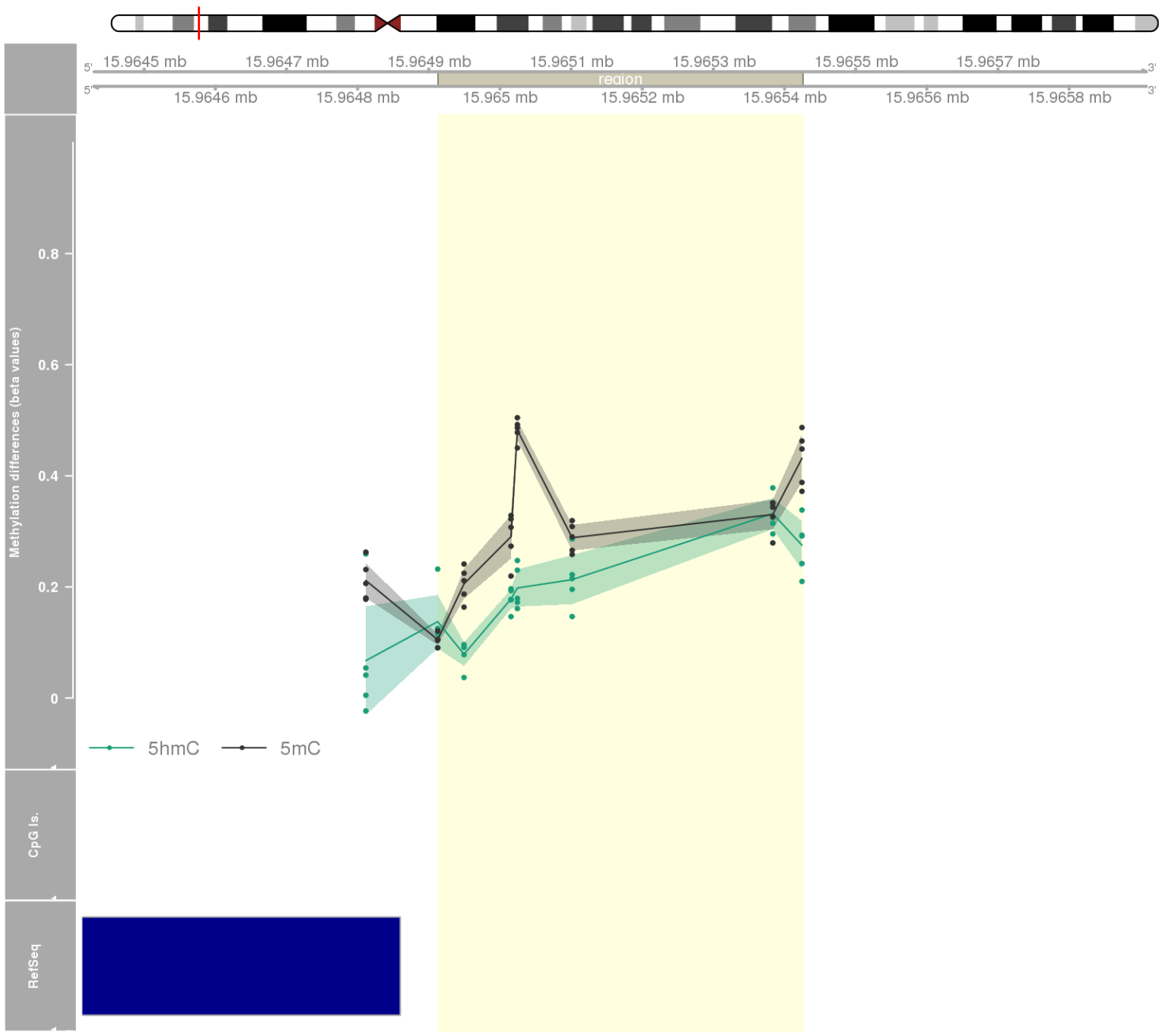
DMR 372 // chr6:31744037-31744735 // 698 pb. (10 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.026 // fwerArea: 1
- genes: VWA7 -



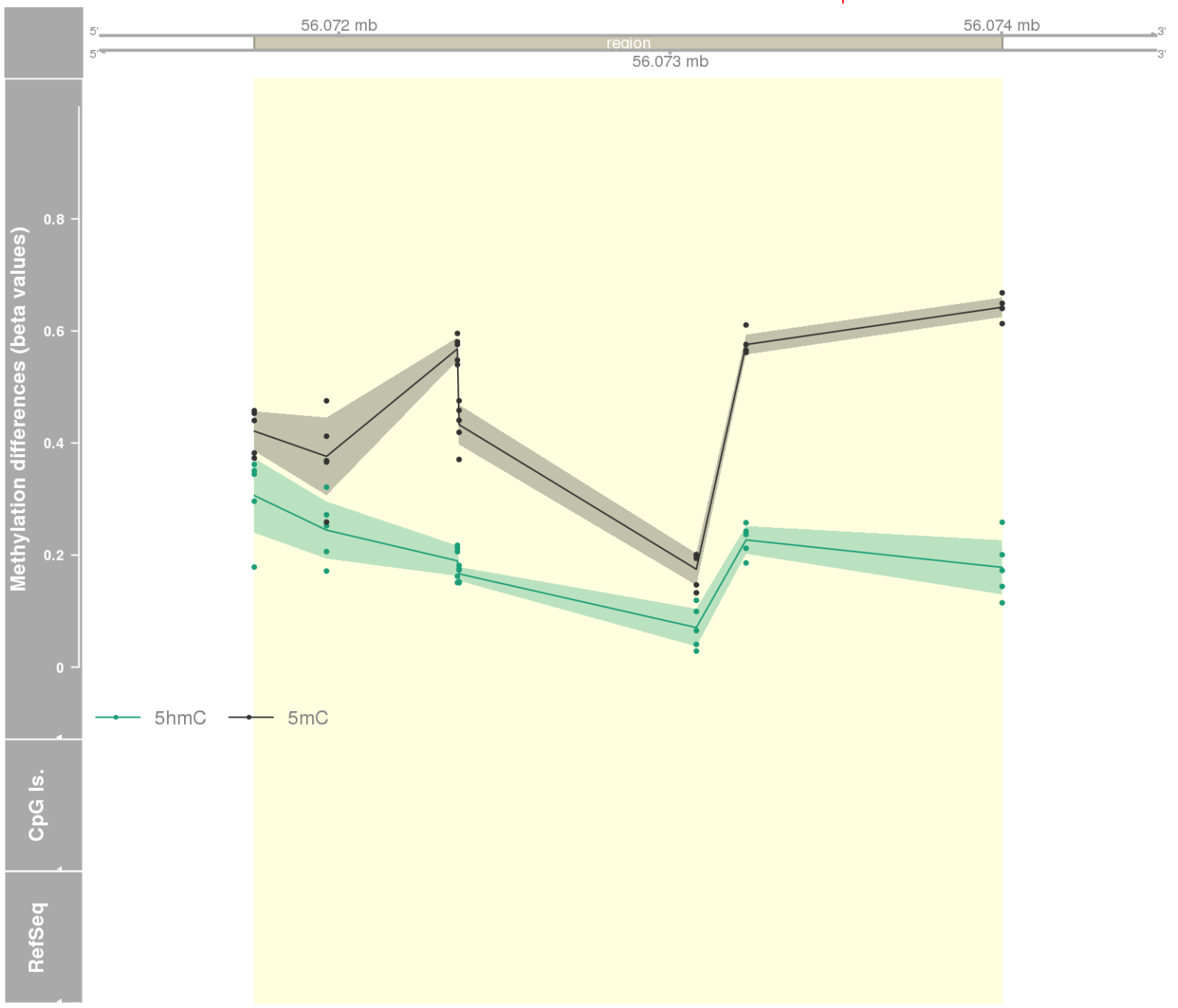
DMR 373 // chr17:1548881-1549896 // 1015 pb. (8 probes) // pvalue: 0.011 // fwer: 0.522 // pvalueArea: 0.026 // fwerArea: 1
- genes: SCARF1 / RILP -



DMR 374 // chr4:15964913-15965425 // 512 pb. (7 probes) // pvalue: 0.008 // fwer: 0.173 // pvalueArea: 0.026 // fwerArea: 1
- genes: FGFBP2 / PROM1 -



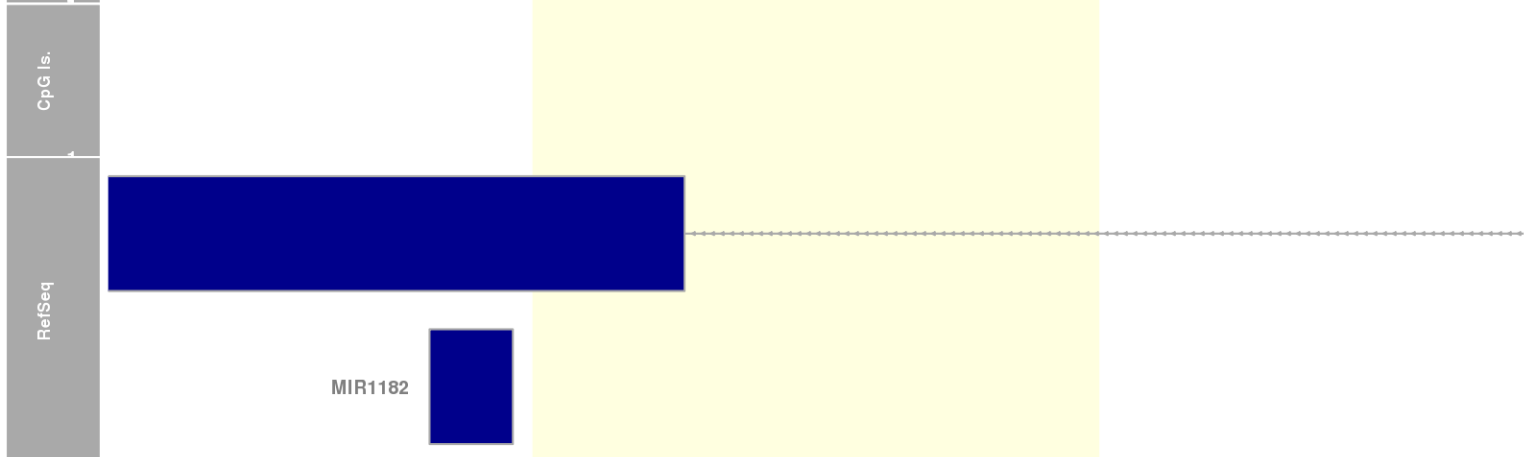
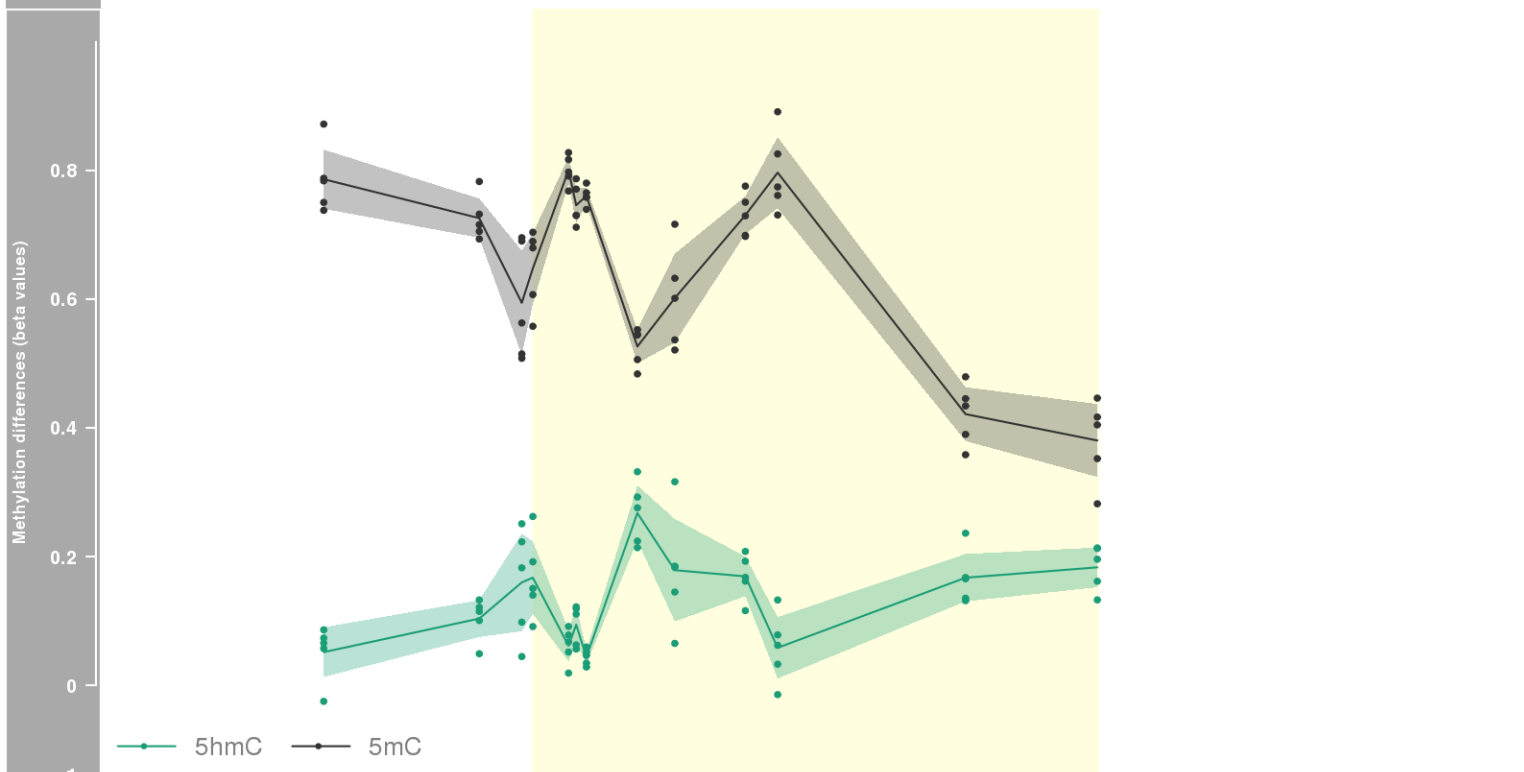
DMR 375 // chr17:56071745-56074002 // 2257 pb. (7 probes) // pvalue: 0.008 // fwer: 0.173 // pvalueArea: 0.026 // fwerArea: 1



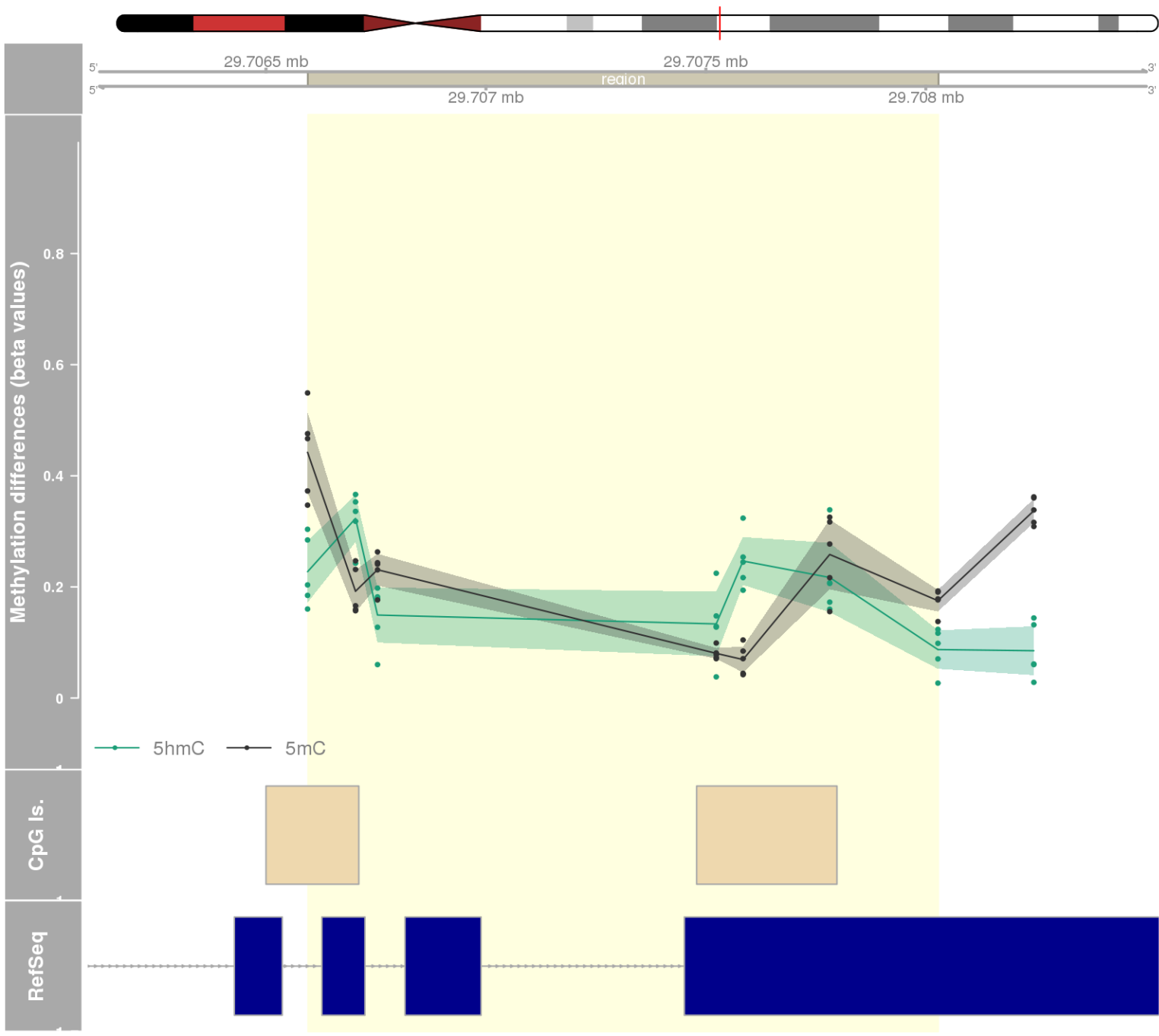
CpG Is.

RefSeq

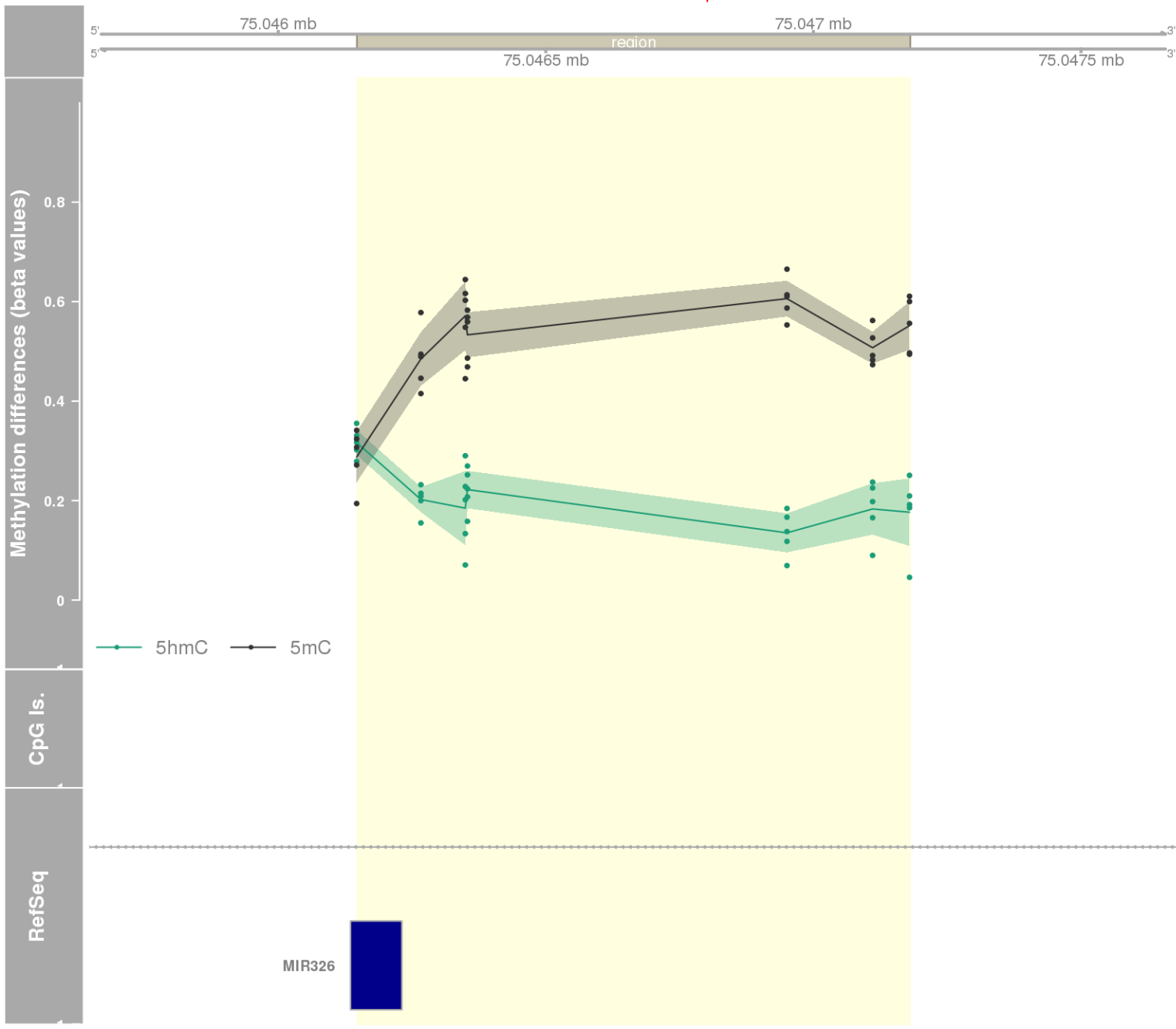
DMR 376 // chr1:231155695-231156359 // 664 pb. (10 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.026 // fwerArea: 1
- genes: FAM89A -



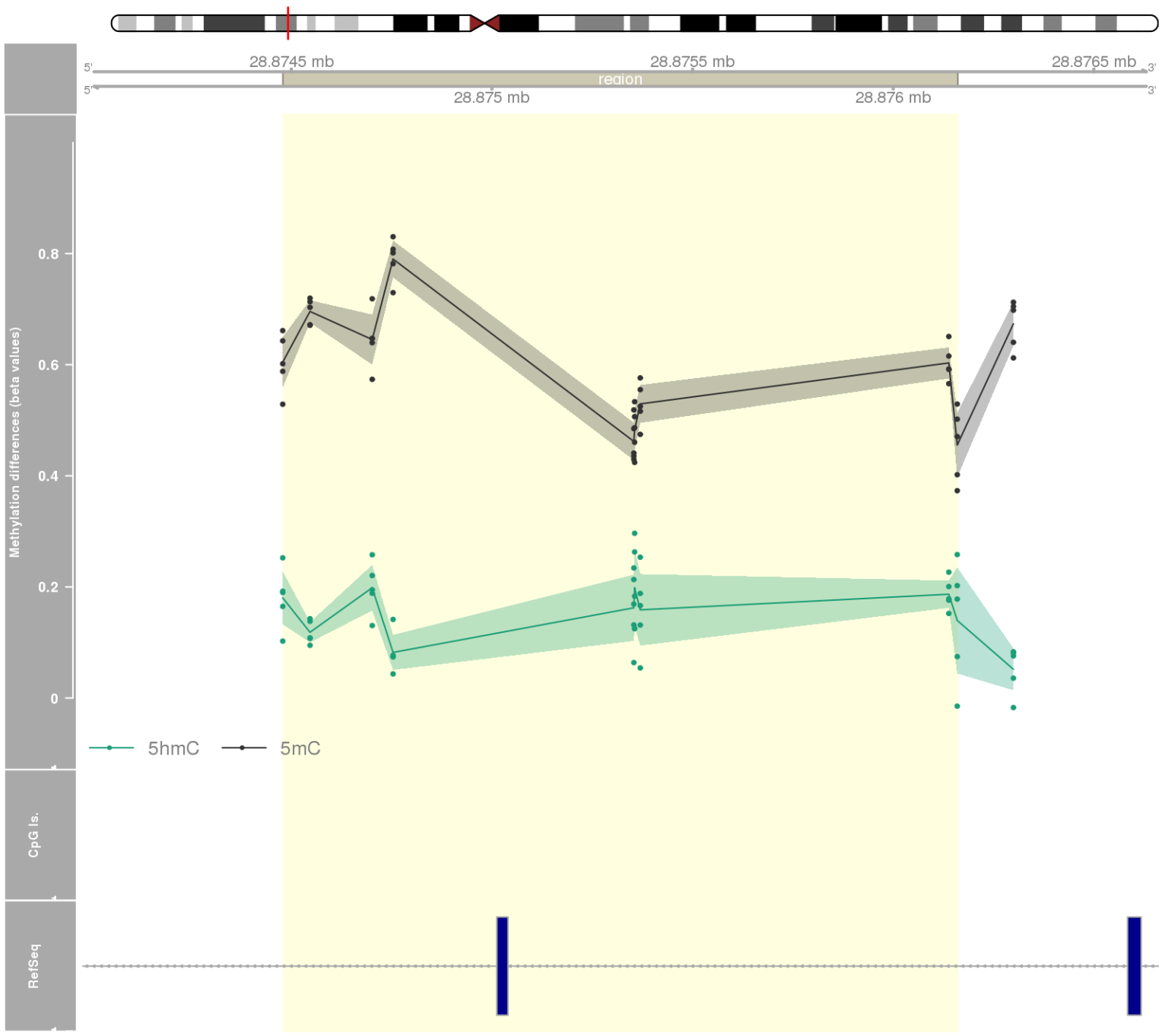
DMR 377 // chr22:29706595-29708028 // 1433 pb. (7 probes) // pvalue: 0.008 // fwer: 0.173 // pvalueArea: 0.026 // fwerArea: 1
- genes: GAS2L1 -



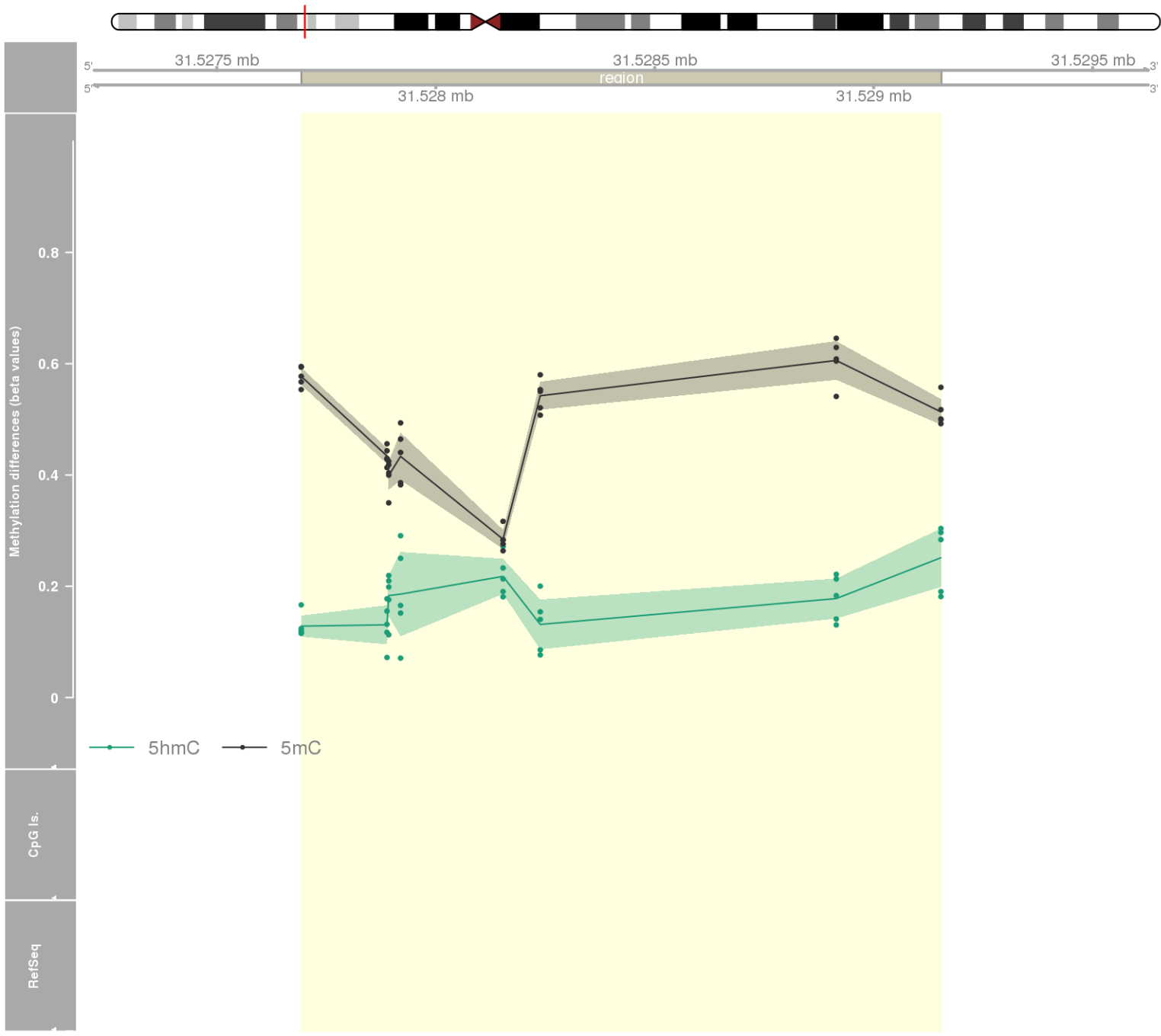
DMR 378 // chr11:75046147-75047180 // 1033 pb. (7 probes) // pvalue: 0.008 // fwer: 0.173 // pvalueArea: 0.026 // fwerArea: 1
- genes: MIR326 / ARRB1 -



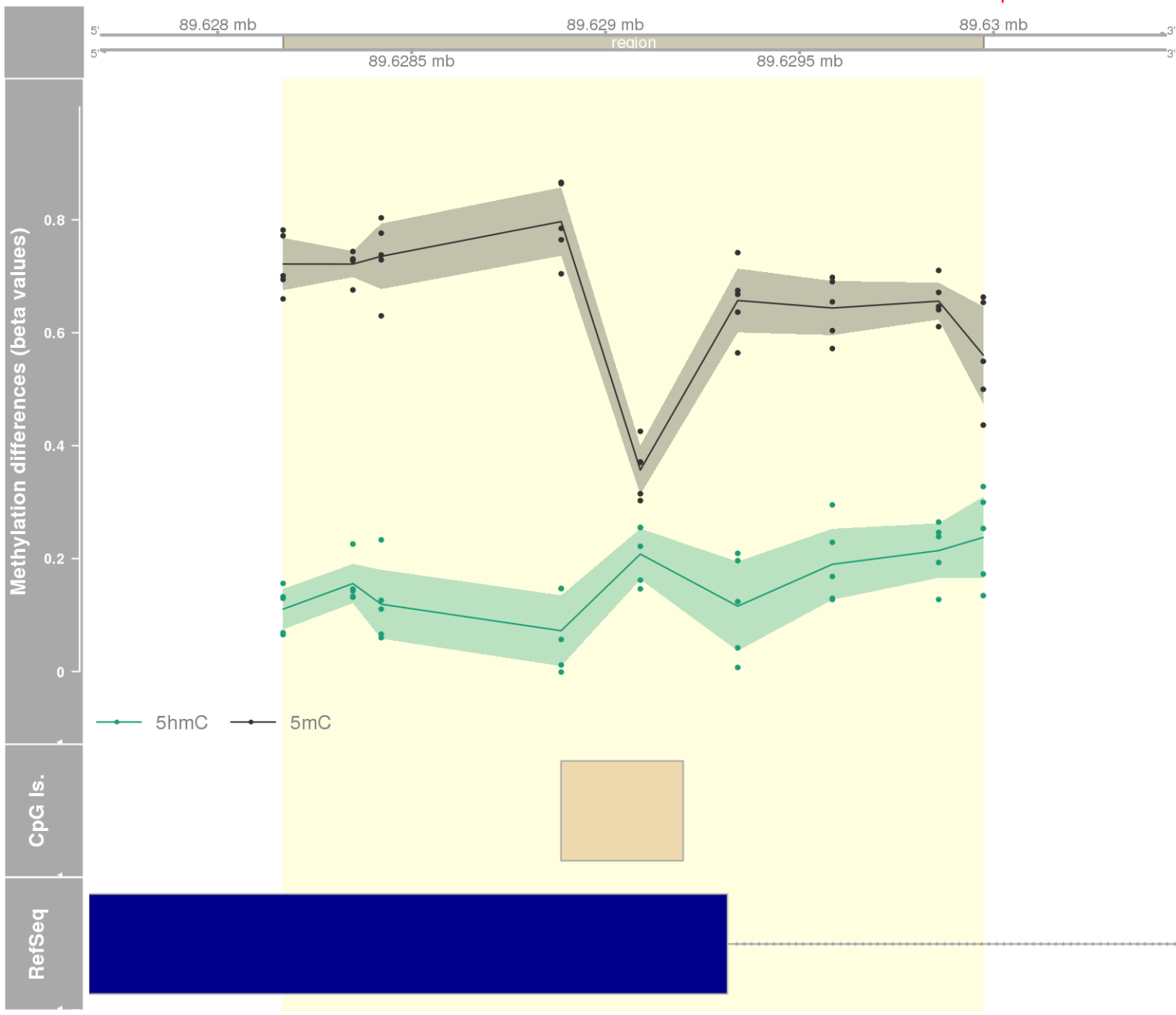
DMR 379 // chr6:28874479-28876160 // 1681 pb. (9 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.026 // fwerArea: 1
- genes: TRIM27 -



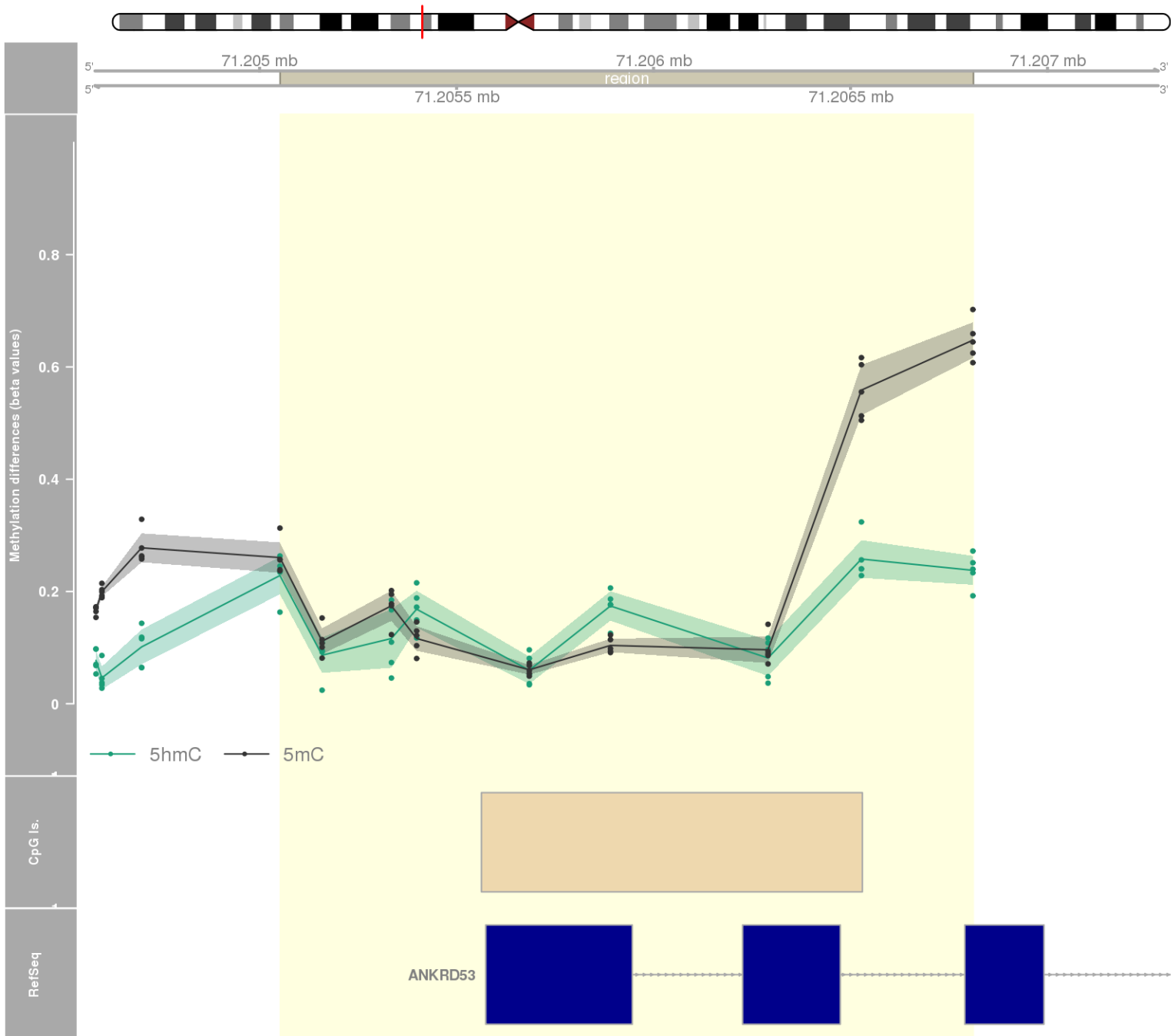
DMR 380 // chr6:31527693-31529154 // 1461 pb. (8 probes) // pvalue: 0.012 // fwer: 0.522 // pvalueArea: 0.026 // fwerArea: 1



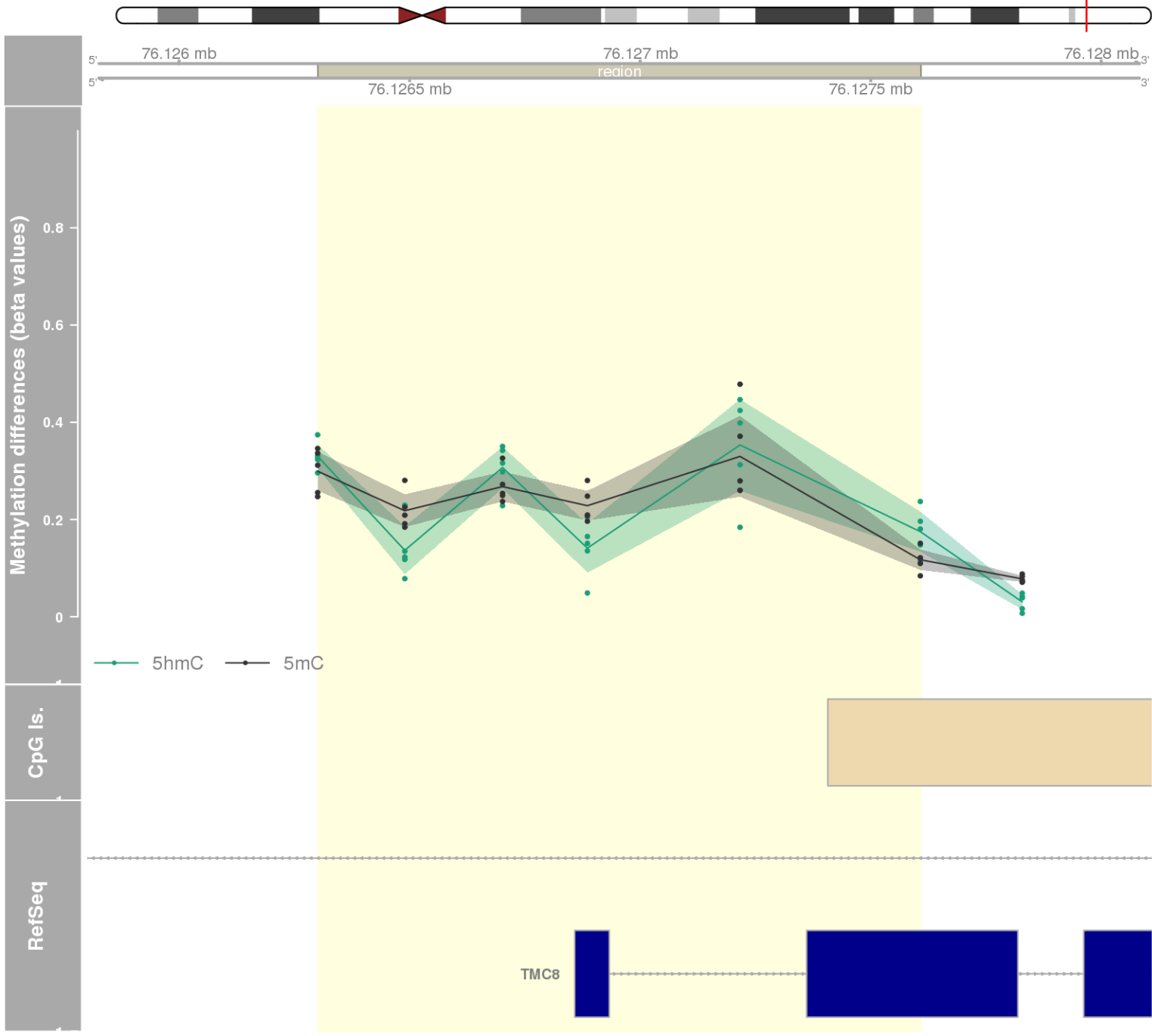
DMR 381 // chr14:89628169-89629974 // 1805 pb. (9 probes) // pvalue: 0.014 // fwer: 0.36 // pvalueArea: 0.027 // fwerArea: 1
- genes: FOXN3 -



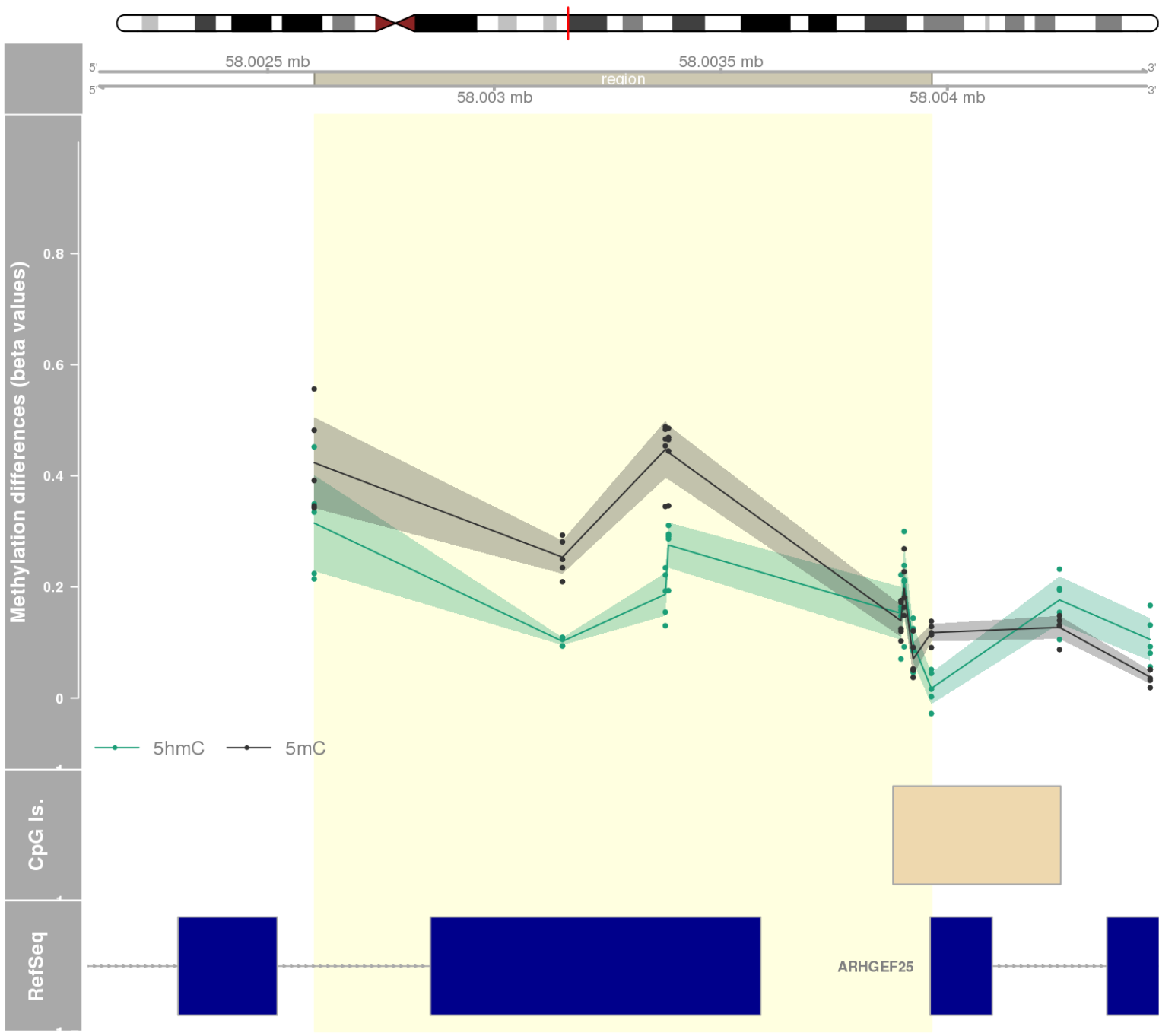
DMR 382 // chr2:71205052-71206811 // 1759 pb. (9 probes) // pvalue: 0.014 // fwer: 0.36 // pvalueArea: 0.027 // fwerArea: 1
- genes: ANKRD53 -



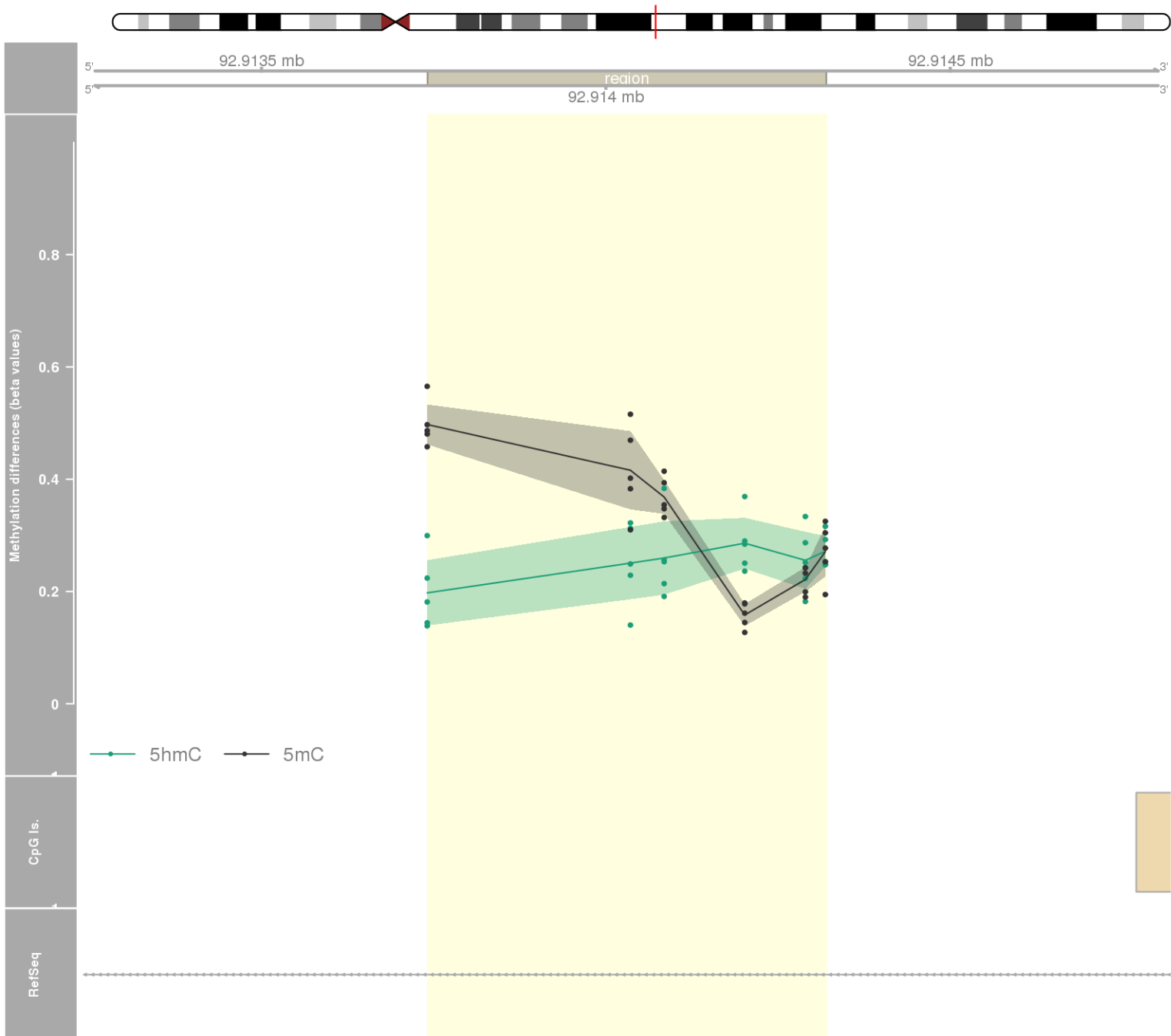
DMR 383 // chr17:76126301-76127608 // 1307 pb. (6 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.027 // fwerArea: 1
- genes: TMC6 / TMC8 -



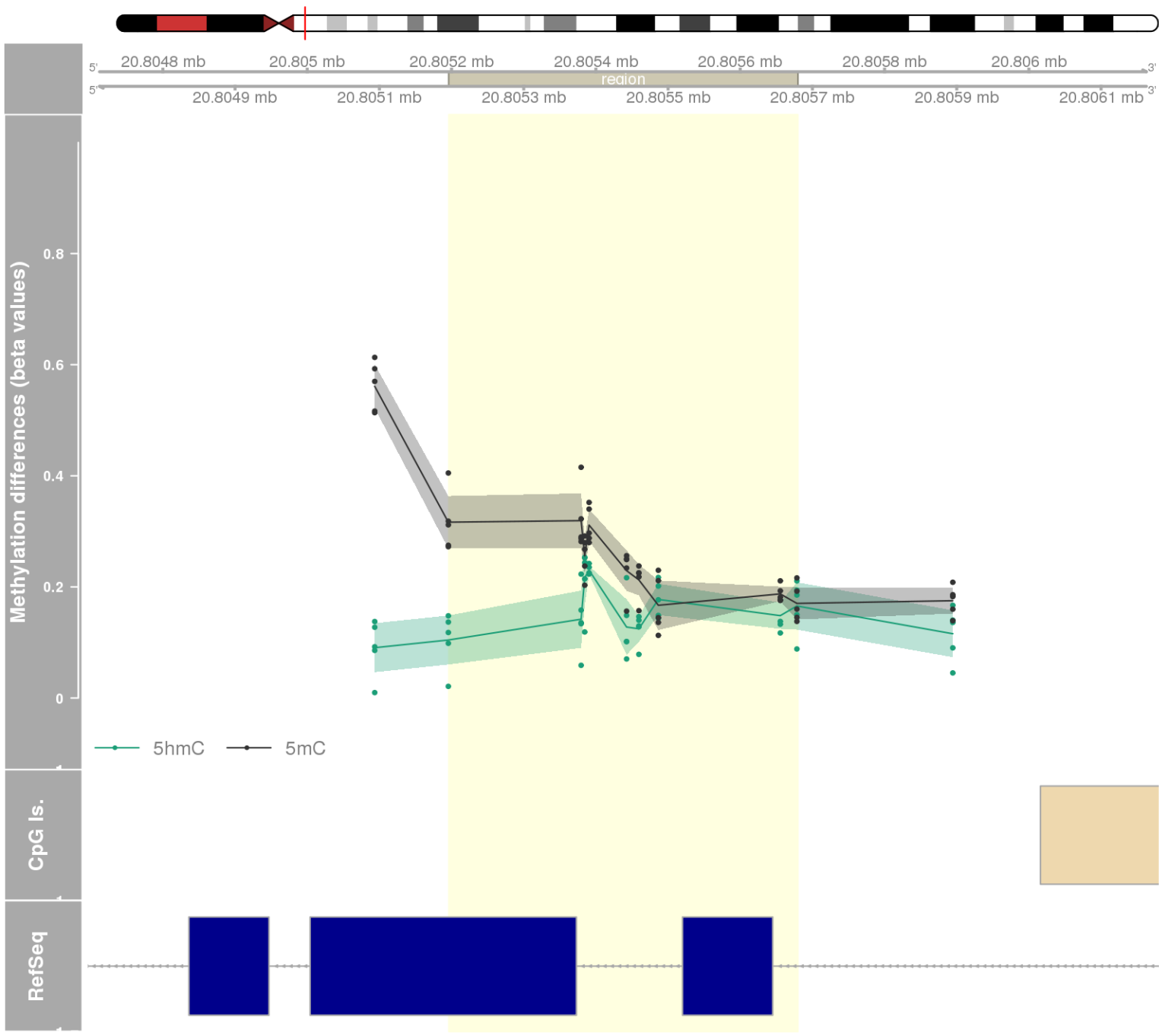
DMR 384 // chr12:58002603-58003965 // 1362 pb. (8 probes) // pvalue: 0.012 // fwer: 0.522 // pvalueArea: 0.027 // fwerArea: 1
- genes: DTX3 / ARHGEF25 -



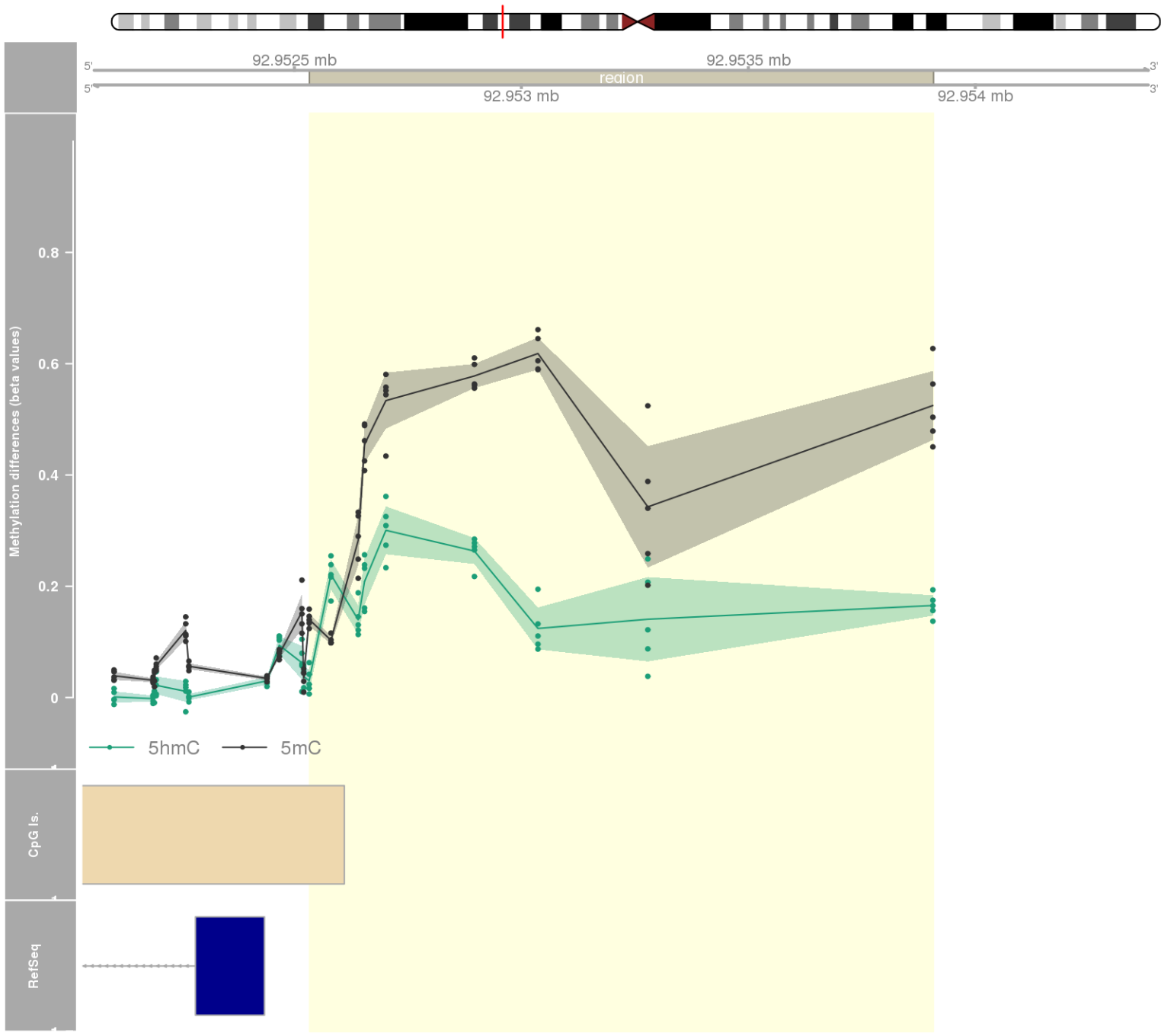
DMR 385 // chr5:92913741-92914319 // 578 pb. (6 probes) // pvalue: 0.002 // fwer: 0.173 // pvalueArea: 0.027 // fwerArea: 1
- genes: NR2F1-AS1 -



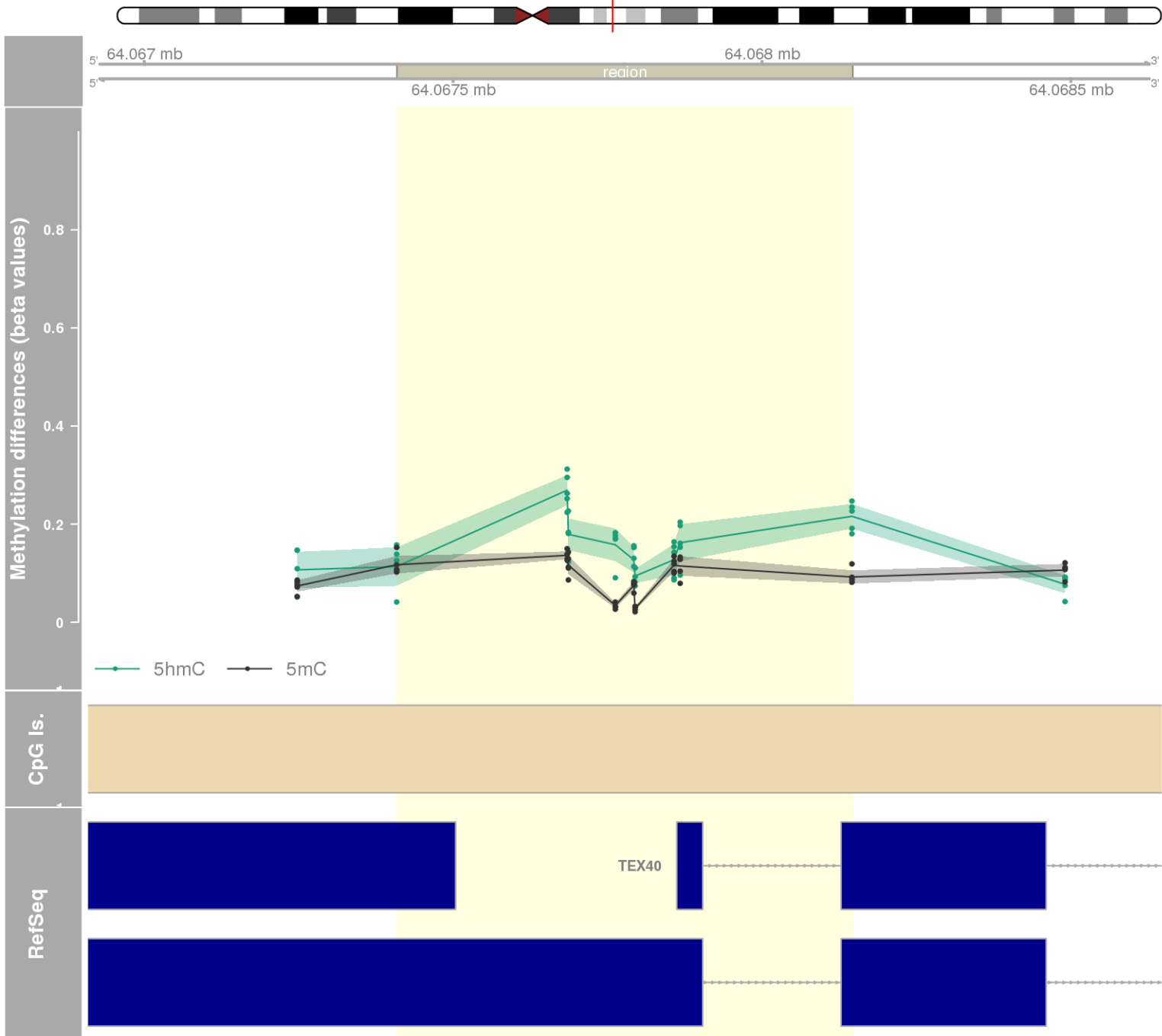
DMR 386 // chr13:20805196-20805679 // 483 pb. (9 probes) // pvalue: 0.014 // fwer: 0.36 // pvalueArea: 0.027 // fwerArea: 1
- genes: GJB6 -



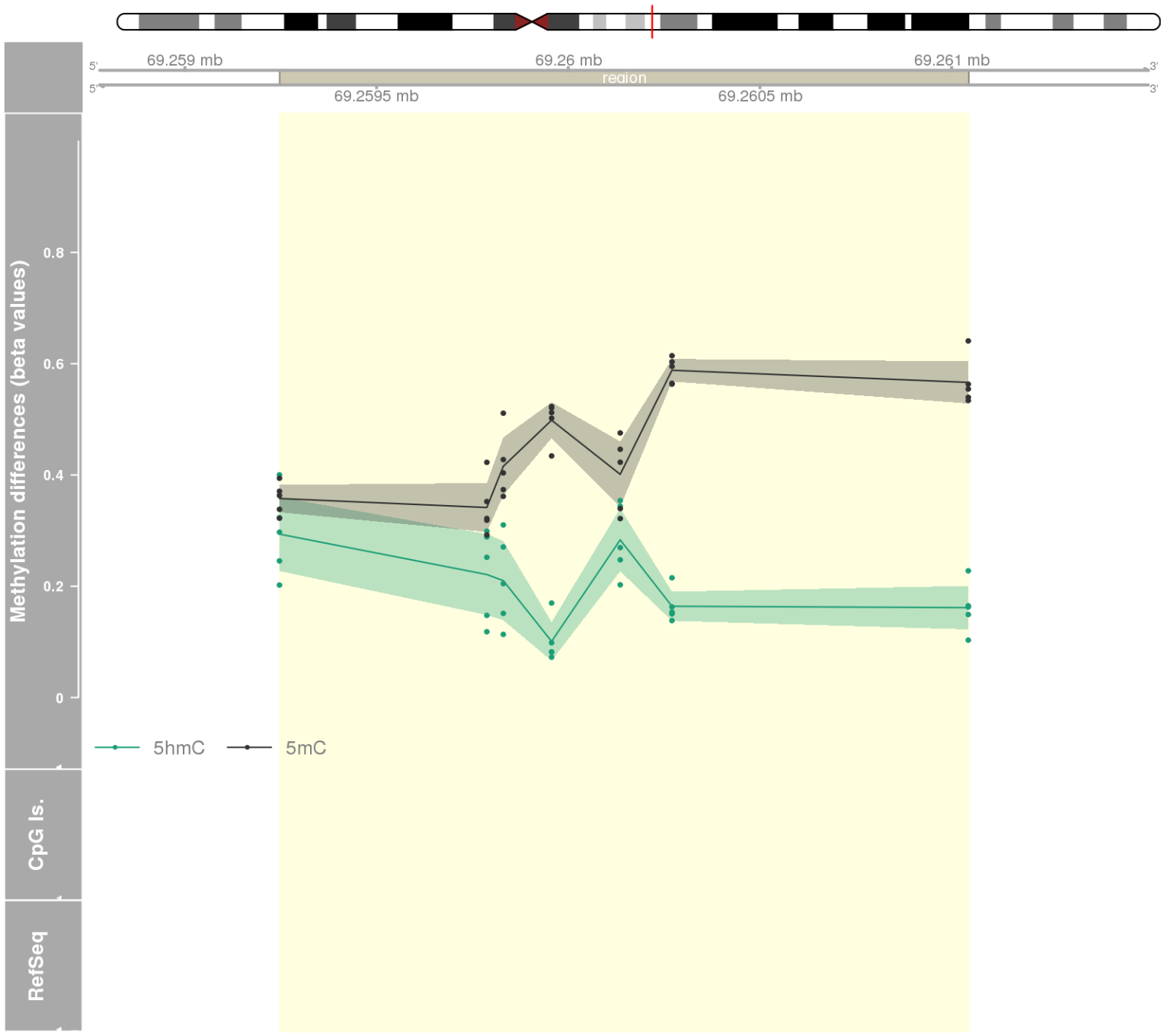
DMR 387 // chr1:92952533-92953907 // 1374 pb. (9 probes) // pvalue: 0.014 // fwer: 0.36 // pvalueArea: 0.027 // fwerArea: 1



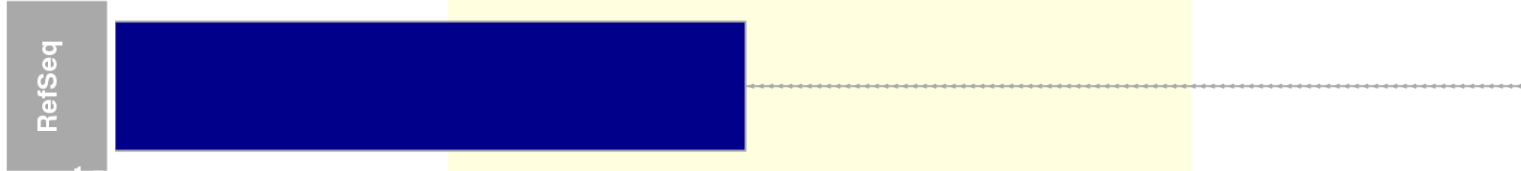
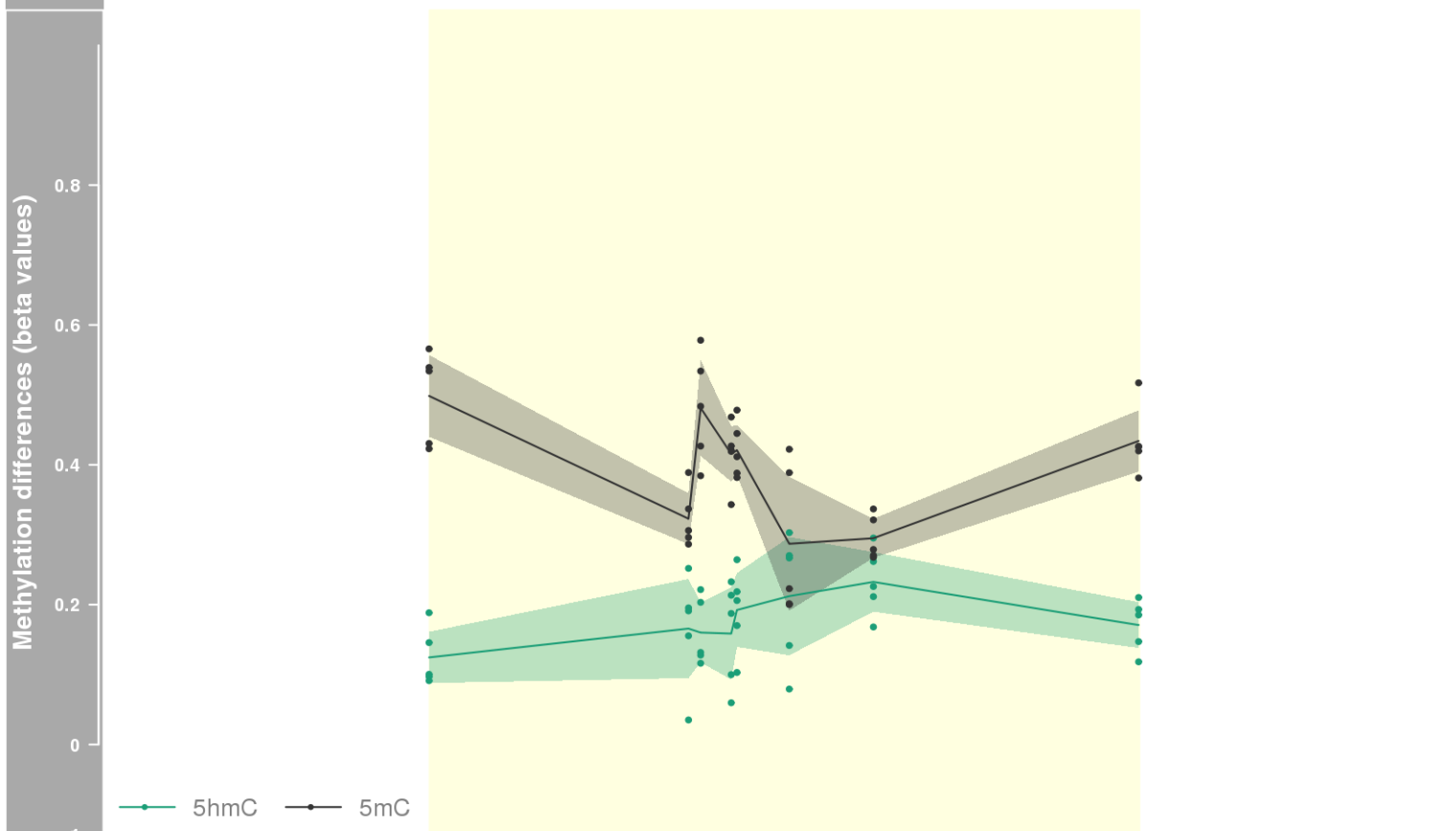
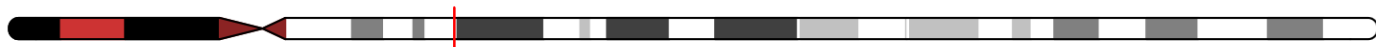
DMR 388 // chr11:64067409-64068146 // 737 pb. (9 probes) // pvalue: 0.014 // fwer: 0.36 // pvalueArea: 0.027 // fwerArea: 1
- genes: KCNK4 / TEX40 -



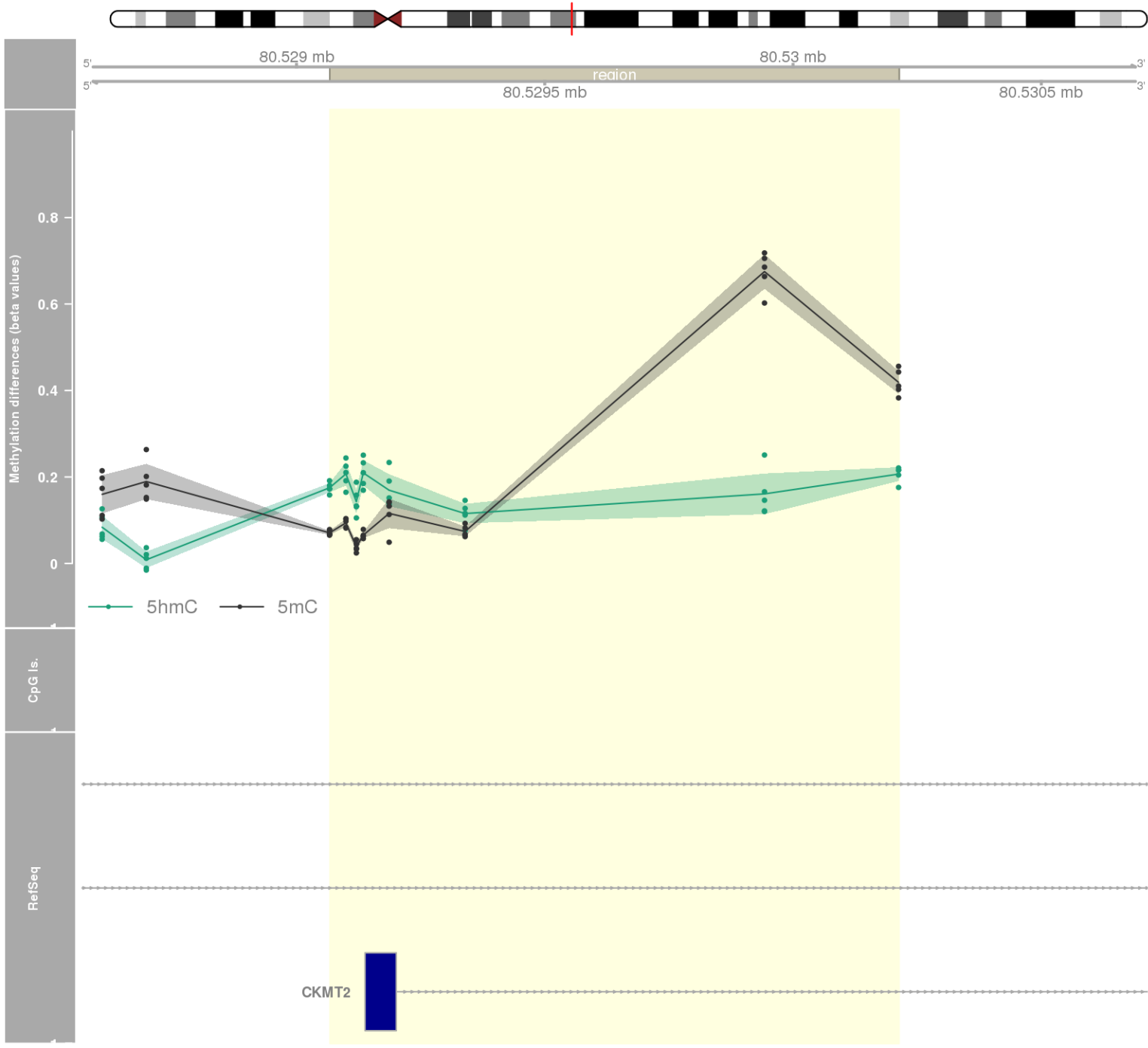
DMR 389 // chr11:69259247-69261044 // 1797 pb. (7 probes) // pvalue: 0.008 // fwer: 0.173 // pvalueArea: 0.027 // fwerArea: 1



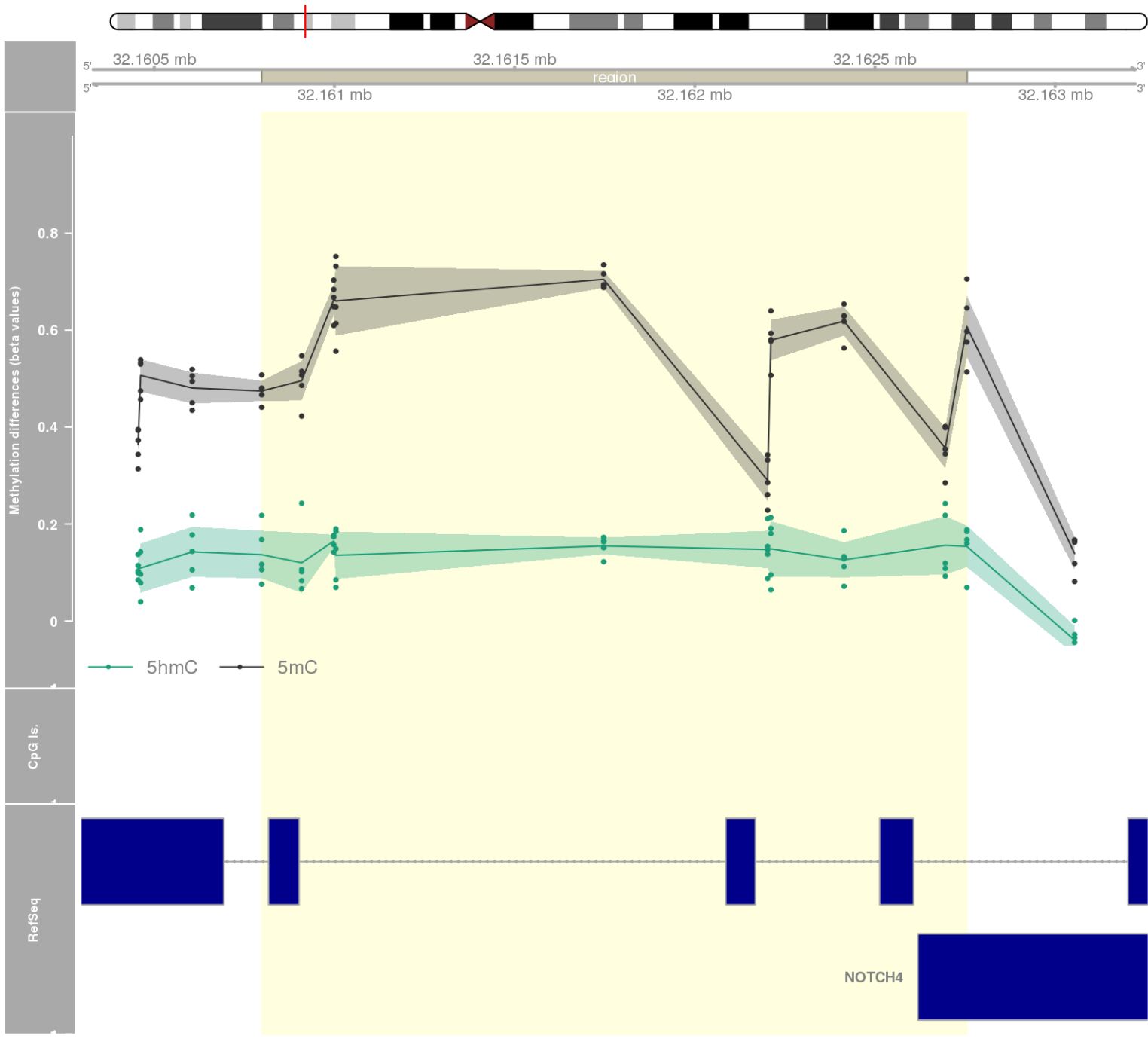
DMR 390 // chr15:33359788-33360901 // 1113 pb. (8 probes) // pvalue: 0.012 // fwer: 0.522 // pvalueArea: 0.027 // fwerArea: 1
- genes: FMN1 -



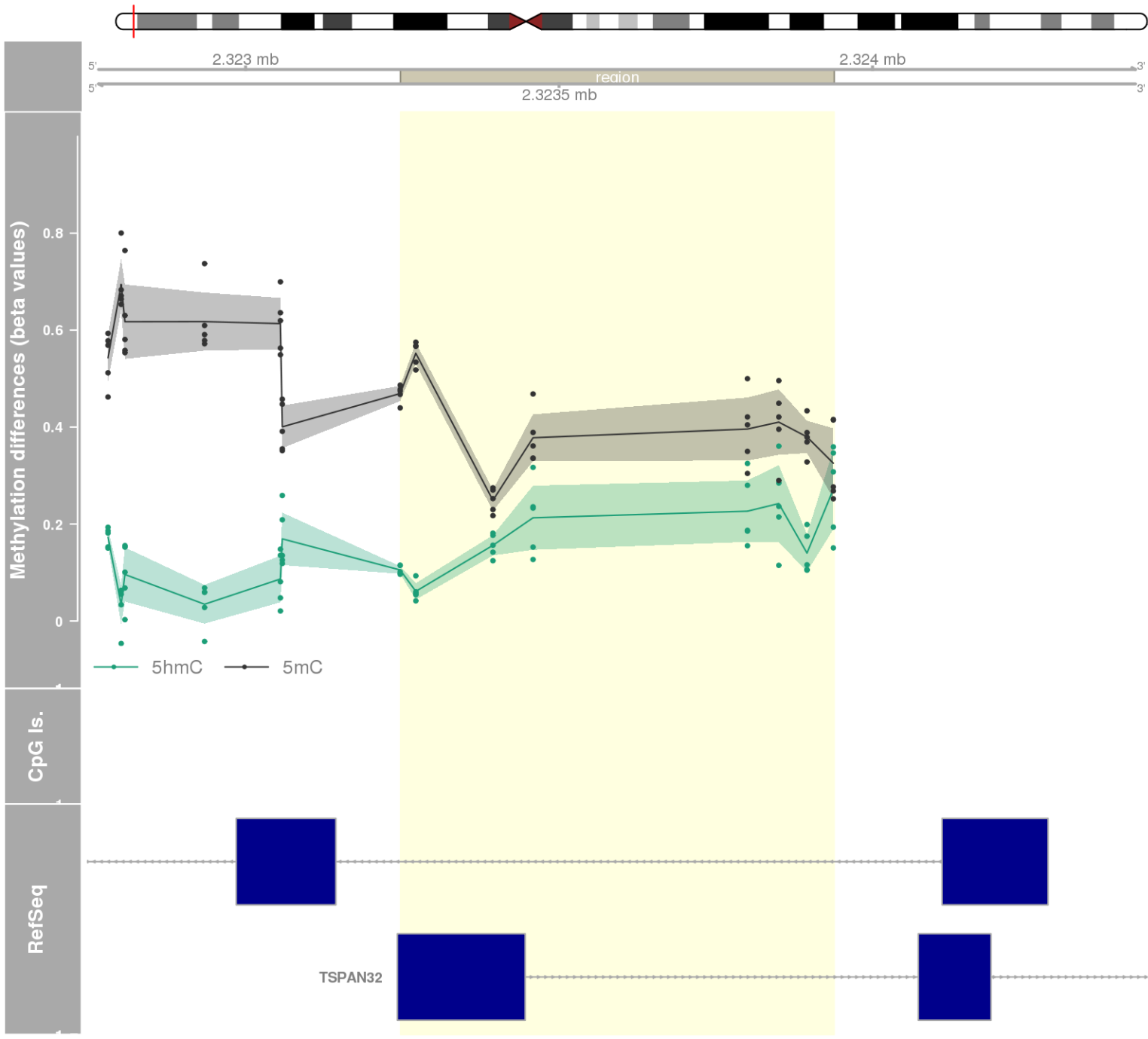
DMR 391 // chr5:80529067-80530213 // 1146 pb. (8 probes) // pvalue: 0.012 // fwer: 0.522 // pvalueArea: 0.027 // fwerArea: 1
- genes: CKMT2-AS1 / CKMT2 -



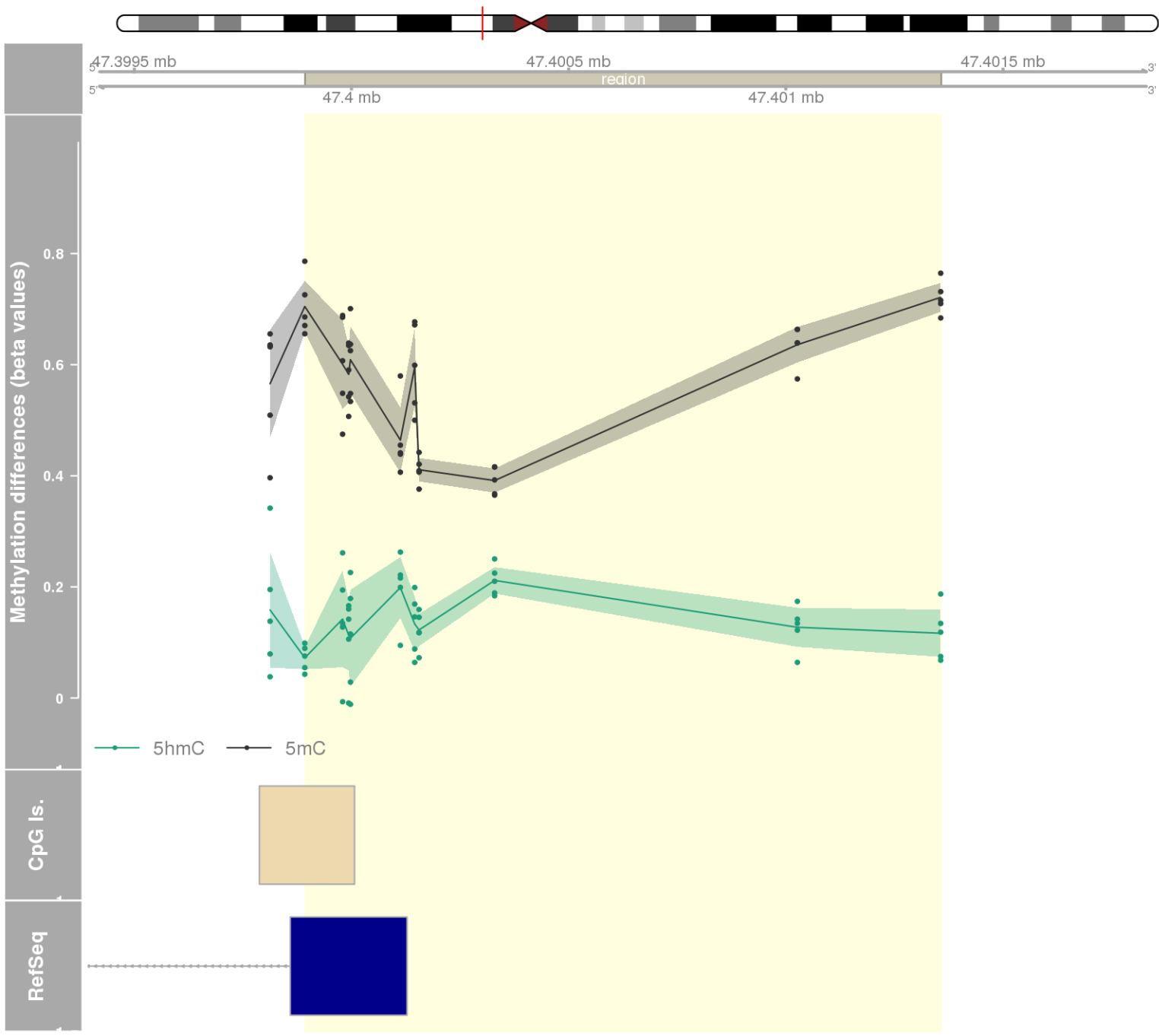
DMR 392 // chr6:32160798-32162755 // 1957 pb. (10 probes) // pvalue: 0.014 // fwer: 0.36 // pvalueArea: 0.027 // fwerArea: 1
- genes: GP5M3 / NOTCH4 -



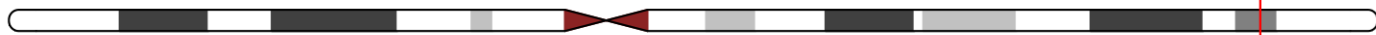
DMR 393 // chr11:2323247-2323938 // 691 pb. (8 probes) // pvalue: 0.012 // fwer: 0.522 // pvalueArea: 0.027 // fwerArea: 1
- genes: C11orf21 / TSPAN32 -



DMR 394 // chr11:47399893-47401357 // 1464 pb. (10 probes) // pvalue: 0.014 // fwer: 0.36 // pvalueArea: 0.027 // fwerArea: 1
- genes: SPI1 -



DMR 395 // chr20:57615398-57617168 // 1770 pb. (6 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.027 // fwerArea: 1
- genes: SLMO2 -



57.615 mb 57.616 mb 57.617 mb
5' 3' region 57.6155 mb 57.6165 mb 57.6175 mb 3'

Methylation differences (beta values)

0.8

0.6

0.4

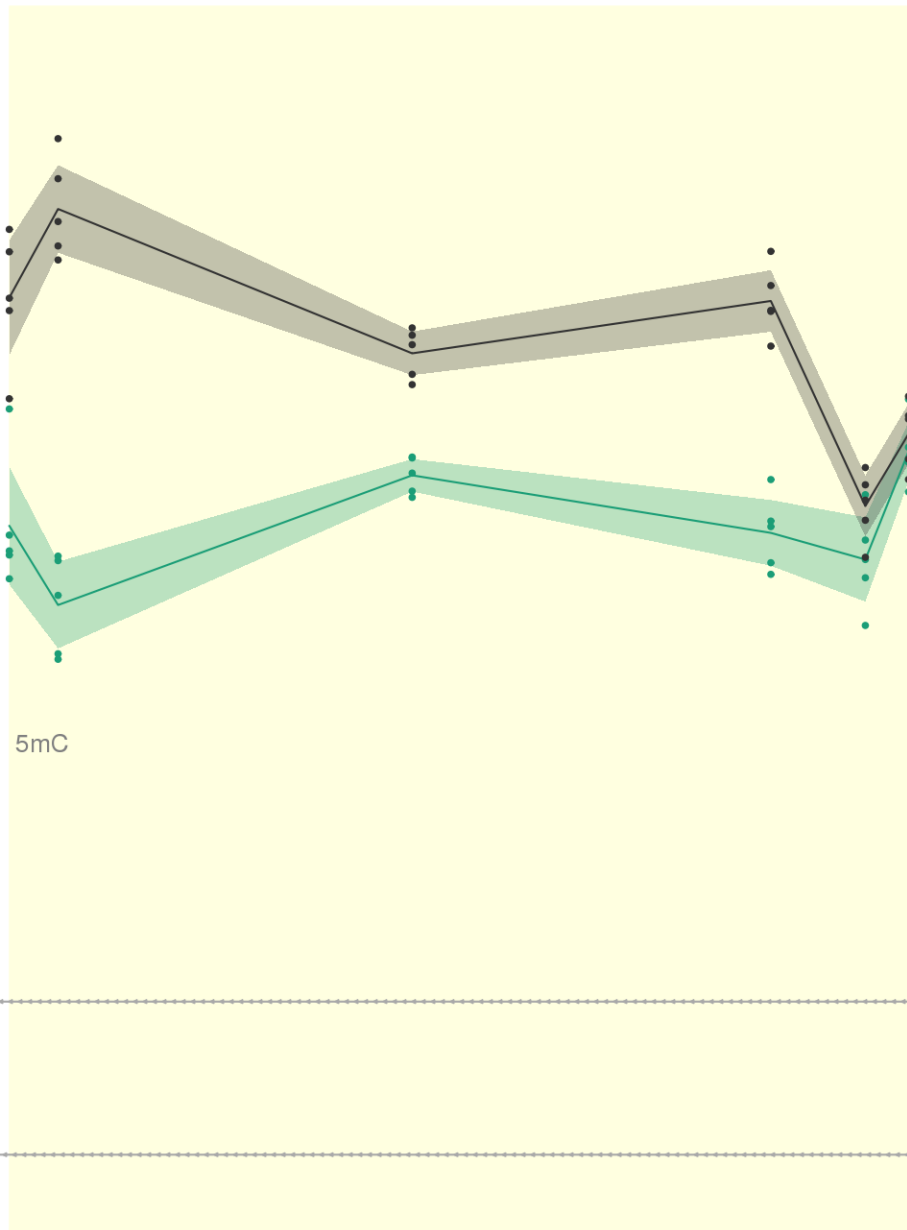
0.2

0

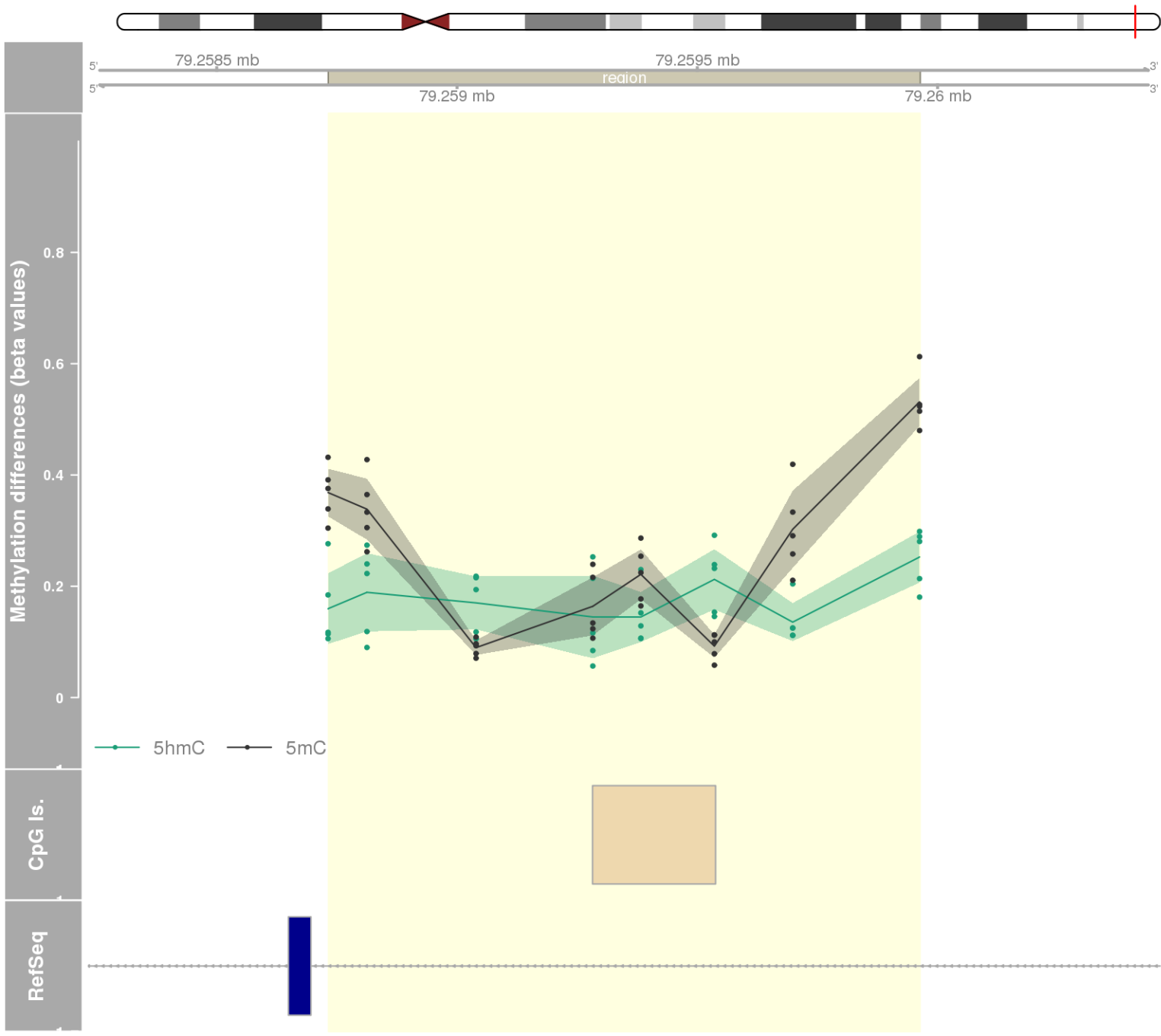
5hmC 5mC

CpG Is.

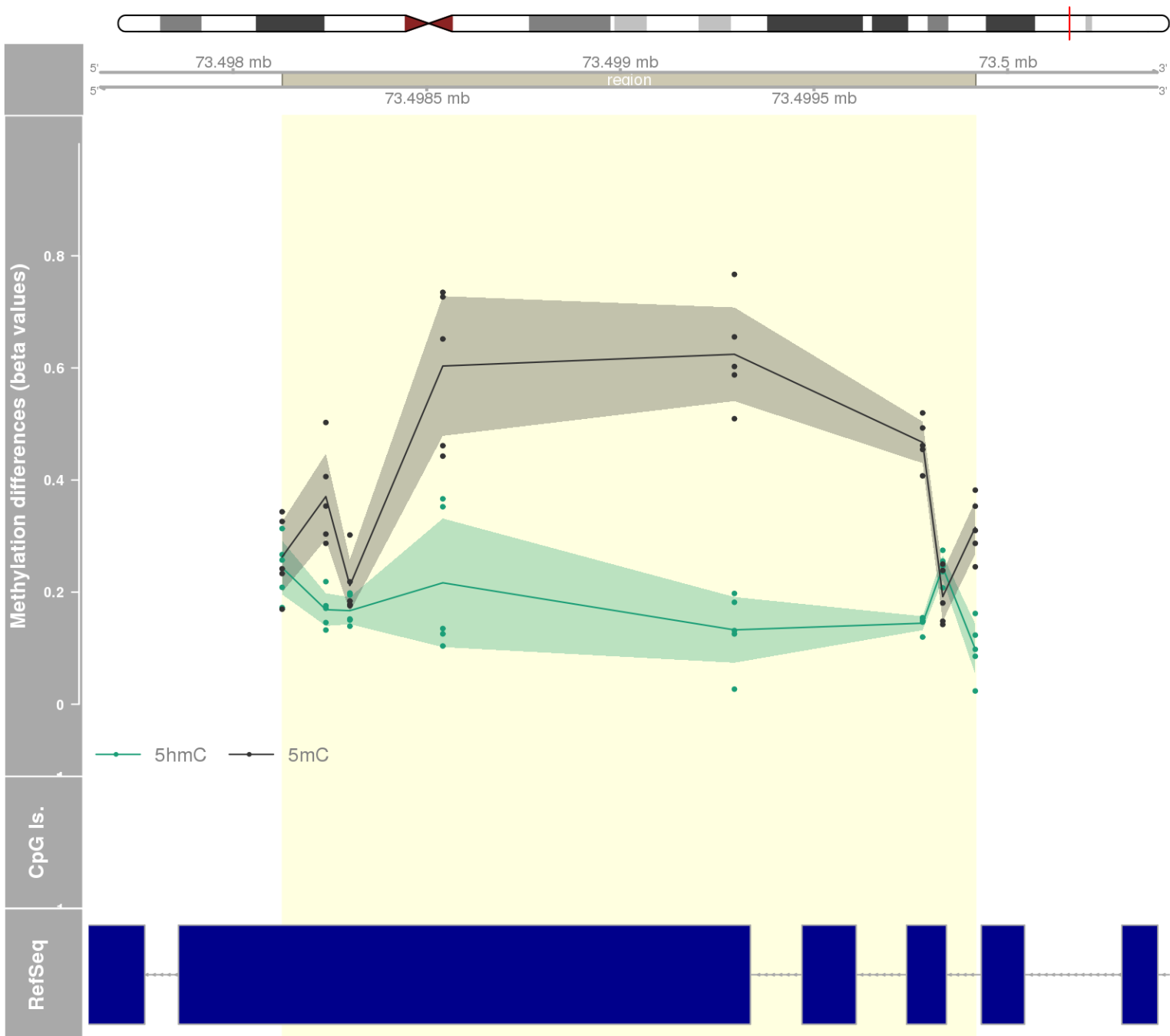
RefSeq



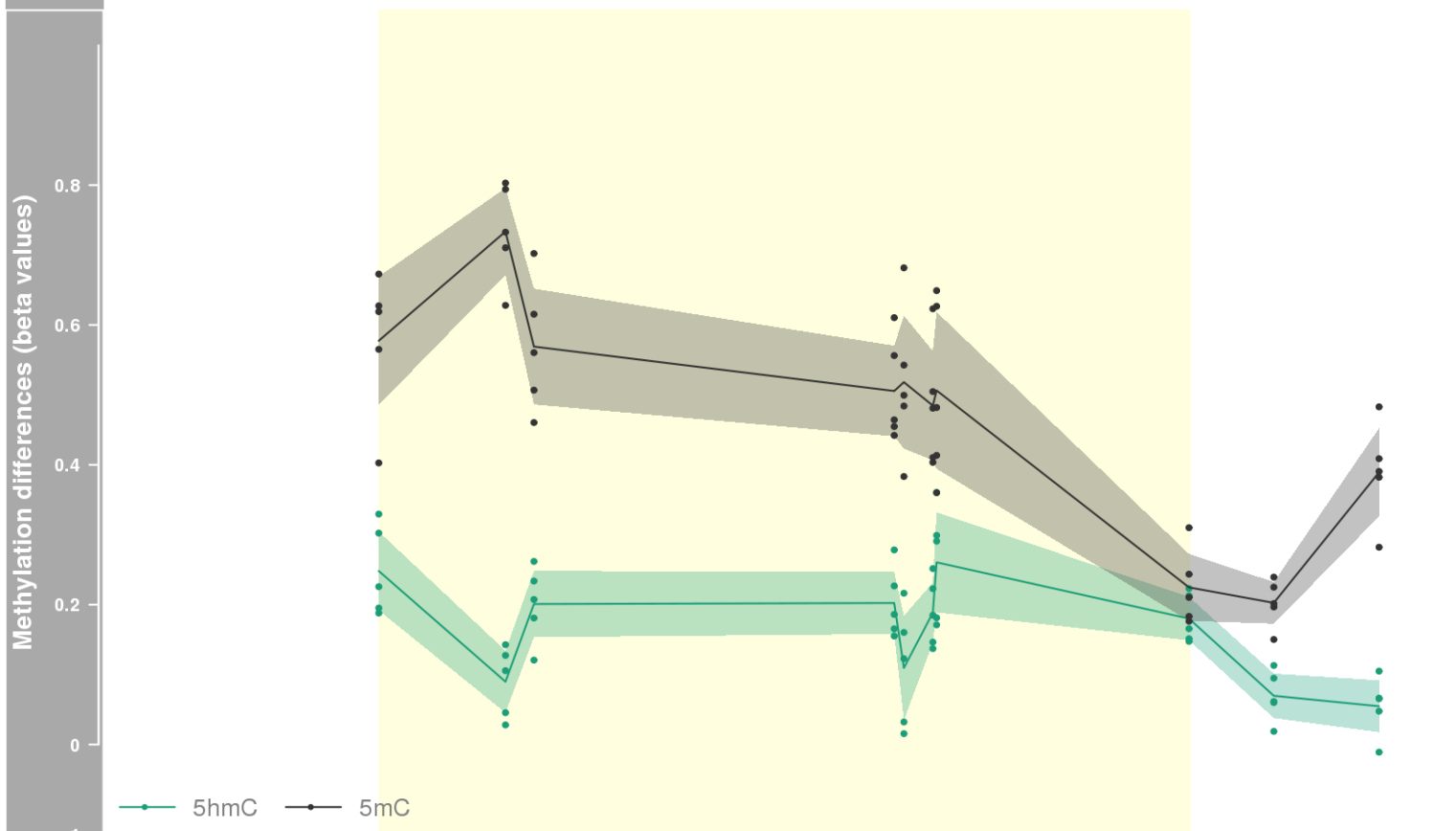
DMR 396 // chr17:79258732-79259963 // 1231 pb. (8 probes) // pvalue: 0.012 // fwer: 0.522 // pvalueArea: 0.028 // fwerArea: 1
- genes: SLC38A10 -



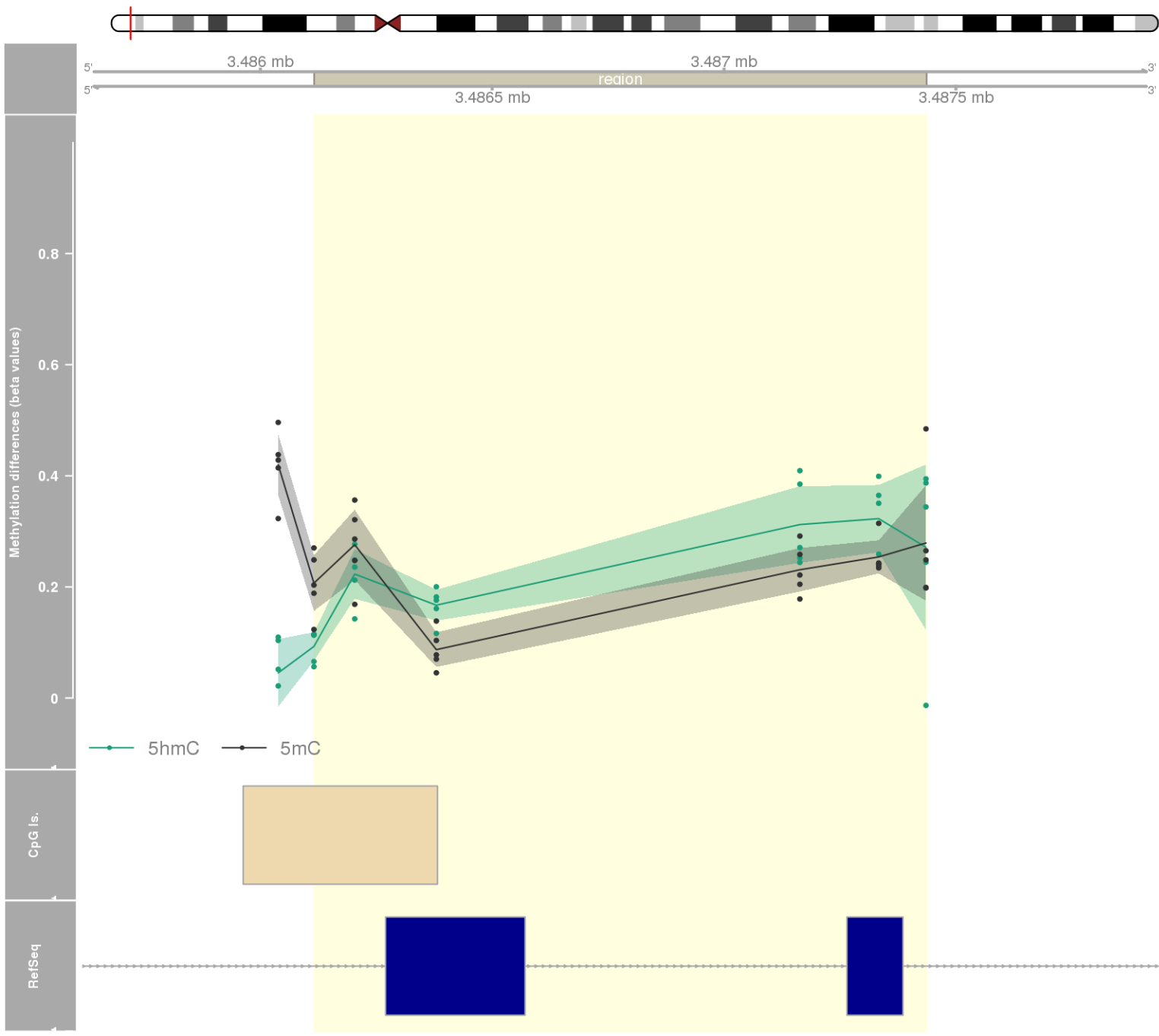
DMR 397 // chr17:73498127-73499917 // 1790 pb. (8 probes) // pvalue: 0.013 // fwer: 0.522 // pvalueArea: 0.028 // fwerArea: 1
- genes: CASKIN2 -



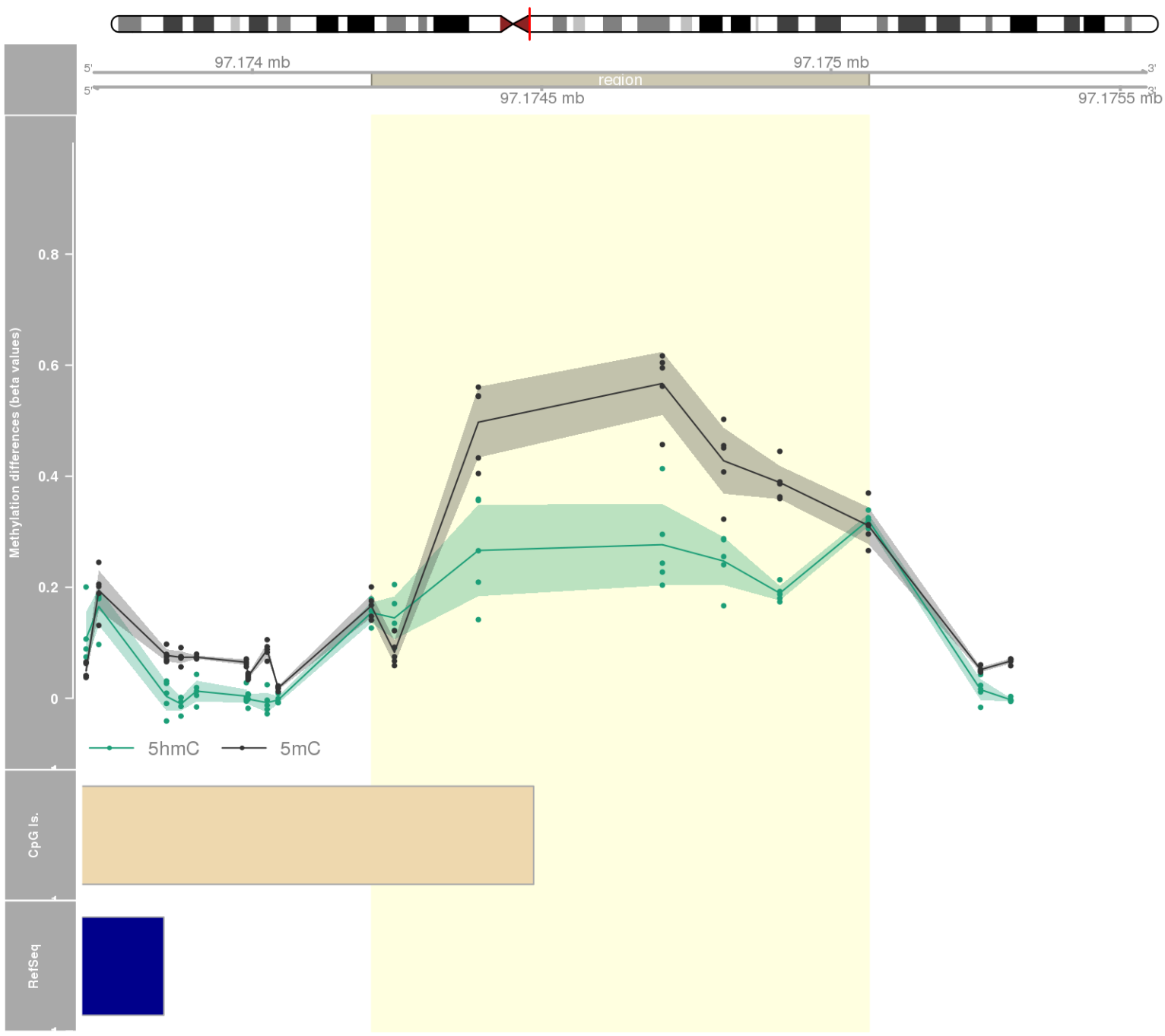
DMR 398 // chr10:21796152-21797661 // 1509 pb. (8 probes) // pvalue: 0.013 // fwer: 0.522 // pvalueArea: 0.028 // fwerArea: 1



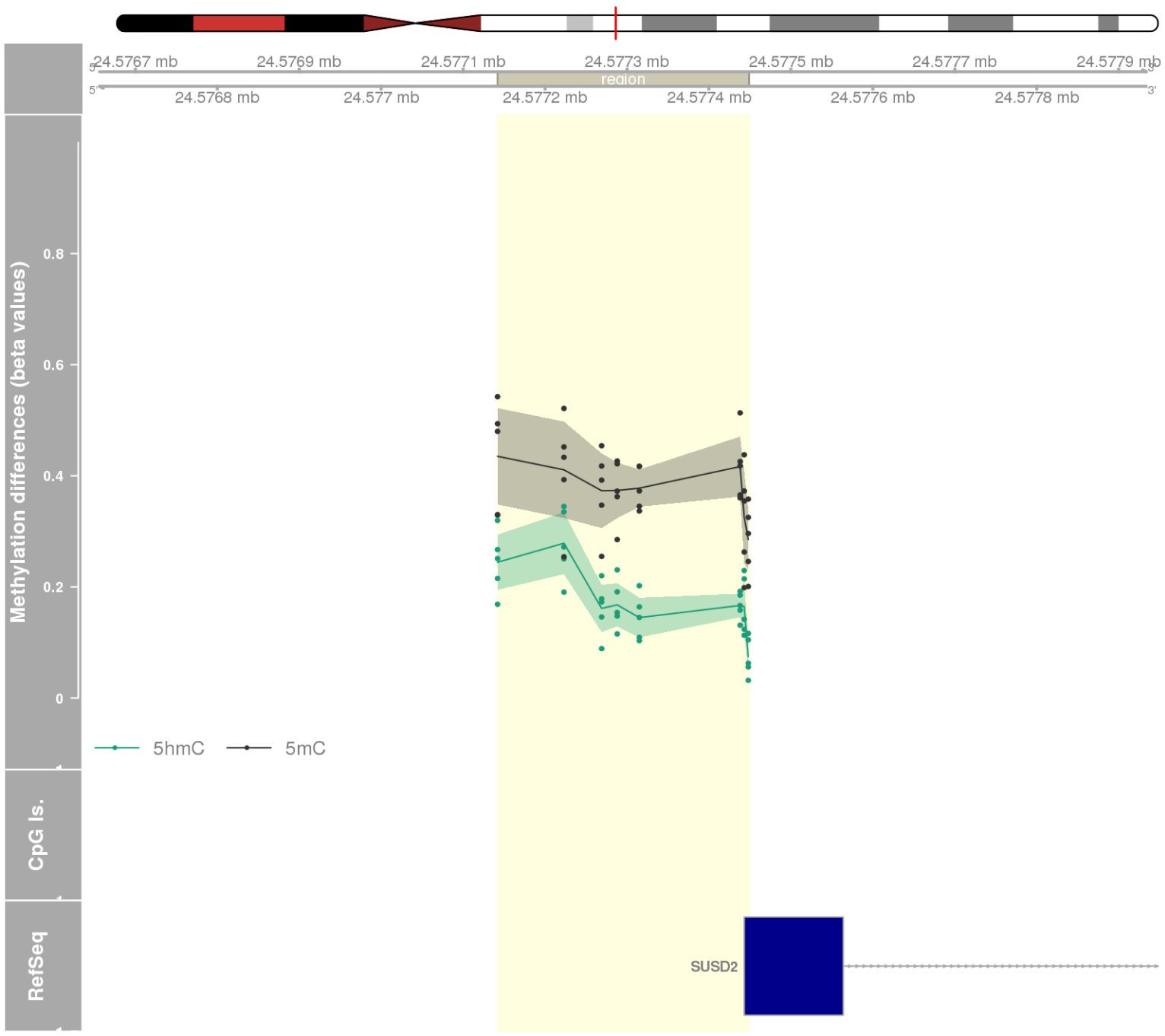
DMR 399 // chr4:3486116-3487436 // 1320 pb. (6 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.028 // fwerArea: 1
- genes: DOK7 -



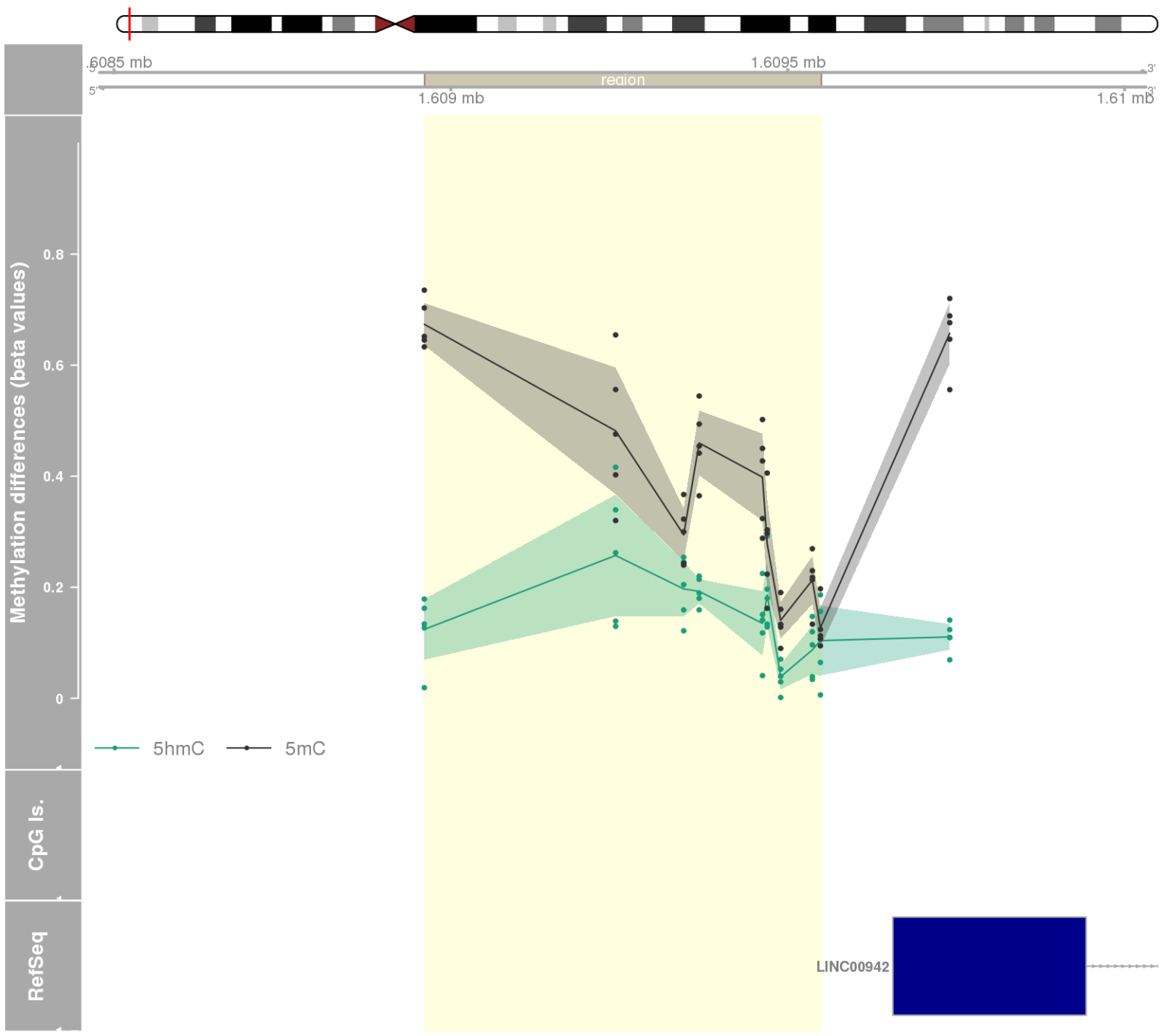
DMR 400 // chr2:97174206-97175065 // 859 pb. (7 probes) // pvalue: 0.008 // fwer: 0.173 // pvalueArea: 0.028 // fwerArea: 1



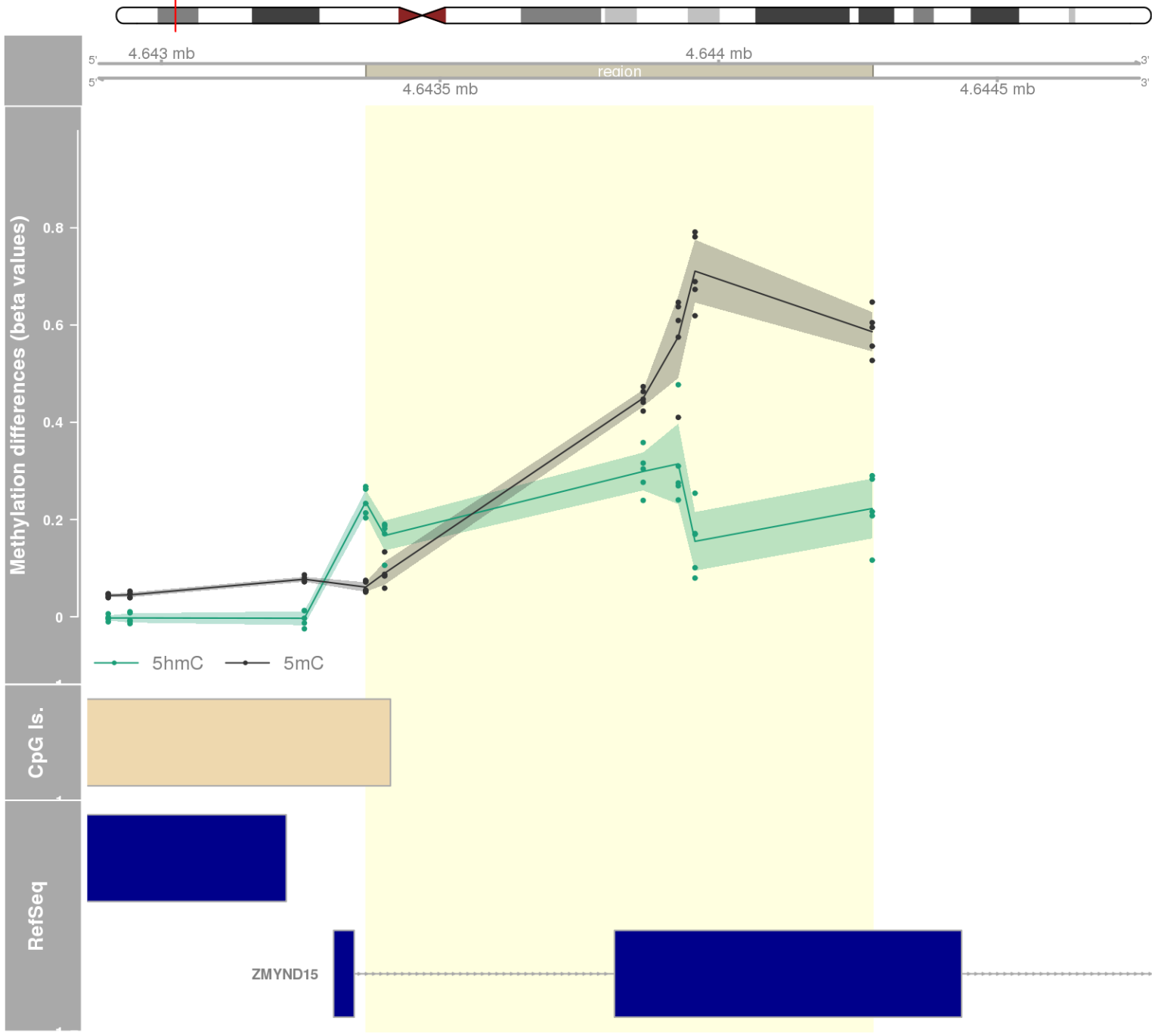
DMR 401 // chr22:24577142-24577448 // 306 pb. (8 probes) // pvalue: 0.013 // fwer: 0.522 // pvalueArea: 0.028 // fwerArea: 1
- genes: SUSD2 -



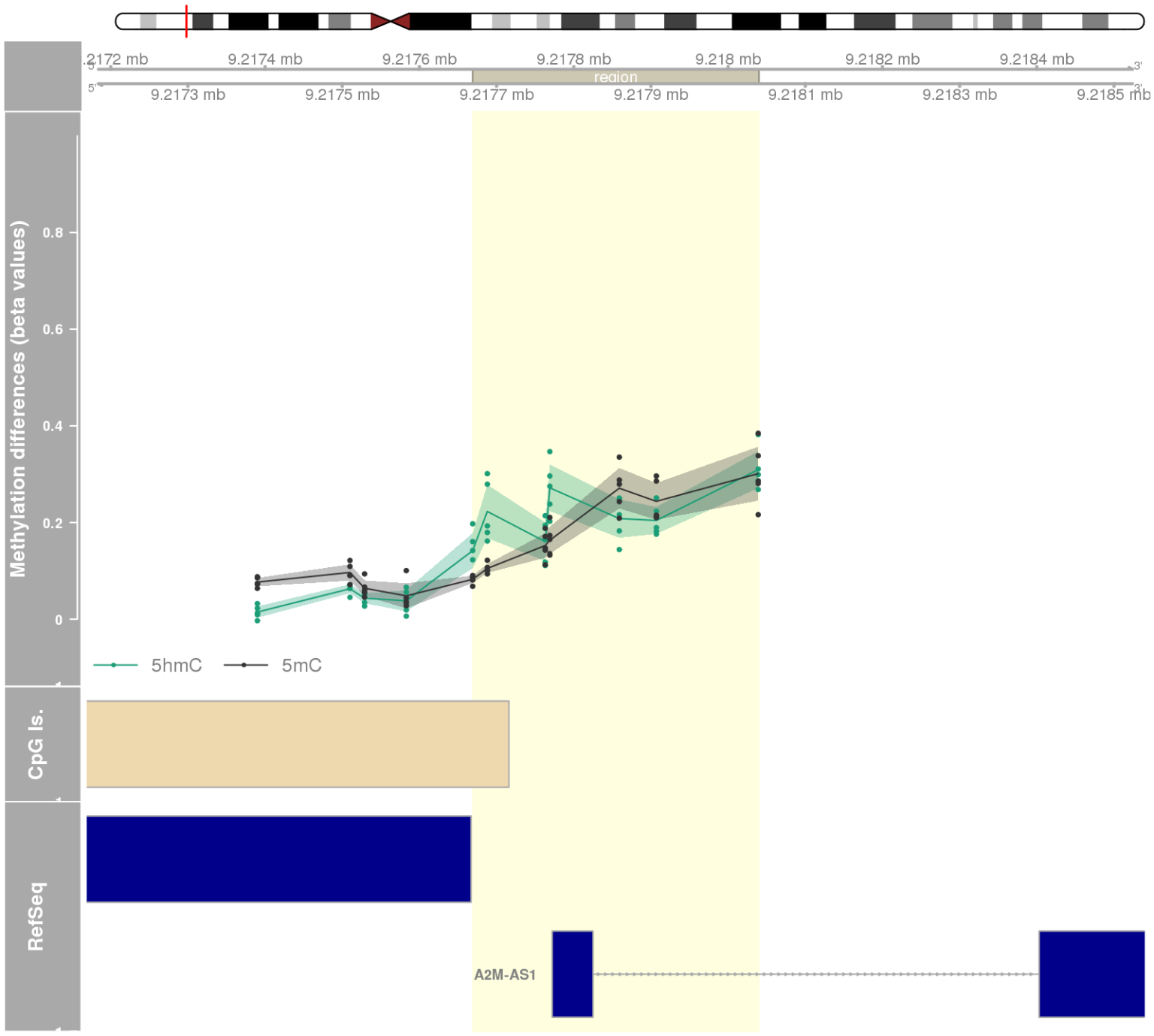
DMR 402 // chr12:1608961-1609549 // 588 pb. (9 probes) // pvalue: 0.014 // fwer: 0.36 // pvalueArea: 0.028 // fwerArea: 1



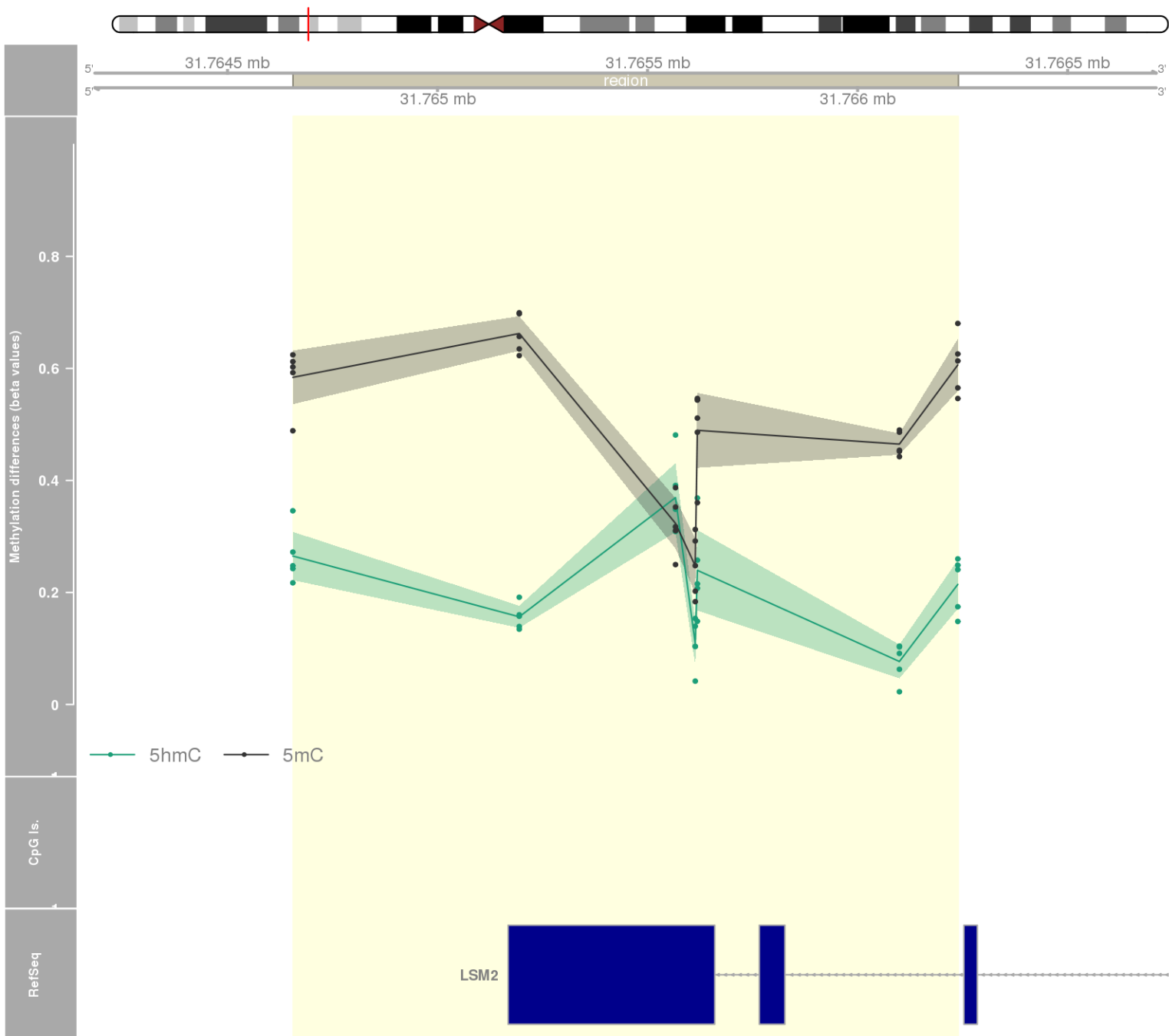
DMR 403 // chr17:4643367-4644276 // 909 pb. (6 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.028 // fwerArea: 1
- genes: ZMYND15 -



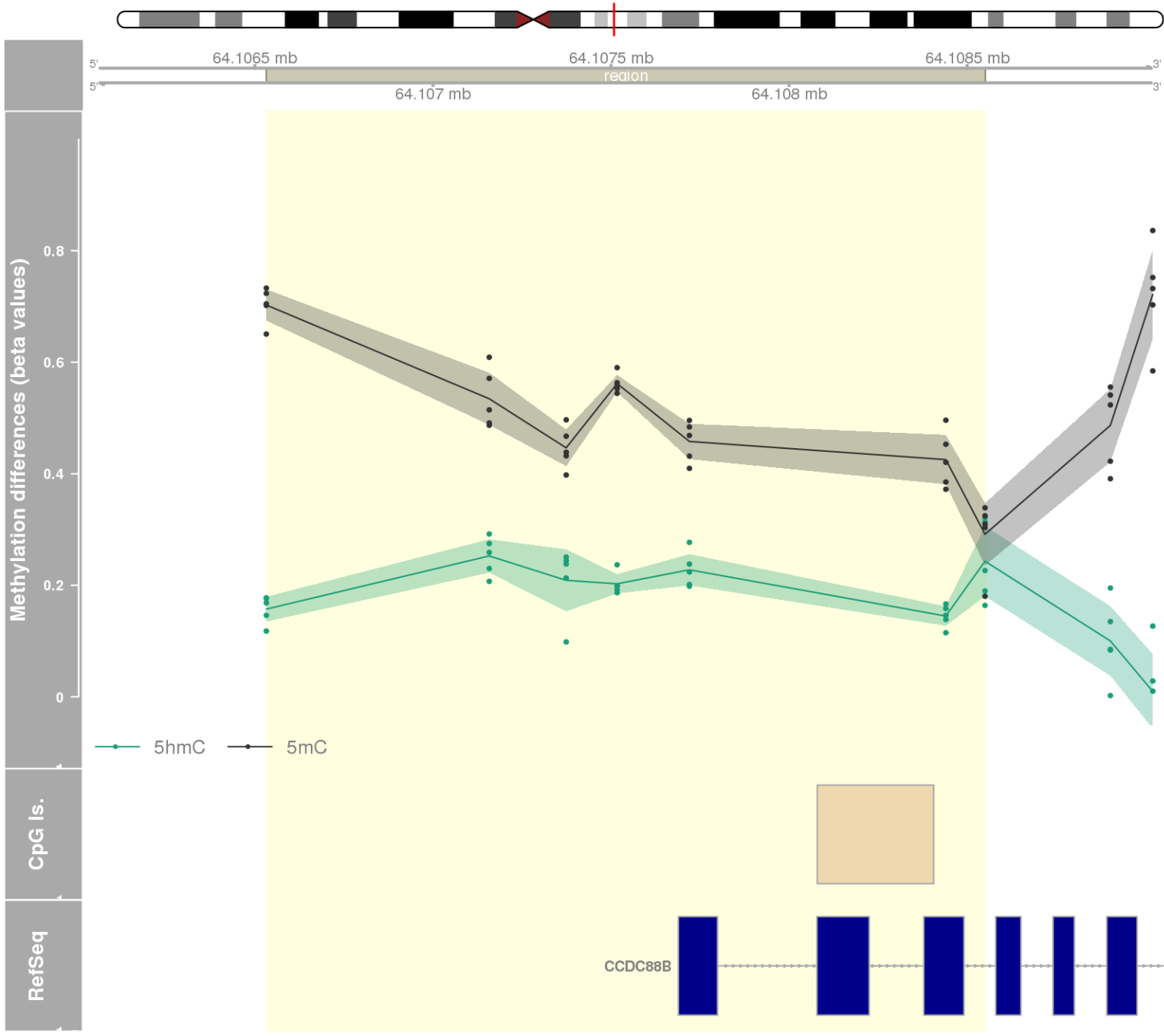
DMR 404 // chr12:9217669-9218039 // 370 pb. (7 probes) // pvalue: 0.009 // fwer: 0.173 // pvalueArea: 0.028 // fwerArea: 1
- genes: A2M-AS1 -



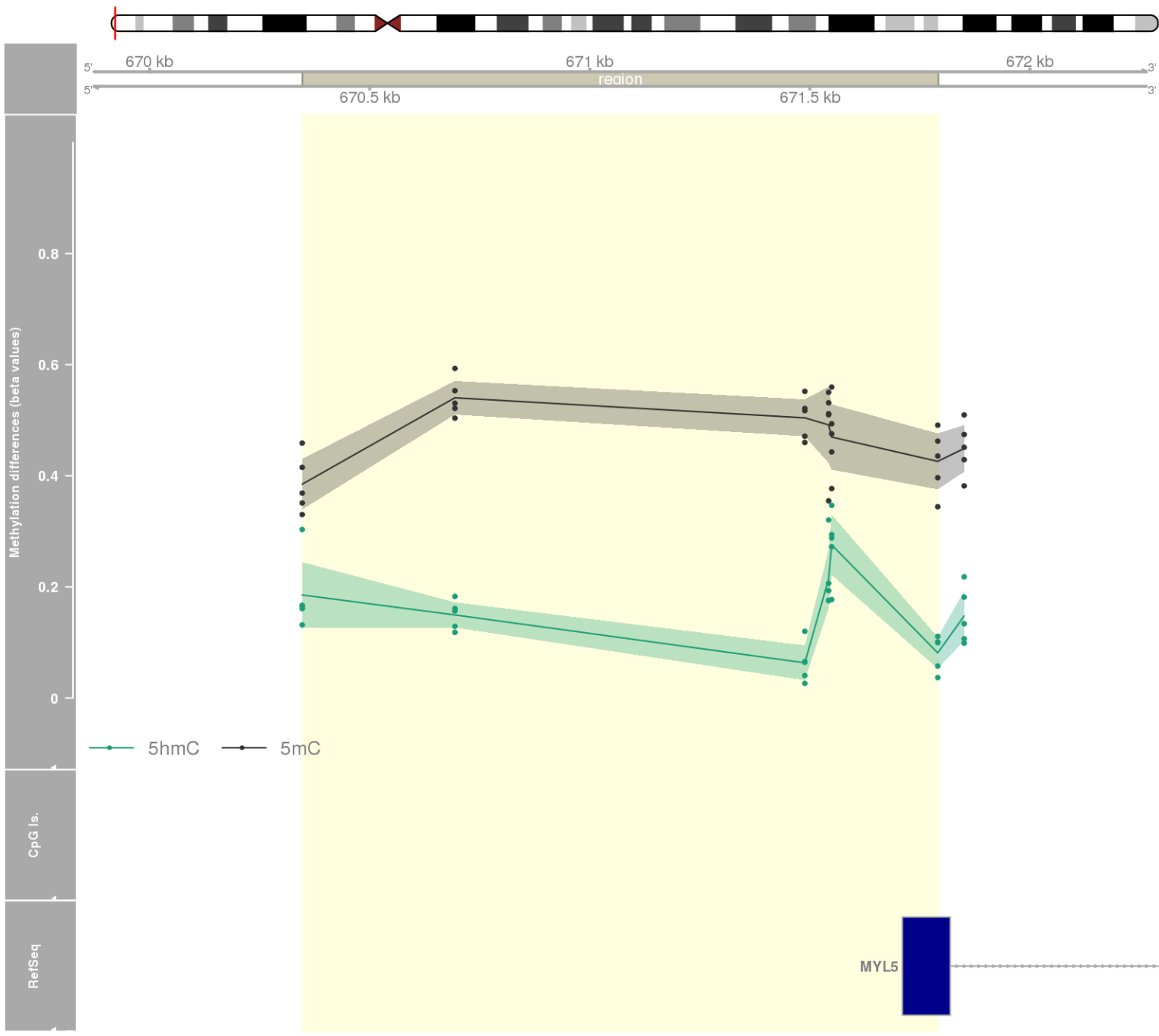
DMR 405 // chr6:31764656-31766239 // 1583 pb. (7 probes) // pvalue: 0.009 // fwer: 0.173 // pvalueArea: 0.028 // fwerArea: 1
- genes: LSM2 -



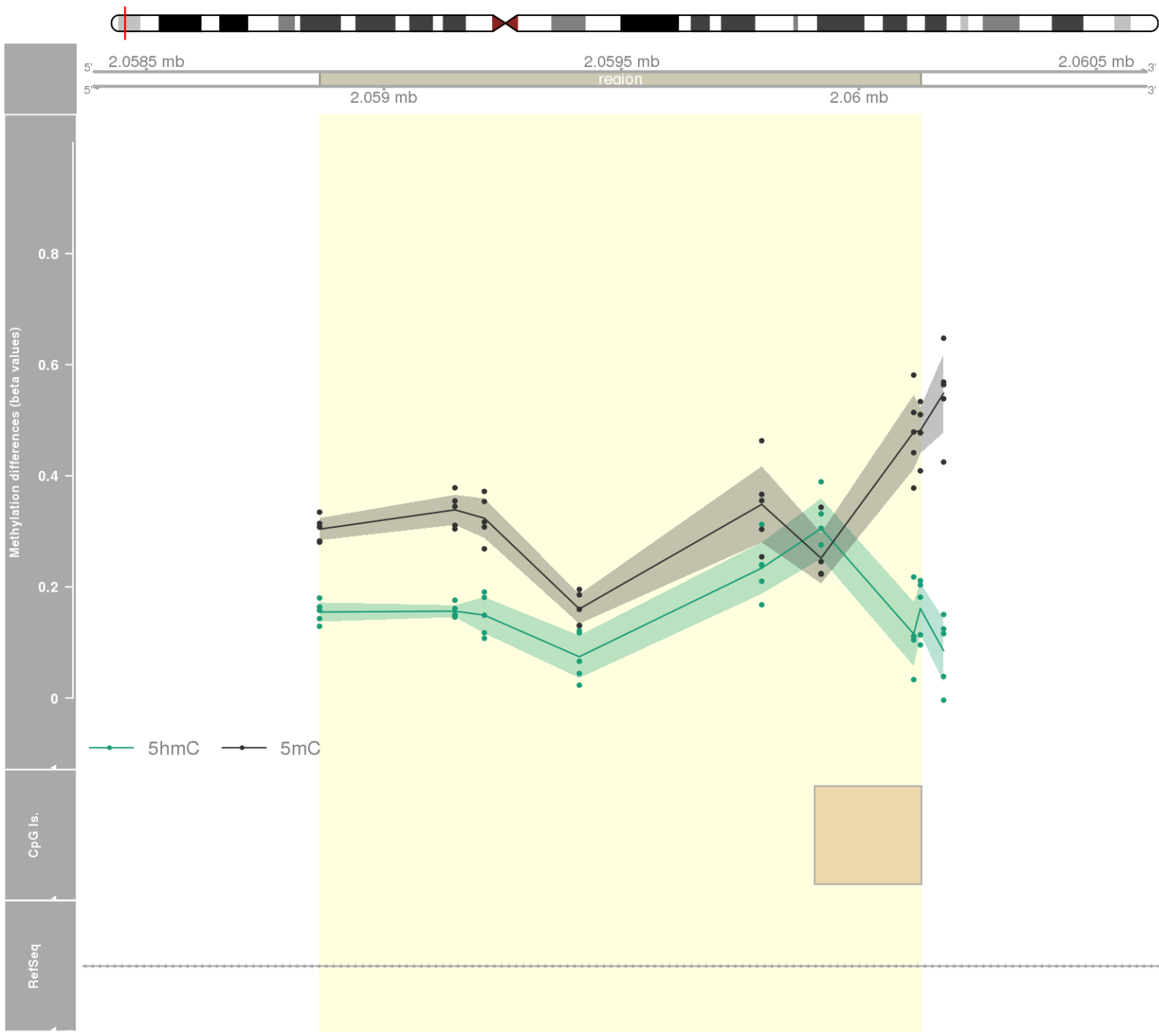
DMR 406 // chr11:64106532-64108550 // 2018 pb. (7 probes) // pvalue: 0.009 // fwer: 0.173 // pvalueArea: 0.028 // fwerArea: 1
- genes: CCDC88B -



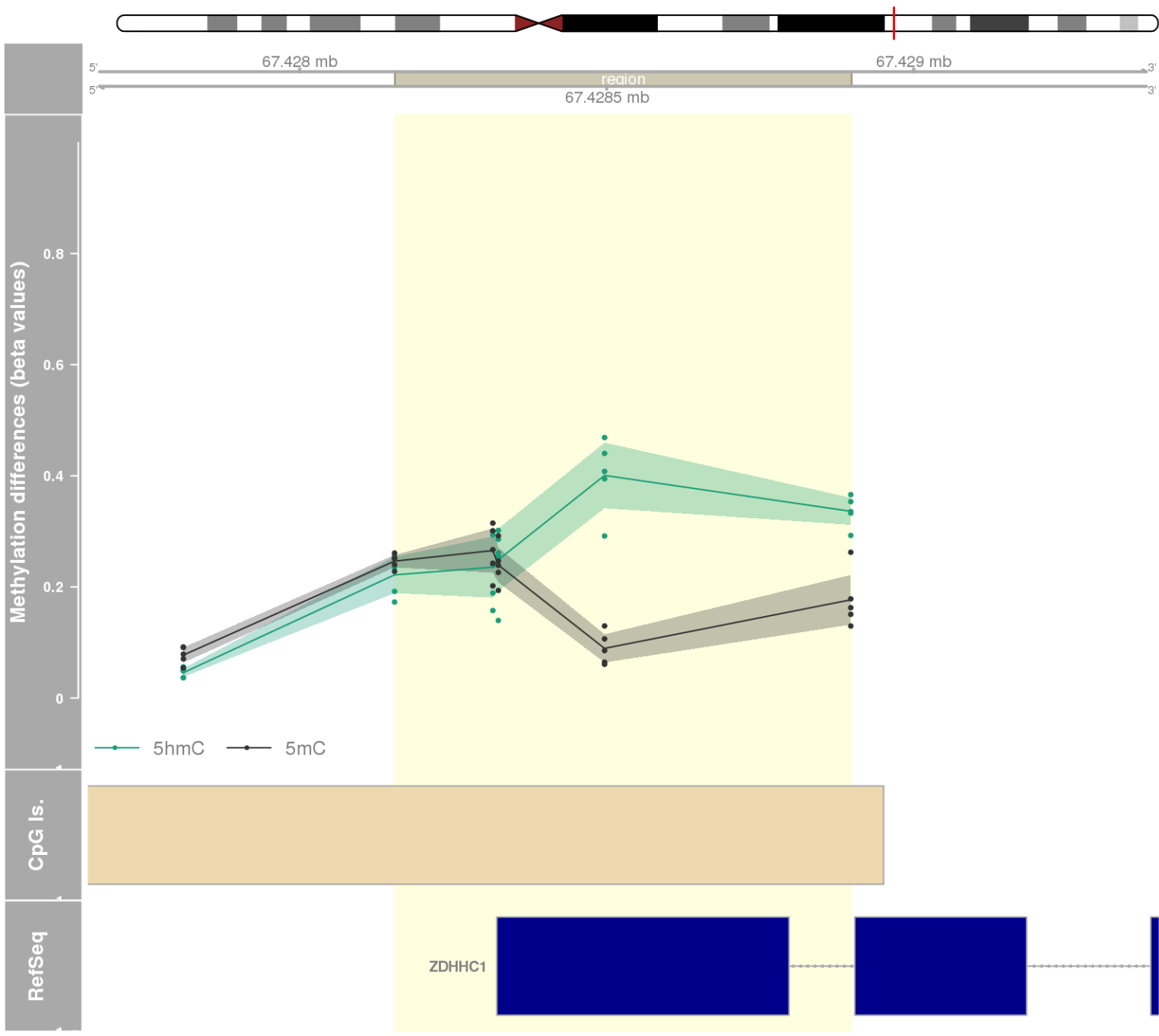
DMR 407 // chr4:670347-671791 // 1444 pb. (6 probes) // pvalue: 0.003 // fwer: 0.173 // pvalueArea: 0.028 // fwerArea: 1
- genes: MYL5 -



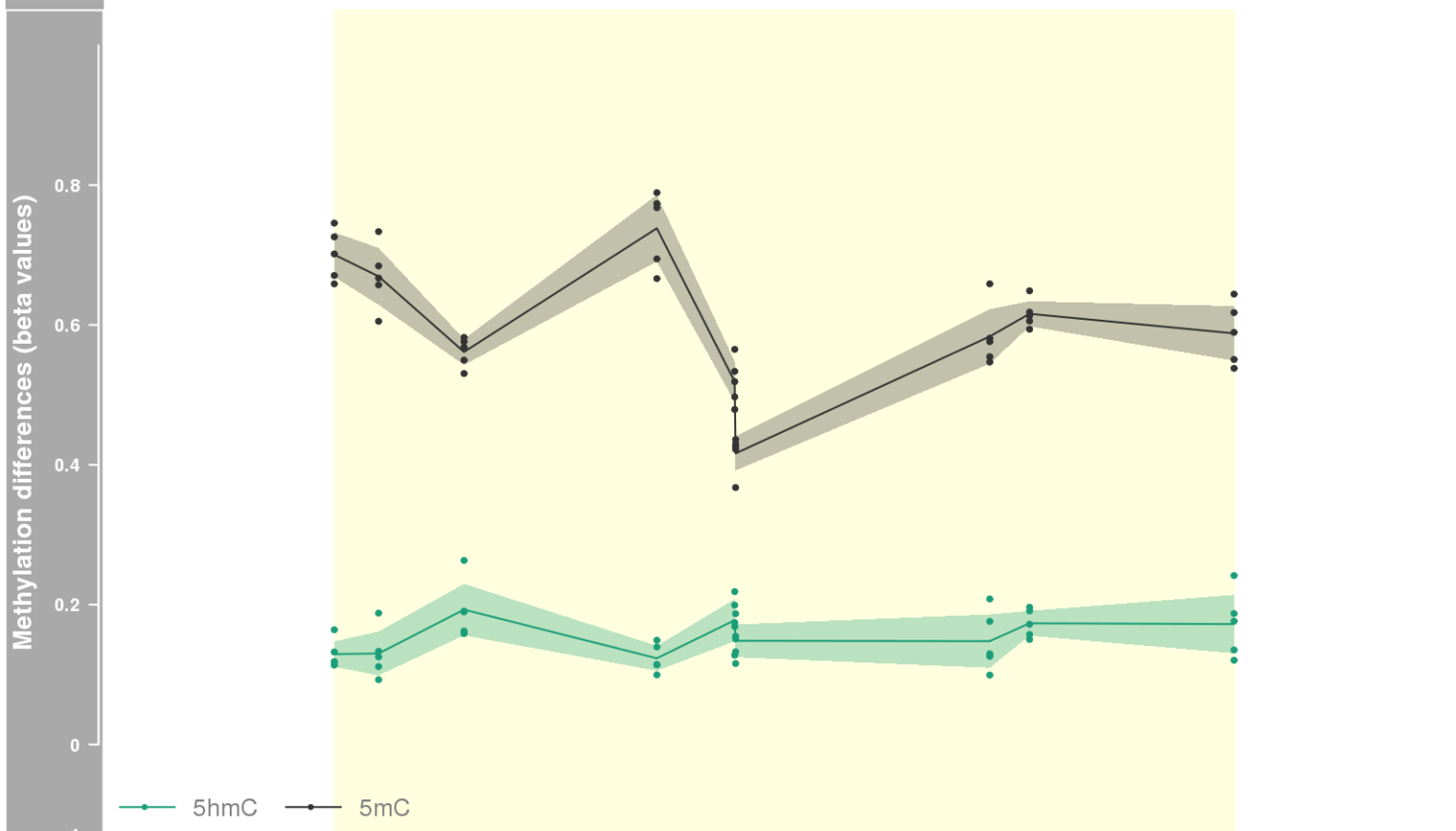
DMR 408 // chr7:2058865-2060130 // 1265 pb. (8 probes) // pvalue: 0.013 // fwer: 0.522 // pvalueArea: 0.028 // fwerArea: 1
- genes: MAD1L1 -



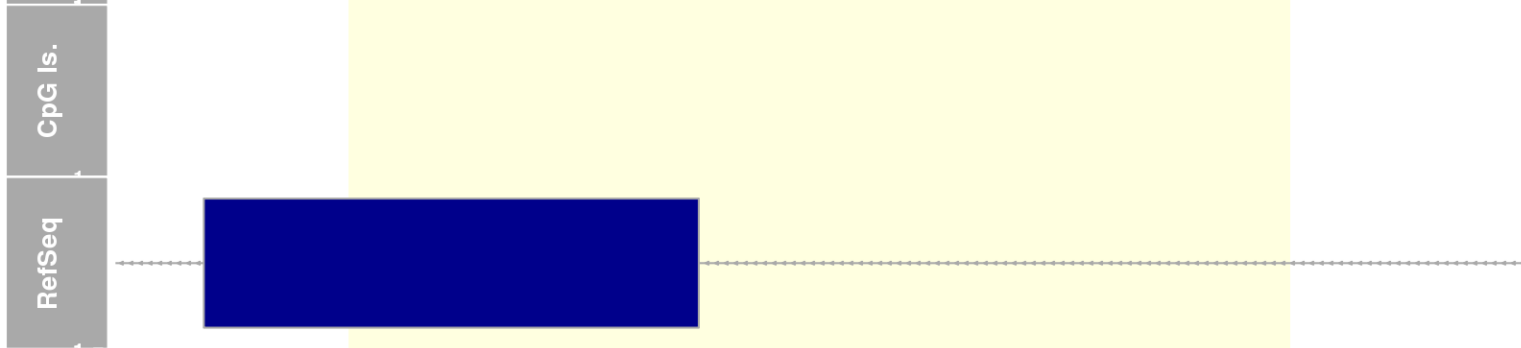
DMR 409 // chr16:67428155-67428898 // 743 pb. (5 probes) // pvalue: 0.001 // fwer: 0.496 // pvalueArea: 0.028 // fwerArea: 1
- genes: ZDHHC1 -



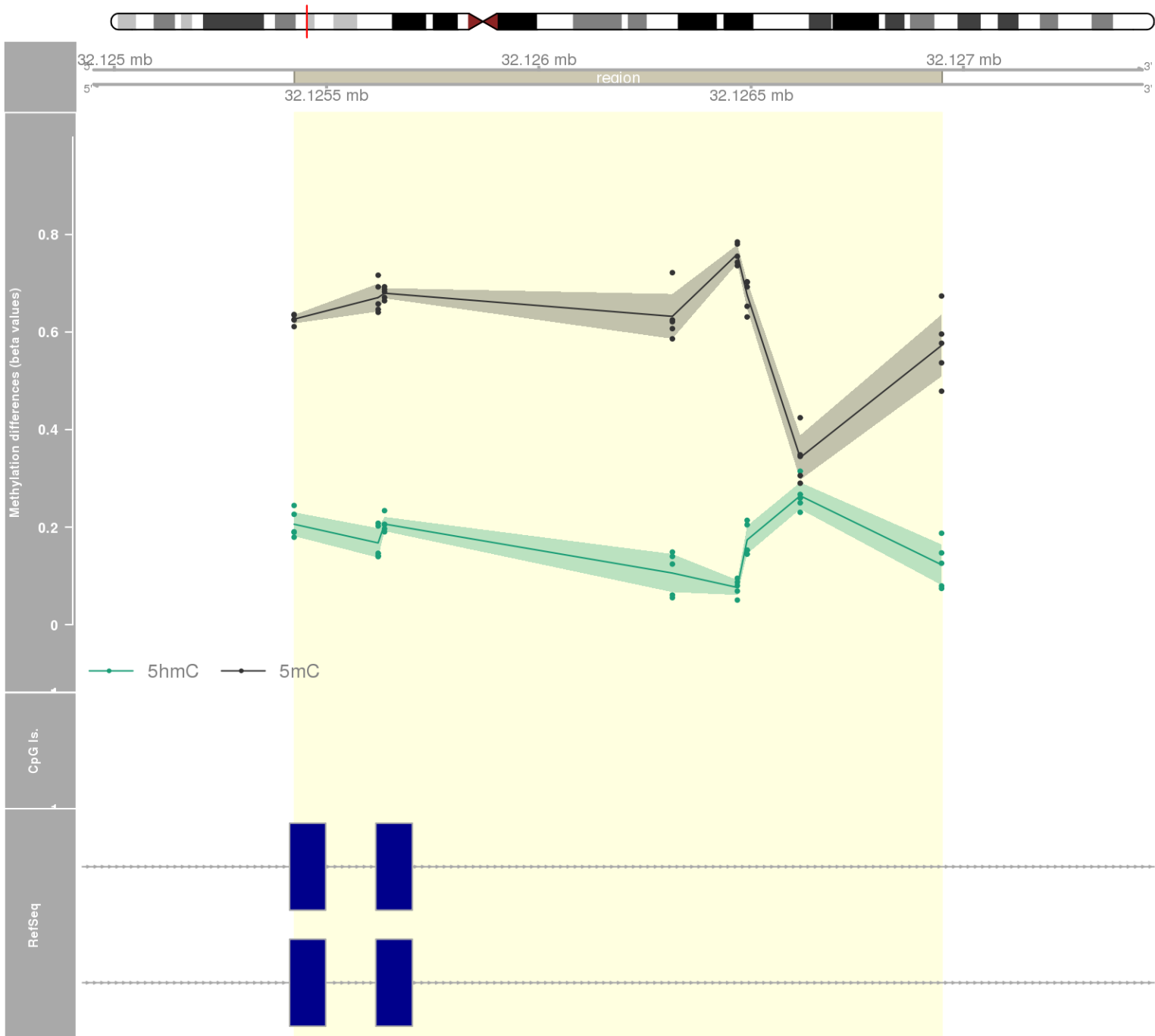
DMR 410 // chr10:125805493-125807504 // 2011 pb. (9 probes) // pvalue: 0.015 // fwer: 0.36 // pvalueArea: 0.029 // fwerArea: 1
- genes: CHST15 -



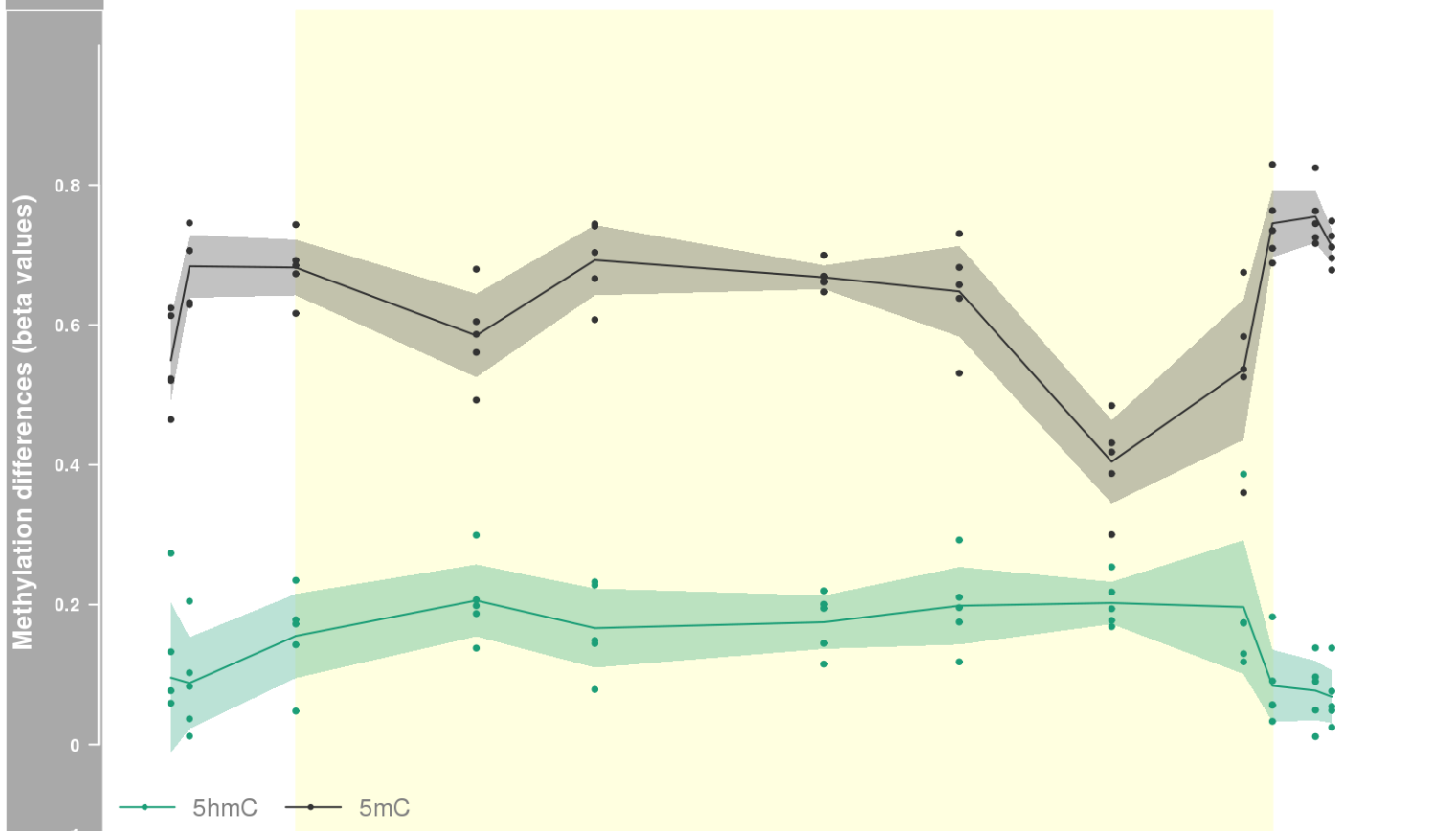
5hmC 5mC



DMR 411 // chr6:32125424-32126949 // 1525 pb. (8 probes) // pvalue: 0.014 // fwer: 0.522 // pvalueArea: 0.029 // fwerArea: 1
- genes: PPT2 / PPT2-EGFL8 -



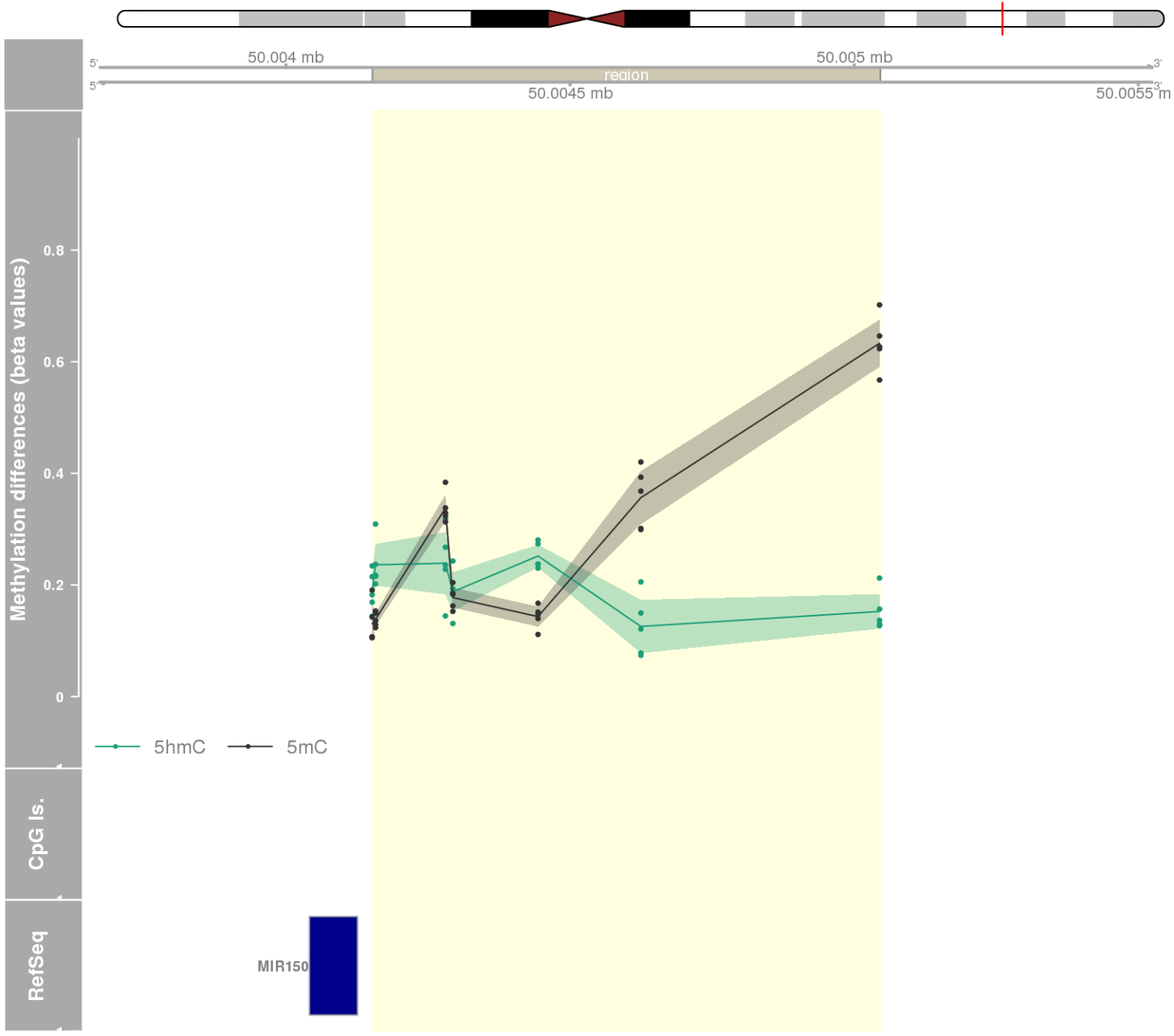
DMR 412 // chr13:113425536-113428172 // 2636 pb. (8 probes) // pvalue: 0.014 // fwer: 0.522 // pvalueArea: 0.029 // fwerArea: 1
- genes: ATP11A -



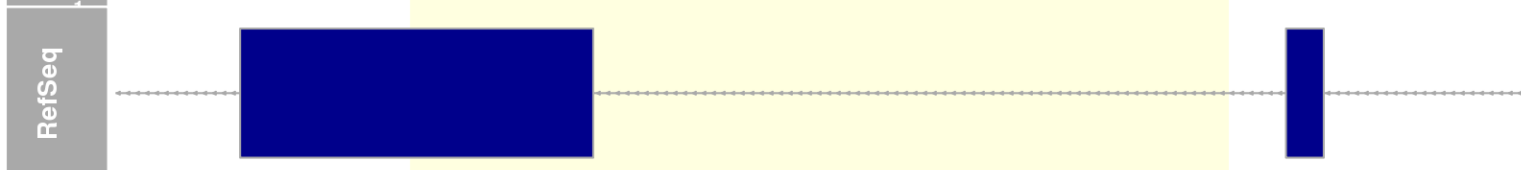
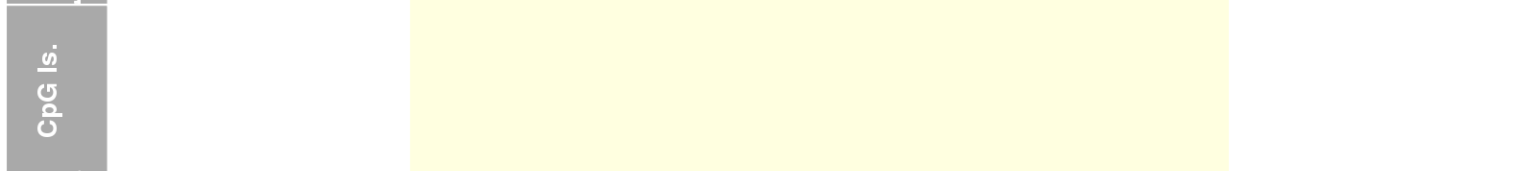
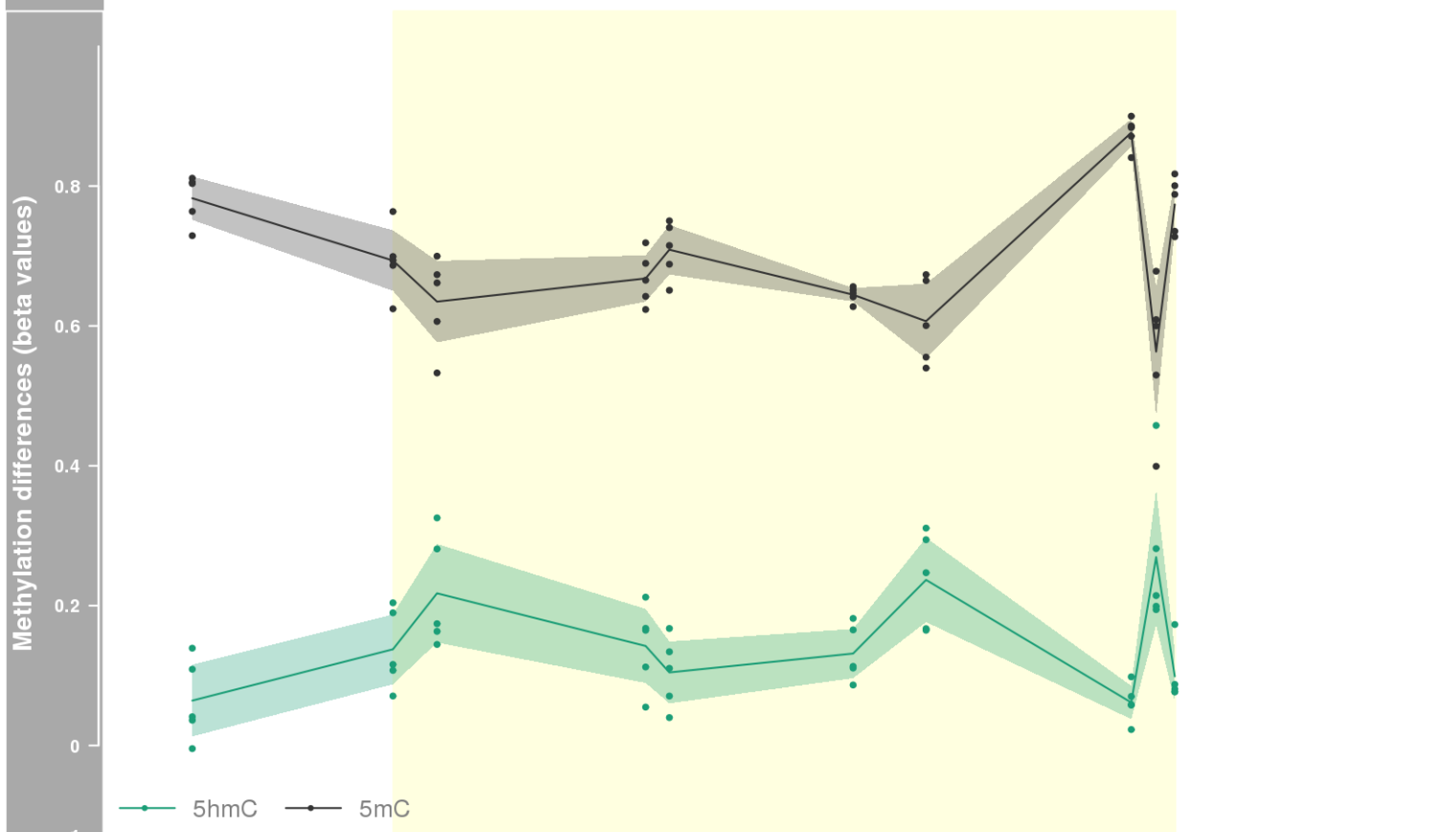
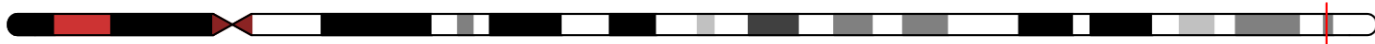
—●— 5hmC —●— 5mC



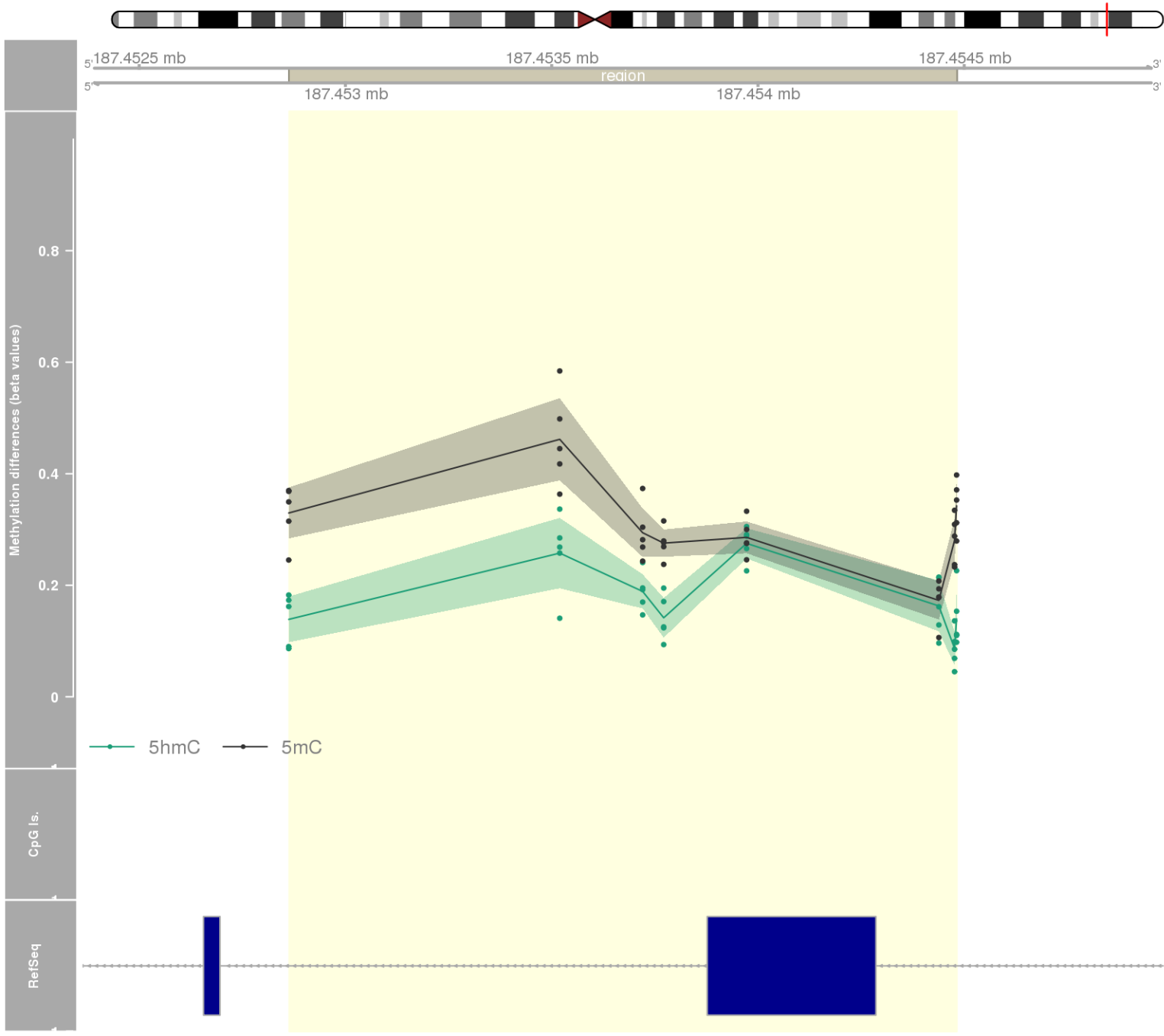
DMR 413 // chr19:50004152-50005045 // 893 pb. (7 probes) // pvalue: 0.009 // fwer: 0.173 // pvalueArea: 0.029 // fwerArea: 1



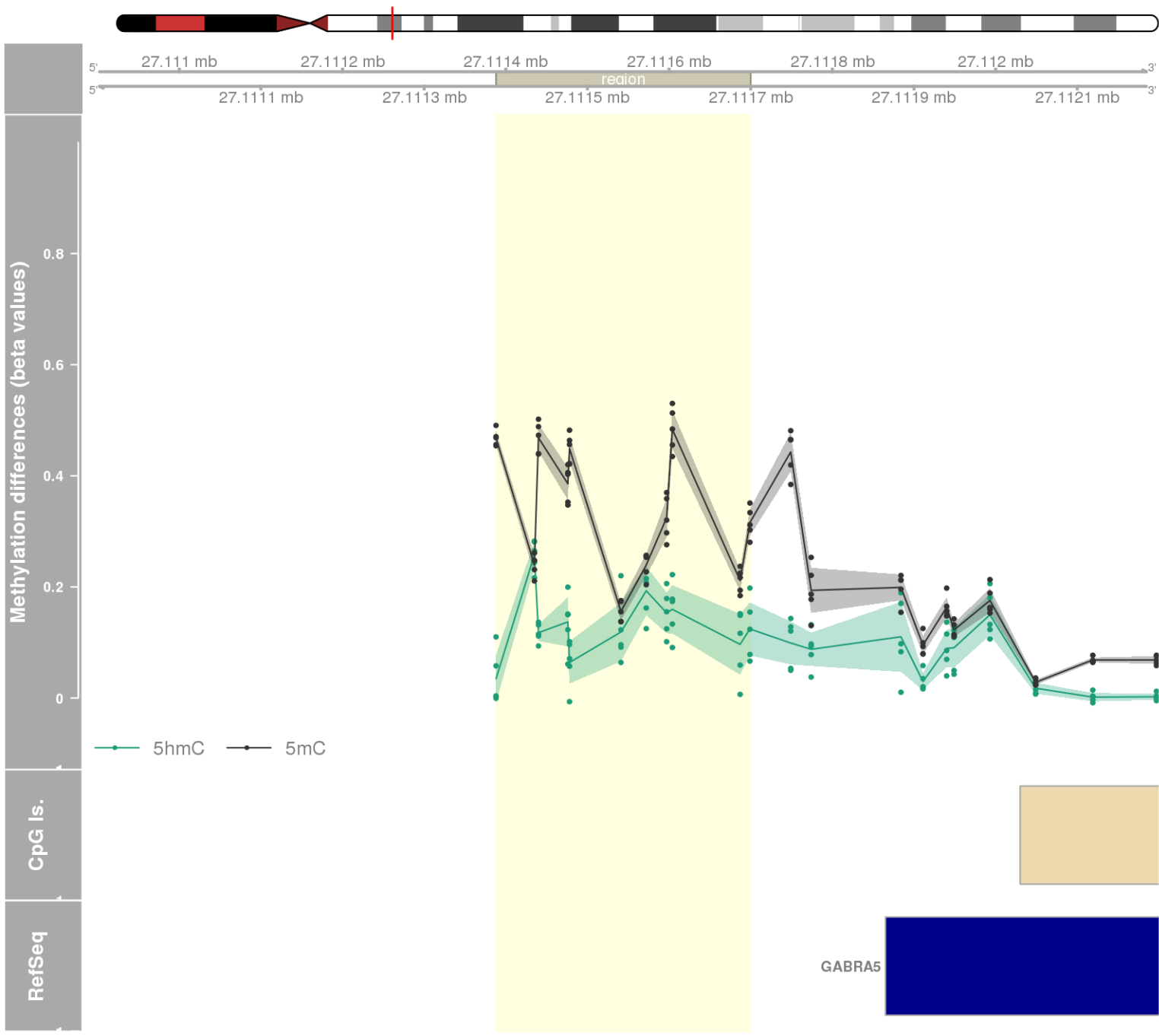
DMR 414 // chr14:103410516-103411899 // 1383 pb. (9 probes) // pvalue: 0.015 // fwer: 0.36 // pvalueArea: 0.029 // fwerArea: 1
- genes: CDC42BPB -



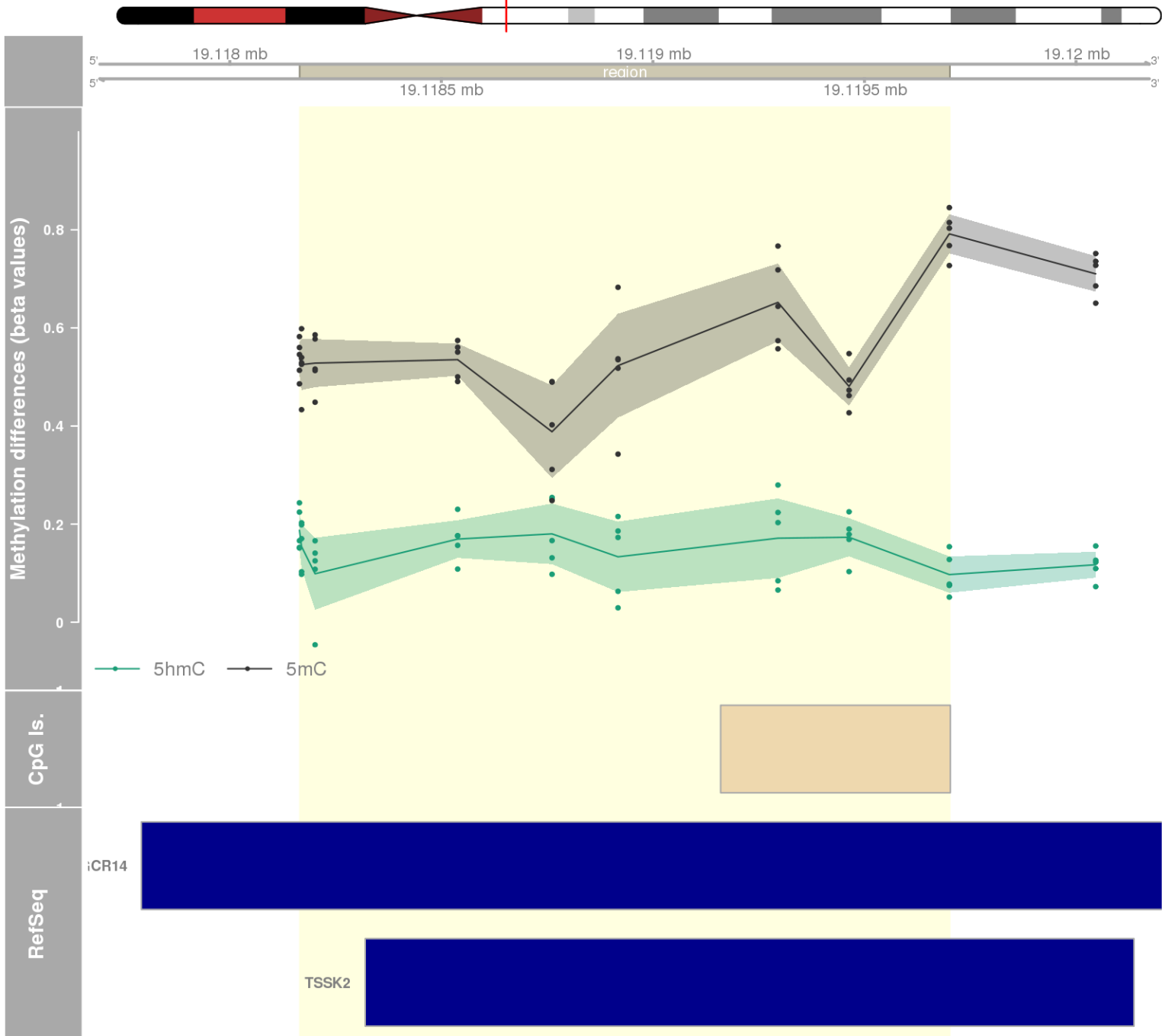
DMR 415 // chr3:187452863-187454482 // 1619 pb. (8 probes) // pvalue: 0.014 // fwer: 0.522 // pvalueArea: 0.029 // fwerArea: 1
- genes: BCL6 -



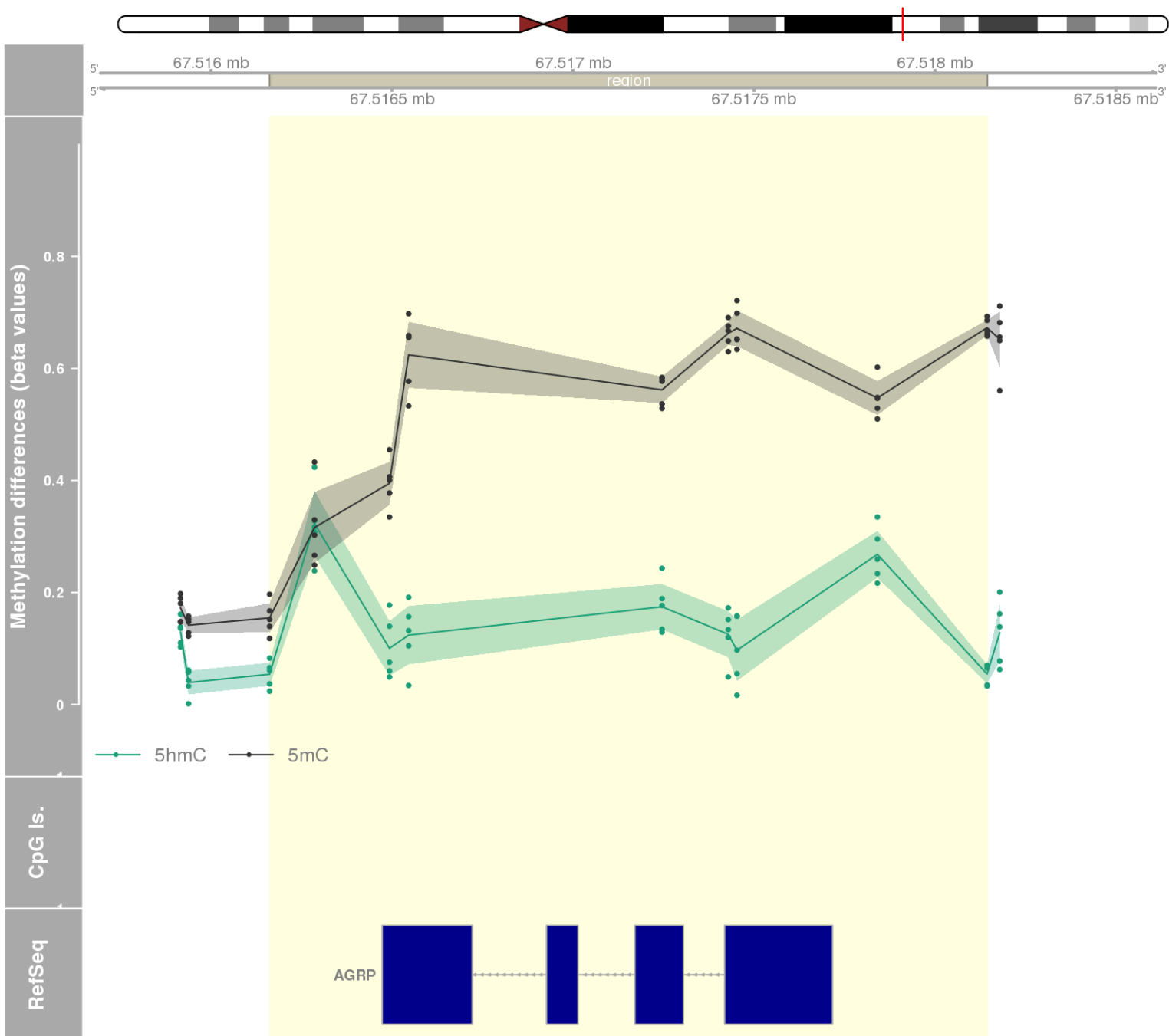
DMR 416 // chr15:27111388-27111699 // 311 pb. (11 probes) // pvalue: 0.011 // fwer: 0.522 // pvalueArea: 0.029 // fwerArea: 1
- genes: GABRB3 / GABRA5 -



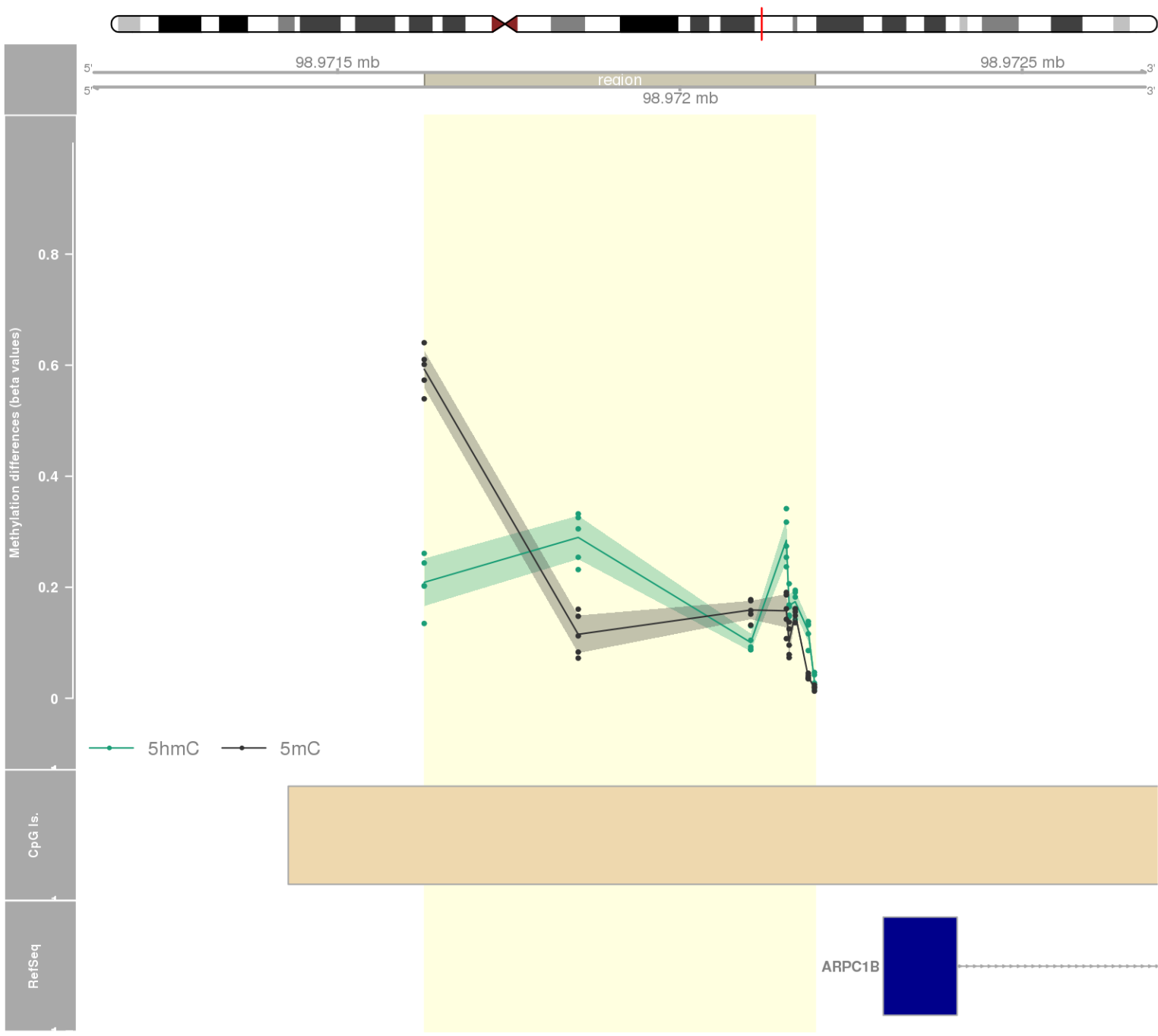
DMR 417 // chr22:19118165-19119701 // 1536 pb. (9 probes) // pvalue: 0.016 // fwer: 0.36 // pvalueArea: 0.029 // fwerArea: 1
- genes: DGCR14 / TSSK2 -



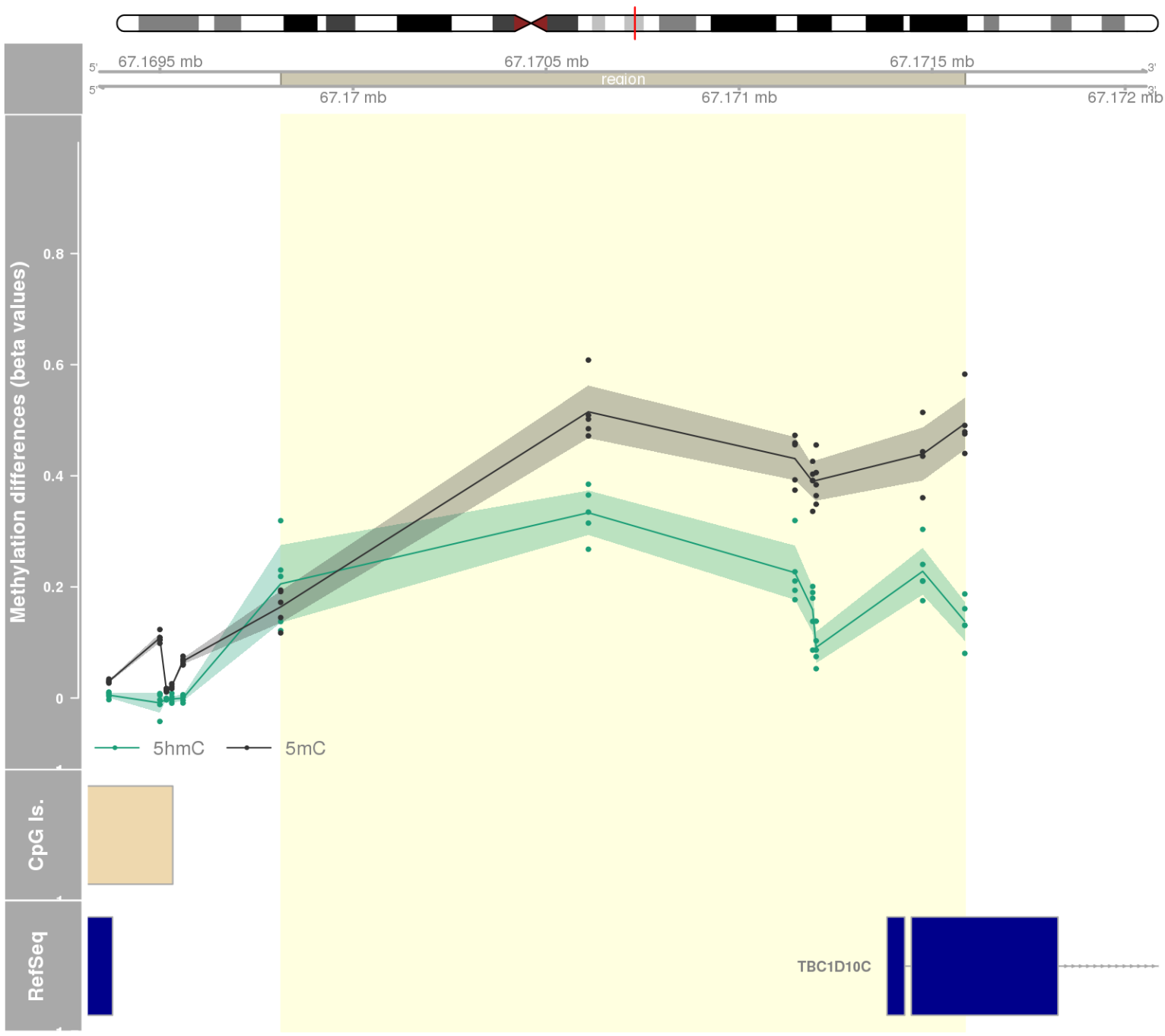
DMR 418 // chr16:67516162-67518144 // 1982 pb. (9 probes) // pvalue: 0.016 // fwer: 0.36 // pvalueArea: 0.029 // fwerArea: 1
- genes: AGRP -



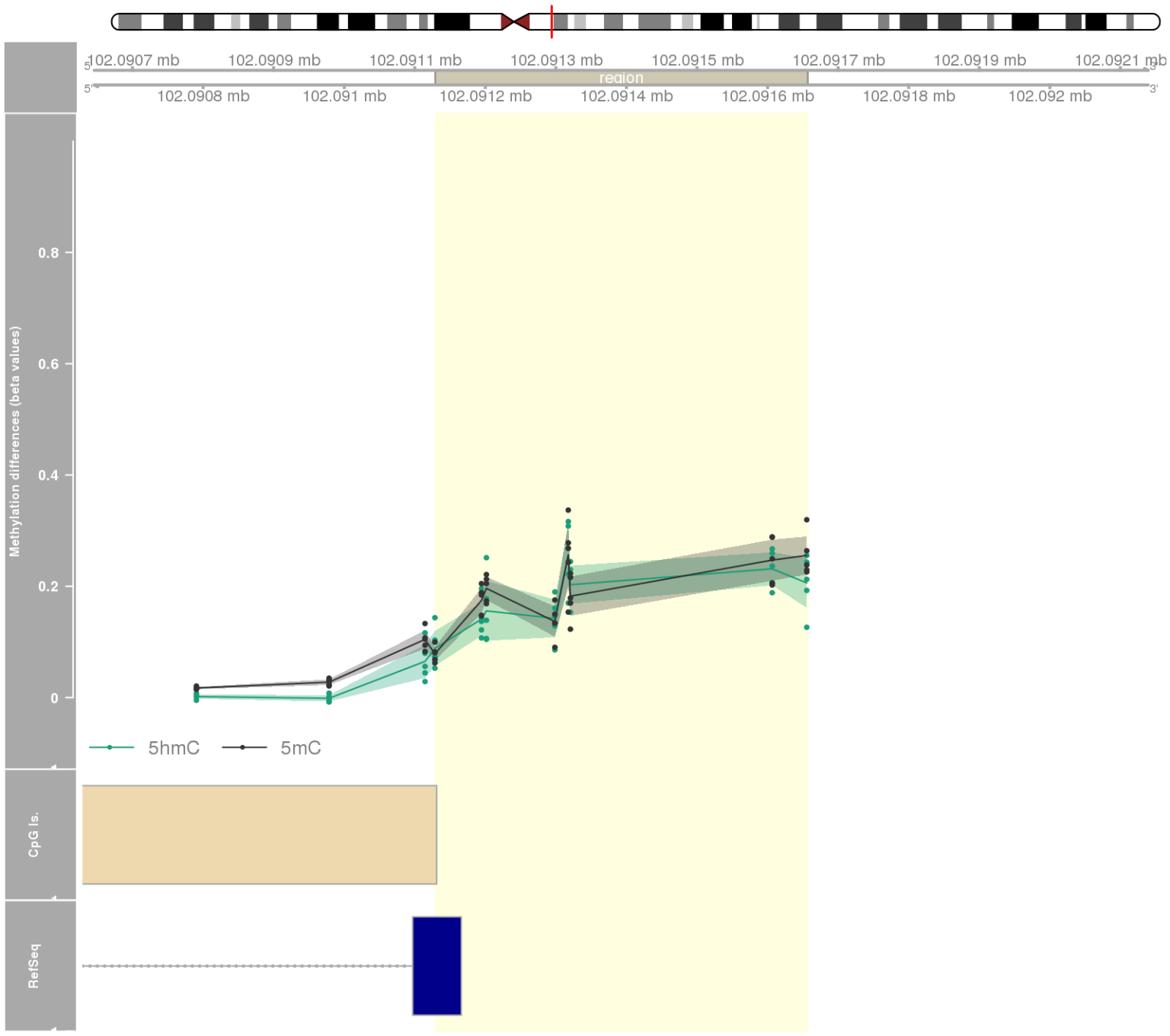
DMR 419 // chr7:98971627-98972197 // 570 pb. (8 probes) // pvalue: 0.014 // fwer: 0.522 // pvalueArea: 0.029 // fwerArea: 1
- genes: ARPC1A / ARPC1B -



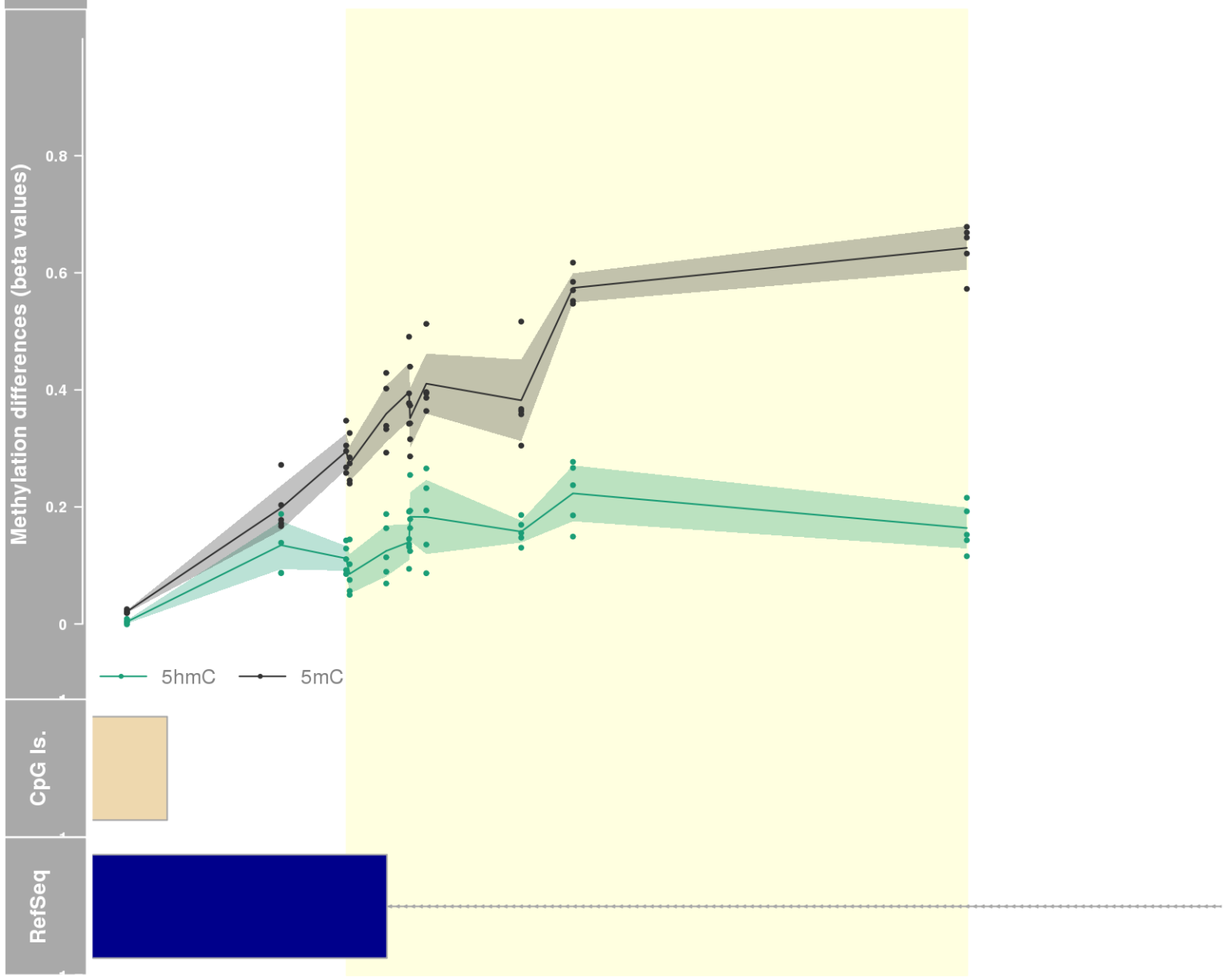
DMR 420 // chr11:67169813-67171585 // 1772 pb. (7 probes) // pvalue: 0.009 // fwer: 0.173 // pvalueArea: 0.029 // fwerArea: 1
- genes: PPP1CA / TBC1D10C -



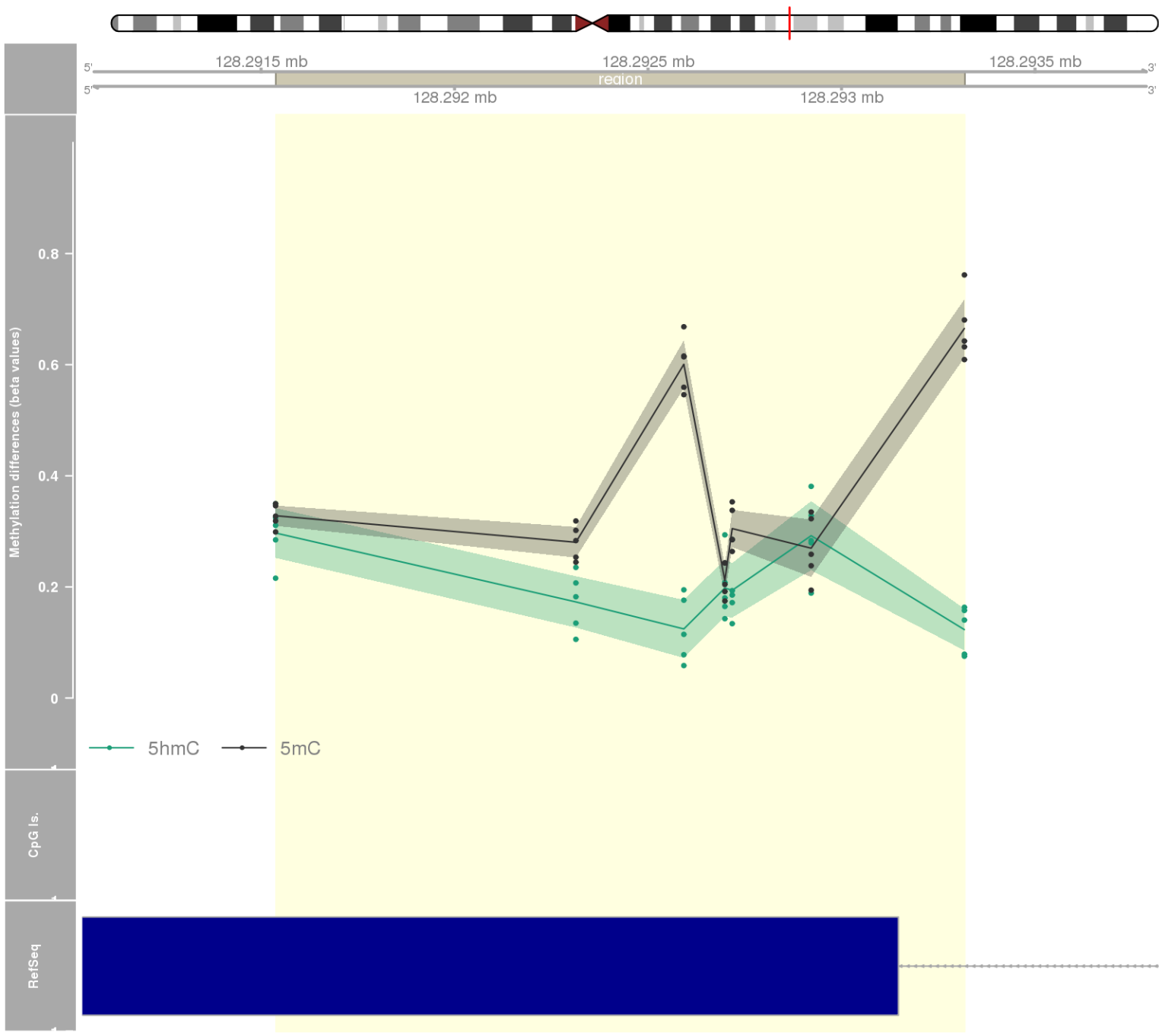
DMR 421 // chr2:102091129-102091656 // 527 pb. (8 probes) // pvalue: 0.014 // fwer: 0.522 // pvalueArea: 0.029 // fwerArea: 1
- genes: RFX8 -



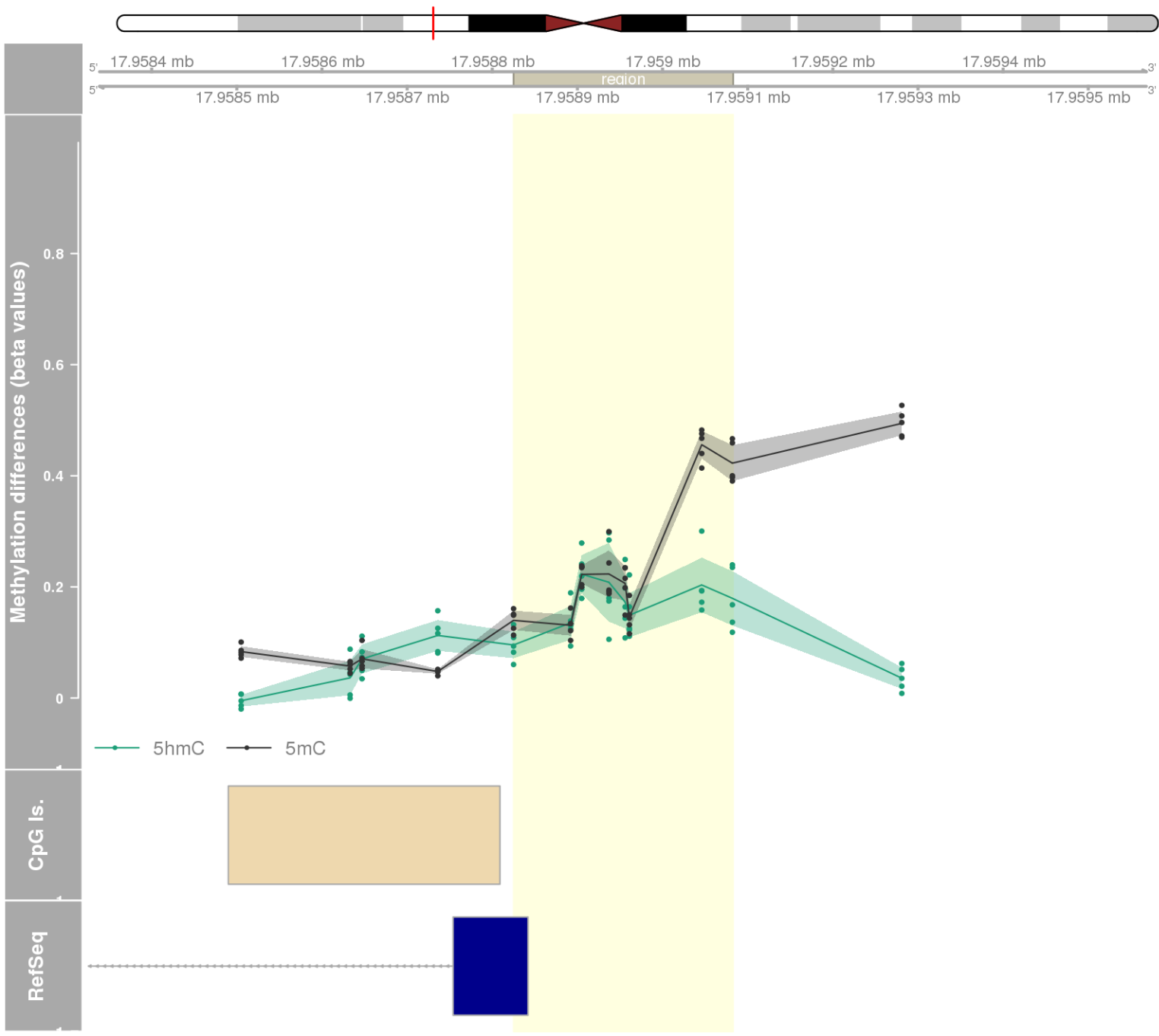
DMR 422 // chr17:19648846-19650069 // 1223 pb. (9 probes) // pvalue: 0.016 // fwer: 0.36 // pvalueArea: 0.029 // fwerArea: 1
- genes: ALDH3A1 -



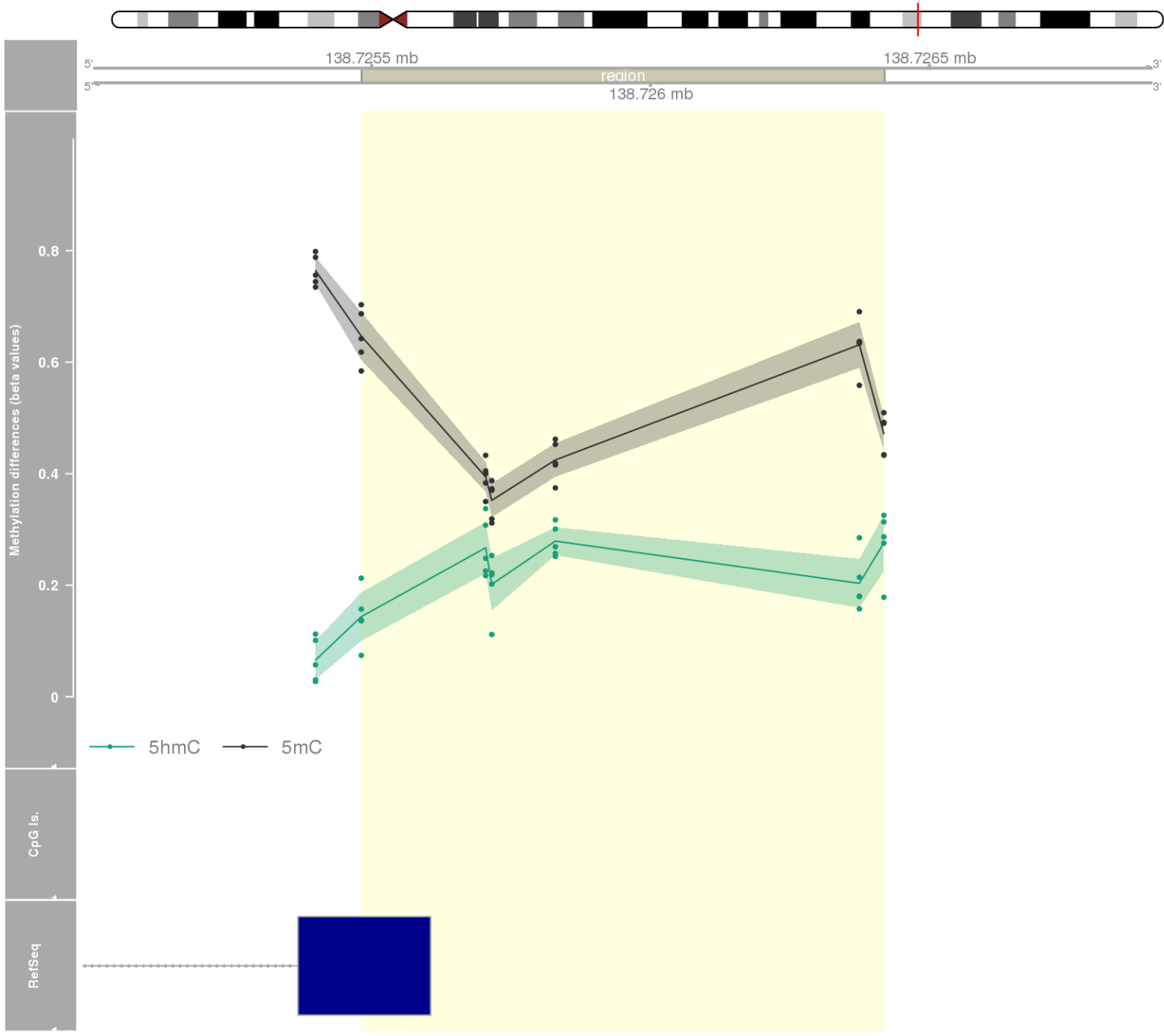
DMR 423 // chr3:128291538-128293318 // 1780 pb. (7 probes) // pvalue: 0.009 // fwer: 0.173 // pvalueArea: 0.029 // fwerArea: 1
- genes: C3orf27 -



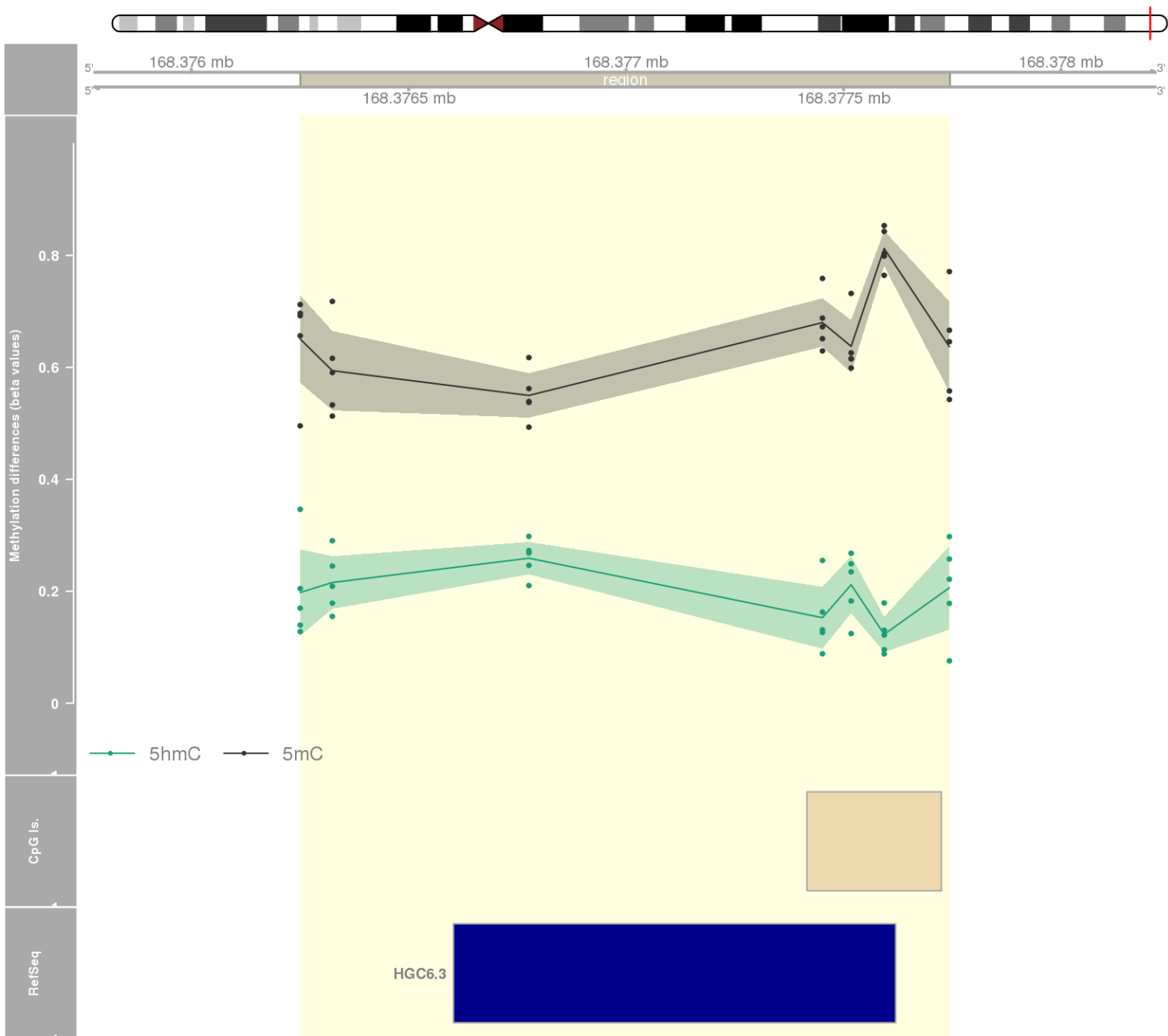
DMR 424 // chr19:17958825-17959082 // 257 pb. (8 probes) // pvalue: 0.015 // fwer: 0.522 // pvalueArea: 0.029 // fwerArea: 1
- genes: JAK3 -



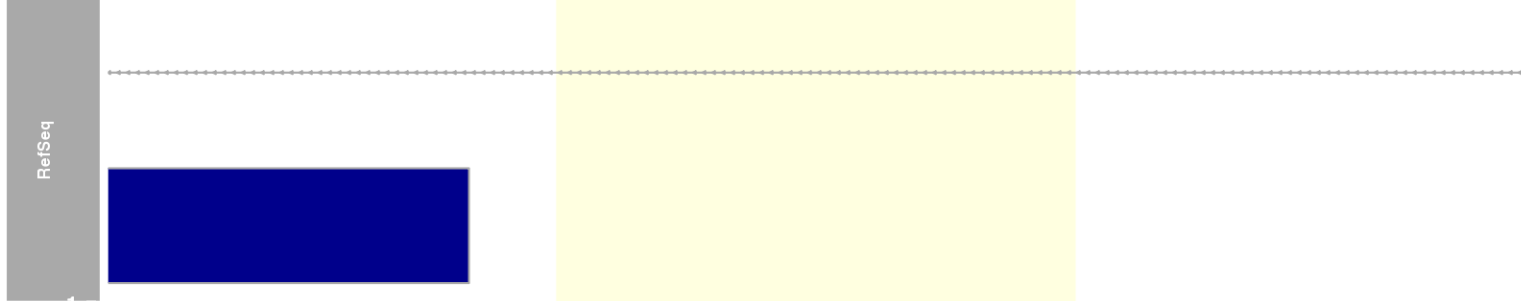
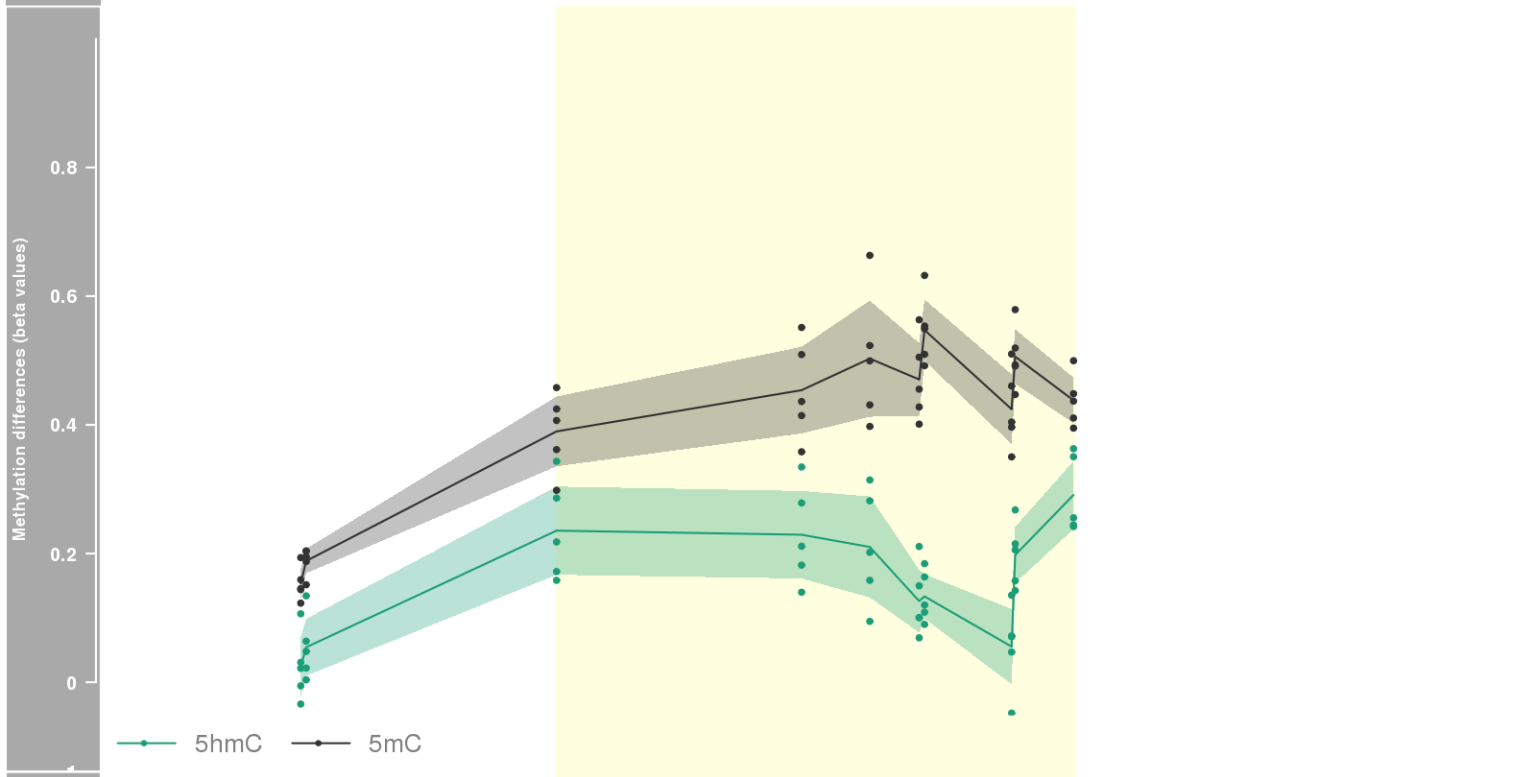
DMR 425 // chr5:138725482-138726419 // 937 pb. (6 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.03 // fwerArea: 1
- genes: MZB1 -



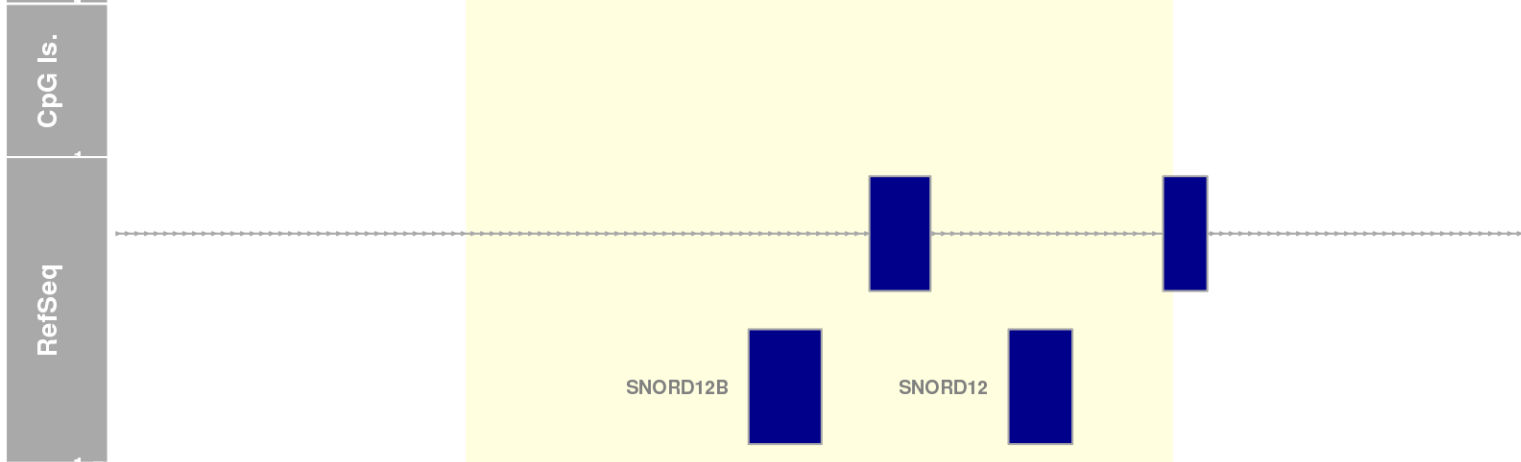
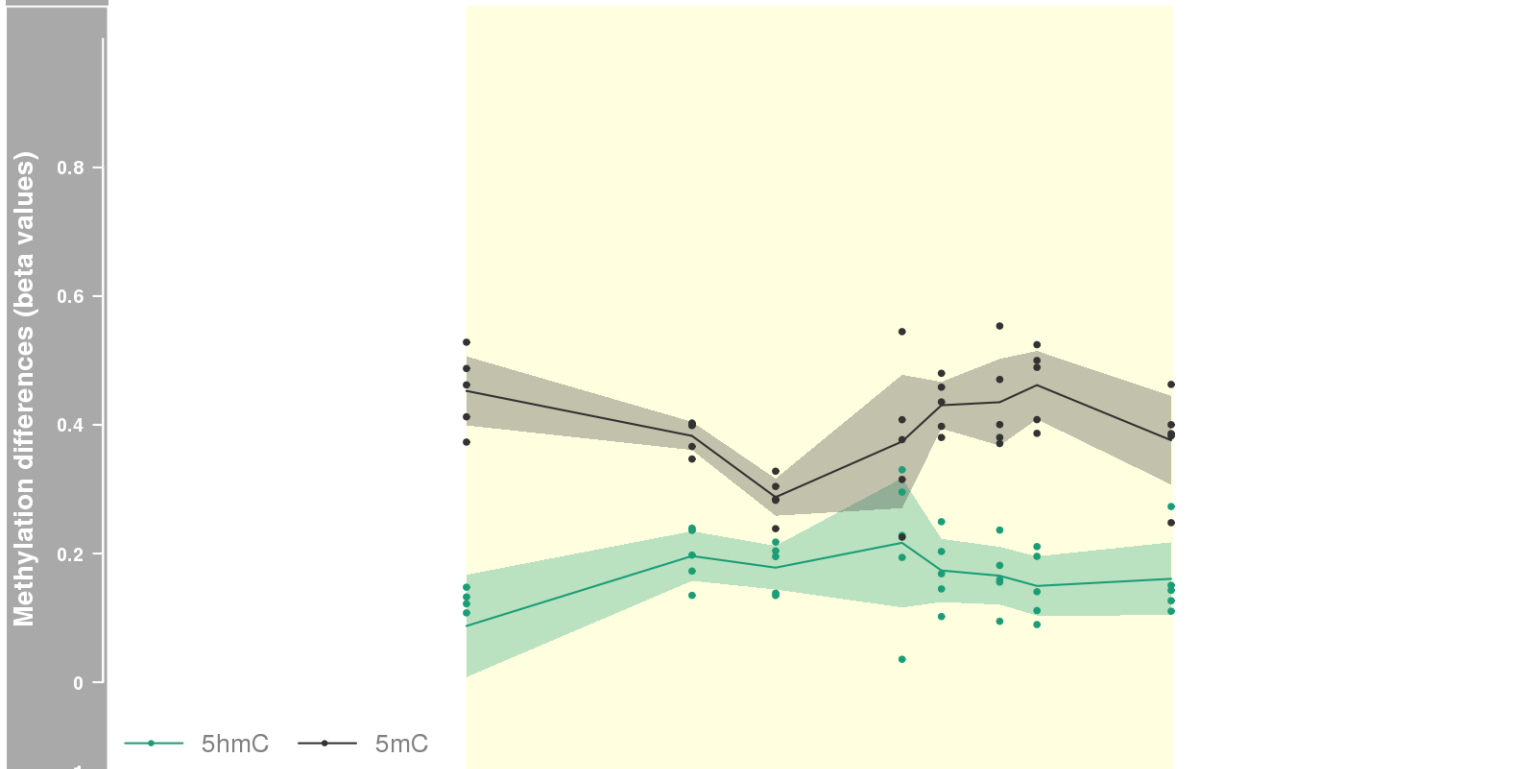
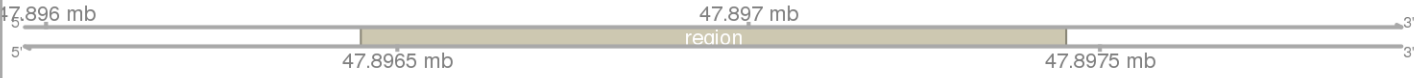
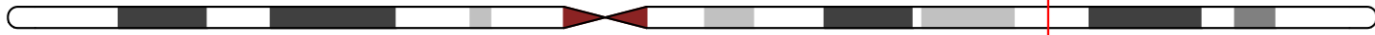
DMR 426 // chr6:168376251-168377744 // 1493 pb. (7 probes) // pvalue: 0.01 // fwer: 0.173 // pvalueArea: 0.03 // fwerArea: 1



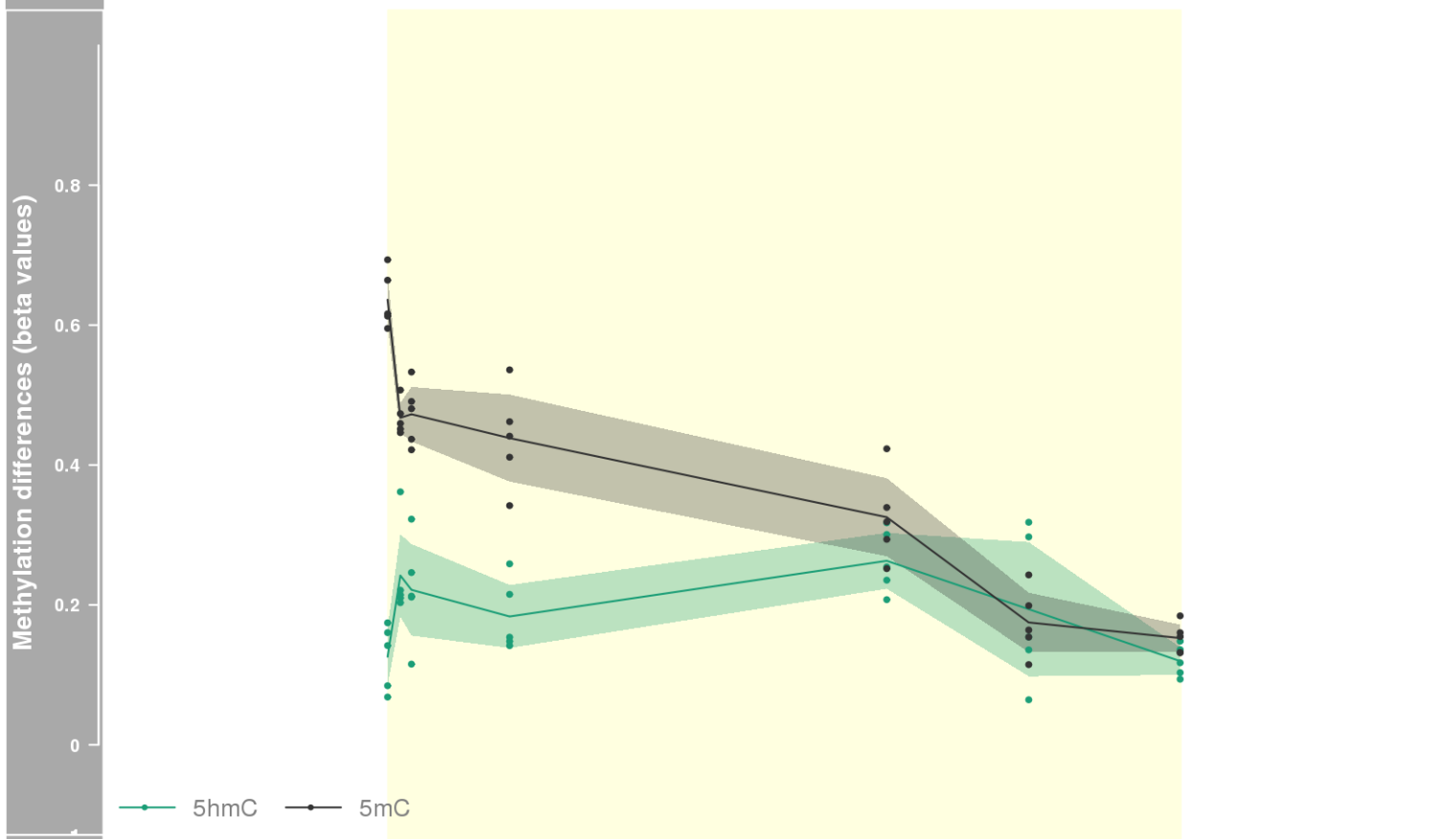
DMR 427 // chr6:31511014-31511590 // 576 pb. (8 probes) // pvalue: 0.015 // fwer: 0.522 // pvalueArea: 0.03 // fwerArea: 1
- genes: ATP6V1G2-DDX39B -



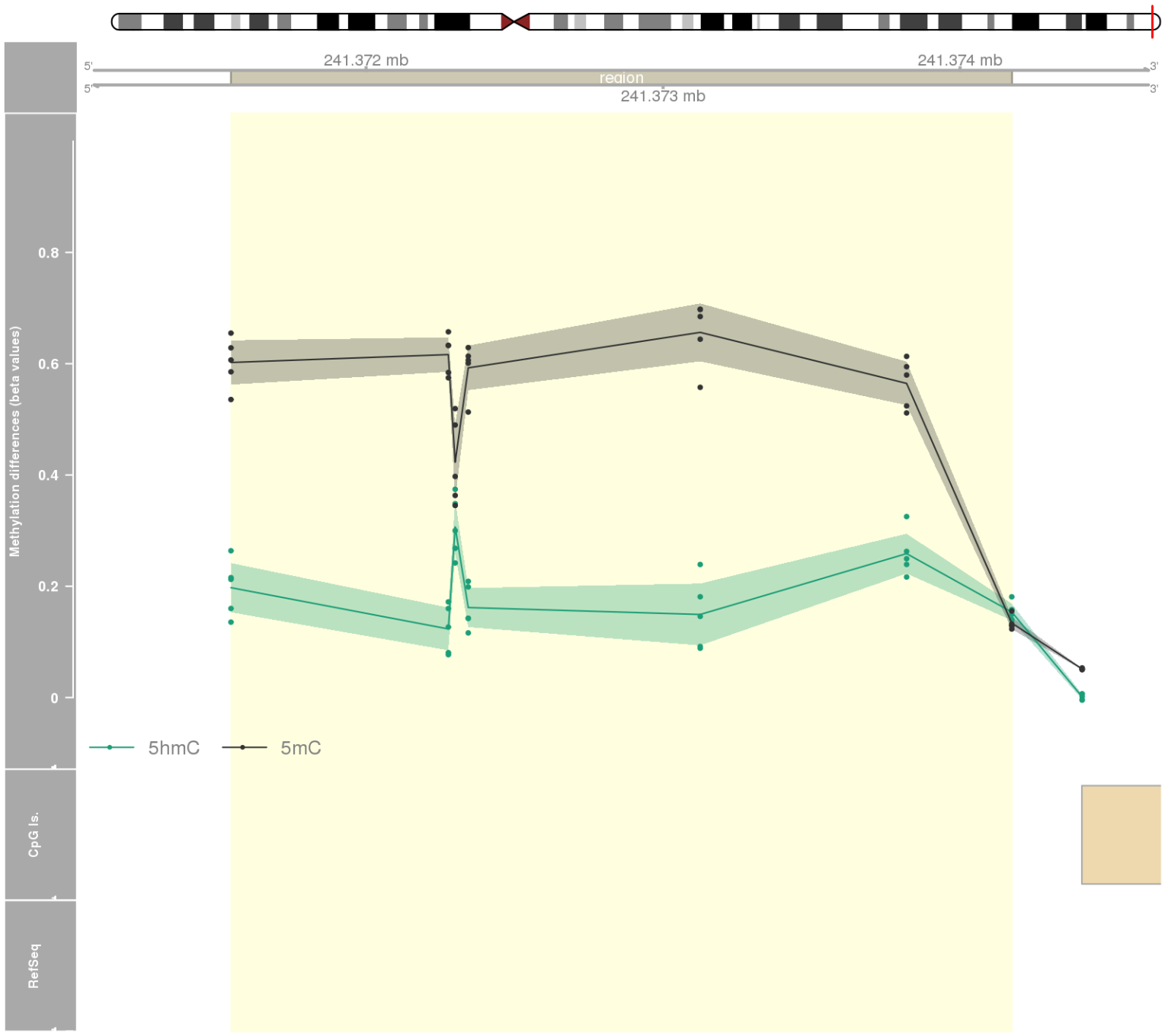
DMR 428 // chr20:47896448-47897451 // 1003 pb. (8 probes) // pvalue: 0.015 // fwer: 0.522 // pvalueArea: 0.03 // fwerArea: 1
- genes: ZFAS1 / SNORD12B / SNORD12 -



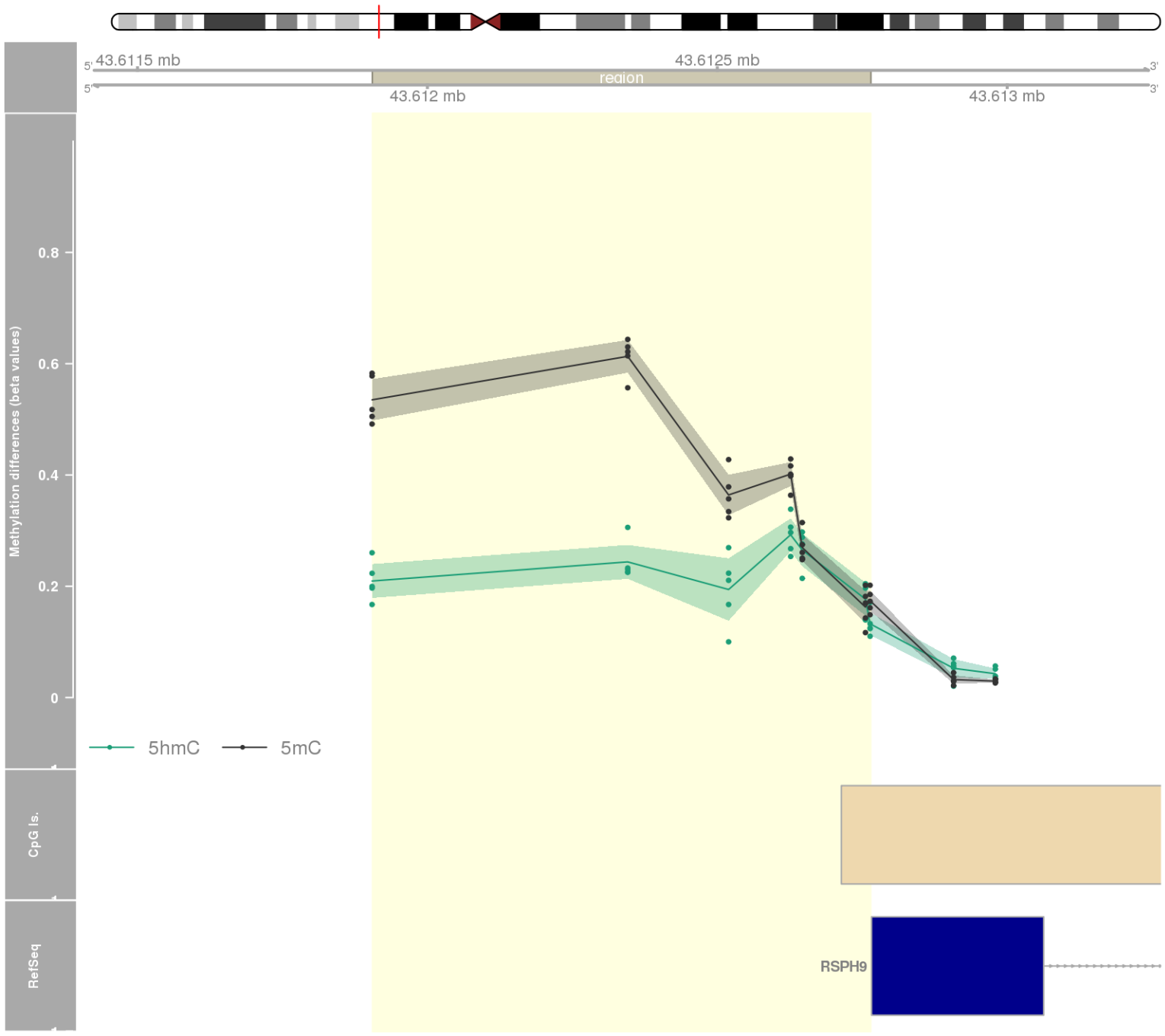
DMR 429 // chr13:88328251-88329680 // 1429 pb. (7 probes) // pvalue: 0.01 // fwer: 0.173 // pvalueArea: 0.03 // fwerArea: 1
- genes: SLITRK5 -



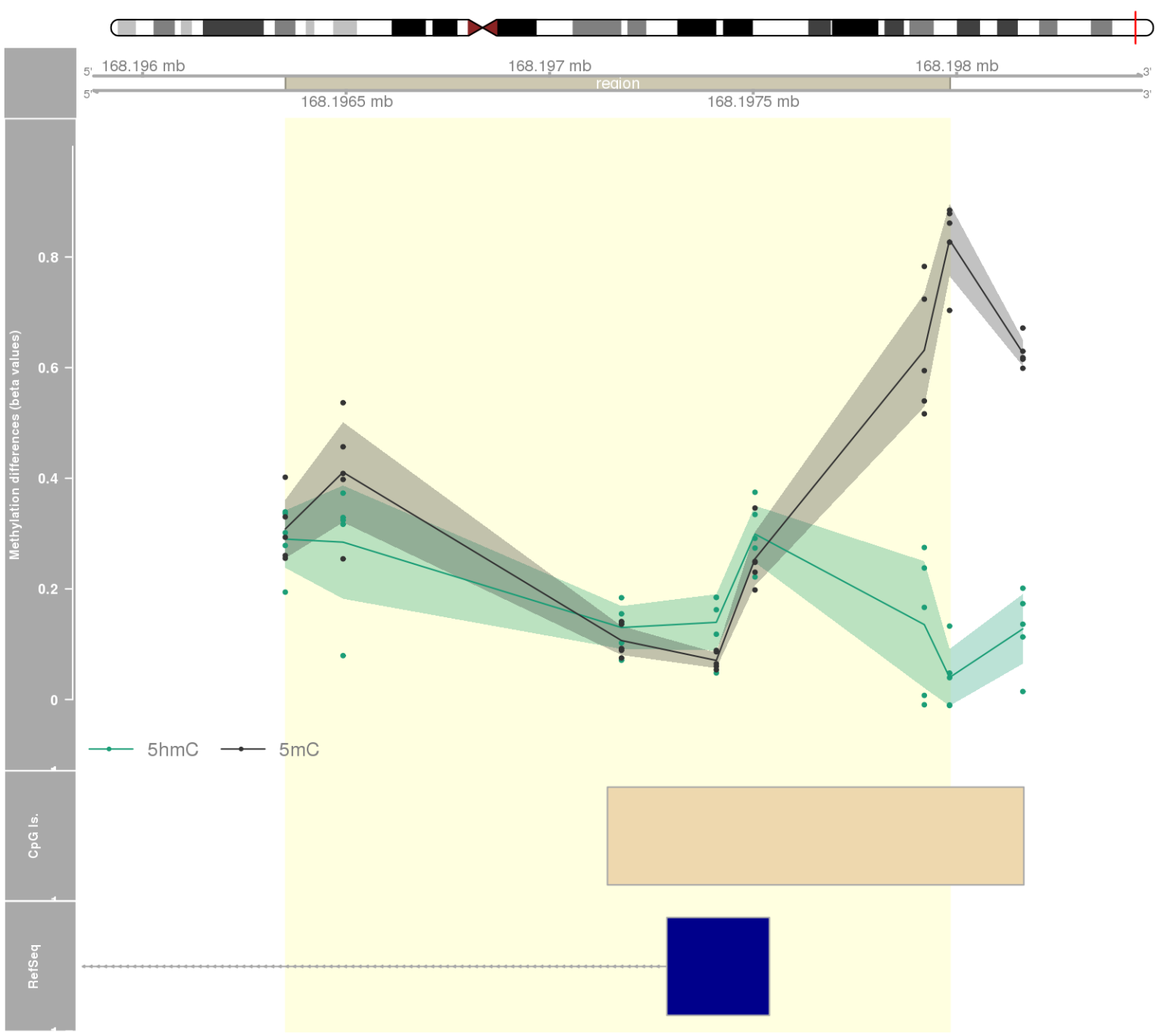
DMR 430 // chr2:241371545-241374174 // 2629 pb. (7 probes) // pvalue: 0.01 // fwer: 0.173 // pvalueArea: 0.03 // fwerArea: 1



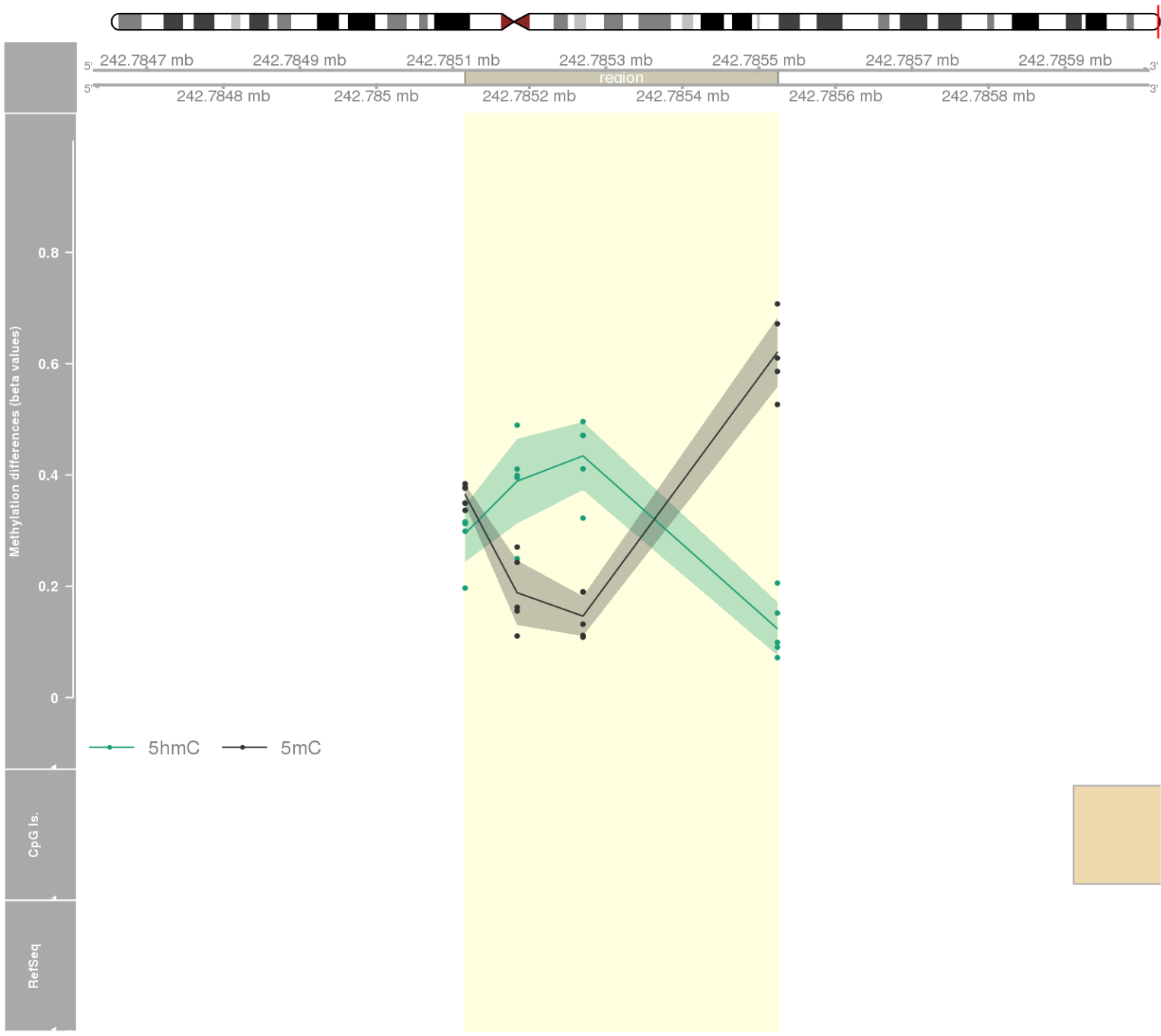
DMR 431 // chr6:43611905-43612764 // 859 pb. (7 probes) // pvalue: 0.01 // fwer: 0.173 // pvalueArea: 0.03 // fwerArea: 1



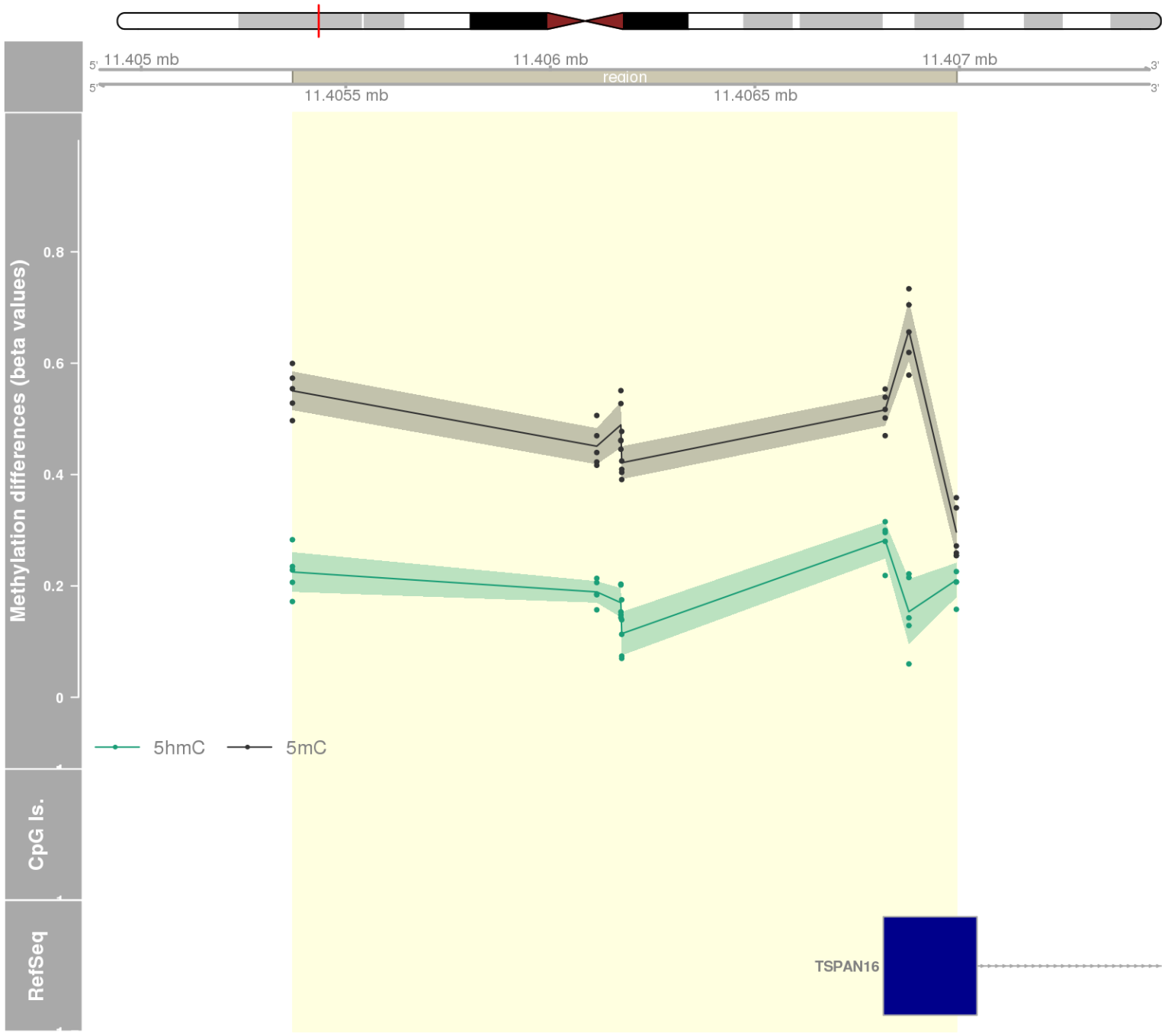
DMR 432 // chr6:168196351-168197983 // 1632 pb. (7 probes) // pvalue: 0.01 // fwer: 0.173 // pvalueArea: 0.03 // fwerArea: 1
- genes: C6orf123 -



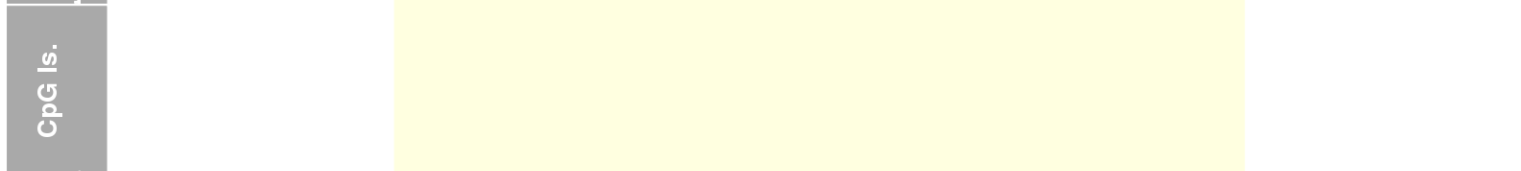
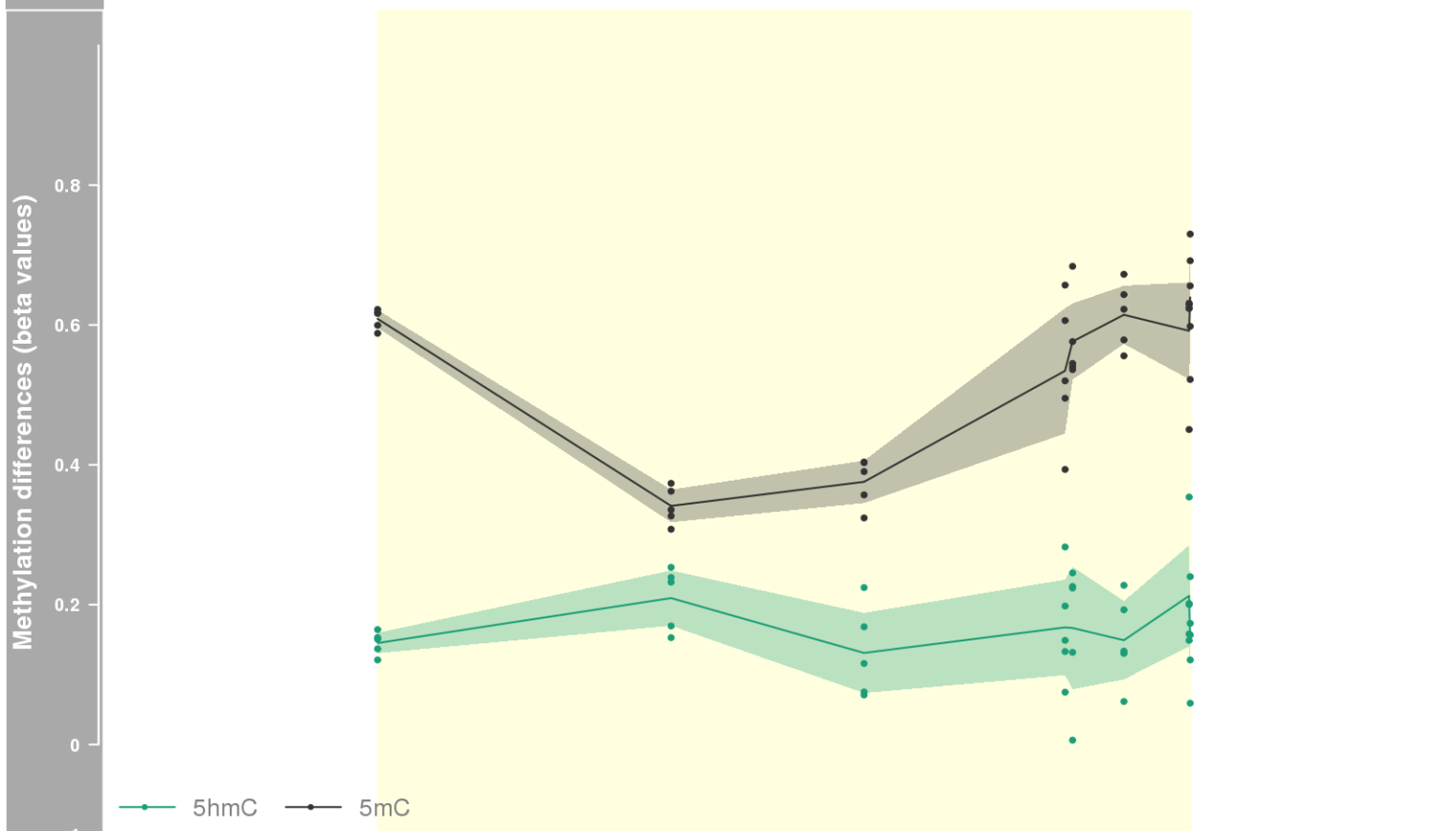
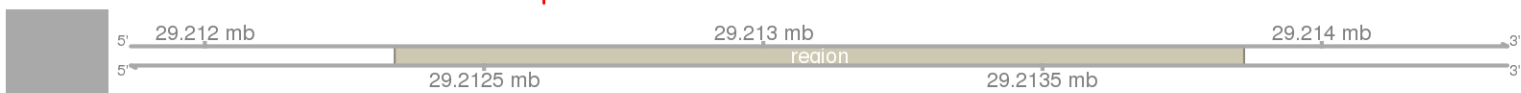
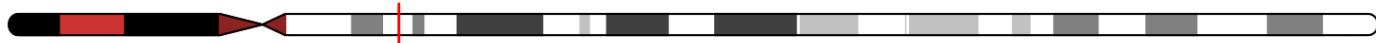
DMR 433 // chr2:242785116-242785524 // 408 pb. (4 probes) // pvalue: 0 // fwer: 0.496 // pvalueArea: 0.03 // fwerArea: 1



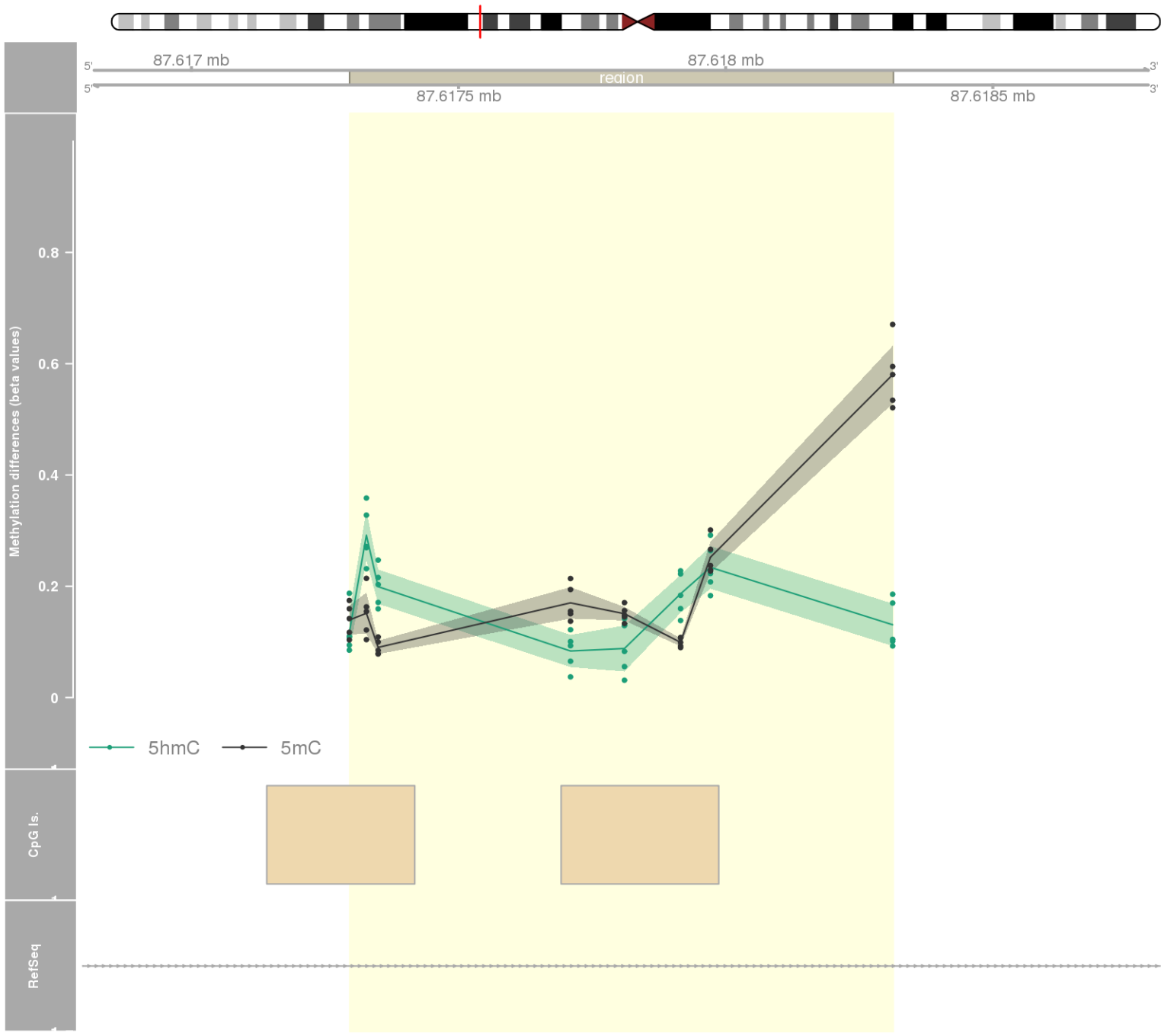
DMR 434 // chr19:11405370-11406993 // 1623 pb. (7 probes) // pvalue: 0.01 // fwer: 0.173 // pvalueArea: 0.03 // fwerArea: 1
- genes: TSPAN16 -



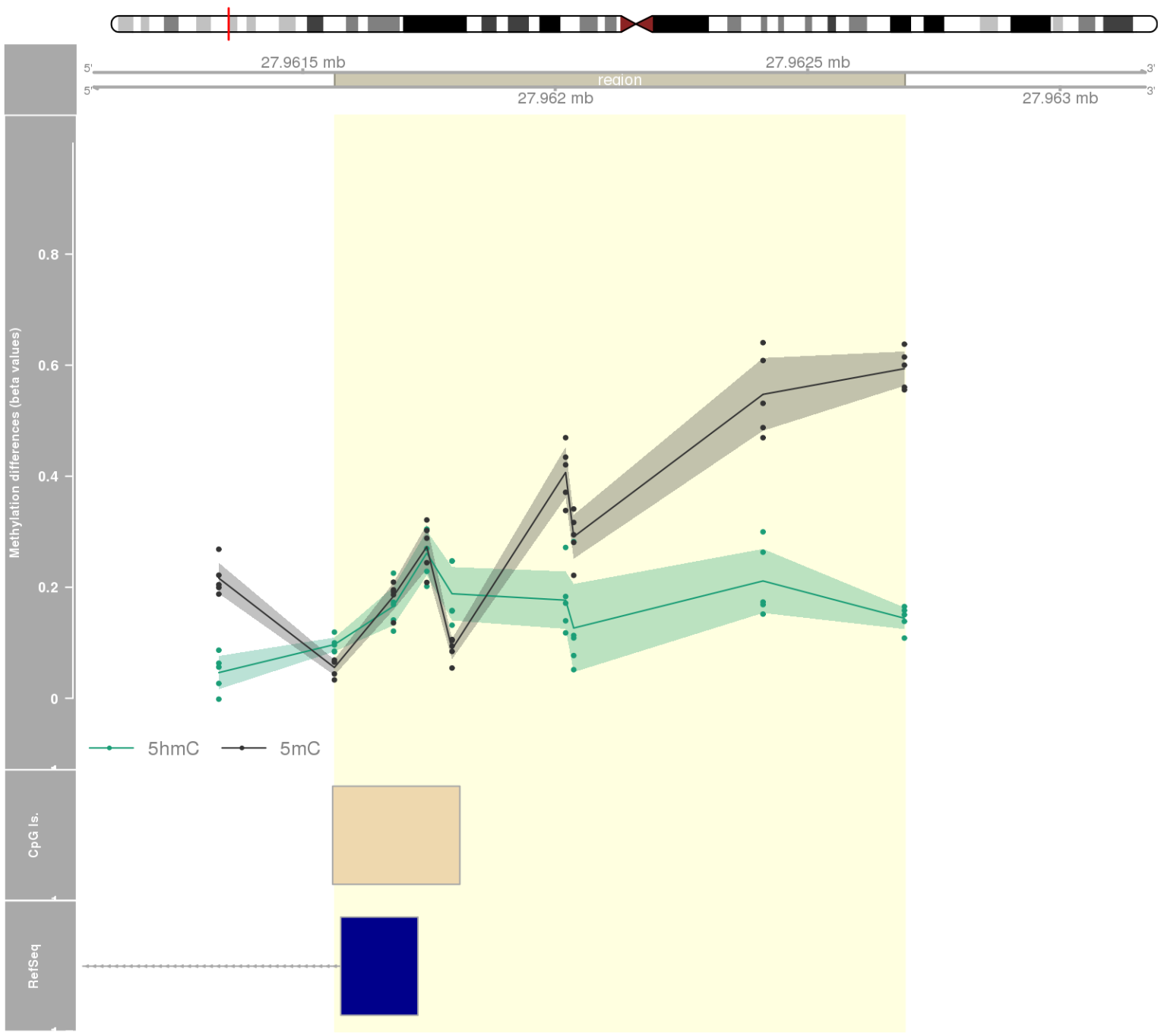
DMR 435 // chr15:29212340-29213860 // 1520 pb. (8 probes) // pvalue: 0.015 // fwer: 0.522 // pvalueArea: 0.03 // fwerArea: 1
- genes: APBA2 -



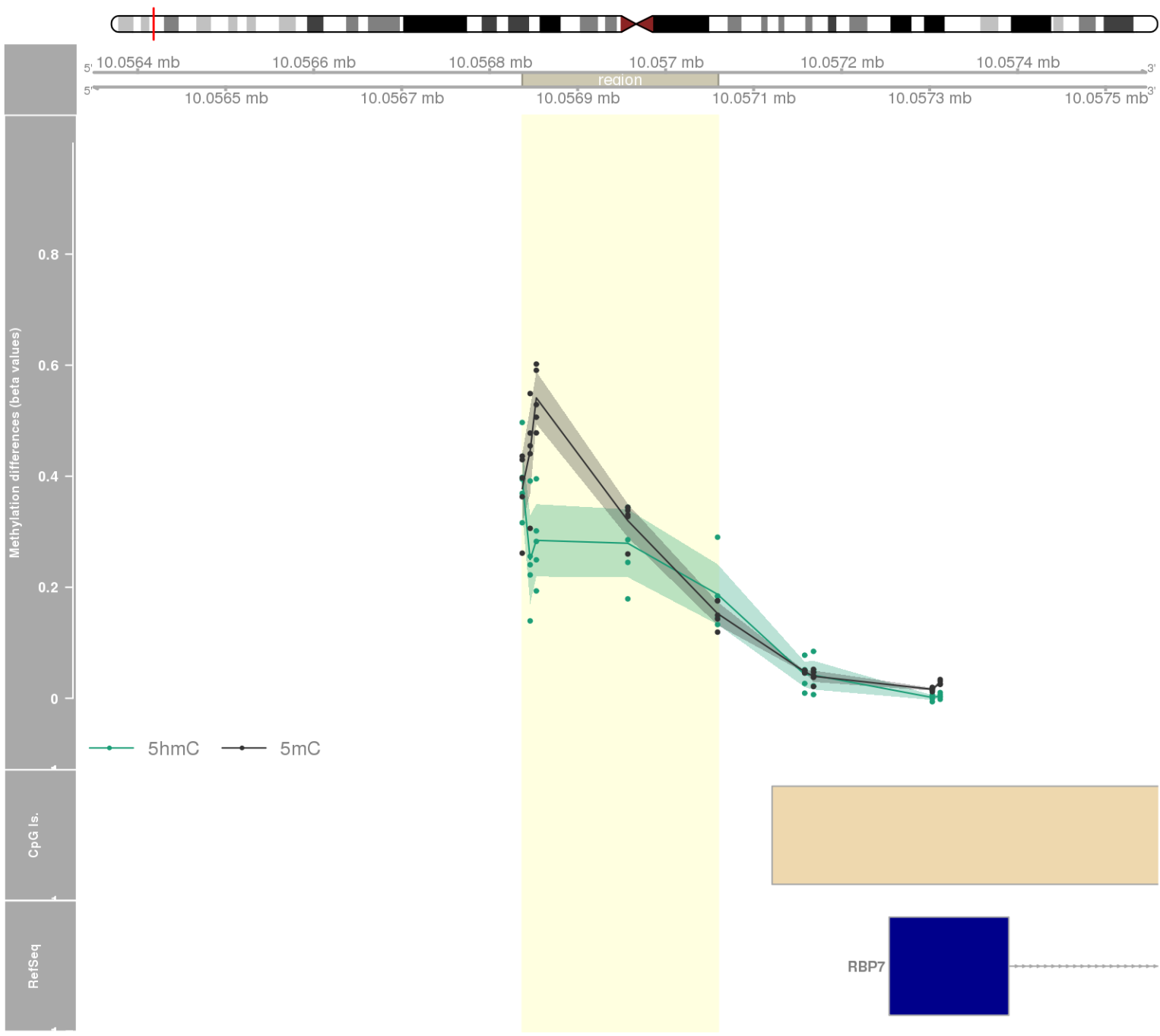
DMR 436 // chr1:87617296-87618313 // 1017 pb. (8 probes) // pvalue: 0.015 // fwer: 0.522 // pvalueArea: 0.03 // fwerArea: 1
- genes: LINC01140 -



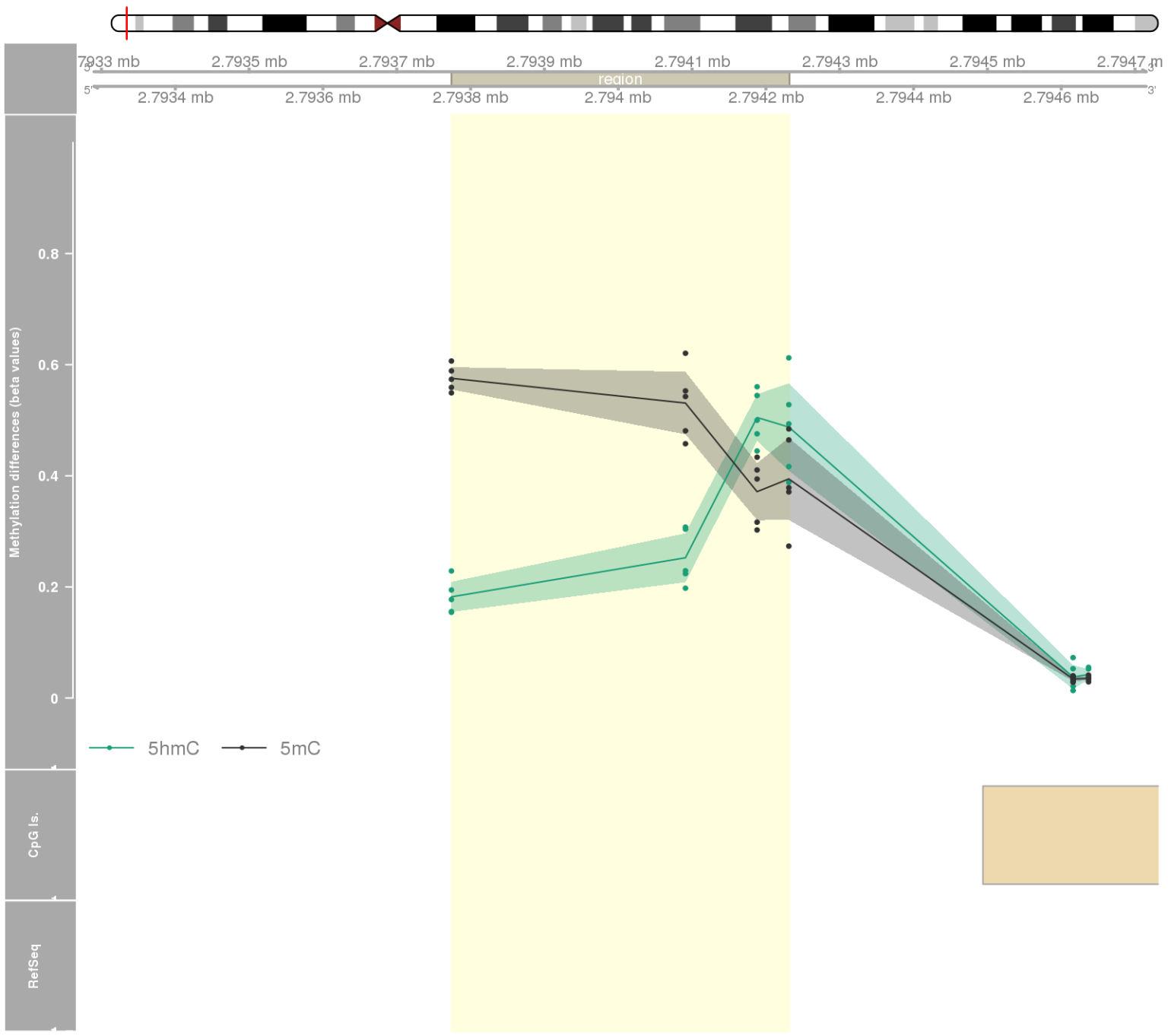
DMR 437 // chr1:27961563-27962692 // 1129 pb. (8 probes) // pvalue: 0.015 // fwer: 0.522 // pvalueArea: 0.03 // fwerArea: 1
- genes: FGR -



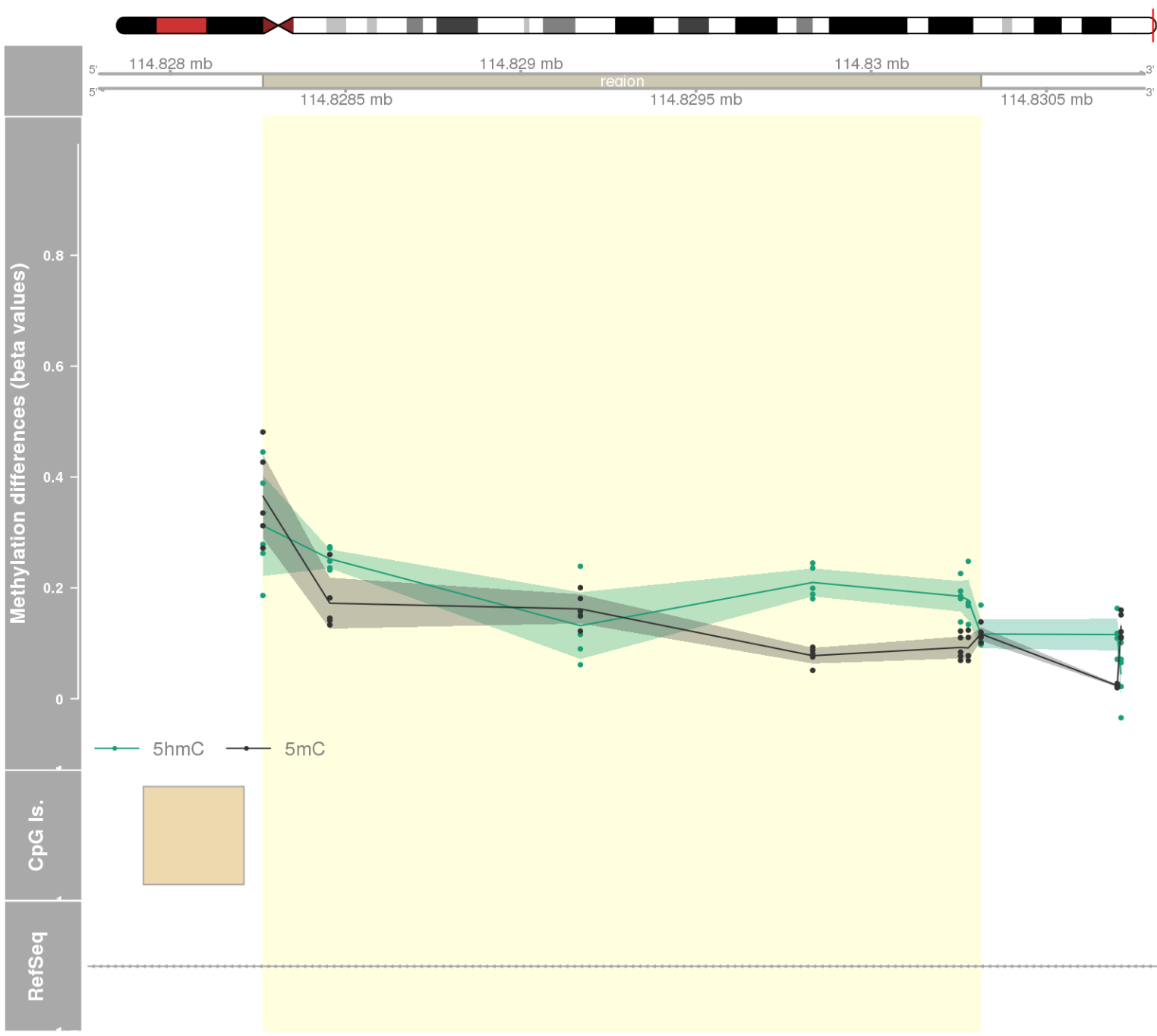
DMR 438 // chr1:10056837-10057059 // 222 pb. (5 probes) // pvalue: 0.001 // fwer: 0.496 // pvalueArea: 0.03 // fwerArea: 1



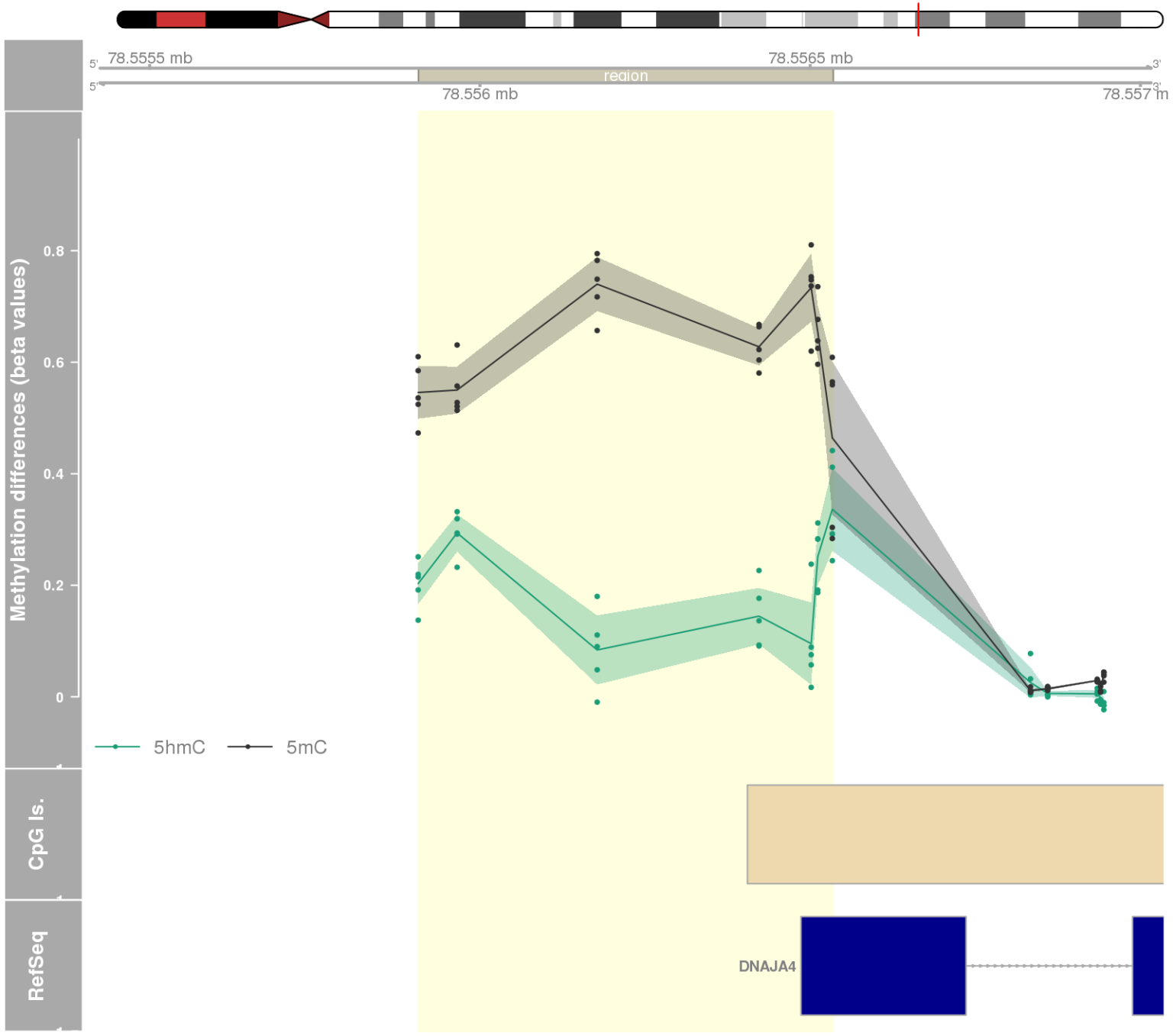
DMR 439 // chr4:2793774-2794231 // 457 pb. (4 probes) // pvalue: 0 // fwer: 0.496 // pvalueArea: 0.031 // fwerArea: 1



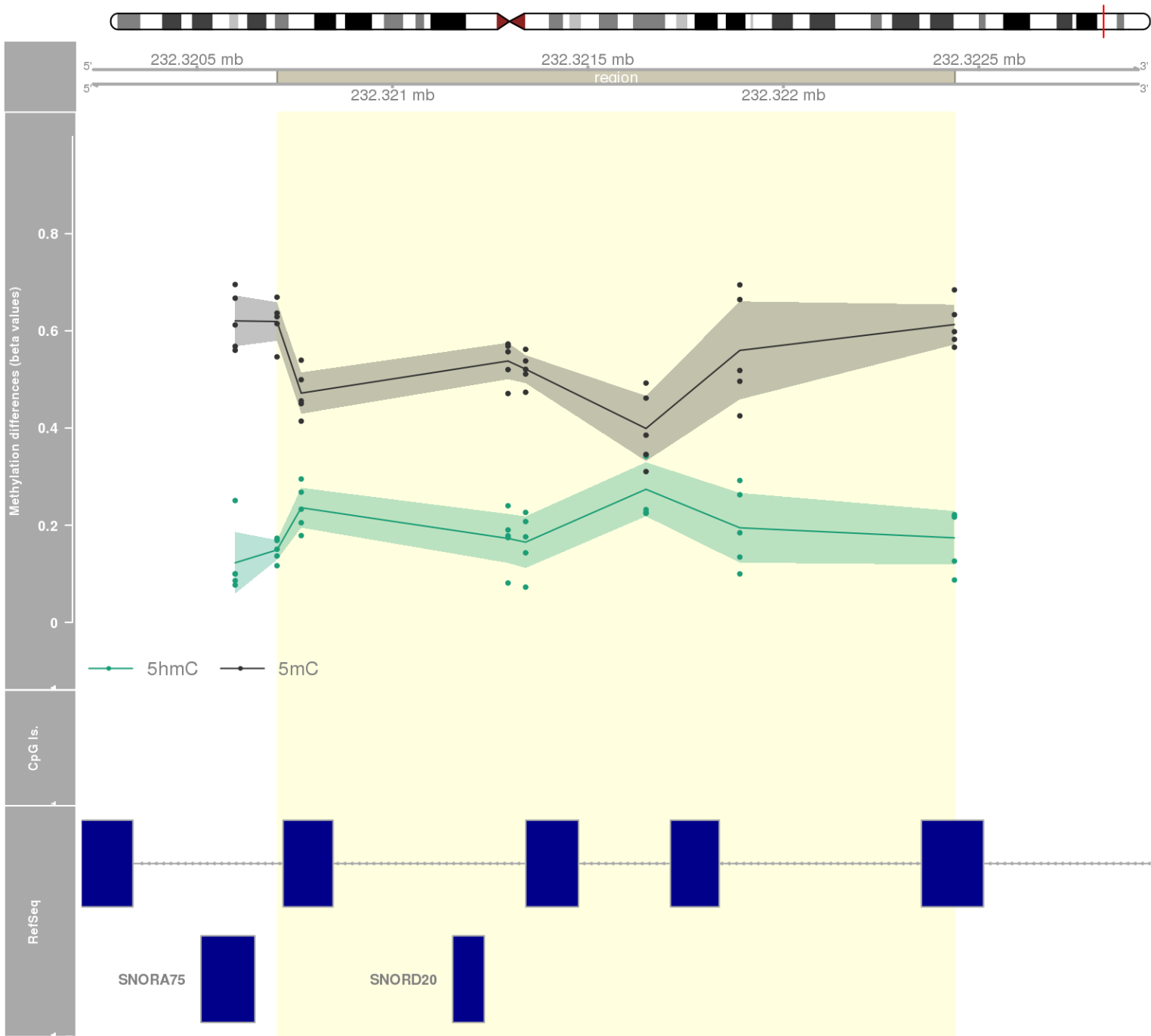
DMR 440 // chr13:114828264-114830313 // 2049 pb. (7 probes) // pvalue: 0.01 // fwer: 0.173 // pvalueArea: 0.031 // fwerArea: 1
- genes: RASA3 -



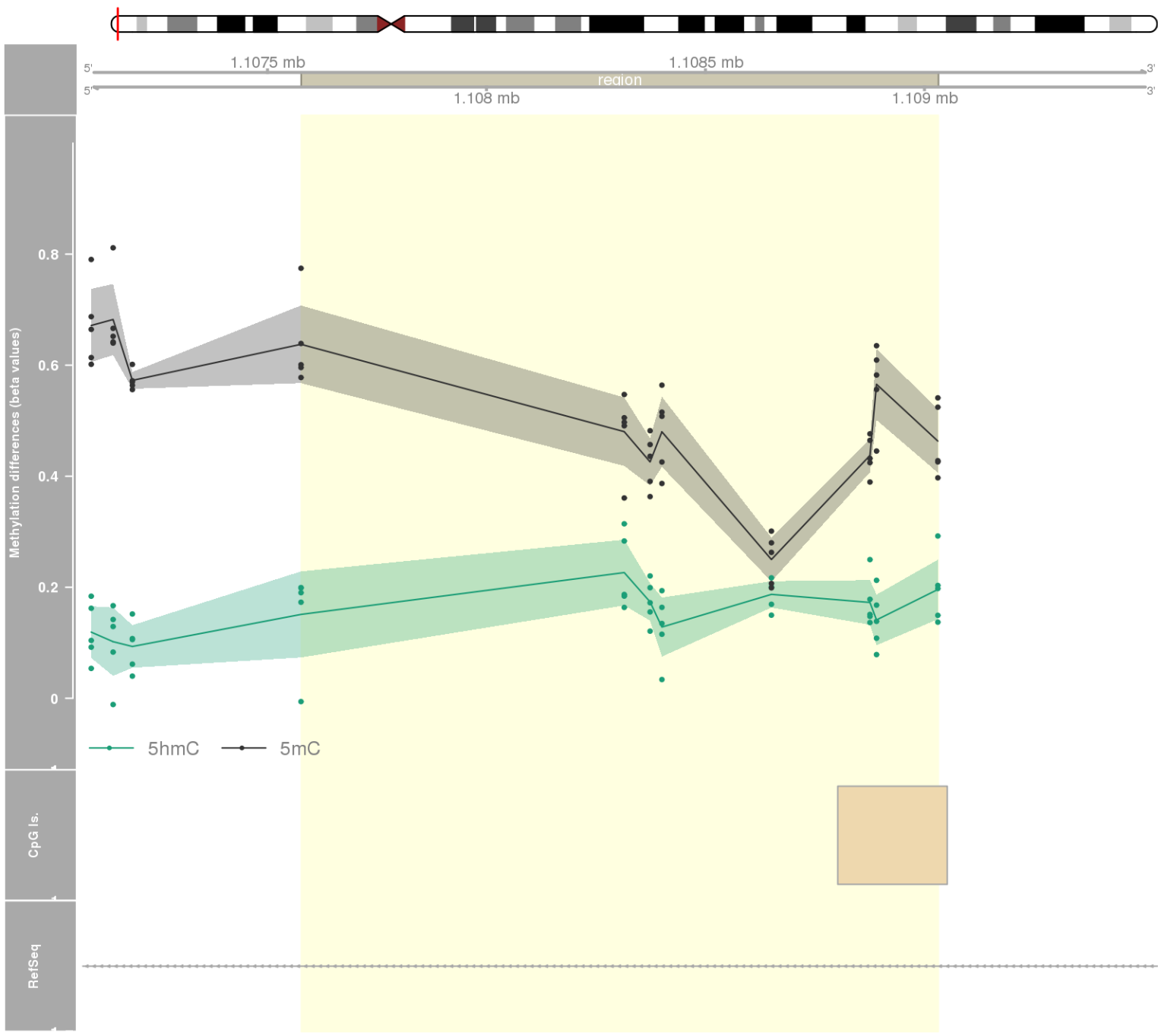
DMR 441 // chr15:78555907-78556534 // 627 pb. (7 probes) // pvalue: 0.01 // fwer: 0.173 // pvalueArea: 0.031 // fwerArea: 1
- genes: DNAJA4 -



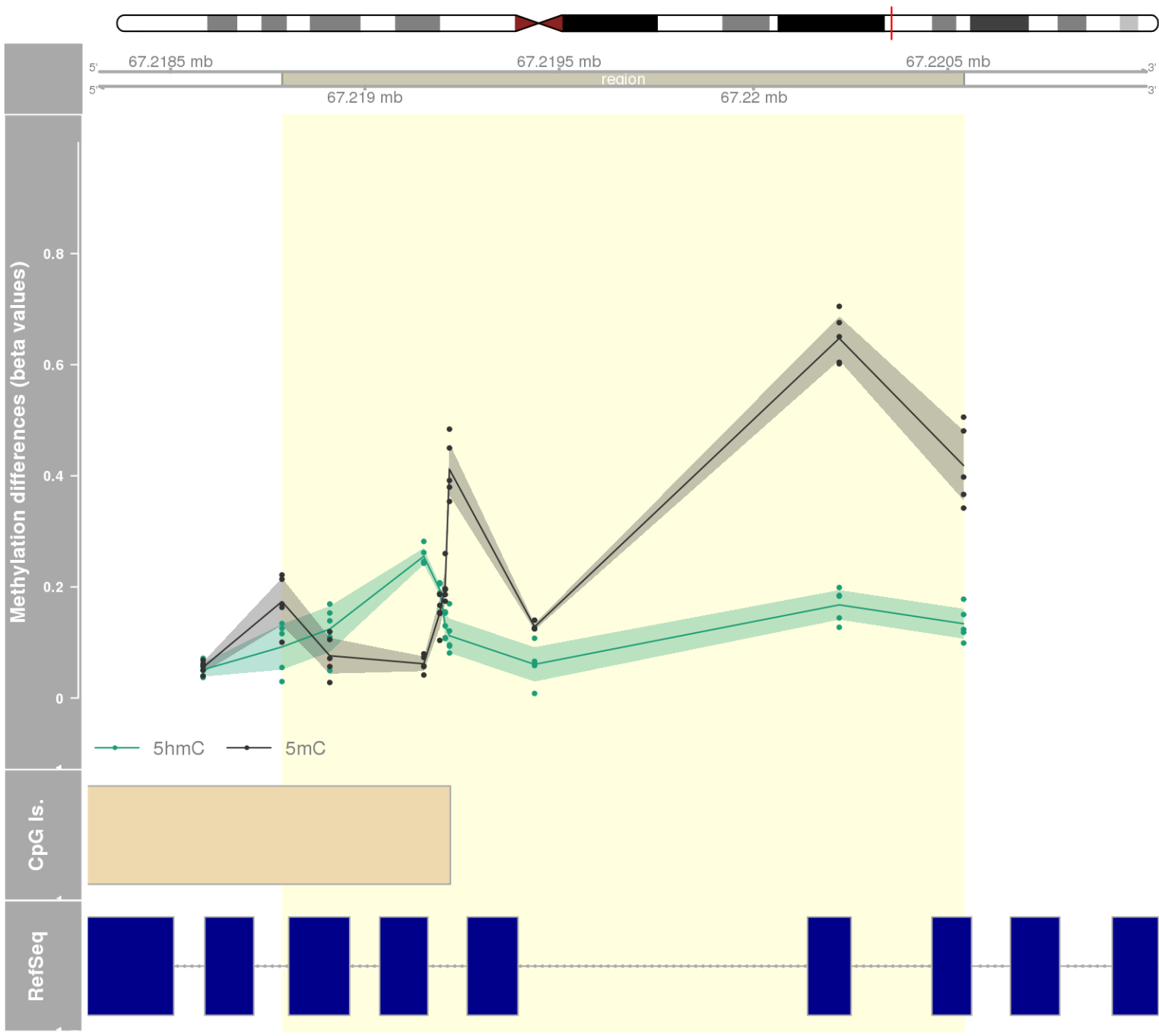
DMR 442 // chr2:232320705-232322438 // 1733 pb. (7 probes) // pvalue: 0.011 // fwer: 0.173 // pvalueArea: 0.031 // fwerArea: 1
- genes: SNORD20 / NCL -



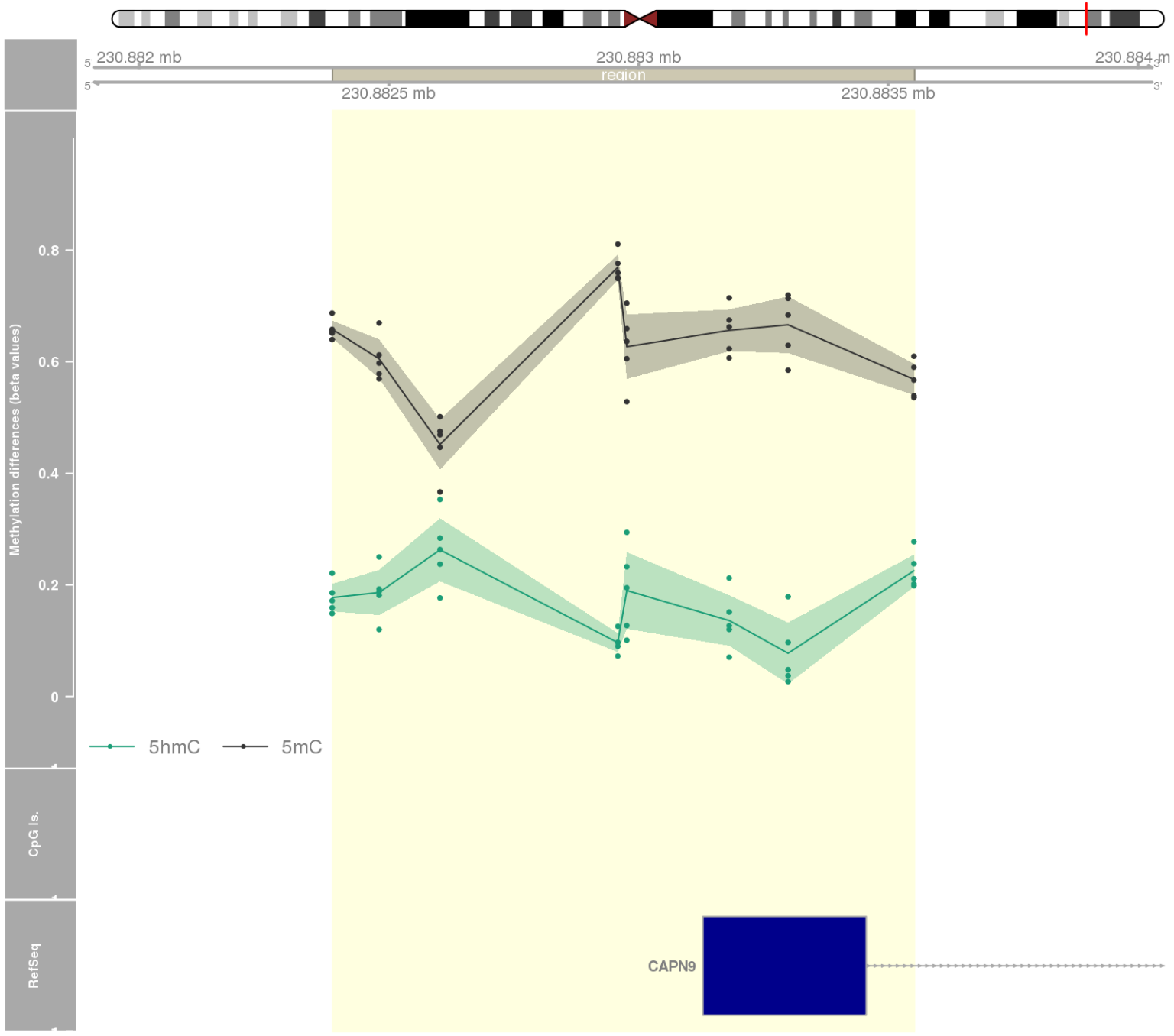
DMR 443 // chr5:1107577-1109031 // 1454 pb. (8 probes) // pvalue: 0.016 // fwer: 0.522 // pvalueArea: 0.031 // fwerArea: 1
- genes: SLC12A7 -



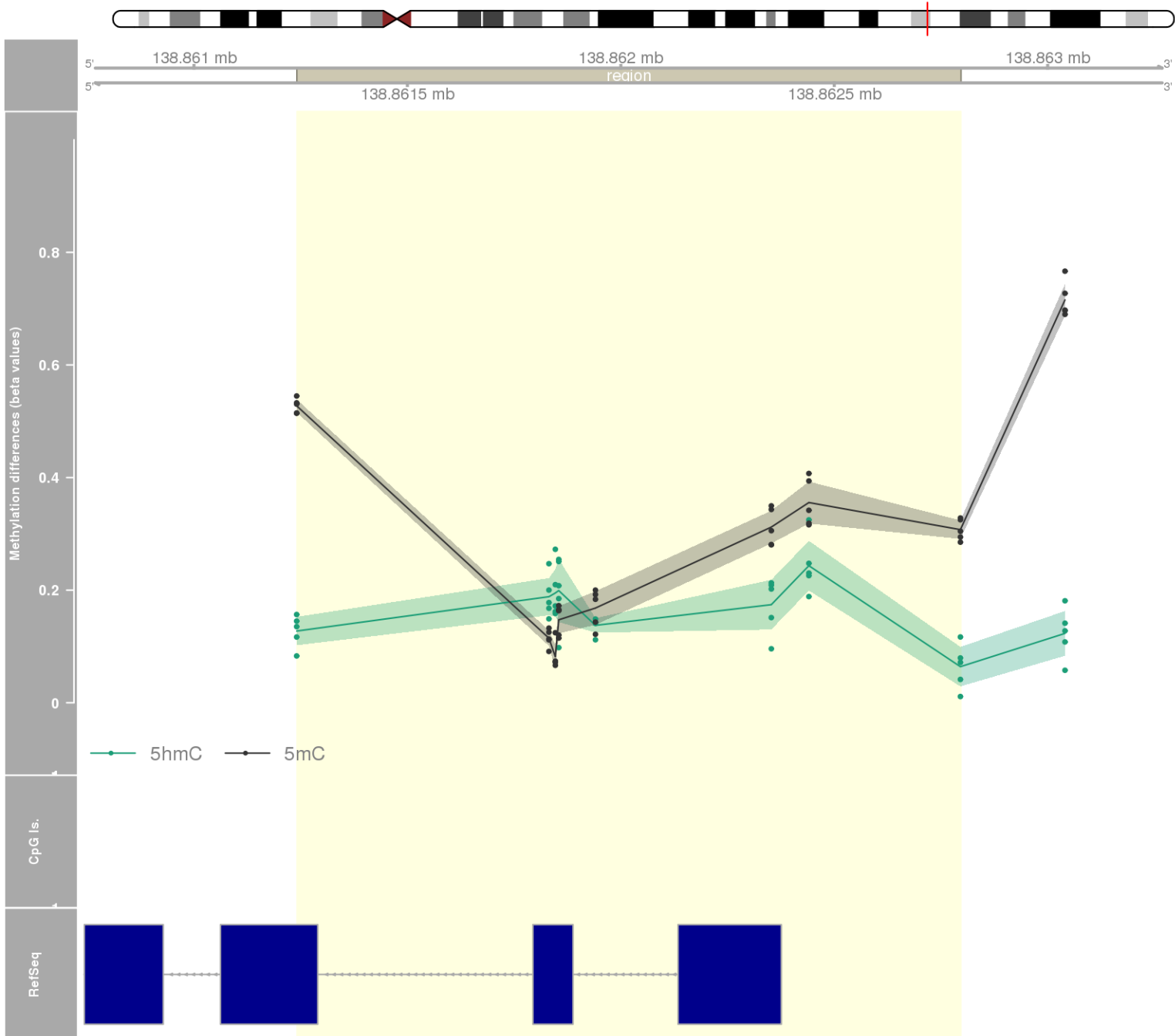
DMR 444 // chr16:67218787-67220541 // 1754 pb. (9 probes) // pvalue: 0.016 // fwer: 0.36 // pvalueArea: 0.031 // fwerArea: 1
- genes: EXOC3L1 -



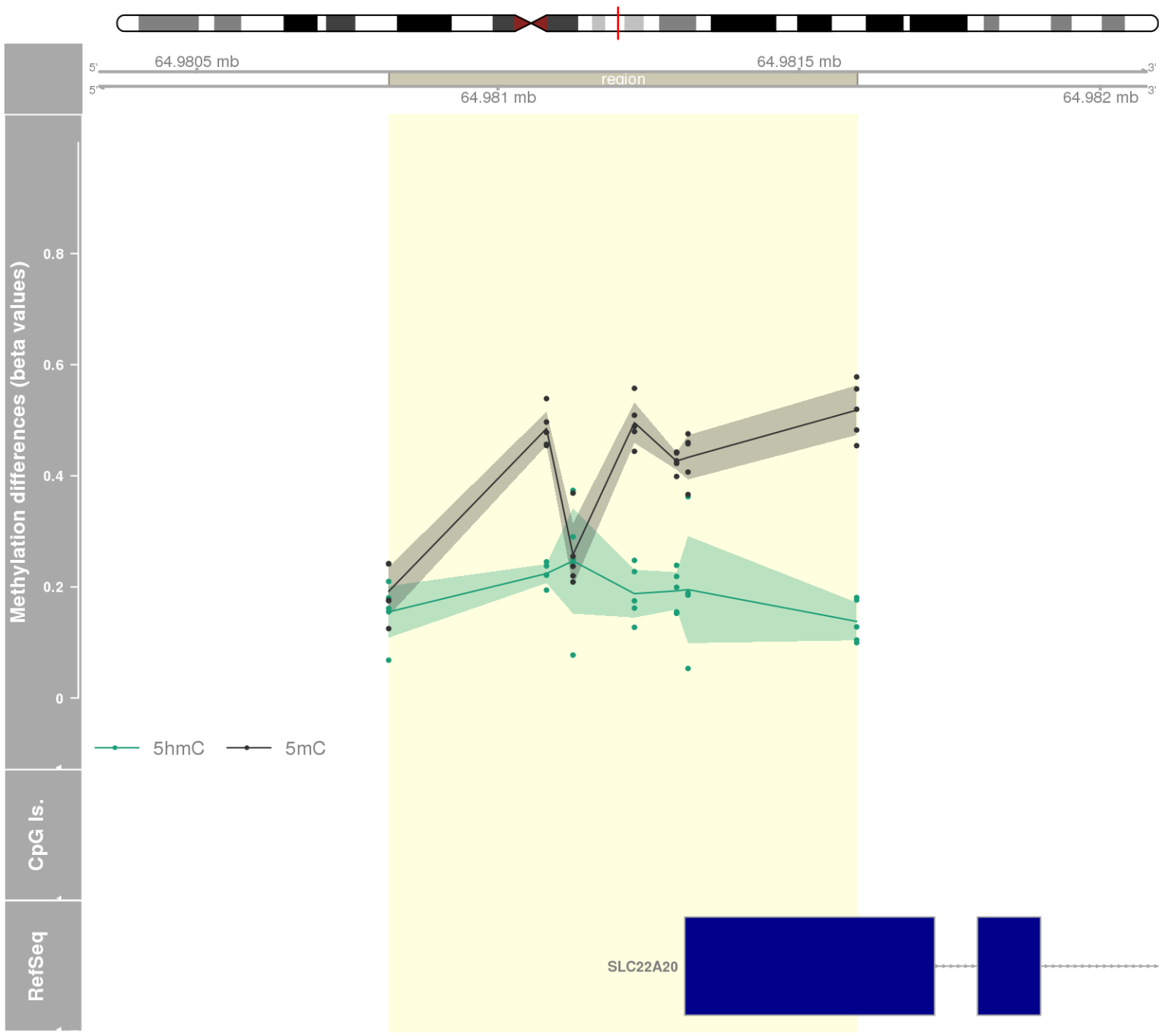
DMR 445 // chr1:230882387-230883552 // 1165 pb. (8 probes) // pvalue: 0.016 // fwer: 0.522 // pvalueArea: 0.031 // fwerArea: 1
- genes: CAPN9 -



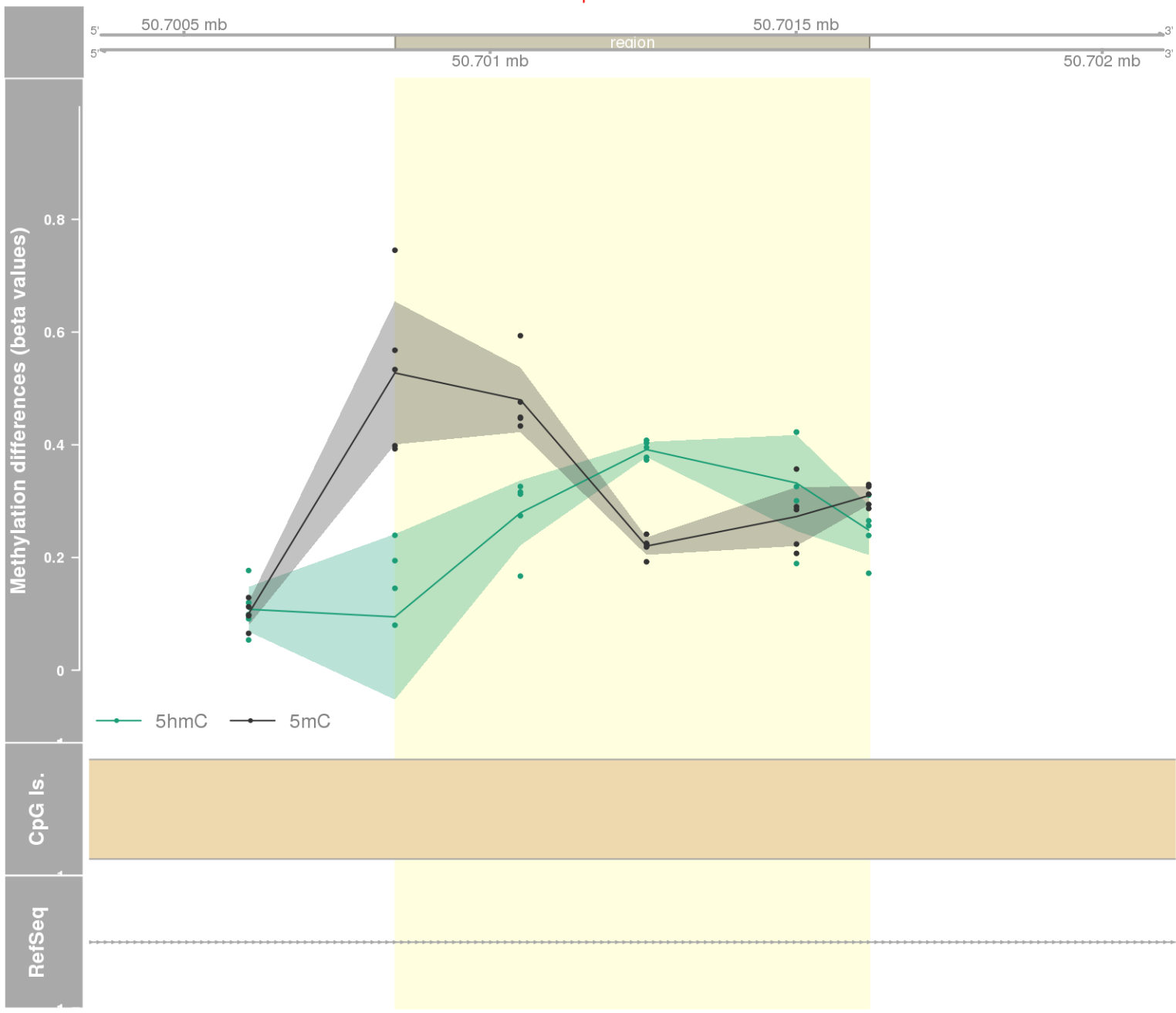
DMR 446 // chr5:138861241-138862796 // 1555 pb. (8 probes) // pvalue: 0.016 // fwer: 0.522 // pvalueArea: 0.031 // fwerArea: 1
- genes: TMEM173 -



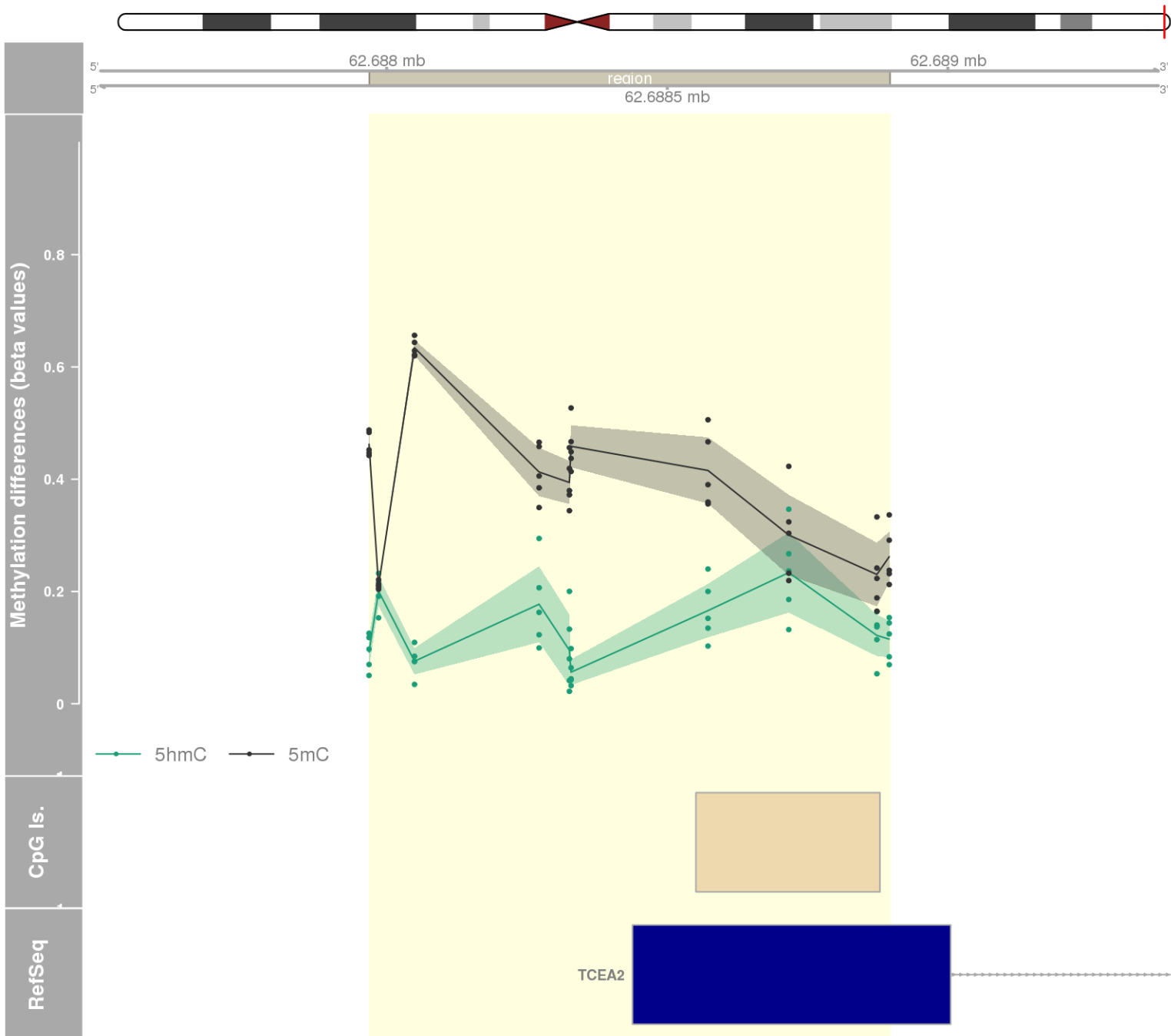
DMR 447 // chr11:64980819-64981596 // 777 pb. (7 probes) // pvalue: 0.011 // fwer: 0.173 // pvalueArea: 0.031 // fwerArea: 1
- genes: SLC22A20 -



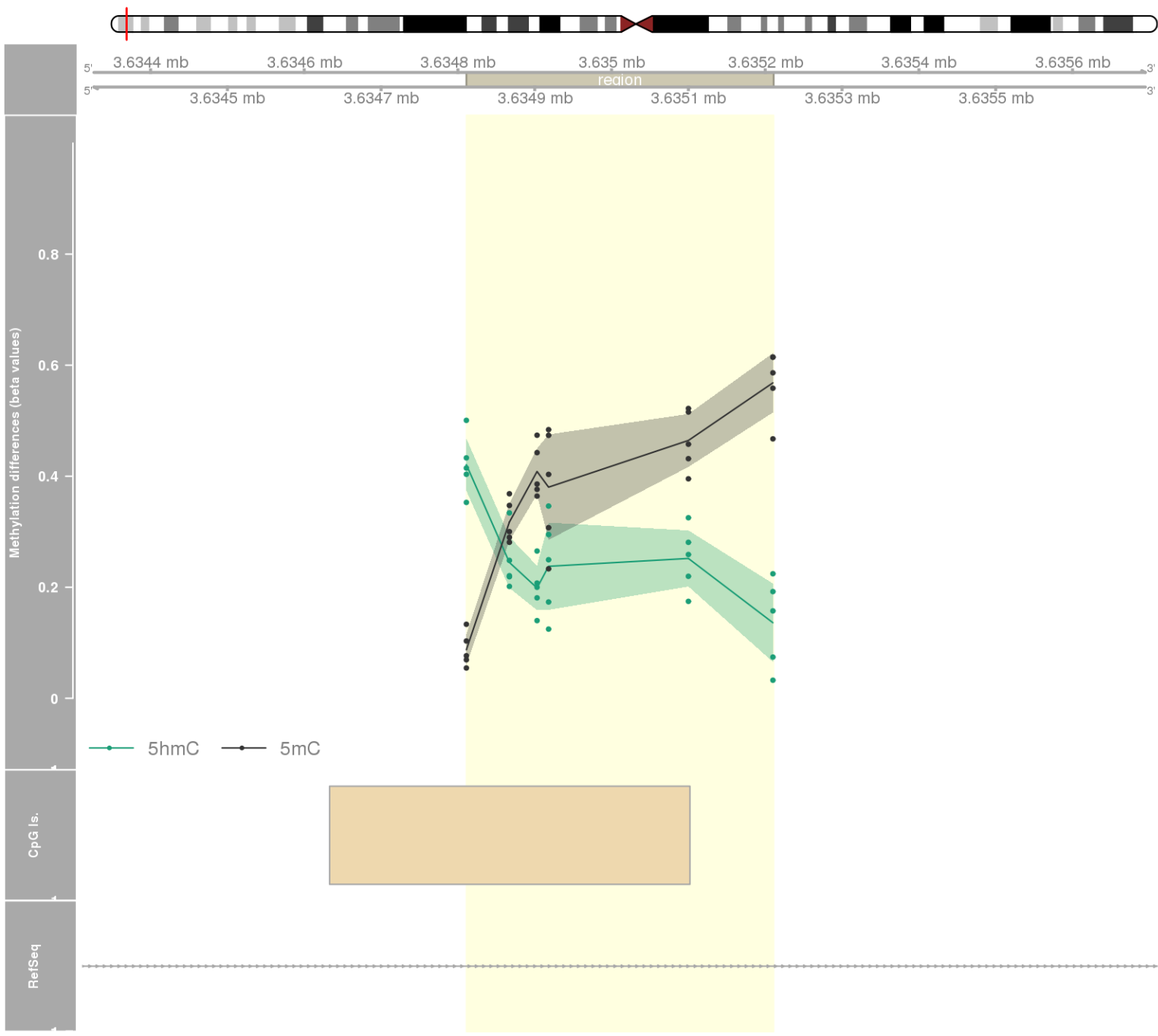
DMR 448 // chr13:50700845-50701619 // 774 pb. (5 probes) // pvalue: 0.001 // fwer: 0.496 // pvalueArea: 0.031 // fwerArea: 1
- genes: DLEU1 -



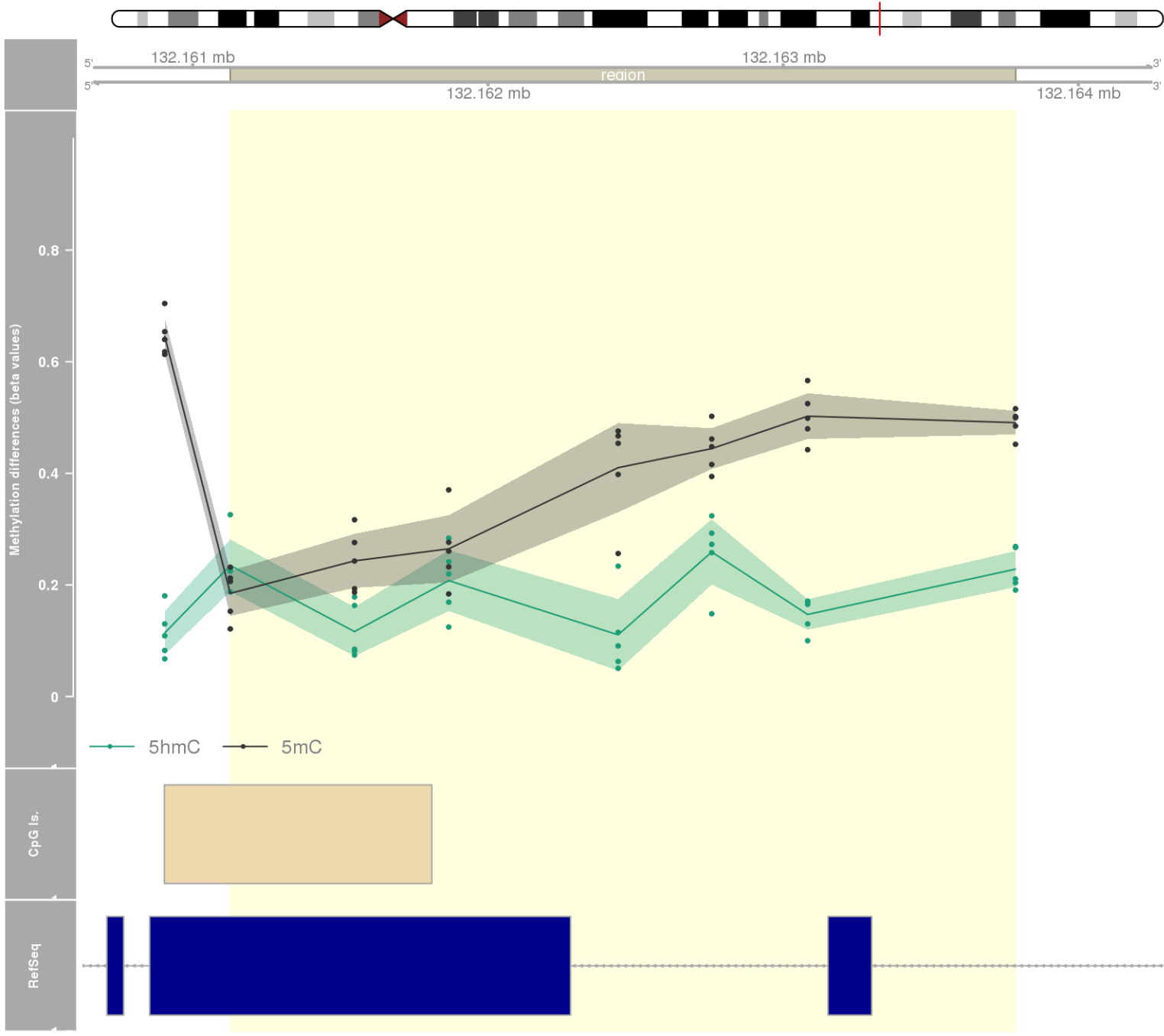
DMR 449 // chr20:62687969-62688896 // 927 pb. (10 probes) // pvalue: 0.015 // fwer: 0.36 // pvalueArea: 0.031 // fwerArea: 1
- genes: TCEA2 -



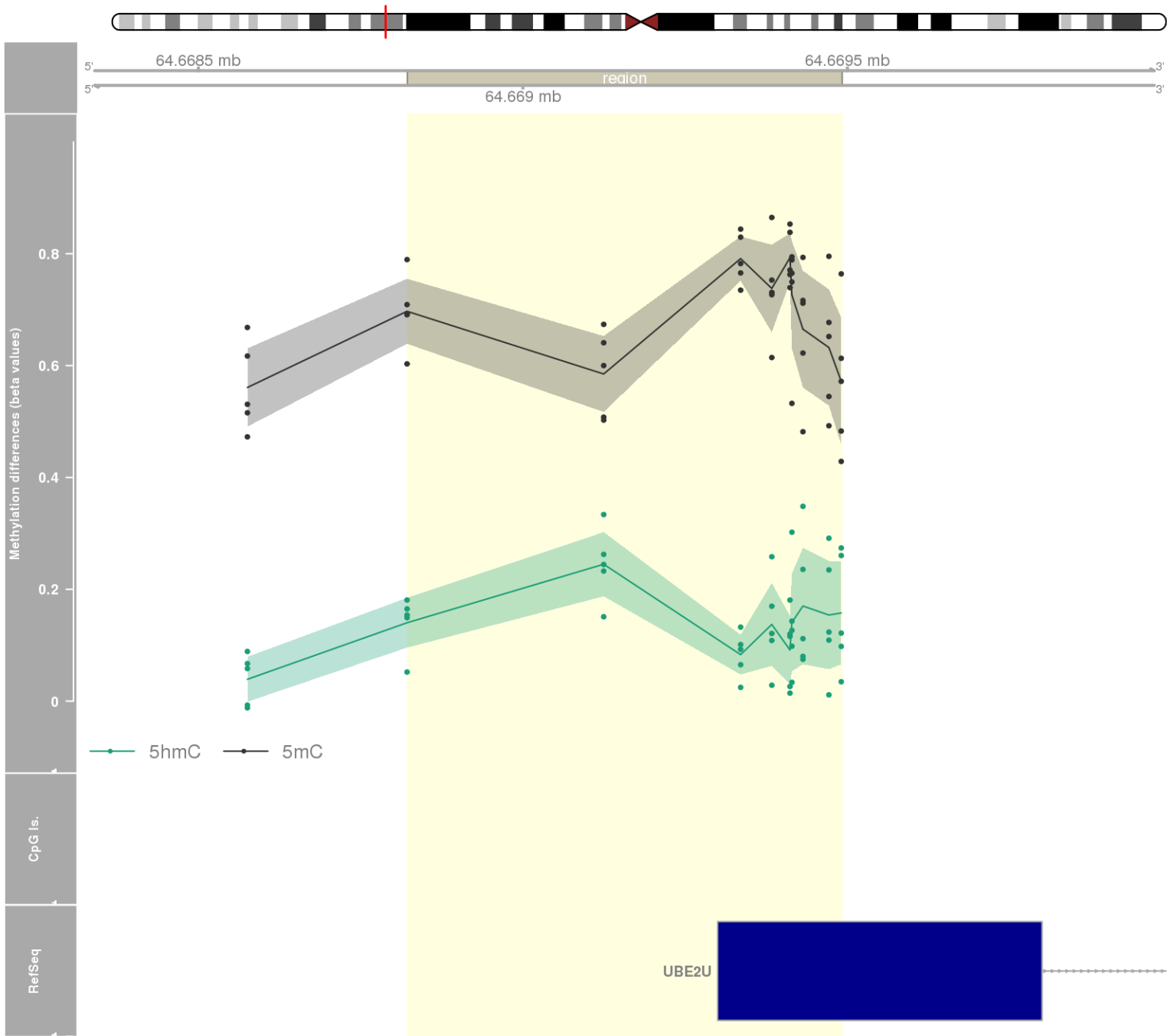
DMR 450 // chr1:3634811-3635210 // 399 pb. (6 probes) // pvalue: 0.004 // fwer: 0.173 // pvalueArea: 0.031 // fwerArea: 1
- genes: TP73 -



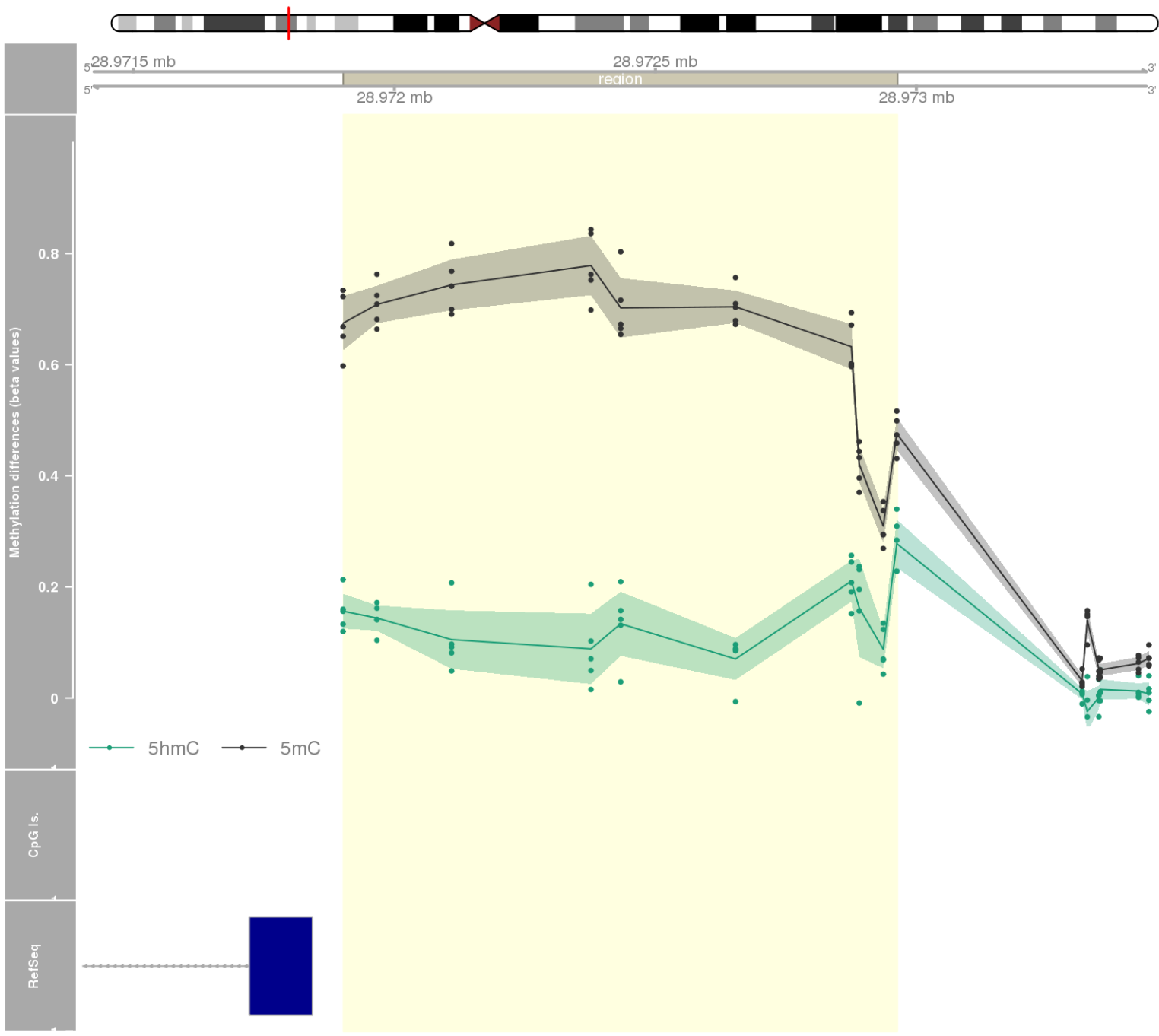
DMR 451 // chr5:132161127-132163787 // 2660 pb. (7 probes) // pvalue: 0.011 // fwer: 0.173 // pvalueArea: 0.031 // fwerArea: 1
- genes: SHROOM1 -



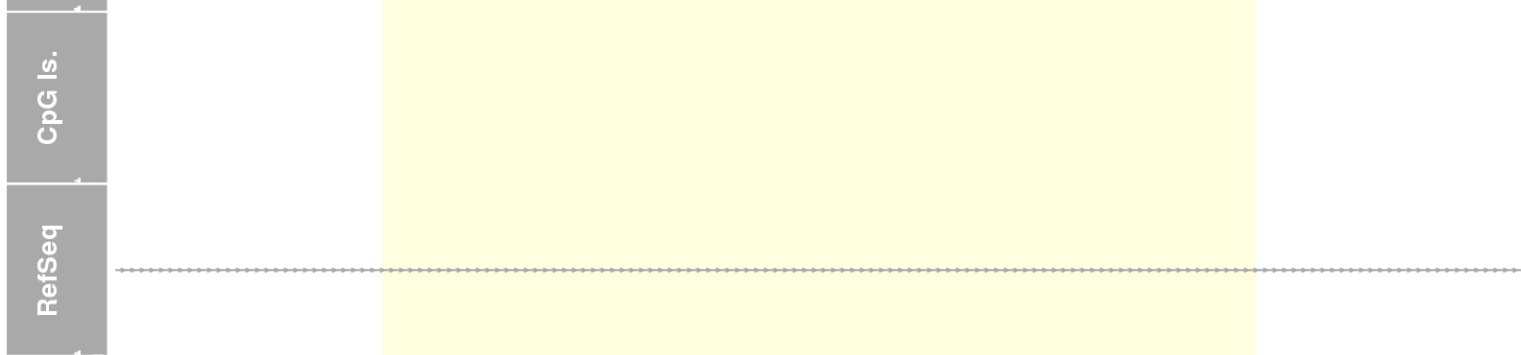
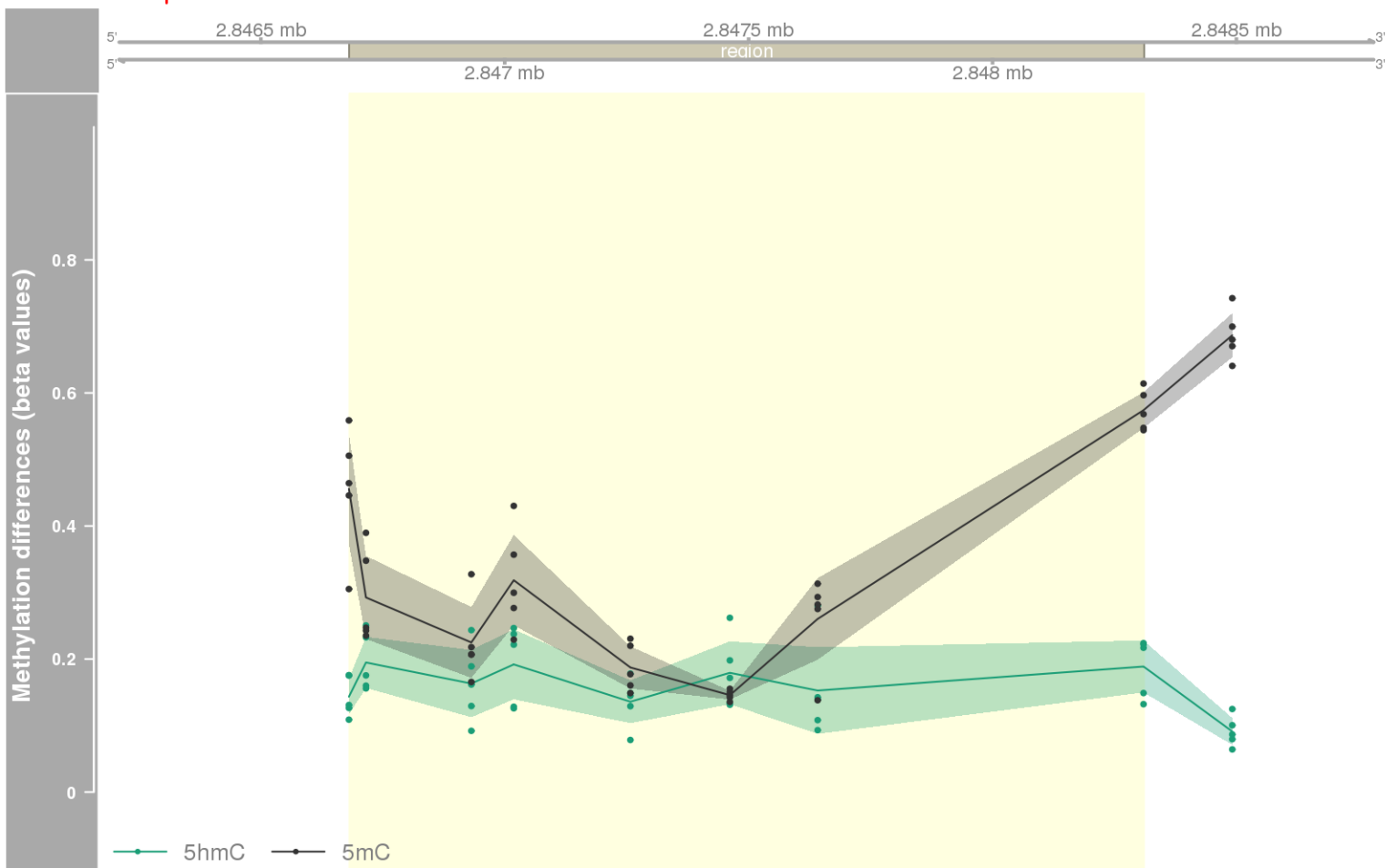
DMR 452 // chr1:64668822-64669491 // 669 pb. (9 probes) // pvalue: 0.017 // fwer: 0.36 // pvalueArea: 0.031 // fwerArea: 1
- genes: UBE2U -



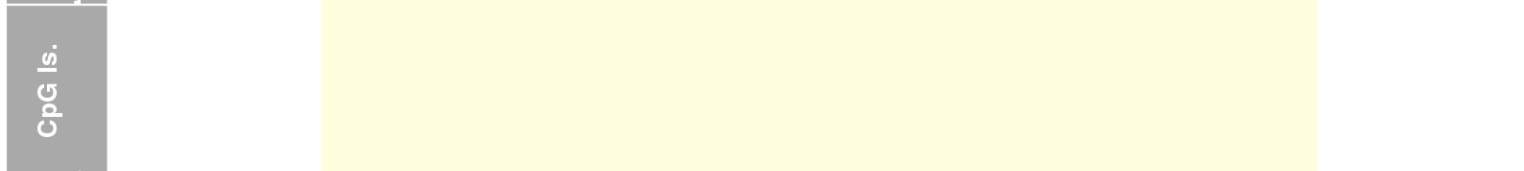
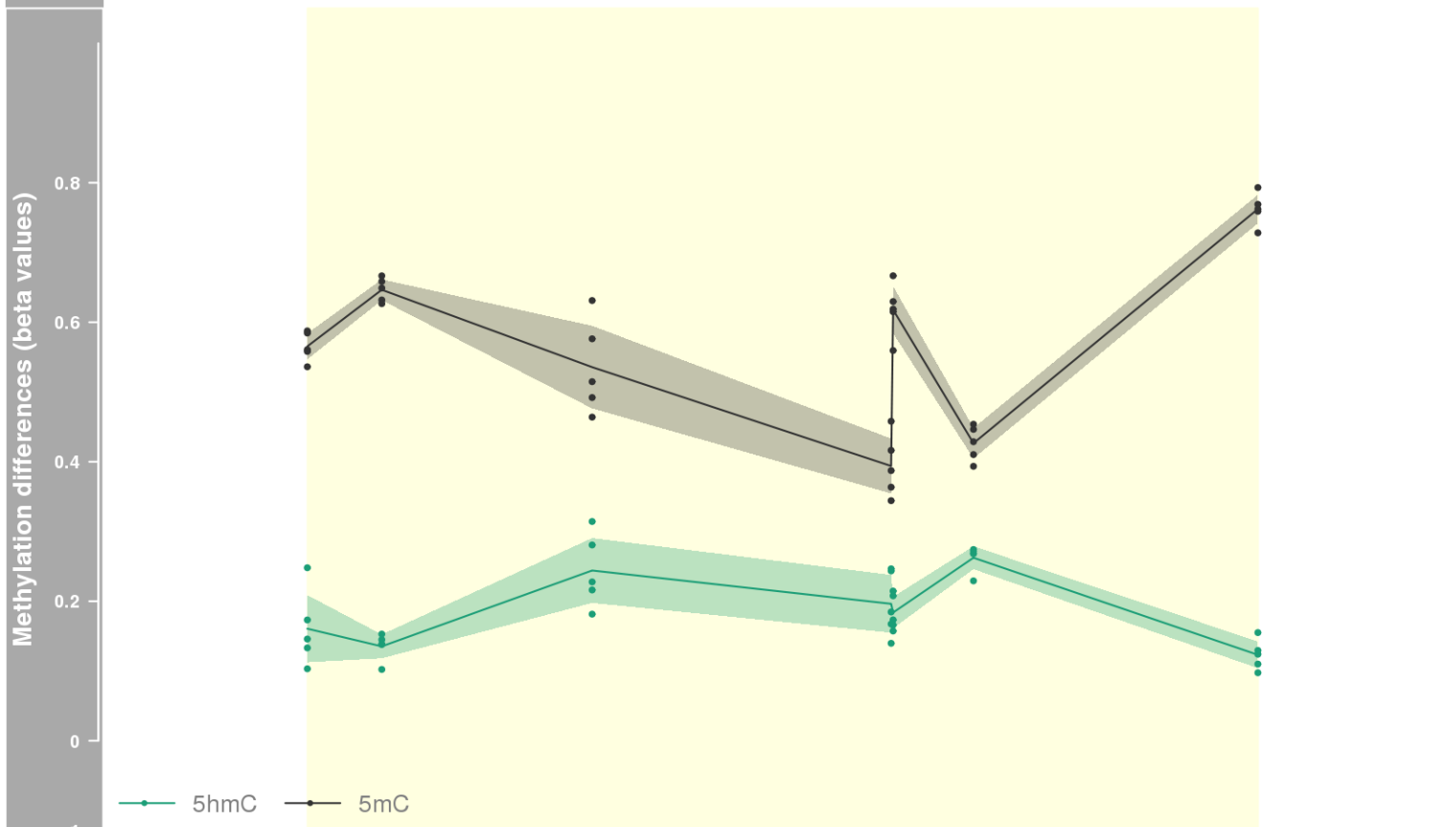
DMR 453 // chr6:28971902-28972963 // 1061 pb. (10 probes) // pvalue: 0.015 // fwer: 0.36 // pvalueArea: 0.031 // fwerArea: 1
- genes: ZNF311 -



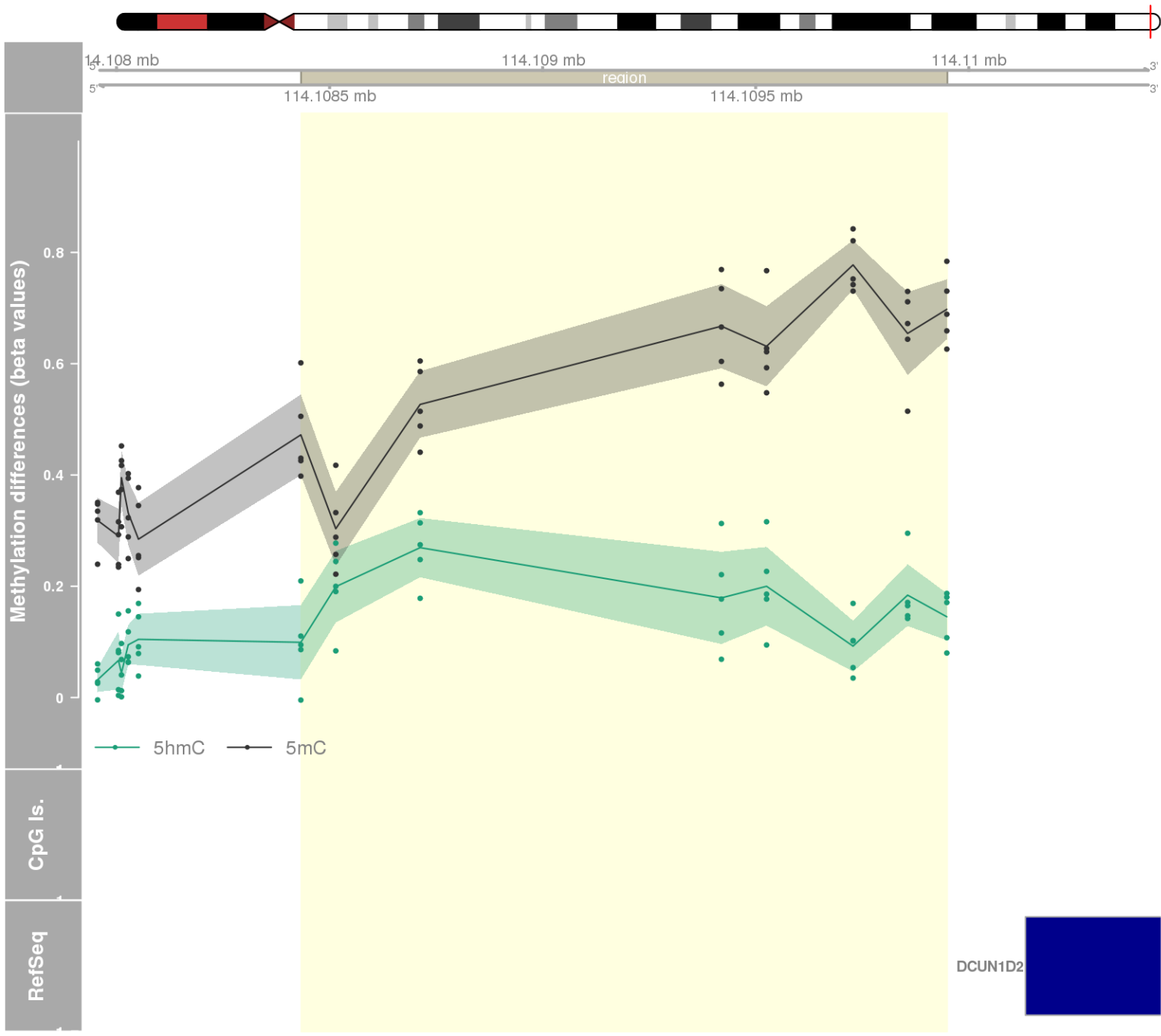
DMR 454 // chr11:2846681-2848310 // 1629 pb. (8 probes) // pvalue: 0.017 // fwer: 0.522 // pvalueArea: 0.032 // fwerArea: 1
- genes: KCNQ1 -



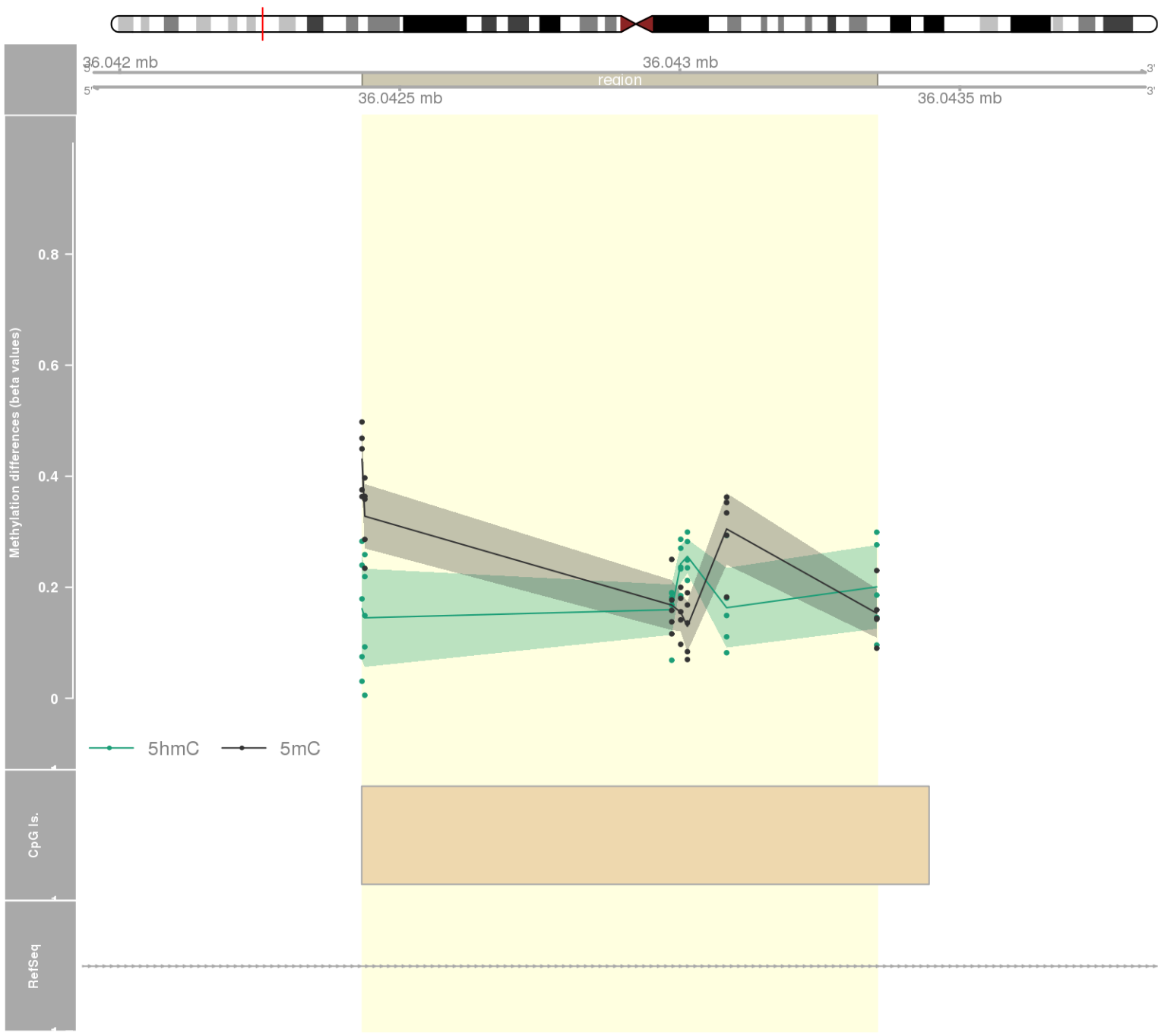
DMR 455 // chr11:73719683-73722096 // 2413 pb. (7 probes) // pvalue: 0.011 // fwer: 0.173 // pvalueArea: 0.032 // fwerArea: 1
- genes: UCP3 -



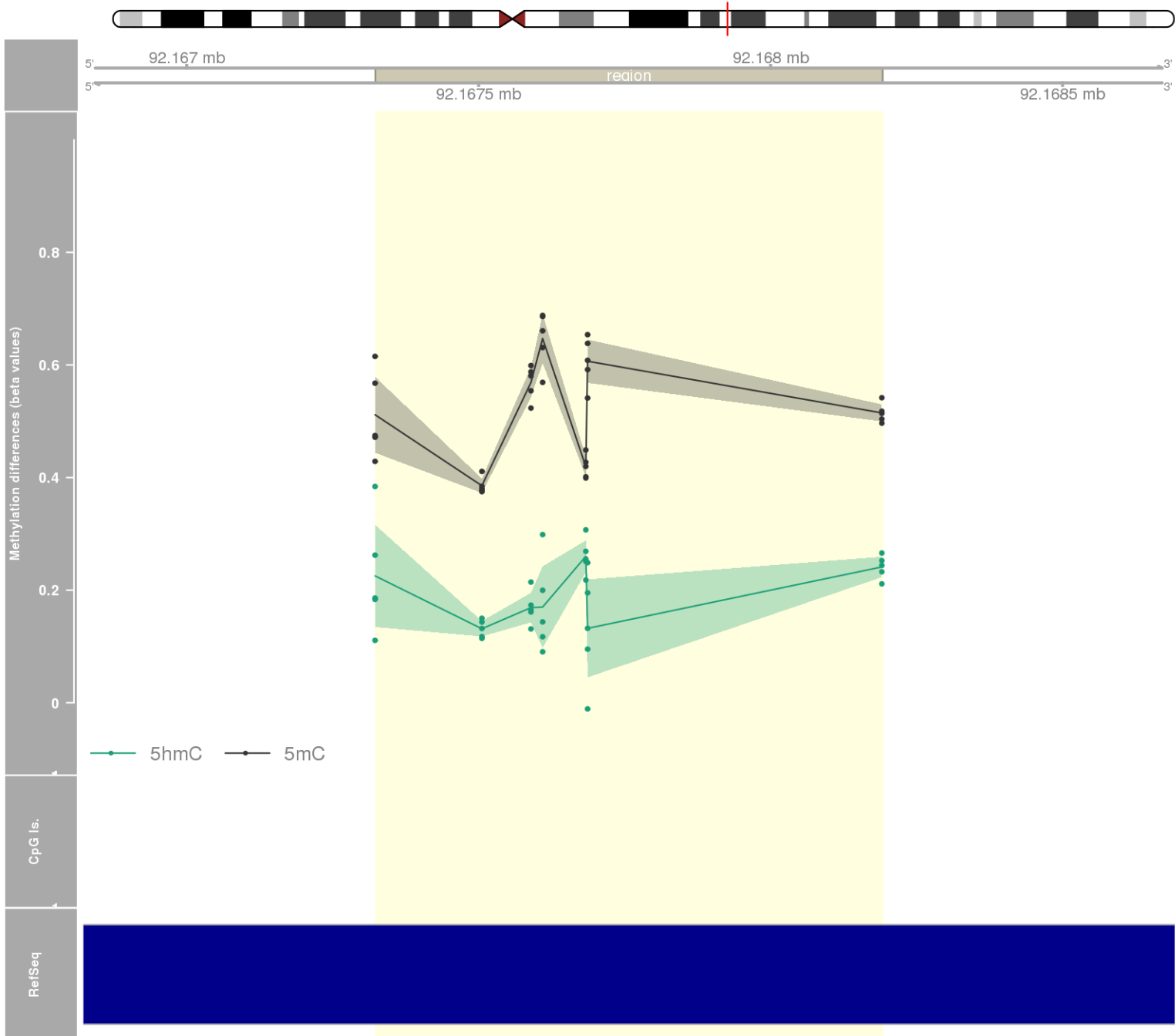
DMR 456 // chr13:114108433-114109949 // 1516 pb. (8 probes) // pvalue: 0.017 // fwer: 0.522 // pvalueArea: 0.032 // fwerArea: 1



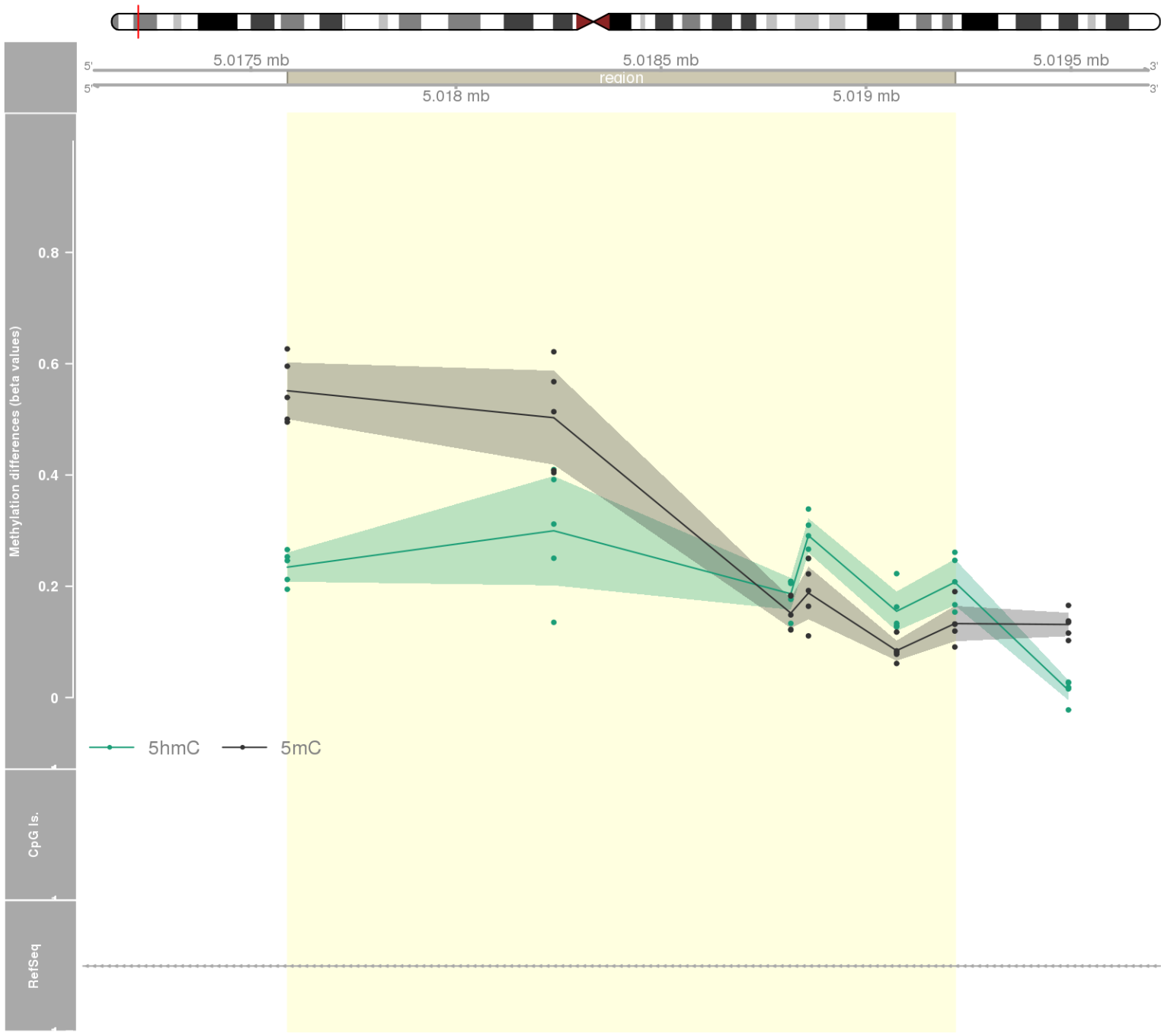
DMR 457 // chr1:36042433-36043352 // 919 pb. (7 probes) // pvalue: 0.011 // fwer: 0.173 // pvalueArea: 0.032 // fwerArea: 1
- genes: TFAP2E -



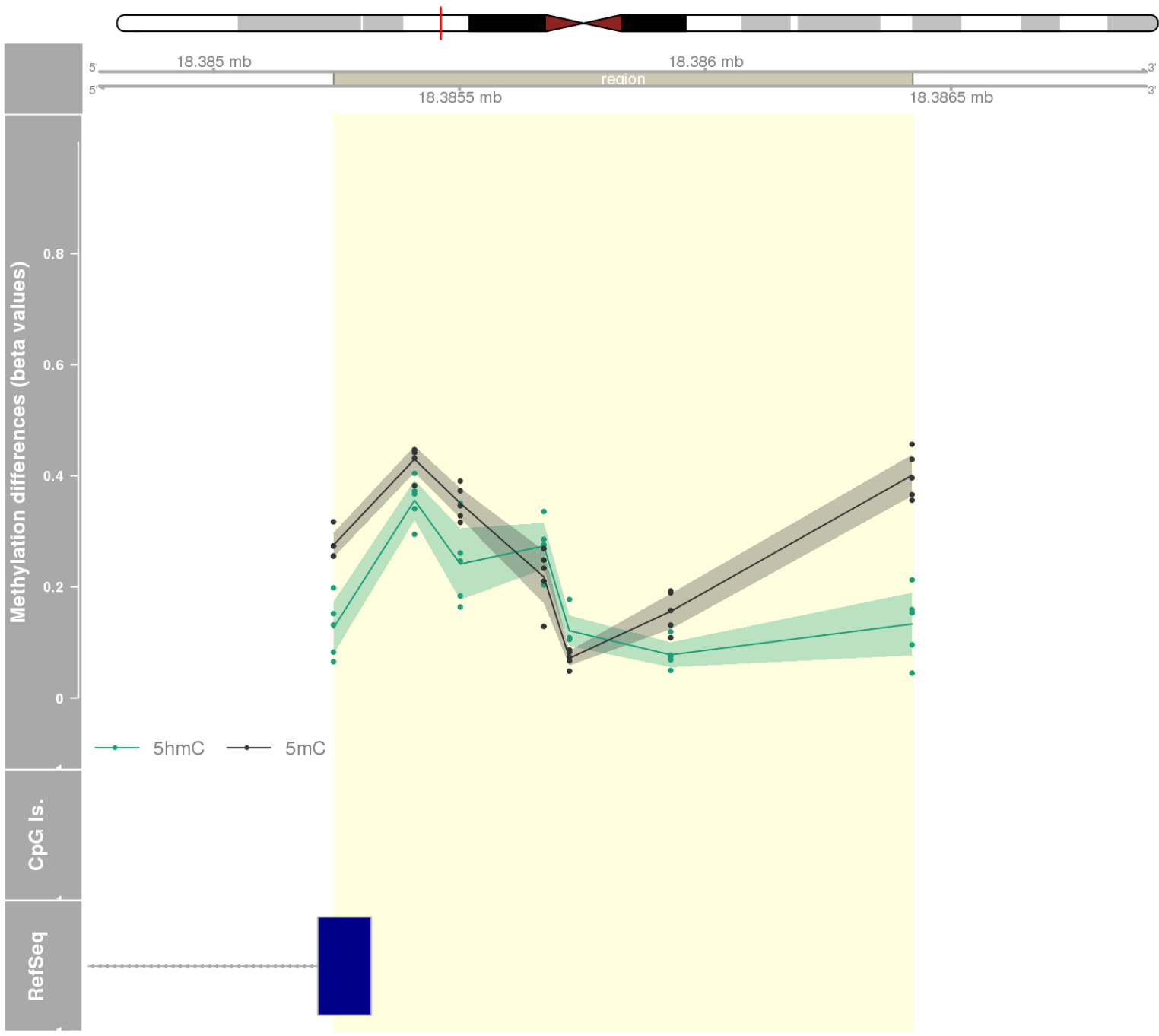
DMR 458 // chr7:92167323-92168191 // 868 pb. (7 probes) // pvalue: 0.011 // fwer: 0.173 // pvalueArea: 0.032 // fwerArea: 1



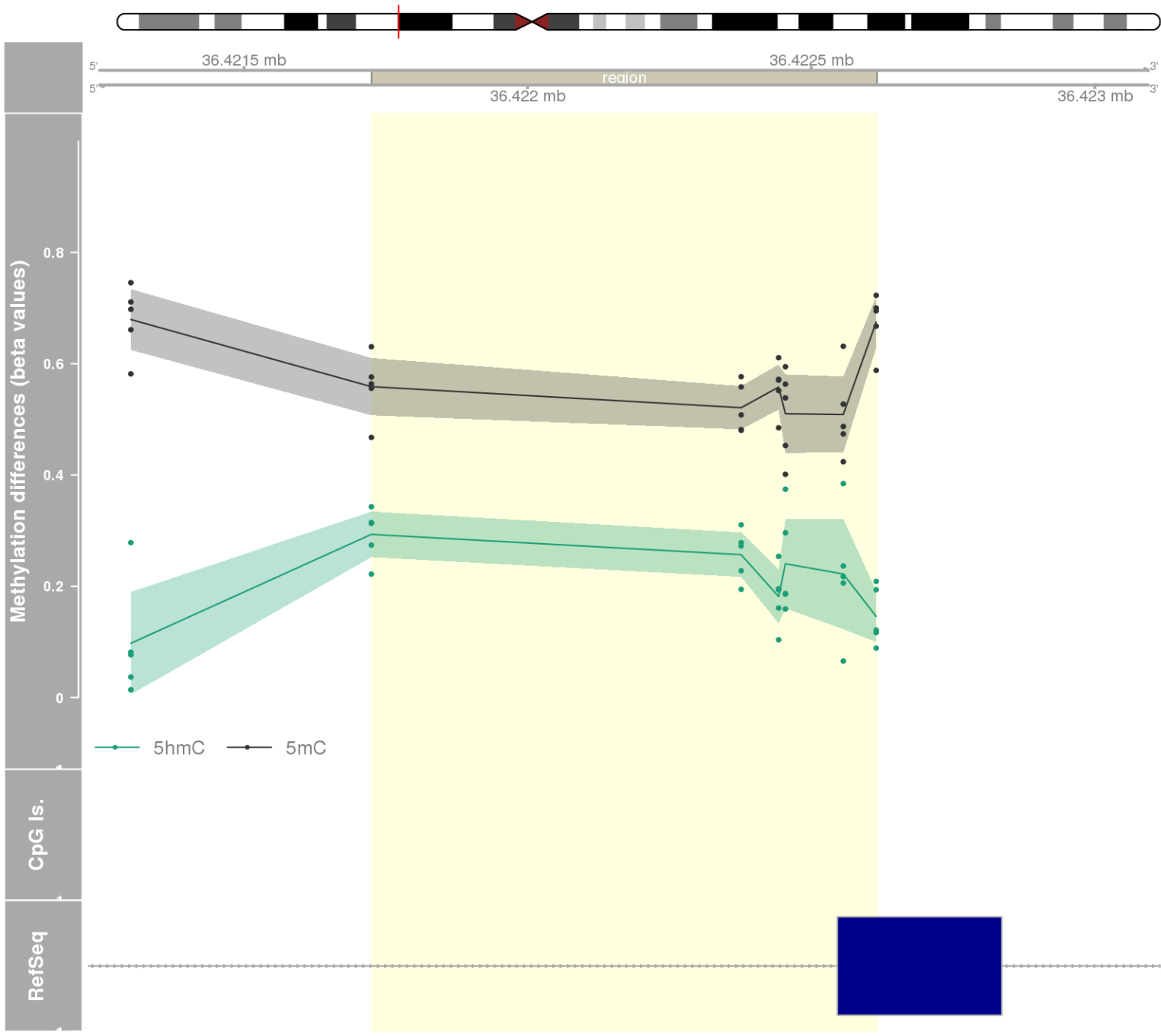
DMR 459 // chr3:5017589-5019217 // 1628 pb. (6 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.032 // fwerArea: 1
- genes: BHLHE40-AS1 -



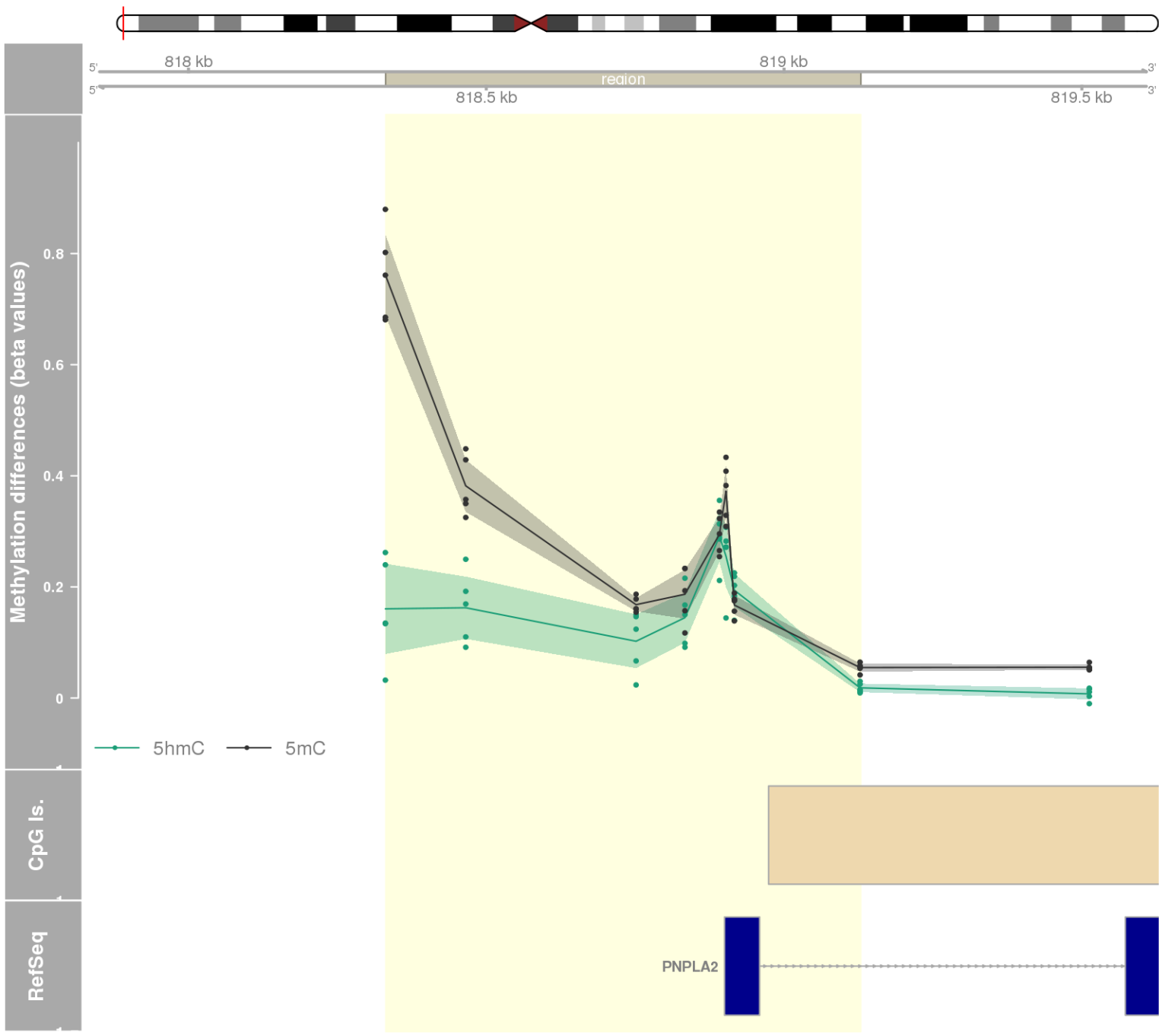
DMR 460 // chr19:18385244-18386421 // 1177 pb. (7 probes) // pvalue: 0.012 // fwer: 0.173 // pvalueArea: 0.032 // fwerArea: 1
- genes: KIAA1683 -



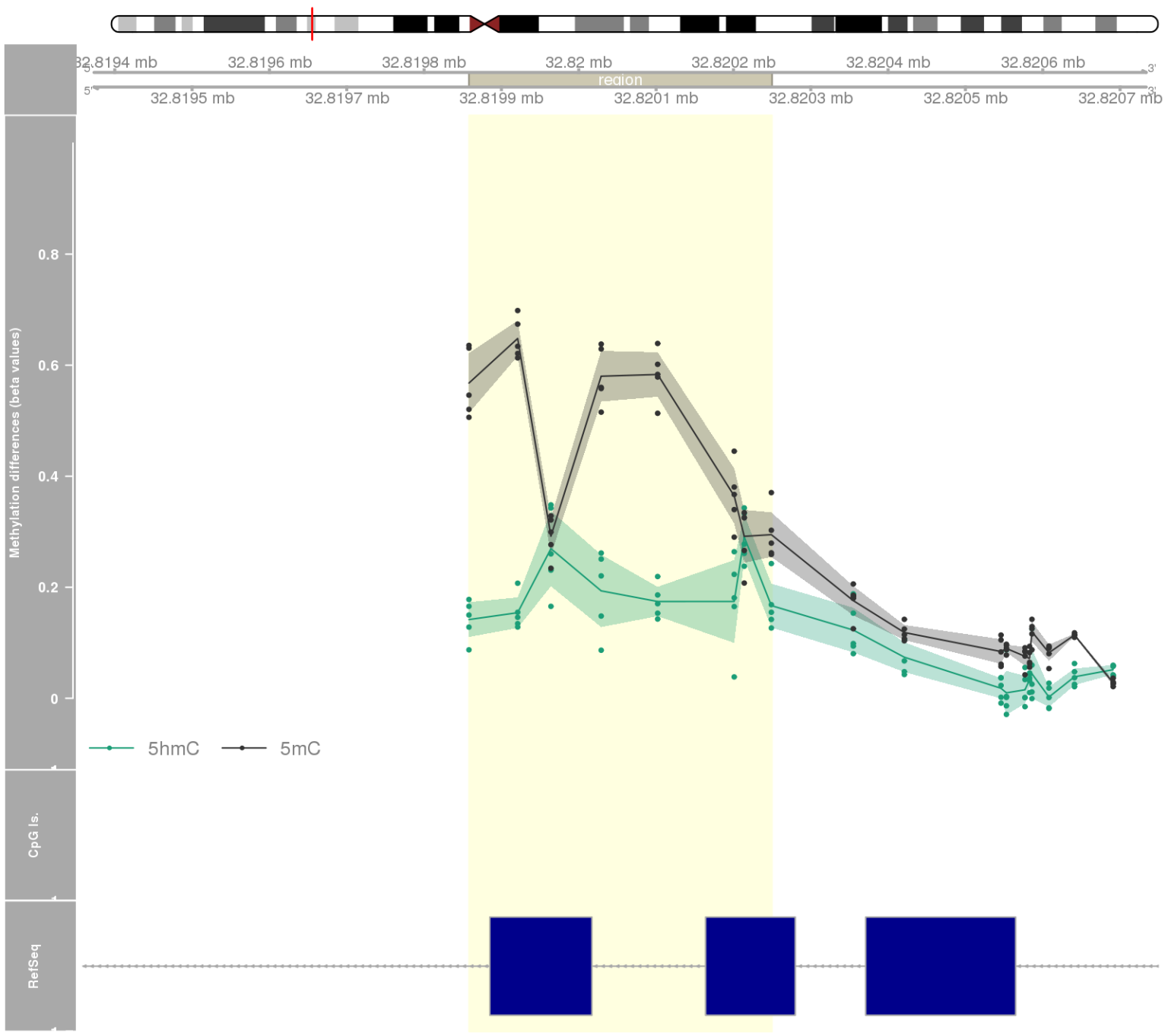
DMR 461 // chr11:36421725-36422615 // 890 pb. (6 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.032 // fwerArea: 1
- genes: PRR5L -



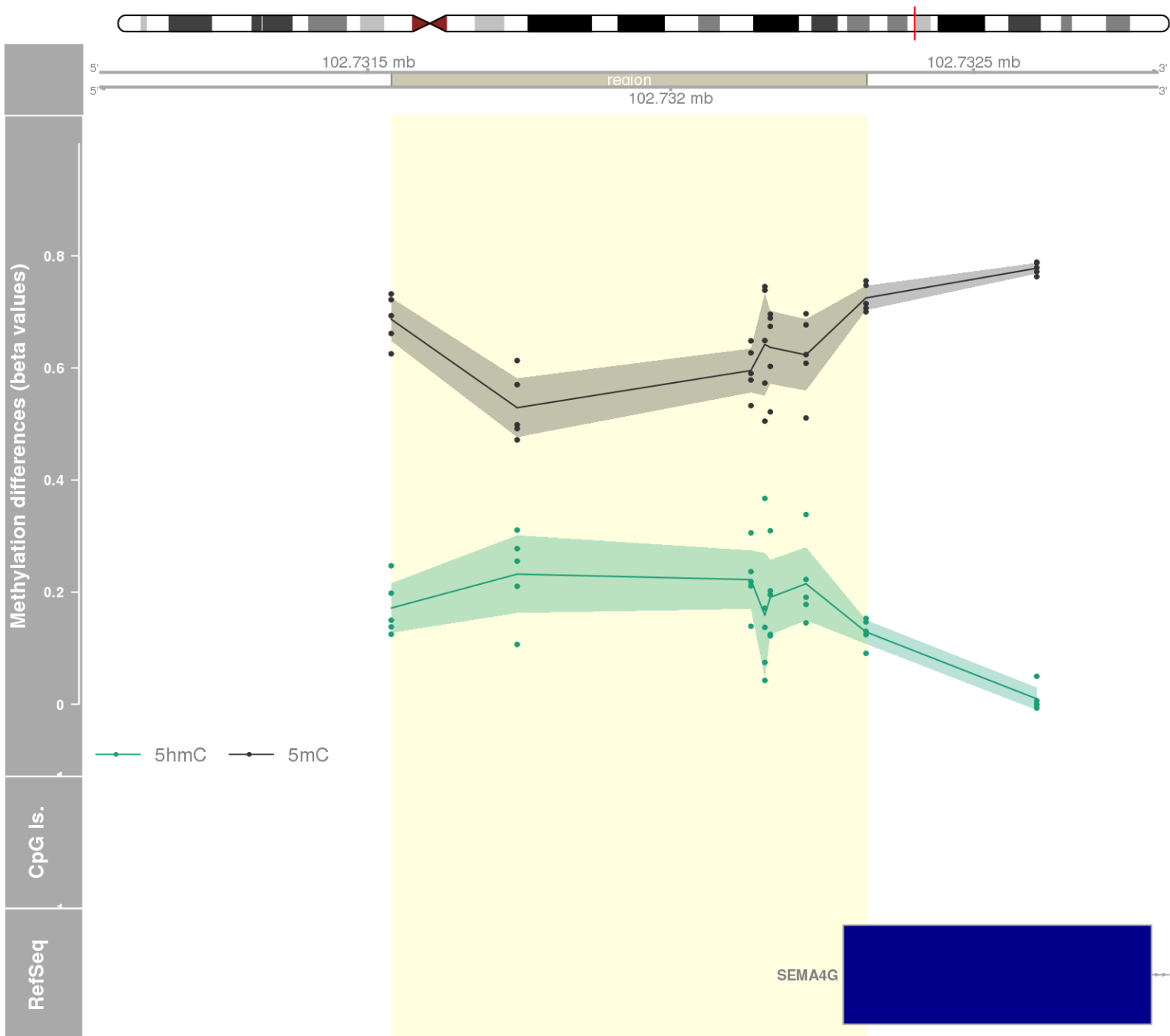
DMR 462 // chr11:818331-819128 // 797 pb. (8 probes) // pvalue: 0.017 // fwer: 0.522 // pvalueArea: 0.032 // fwerArea: 1
- genes: PNPLA2 -



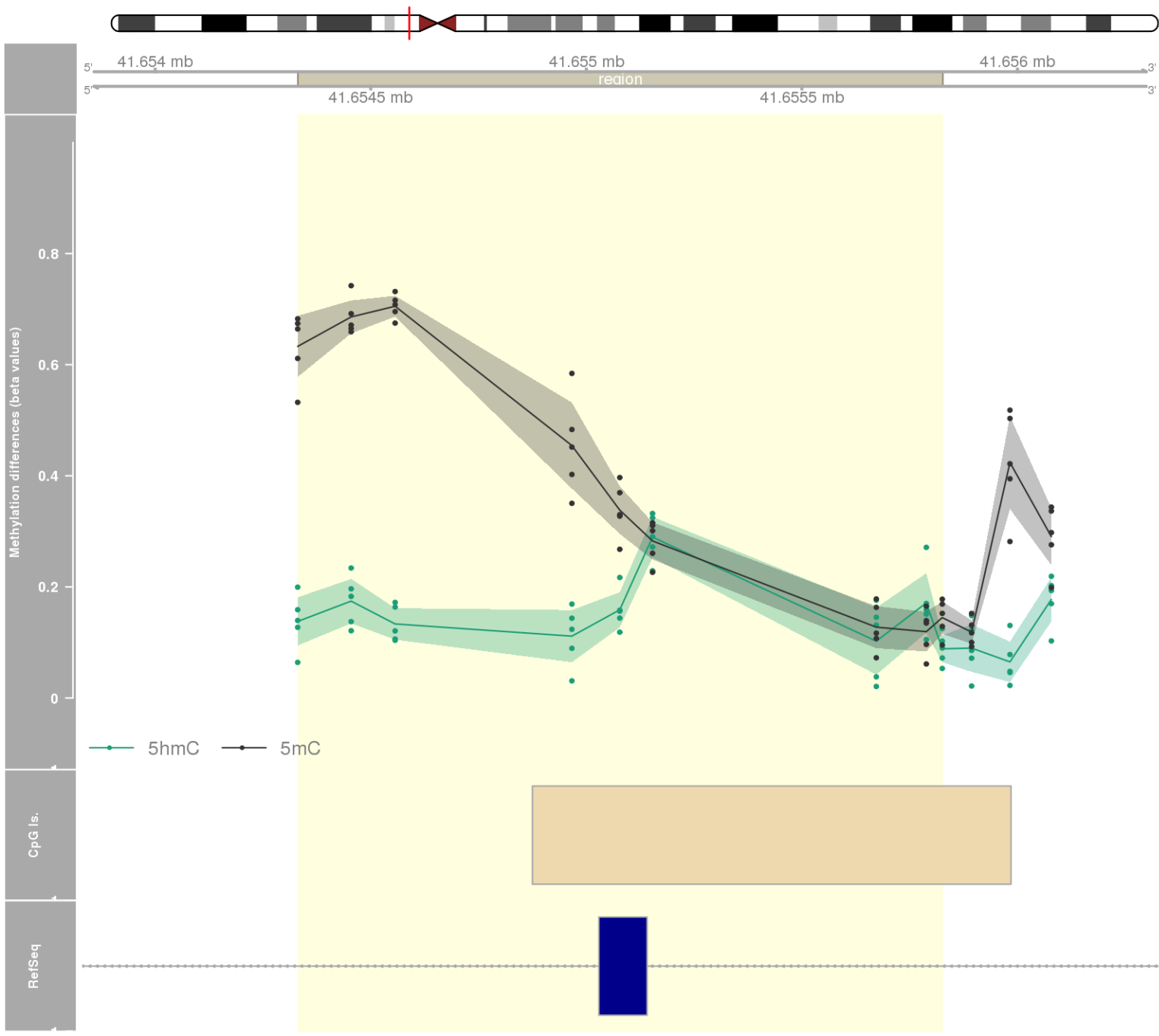
DMR 463 // chr6:32819858-32820249 // 391 pb. (8 probes) // pvalue: 0.017 // fwer: 0.522 // pvalueArea: 0.032 // fwerArea: 1
- genes: PSMB9 / TAP1 -



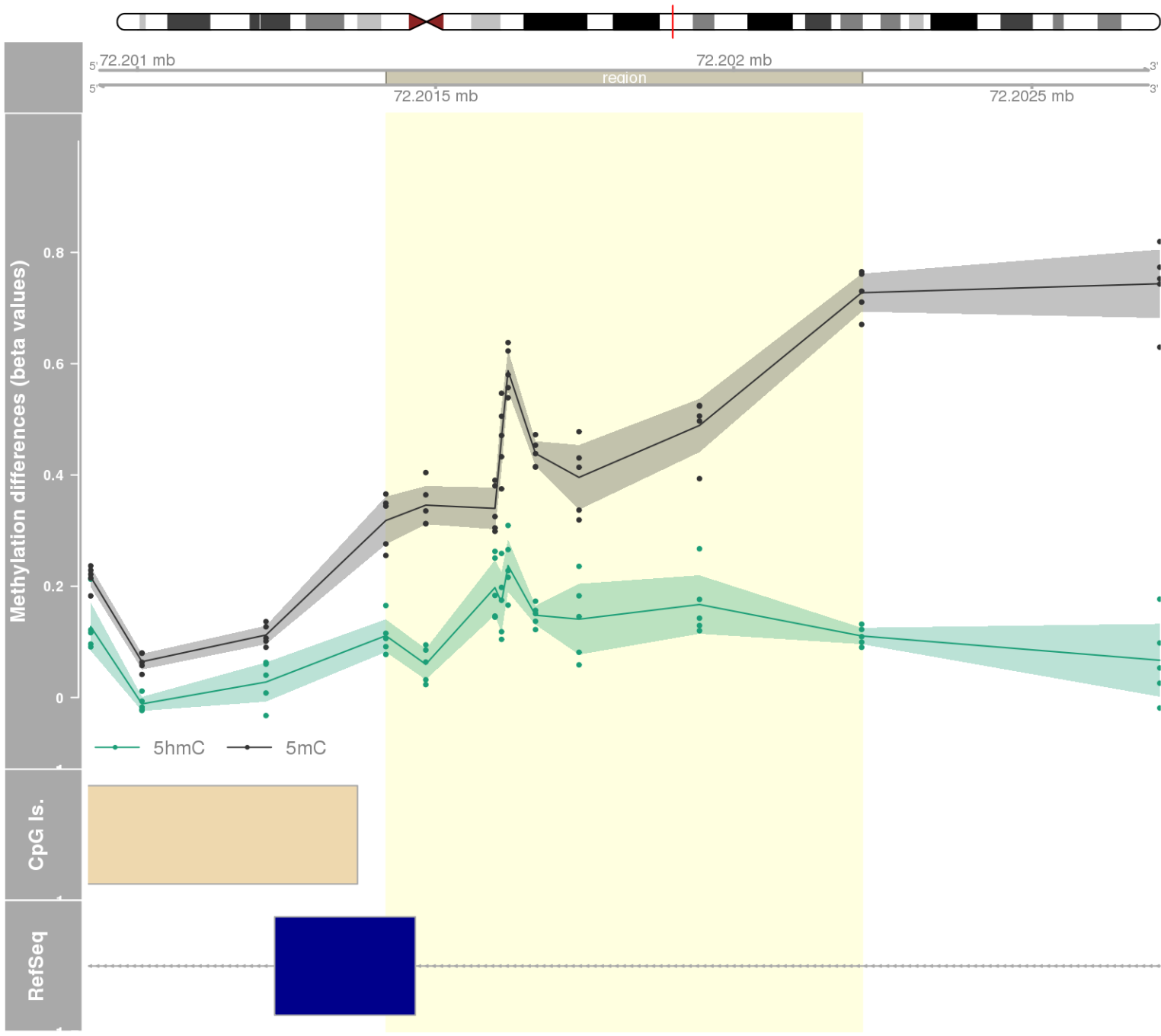
DMR 464 // chr10:102731539-102732323 // 784 pb. (7 probes) // pvalue: 0.012 // fwer: 0.173 // pvalueArea: 0.032 // fwerArea: 1
- genes: MRPL43 / SEMA4G -



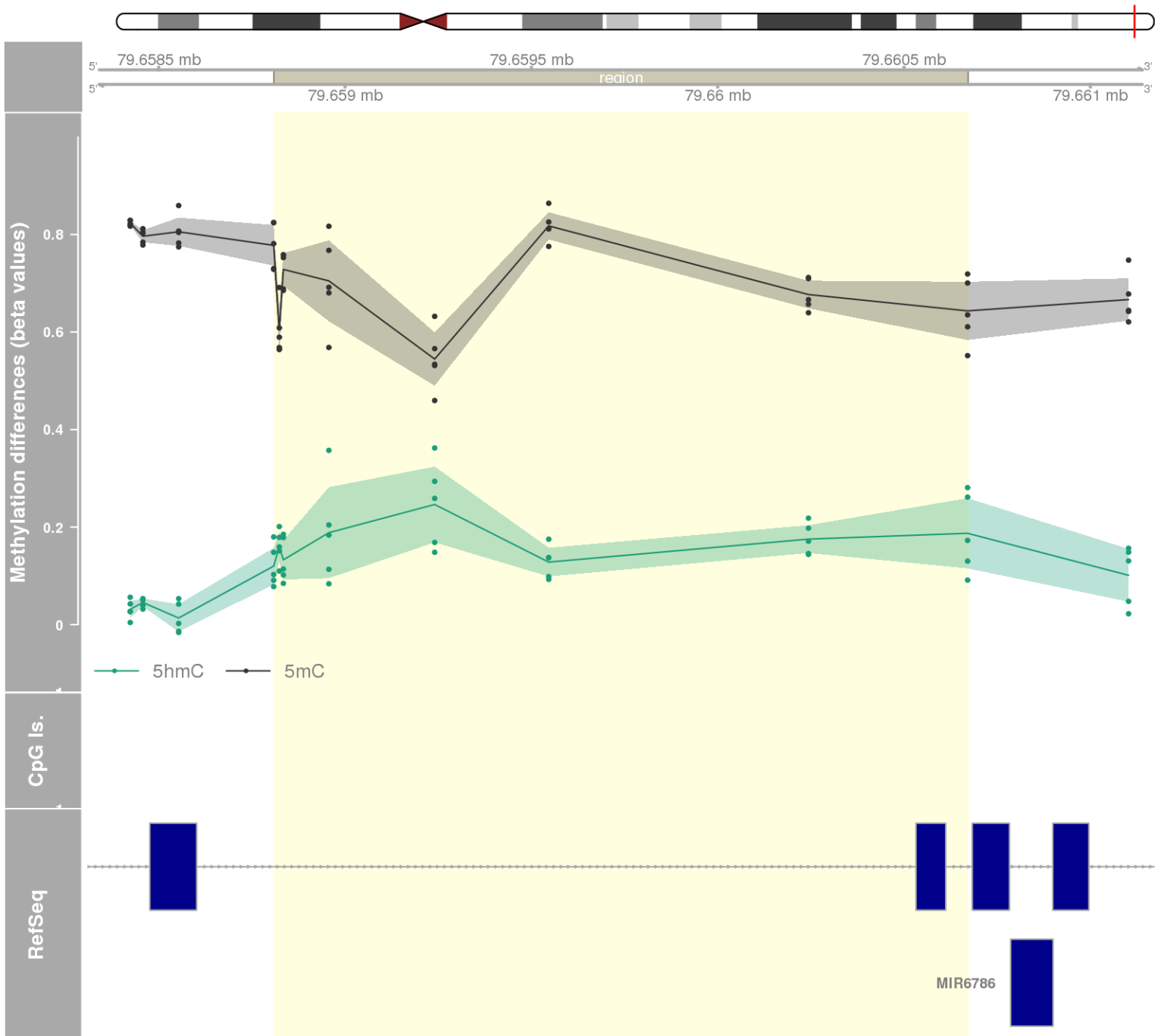
DMR 465 // chr8:41654331-41655826 // 1495 pb. (9 probes) // pvalue: 0.017 // fwer: 0.36 // pvalueArea: 0.032 // fwerArea: 1
- genes: ANK1 -



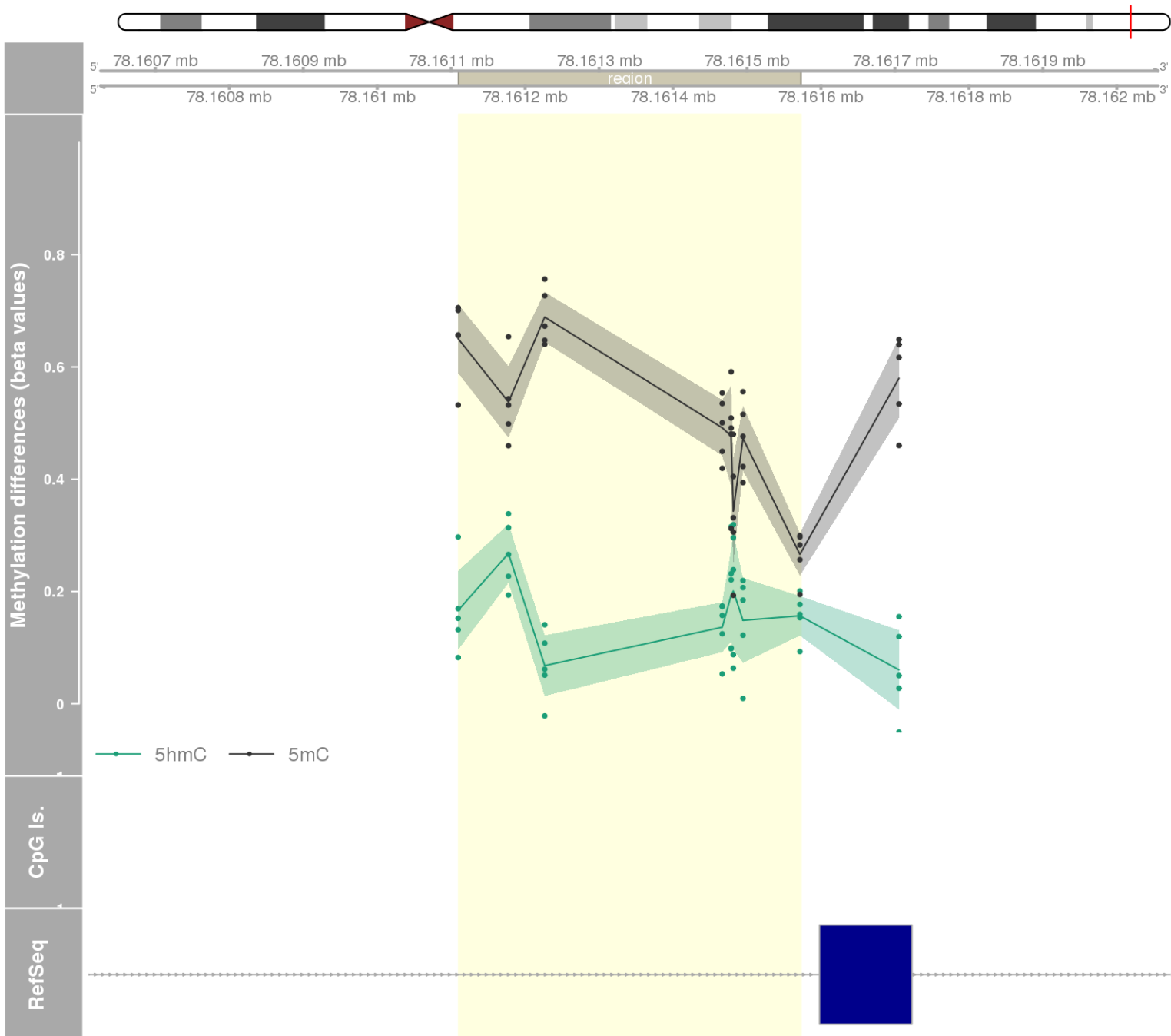
DMR 466 // chr10:72201417-72202215 // 798 pb. (9 probes) // pvalue: 0.017 // fwer: 0.36 // pvalueArea: 0.032 // fwerArea: 1
- genes: NODAL -



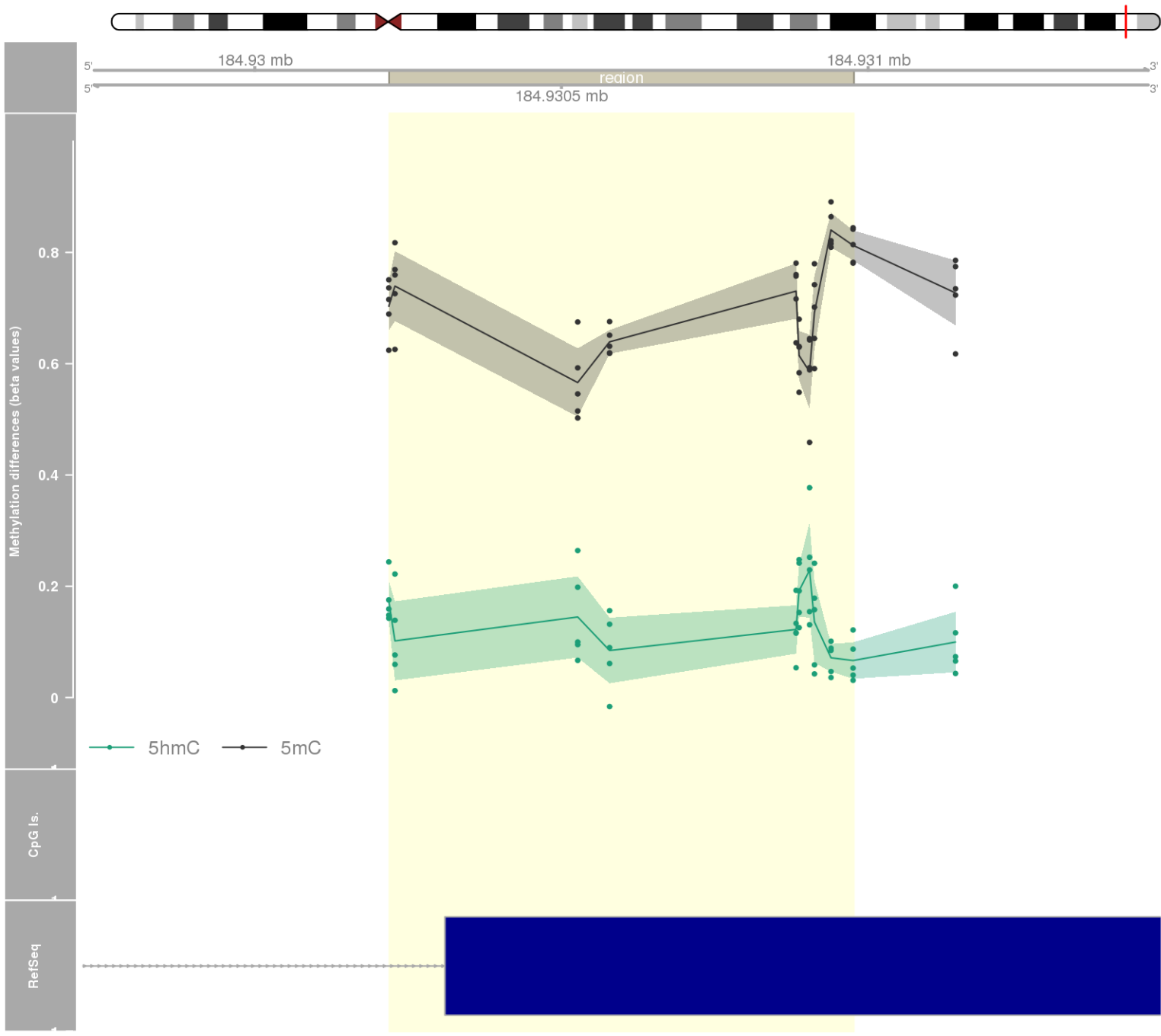
DMR 467 // chr17:79658809-79660671 // 1862 pb. (8 probes) // pvalue: 0.018 // fwer: 0.522 // pvalueArea: 0.032 // fwerArea: 1
- genes: HGS -



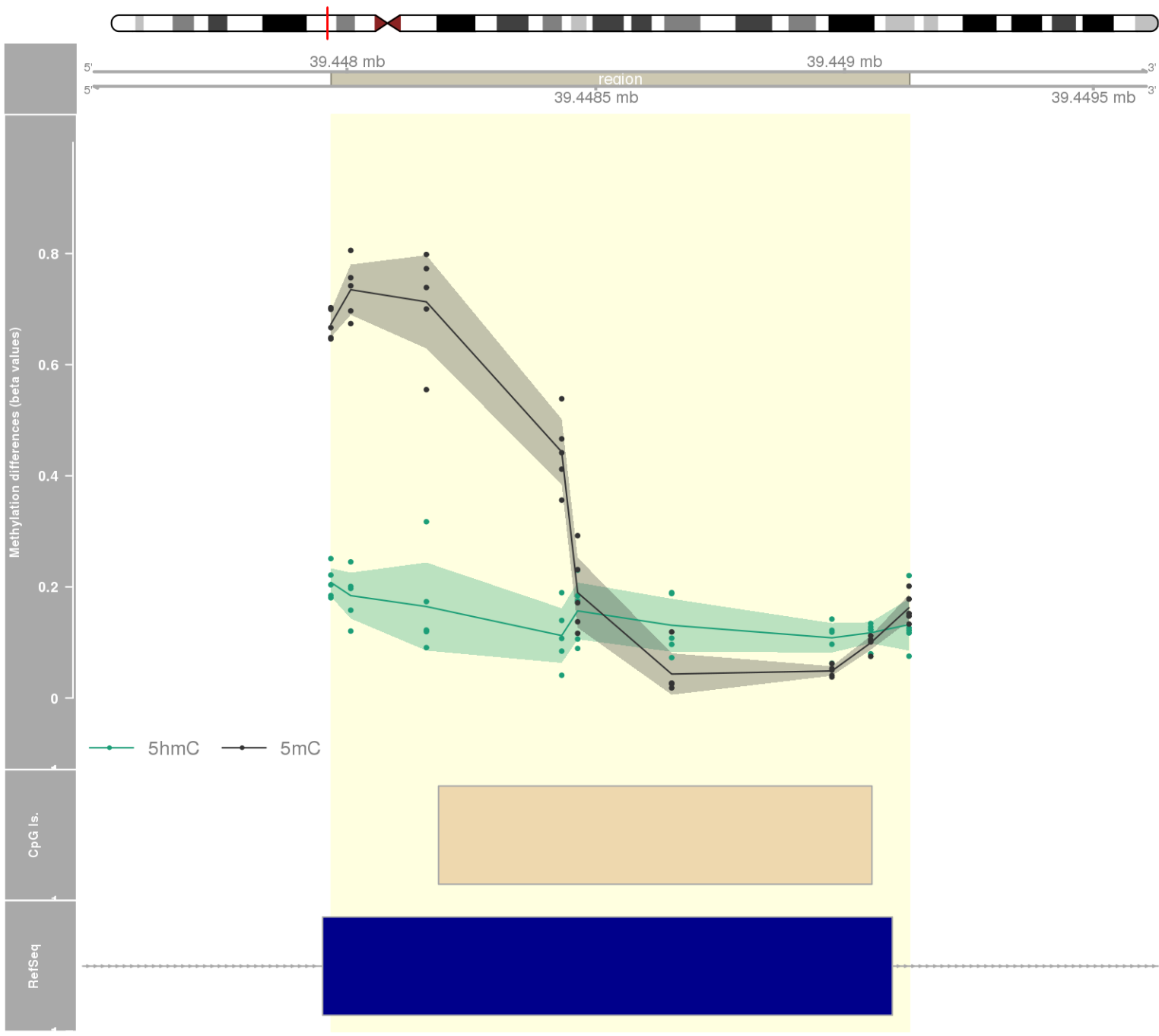
DMR 468 // chr17:78161110-78161572 // 462 pb. (8 probes) // pvalue: 0.018 // fwer: 0.522 // pvalueArea: 0.032 // fwerArea: 1
- genes: CARD14 -



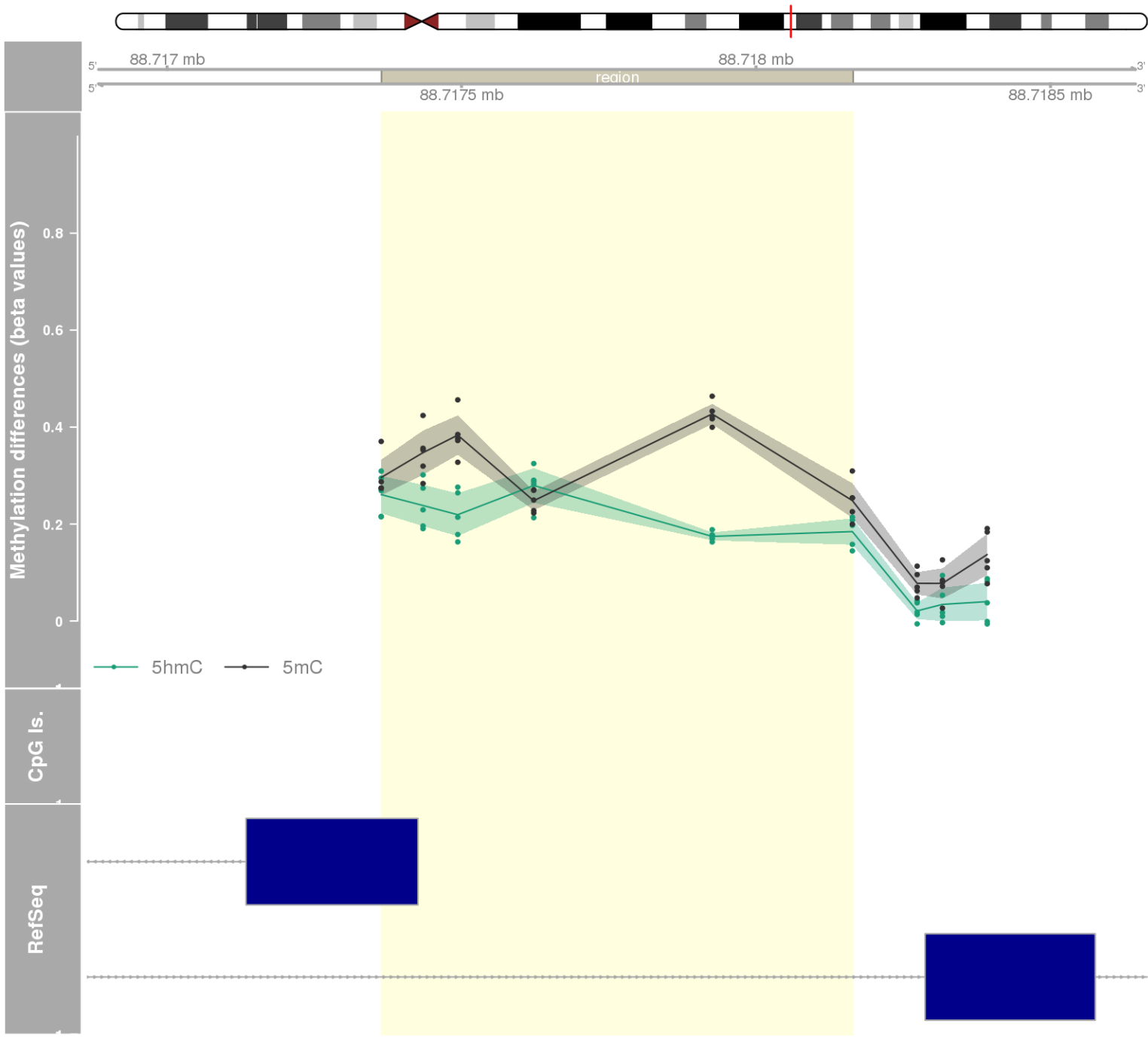
DMR 469 // chr4:184930219-184930976 // 757 pb. (10 probes) // pvalue: 0.015 // fwer: 0.36 // pvalueArea: 0.032 // fwerArea: 1
- genes: STOX2 -



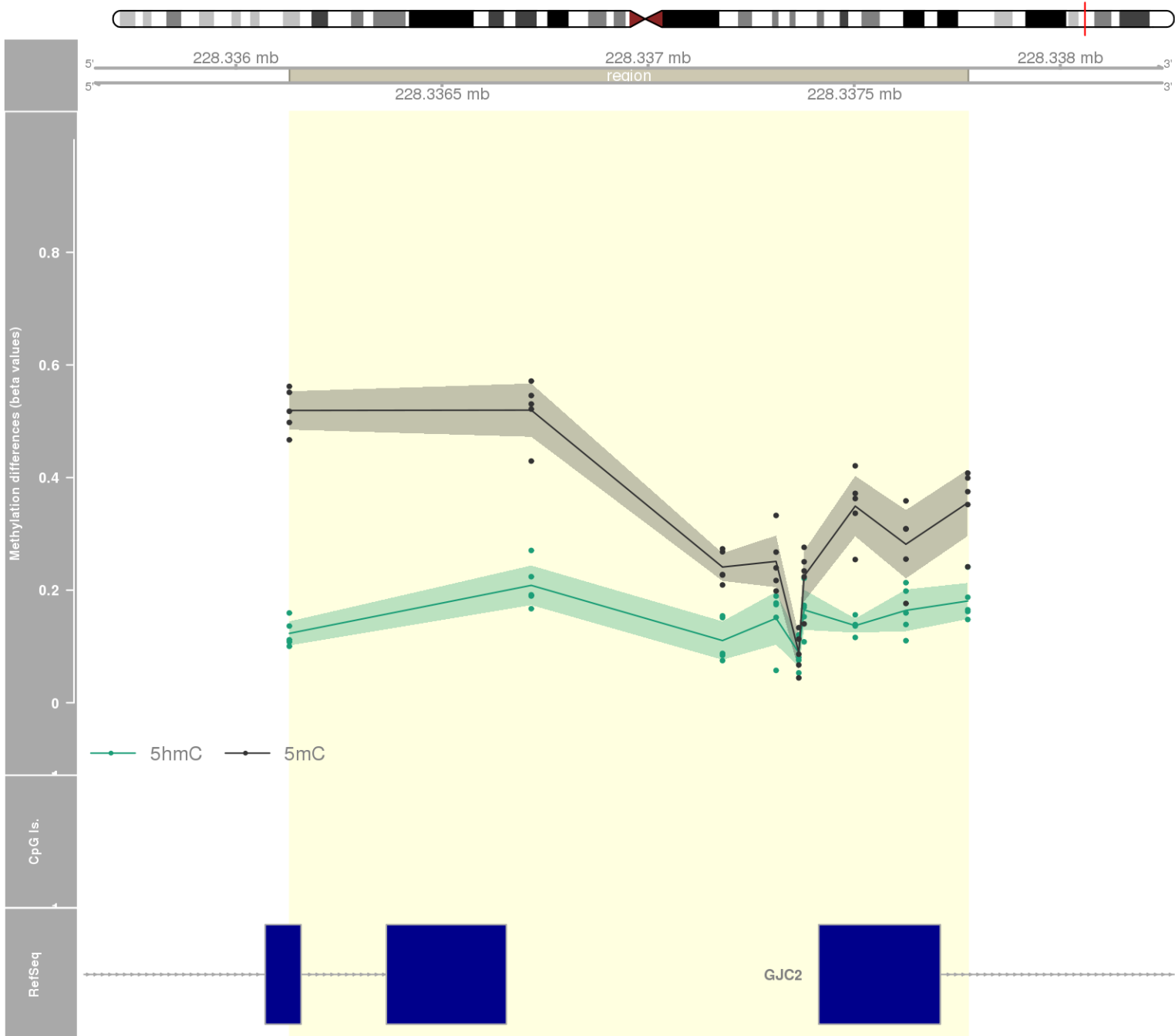
DMR 470 // chr4:39447968-39449130 // 1162 pb. (9 probes) // pvalue: 0.017 // fwer: 0.36 // pvalueArea: 0.033 // fwerArea: 1
- genes: KLB -



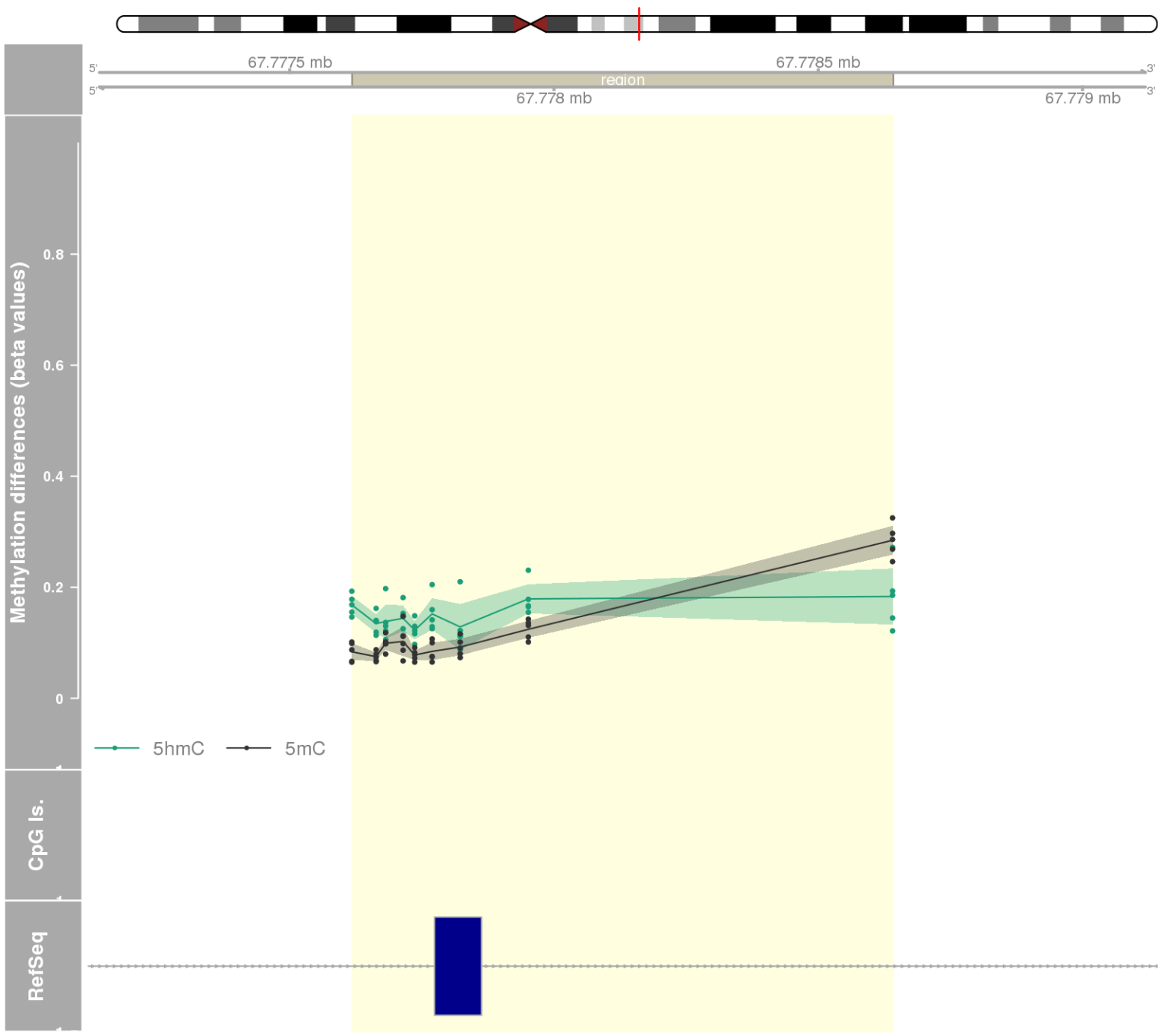
DMR 471 // chr10:88717364-88718164 // 800 pb. (6 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.033 // fwerArea: 1
- genes: MMRN2 -



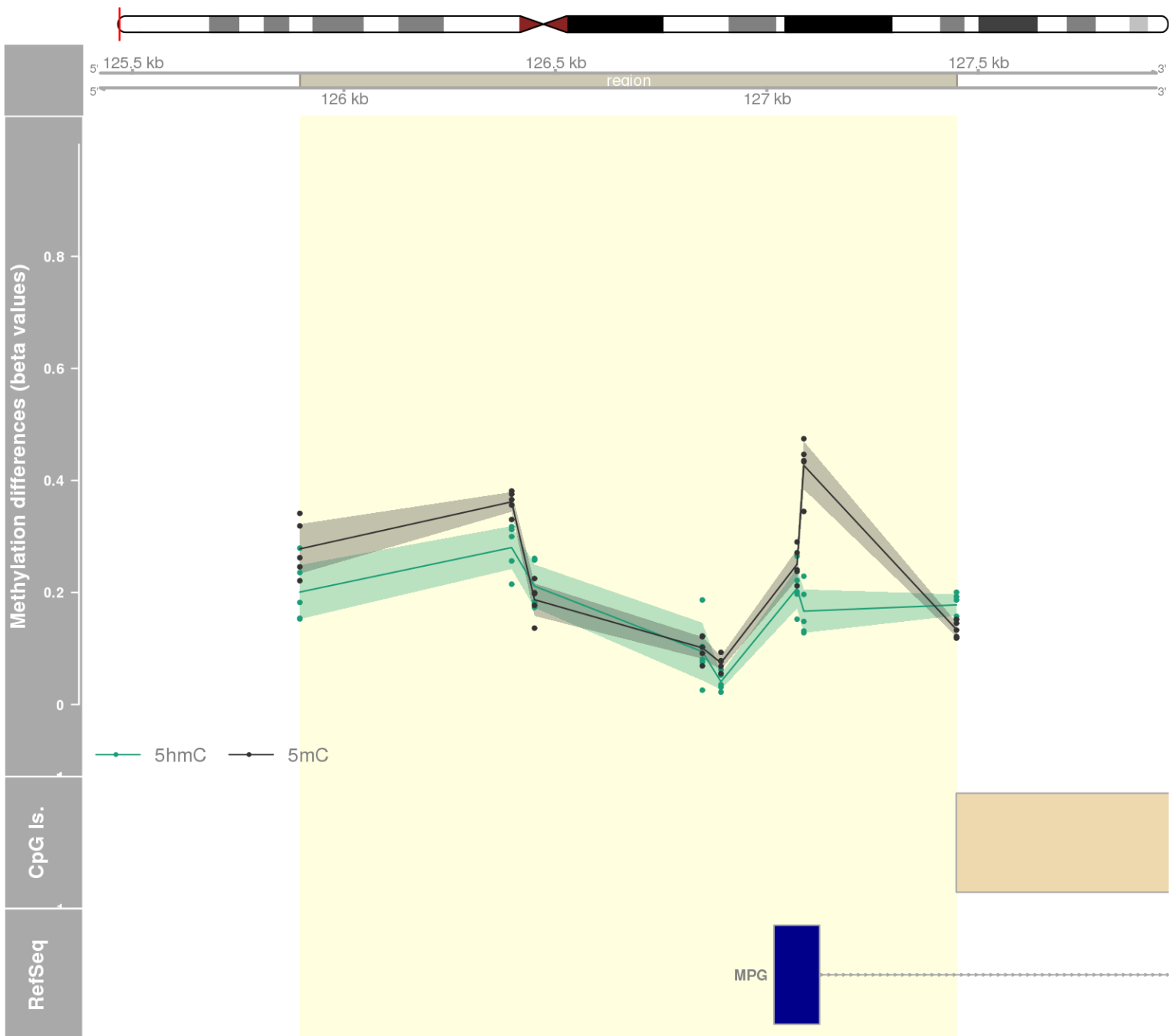
DMR 472 // chr1:228336130-228337776 // 1646 pb. (9 probes) // pvalue: 0.017 // fwer: 0.36 // pvalueArea: 0.033 // fwerArea: 1
- genes: GUK1 / GJC2 -



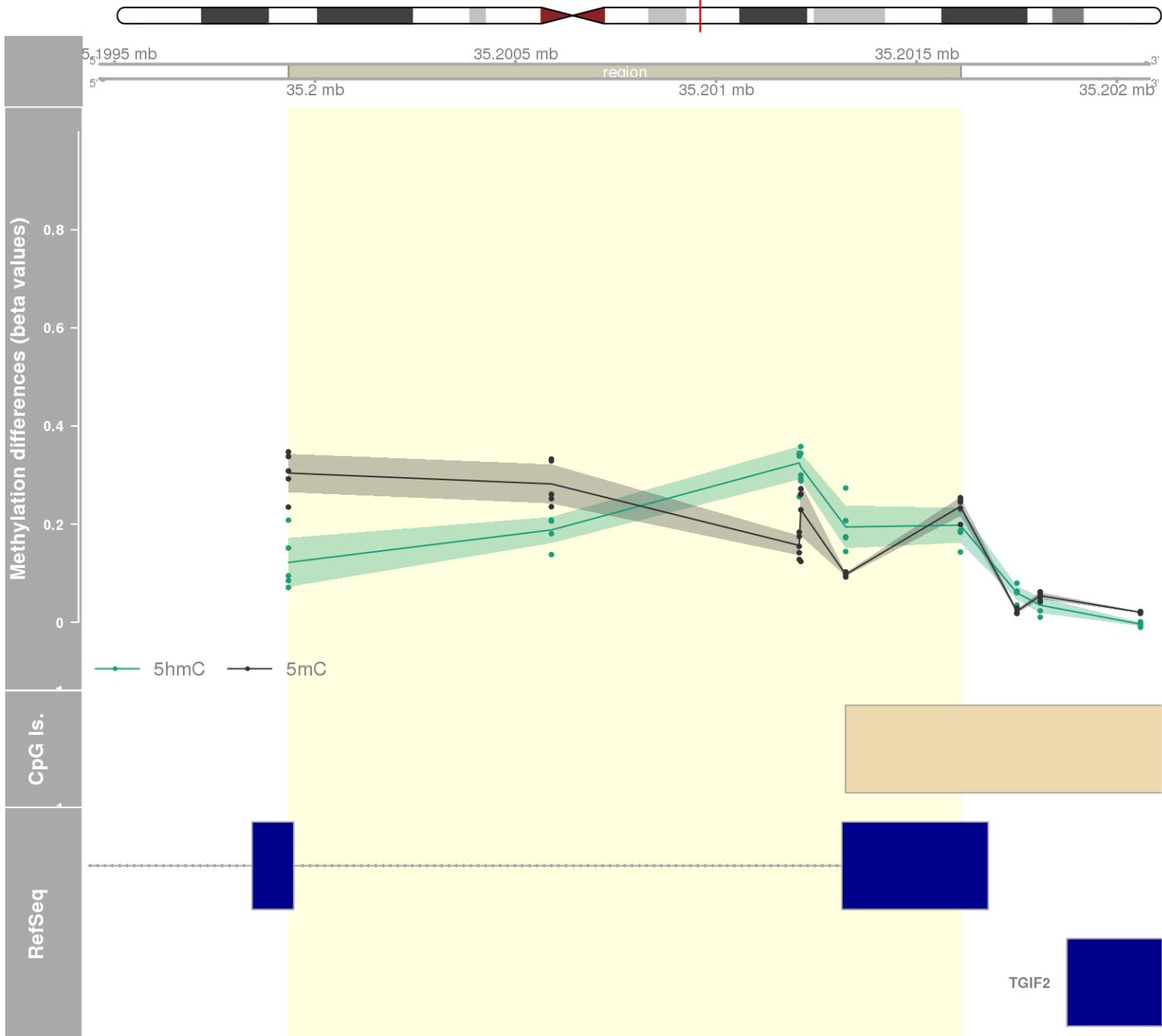
DMR 473 // chr11:67777618-67778641 // 1023 pb. (9 probes) // pvalue: 0.018 // fwer: 0.36 // pvalueArea: 0.033 // fwerArea: 1
- genes: ALDH3B1 -



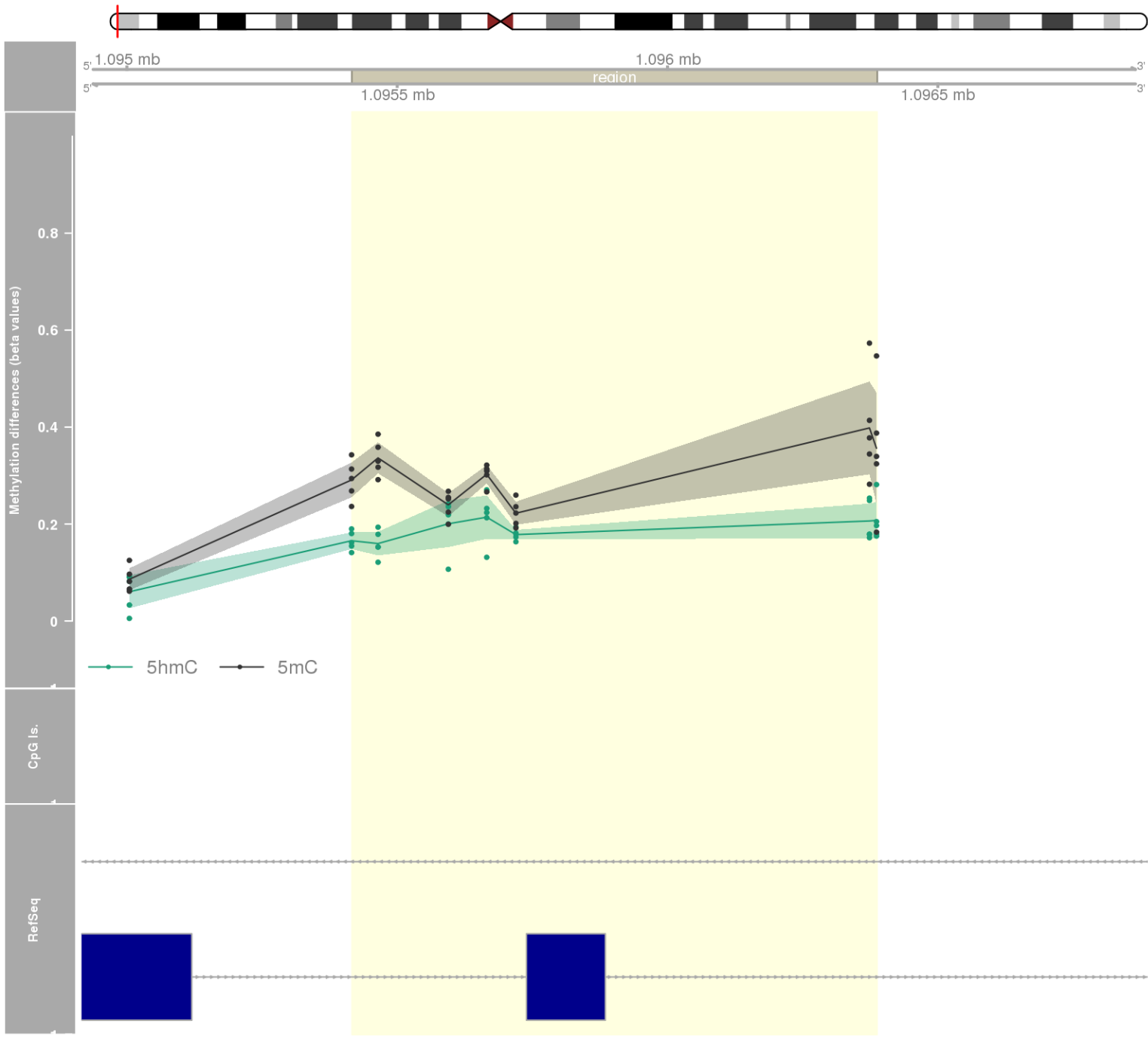
DMR 474 // chr16:125896-127449 // 1553 pb. (8 probes) // pvalue: 0.018 // fwer: 0.522 // pvalueArea: 0.033 // fwerArea: 1
- genes: RHBDF1 / MPG -



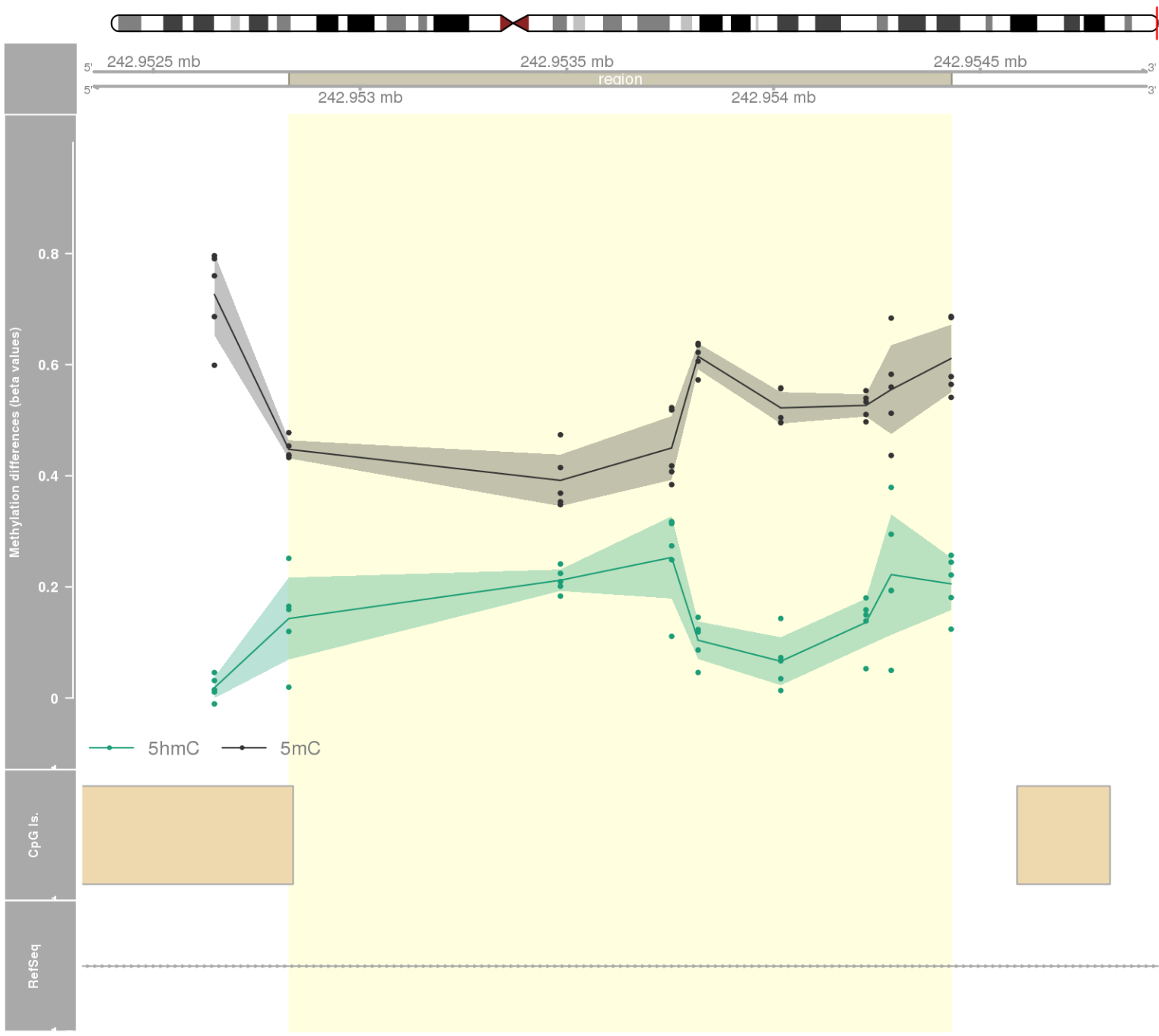
DMR 475 // chr20:35199934-35201610 // 1676 pb. (6 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.033 // fwerArea: 1



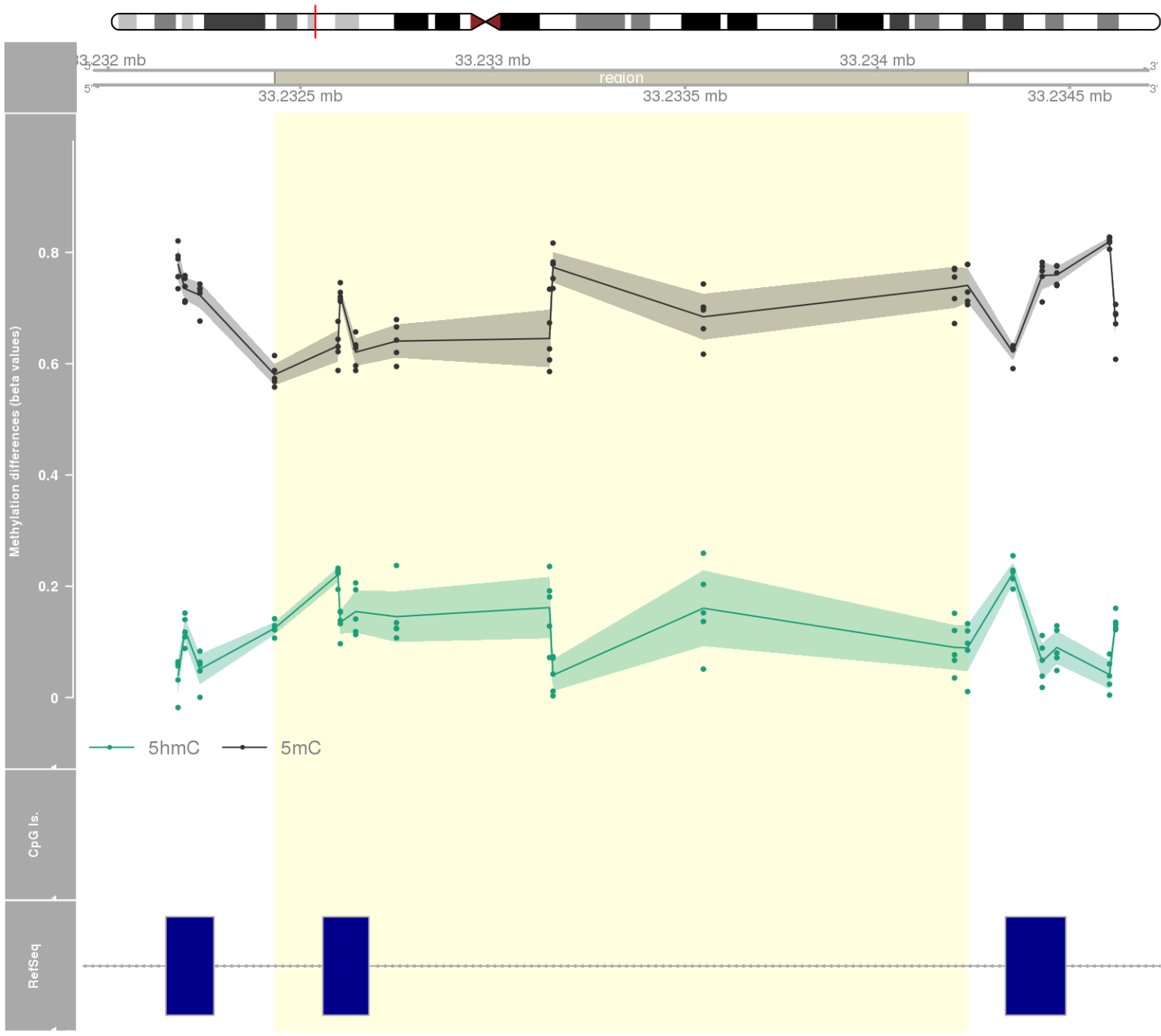
DMR 476 // chr7:1095416-1096387 // 971 pb. (7 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.033 // fwerArea: 1
- genes: C7orf50 / GPR146 -



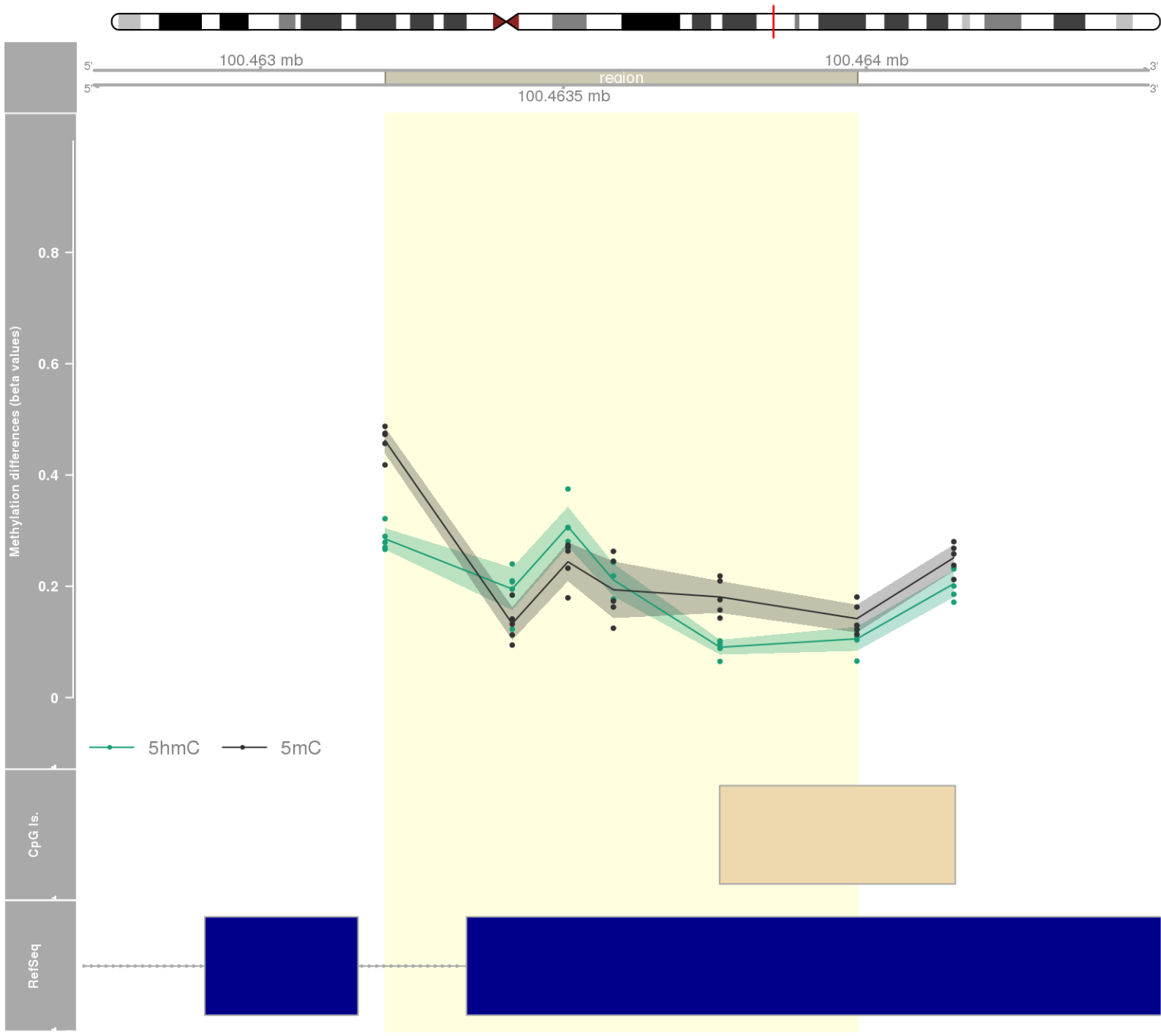
DMR 477 // chr2:242952828-242954430 // 1602 pb. (8 probes) // pvalue: 0.018 // fwer: 0.522 // pvalueArea: 0.033 // fwerArea: 1



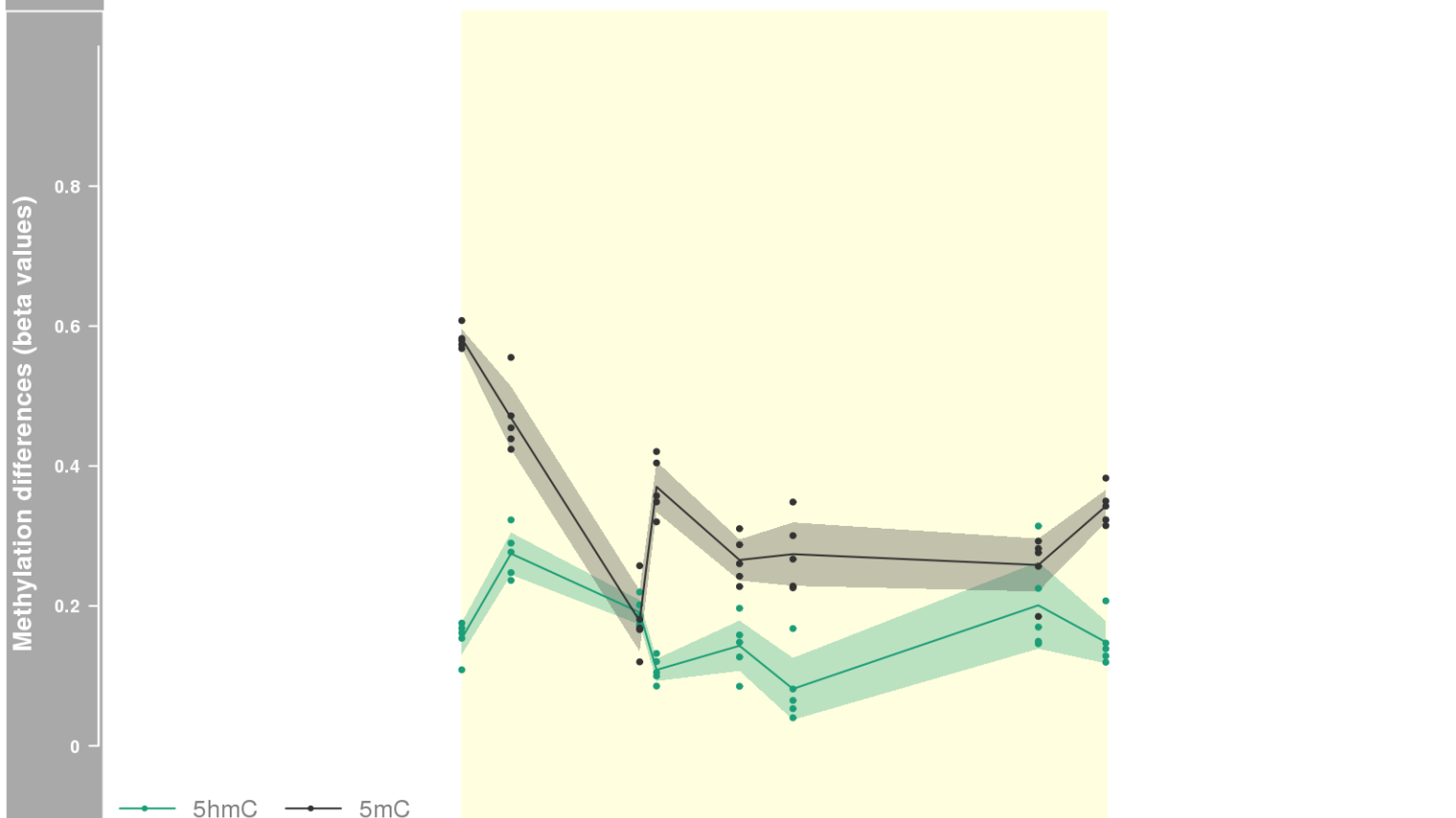
DMR 478 // chr6:33232433-33234235 // 1802 pb. (10 probes) // pvalue: 0.015 // fwer: 0.36 // pvalueArea: 0.033 // fwerArea: 1
- genes: VPS52 -



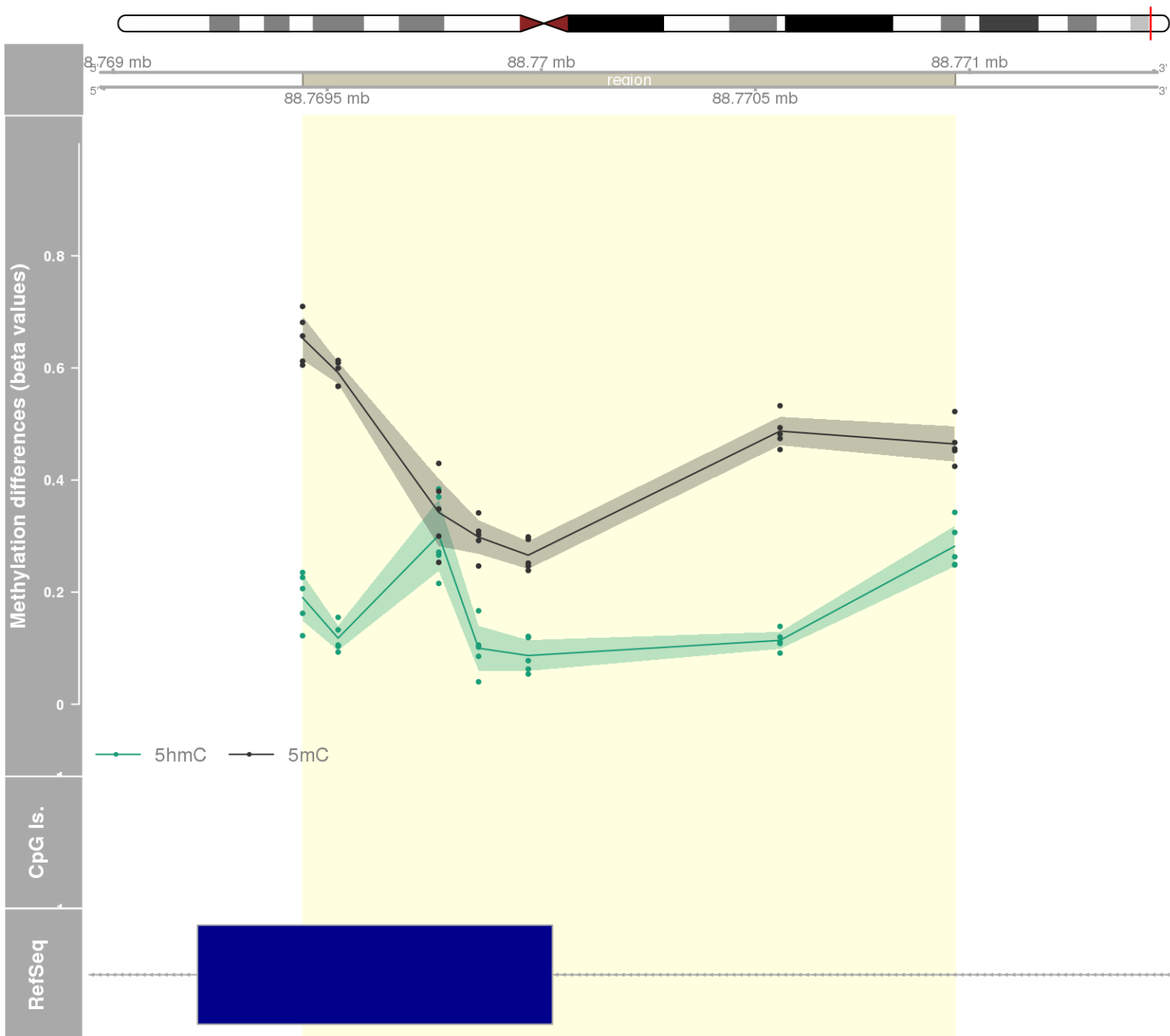
DMR 479 // chr7:100463206-100463985 // 779 pb. (6 probes) // pvalue: 0.005 // fwer: 0.173 // pvalueArea: 0.033 // fwerArea: 1
- genes: SLC12A9 -



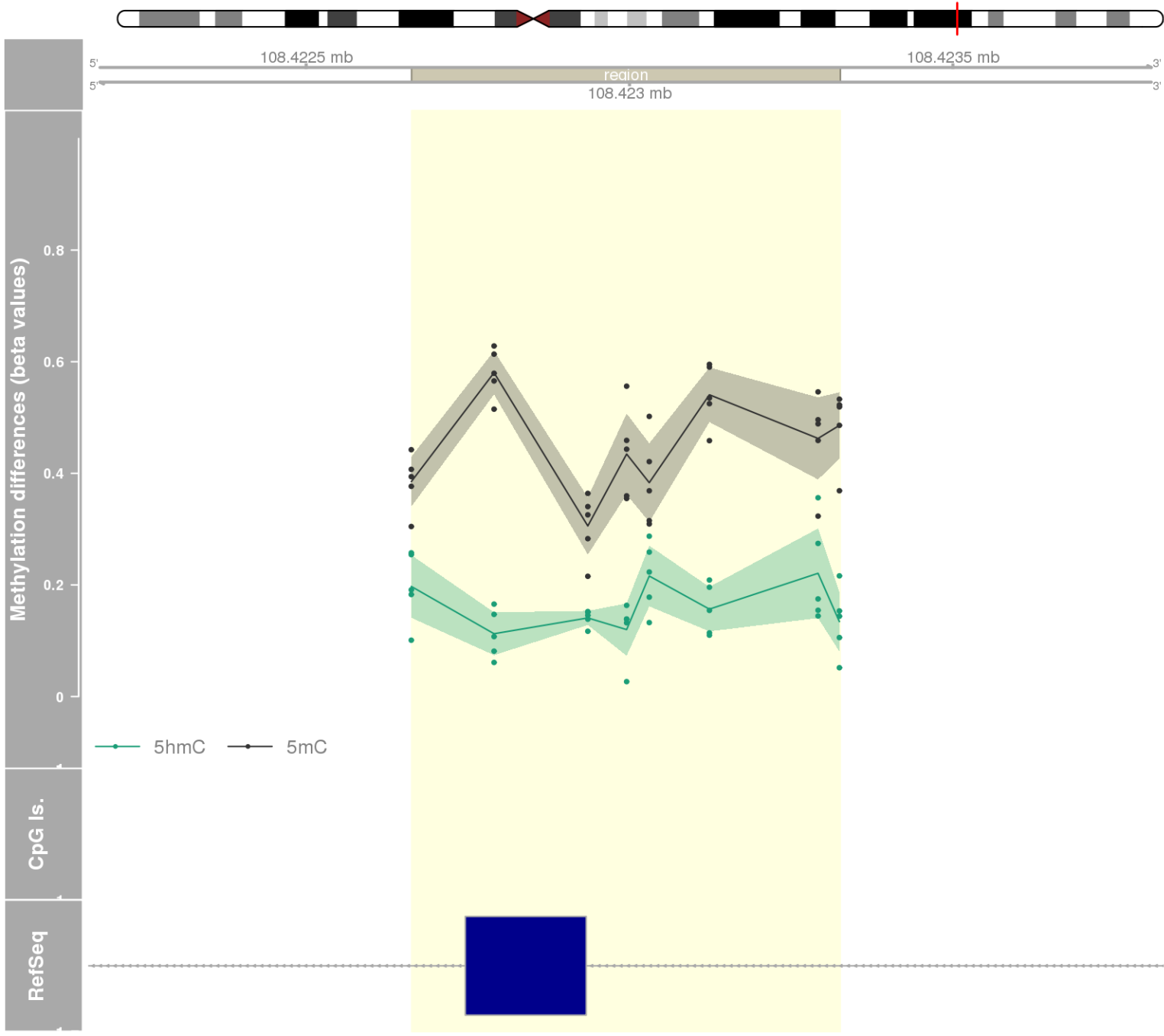
DMR 480 // chr17:72462164-72463080 // 916 pb. (8 probes) // pvalue: 0.018 // fwer: 0.522 // pvalueArea: 0.033 // fwerArea: 1
- genes: CD300A -



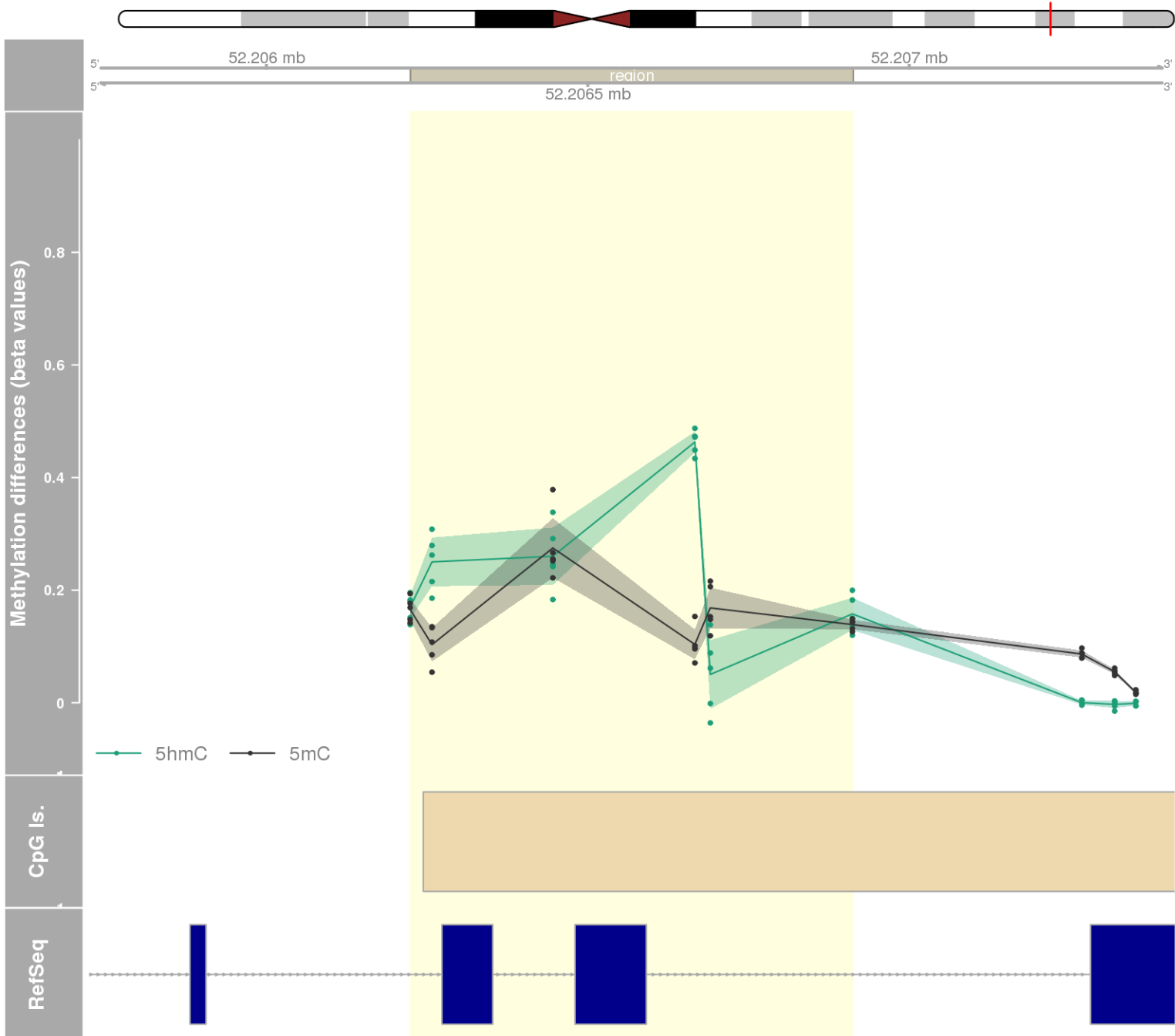
DMR 481 // chr16:88769443-88770966 // 1523 pb. (7 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.033 // fwerArea: 1
- genes: RNF166 -



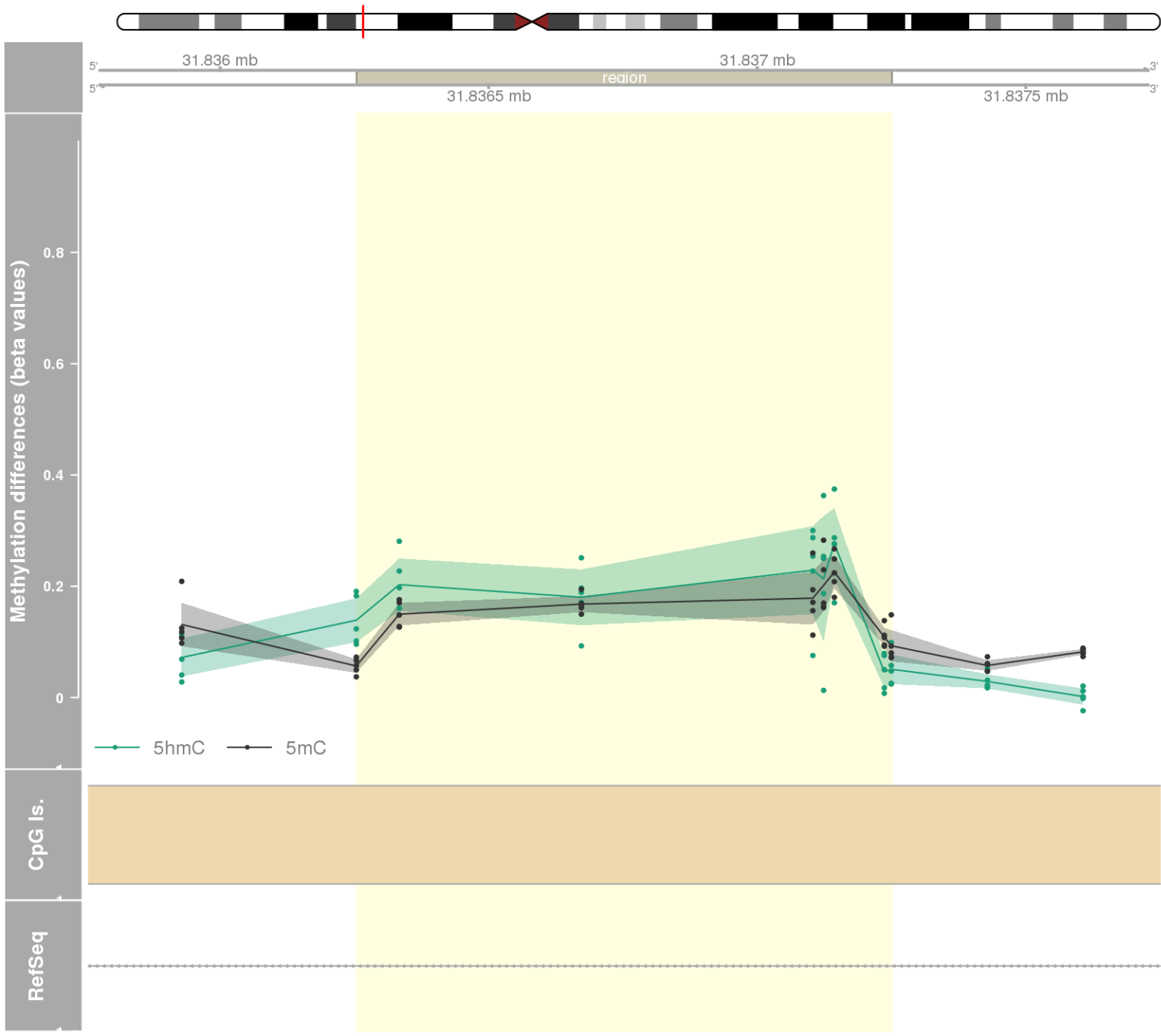
DMR 482 // chr11:108422663-108423325 // 662 pb. (8 probes) // pvalue: 0.018 // fwer: 0.522 // pvalueArea: 0.033 // fwerArea: 1
- genes: EXPH5 -



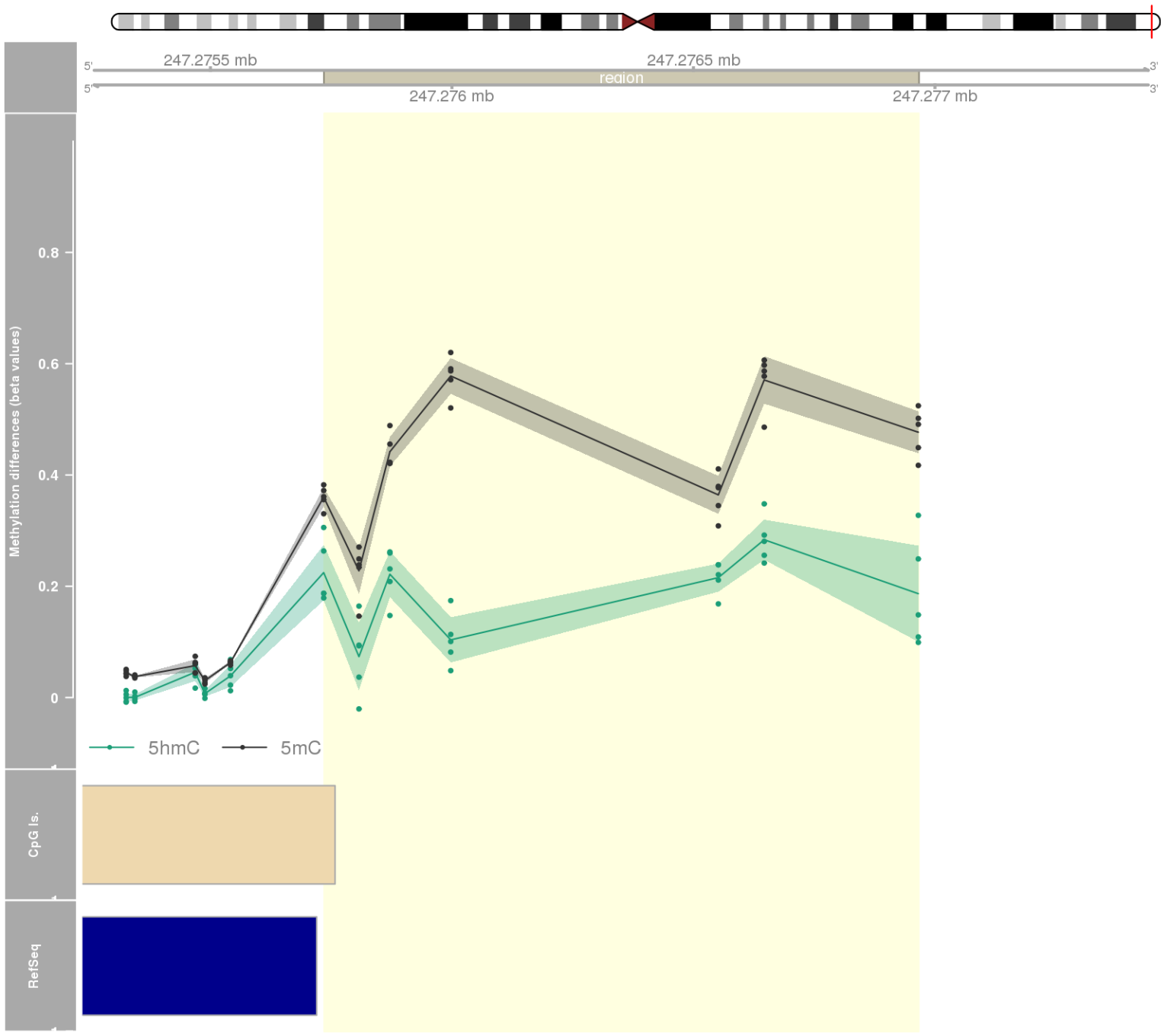
DMR 483 // chr19:52206224-52206912 // 688 pb. (6 probes) // pvalue: 0.006 // fwer: 0.173 // pvalueArea: 0.033 // fwerArea: 1
- genes: SPACA6P -



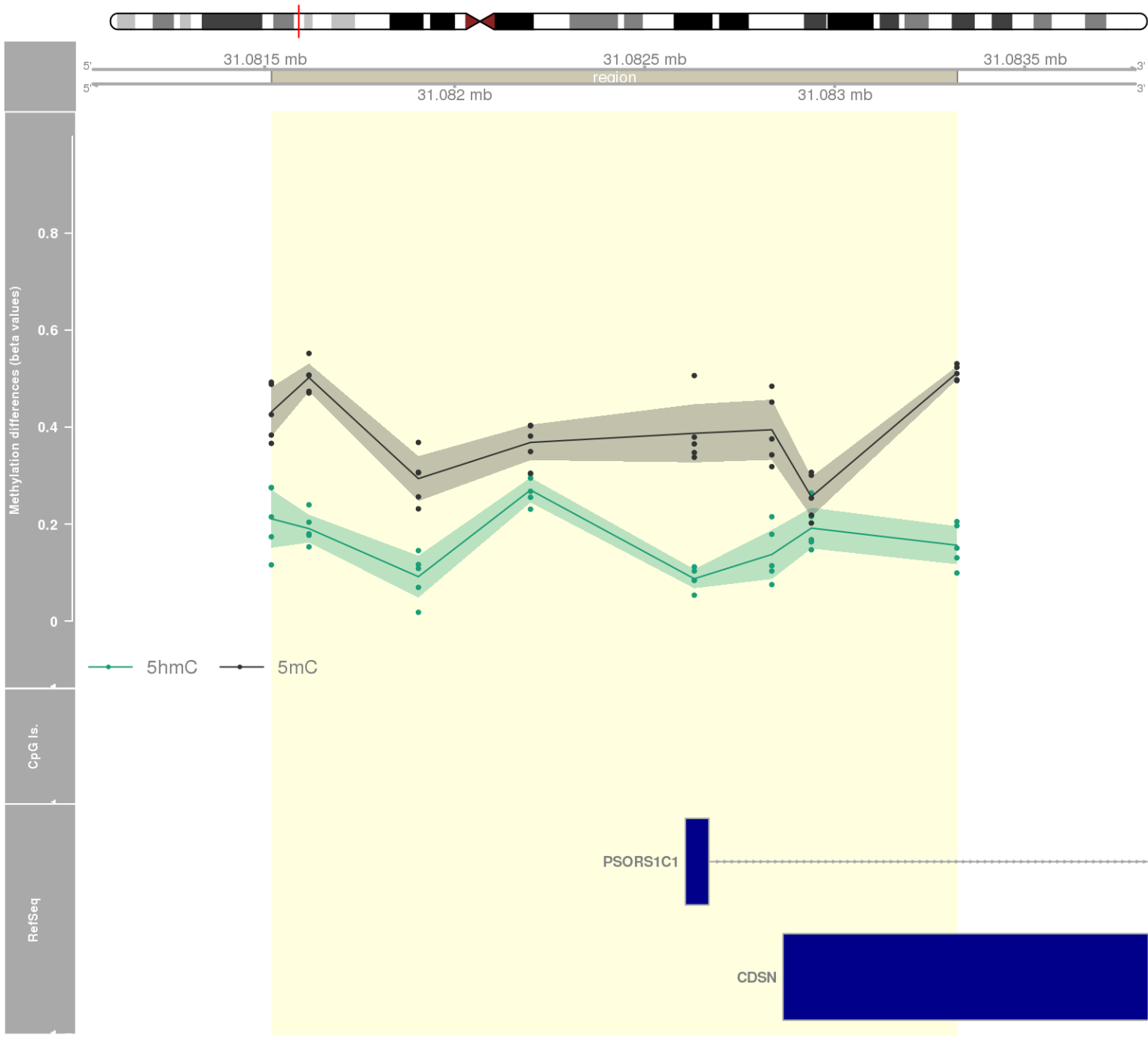
DMR 484 // chr11:31836254-31837250 // 996 pb. (8 probes) // pvalue: 0.018 // fwer: 0.522 // pvalueArea: 0.033 // fwerArea: 1
- genes: PAX6 / RCN1 -



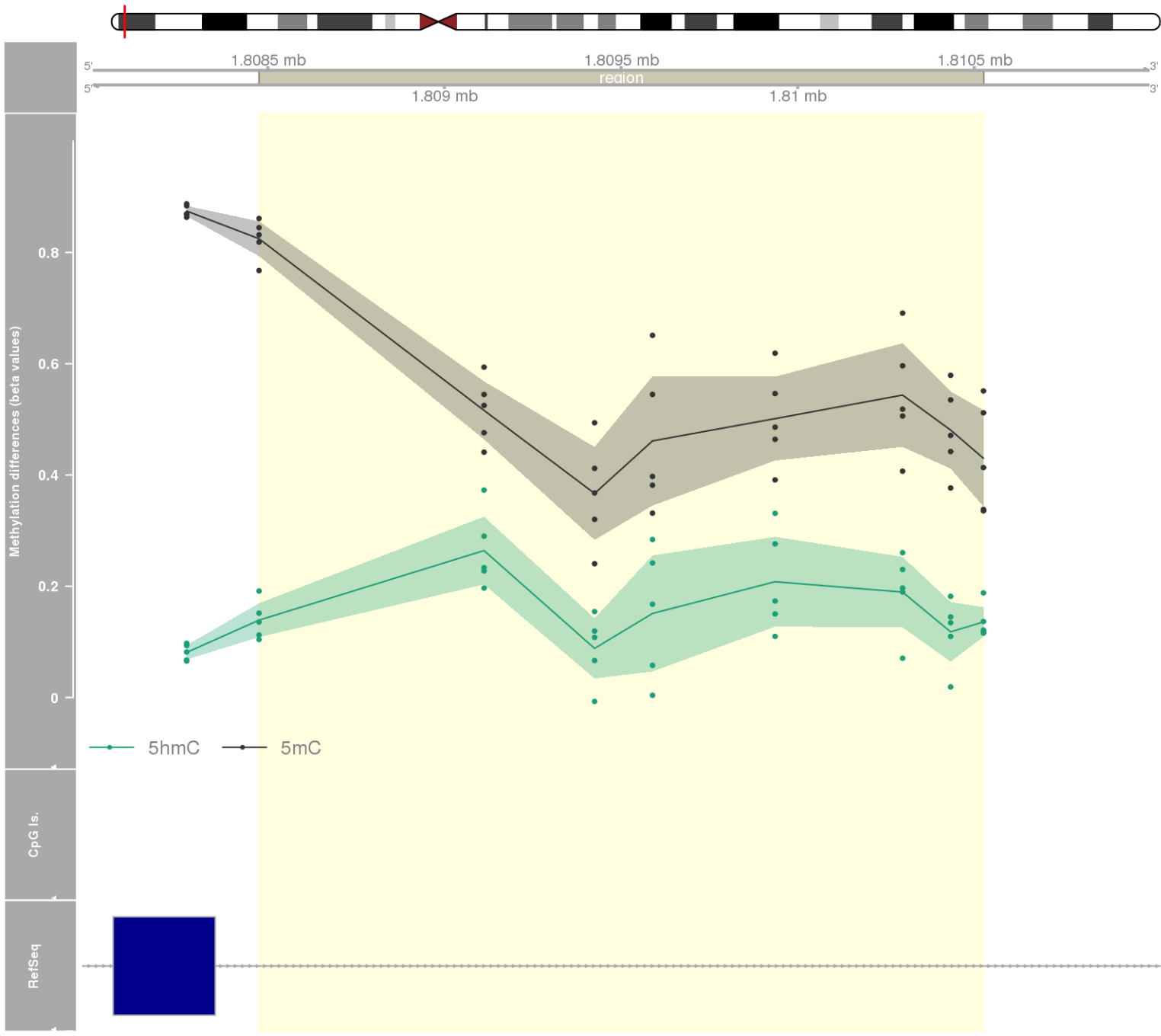
DMR 485 // chr1:247275735-247276966 // 1231 pb. (7 probes) // pvalue: 0.013 // fwer: 0.36 // pvalueArea: 0.034 // fwerArea: 1



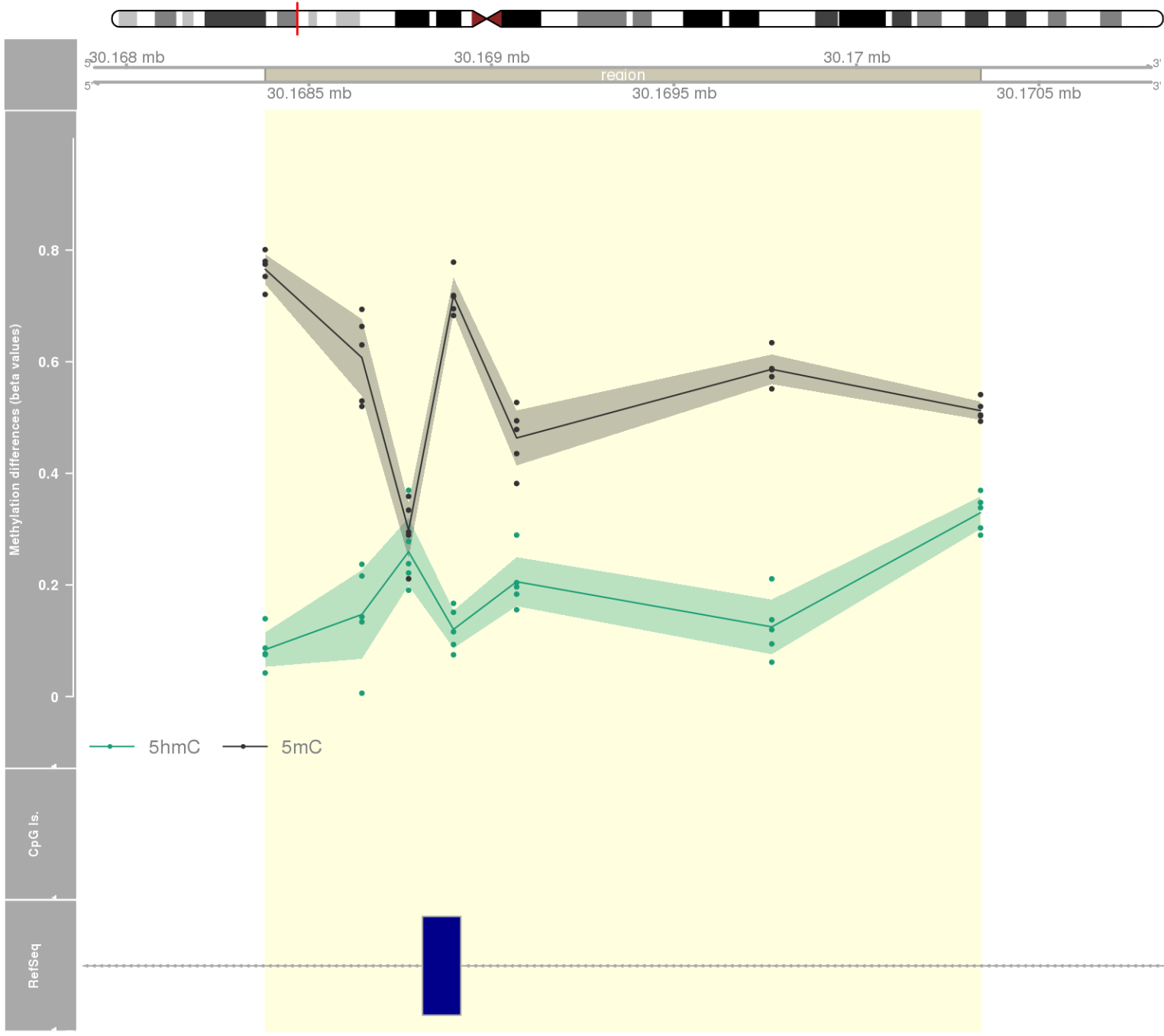
DMR 486 // chr6:31081518-31083322 // 1804 pb. (8 probes) // pvalue: 0.019 // fwer: 0.522 // pvalueArea: 0.034 // fwerArea: 1
- genes: PSORS1C1 / CDSN -



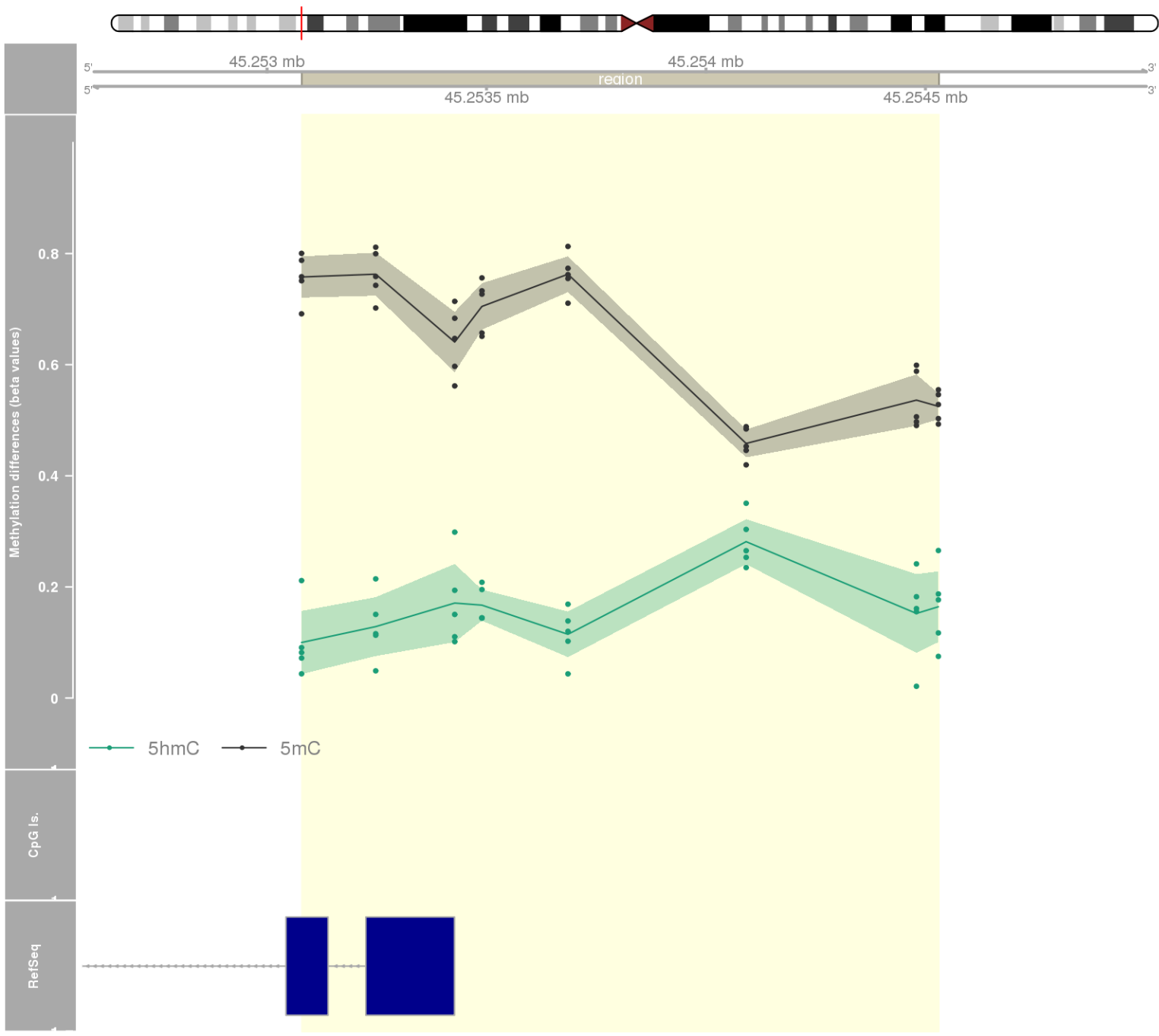
DMR 487 // chr8:1808475-1810526 // 2051 pb. (8 probes) // pvalue: 0.019 // fwer: 0.522 // pvalueArea: 0.034 // fwerArea: 1
- genes: ARHGEF10 -



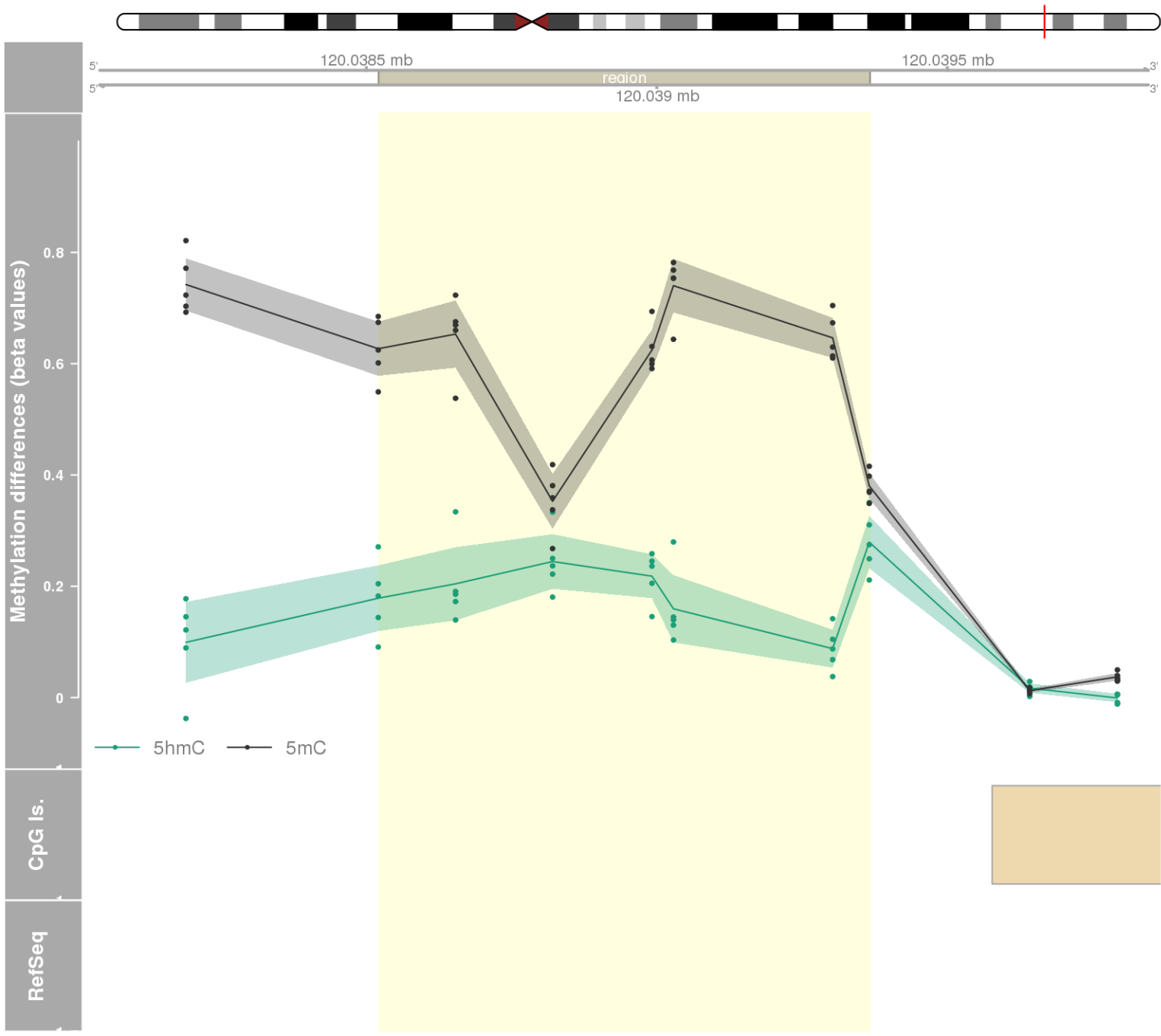
DMR 488 // chr6:30168380-30170340 // 1960 pb. (7 probes) // pvalue: 0.014 // fwer: 0.36 // pvalueArea: 0.034 // fwerArea: 1
- genes: TRIM26 -



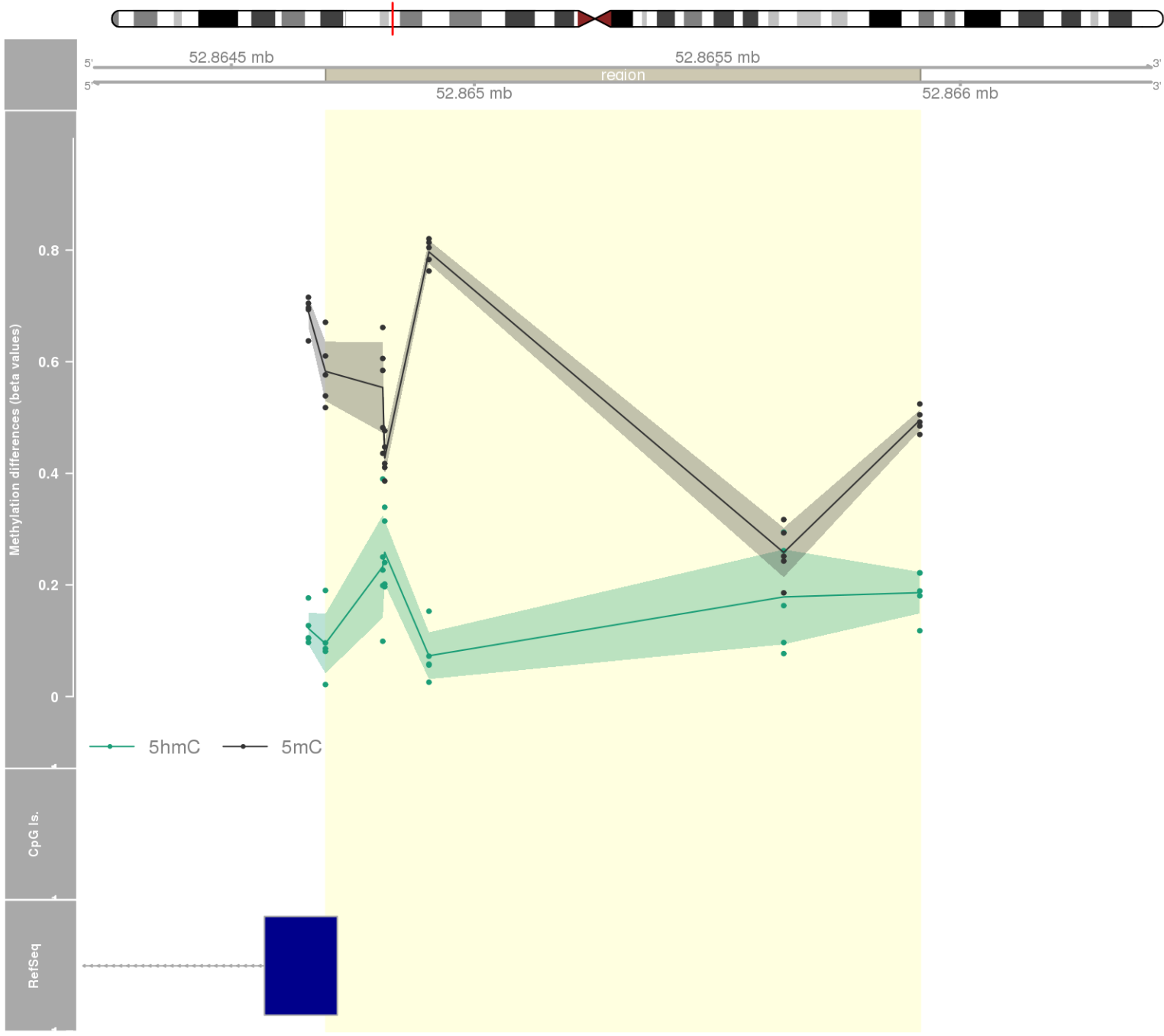
DMR 489 // chr1:45253079-45254530 // 1451 pb. (8 probes) // pvalue: 0.019 // fwer: 0.522 // pvalueArea: 0.034 // fwerArea: 1
- genes: BEST4 -



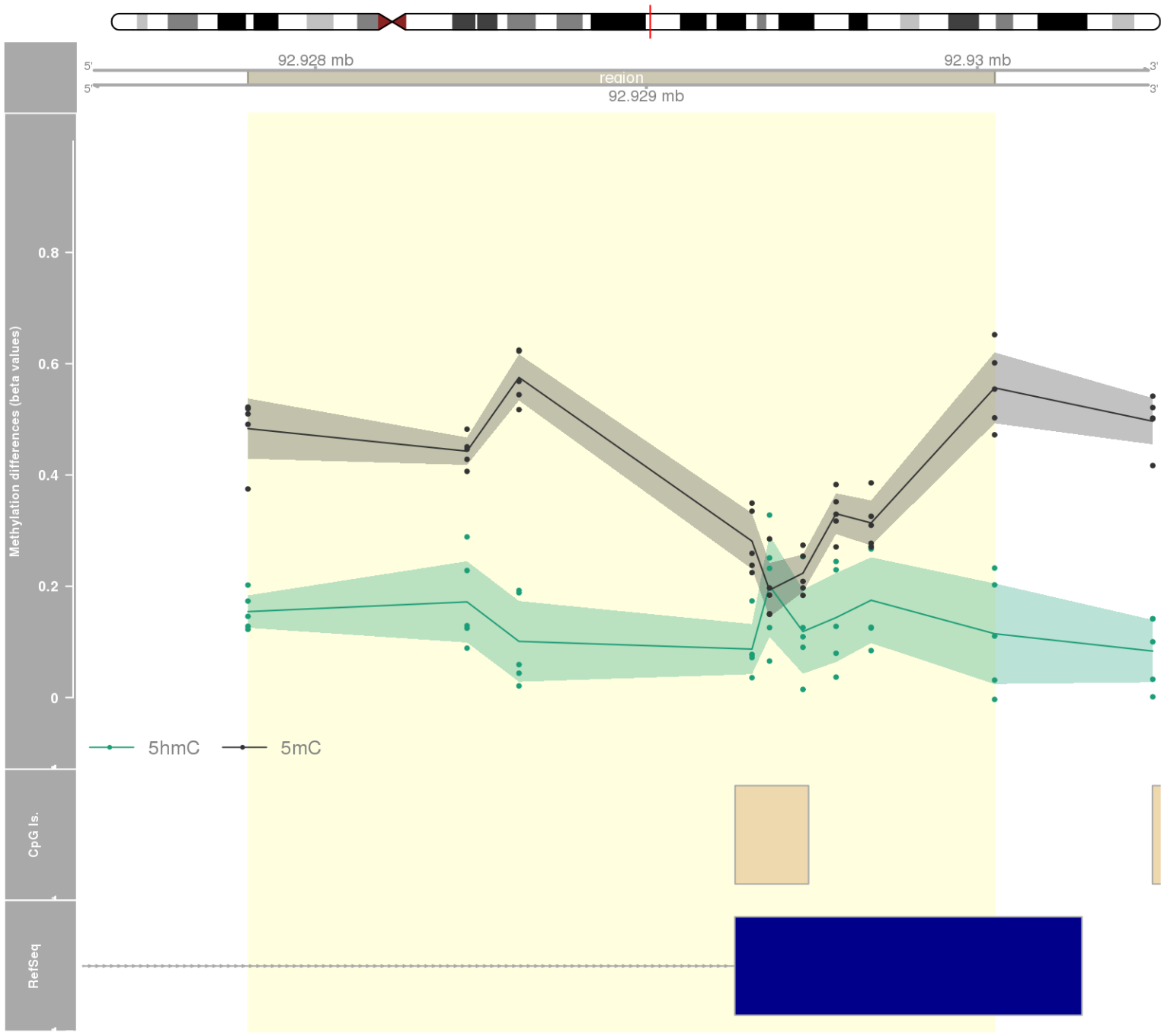
DMR 490 // chr11:120038521-120039366 // 845 pb. (7 probes) // pvalue: 0.014 // fwer: 0.36 // pvalueArea: 0.034 // fwerArea: 1
- genes: TRIM29 -



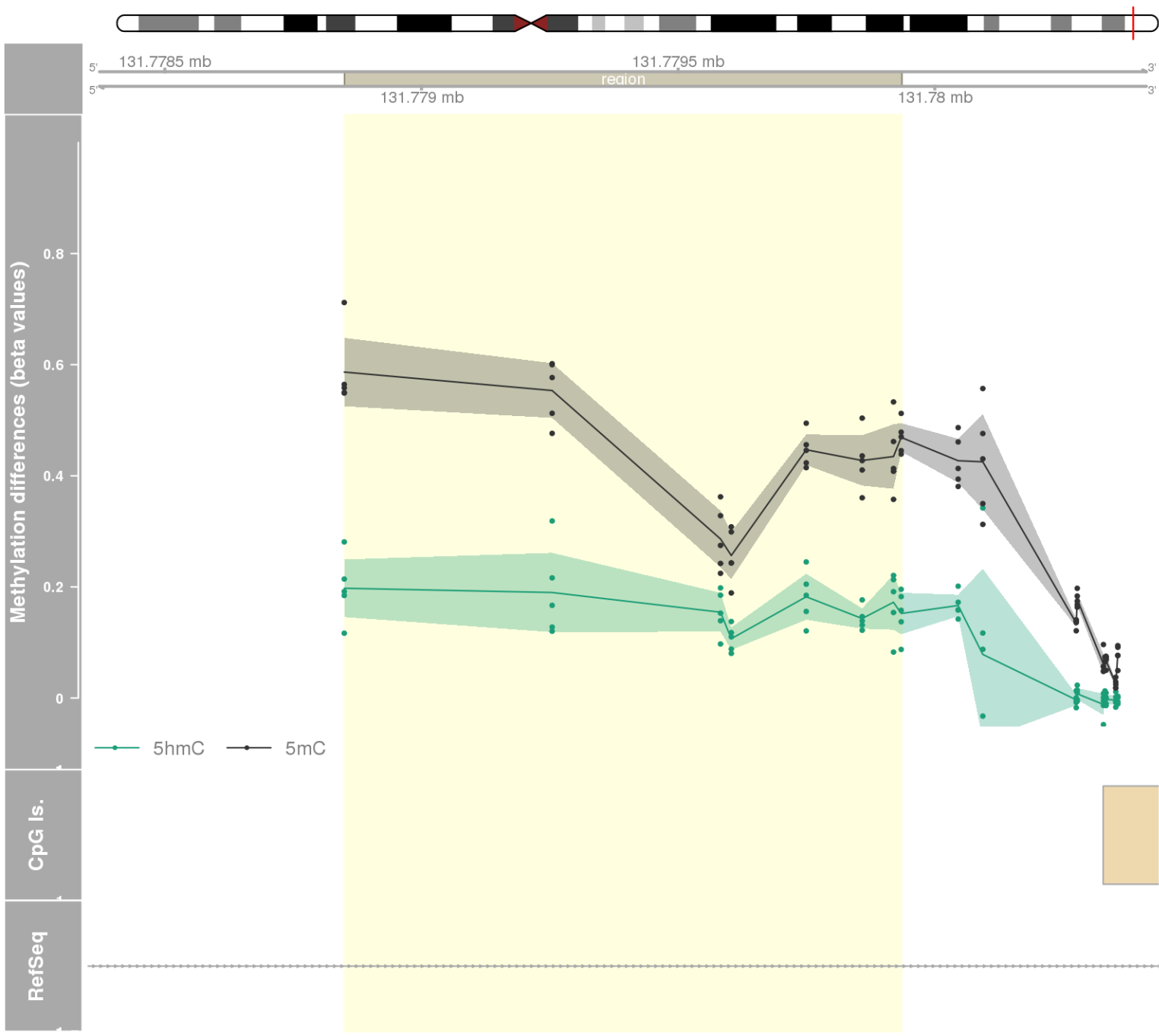
DMR 491 // chr3:52864694-52865917 // 1223 pb. (6 probes) // pvalue: 0.006 // fwer: 0.173 // pvalueArea: 0.034 // fwerArea: 1
- genes: ITIH4 -



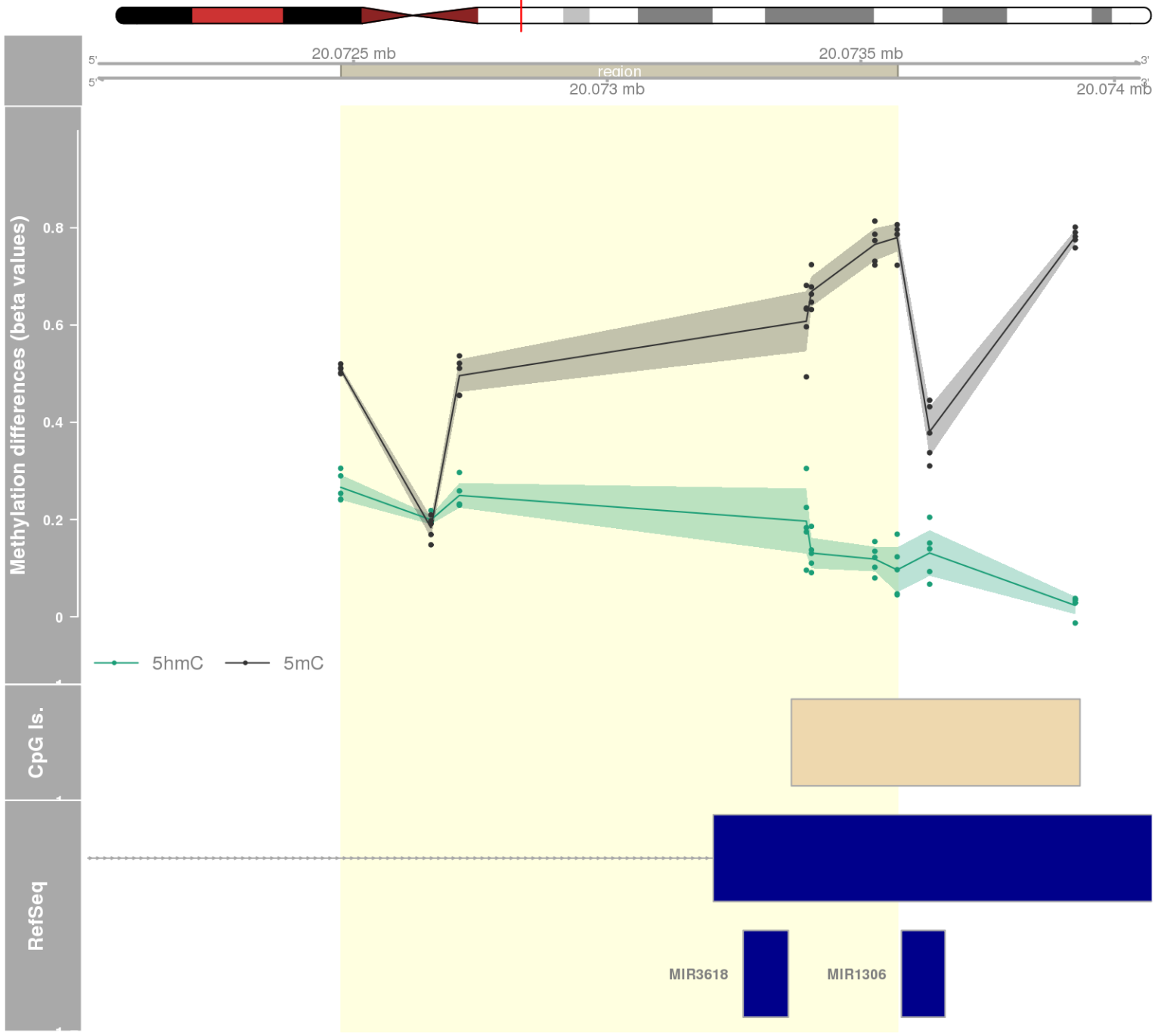
DMR 492 // chr5:92927796-92930052 // 2256 pb. (9 probes) // pvalue: 0.018 // fwer: 0.36 // pvalueArea: 0.034 // fwerArea: 1
- genes: NR2F1 -



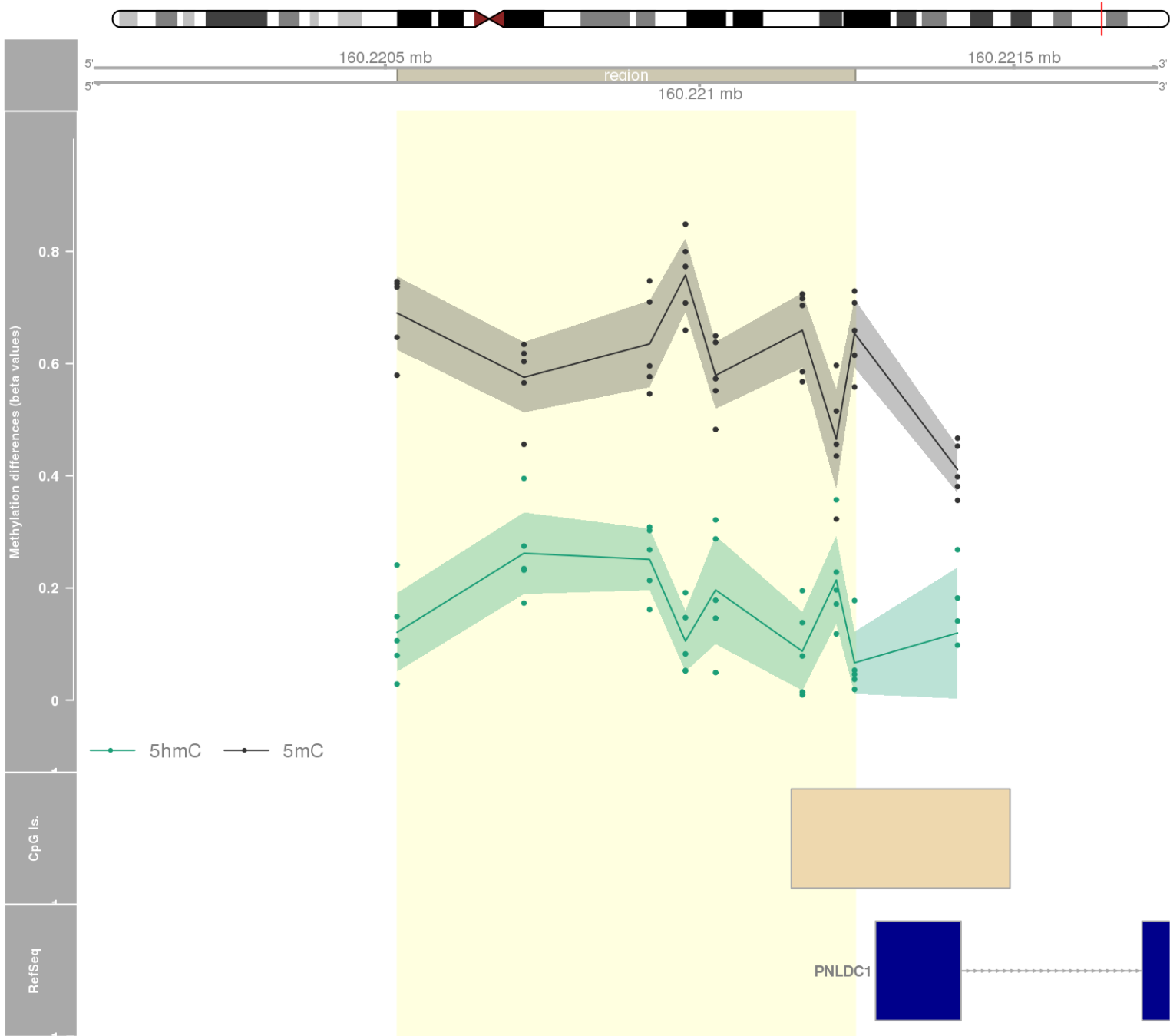
DMR 493 // chr11:131778850-131779935 // 1085 pb. (8 probes) // pvalue: 0.019 // fwer: 0.522 // pvalueArea: 0.034 // fwerArea: 1
- genes: NTM -



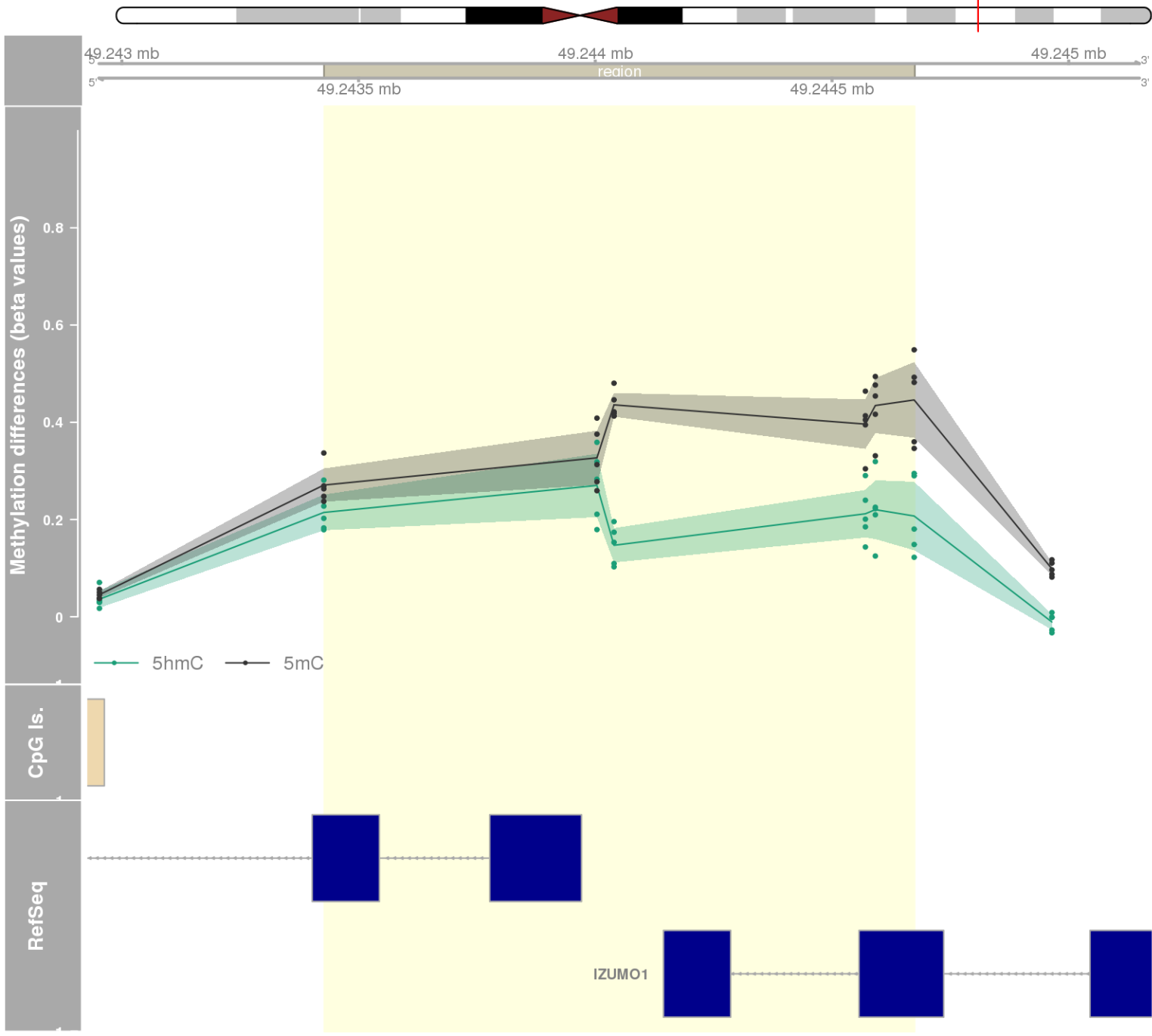
DMR 494 // chr22:20072475-20073572 // 1097 pb. (7 probes) // pvalue: 0.014 // fwer: 0.36 // pvalueArea: 0.034 // fwerArea: 1
- genes: MIR3618 / DGCR8 -



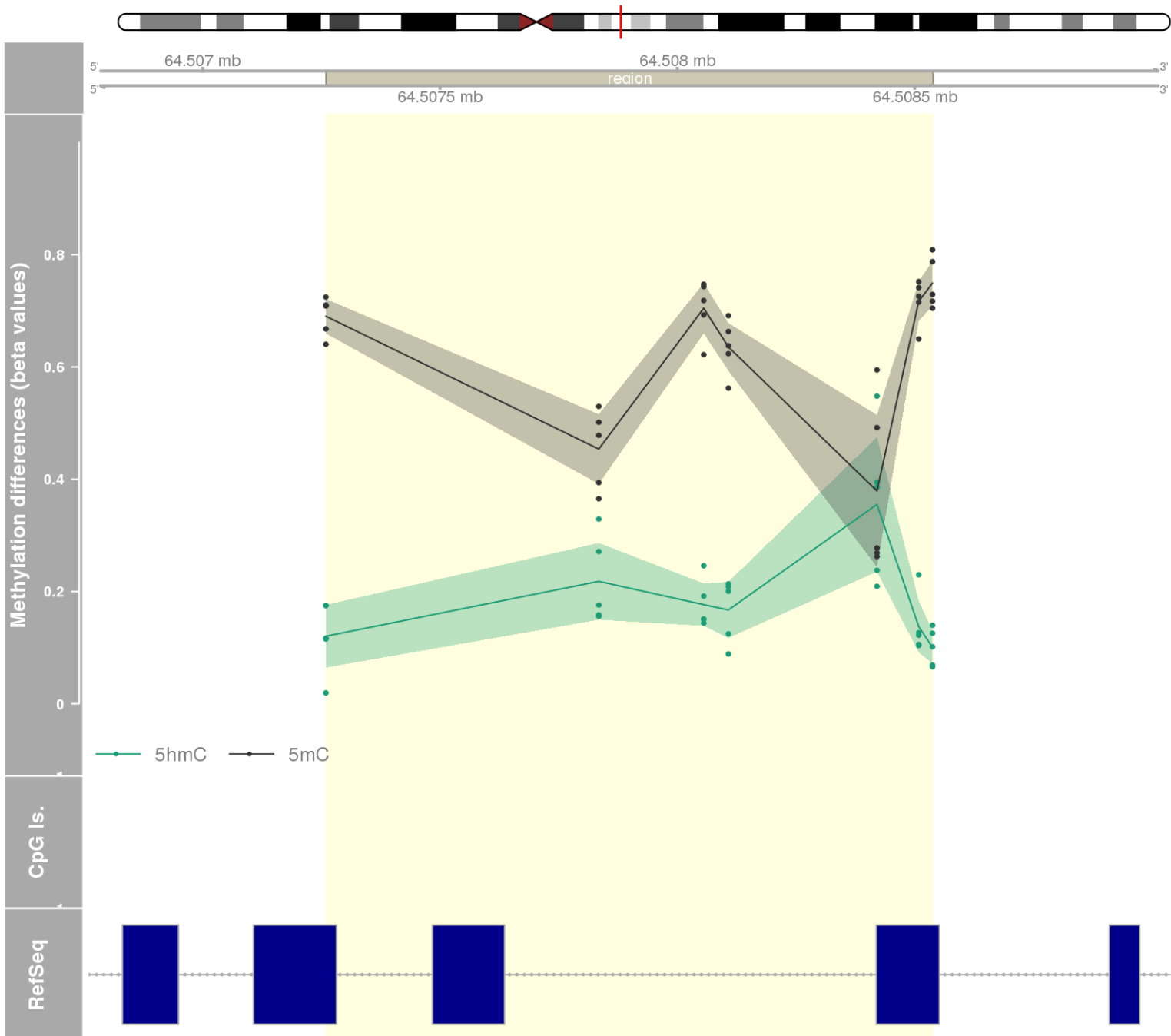
DMR 495 // chr6:160220519-160221247 // 728 pb. (8 probes) // pvalue: 0.019 // fwer: 0.522 // pvalueArea: 0.034 // fwerArea: 1



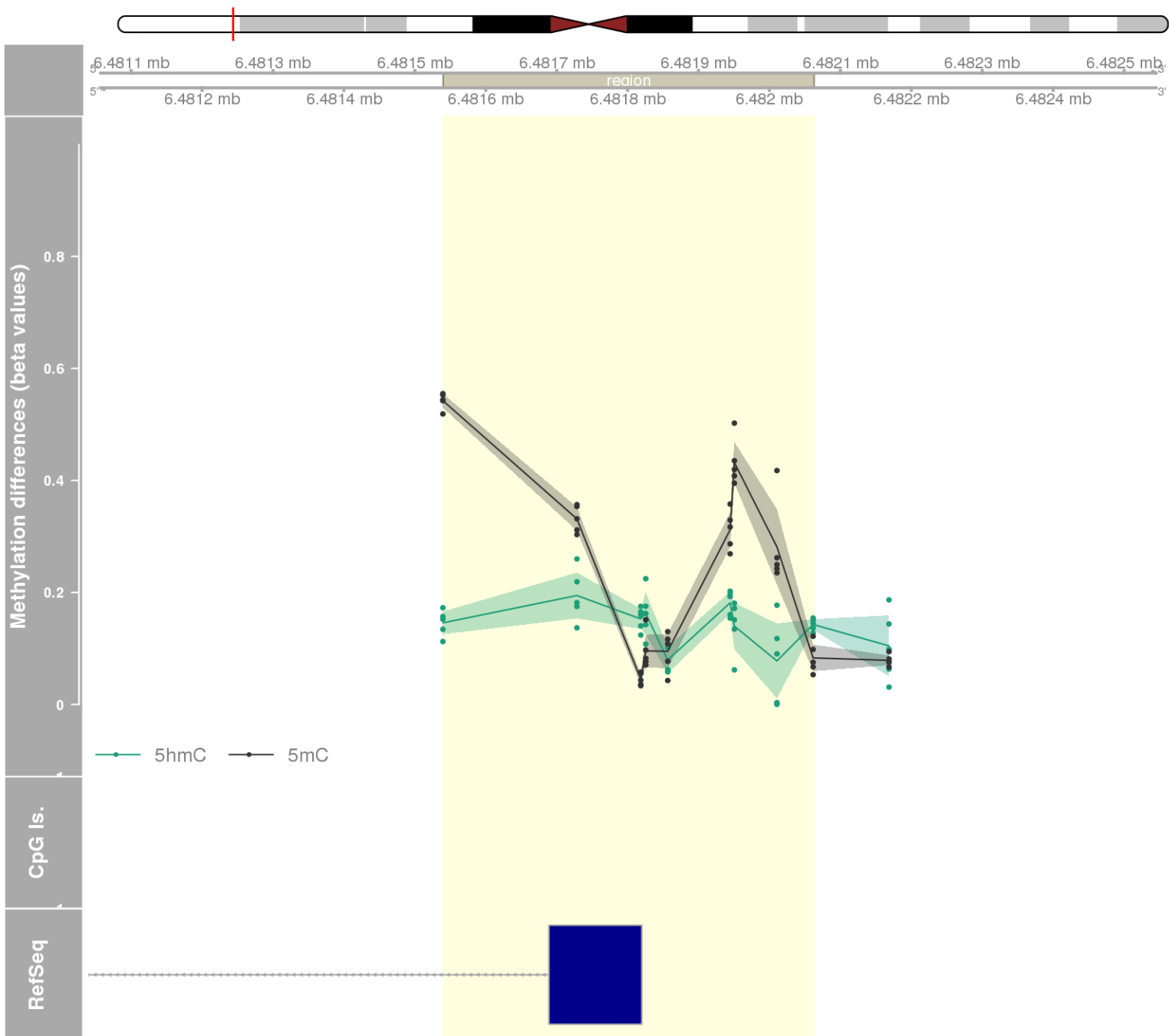
DMR 496 // chr19:49243427-49244674 // 1247 pb. (6 probes) // pvalue: 0.007 // fwer: 0.173 // pvalueArea: 0.034 // fwerArea: 1
- genes: RASIP1 / IZUMO1 -



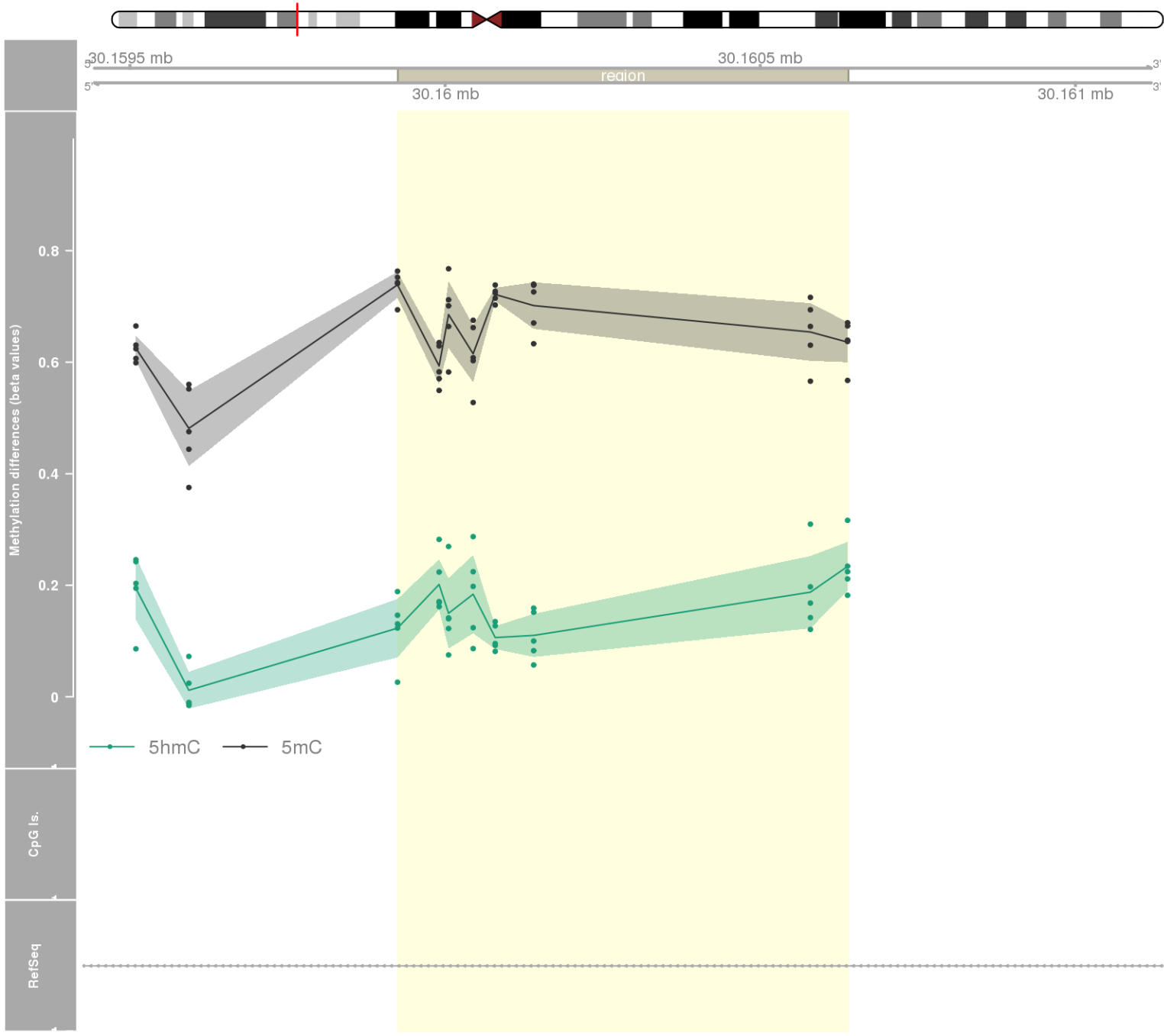
DMR 497 // chr11:64507260-64508538 // 1278 pb. (7 probes) // pvalue: 0.015 // fwer: 0.36 // pvalueArea: 0.034 // fwerArea: 1
- genes: RASGRP2 -



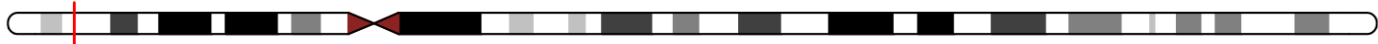
DMR 498 // chr19:6481540-6482062 // 522 pb. (9 probes) // pvalue: 0.018 // fwer: 0.36 // pvalueArea: 0.034 // fwerArea: 1
- genes: DENND1C -



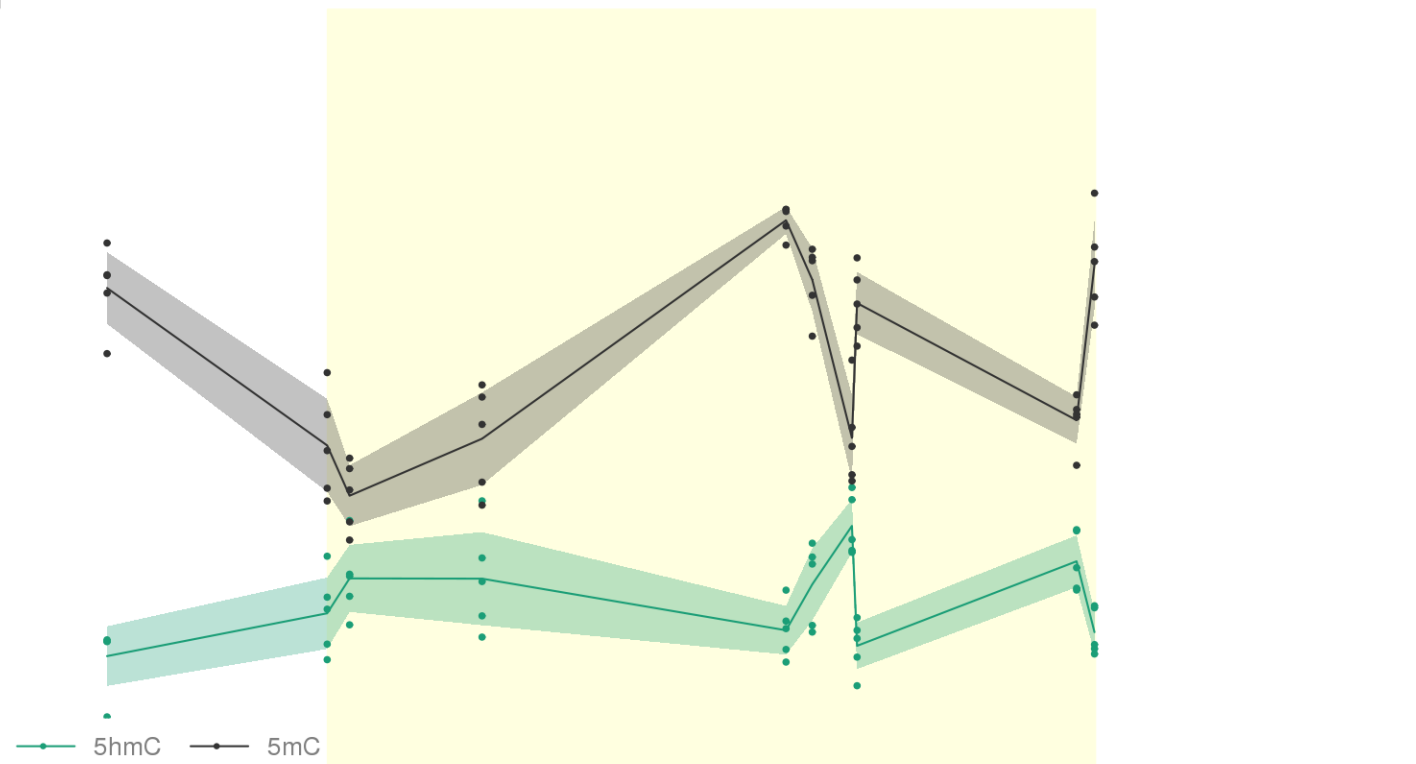
DMR 499 // chr6:30159925-30160639 // 714 pb. (8 probes) // pvalue: 0.02 // fwer: 0.522 // pvalueArea: 0.034 // fwerArea: 1
- genes: TRIM26 -



DMR 500 // chr12:6485881-6487080 // 1199 pb. (9 probes) // pvalue: 0.019 // fwer: 0.36 // pvalueArea: 0.035 // fwerArea: 1
- genes: SCNN1A / LTBR -



Methylation differences (beta values)



CpG Is.

RefSeq

