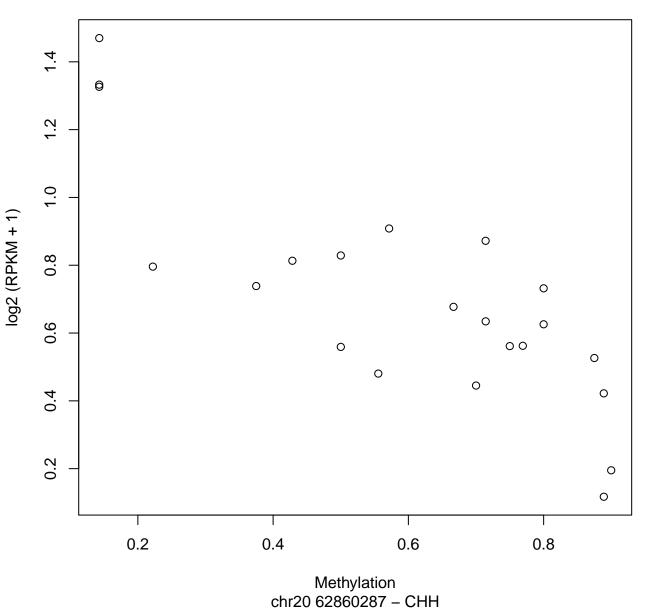
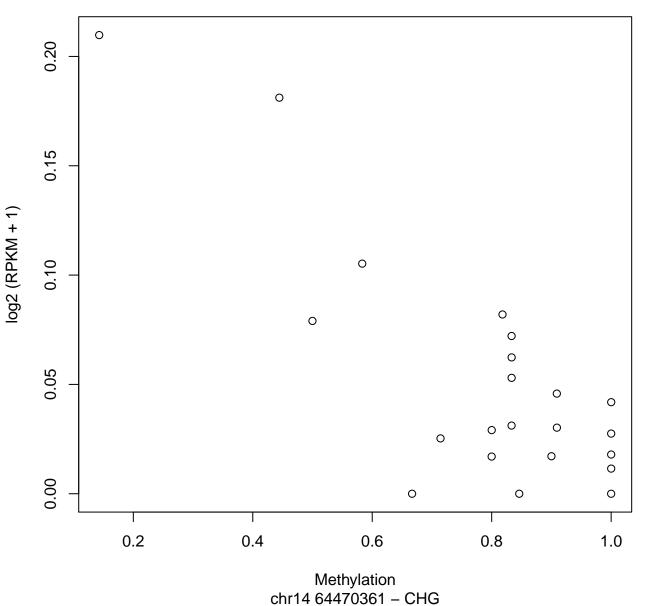
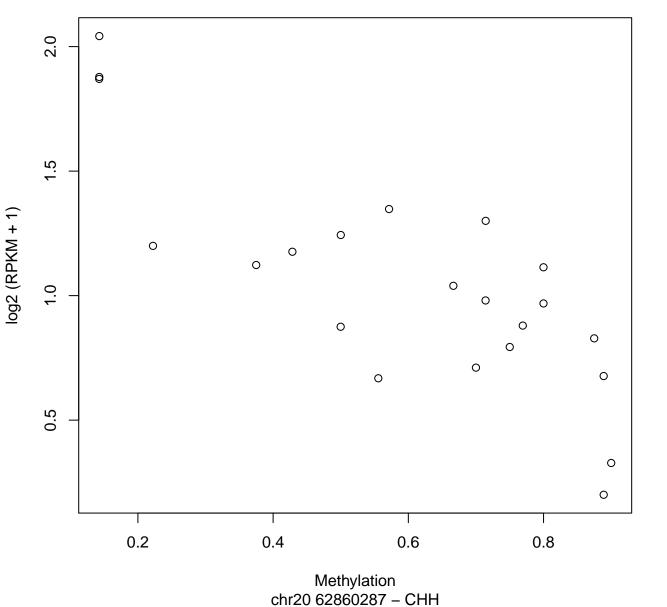
exon ENSE00003742259.1_1 FDR 0.00497 MYT1



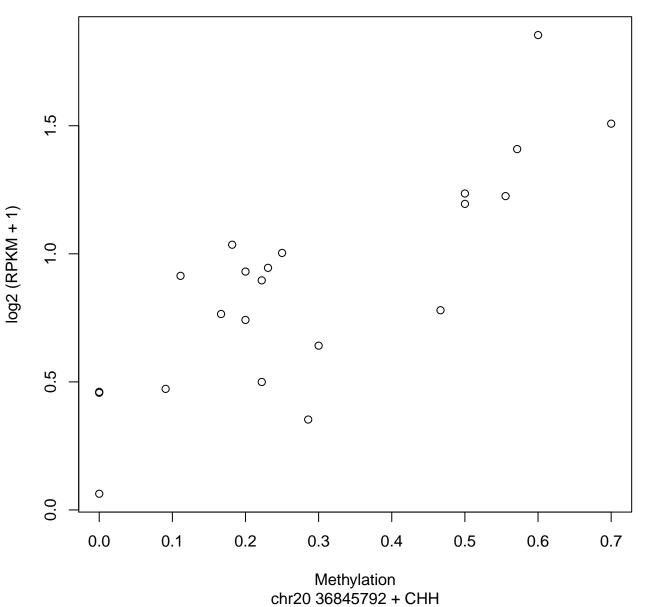
exon ENSE00003752734.1_1 FDR 0.00814 SYNE2



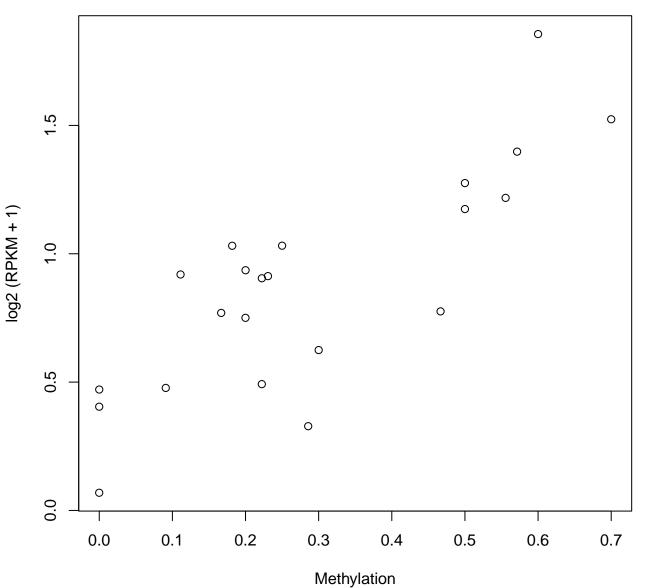
exon ENSE00001171090.1_1 FDR 0.0083 MYT1



exon ENSE00001948043.1_1 FDR 0.0094 KIAA1755

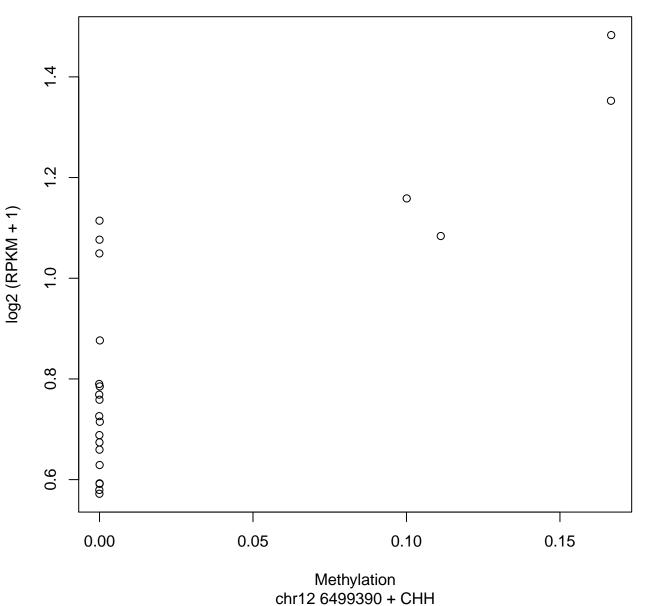


exon ENSE00001950686.1_1 FDR 0.0103 KIAA1755

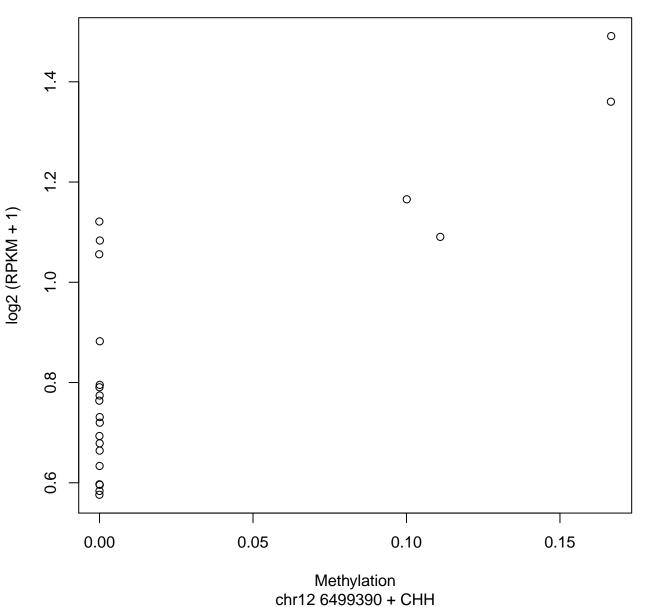


chr20 36845792 + CHH

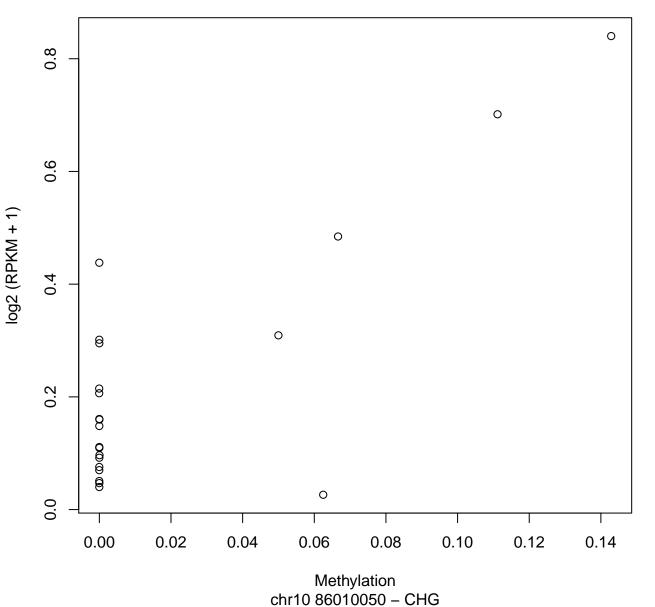
exon ENSE00000866904.6_1 FDR 0.0126 LTBR



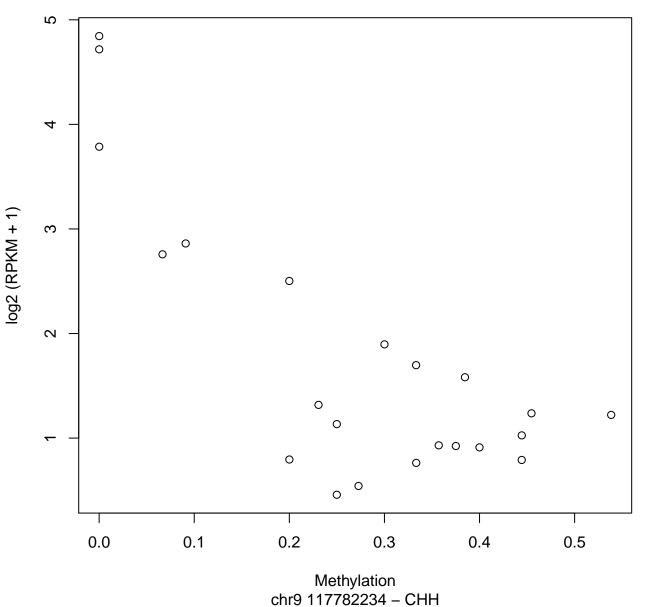
exon ENSE00002212537.1_1 FDR 0.0127 LTBR



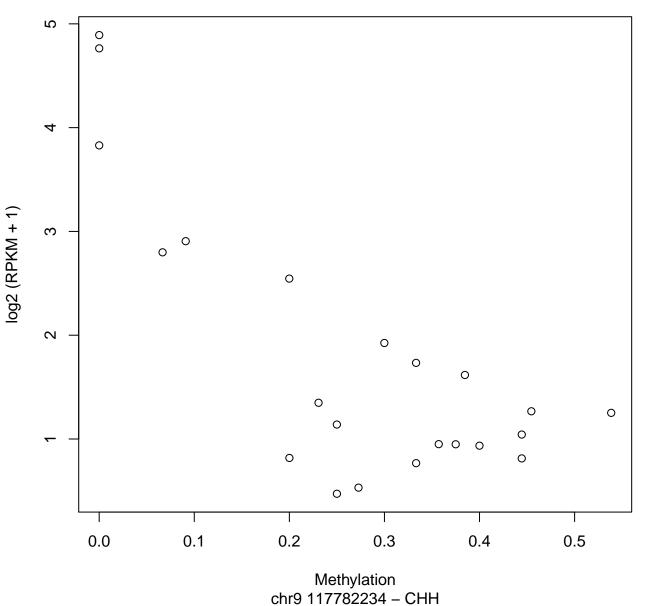
exon ENSE00001456895.3_1 FDR 0.0144 RGR



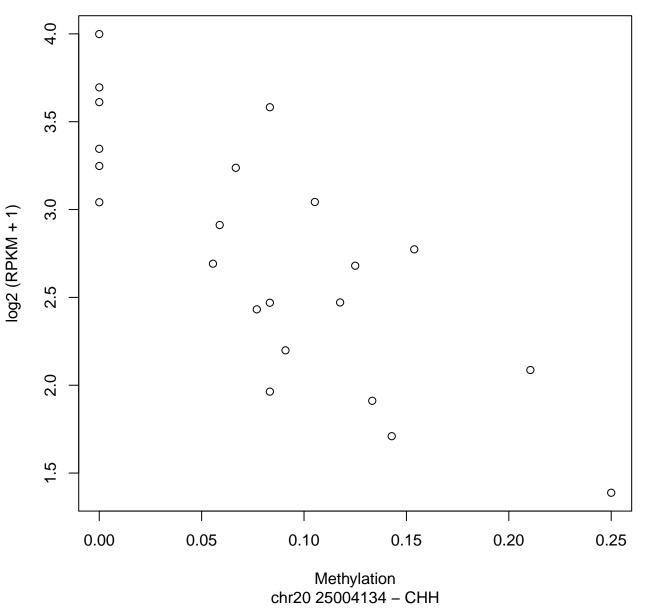
exon ENSE00001948855.1_1 FDR 0.0192 TNC



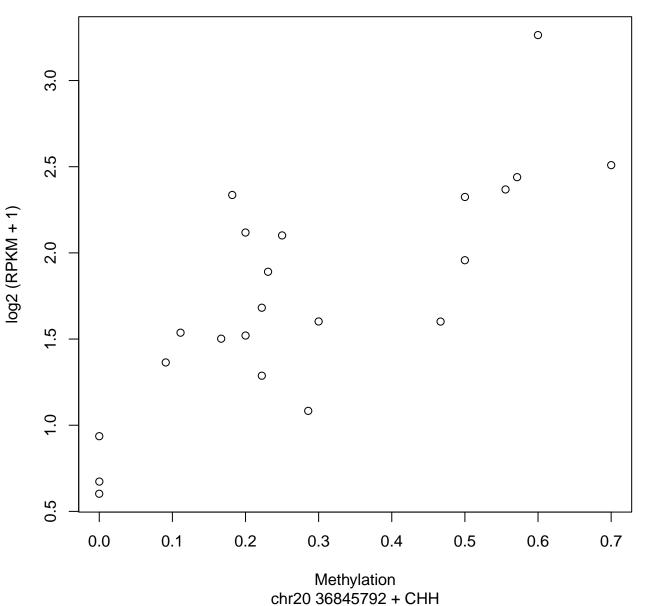
exon ENSE00002283977.1_1 FDR 0.0201 TNC



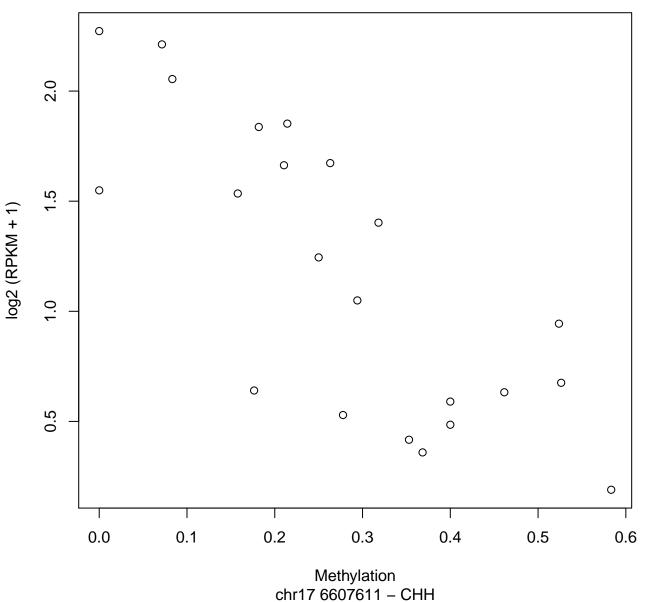
exon ENSE00001203636.1_1 FDR 0.021 ACSS1



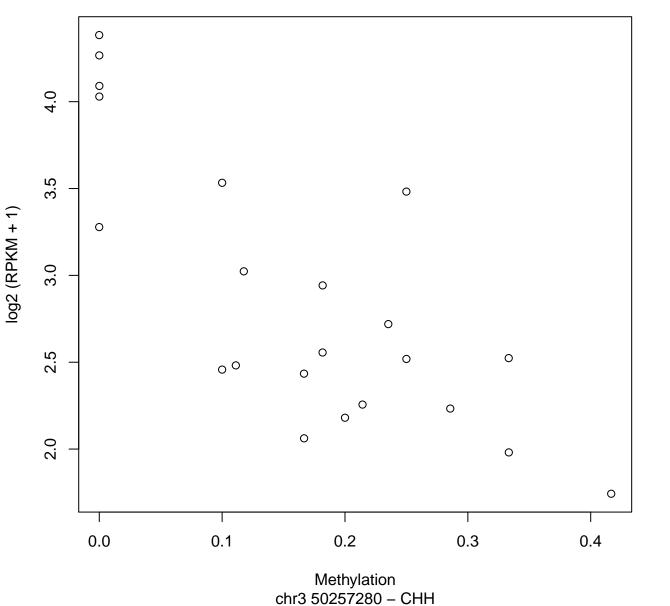
exon ENSE00003472816.1_1 FDR 0.0213 KIAA1755



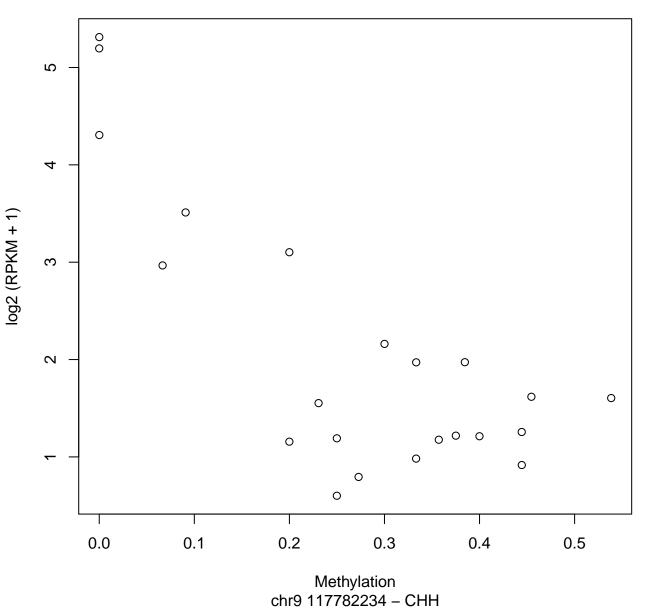
exon ENSE00002643163.1_1 FDR 0.0229 SLC13A5



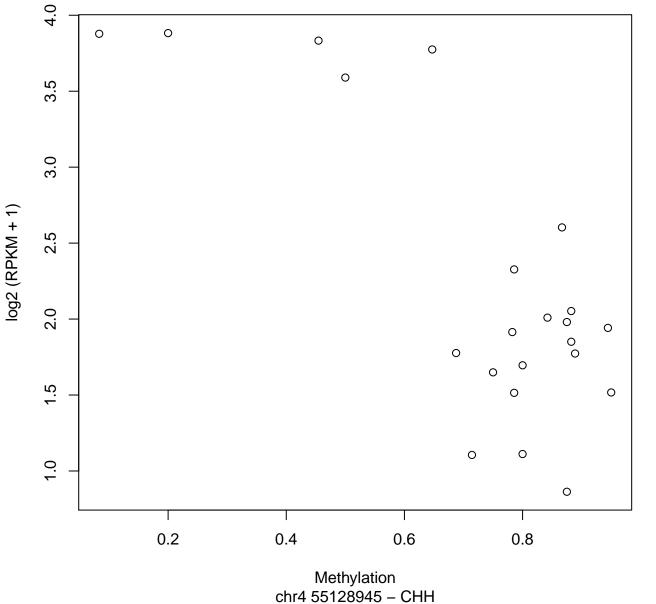
exon ENSE00003738580.1_1 FDR 0.0247 SLC38A3



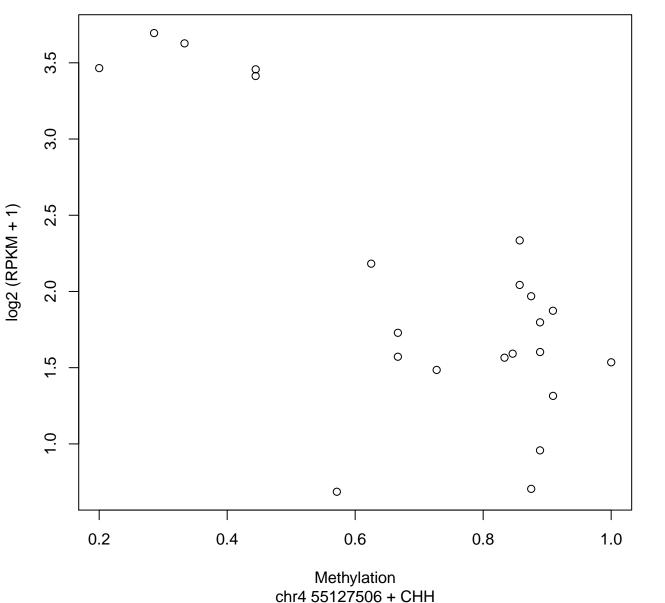
exon ENSE00002278148.1_1 FDR 0.0259 TNC



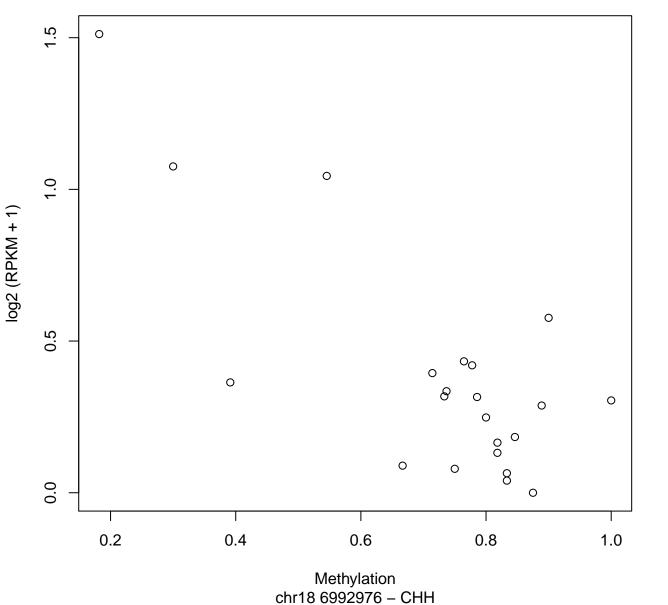
exon ENSE00002050497.1_1 FDR 0.0318 PDGFRA



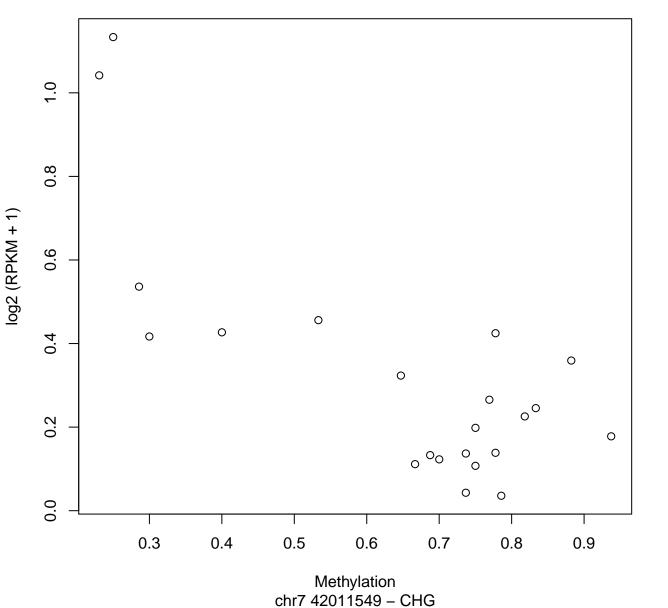
exon ENSE00003785223.1_1 FDR 0.0332 PDGFRA



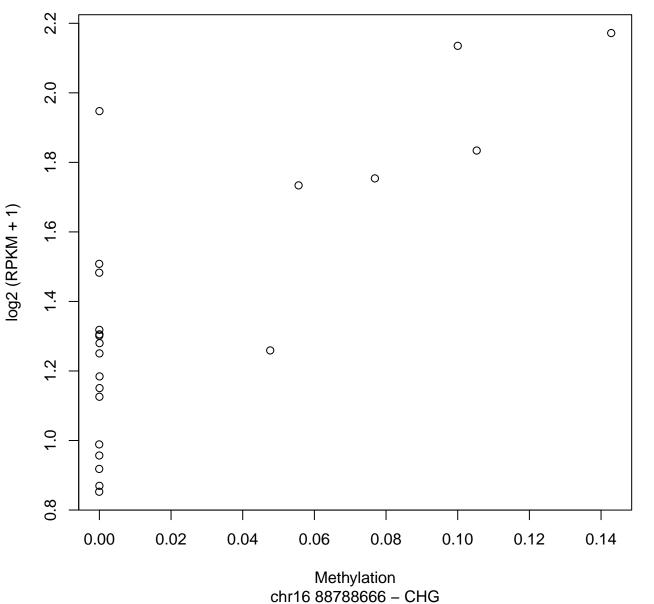
exon ENSE00003666767.1_1 FDR 0.0341 LAMA1



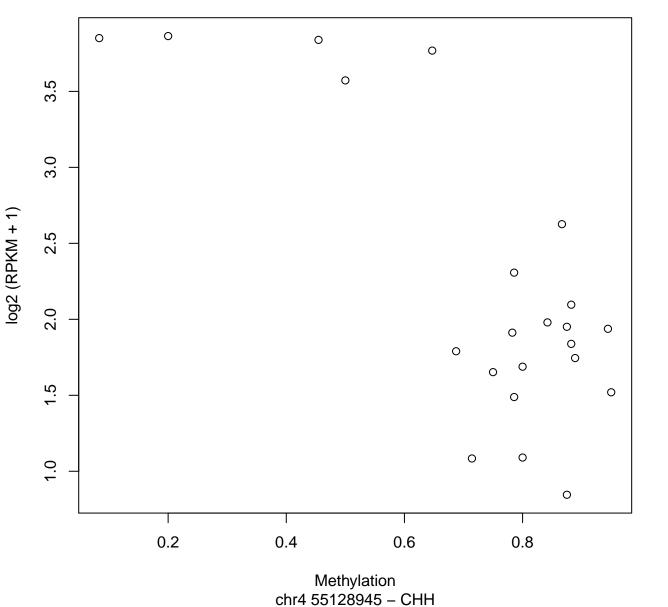
exon ENSE00003537008.1_1 FDR 0.0347 GLI3



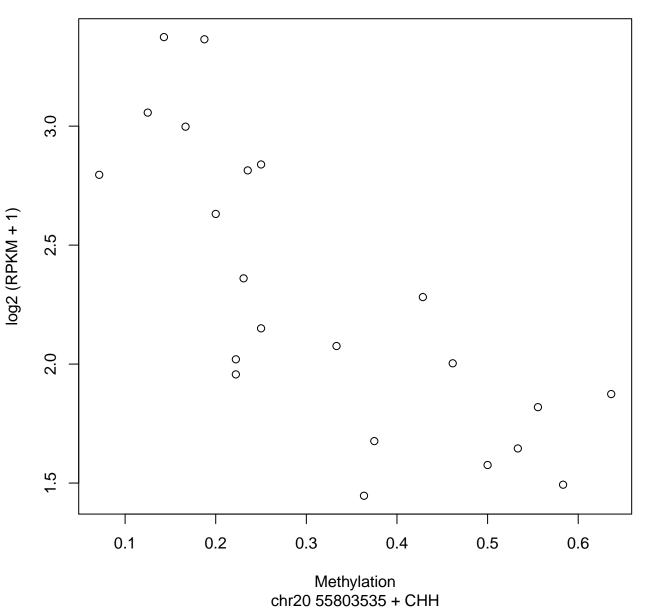
exon ENSE00001262469.1_1 FDR 0.0355 PIEZO1



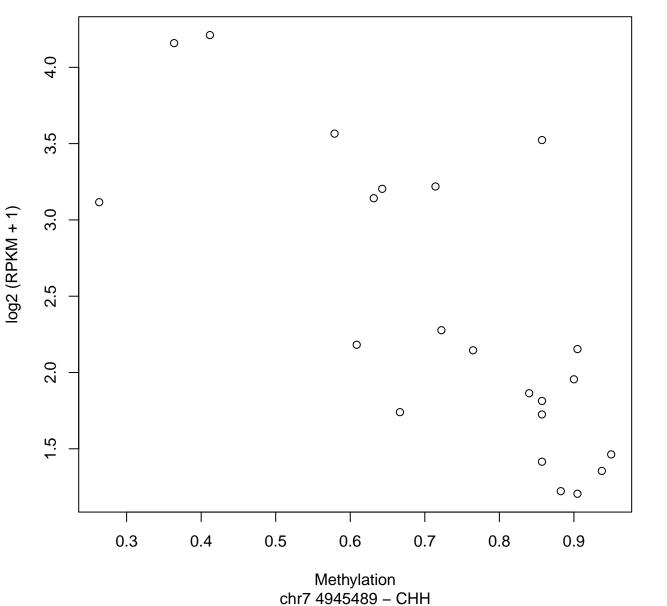
exon ENSE00002062670.1_1 FDR 0.0356 PDGFRA



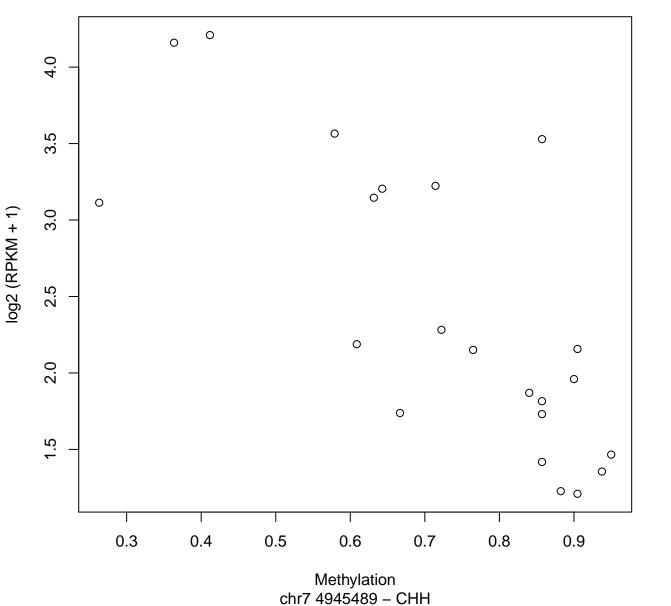
exon ENSE00003463644.1_1 FDR 0.0362 BMP7



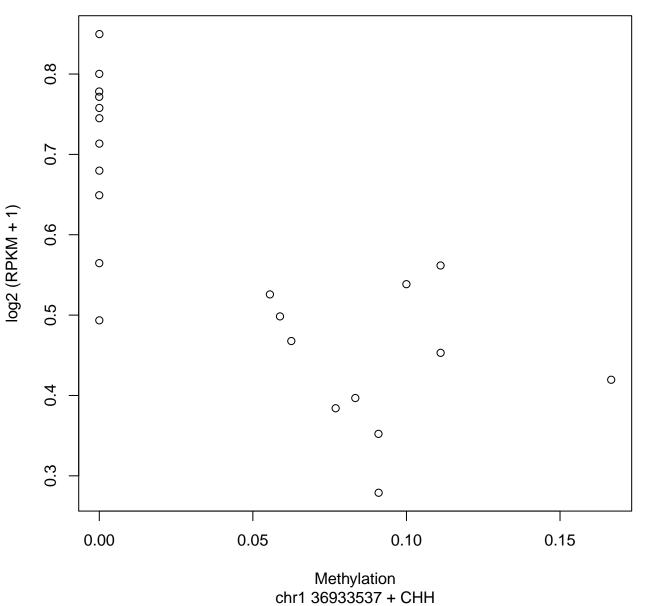
exon ENSE00003688298.1_1 FDR 0.0428 MMD2



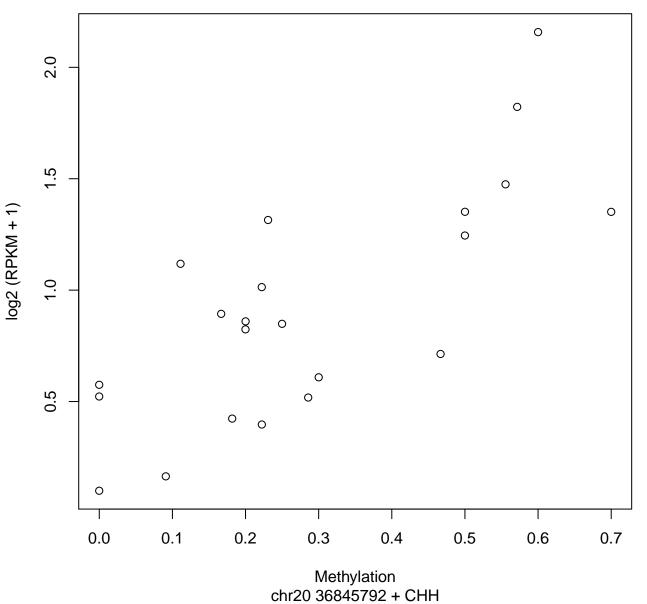
exon ENSE00001831476.1_1 FDR 0.0437 MMD2



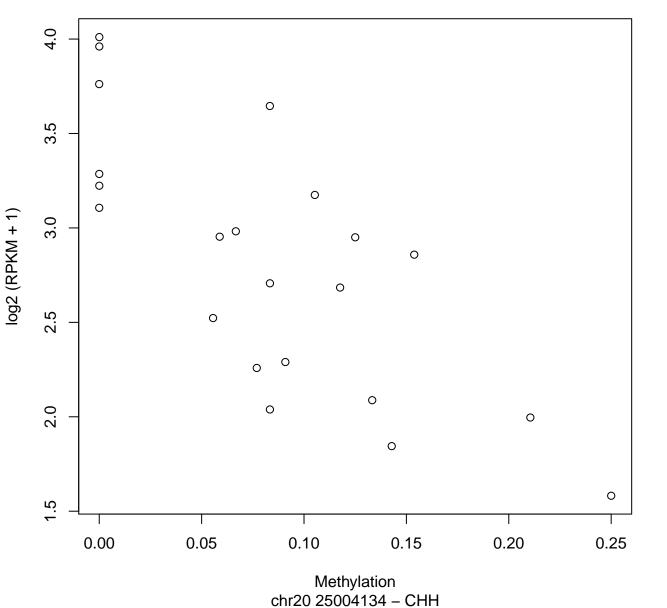
exon ENSE00001913571.2_1 FDR 0.0439 CSF3R



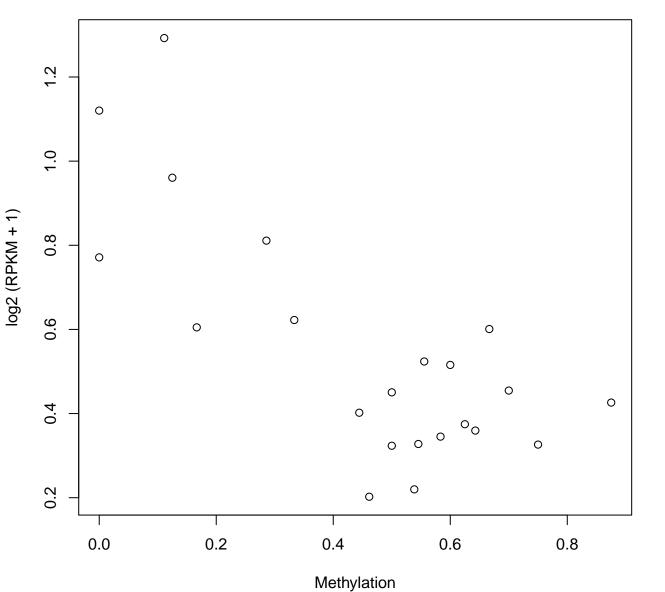
exon ENSE00001921792.1_1 FDR 0.045 KIAA1755



exon ENSE00001299793.1_1 FDR 0.0469 ACSS1



exon ENSE00002326770.1_1 FDR 0.0498 PARP14



chr3 122420609 + CHH

exon ENSE00000880104.1_1 FDR 0.0498 TST

