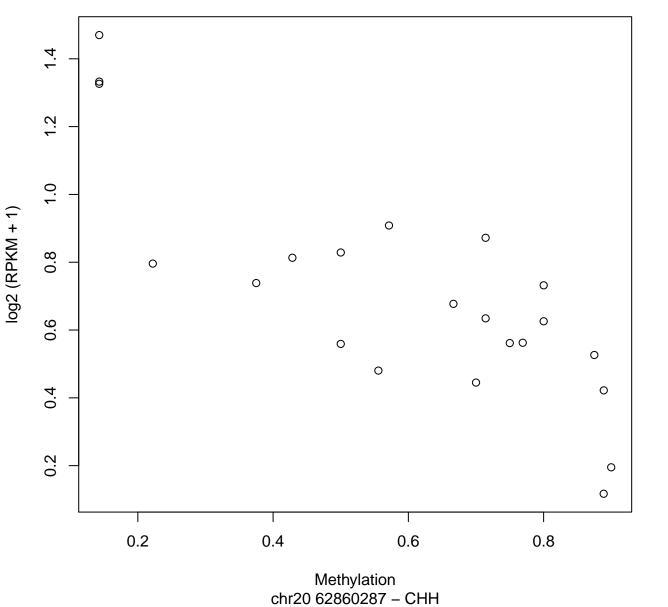
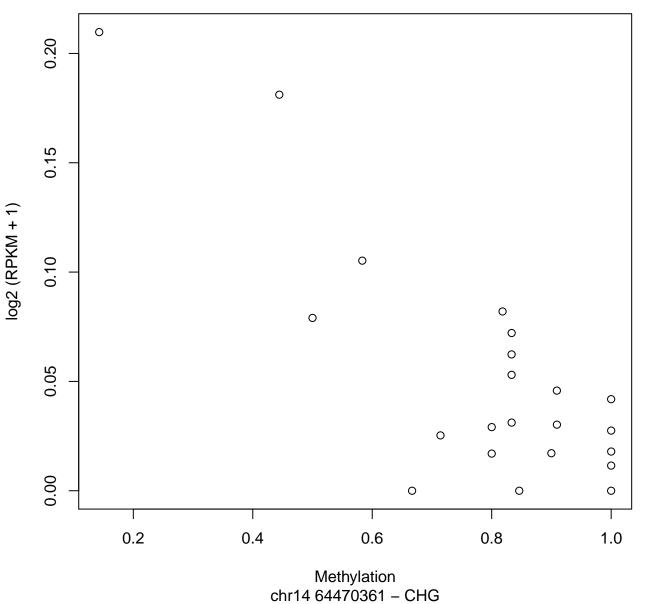
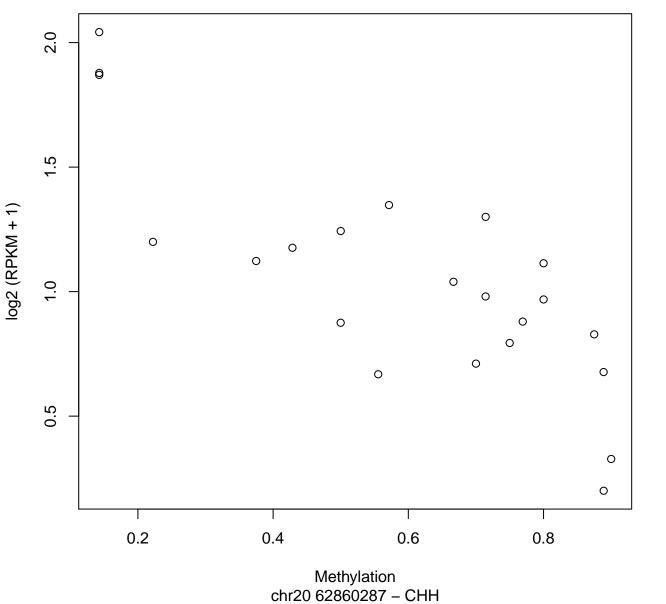
exon e1092329 FDR 0.00497 MYT1



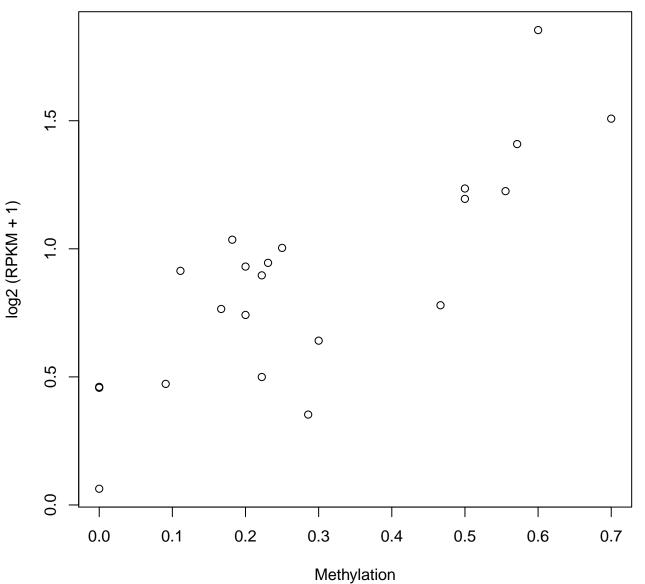
exon e778085 FDR 0.00814 SYNE2



exon e1092347 FDR 0.0083 MYT1

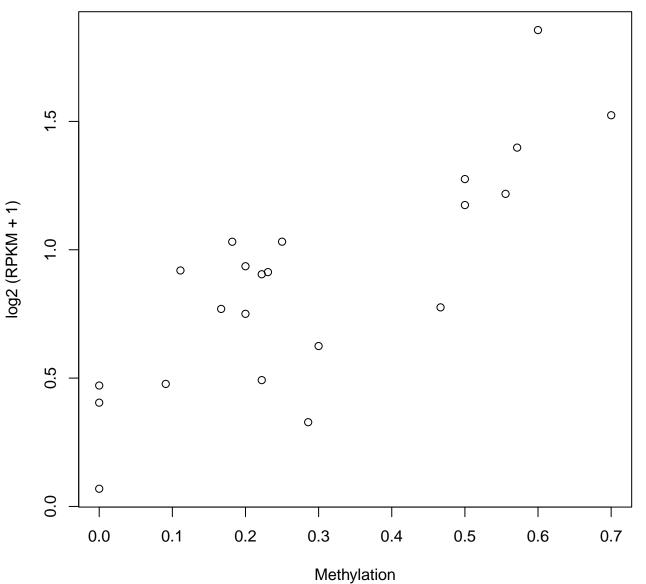


exon e1080751 FDR 0.0094 KIAA1755



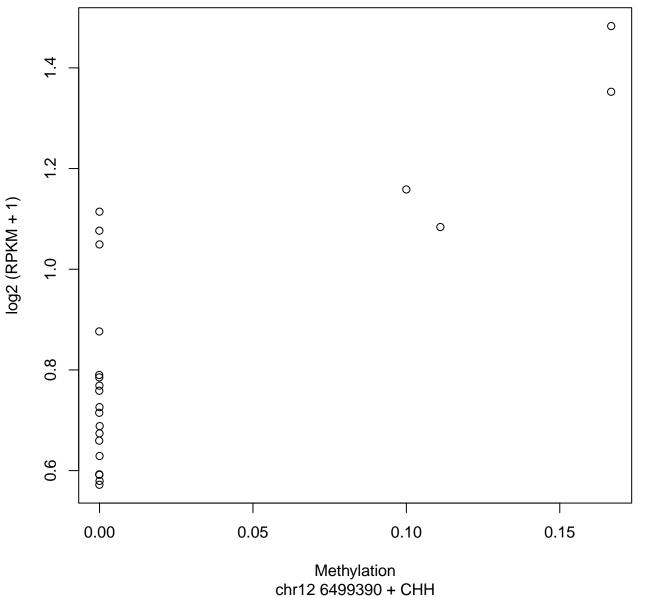
chr20 36845792 + CHH

exon e1080759 FDR 0.0103 KIAA1755

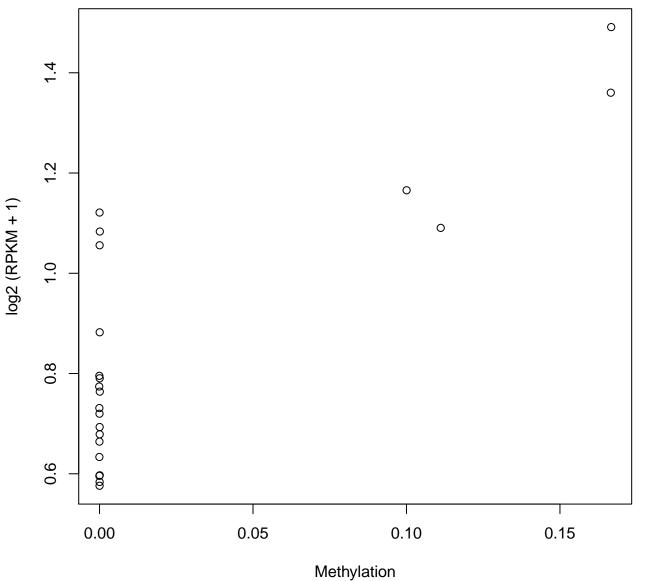


chr20 36845792 + CHH

exon e675117 FDR 0.0126 LTBR

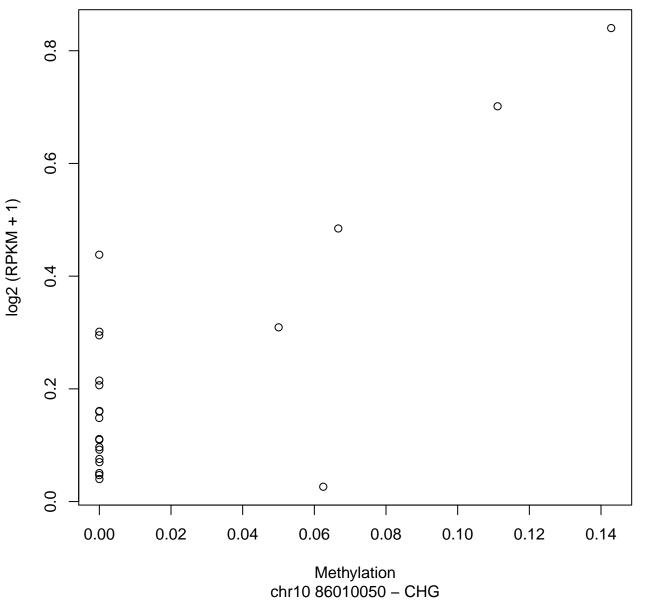


exon e675072 FDR 0.0127 LTBR

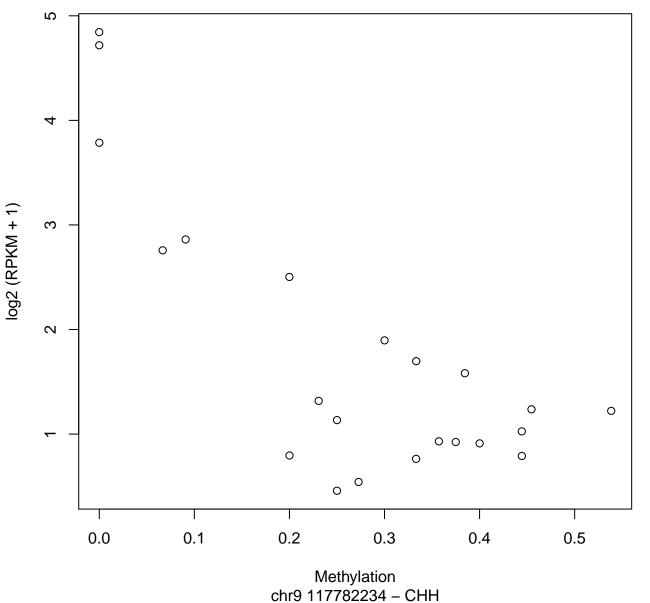


chr12 6499390 + CHH

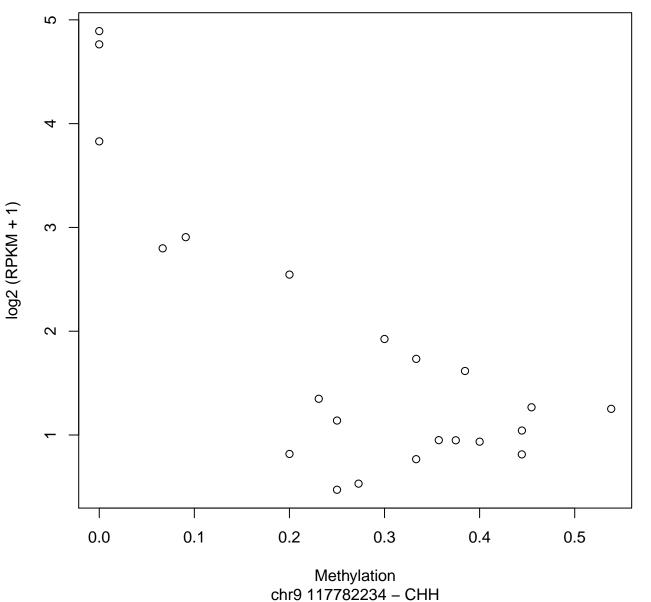
exon e583287 FDR 0.0144 RGR



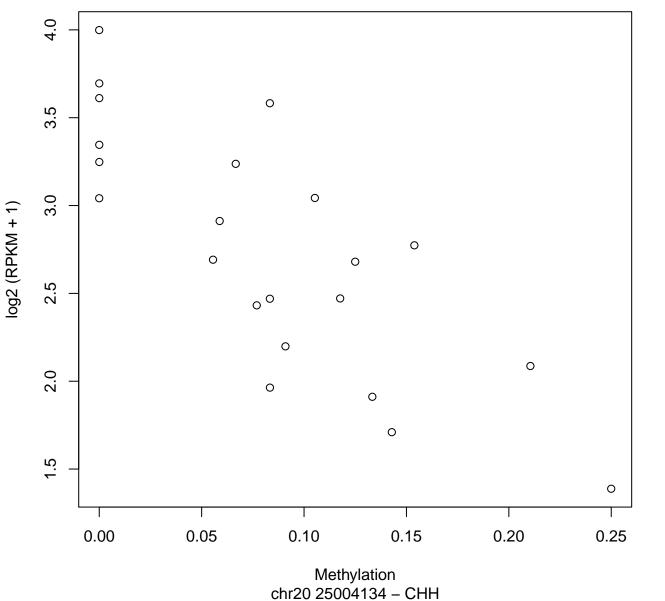
exon e542983 FDR 0.0192 TNC



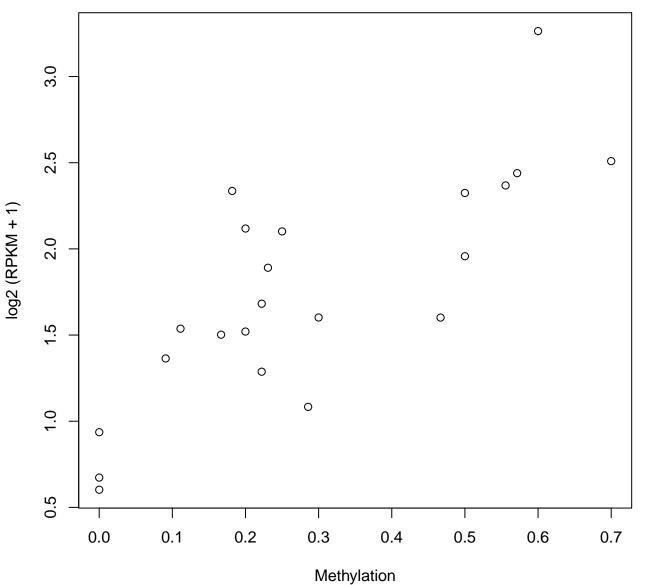
exon e543008 FDR 0.0201 TNC



exon e1074247 FDR 0.021 ACSS1

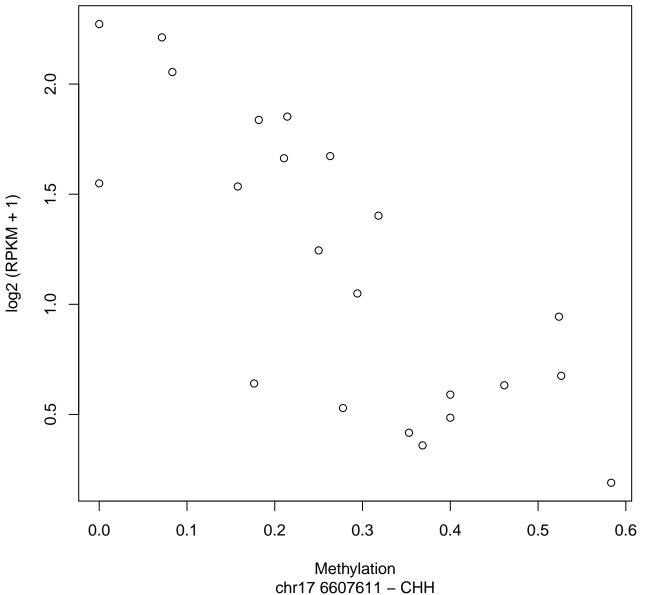


exon e1080748 FDR 0.0213 KIAA1755

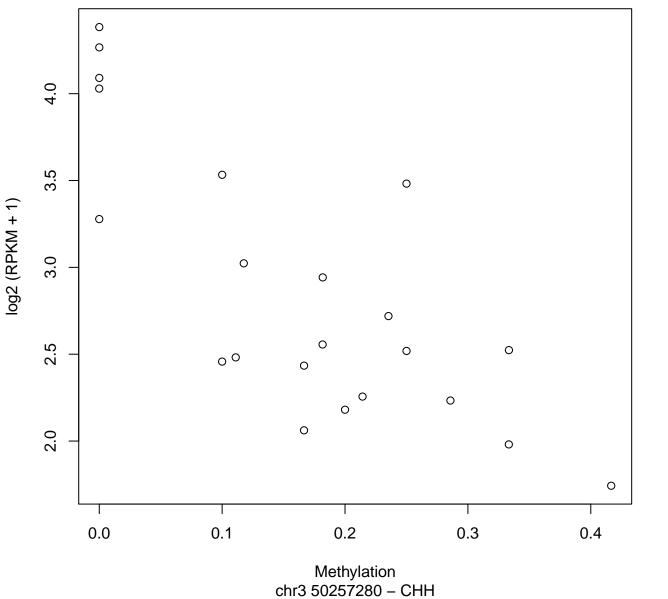


chr20 36845792 + CHH

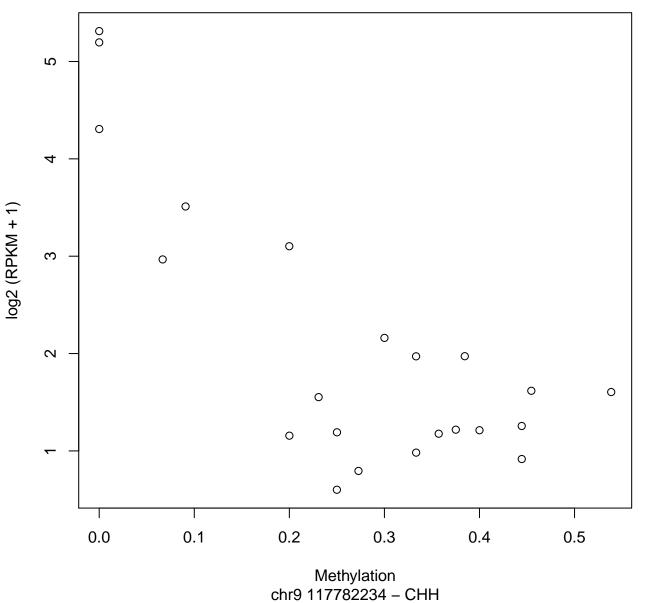
exon e908676 FDR 0.0229 SLC13A5



exon e220142 FDR 0.0247 SLC38A3



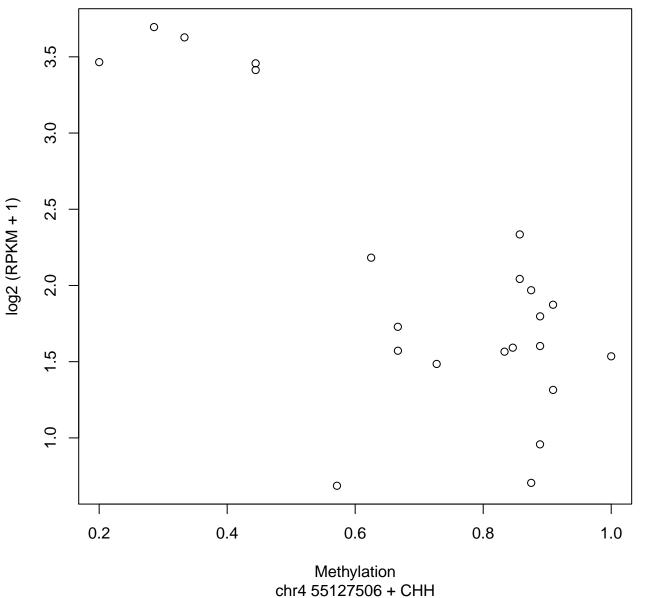
exon e543033 FDR 0.0259 TNC



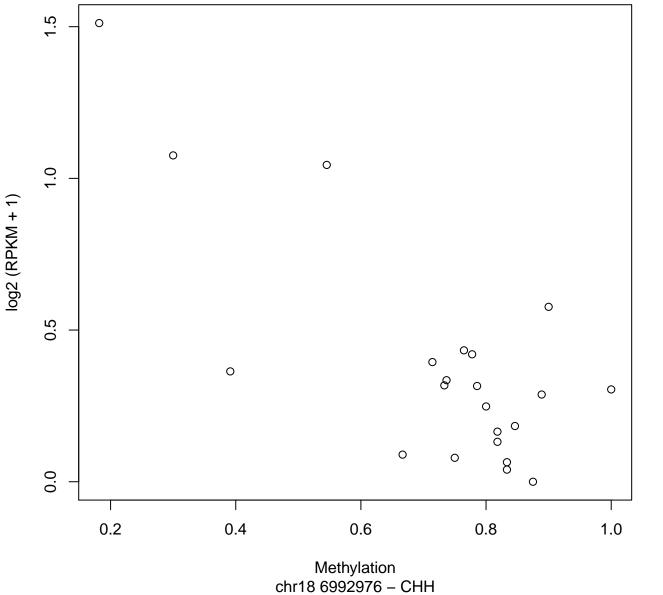
exon e284534 FDR 0.0318 **PDGFRA** 4.0 0 0 0 0 0 3.5 3.0 log2 (RPKM + 1) 0 2.5 0 2.0 0 0 0 0 00 0 0 1.5 0 0 0 0 1.0 0 0.2 0.4 0.6 8.0

Methylation chr4 55128945 – CHH

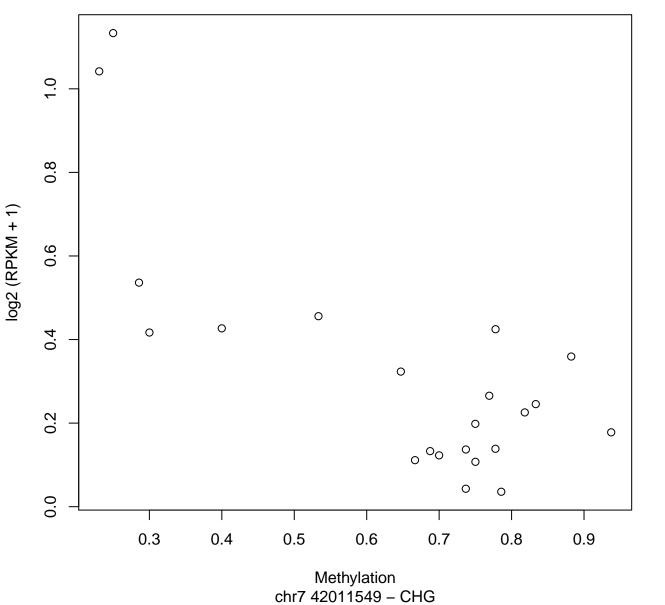
exon e284465 FDR 0.0332 PDGFRA



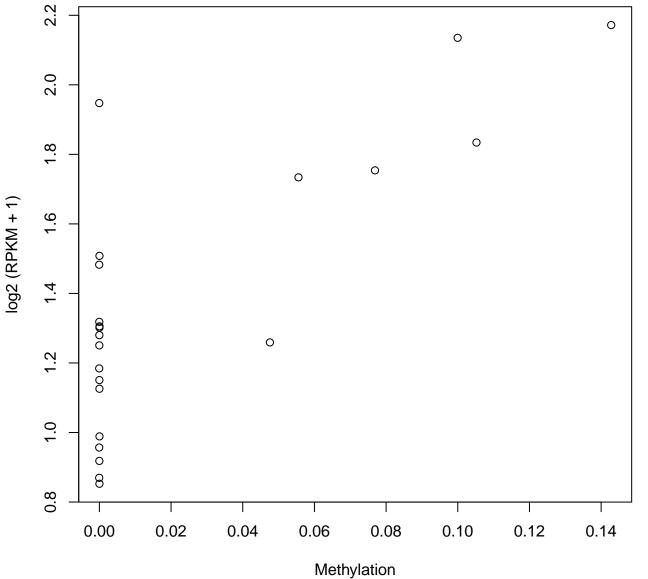
exon e978496 FDR 0.0341 LAMA1



exon e432062 FDR 0.0347 GLI3

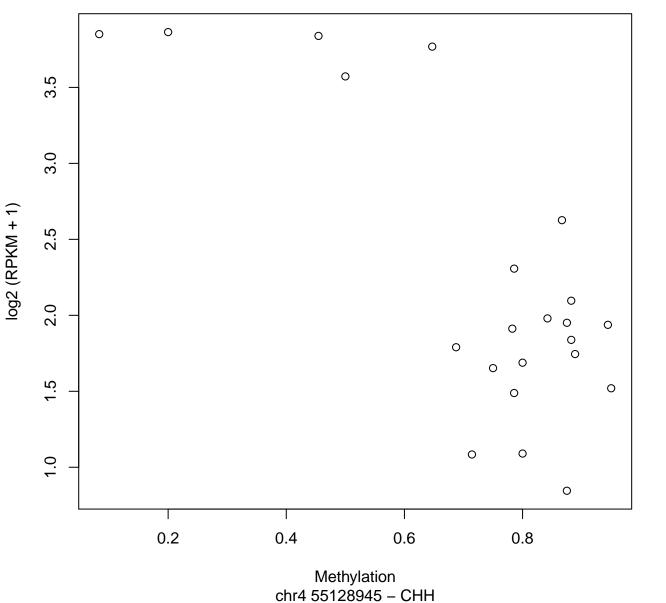


exon e898855 FDR 0.0355 PIEZO1

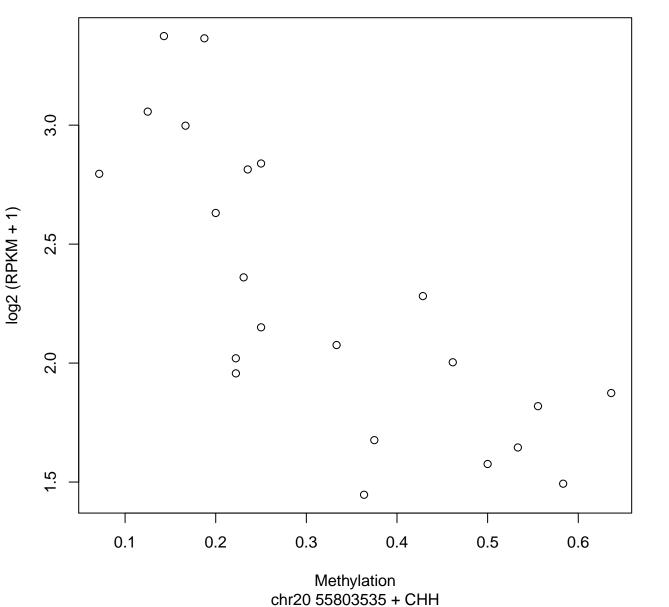


chr16 88788666 - CHG

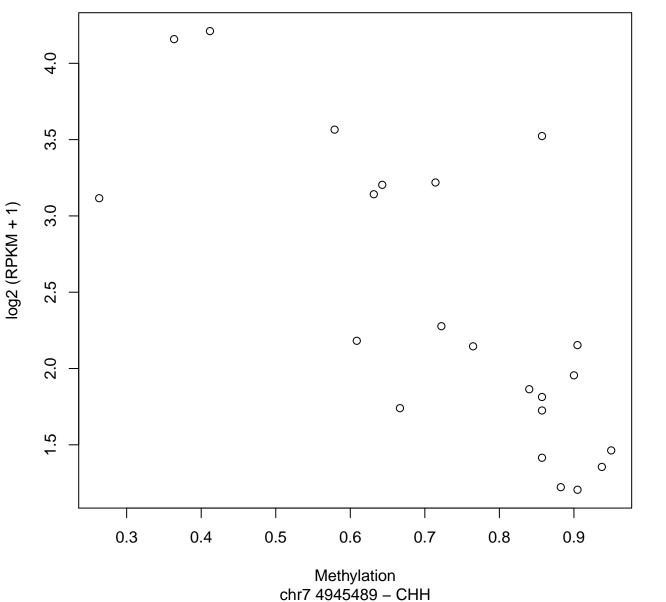
exon e284530 FDR 0.0356 PDGFRA



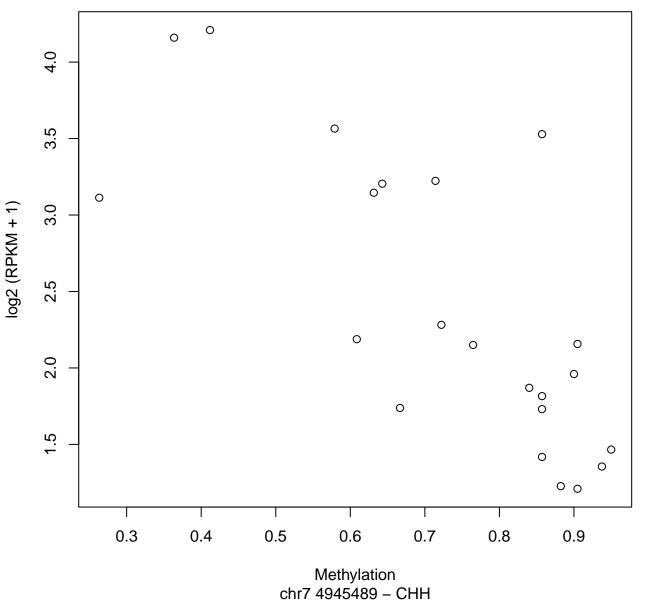
exon e1087421 FDR 0.0362 BMP7



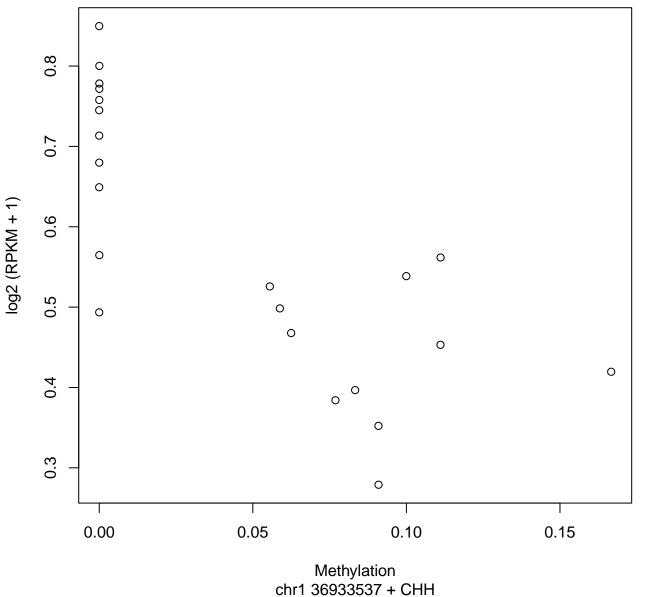
exon e421583 FDR 0.0428 MMD2



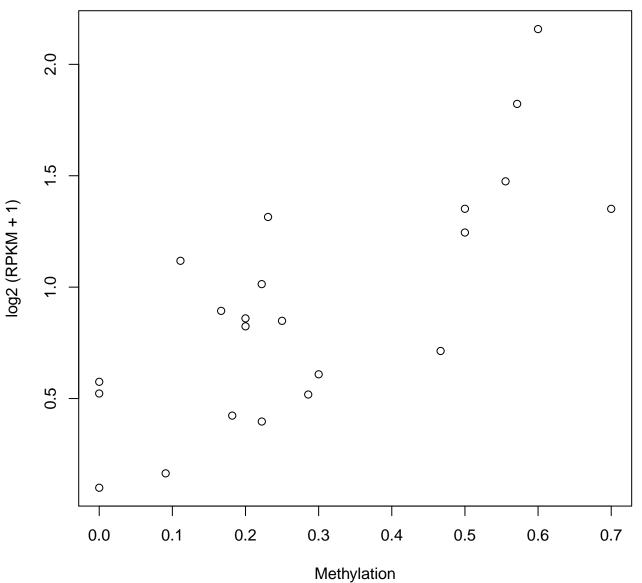
exon e421597 FDR 0.0437 MMD2



exon e26173 FDR 0.0439 CSF3R

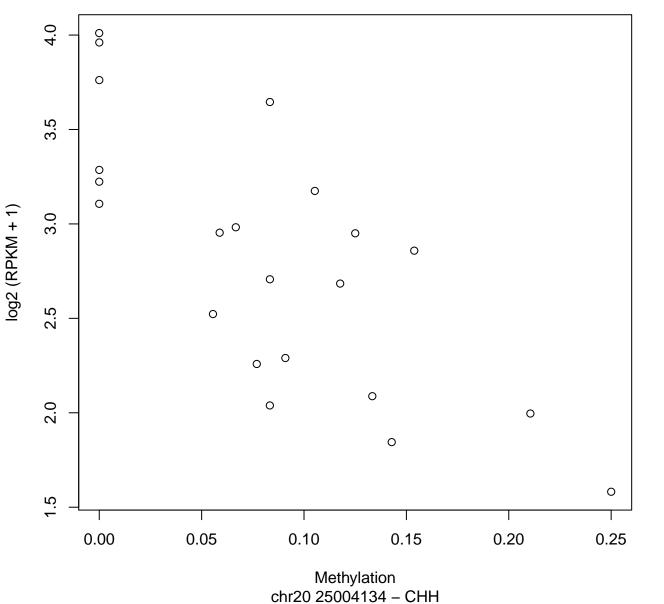


exon e1080752 FDR 0.045 KIAA1755

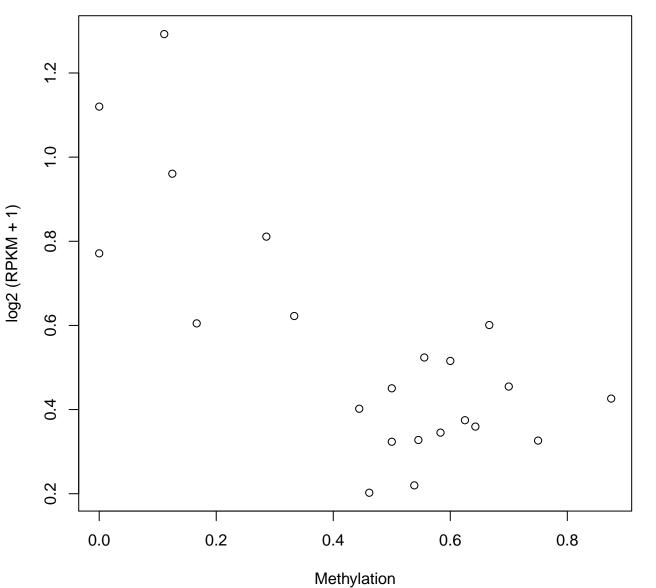


chr20 36845792 + CHH

exon e1074246 FDR 0.0469 ACSS1

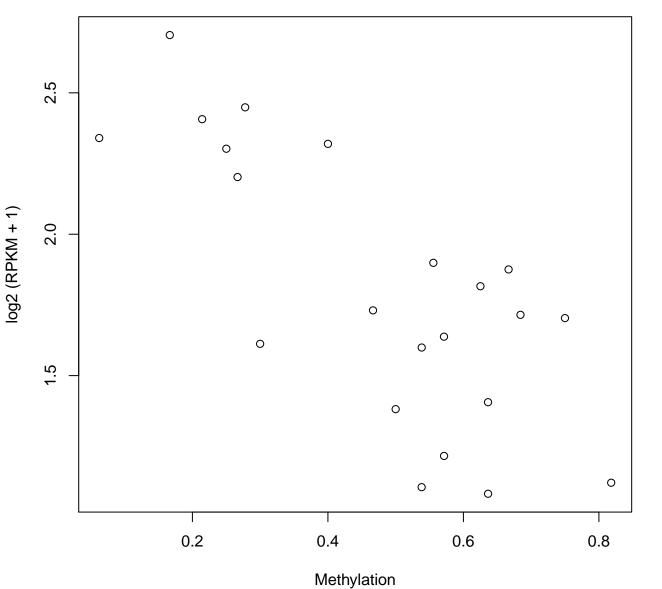


exon e239711 FDR 0.0498 PARP14



chr3 122420609 + CHH

exon e1122004 FDR 0.0498 TST



chr22 37414408 - CHH