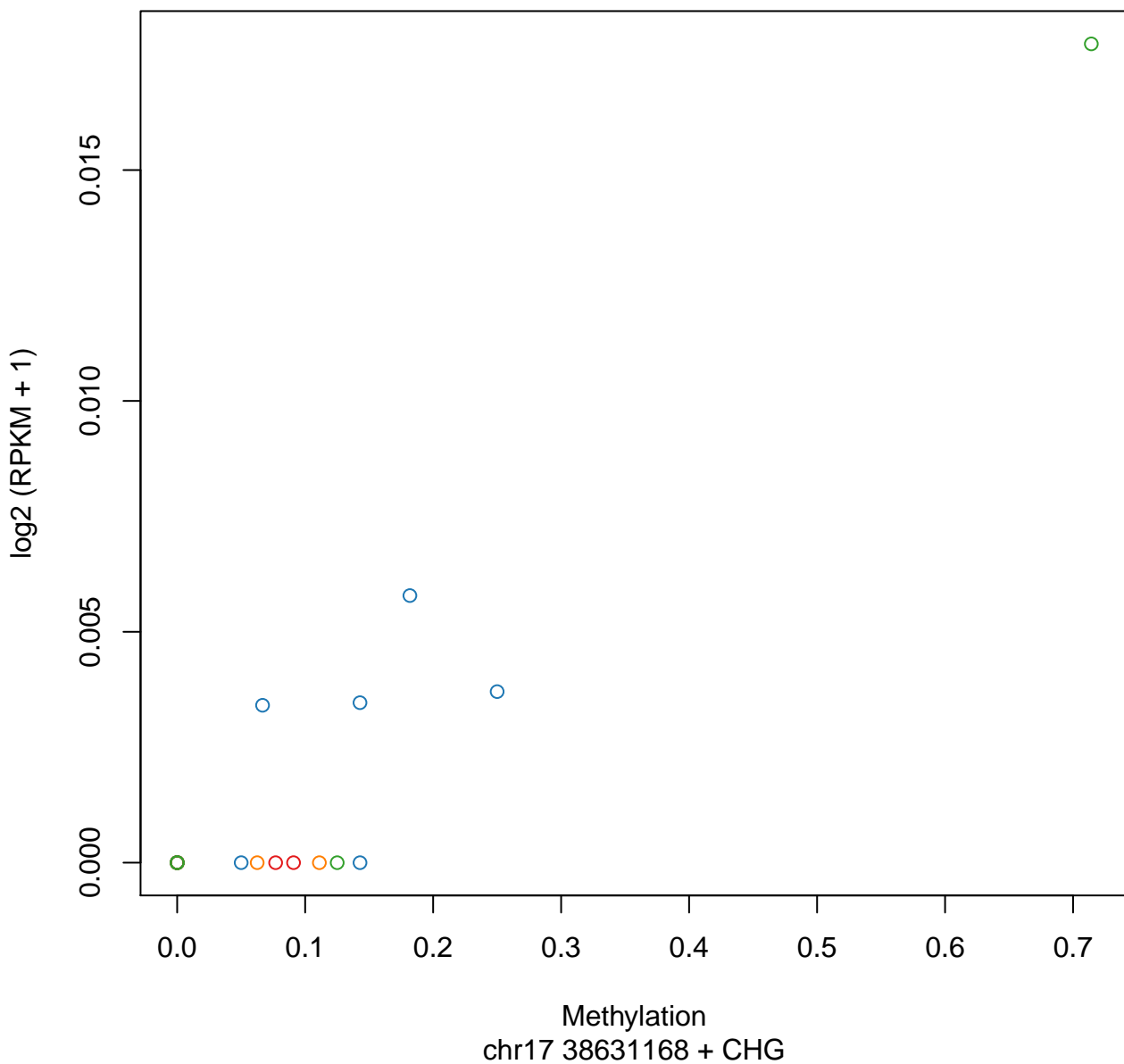


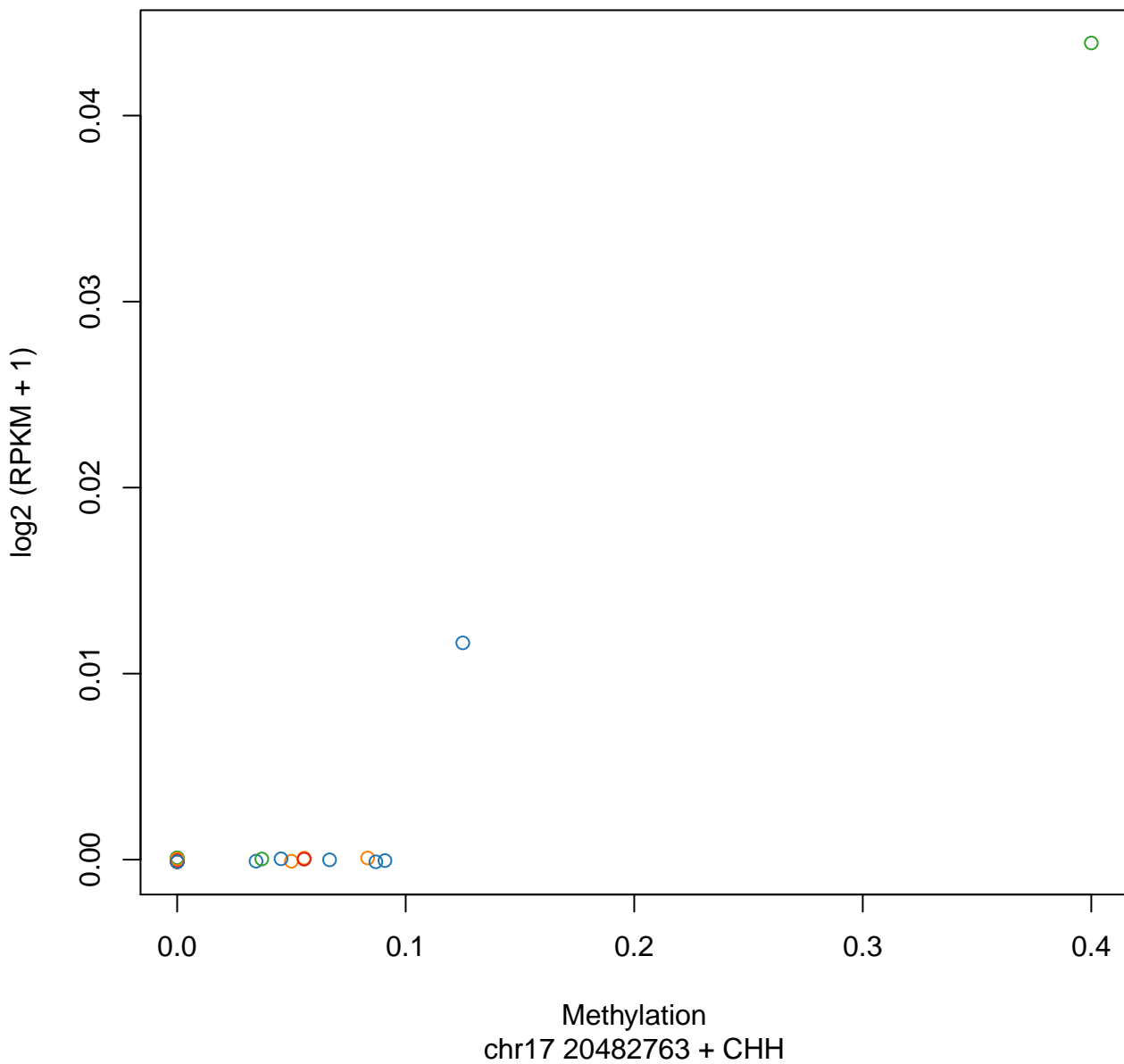
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TNS4



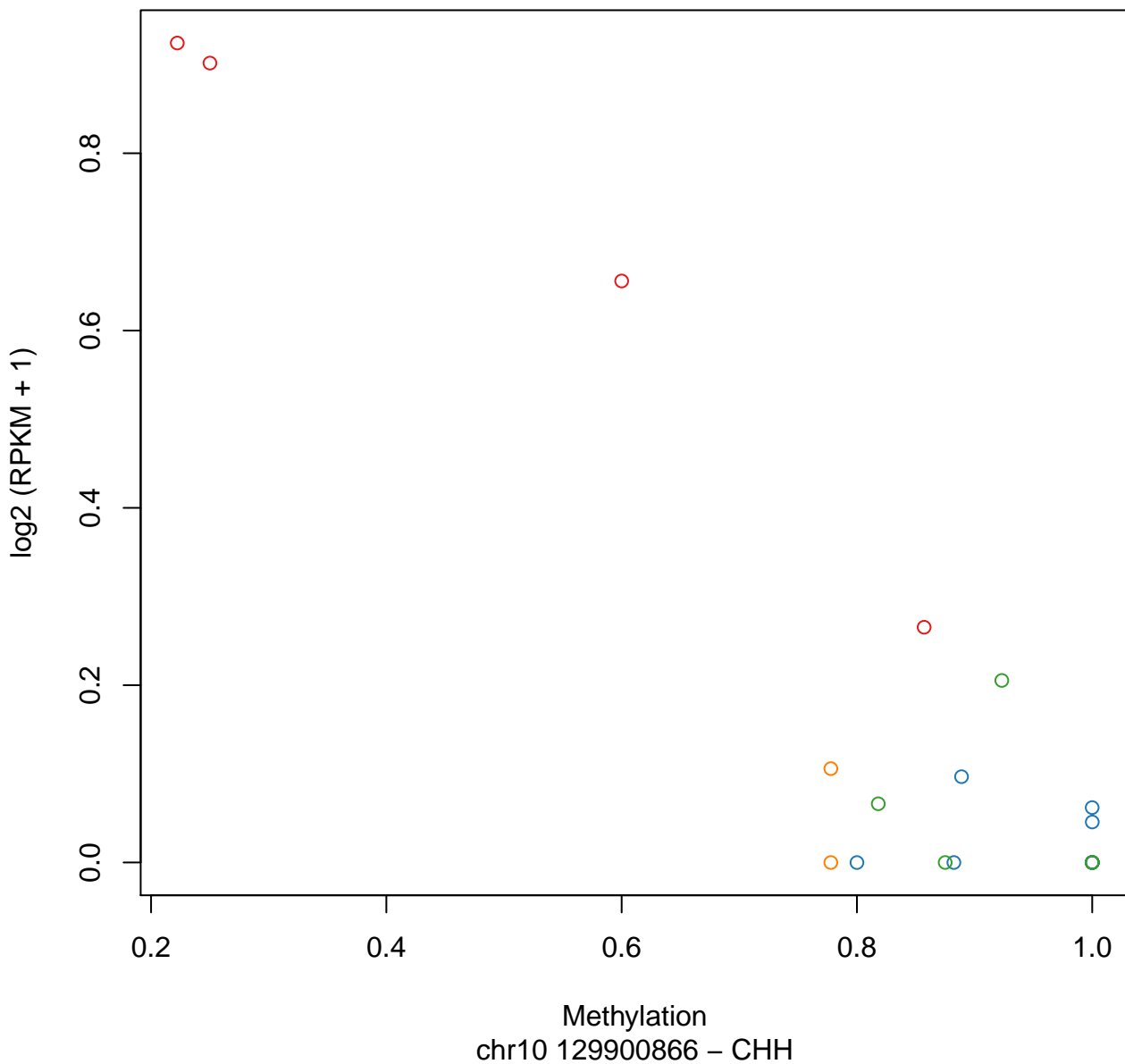
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CDRT15L2



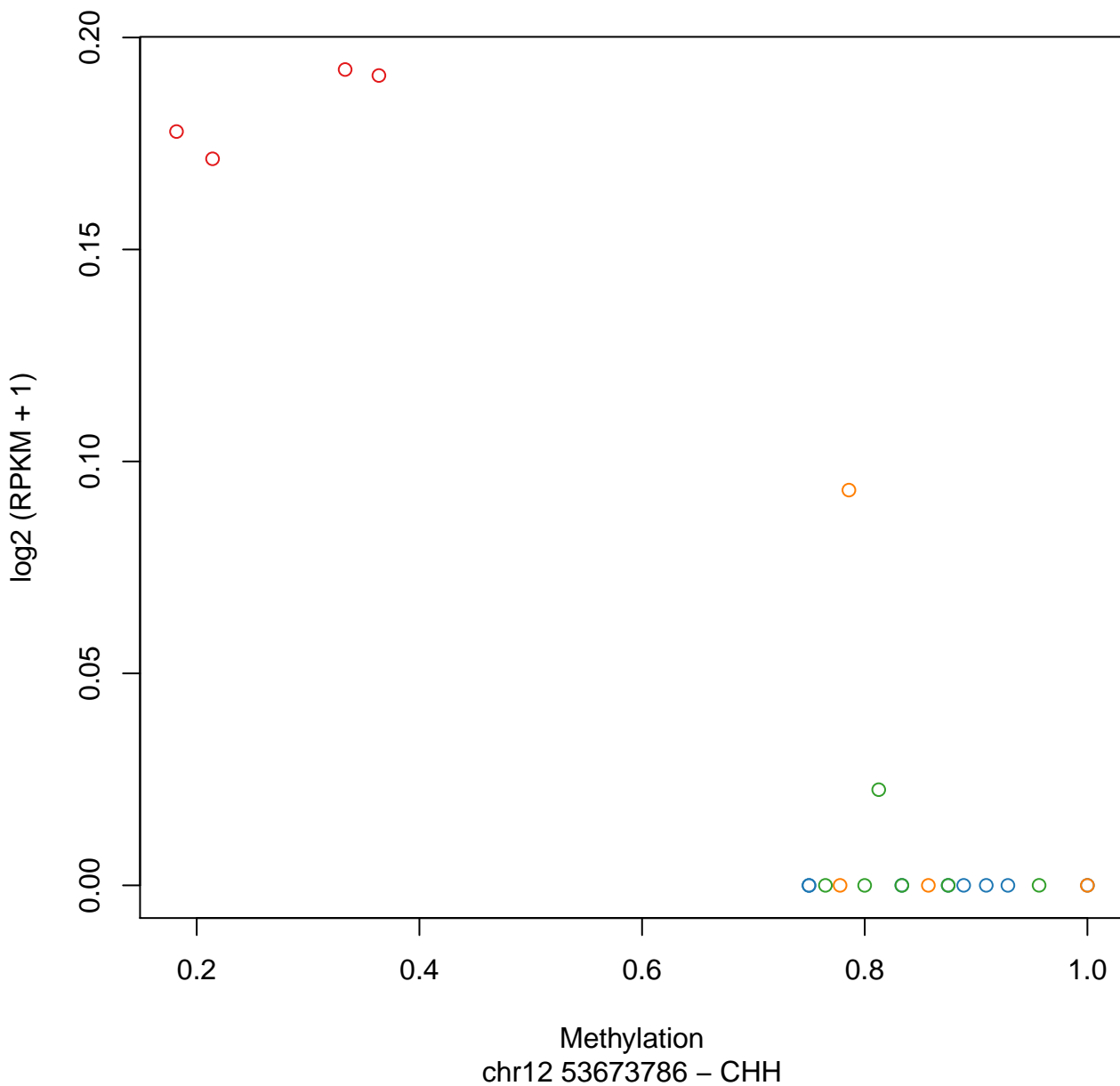
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MKI67

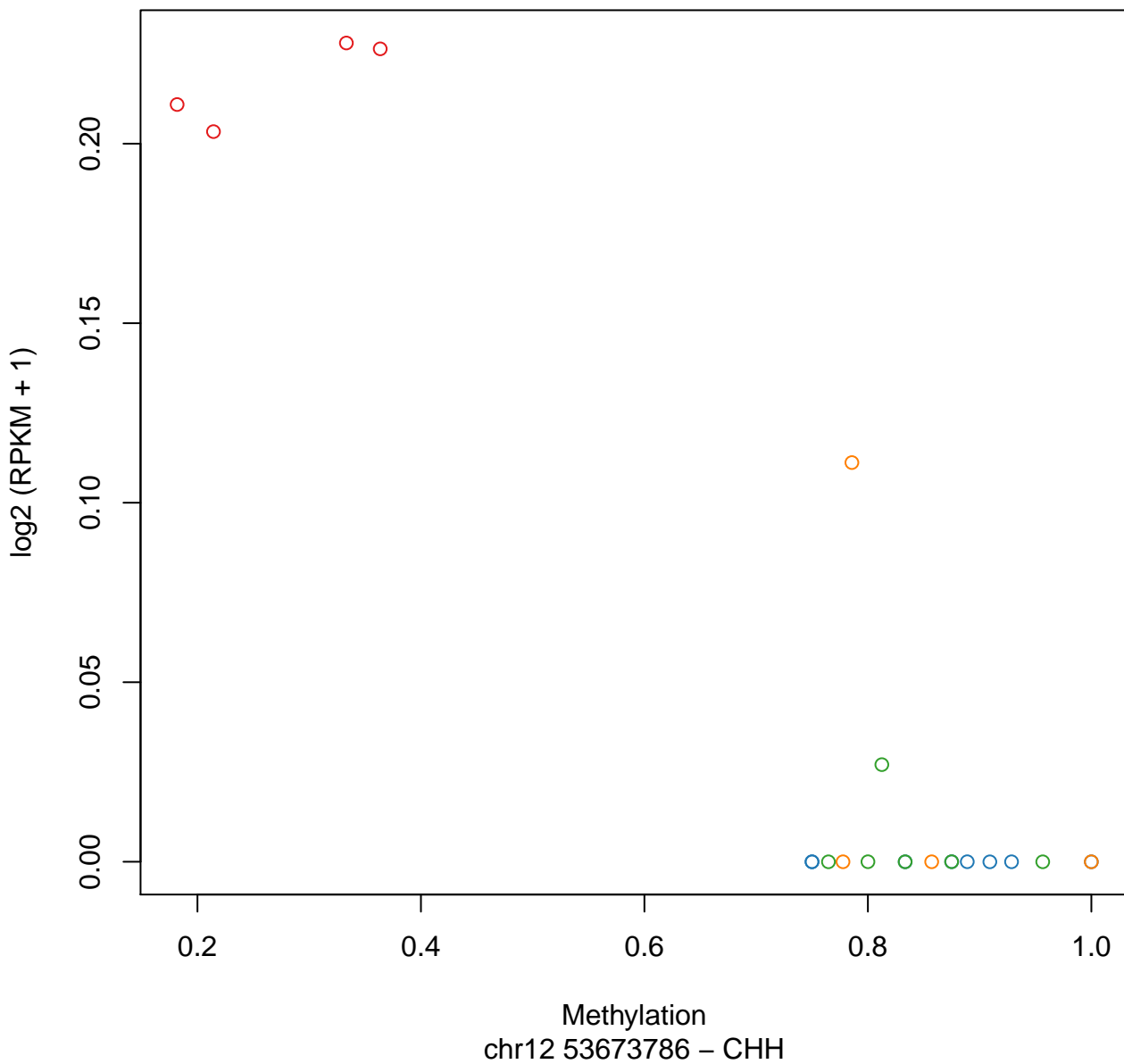


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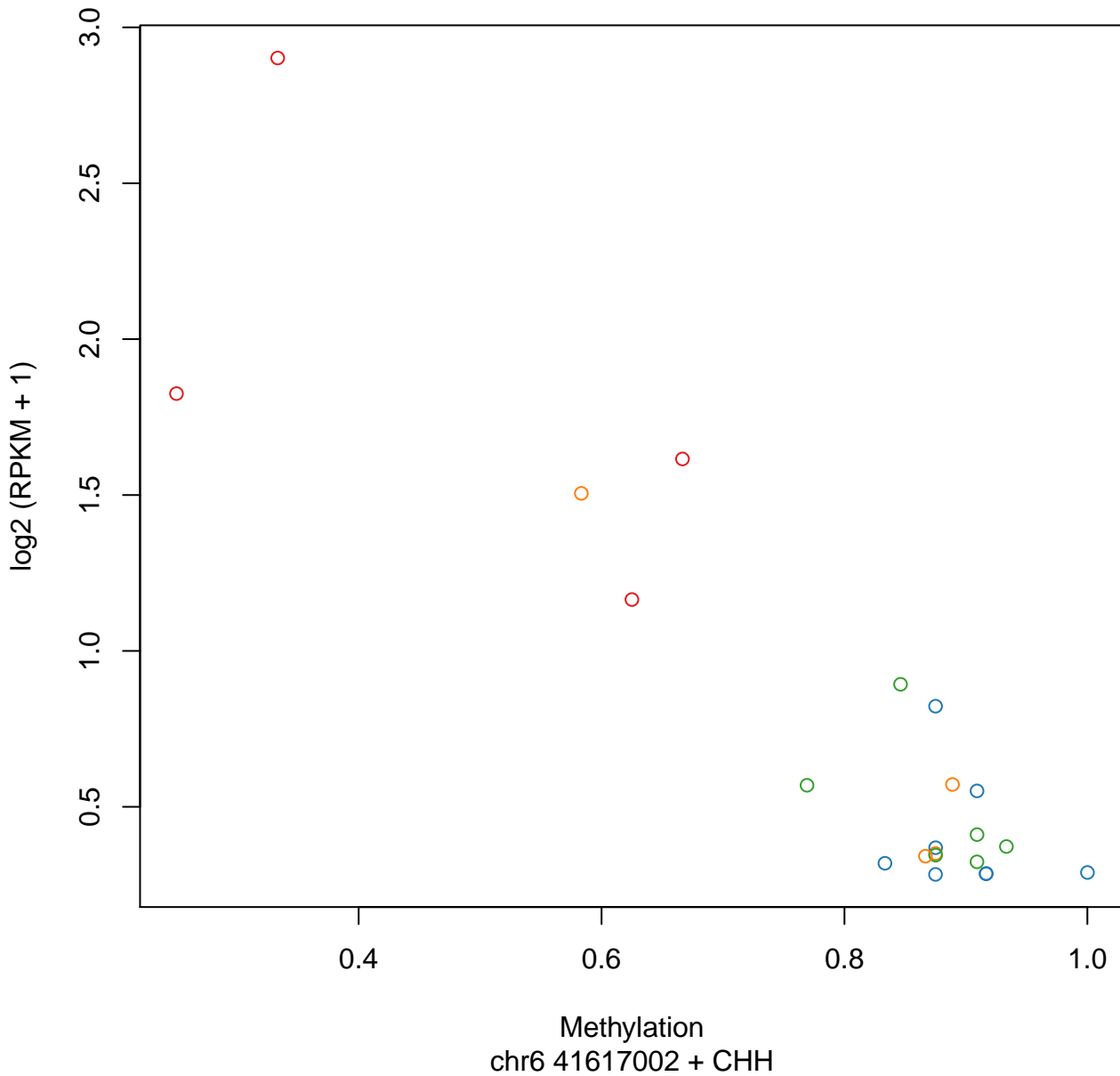
ESPL1



exon ENSE00003520699.1_1 FDR 9.71e-06
ESPL1

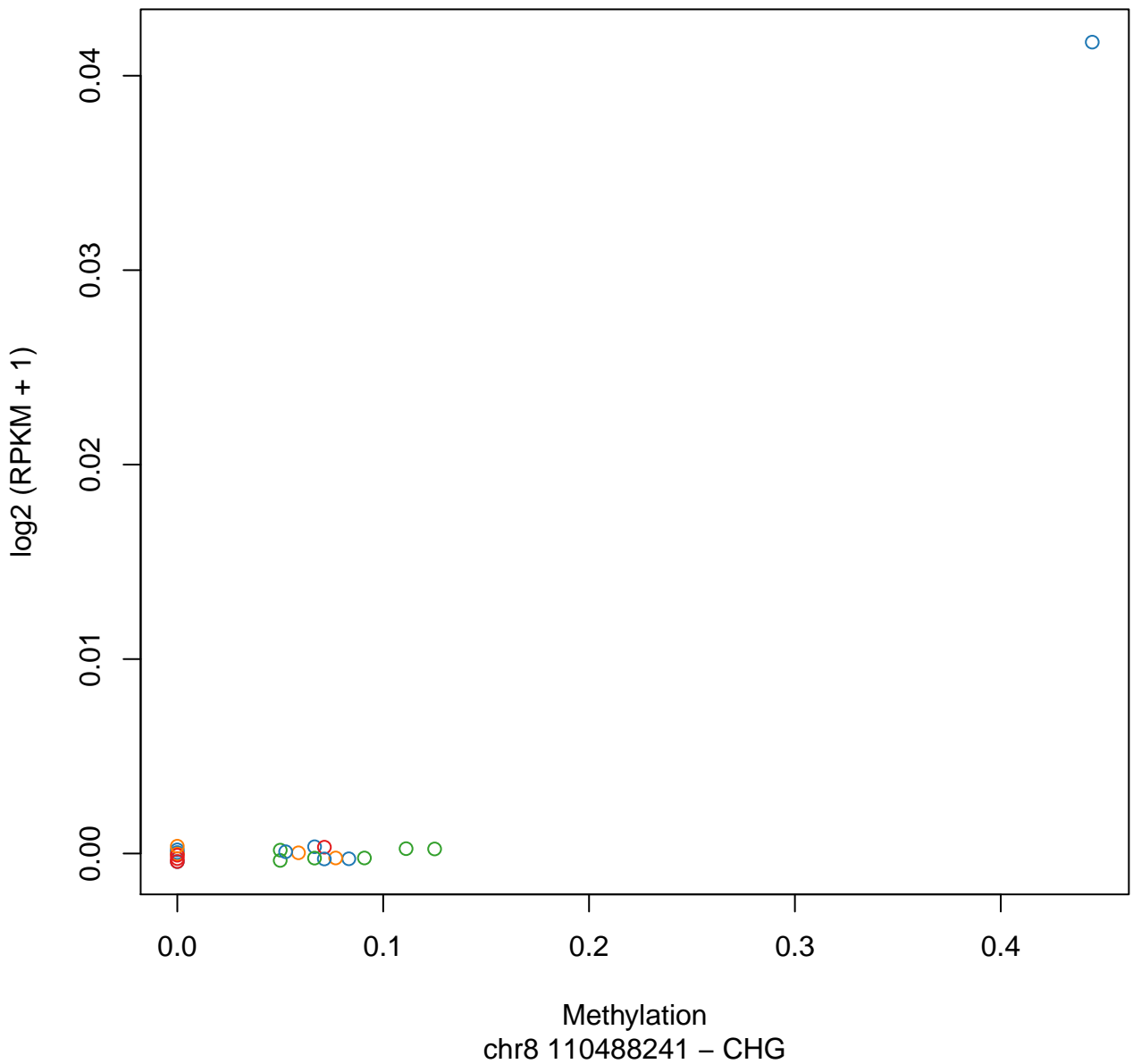


exon ENSE00001913030.1_1 FDR 4.68e-05
MDFI

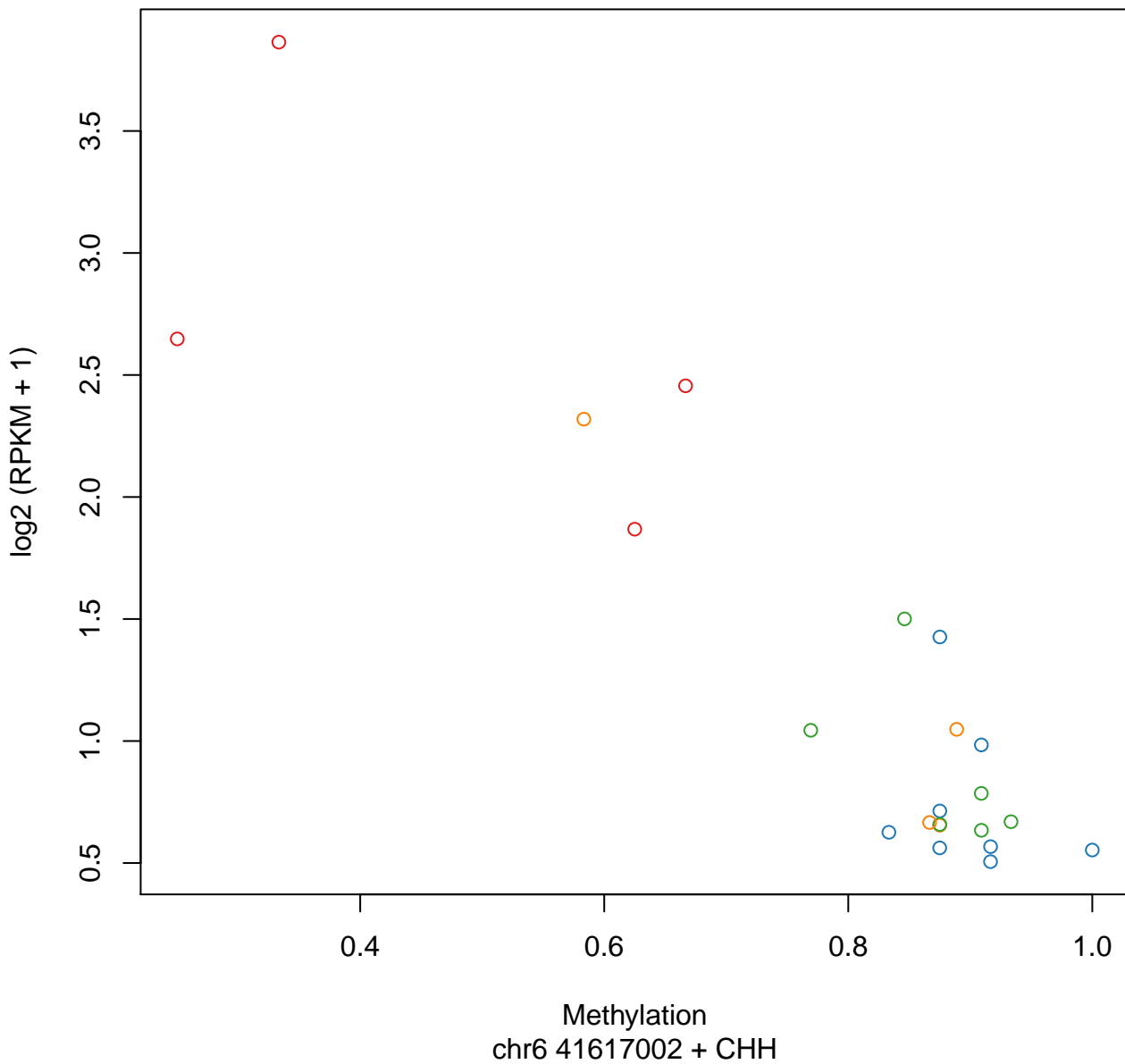


exon ENSE00001477397.1_1 FDR 5.3e-05

PKHD1L1

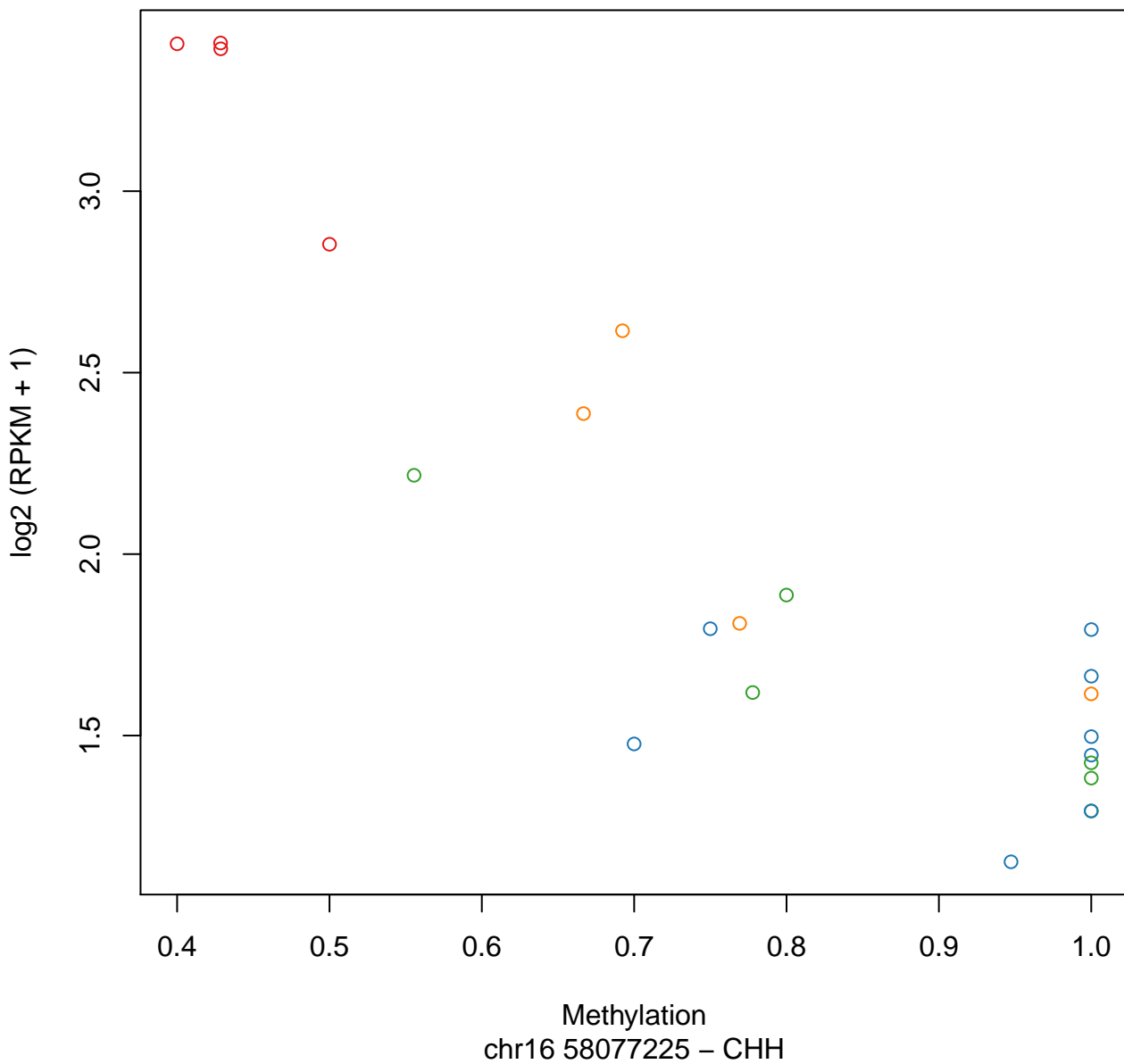


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MDFI

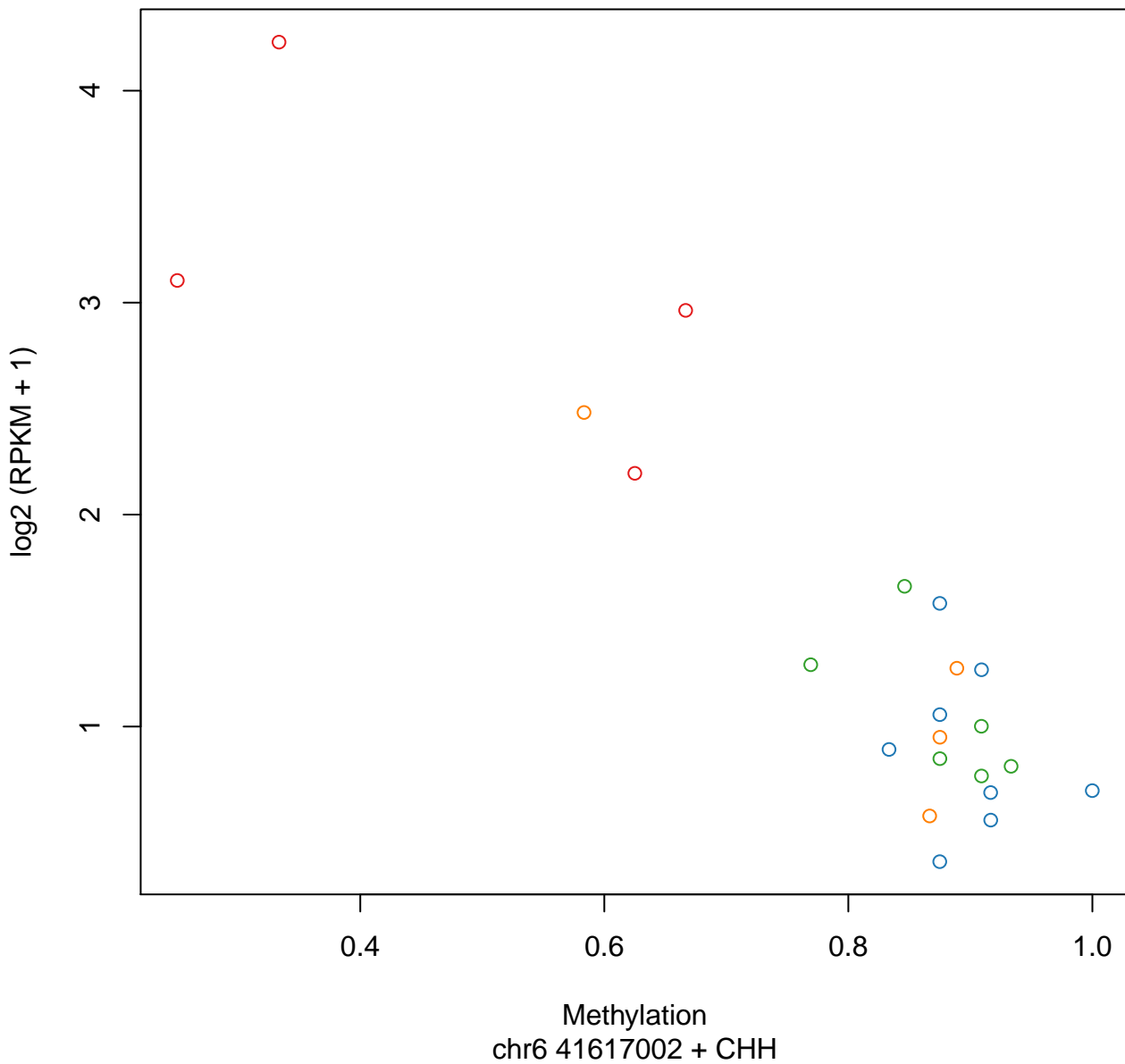


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MMP15

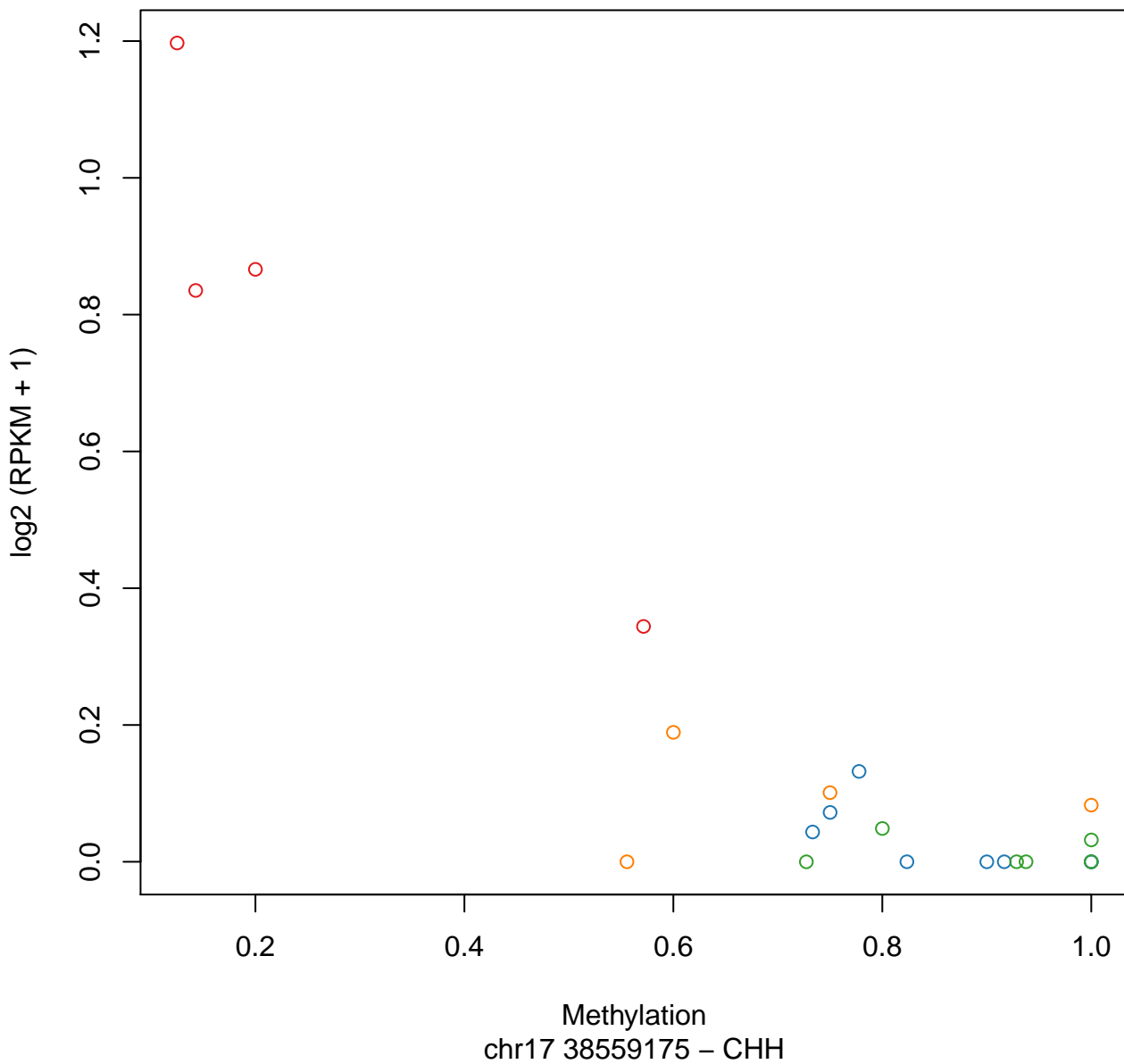


exon ENSE00001728638.1_1 FDR 8.06e-05
MDFI



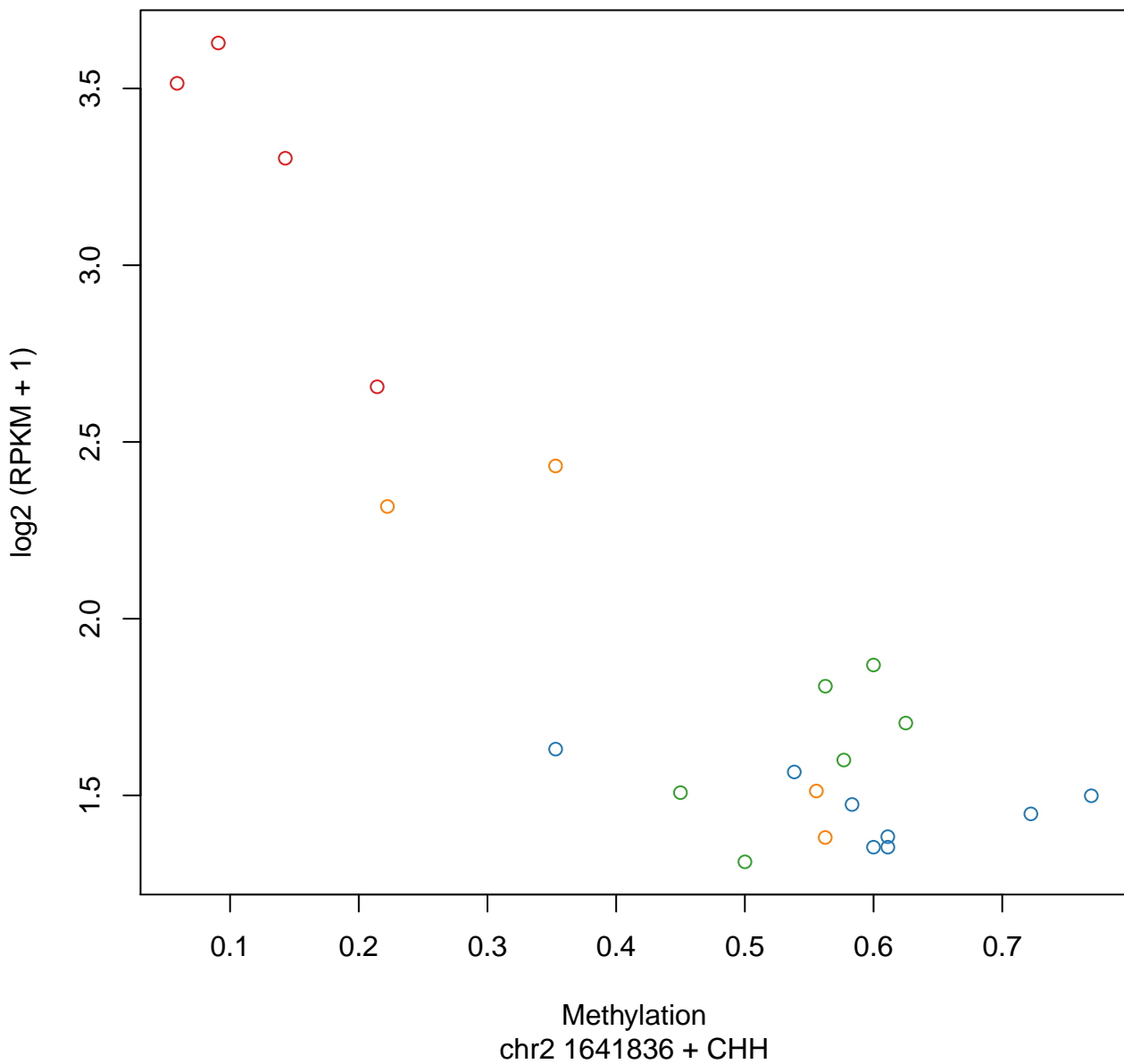
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TOP2A



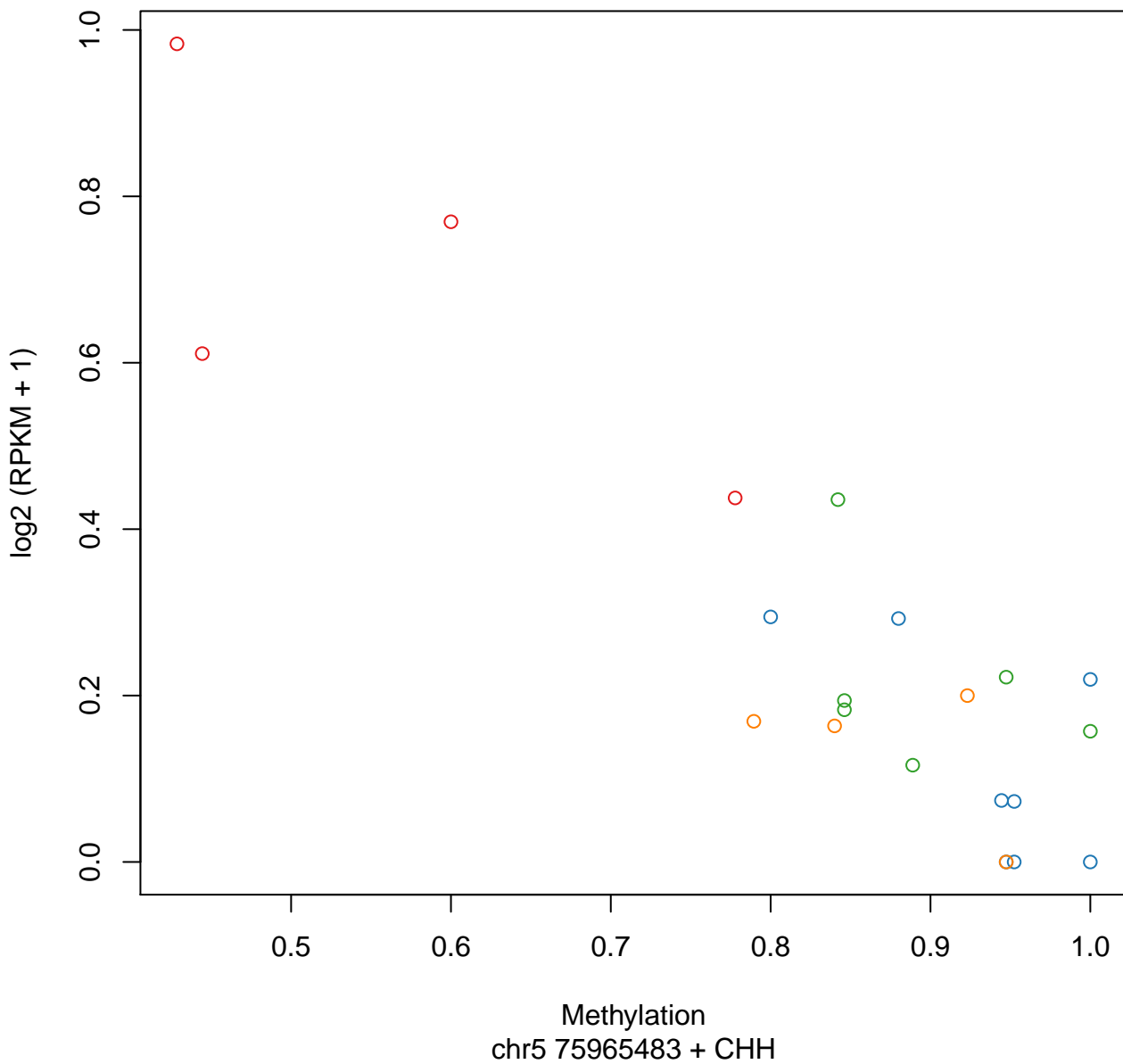
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PXDN

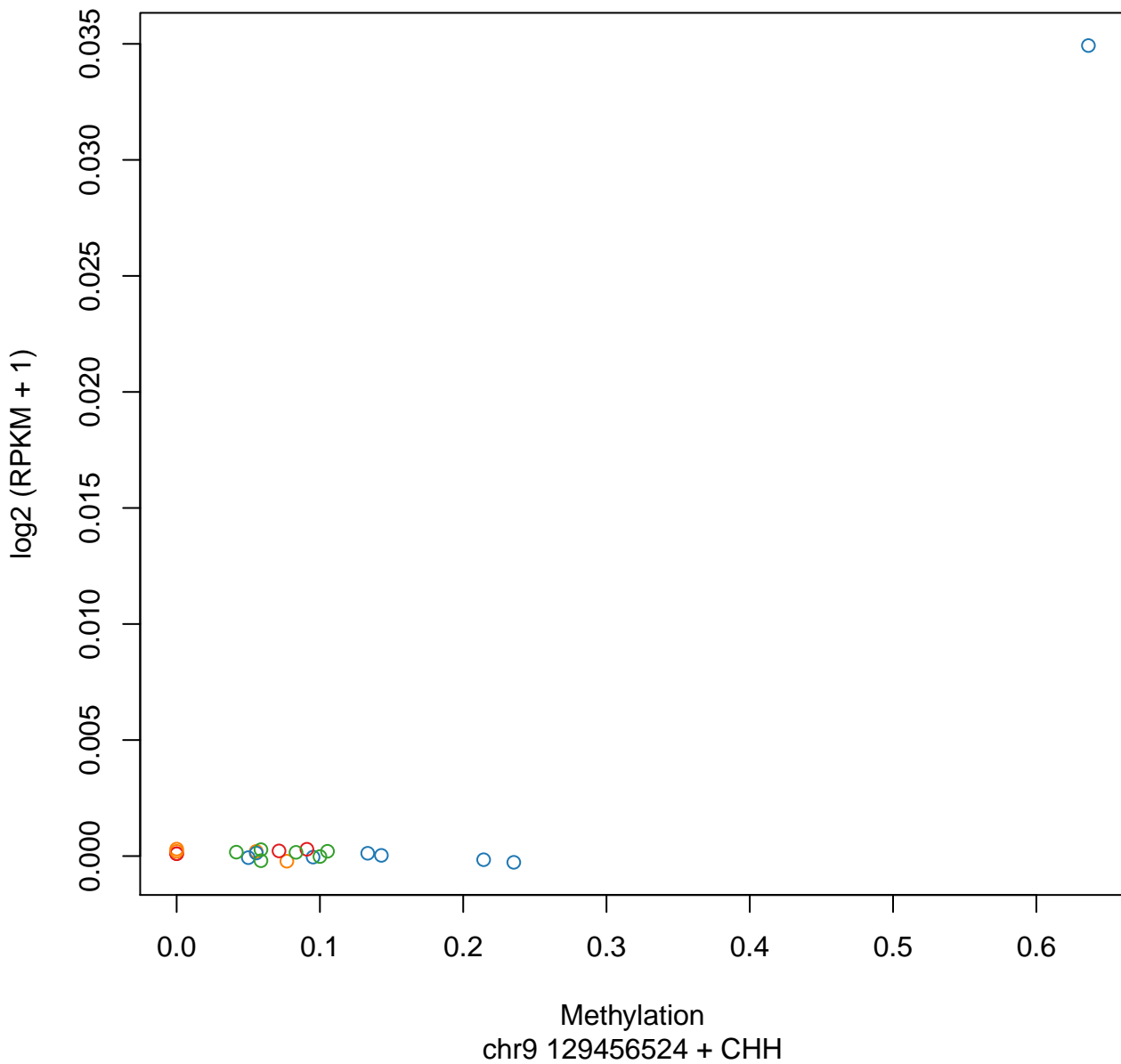


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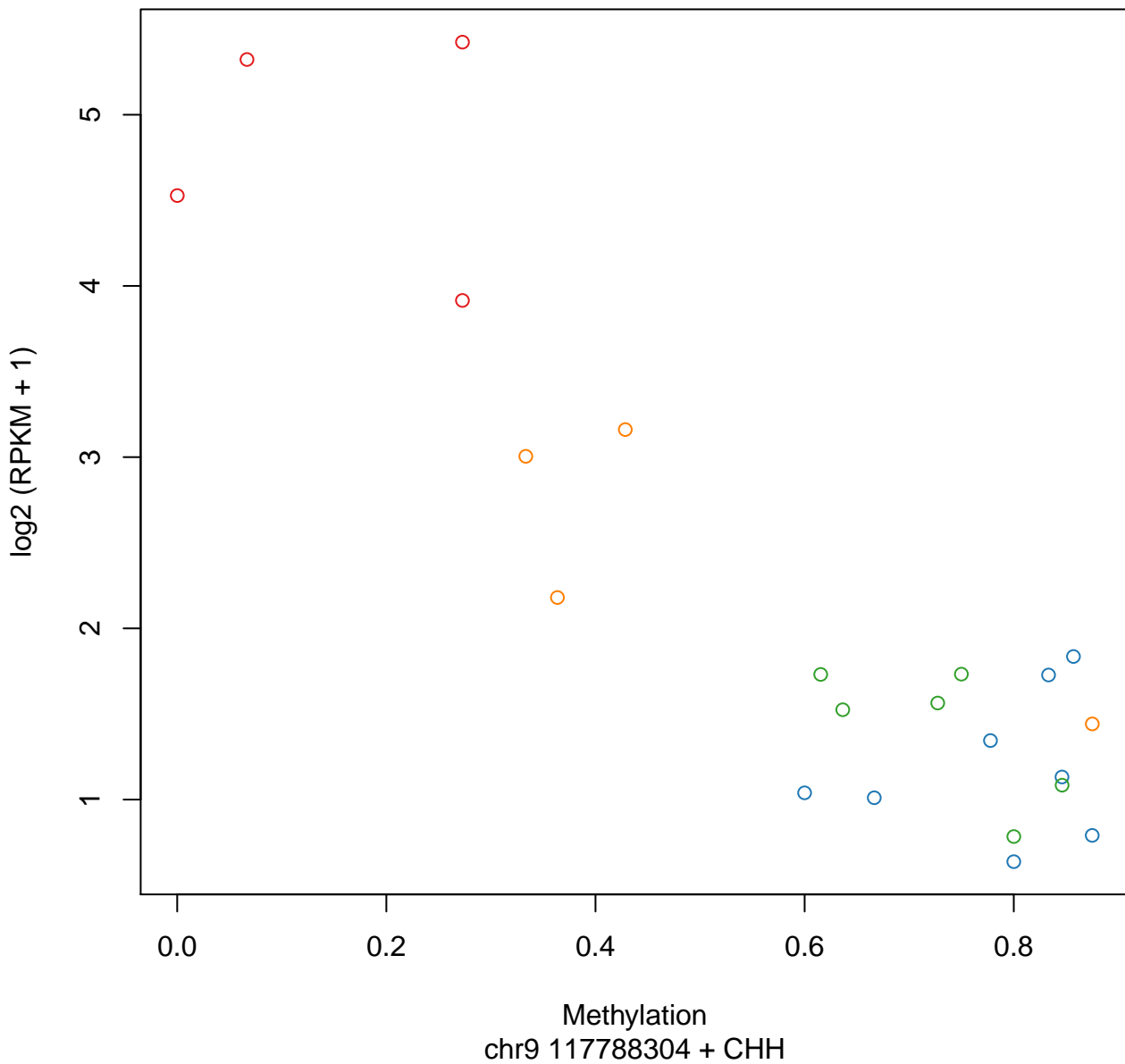
IQGAP2



LMX1B

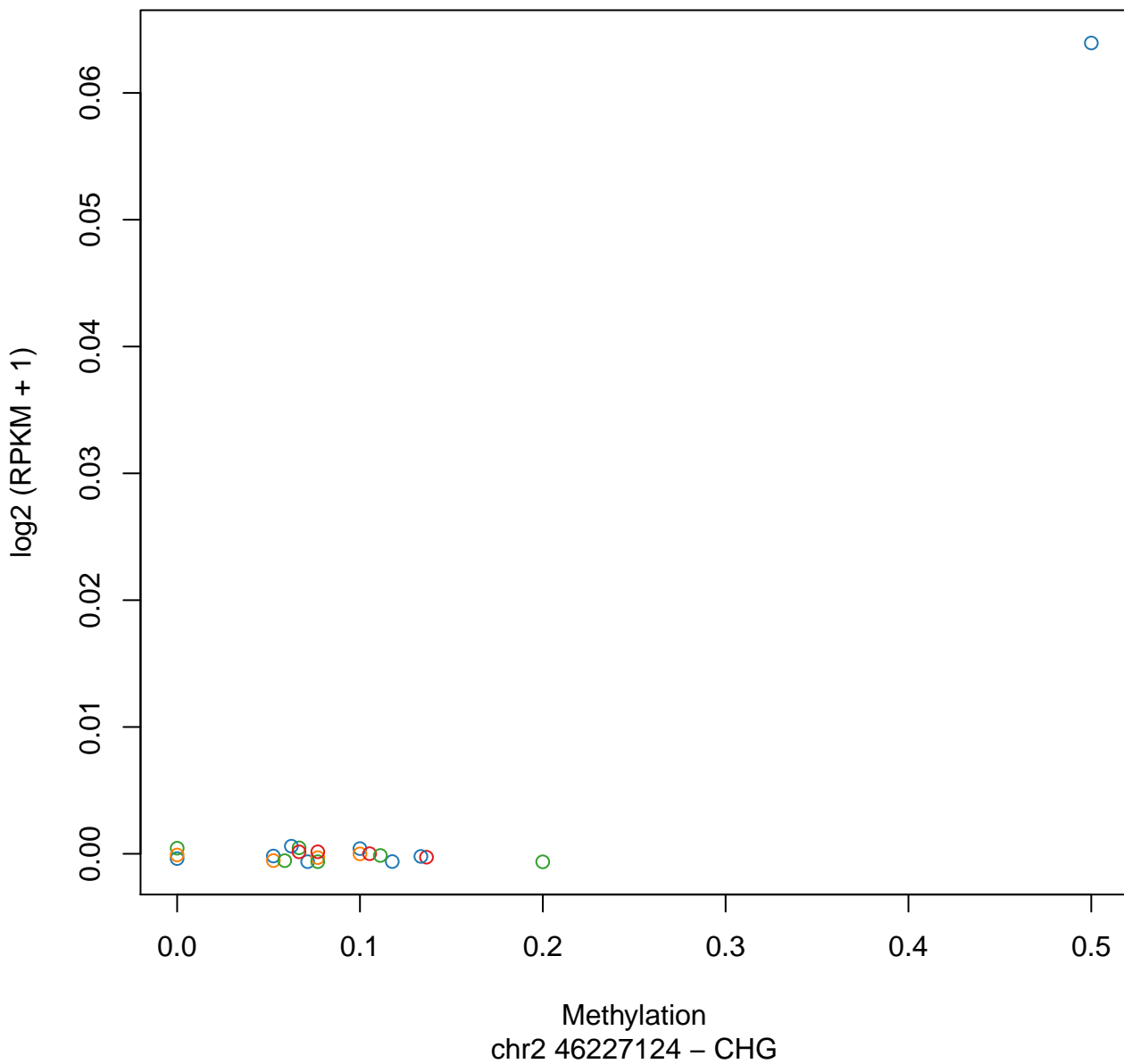


exon ENSE00000926914.1_1 FDR 0.000206
TNC



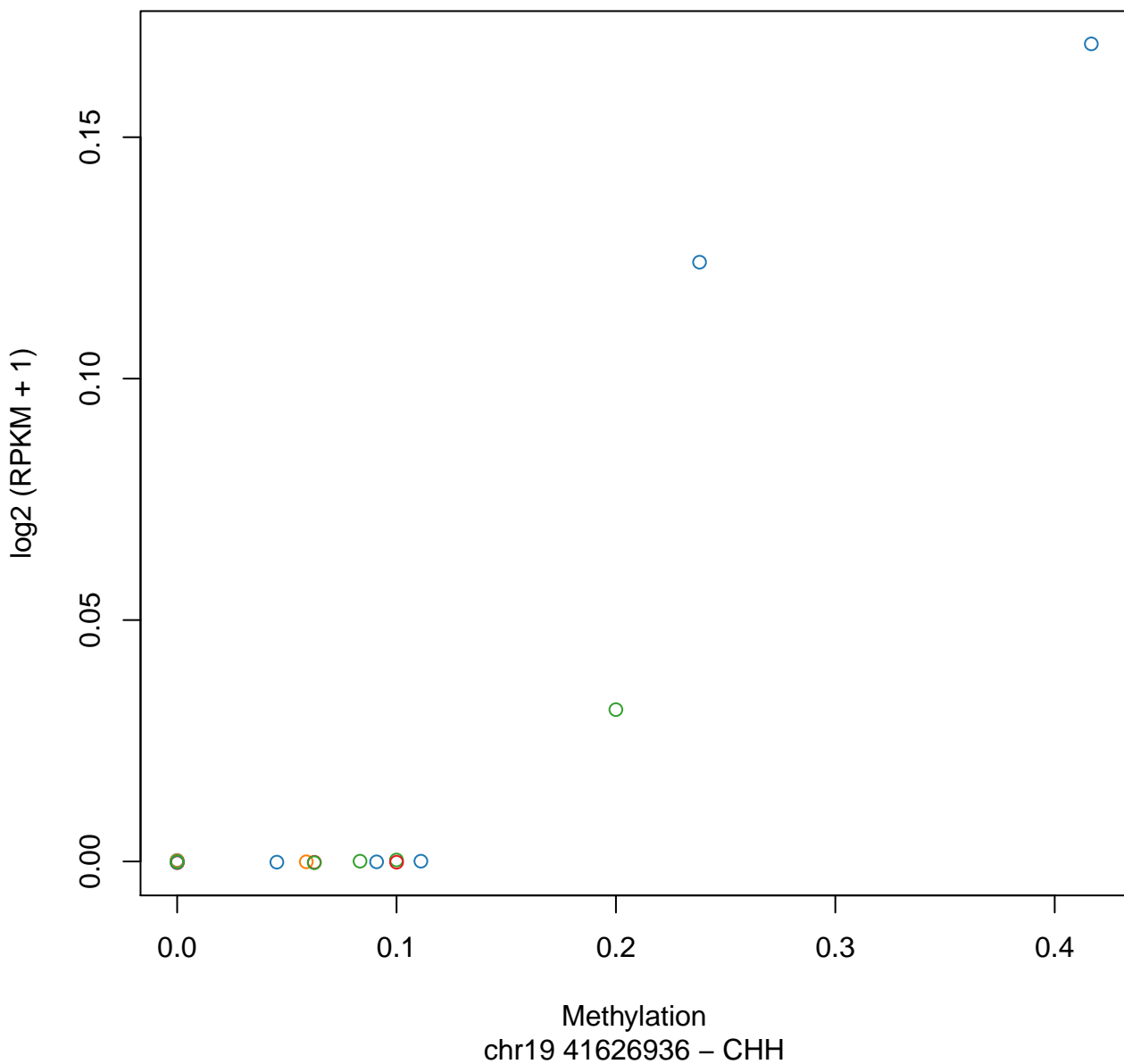
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PRKCE



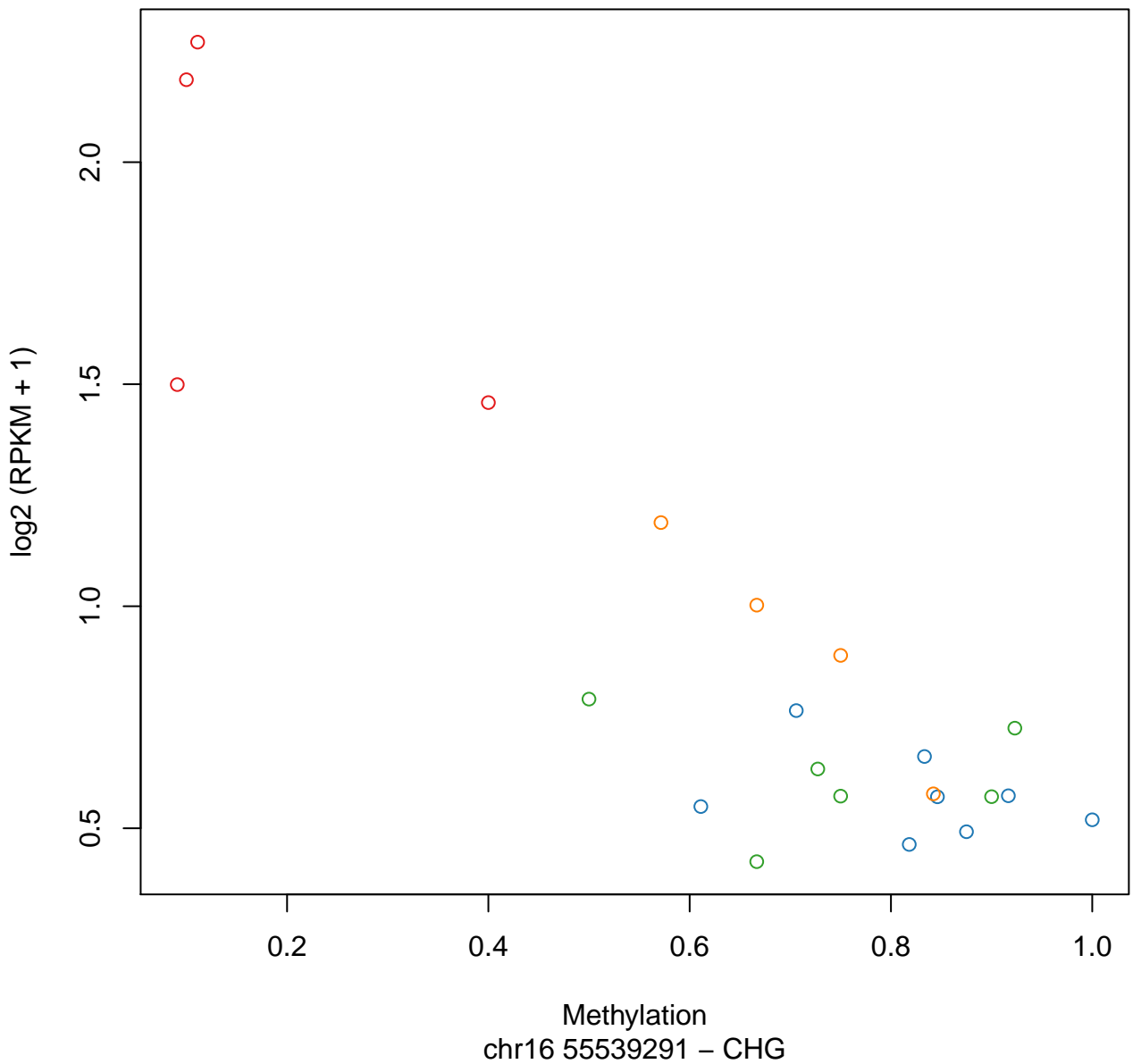
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CYP2F1



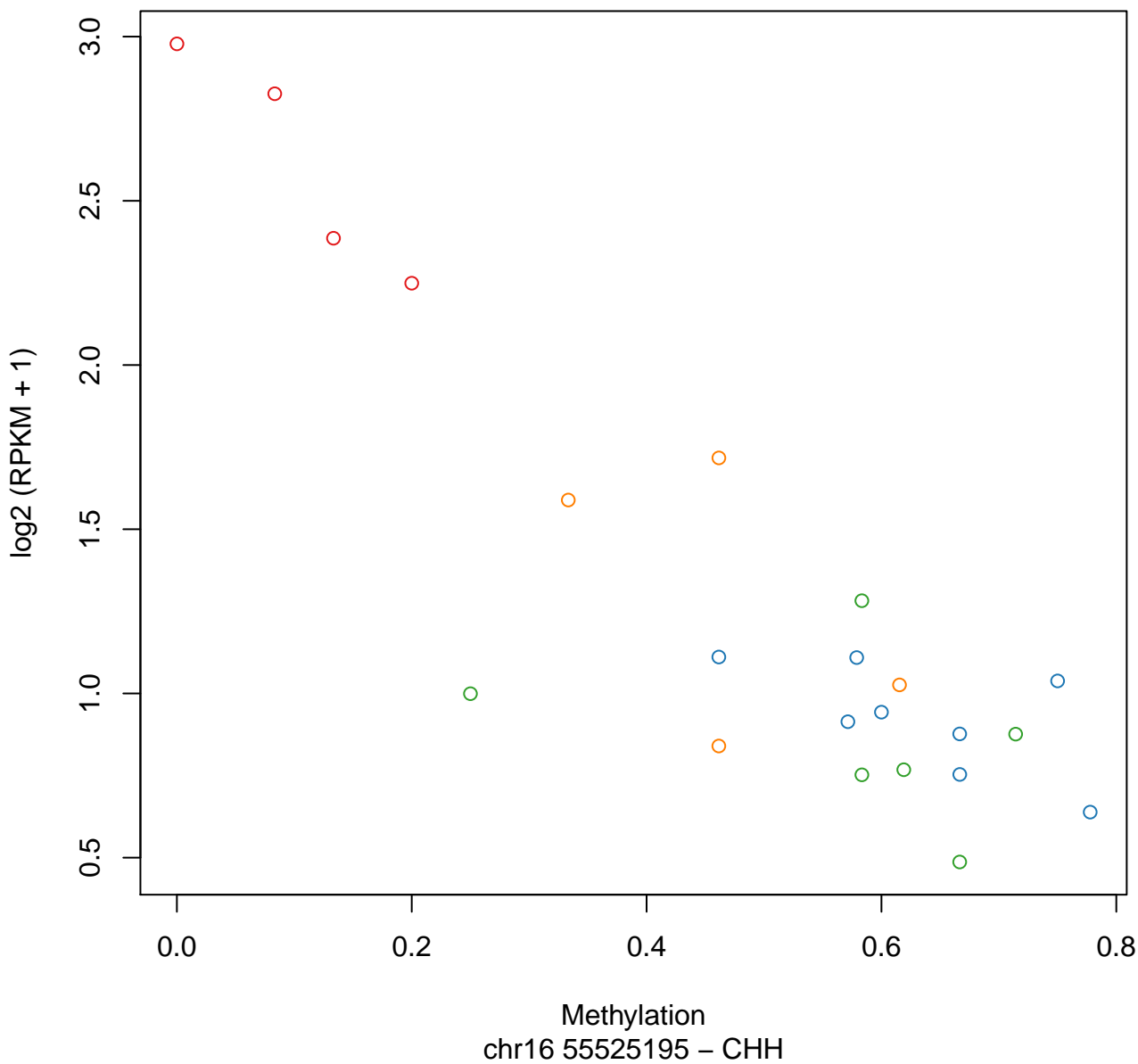
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MMP2

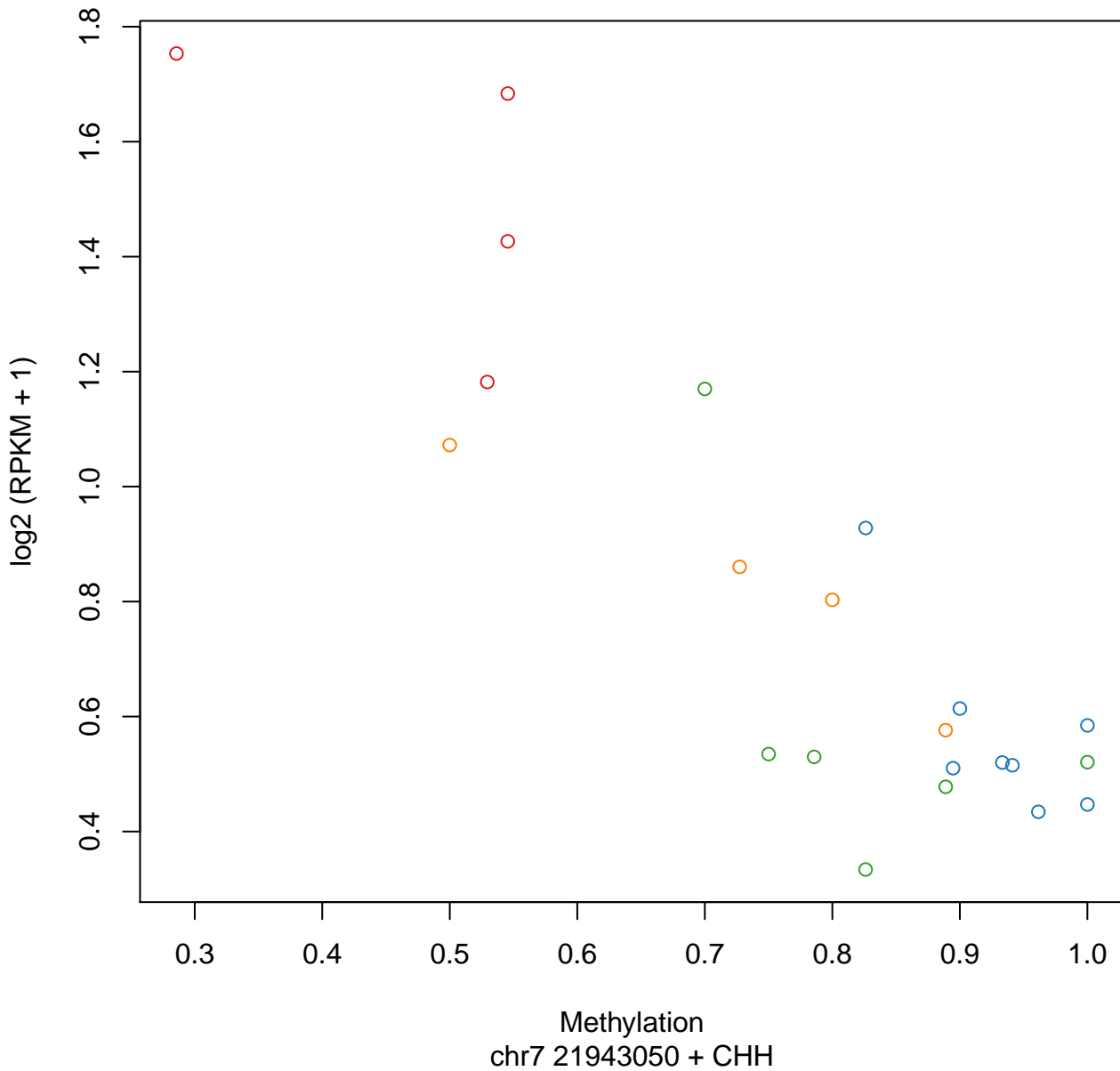


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MMP2

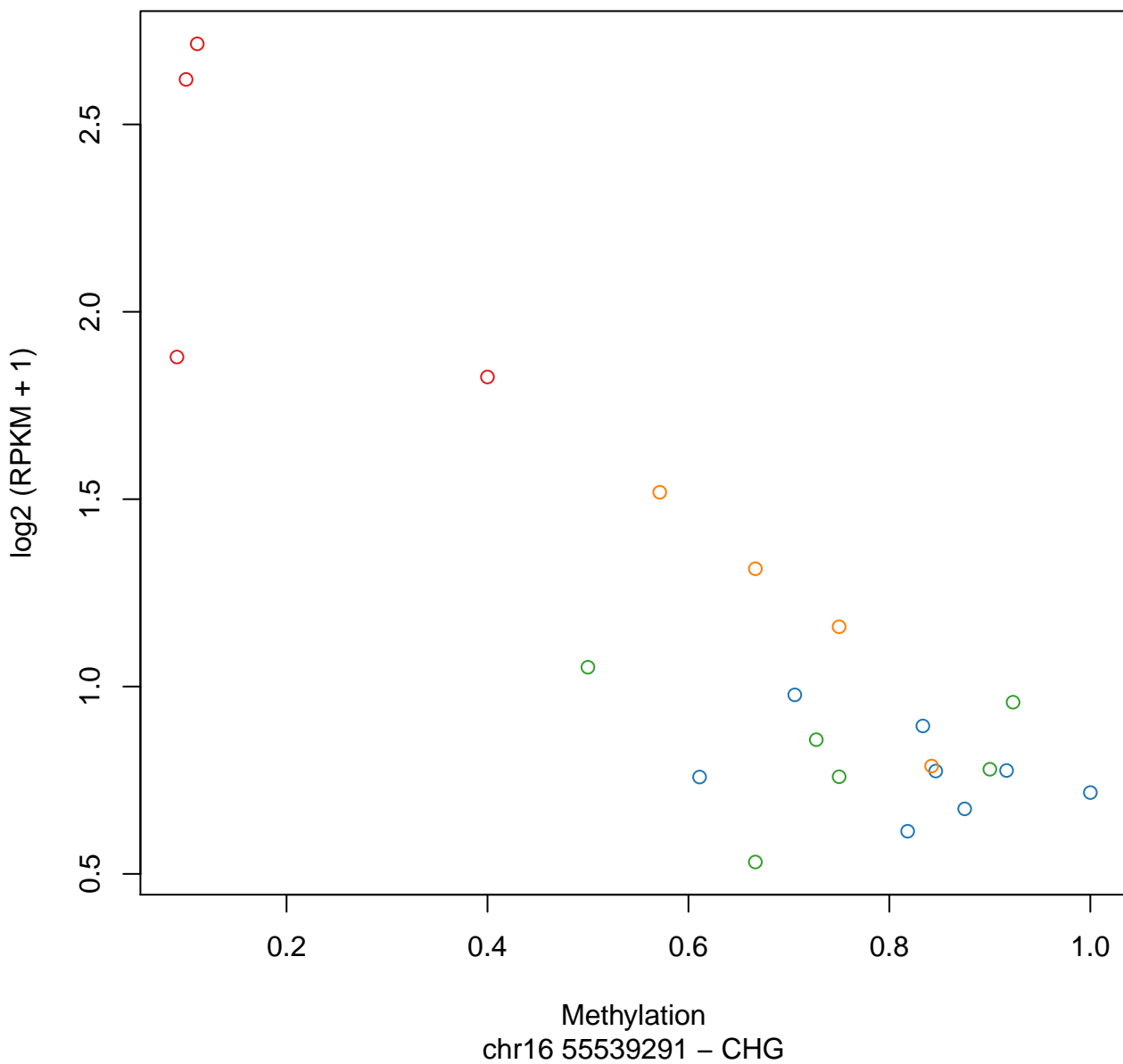


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CDCA7L



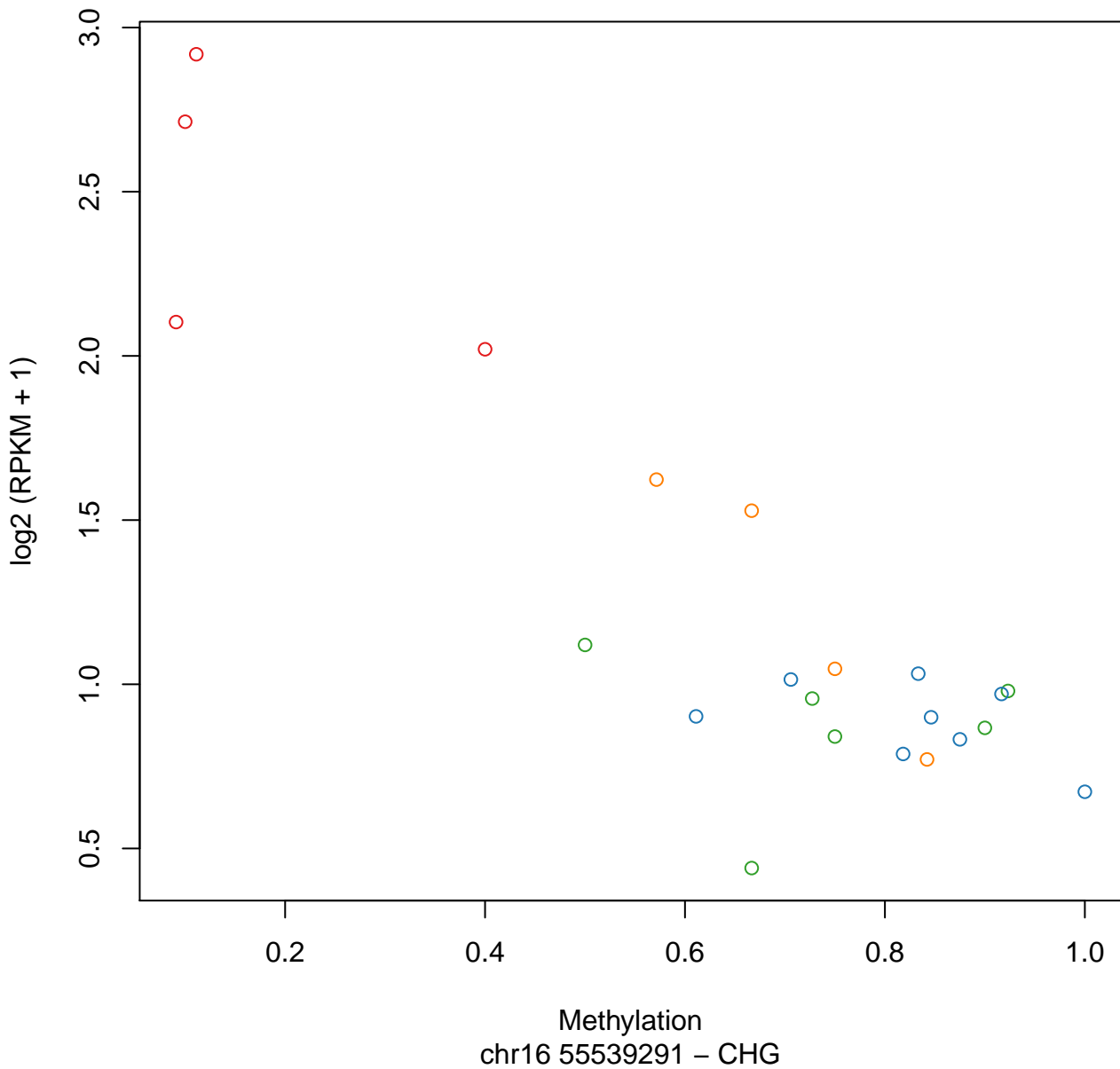
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MMP2



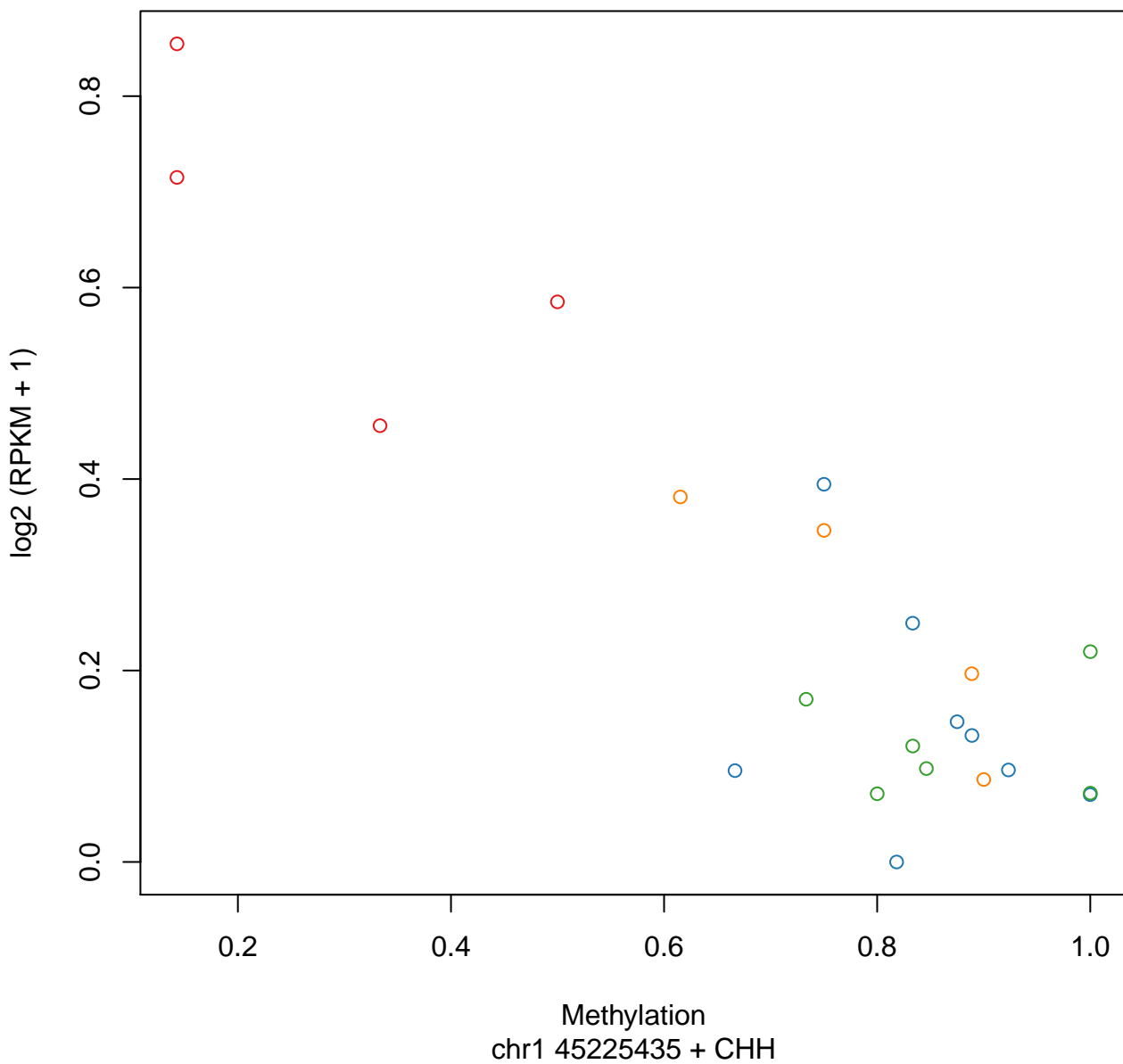
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MMP2



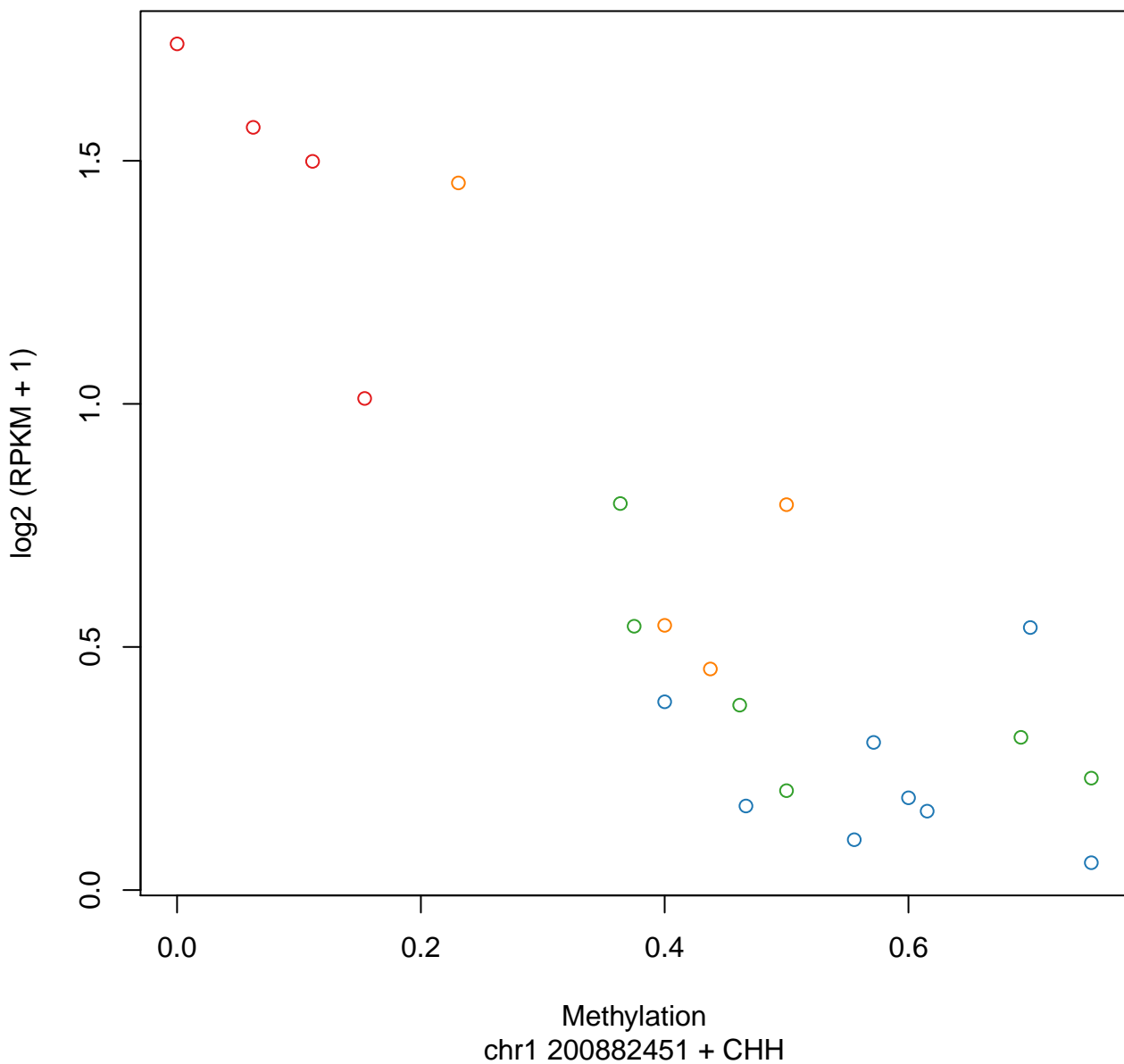
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KIF2C



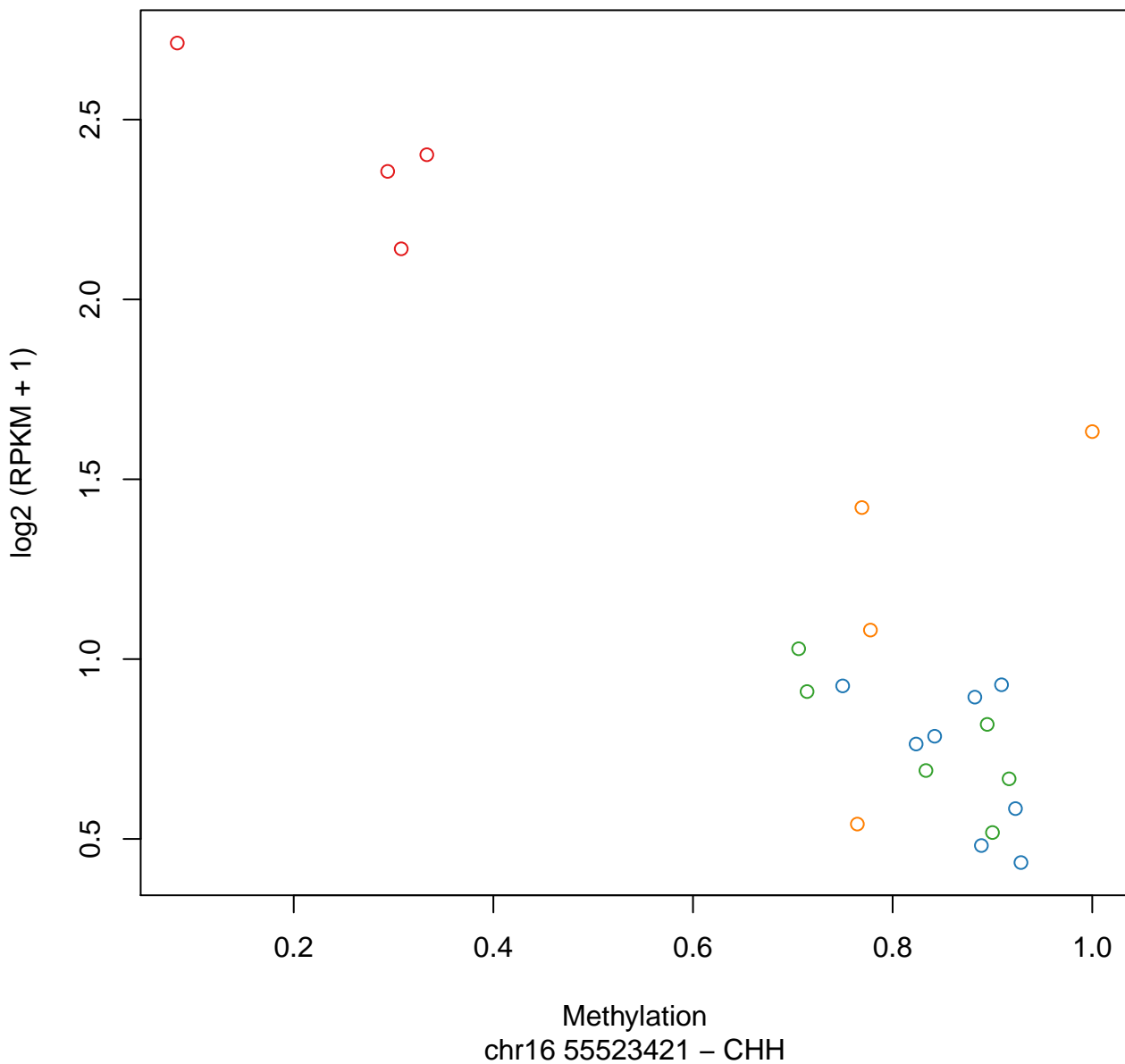
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C1orf106



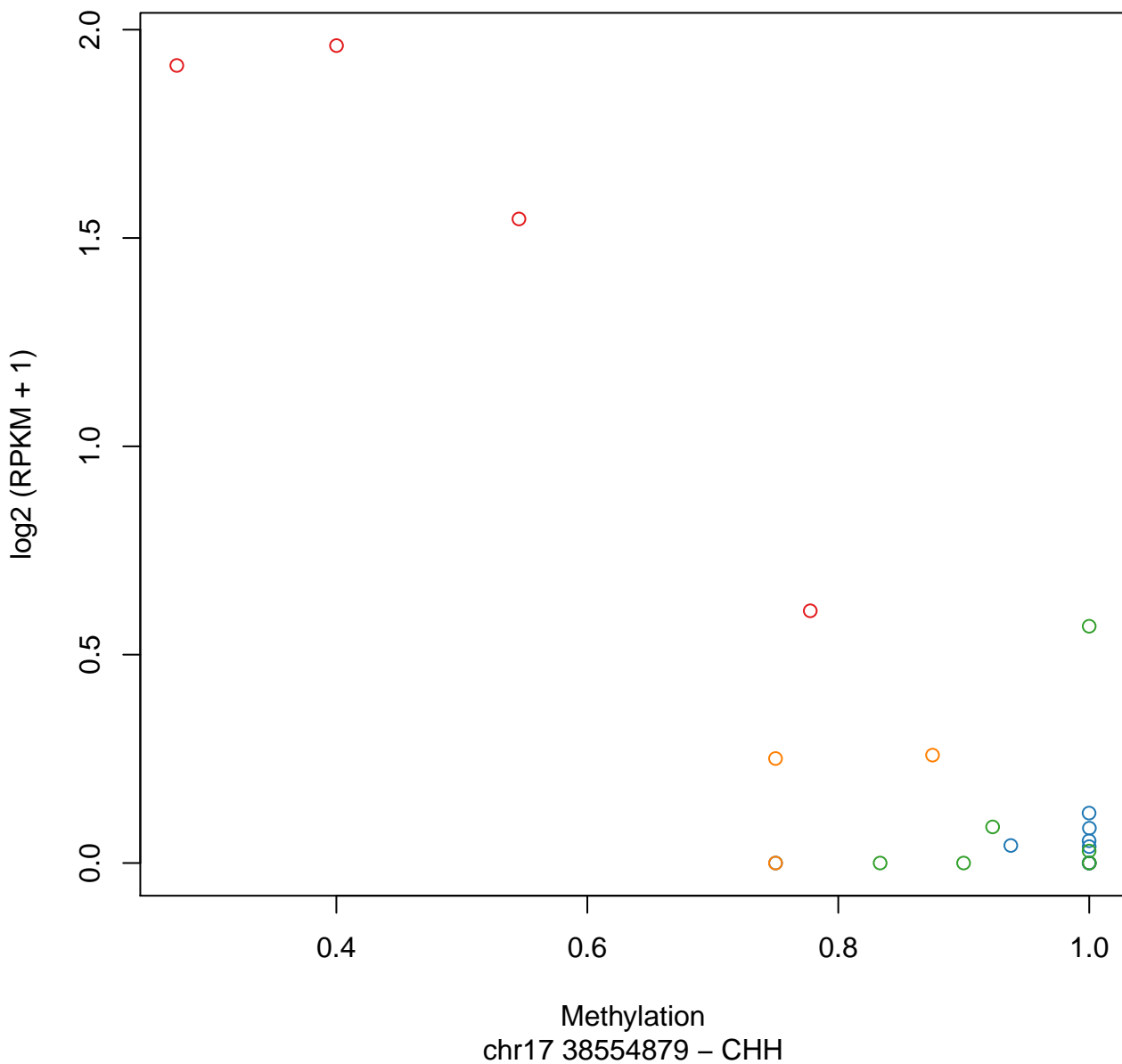
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MMP2



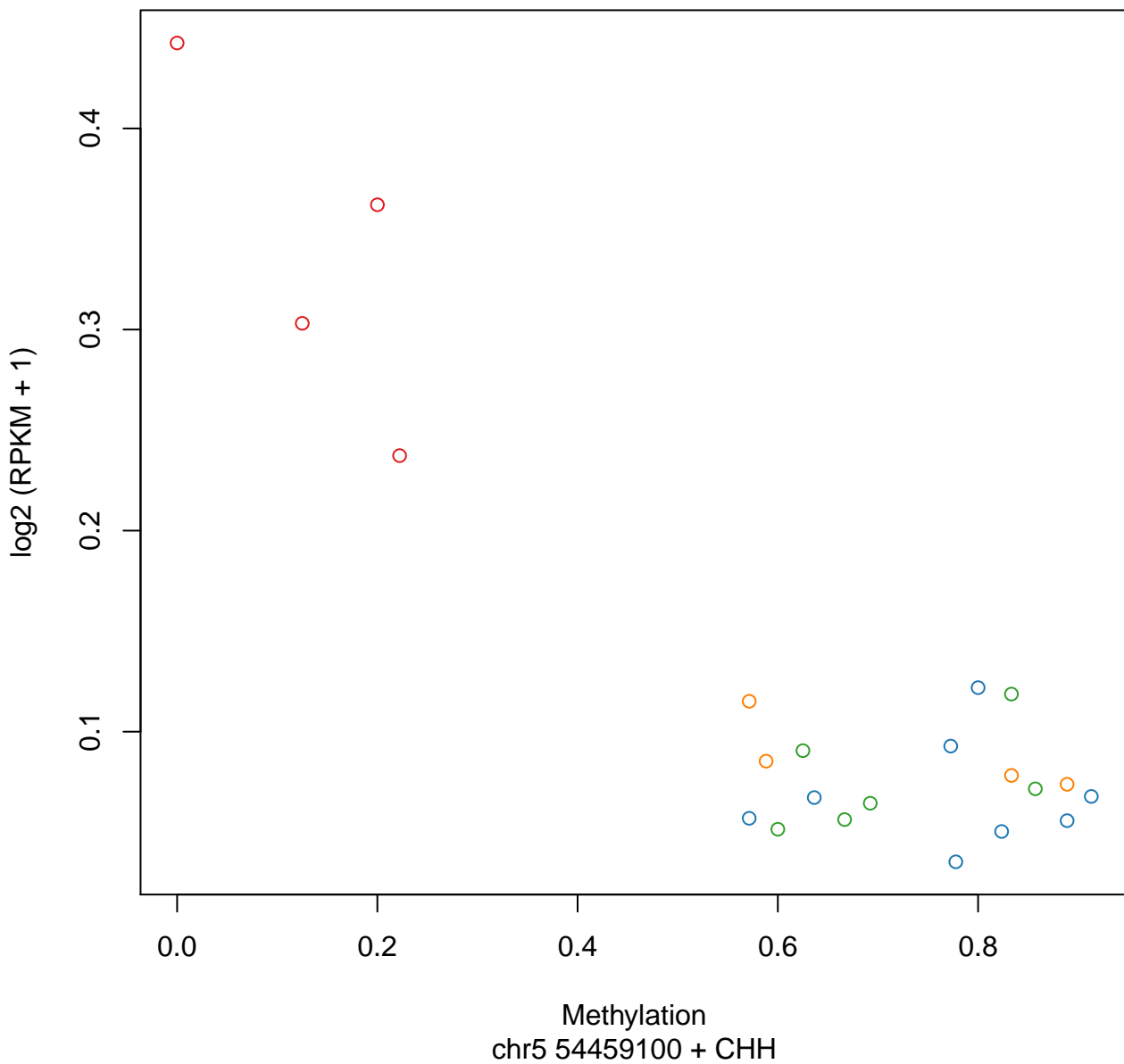
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TOP2A



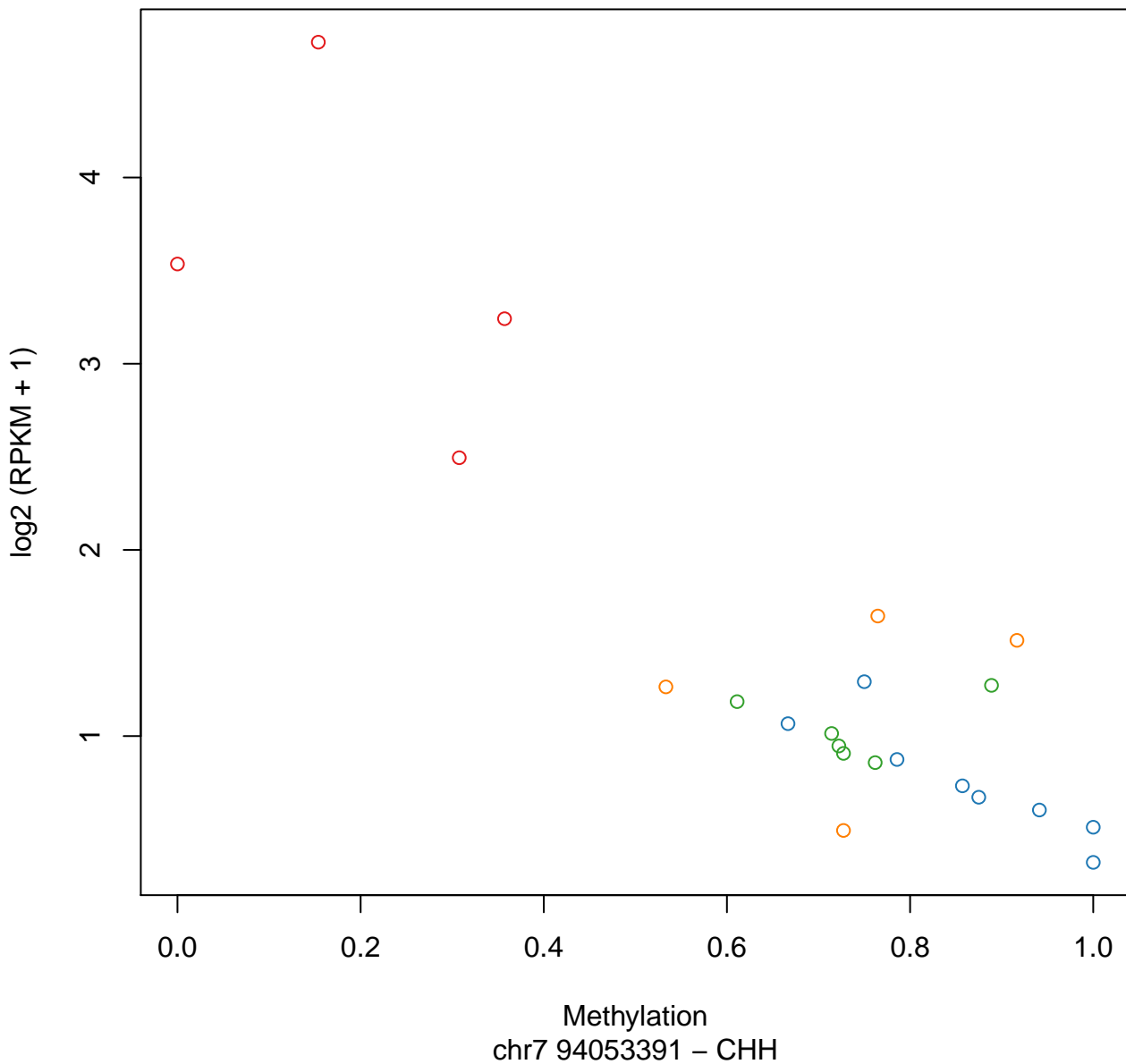
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GPX8



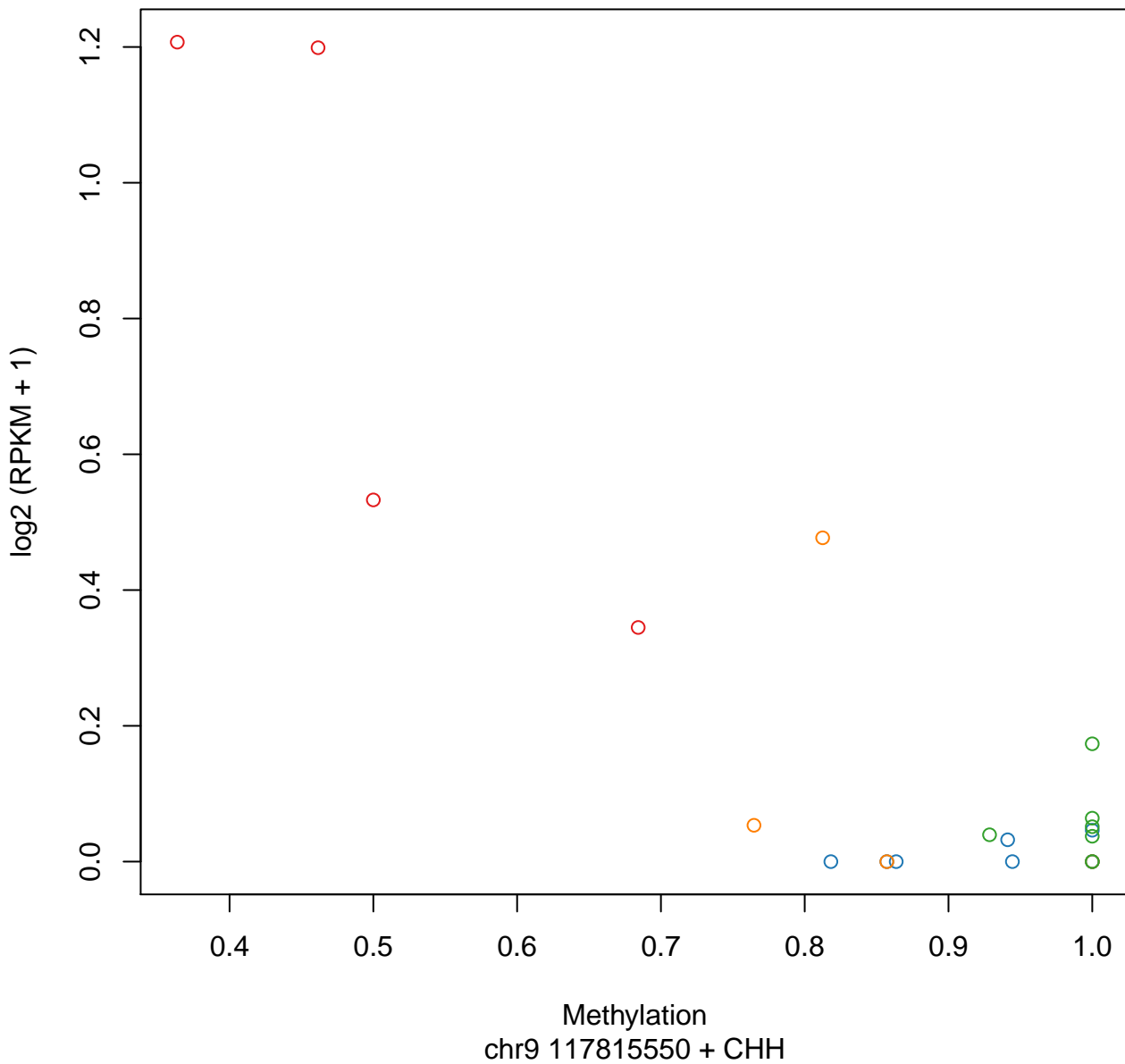
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COL1A2



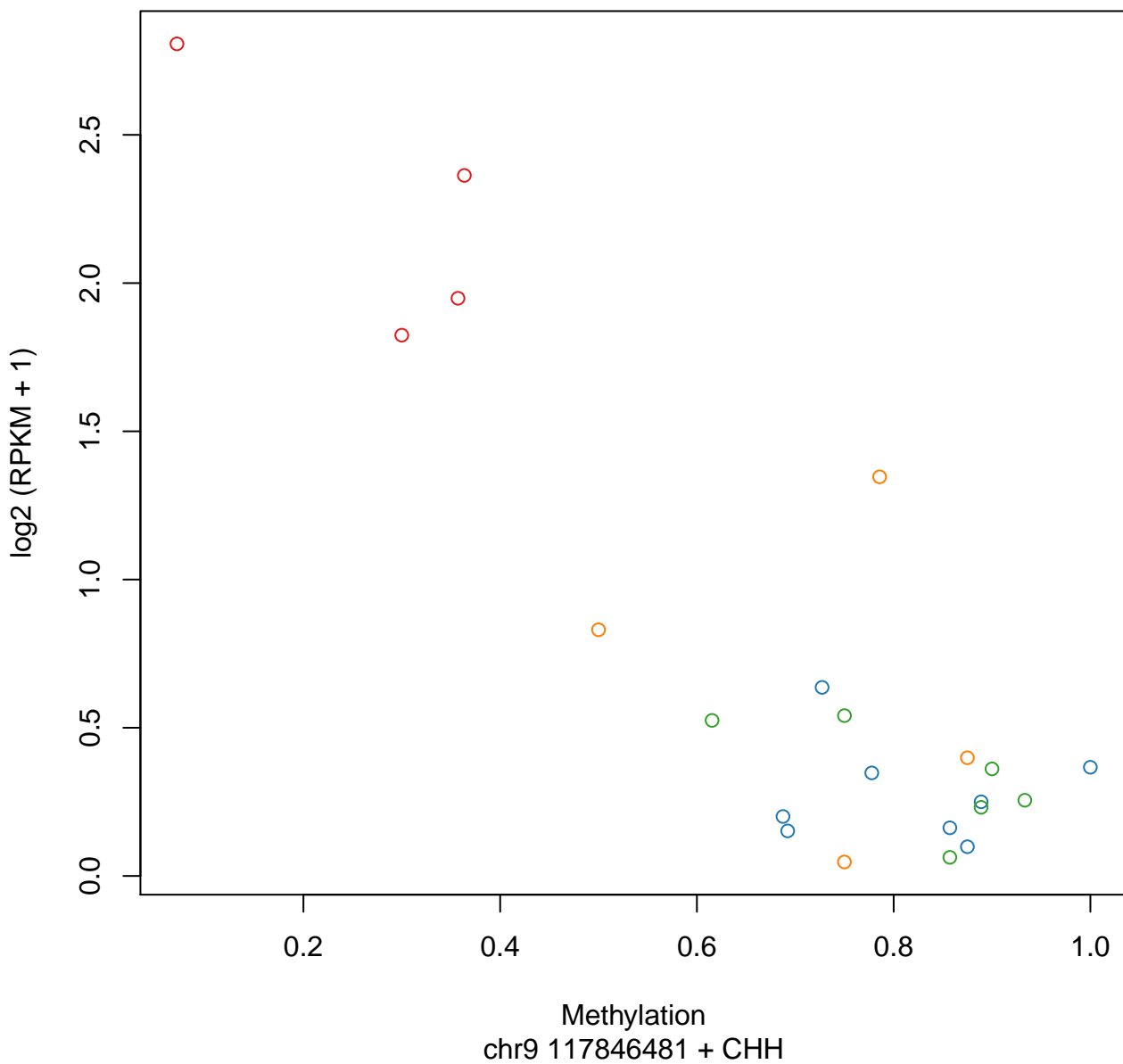
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TNC



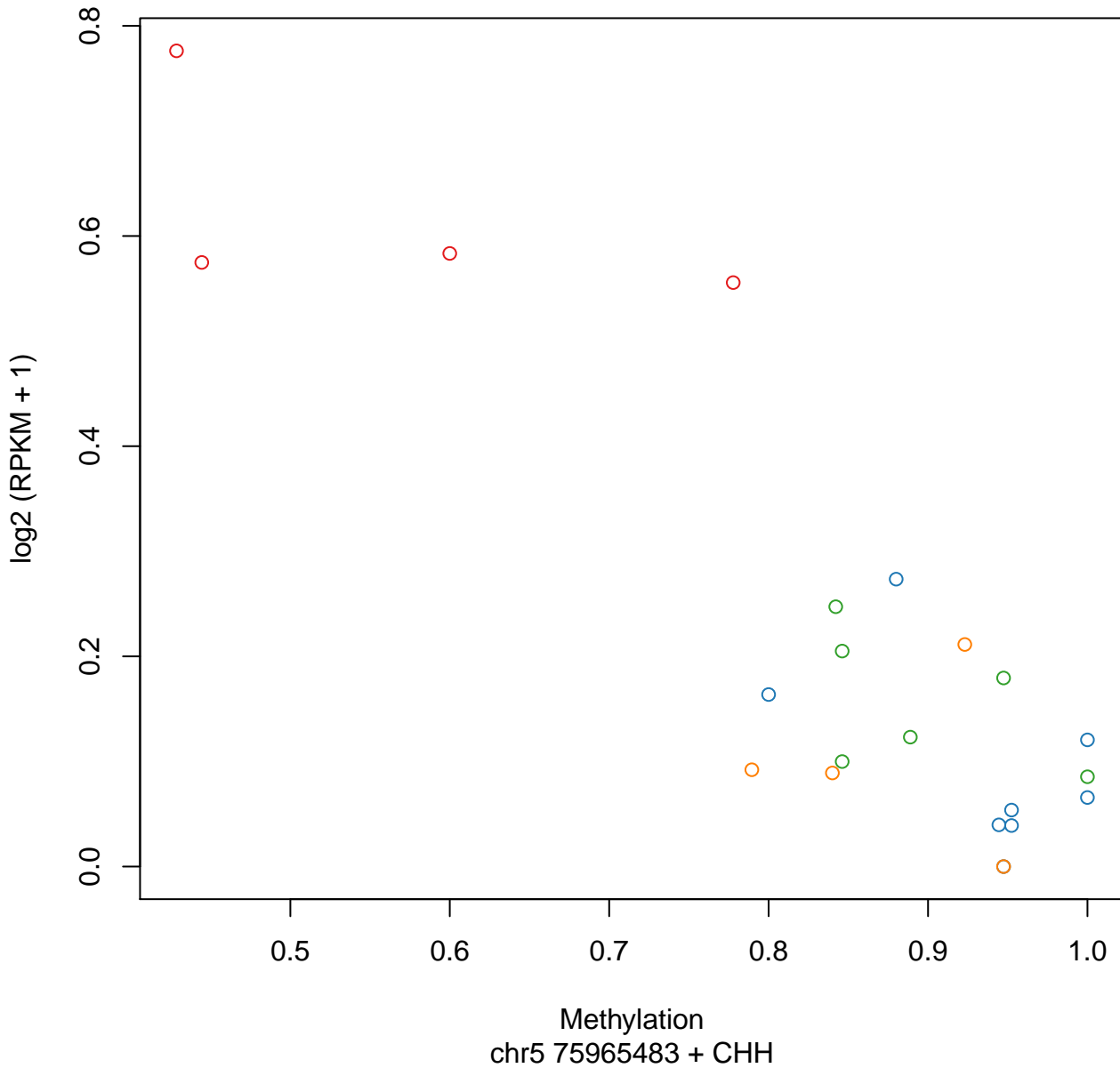
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TNC



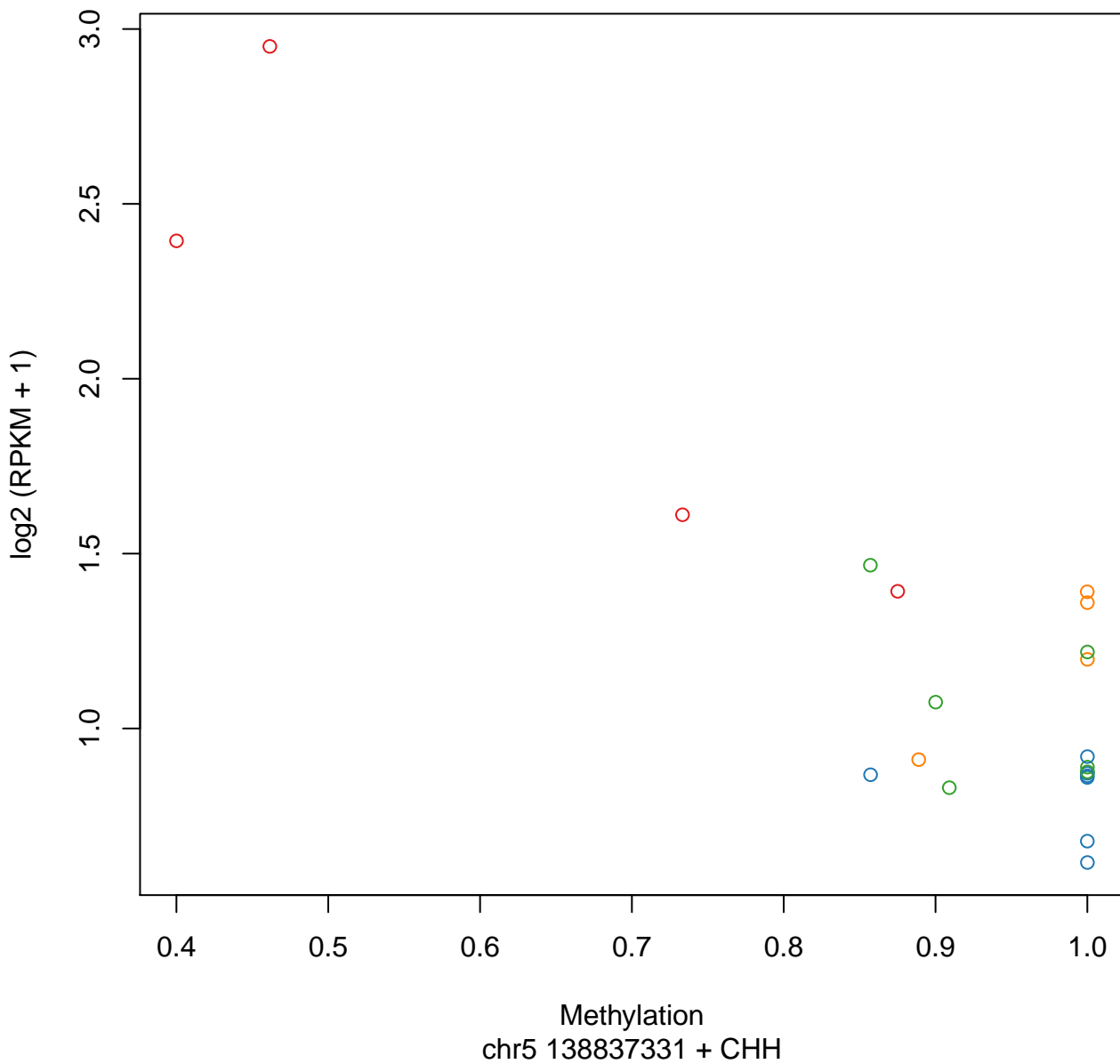
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IQGAP2

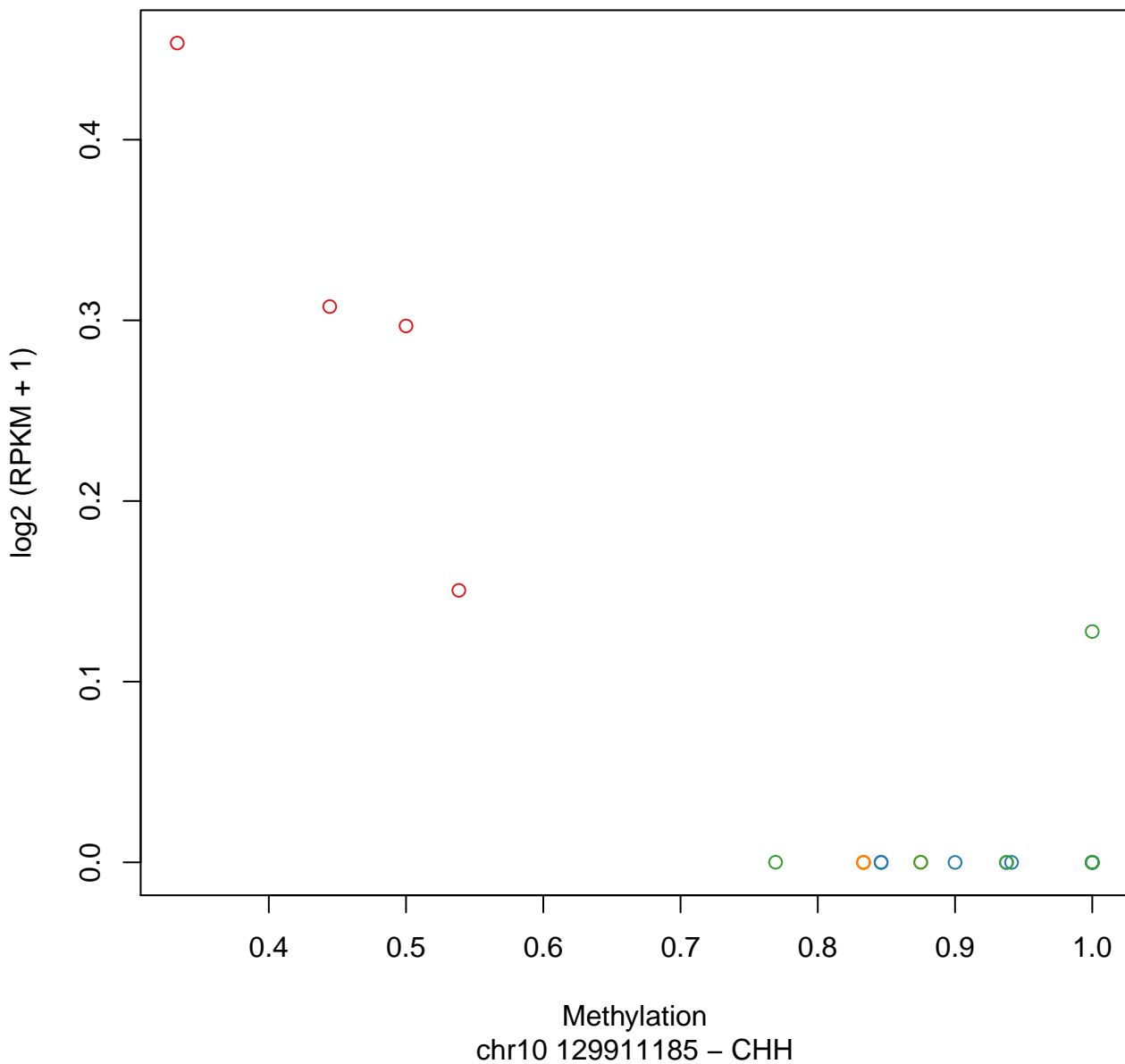


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ECSCR

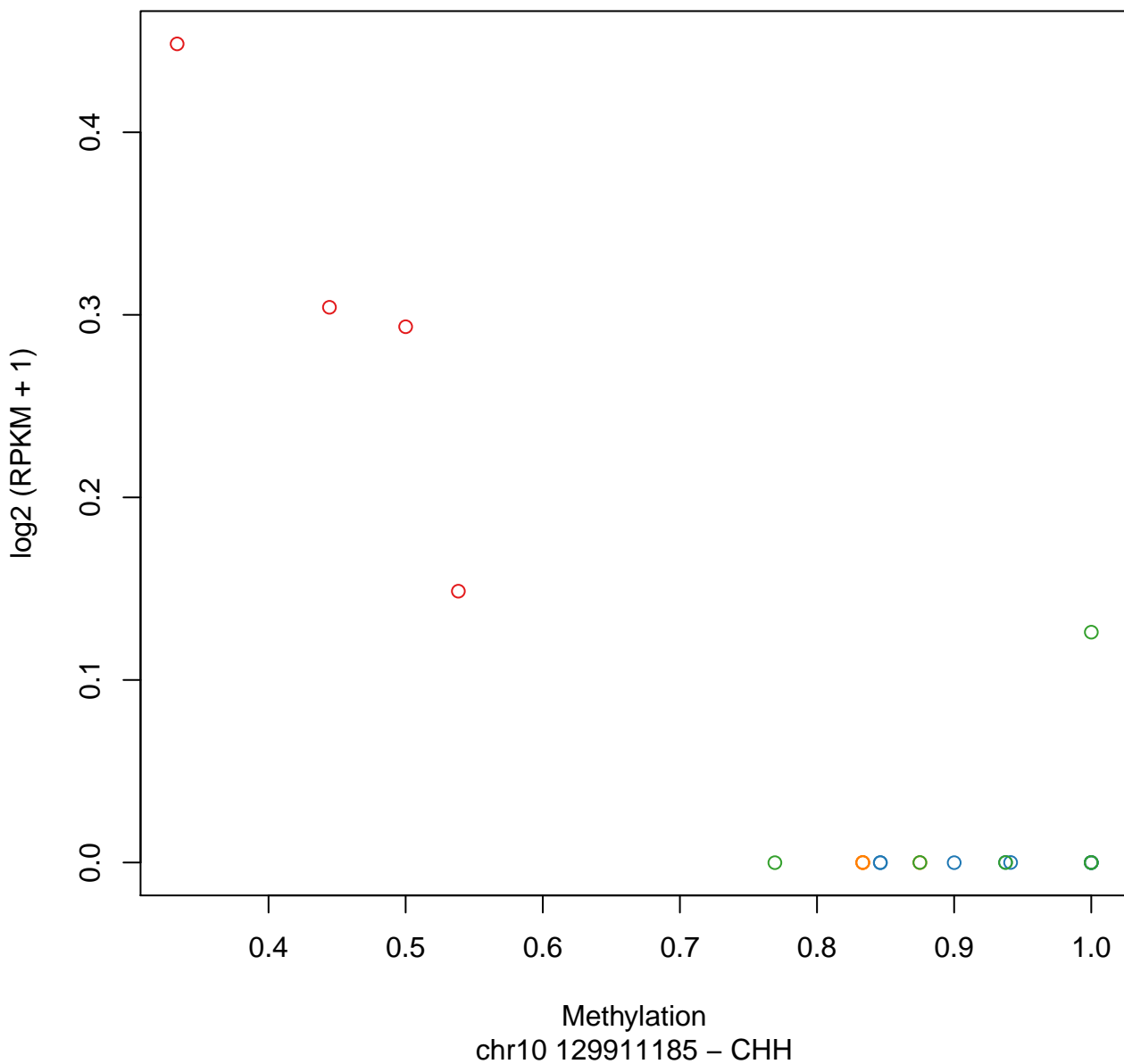


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MKI67



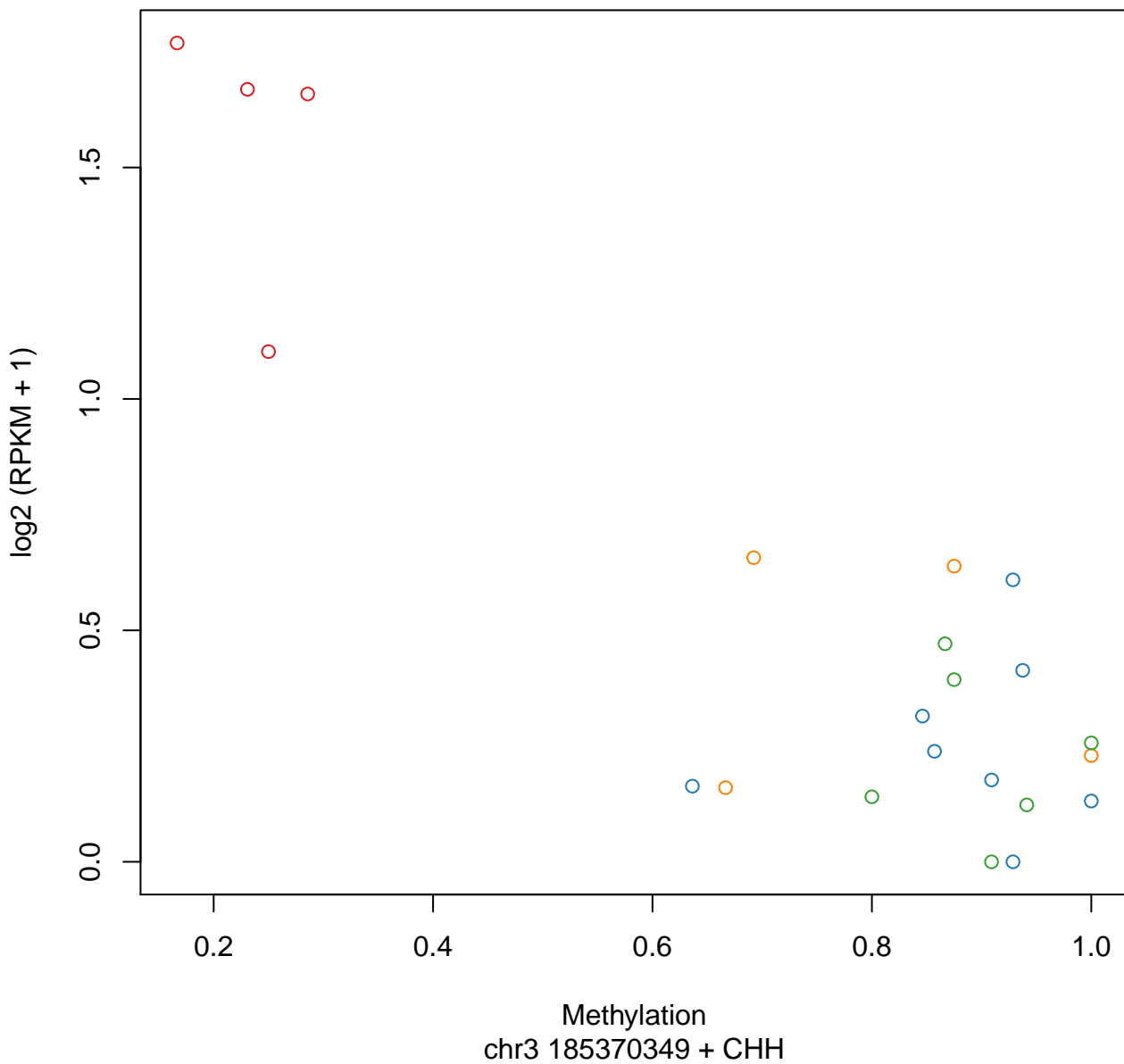
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MKI67



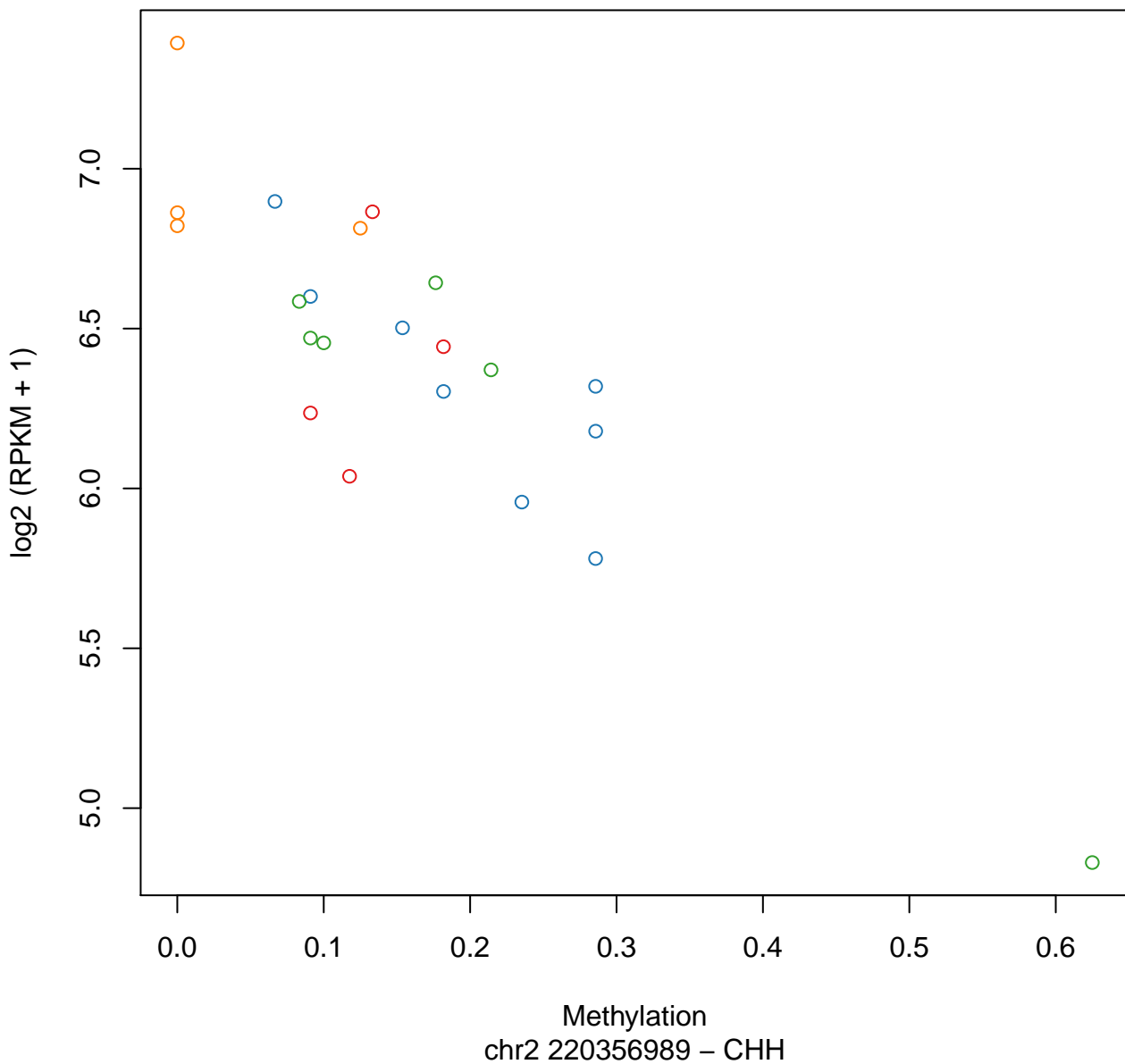
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IGF2BP2



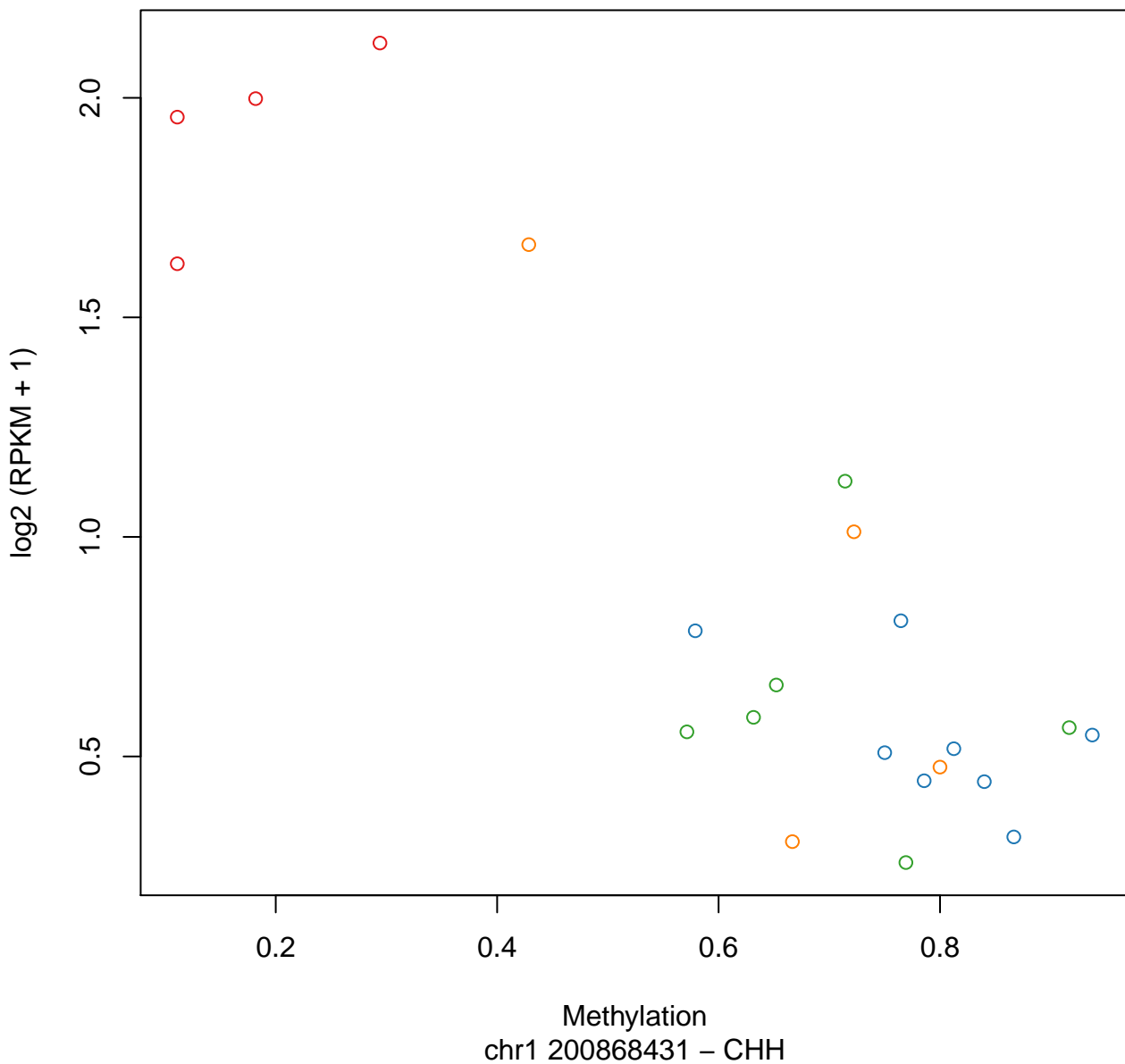
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SPEG

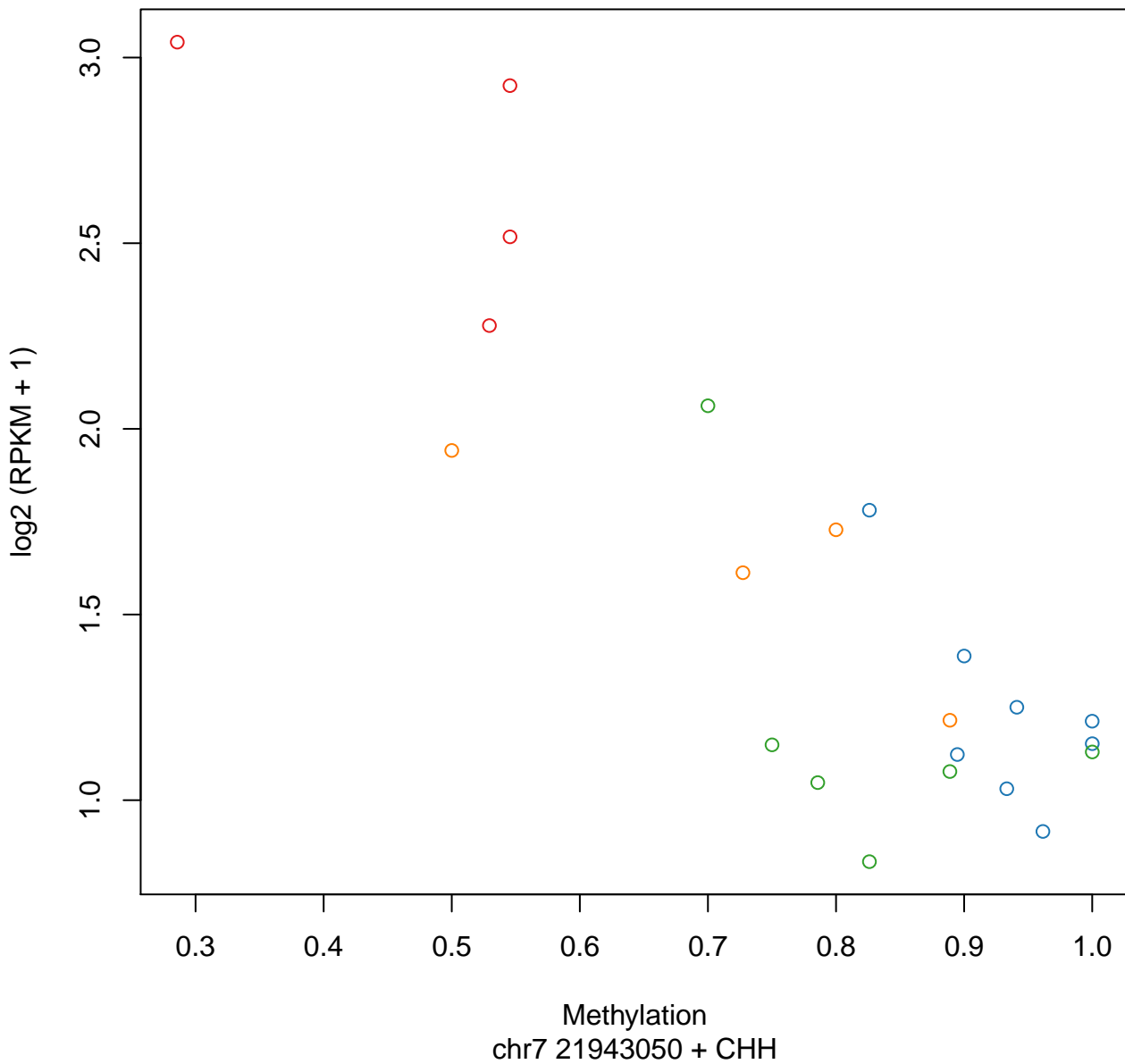


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C1orf106

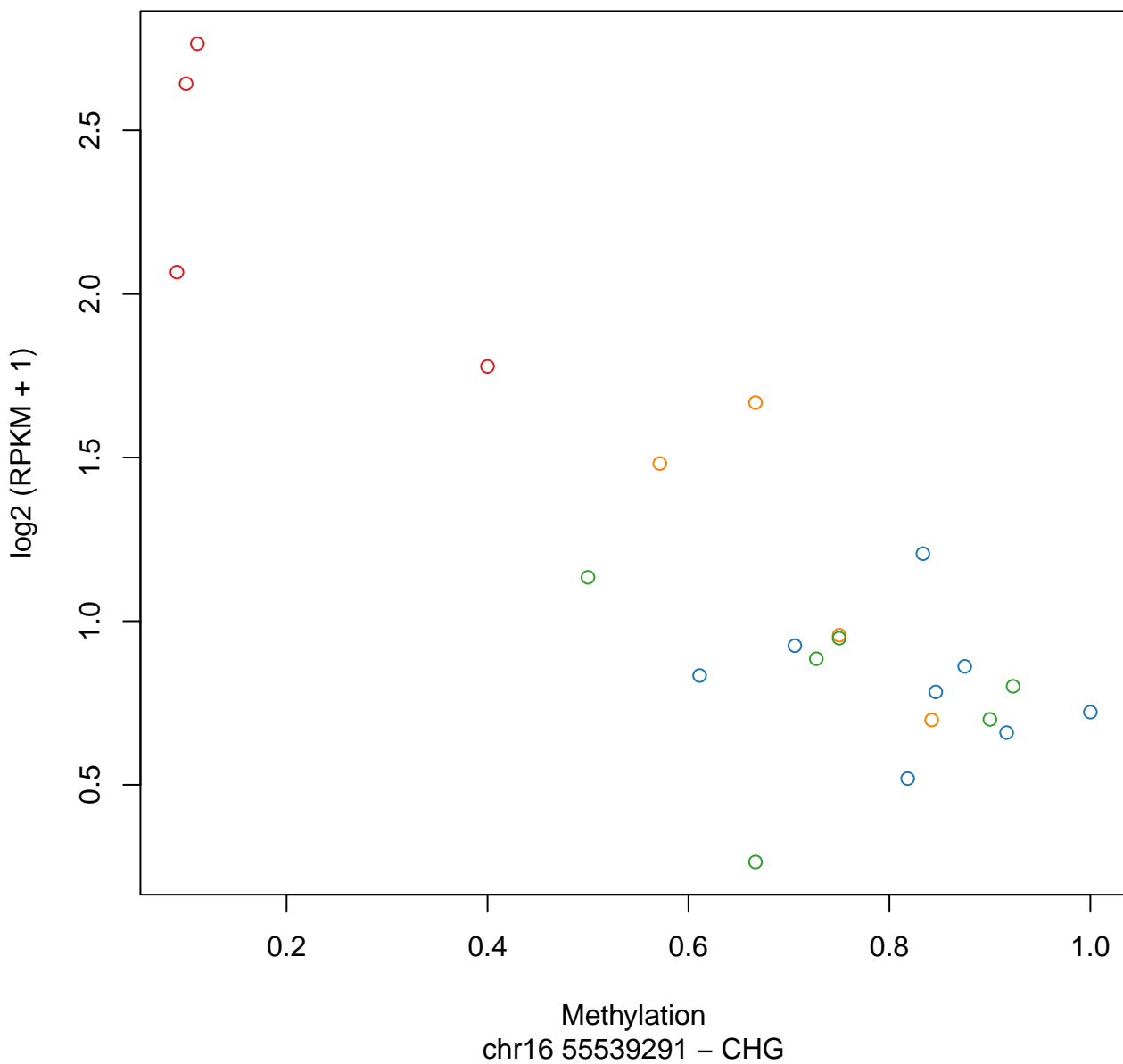


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CDCA7L



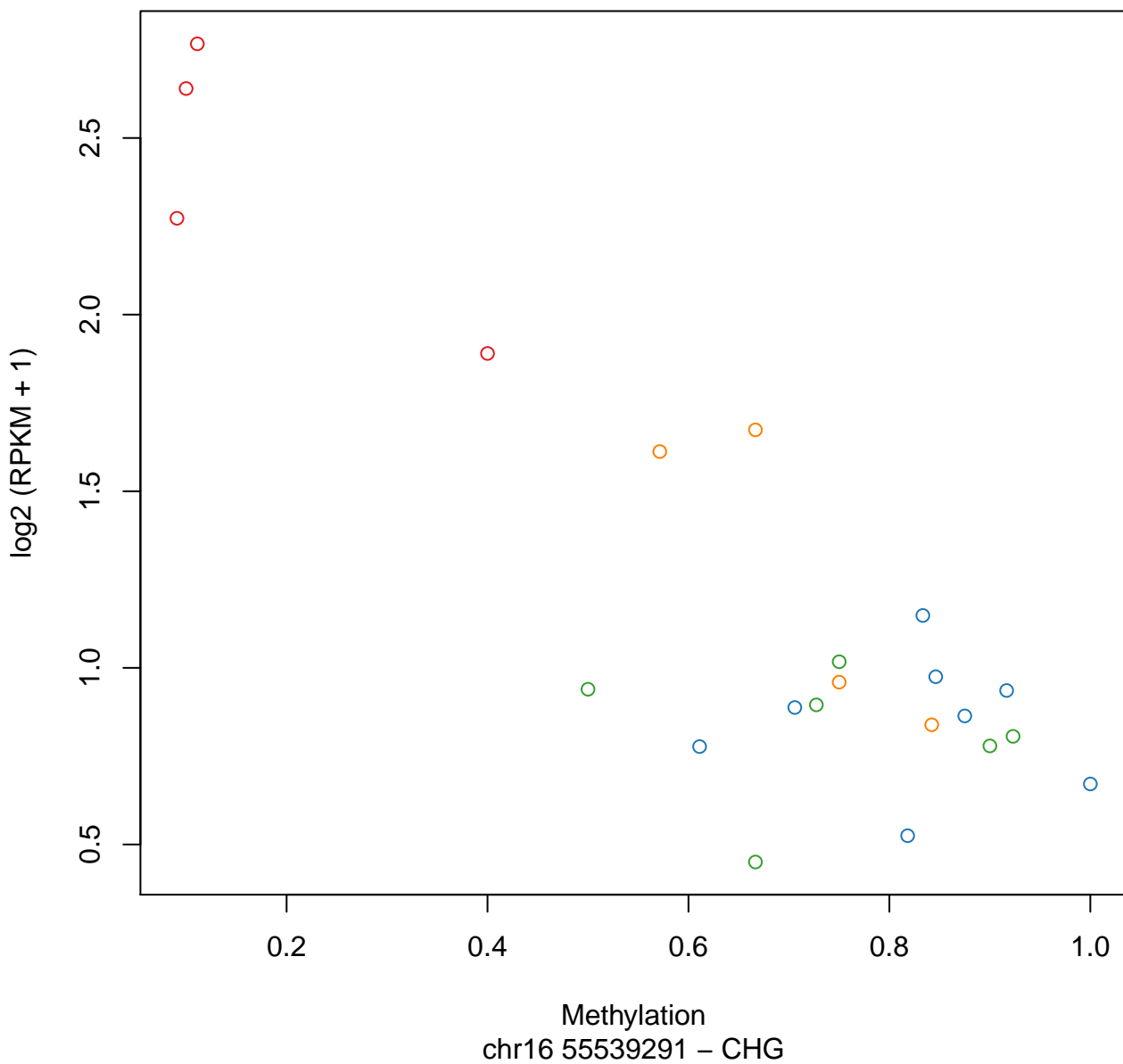
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MMP2



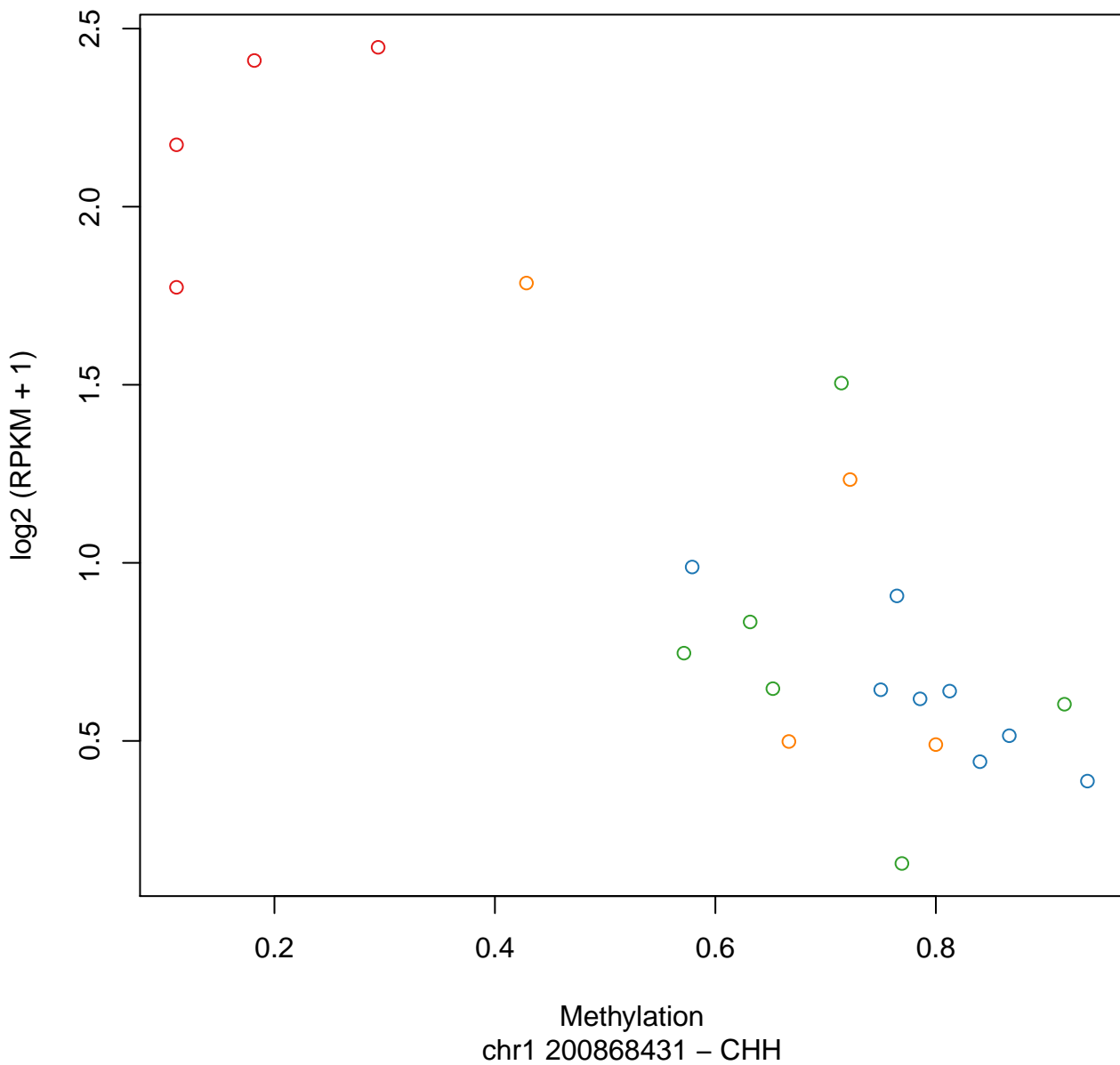
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MMP2



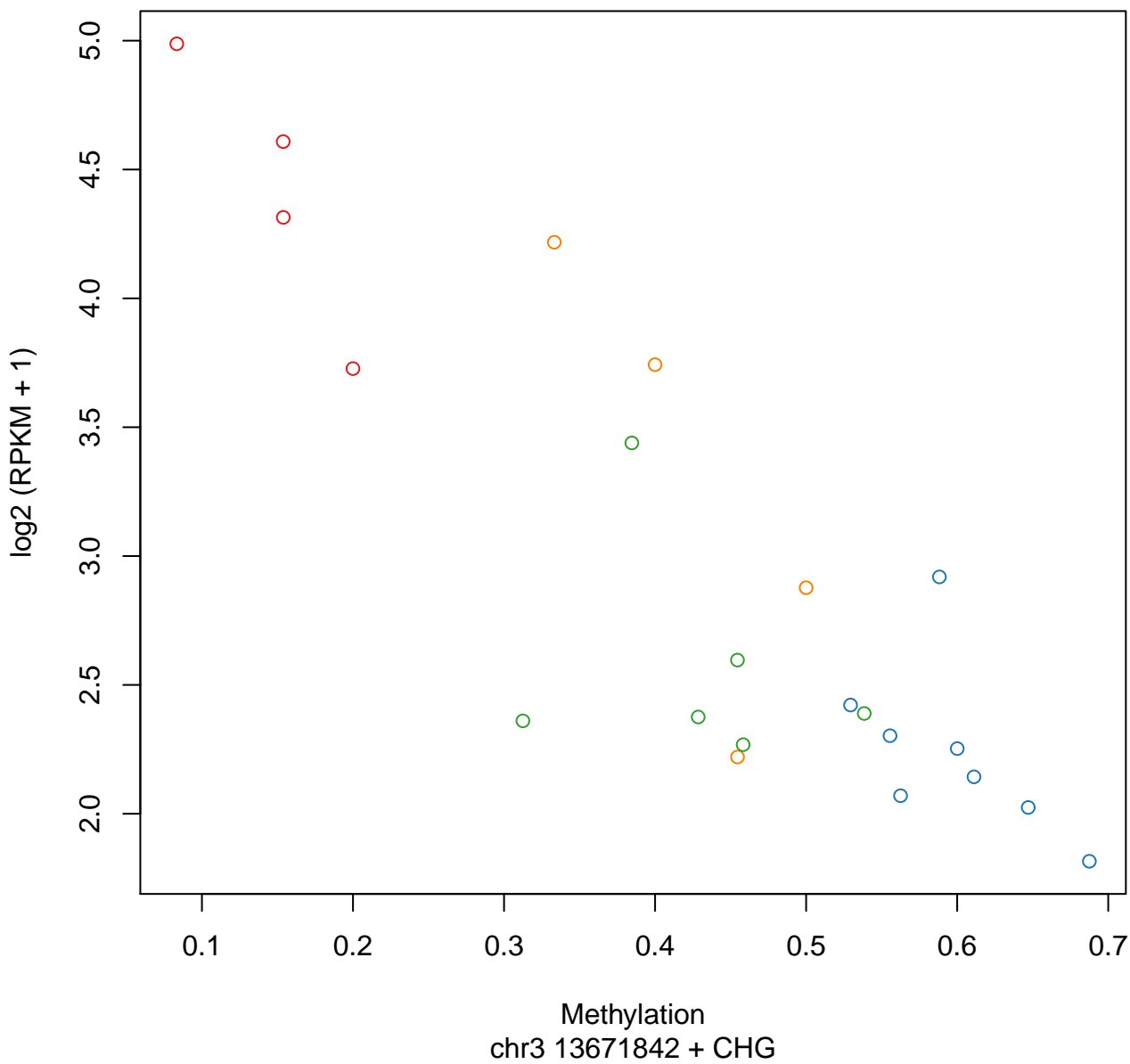
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C1orf106

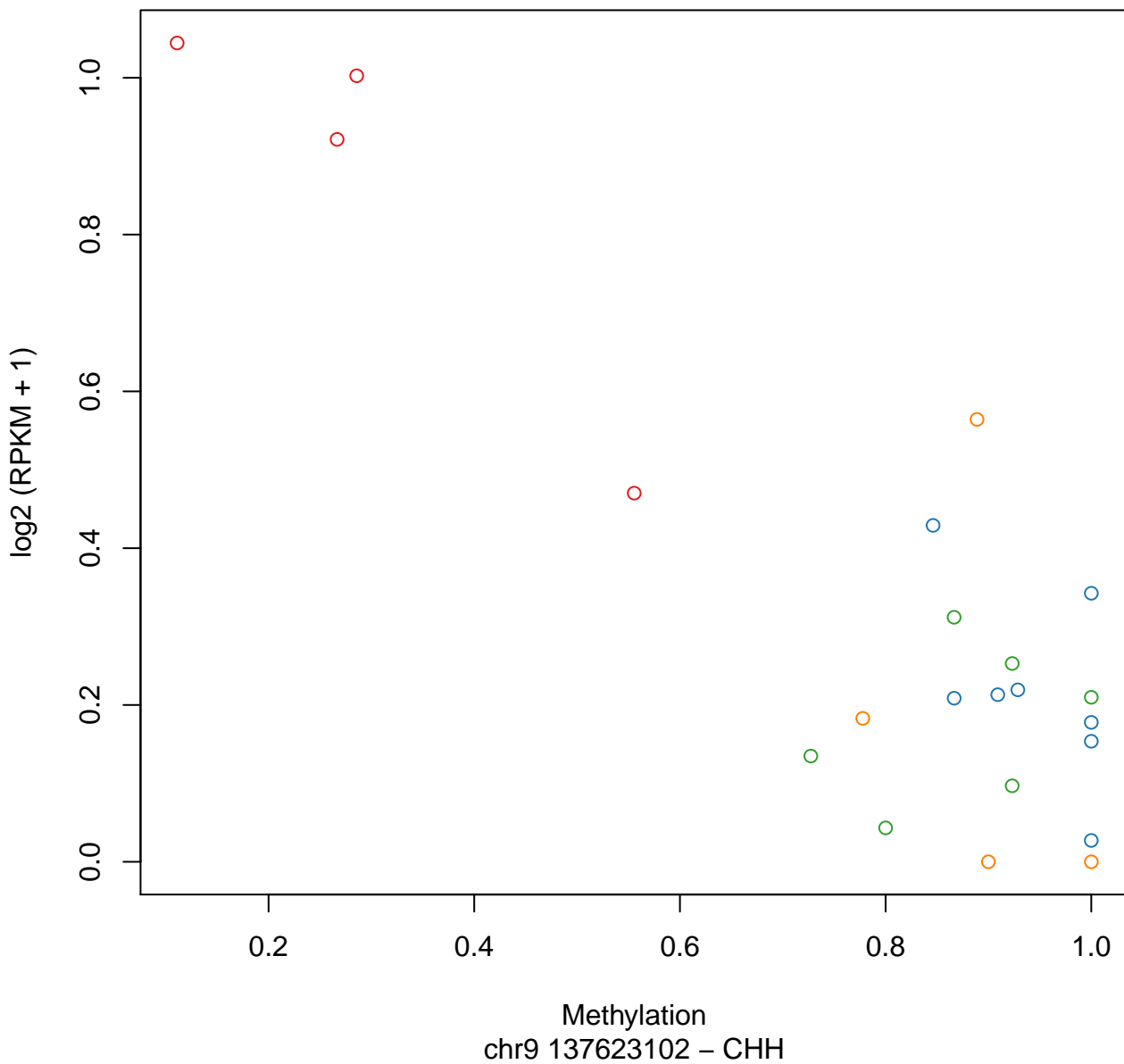


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FBLN2

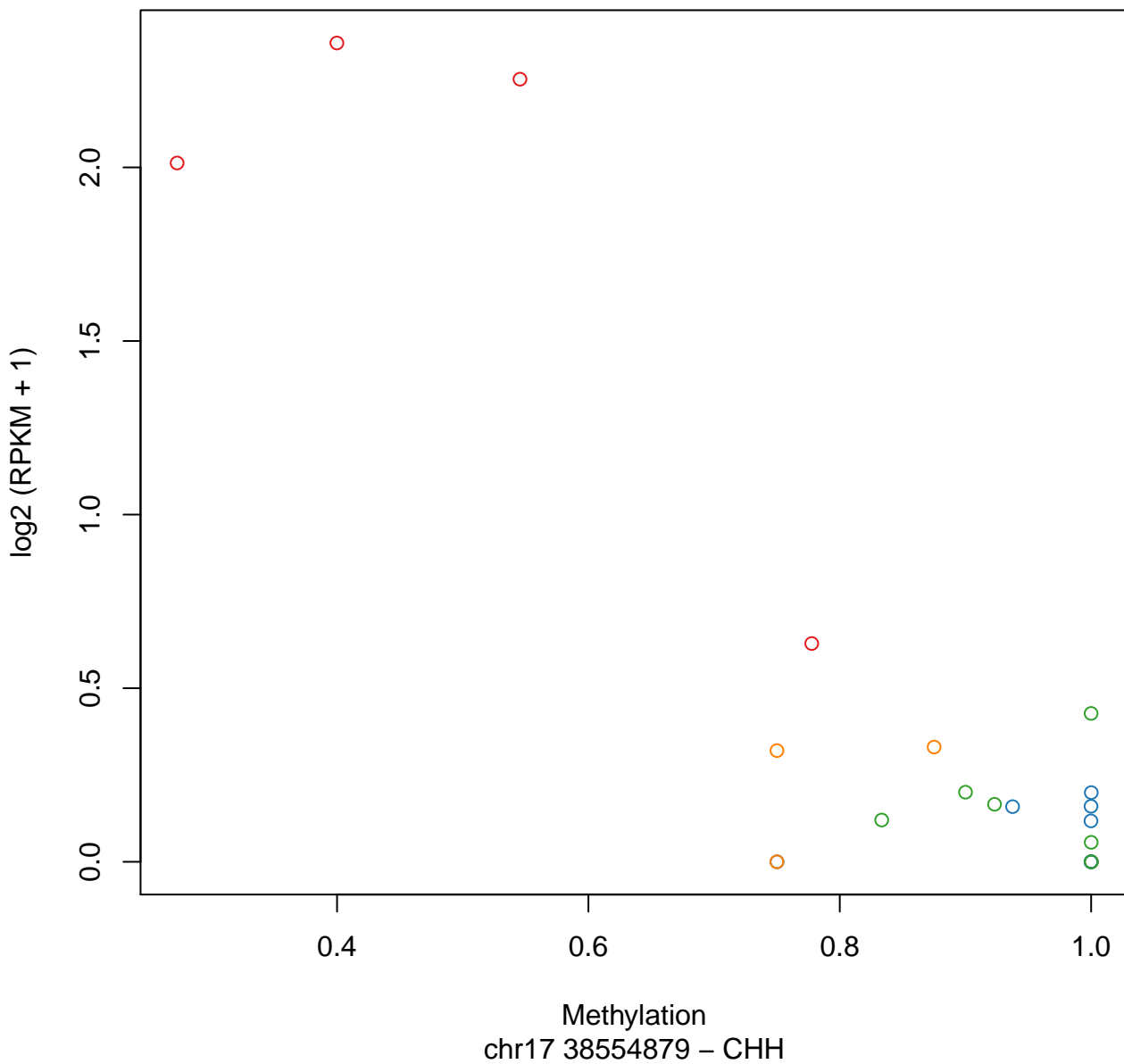


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COL5A1



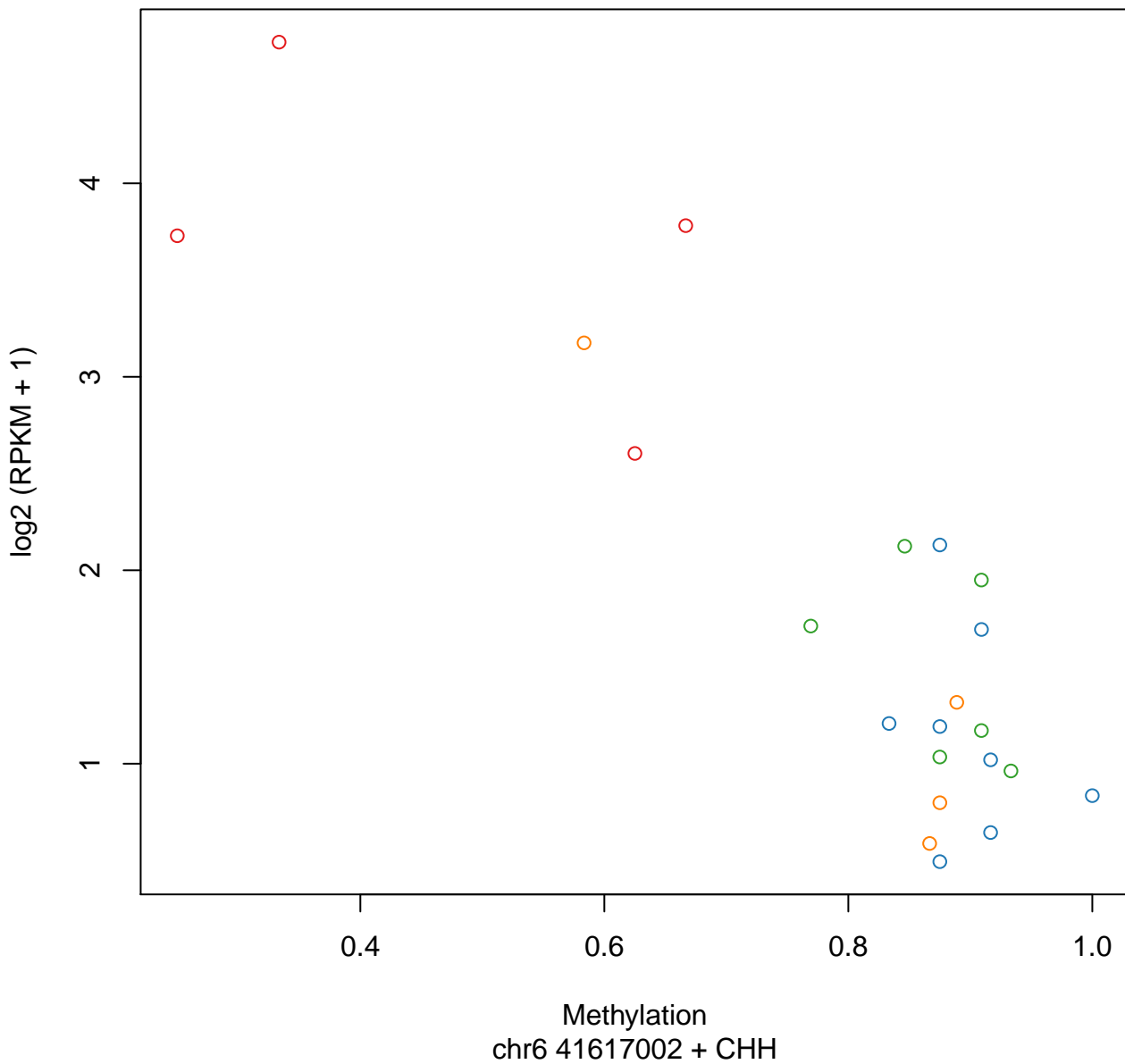
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TOP2A



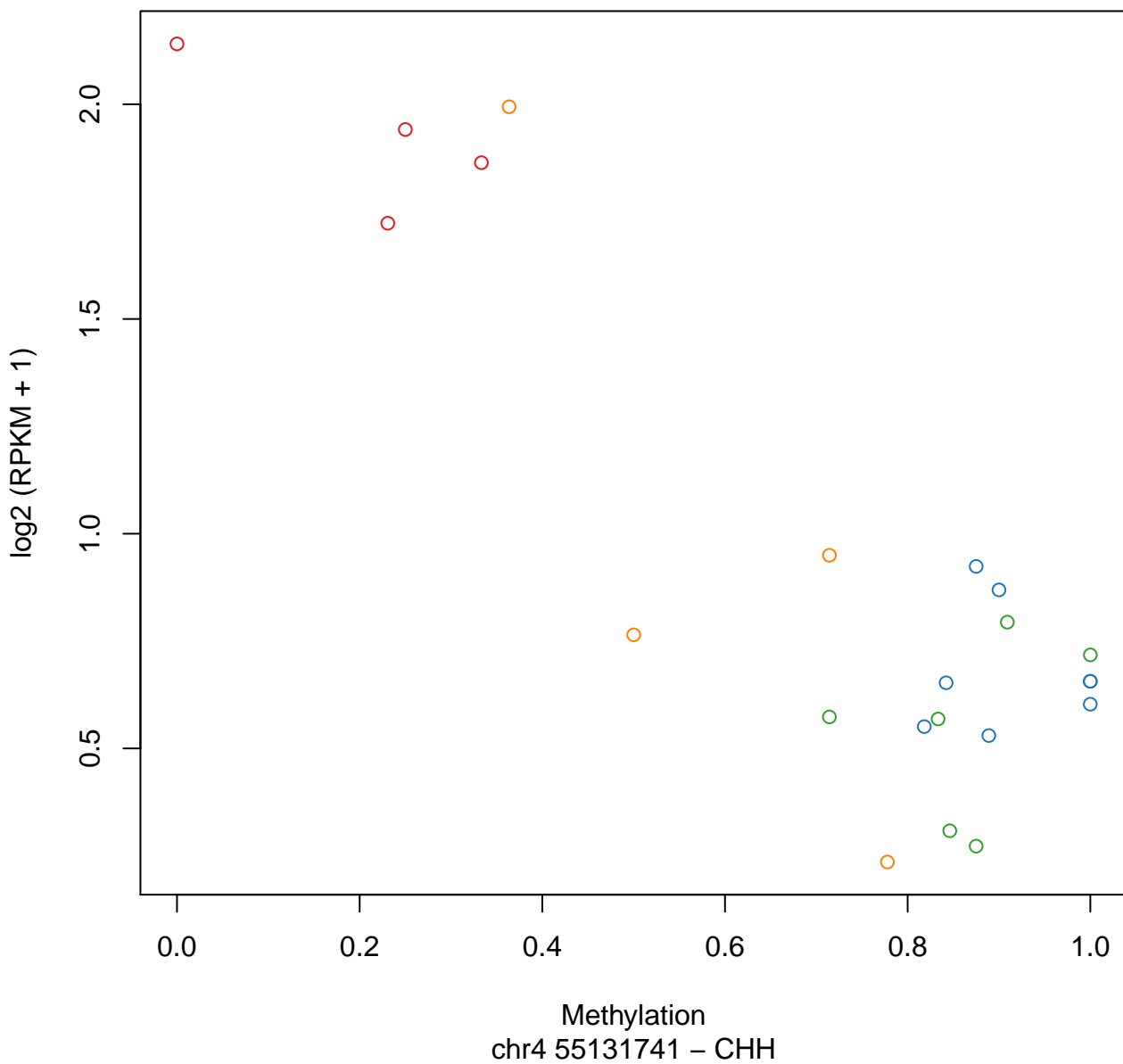
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MDFI



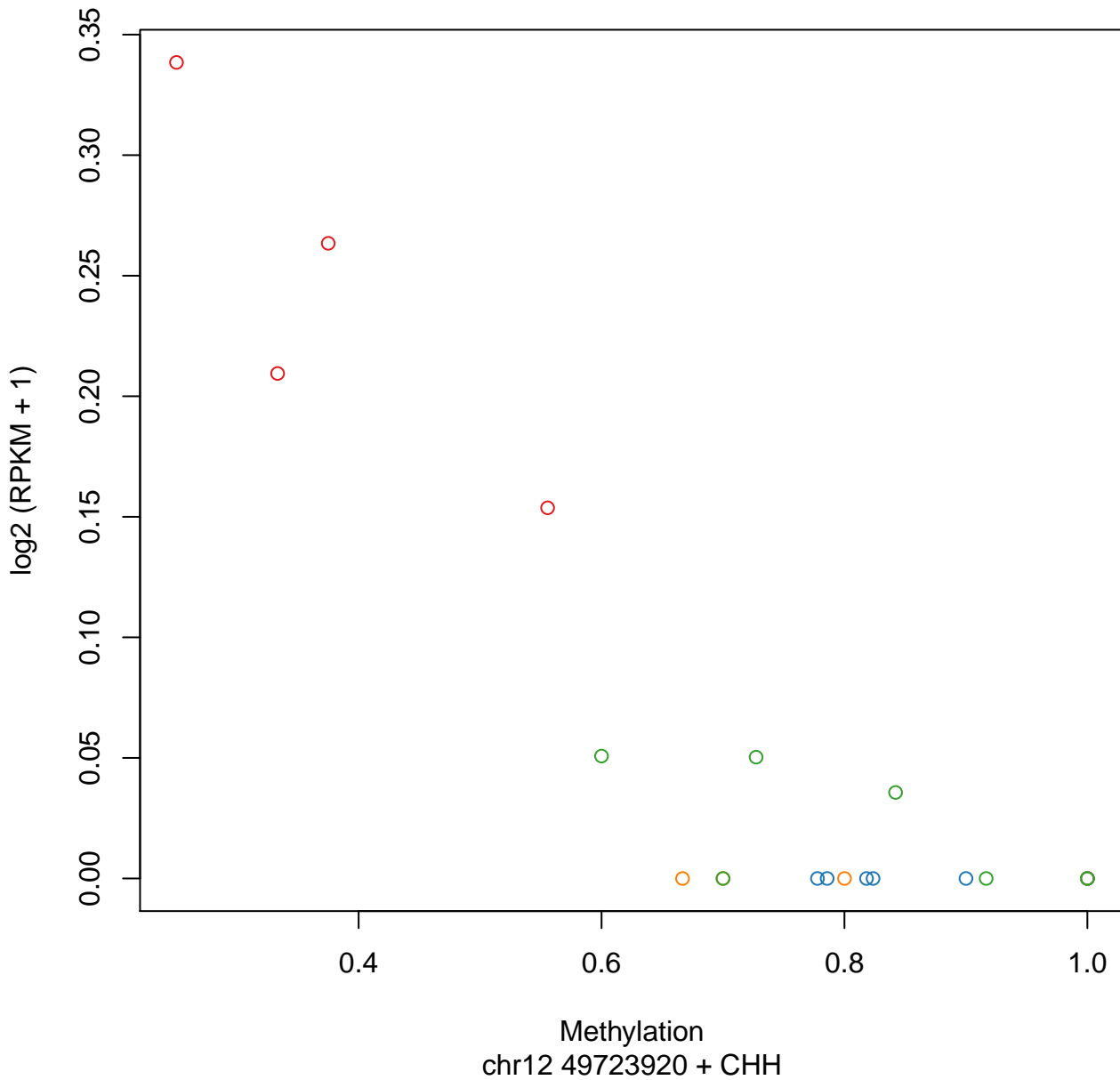
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PDGFRA



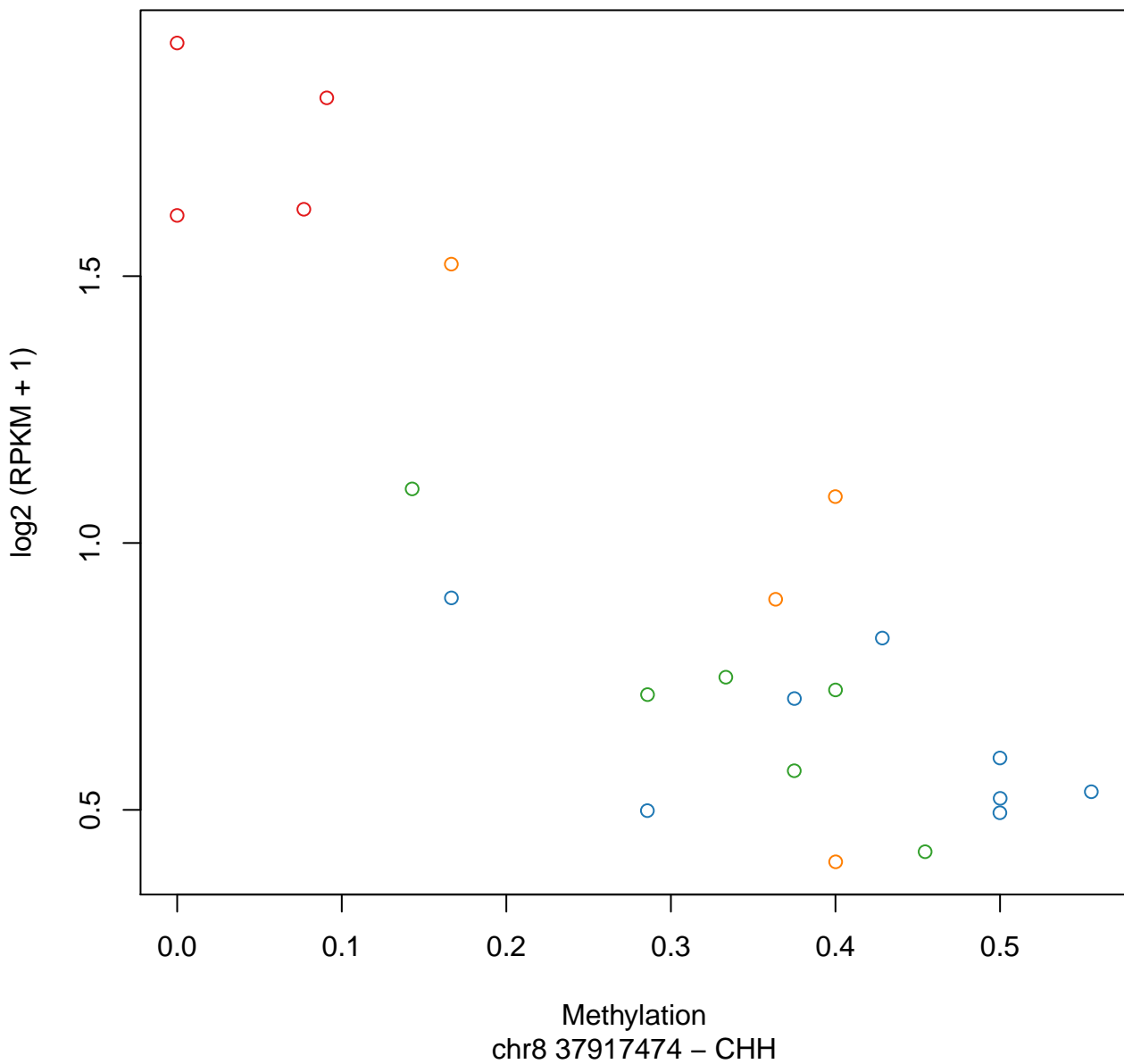
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TROAP



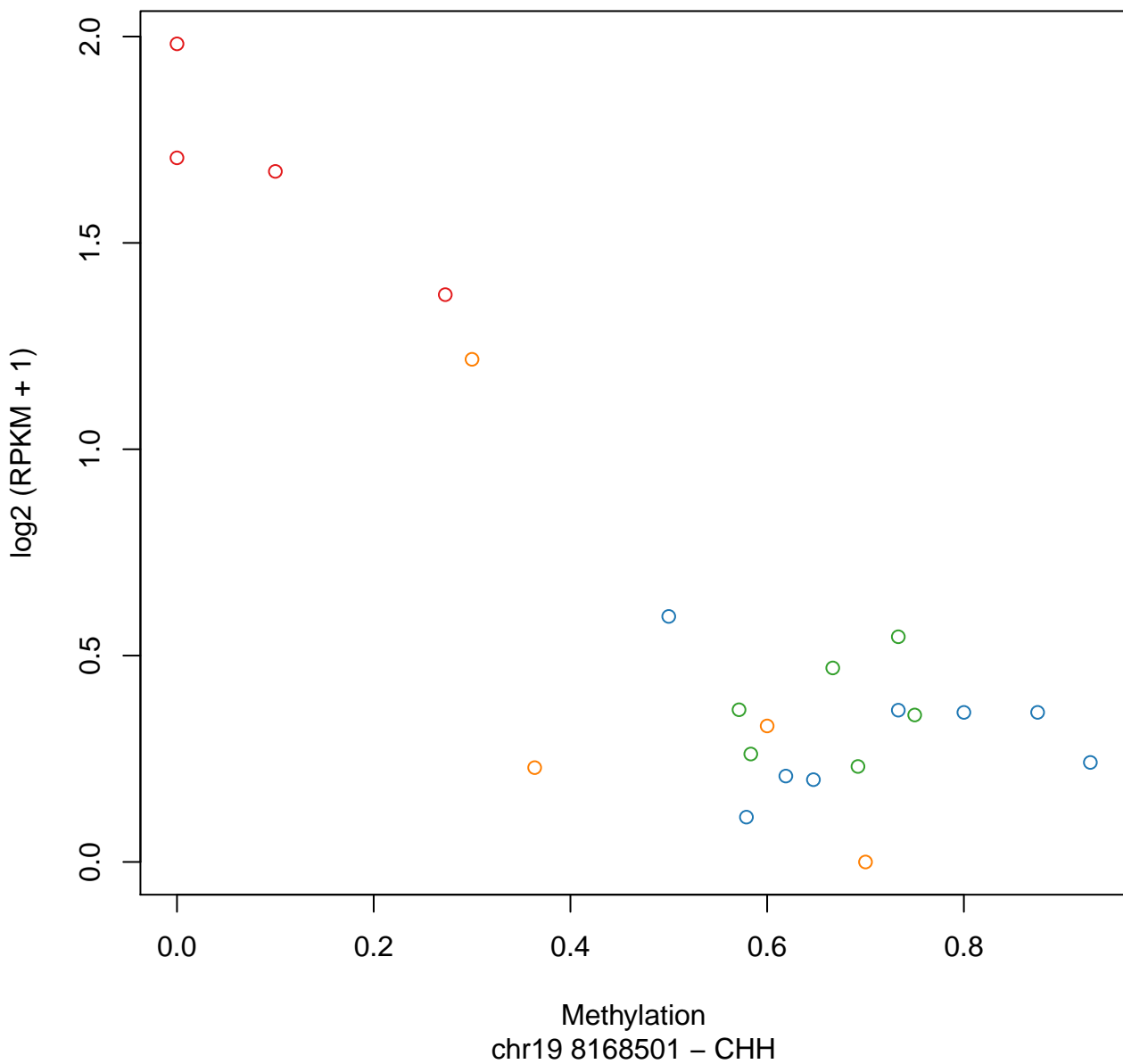
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EIF4EBP1

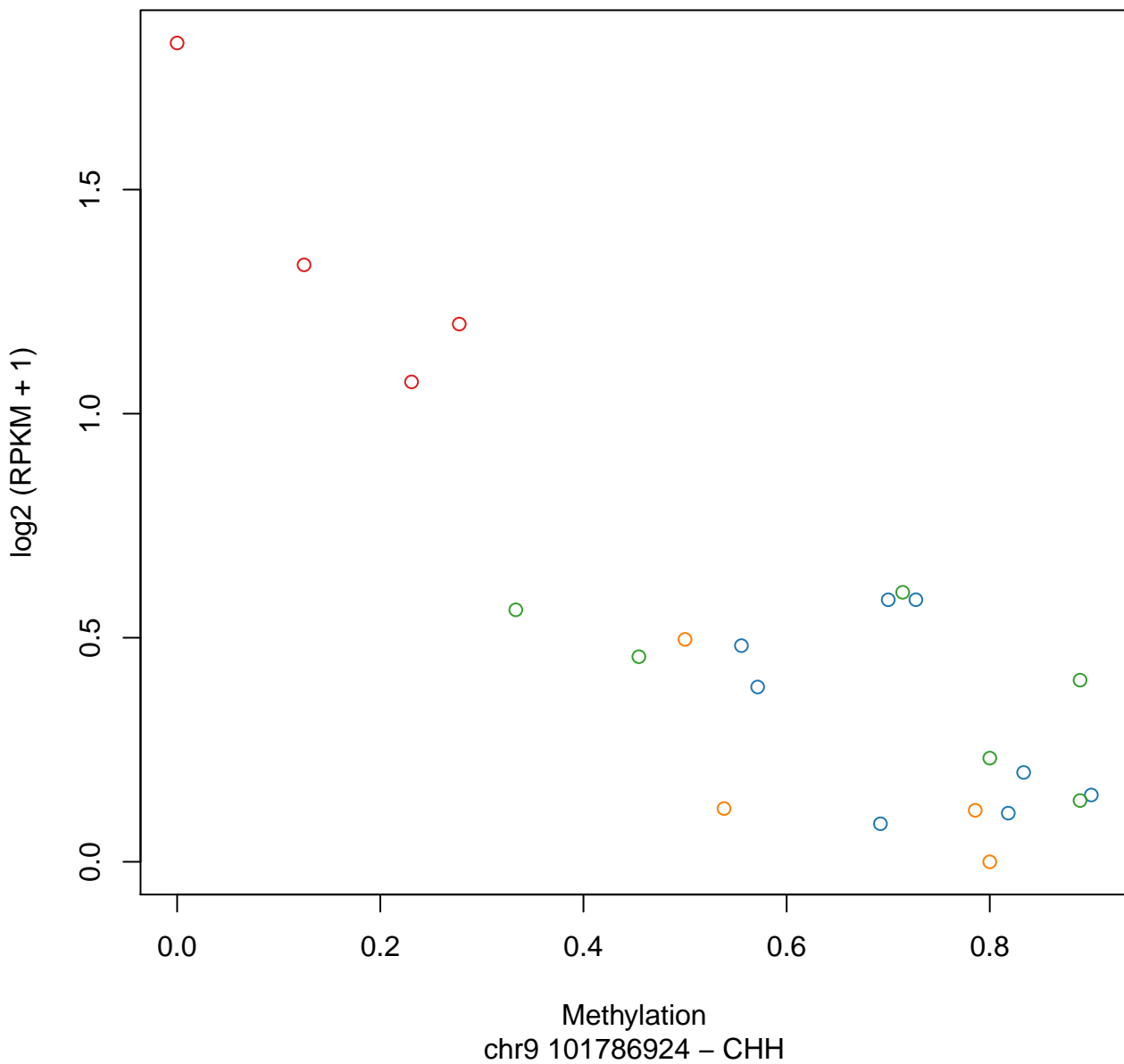


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FBN3

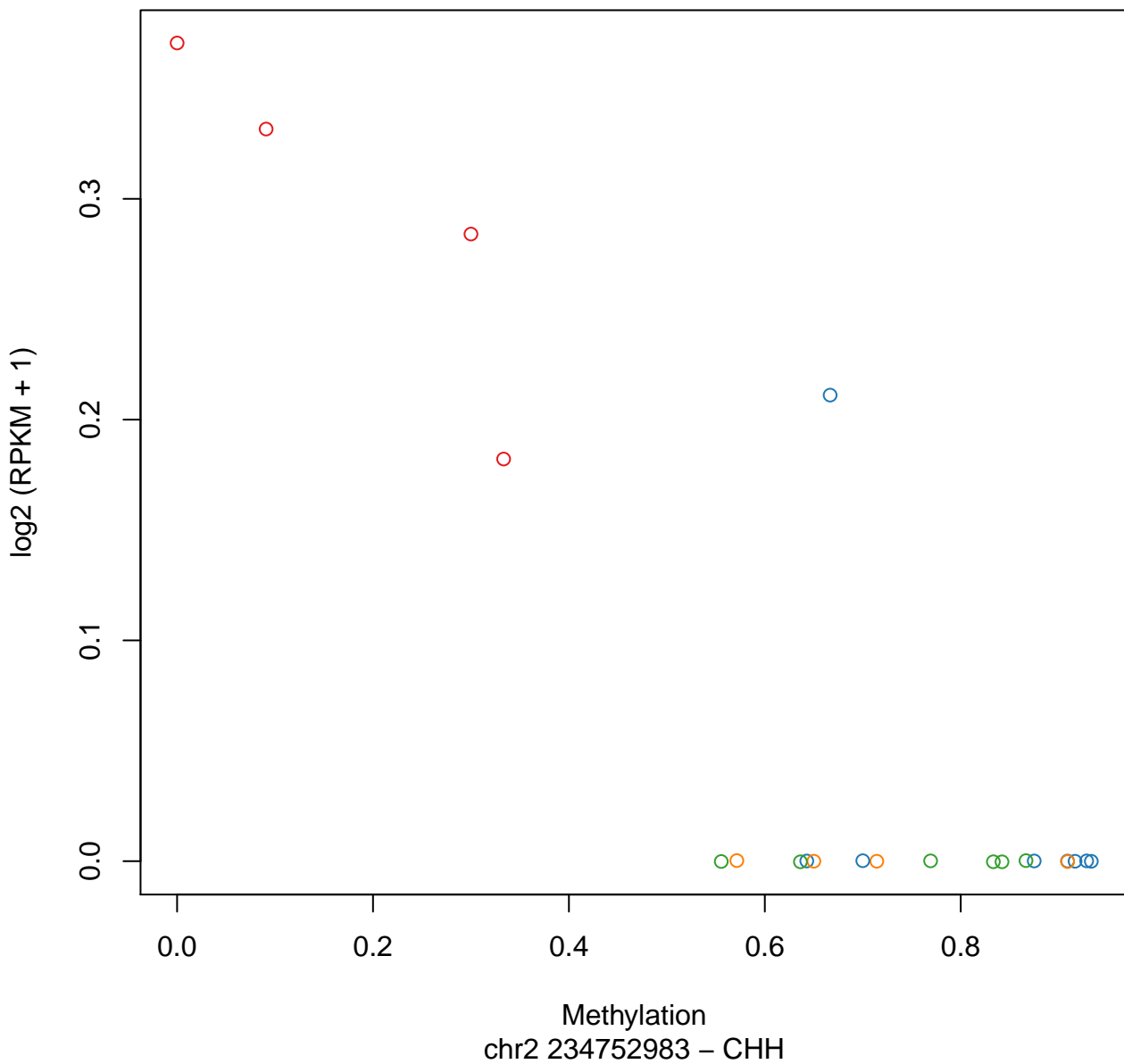


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COL15A1



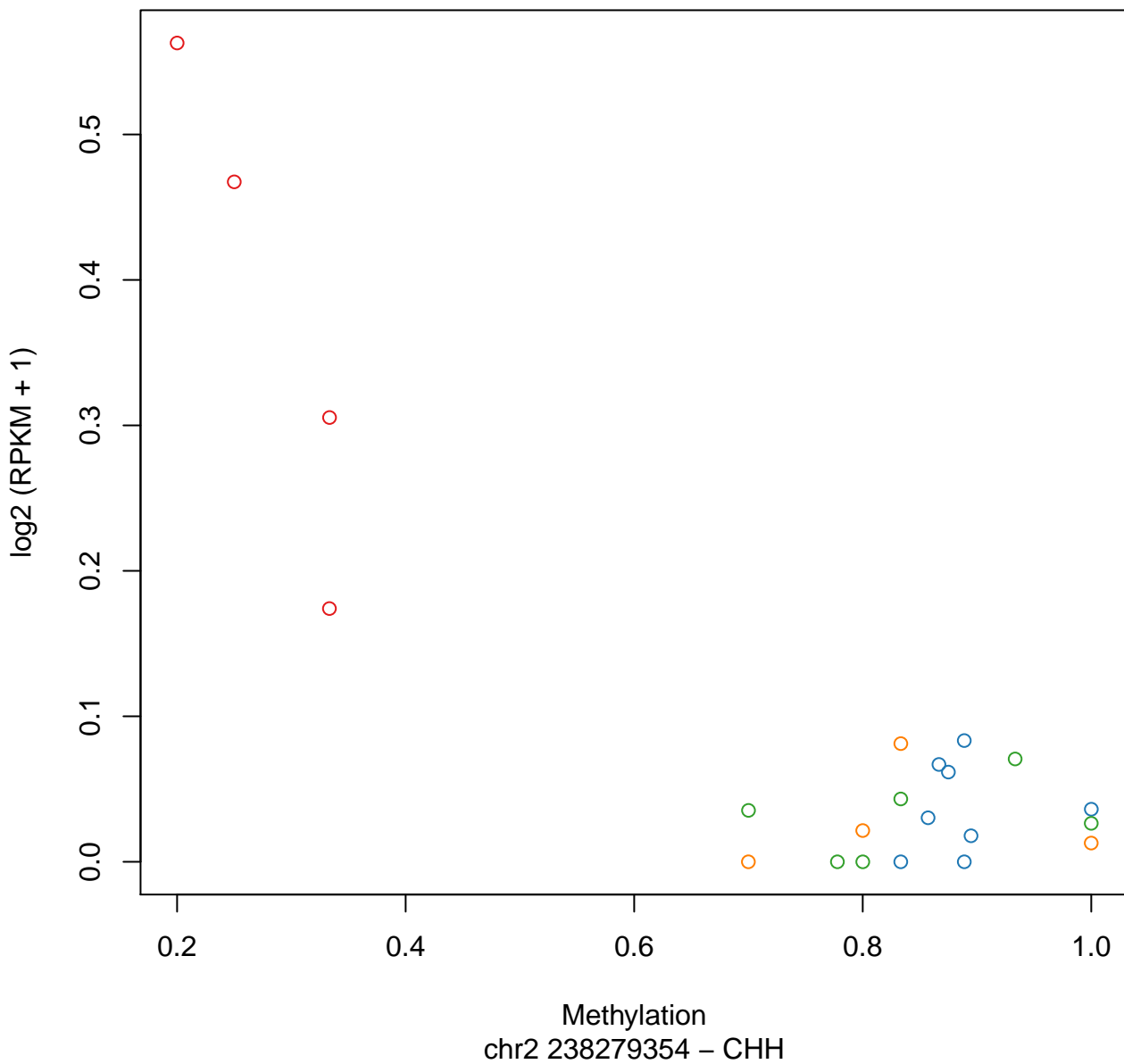
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HJURP



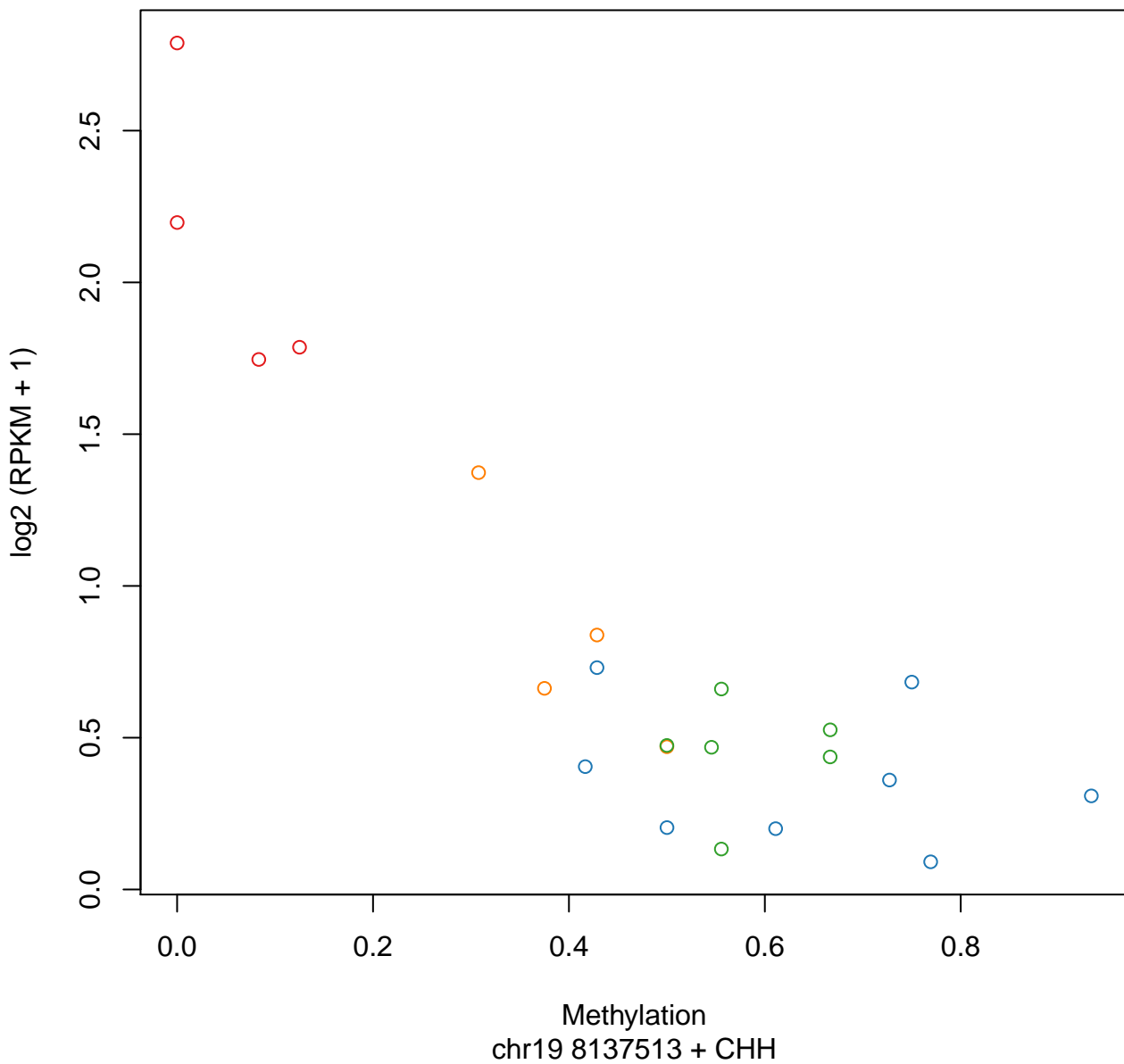
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COL6A3

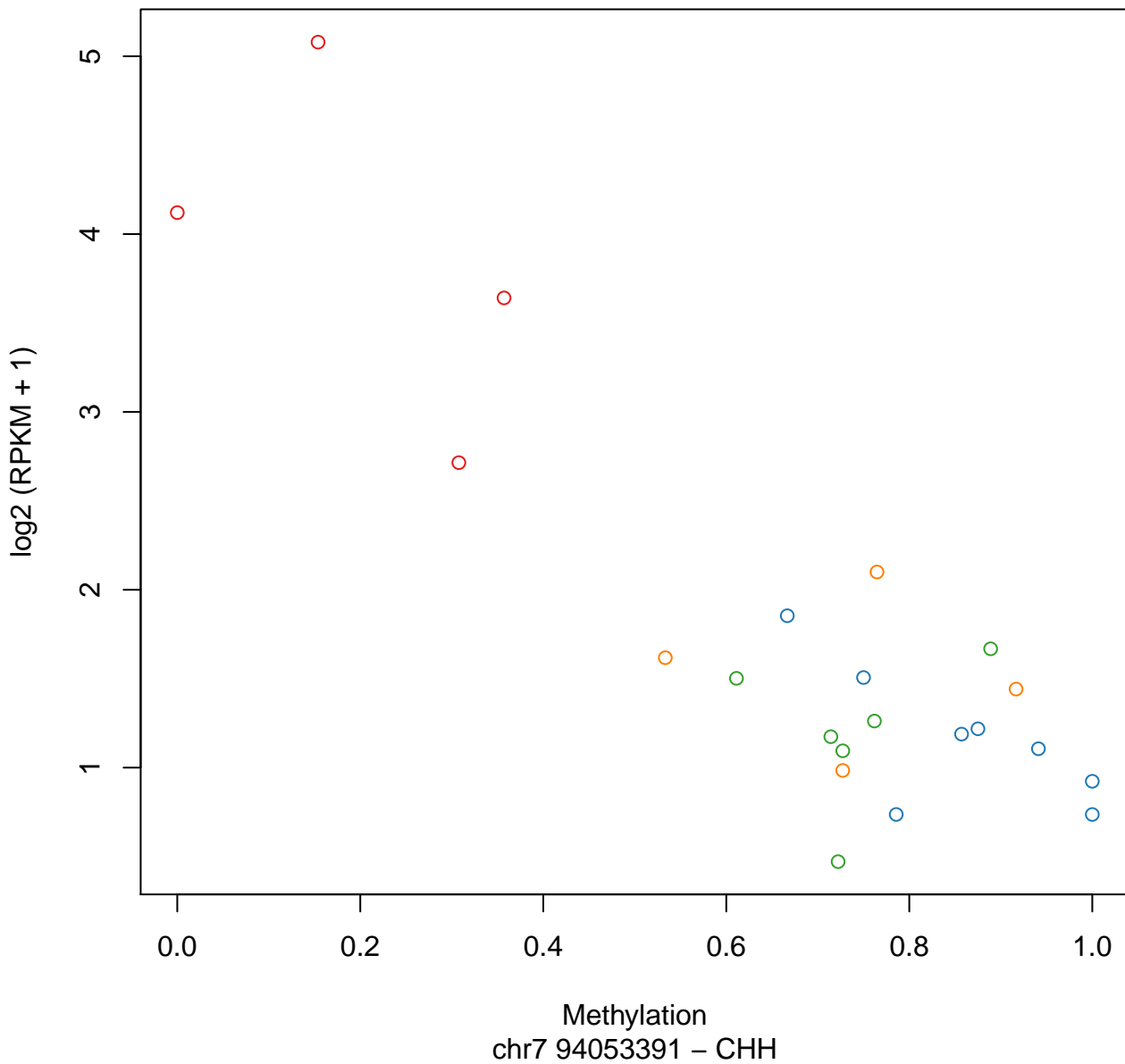


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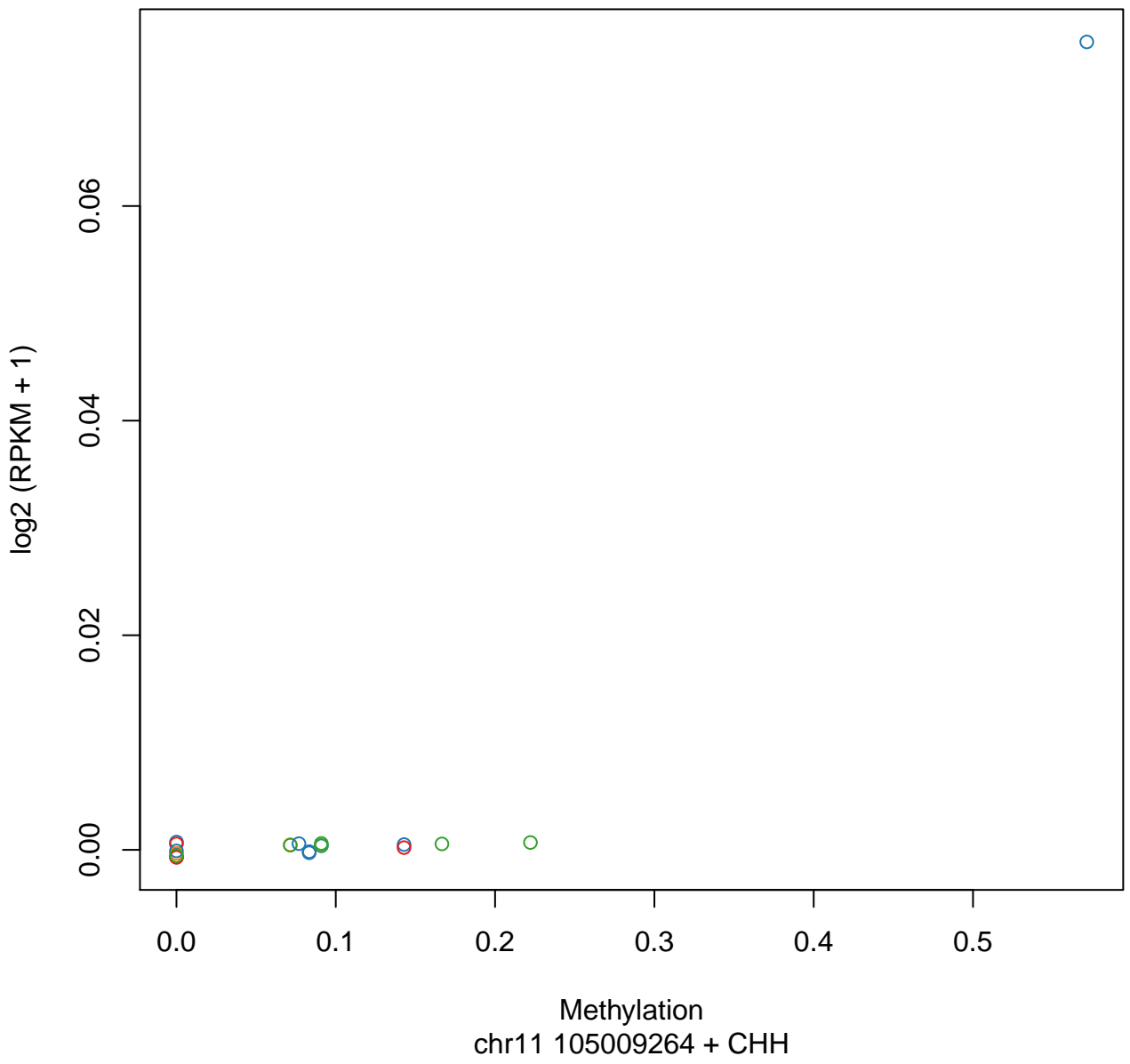
FBN3



exon ENSE00003462782.1_1 FDR 0.000962
COL1A2

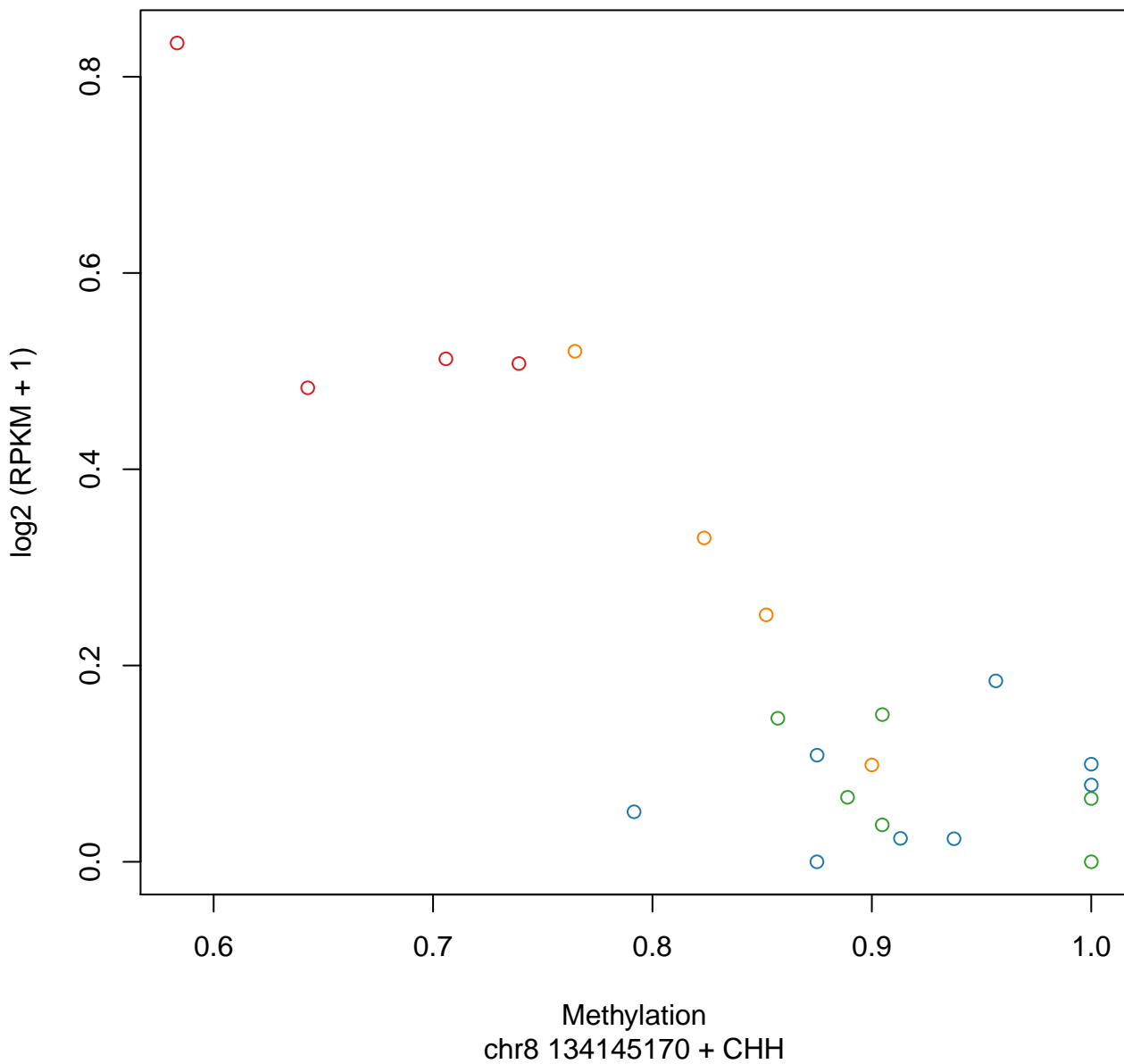


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CARD18



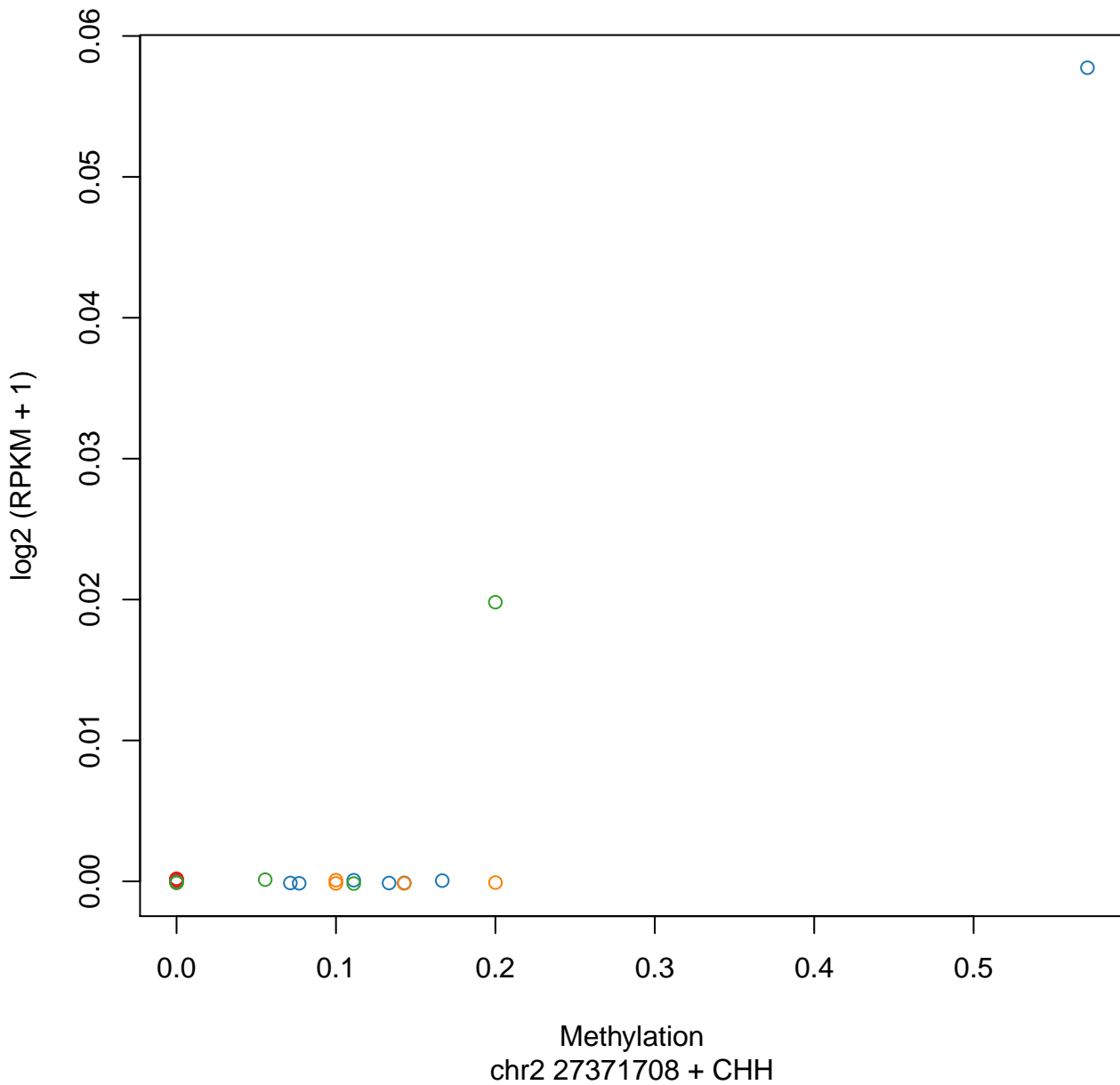
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TG

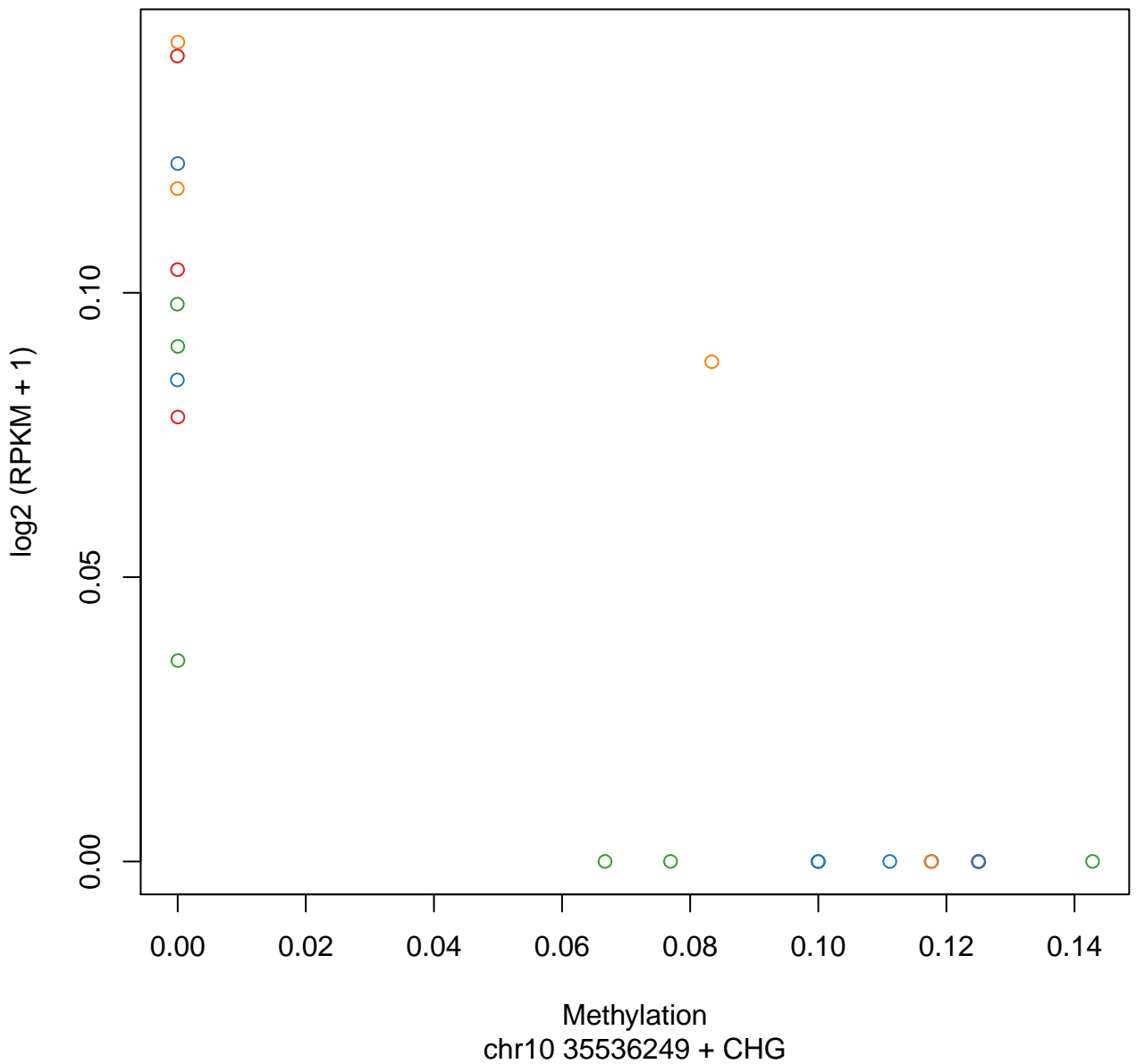


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TCF23

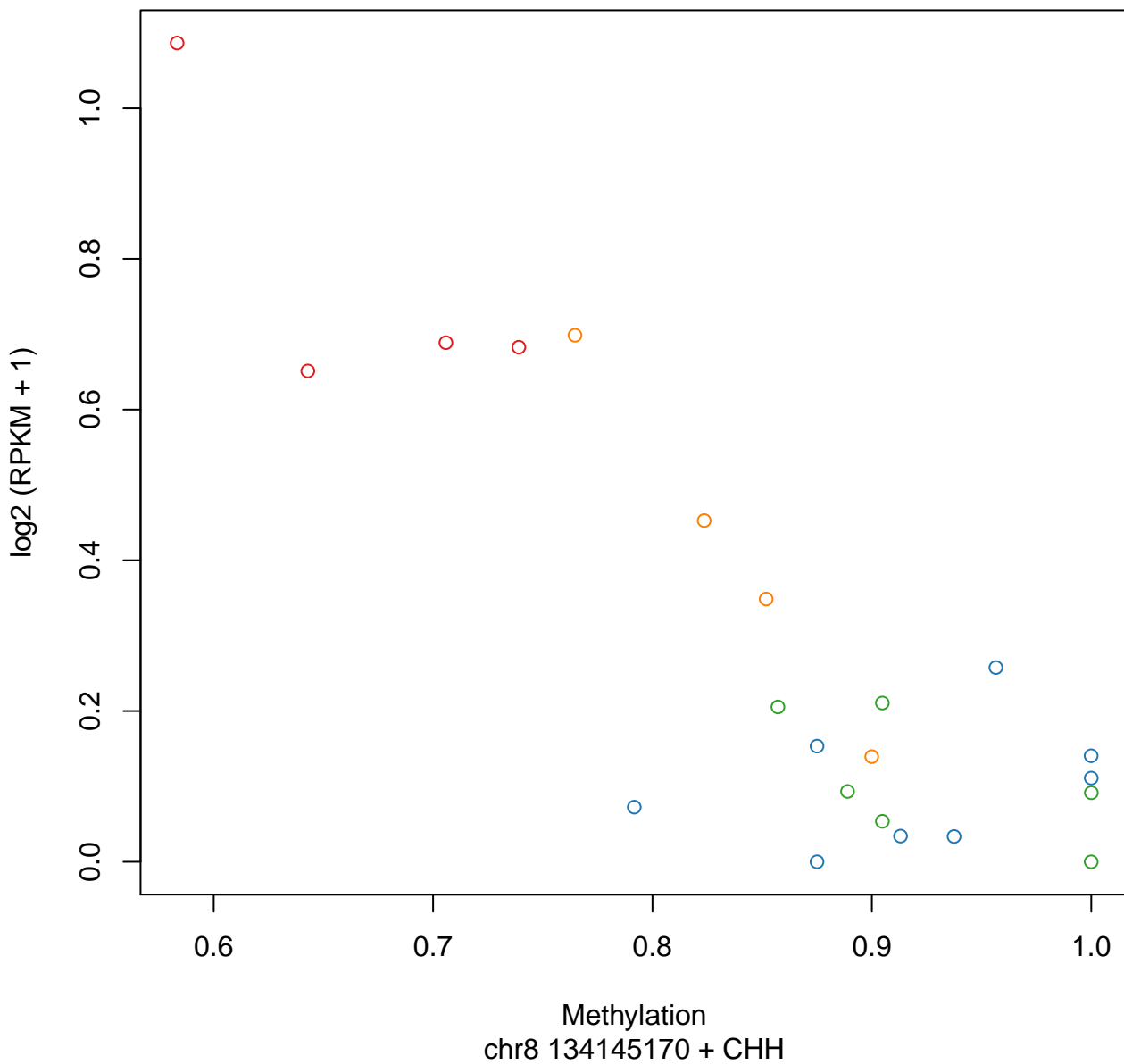


exon ENSE00001464397.1_1 FDR 0.00118
CCNY

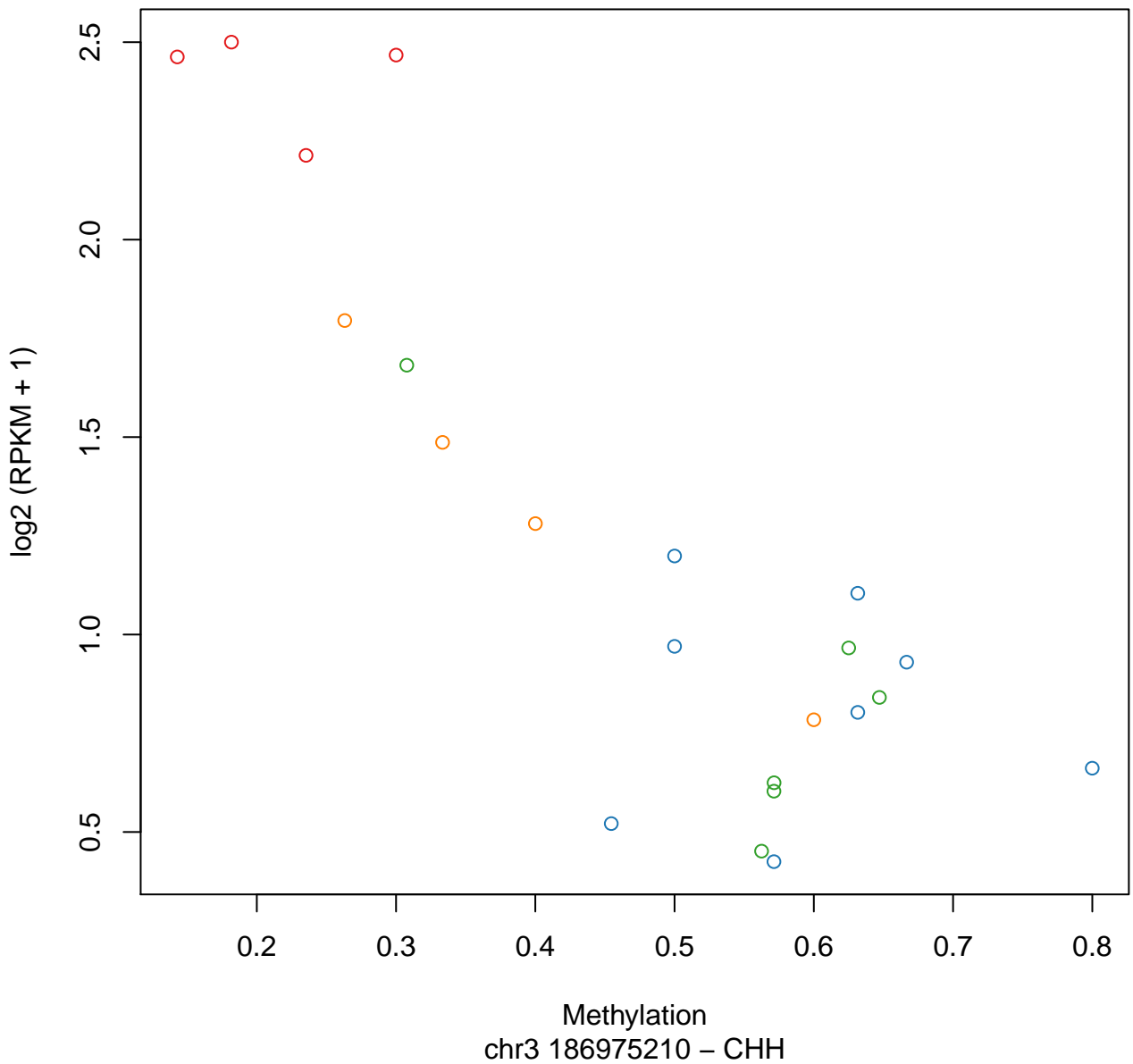


exon ENSE00003582256.1_1 FDR 0.00121

TG

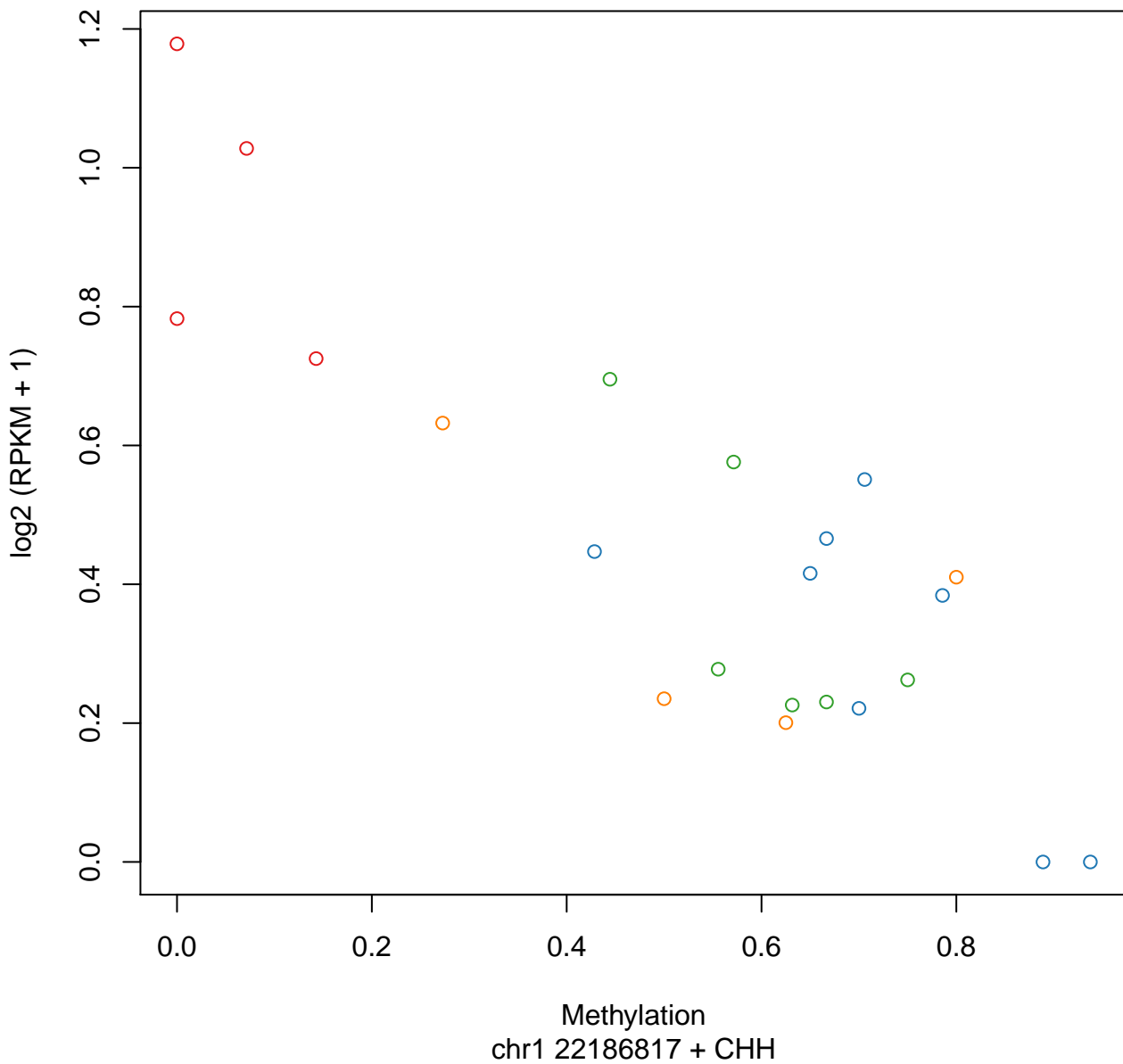


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MASP1



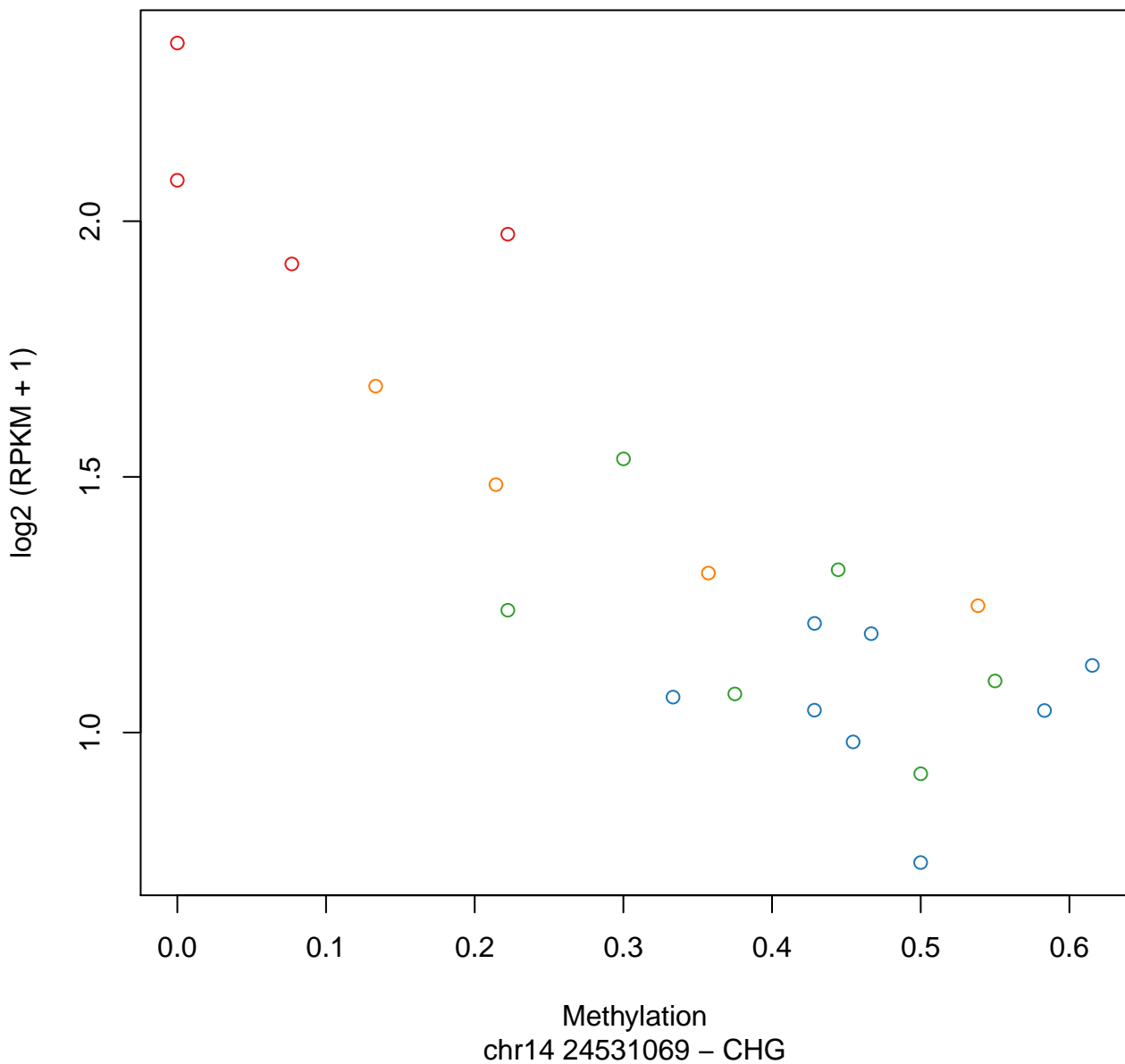
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HSPG2



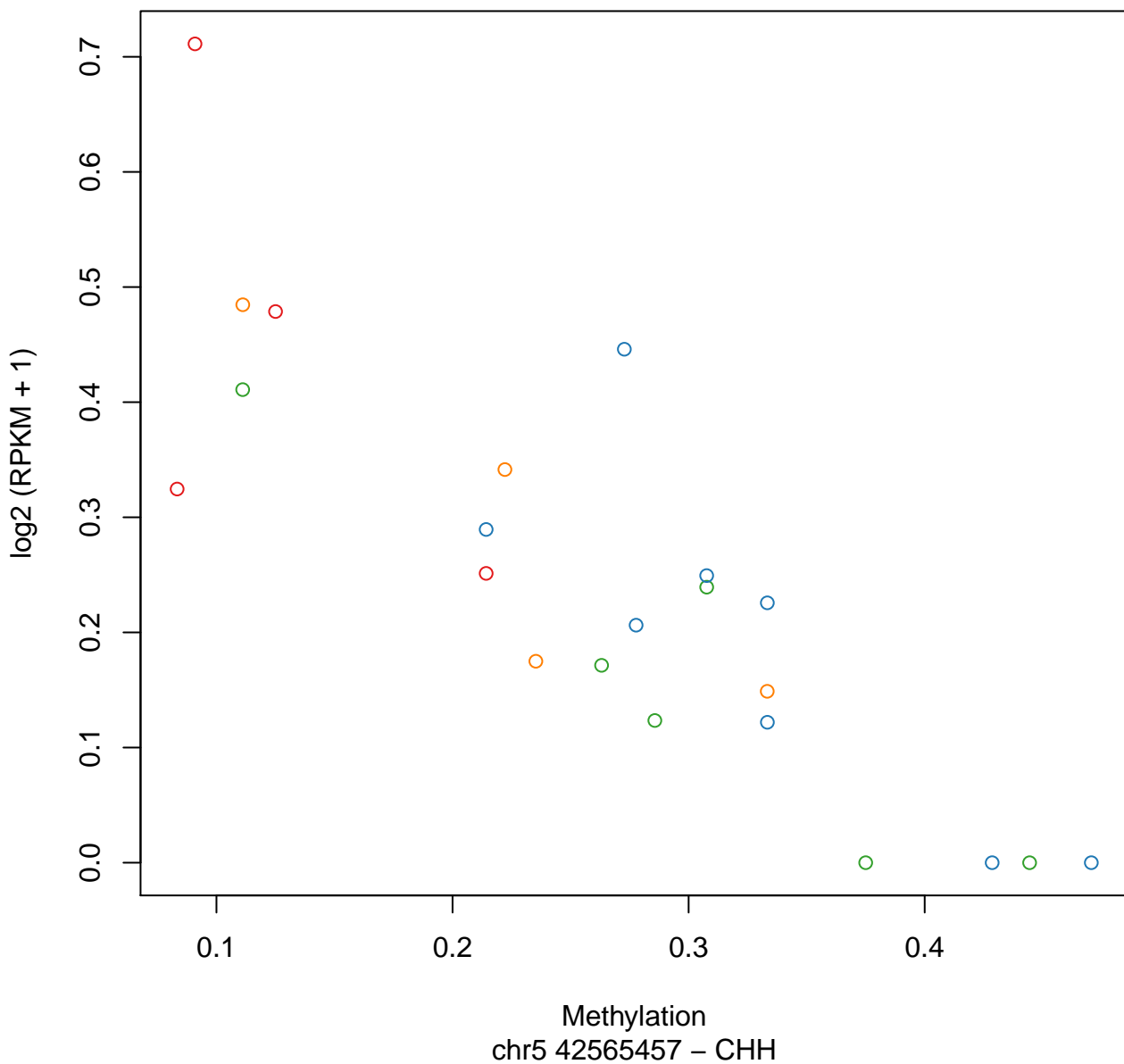
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CARMIL3

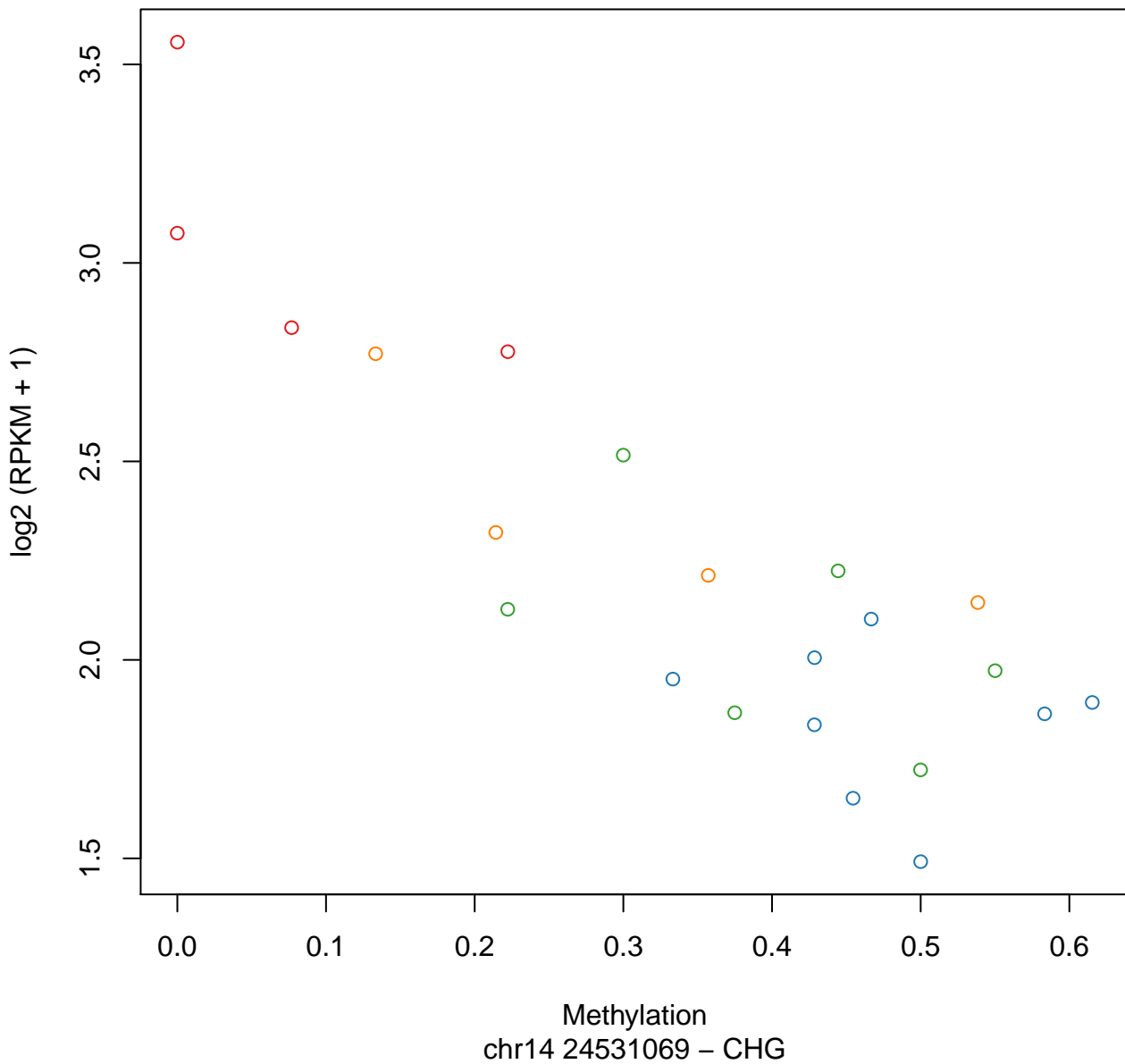


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GHR

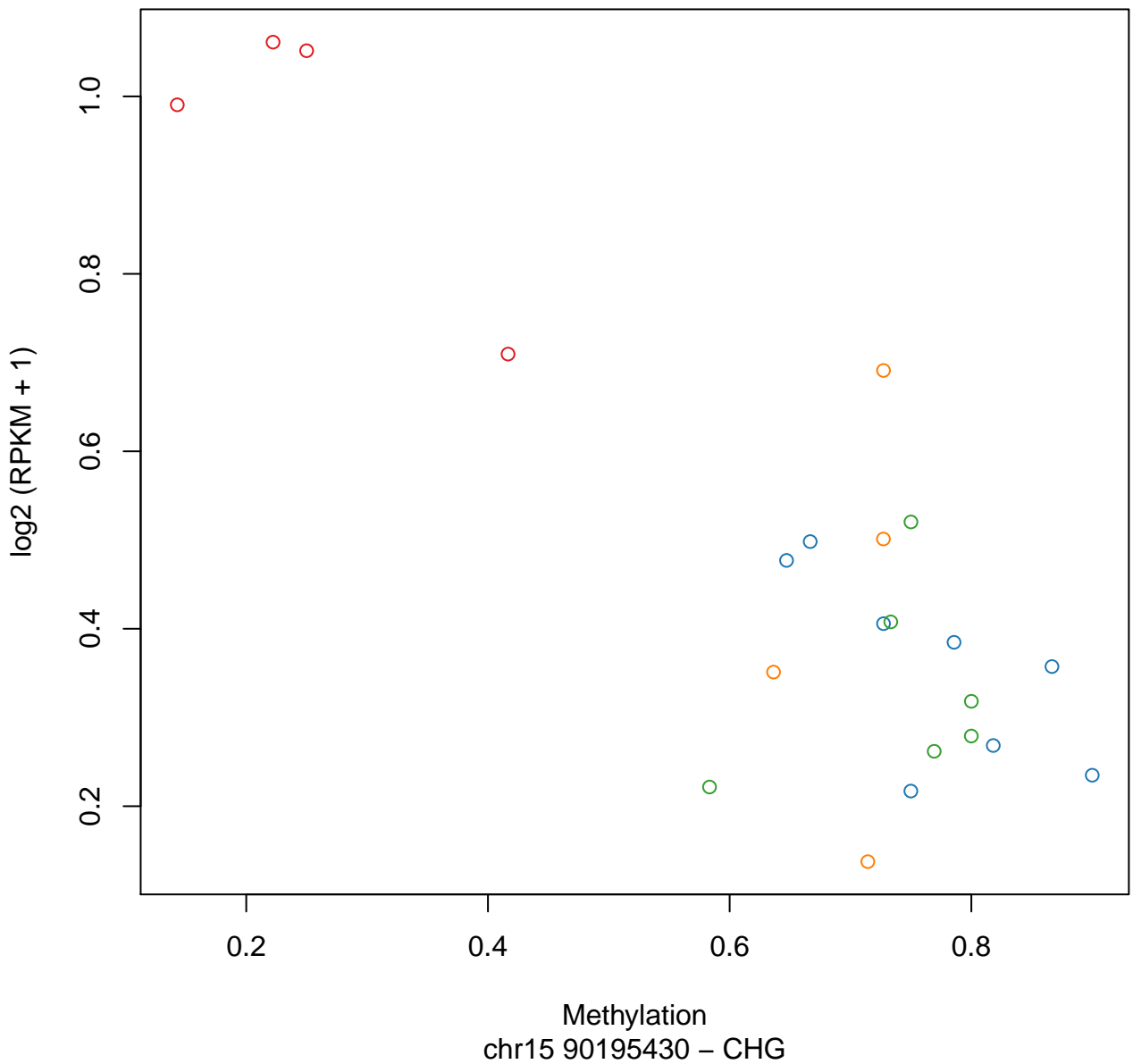


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CARMIL3



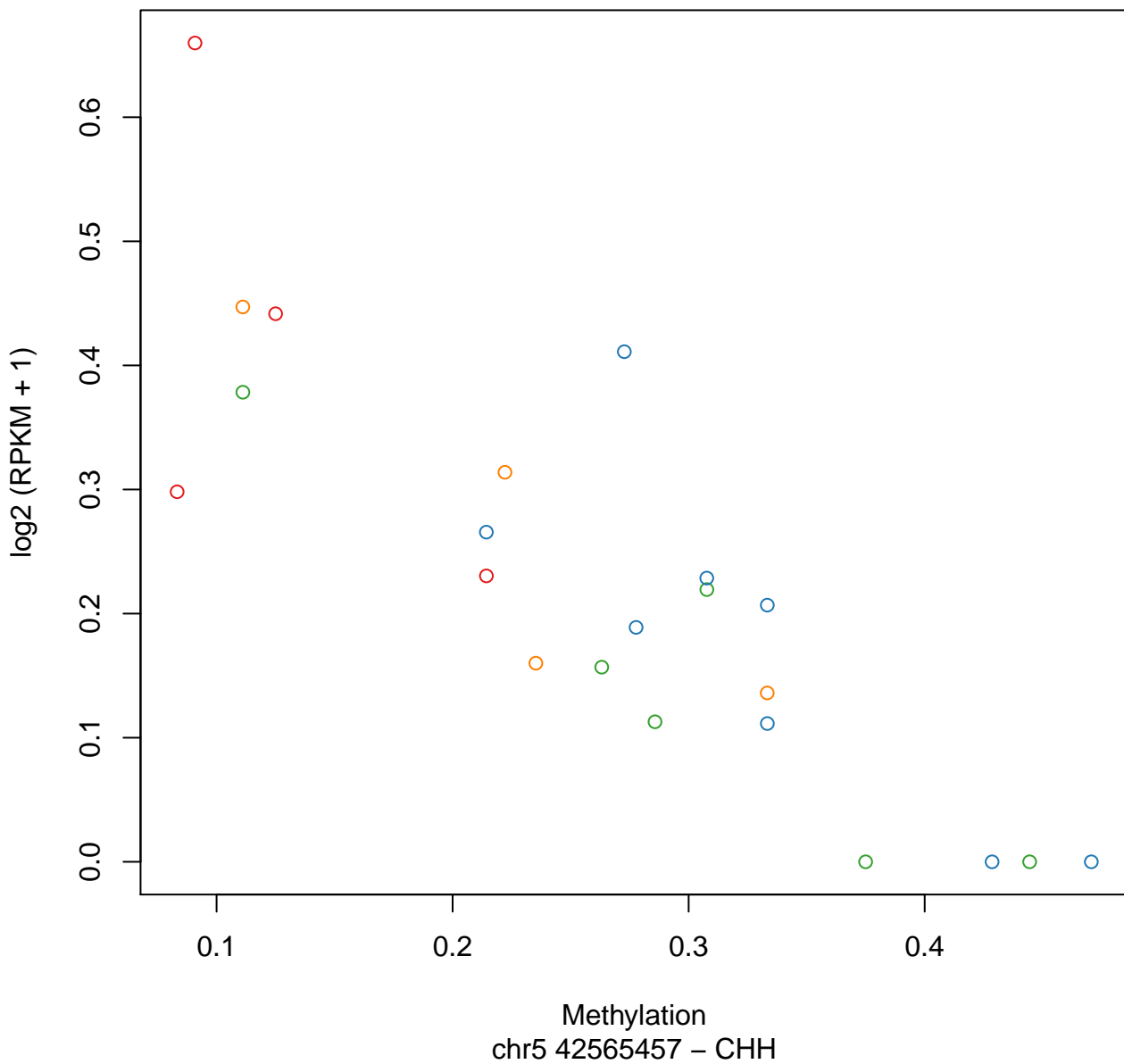
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KIF7



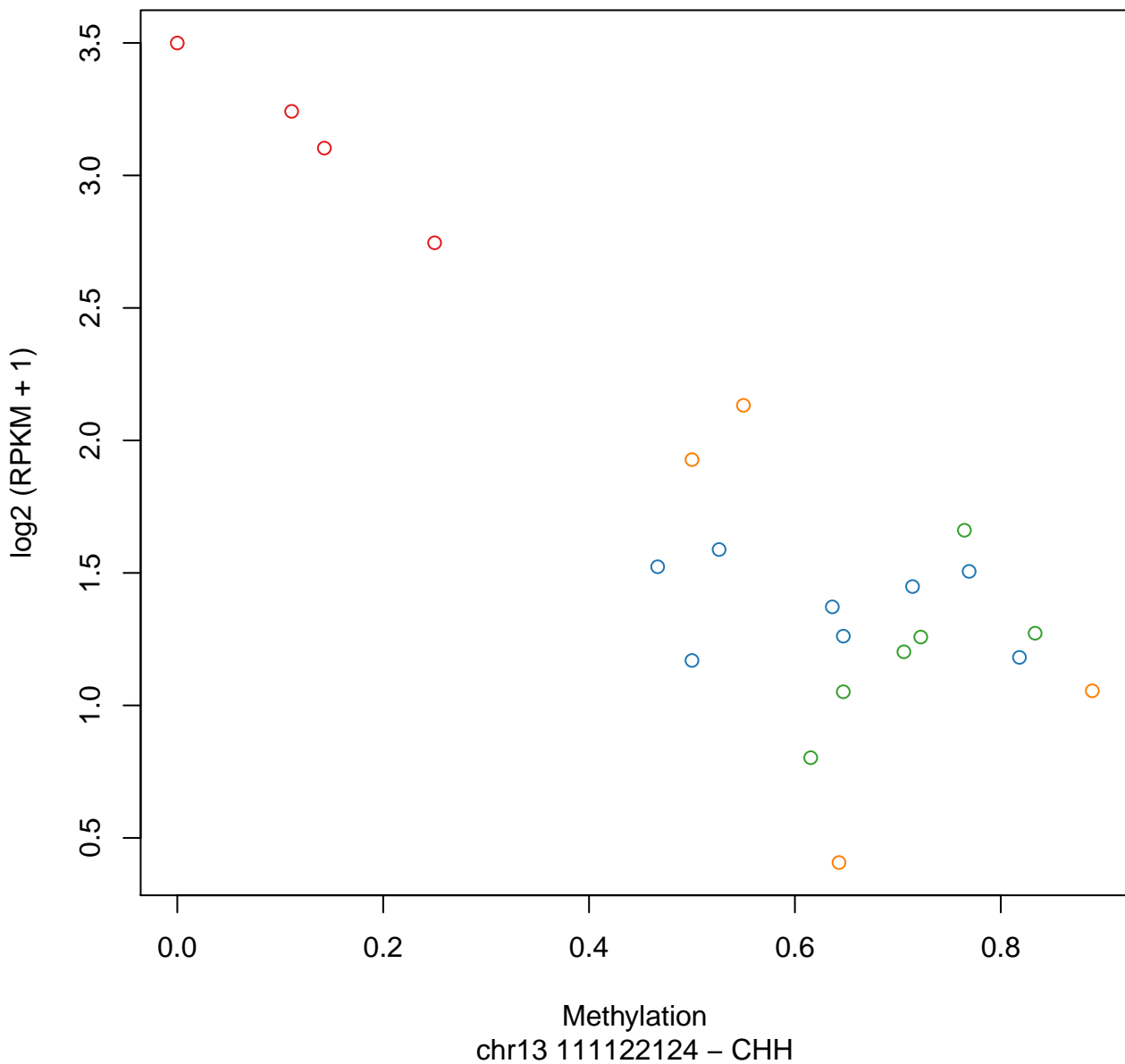
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GHR



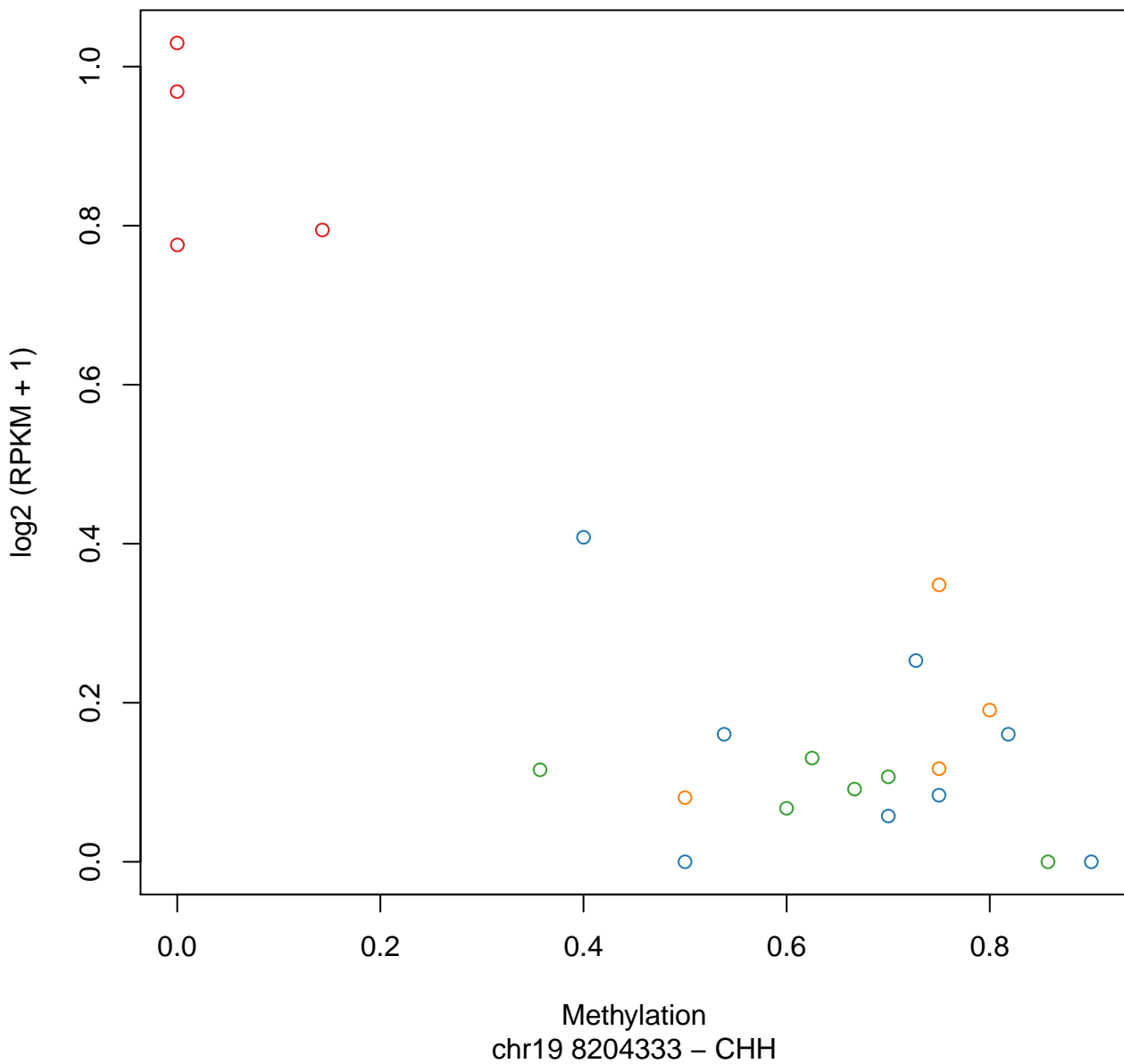
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COL4A2

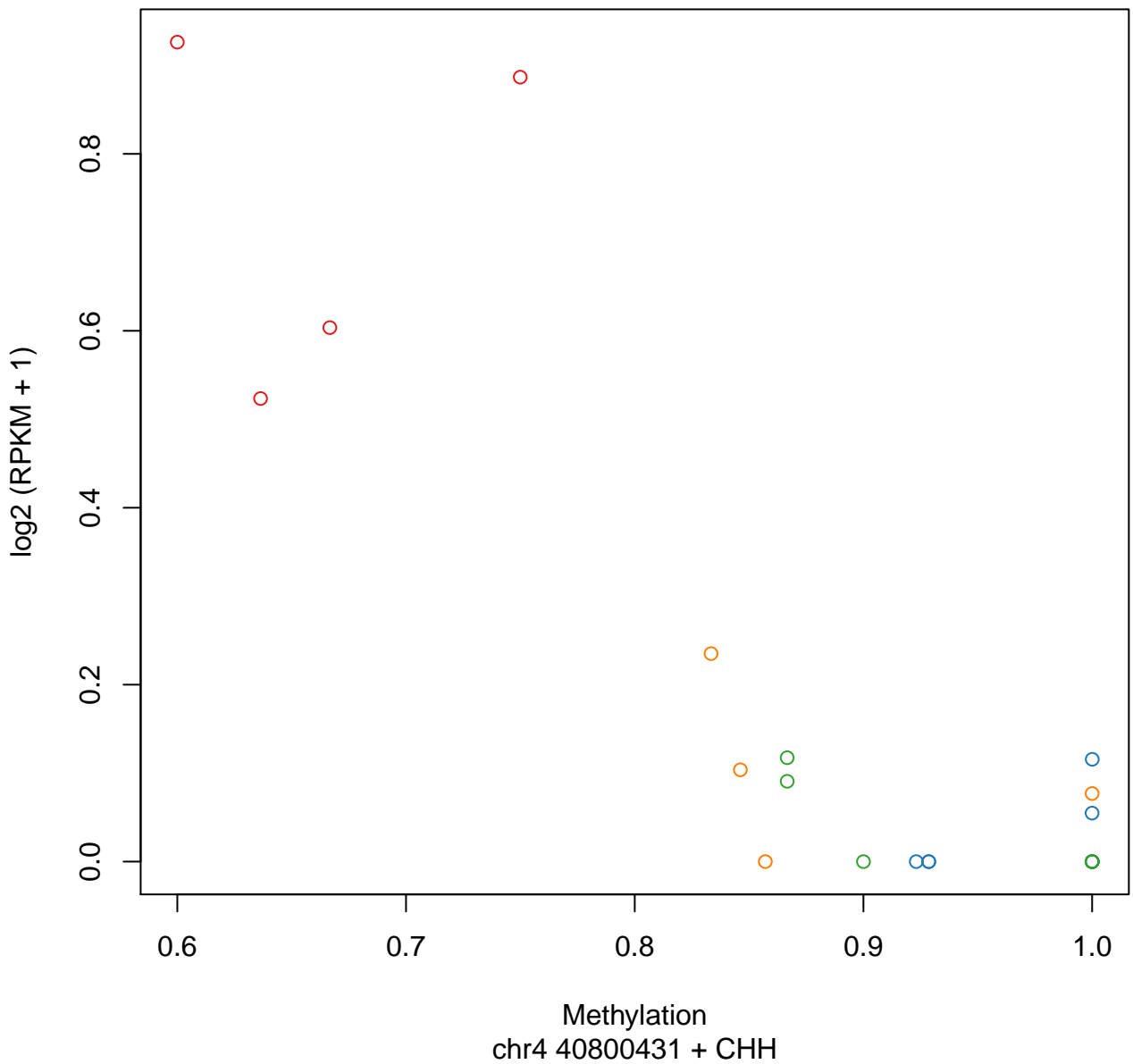


exon ENSE00001734041.1_1 FDR 0.00143

FBN3

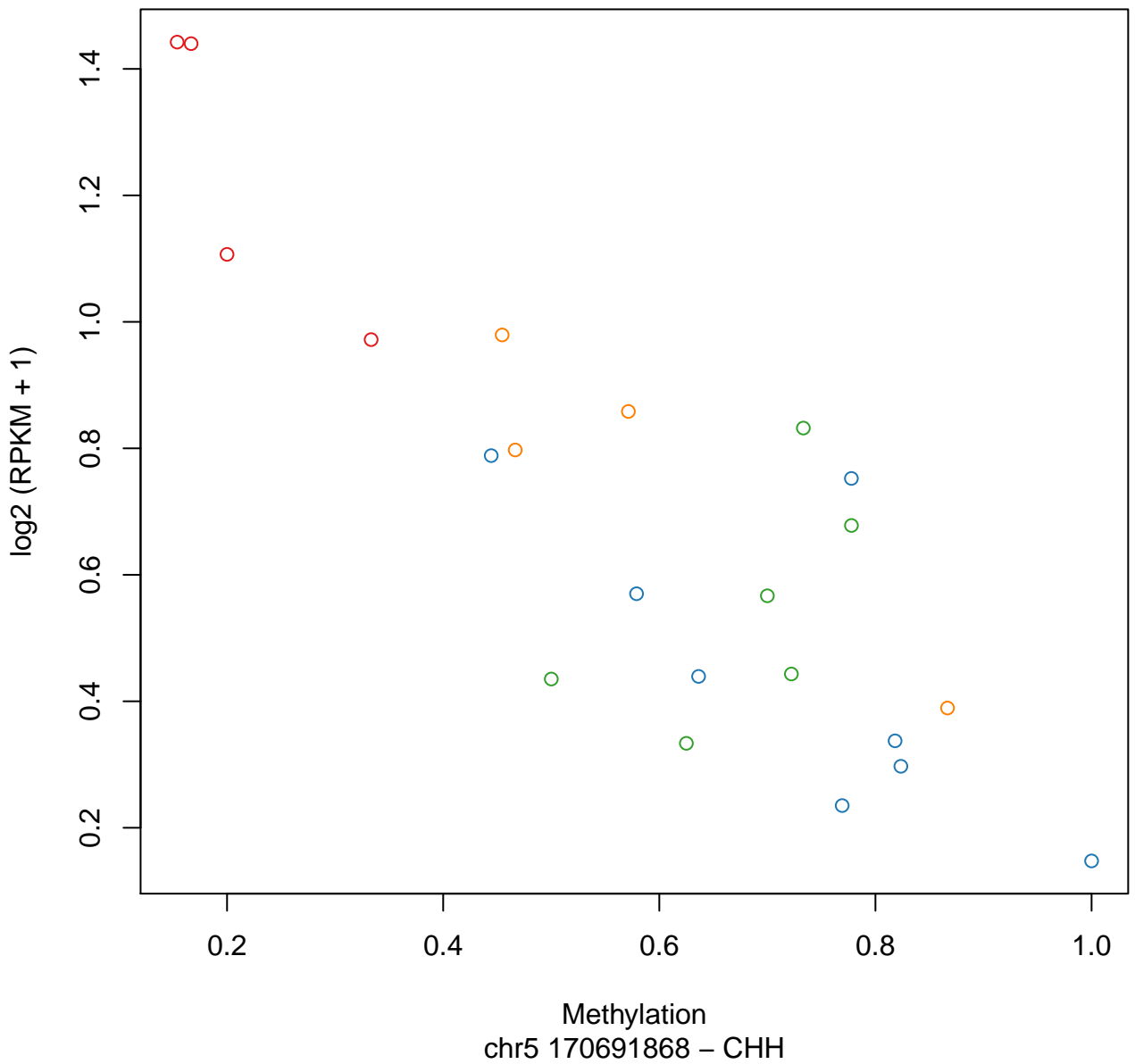


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NSUN7



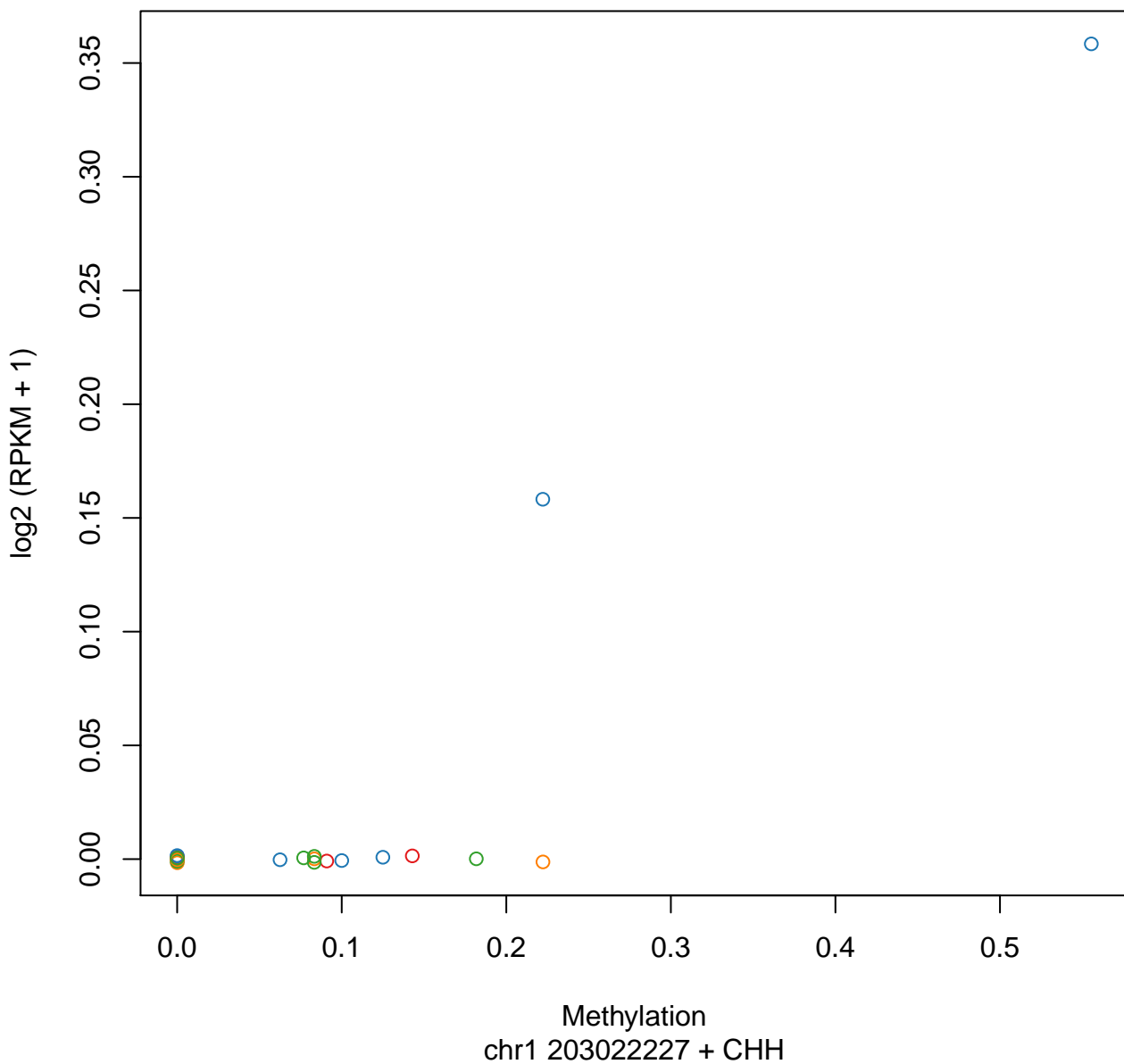
exon ENSE00003681284.1_1 FDR 0.00153

RANBP17

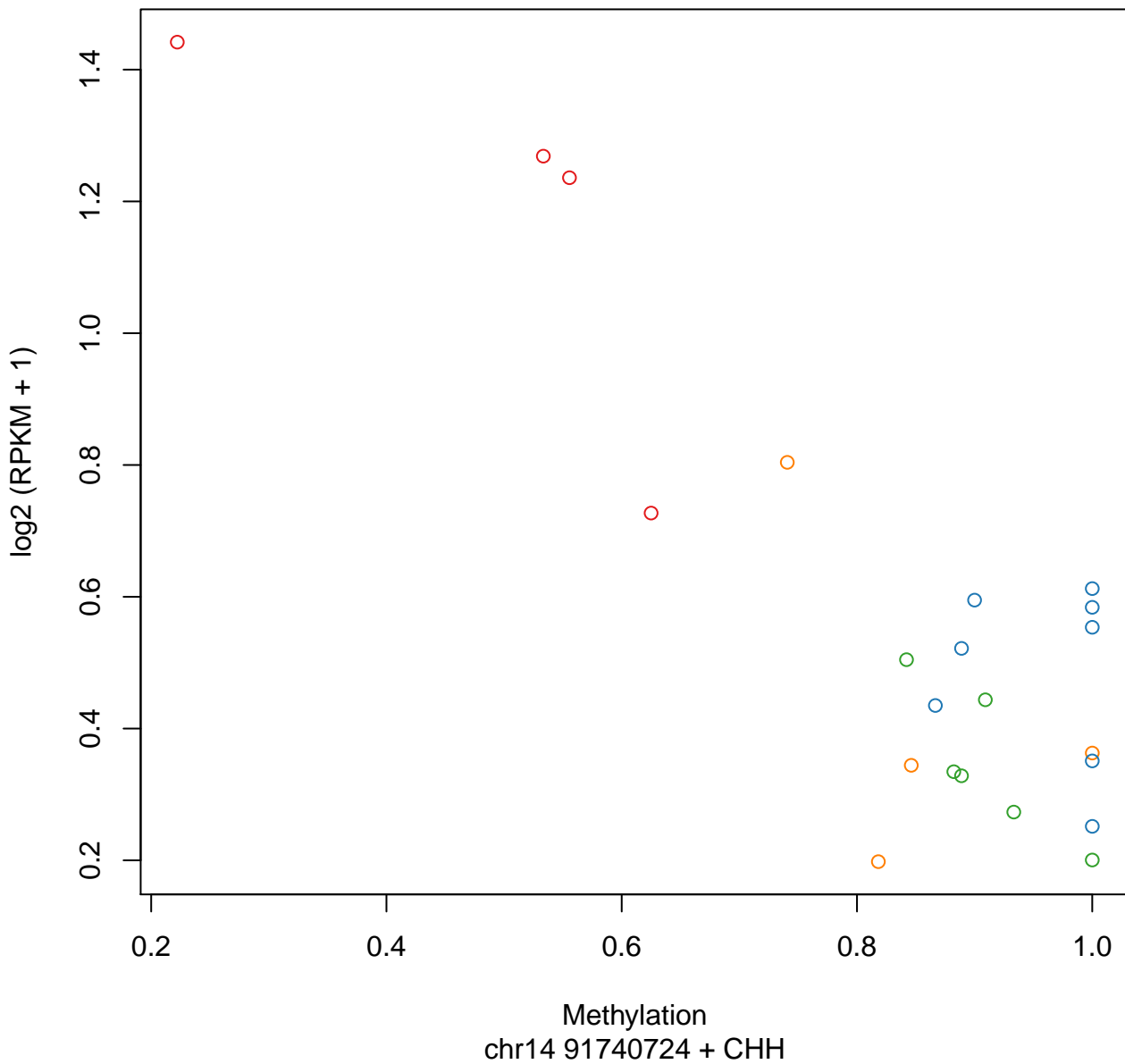


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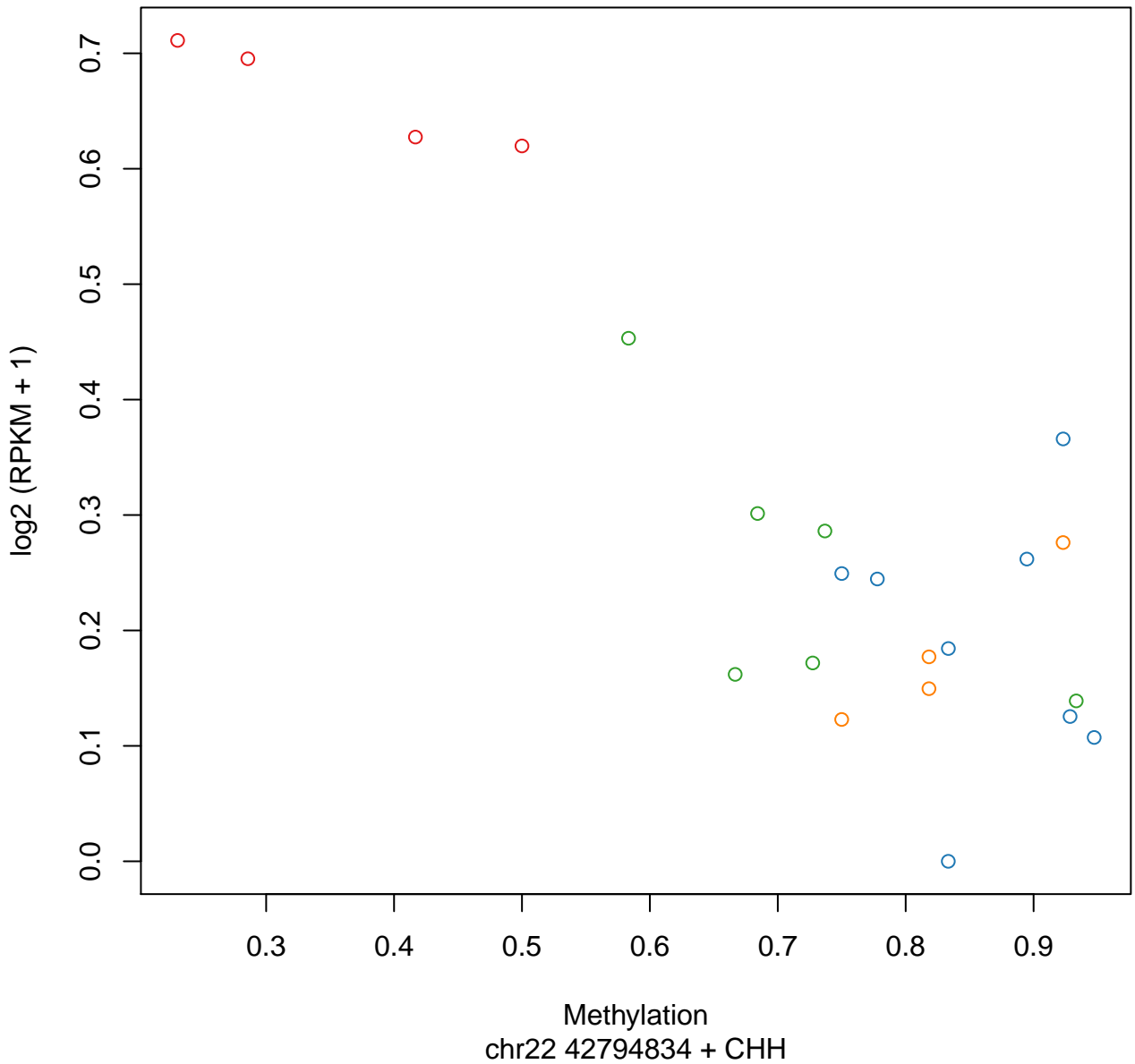
PPFIA4



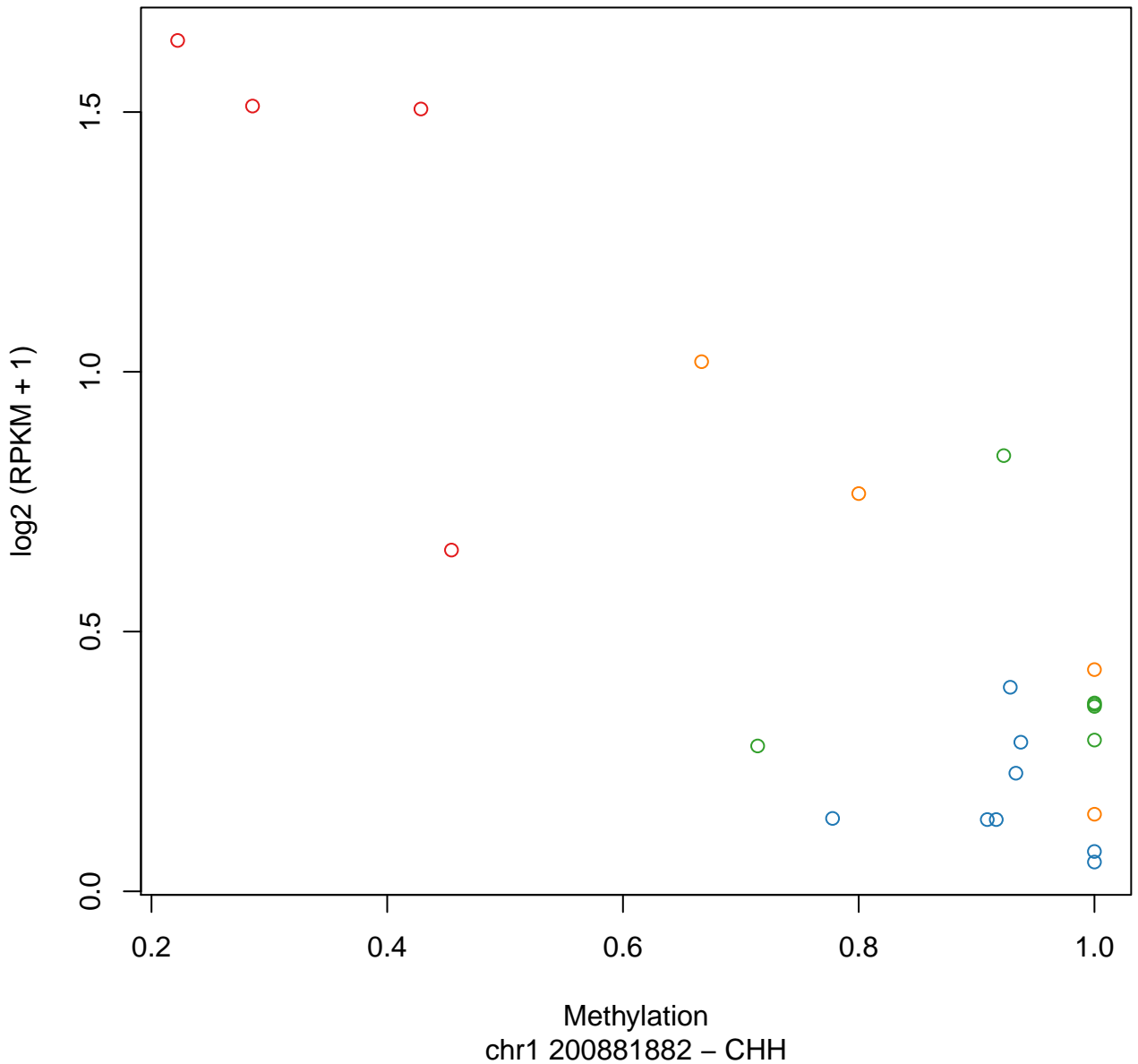
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CCDC88C



exon ENSE00001722461.1_1 FDR 0.00161
NFAM1

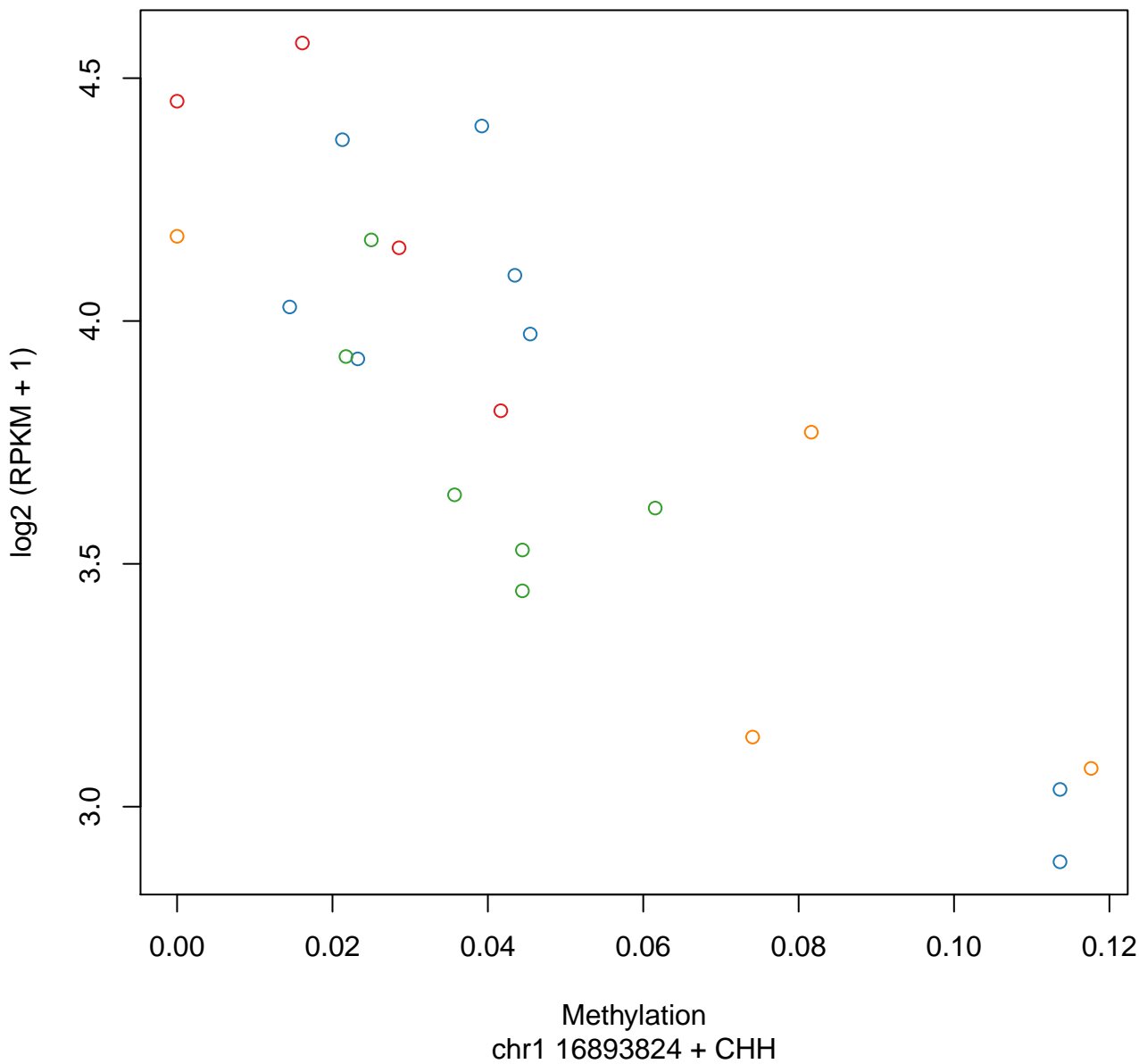


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C1orf106



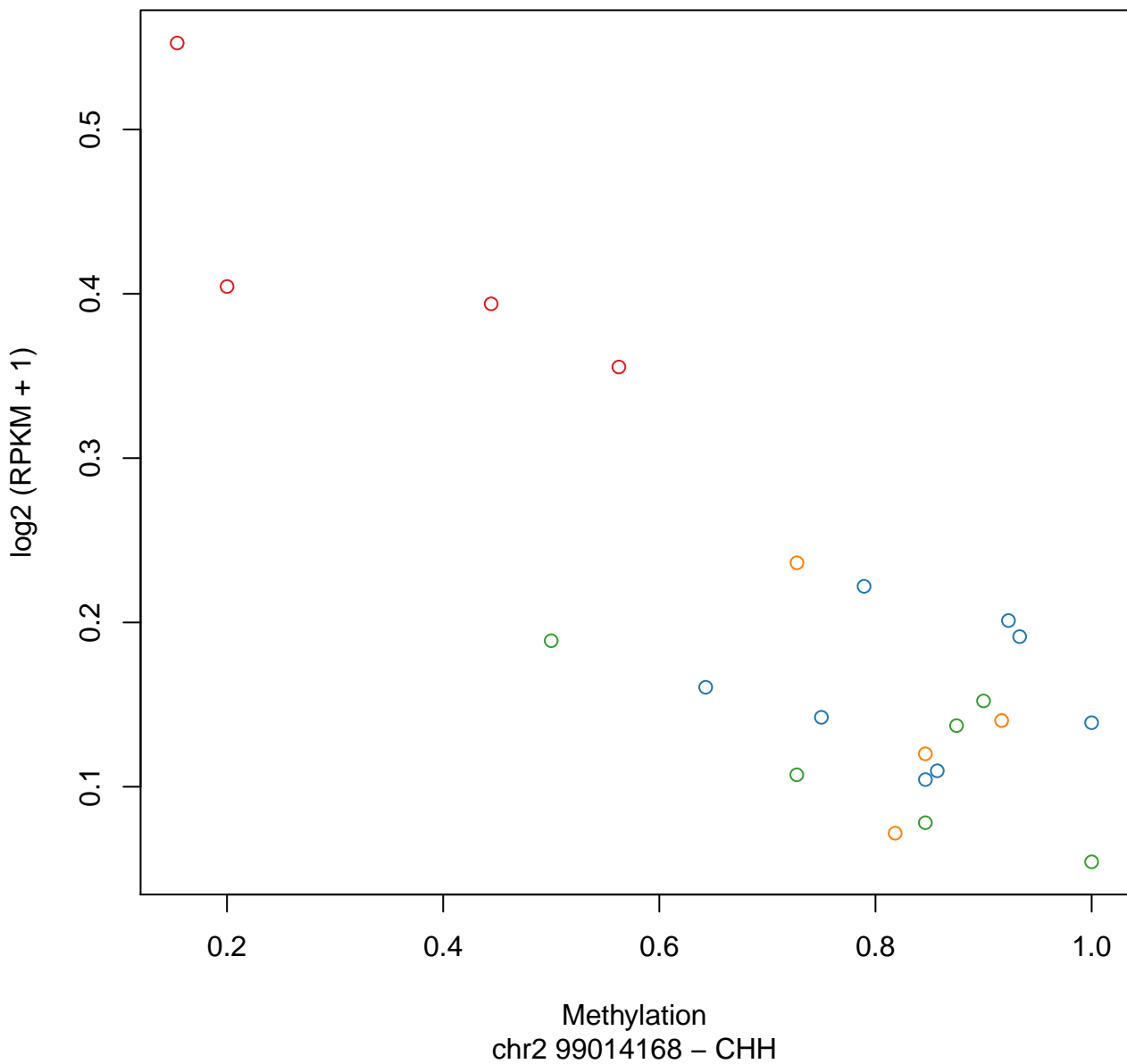
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NBPF1

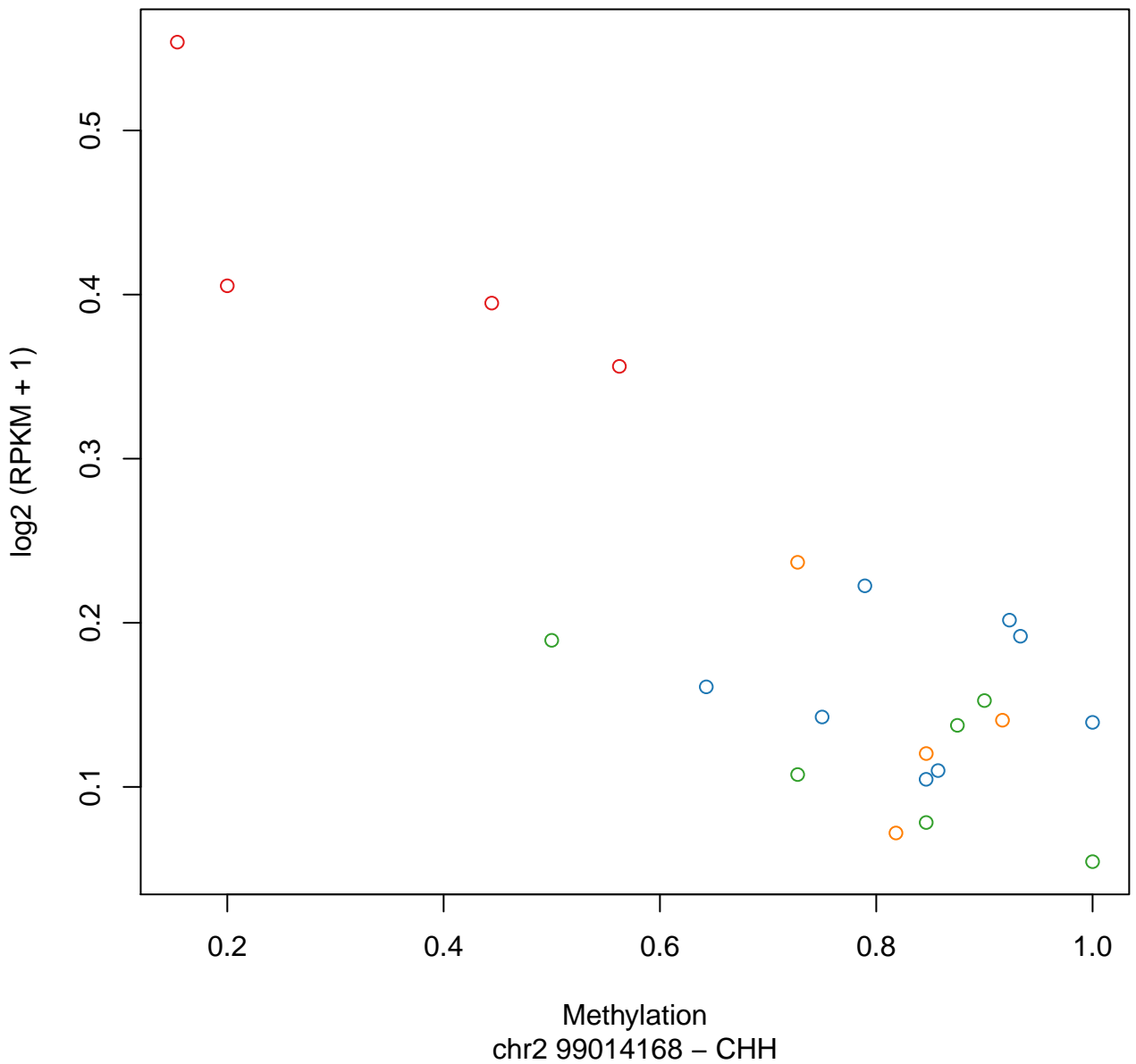


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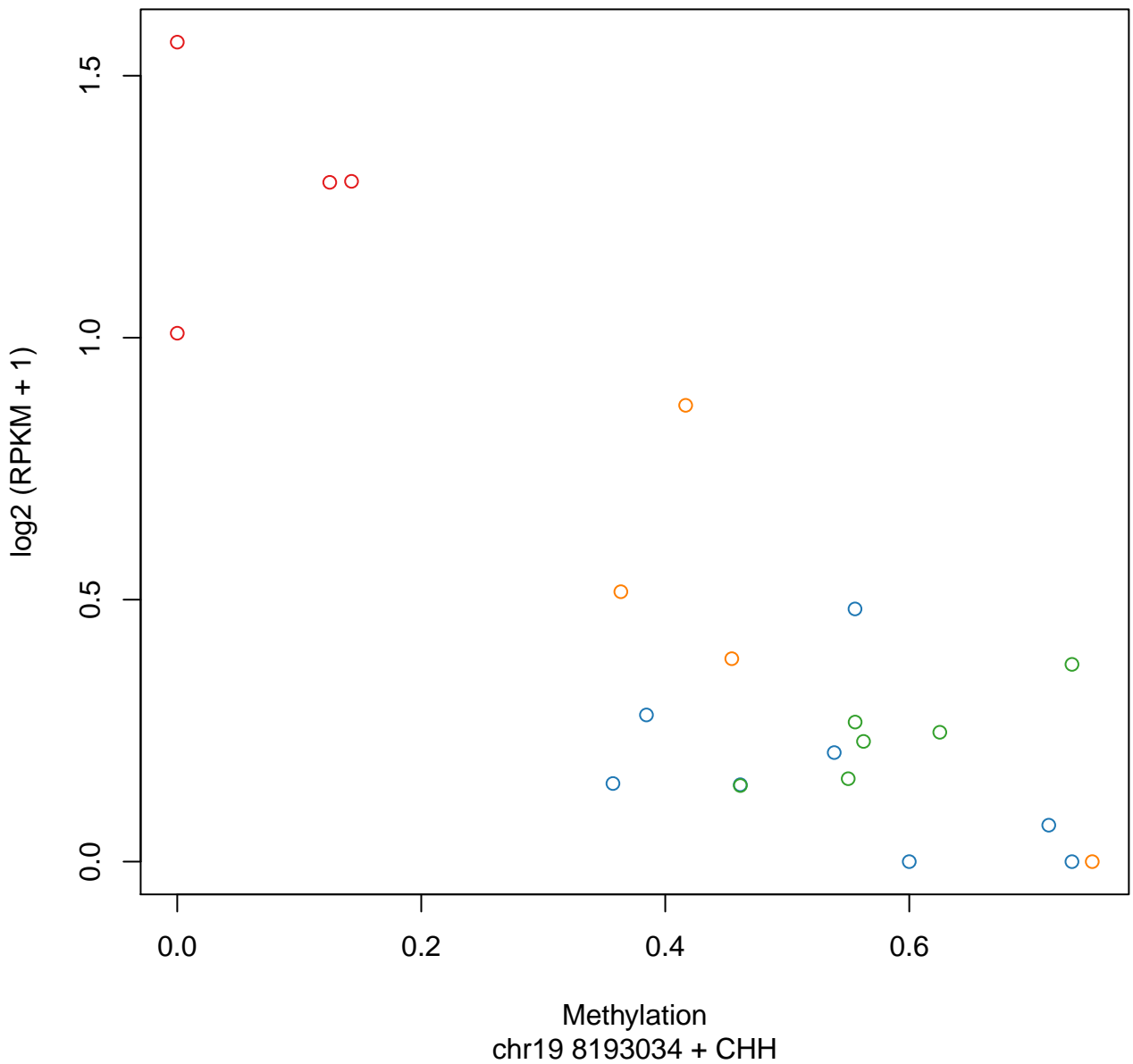
CNGA3



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CNGA3

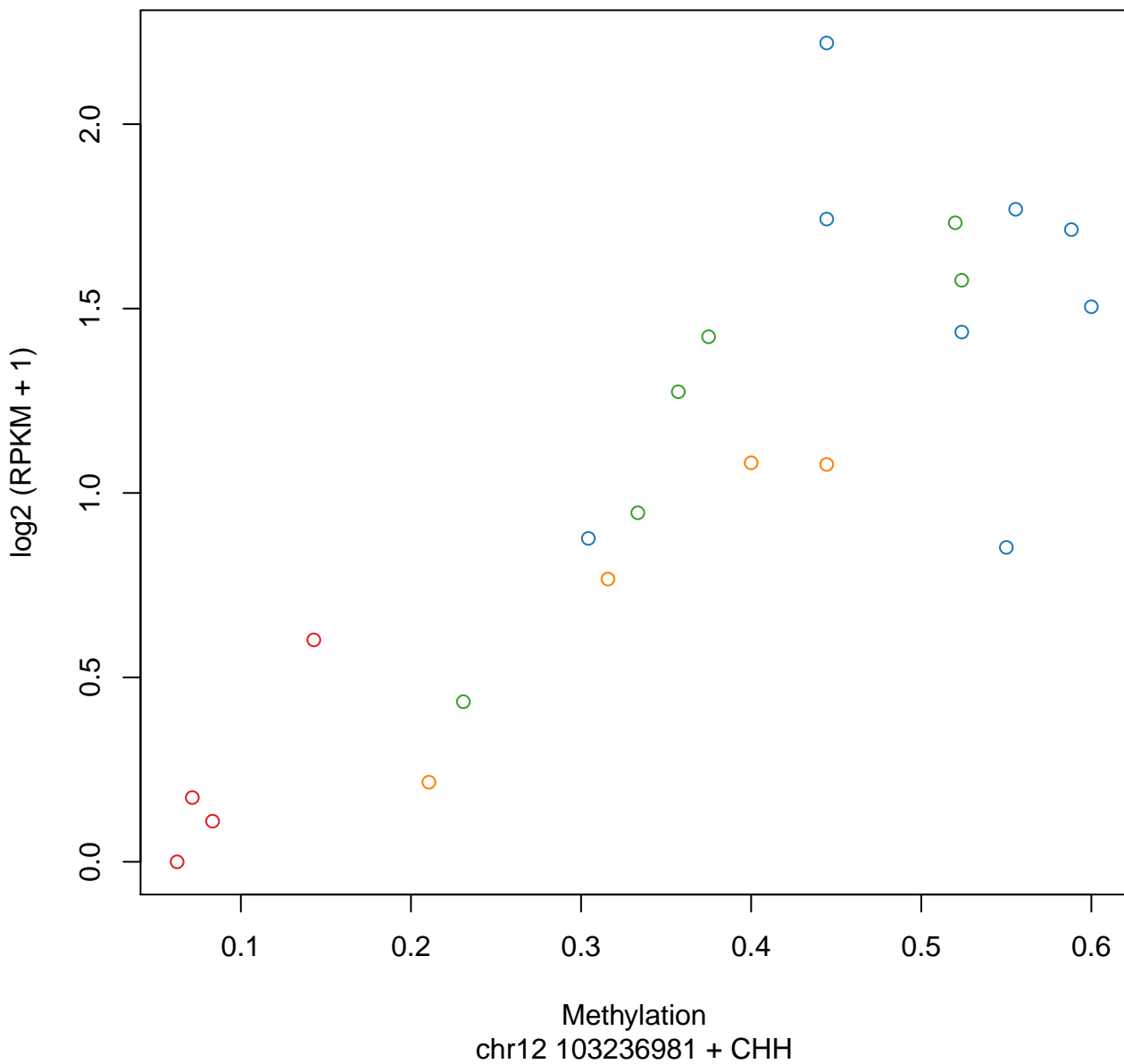


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FBN3



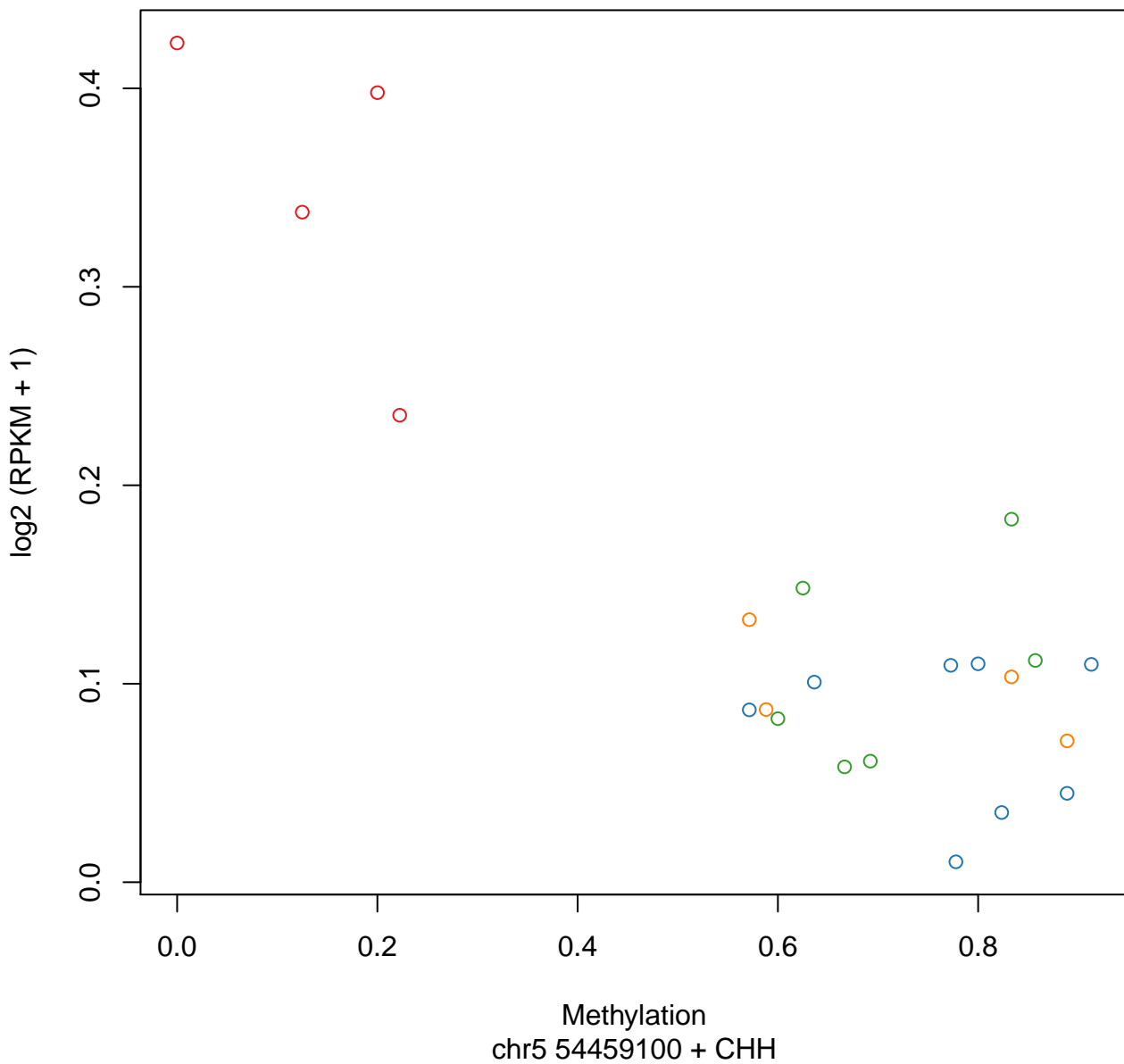
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PAH

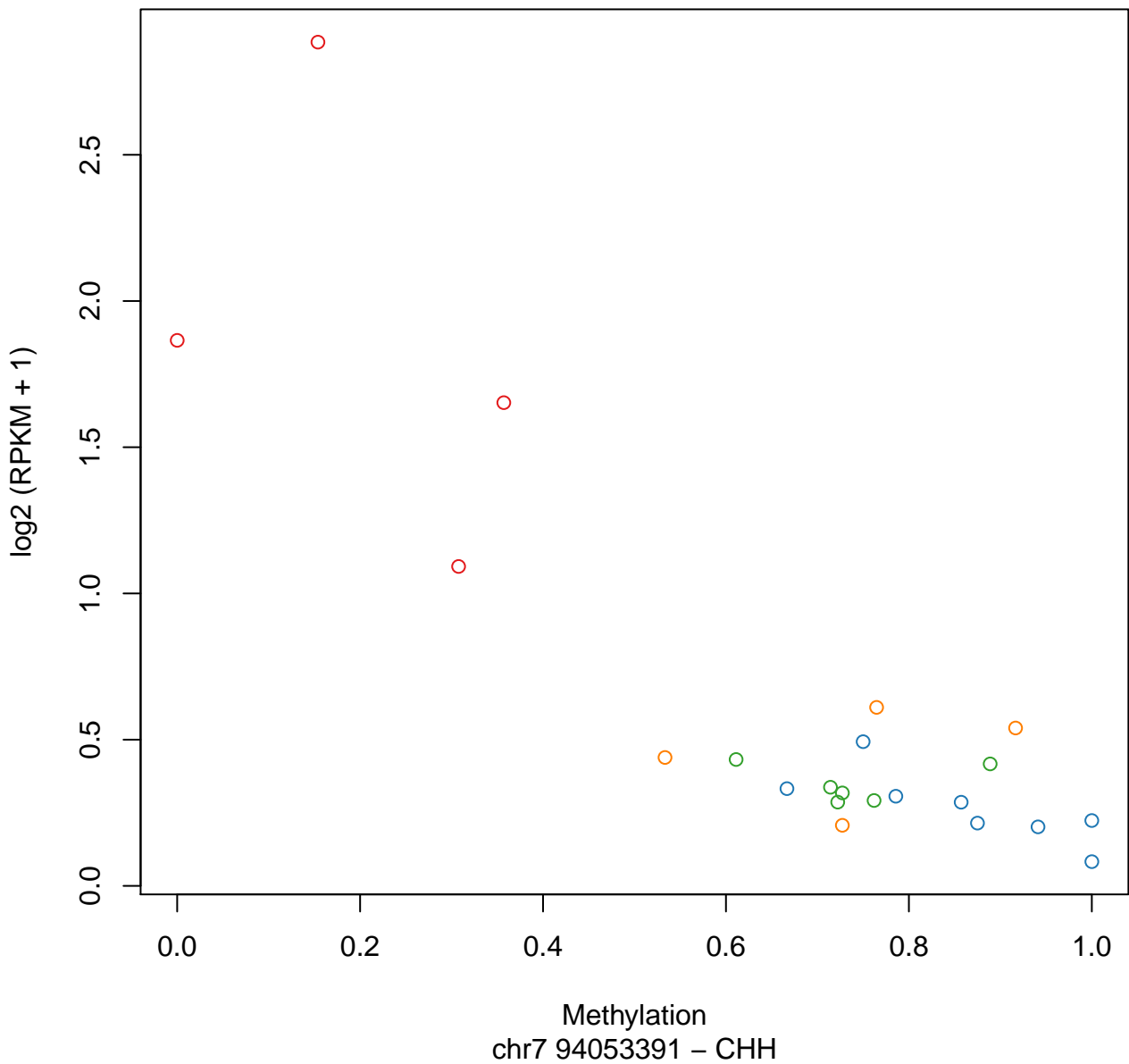


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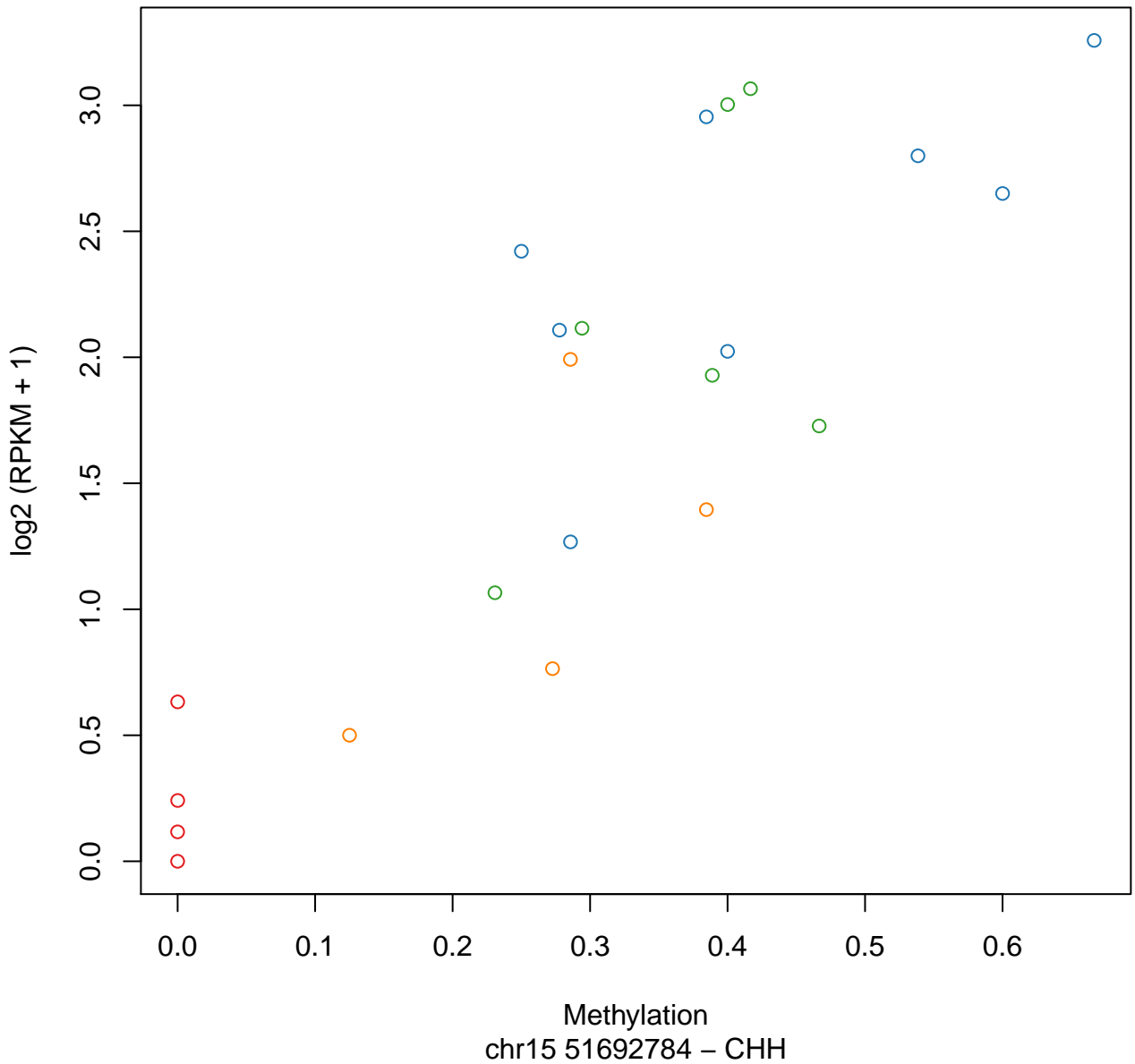


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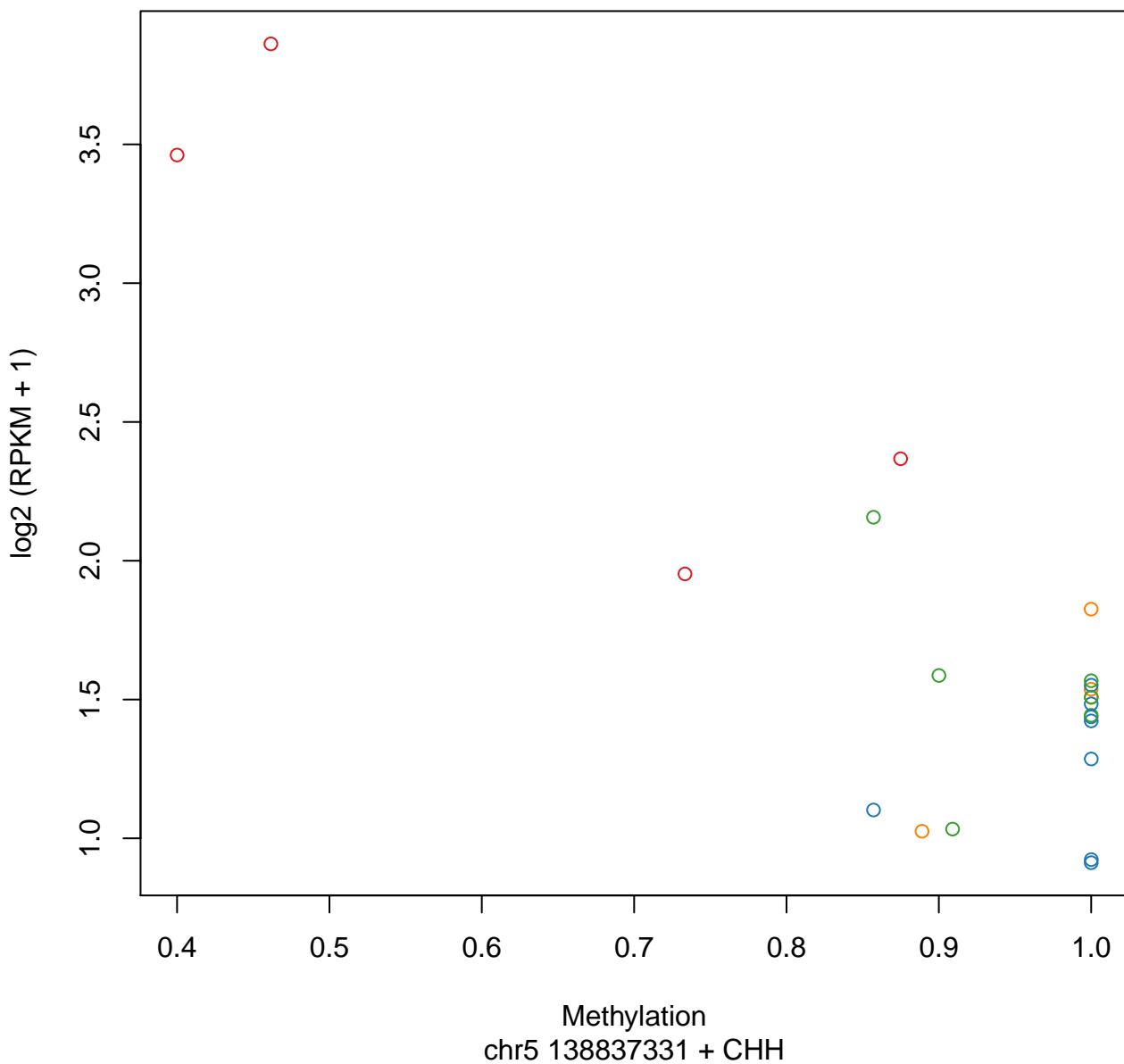


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GLDN

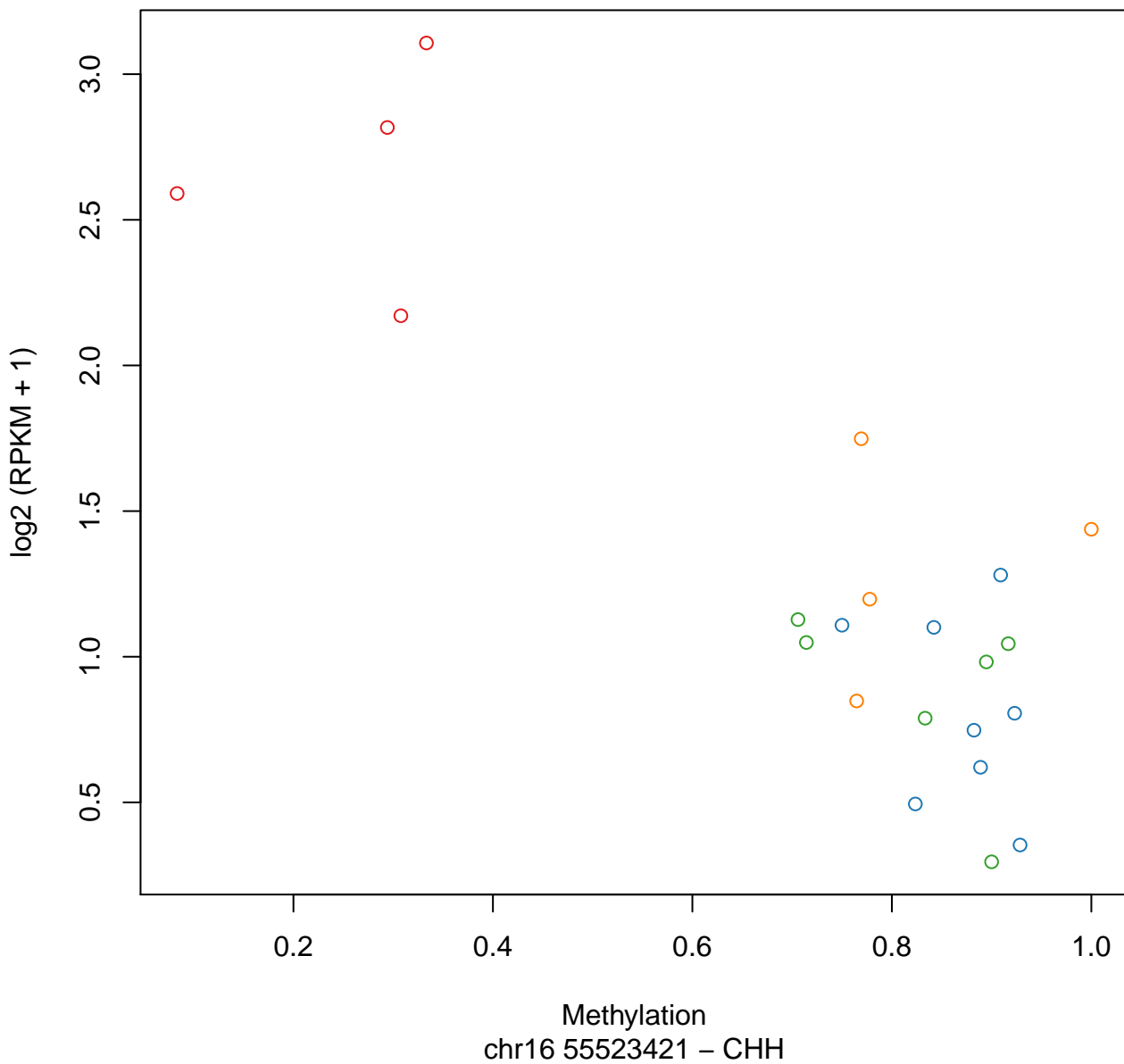


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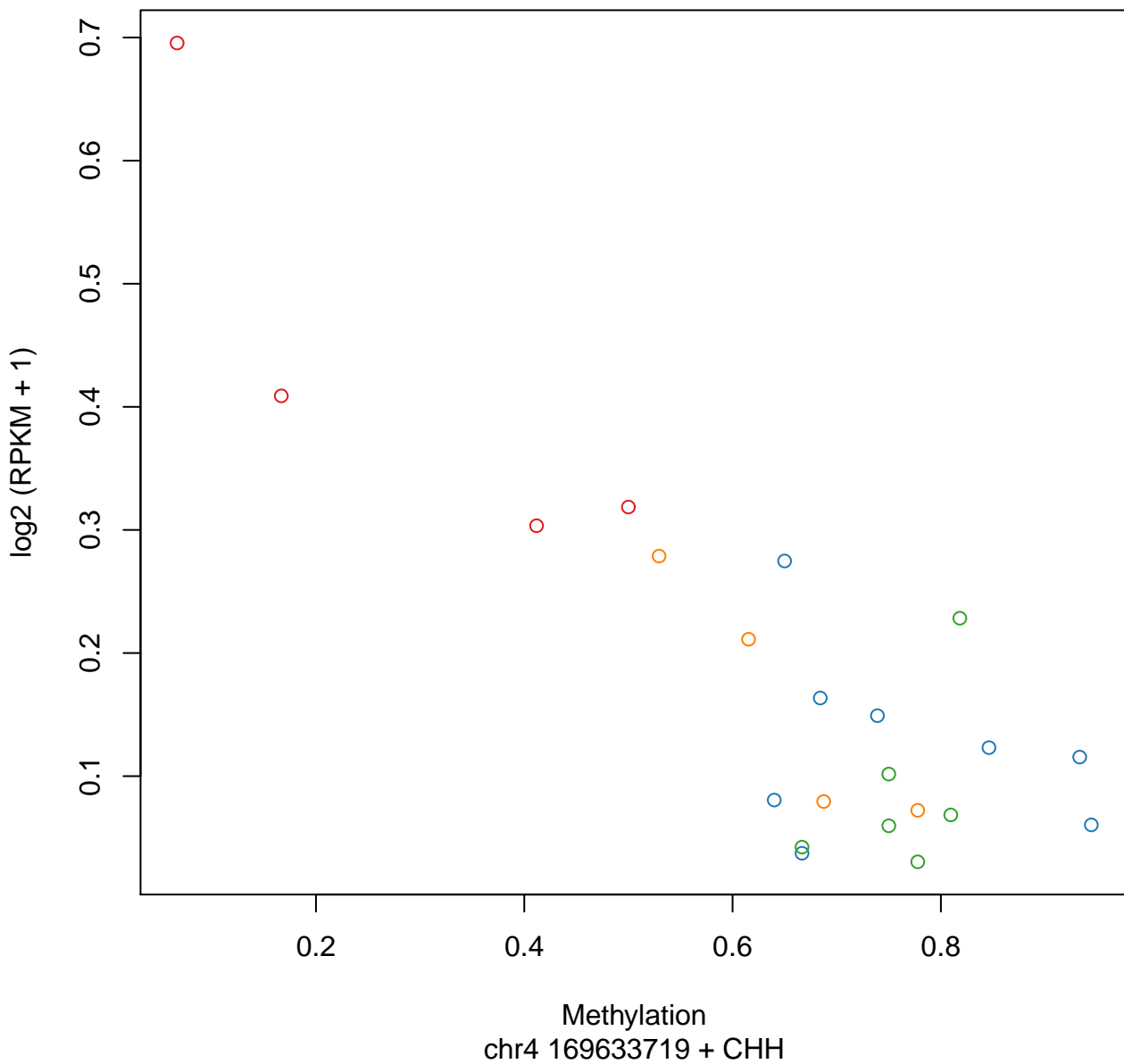
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MMP2



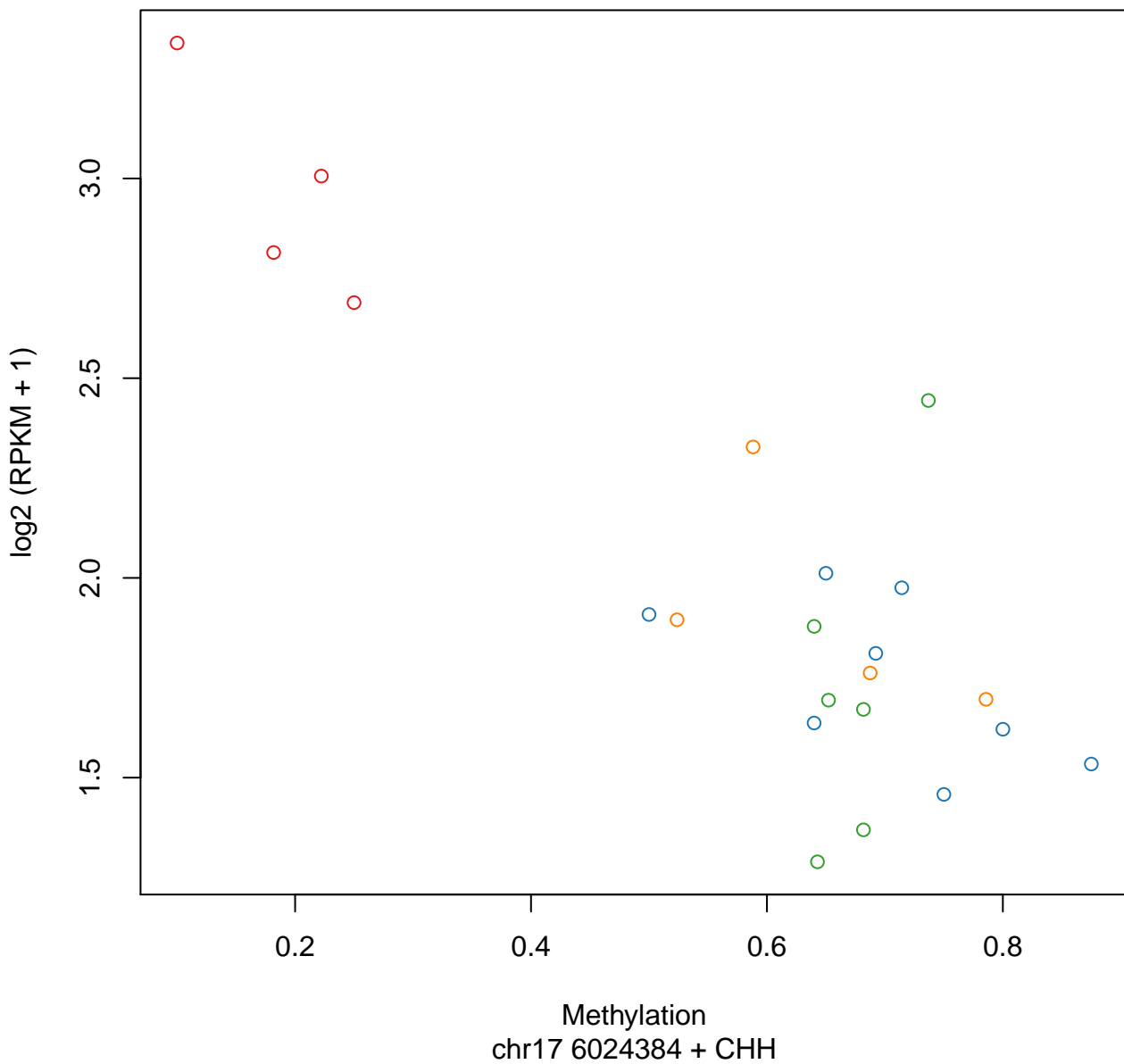
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PALLD



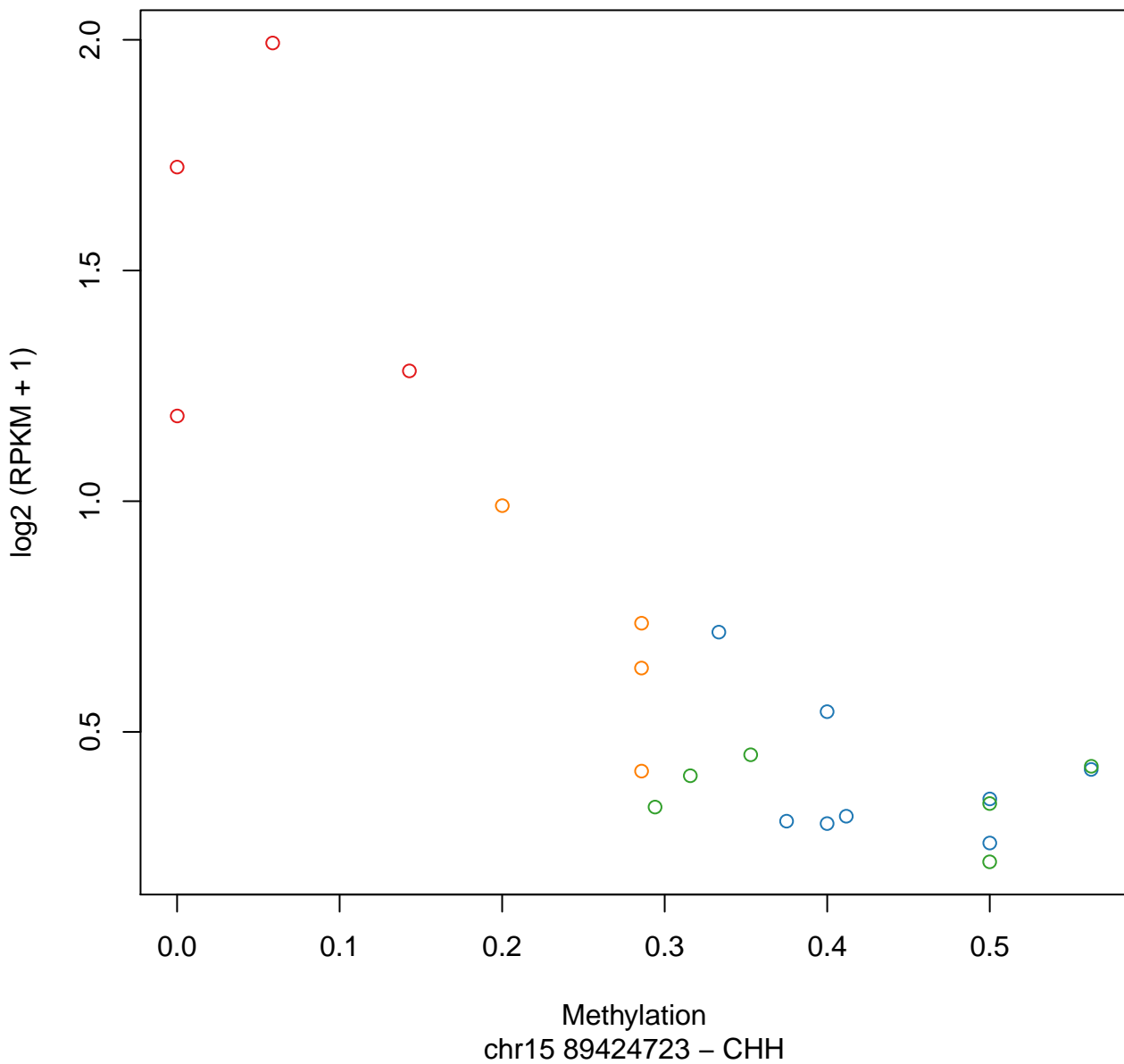
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WSCD1

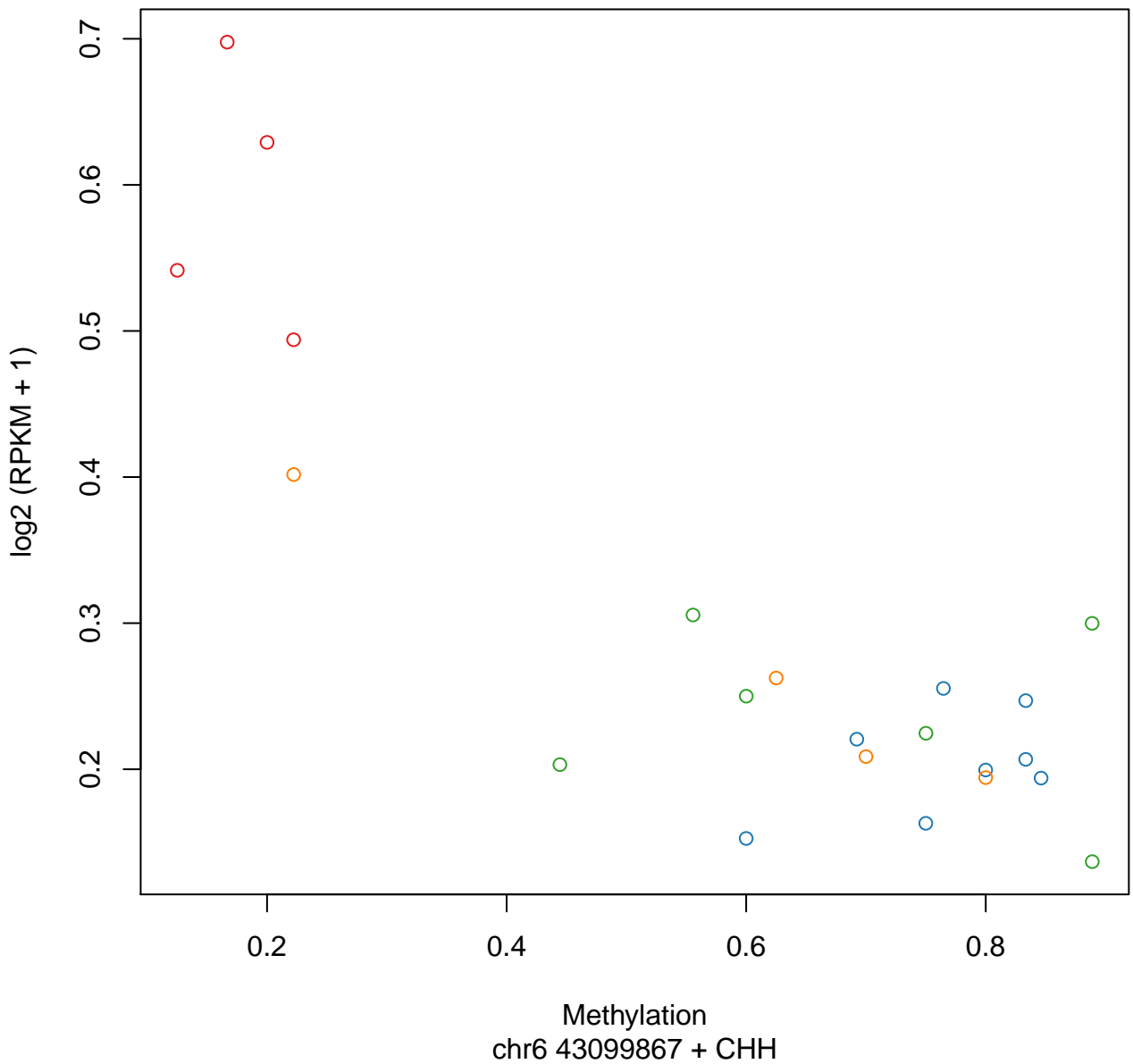


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HAPLN3

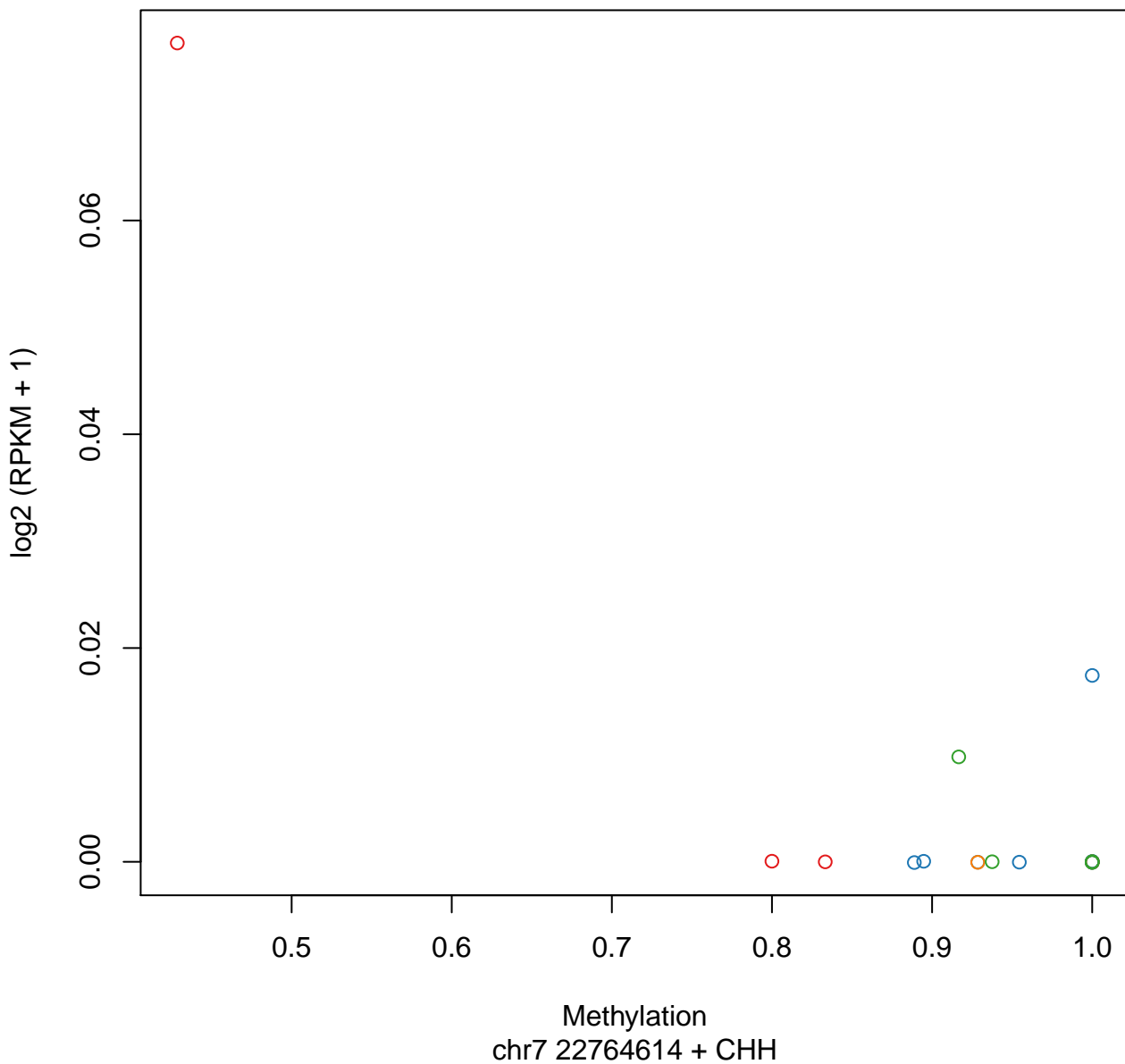


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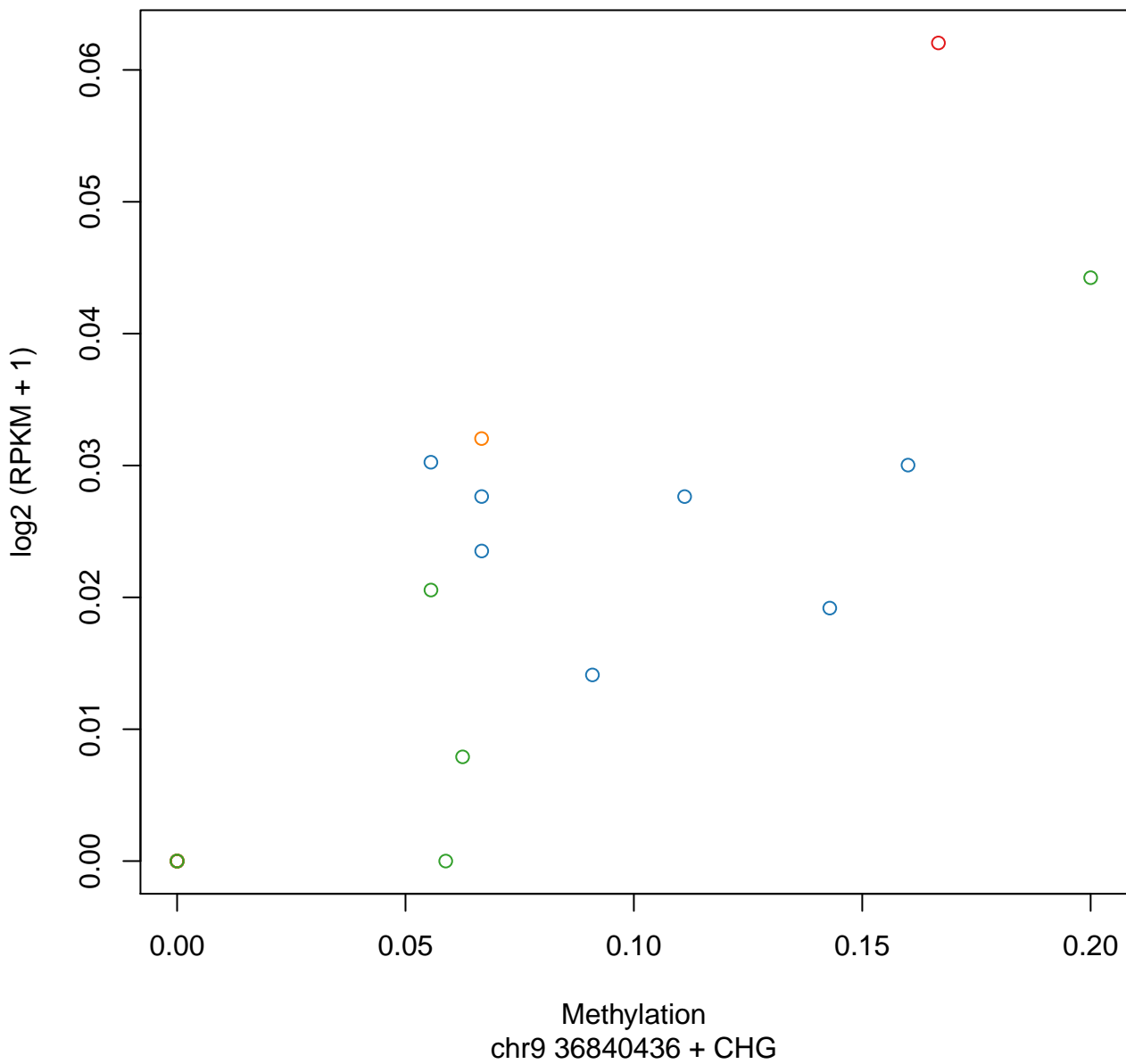
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IL6



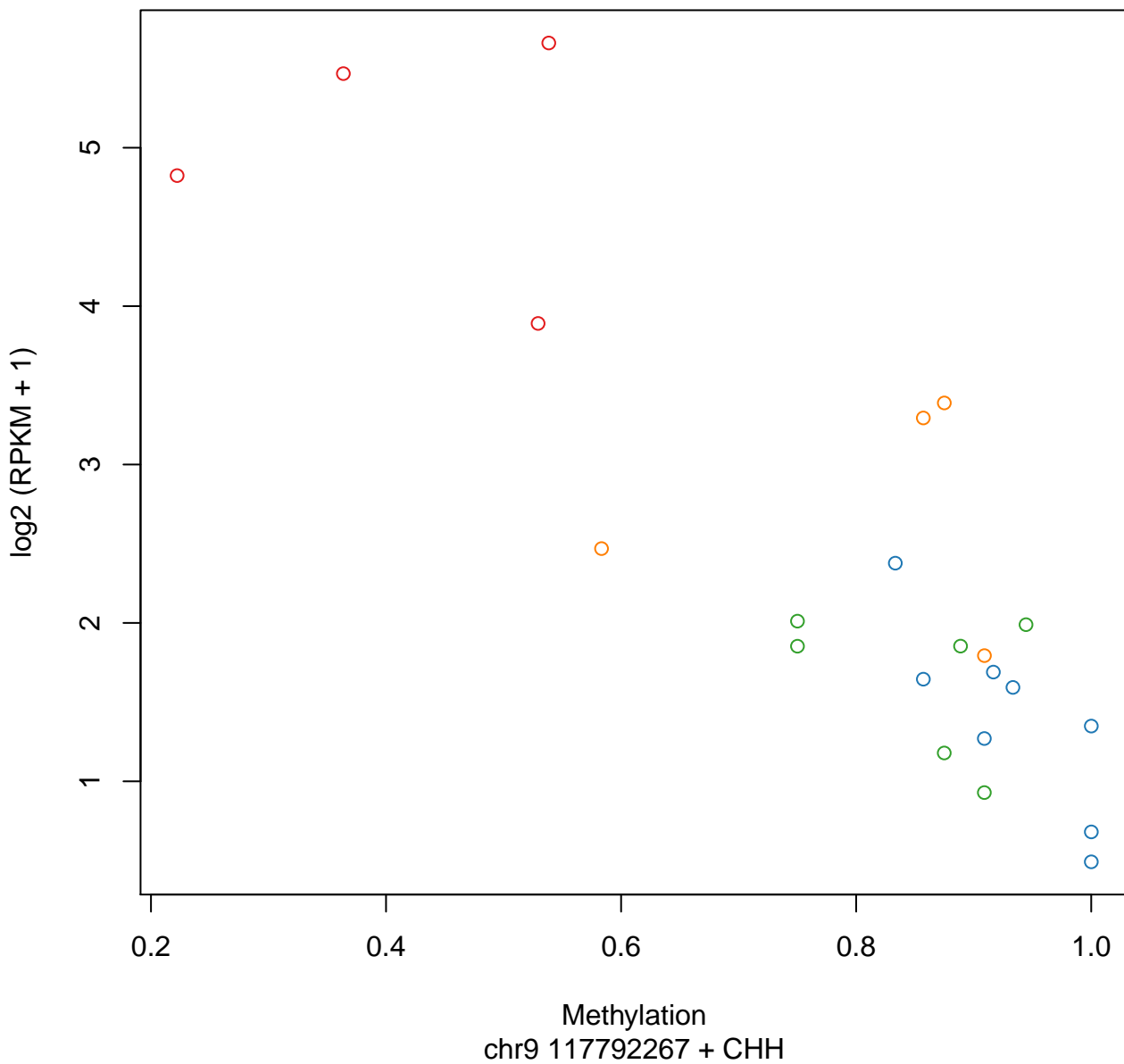
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PAX5

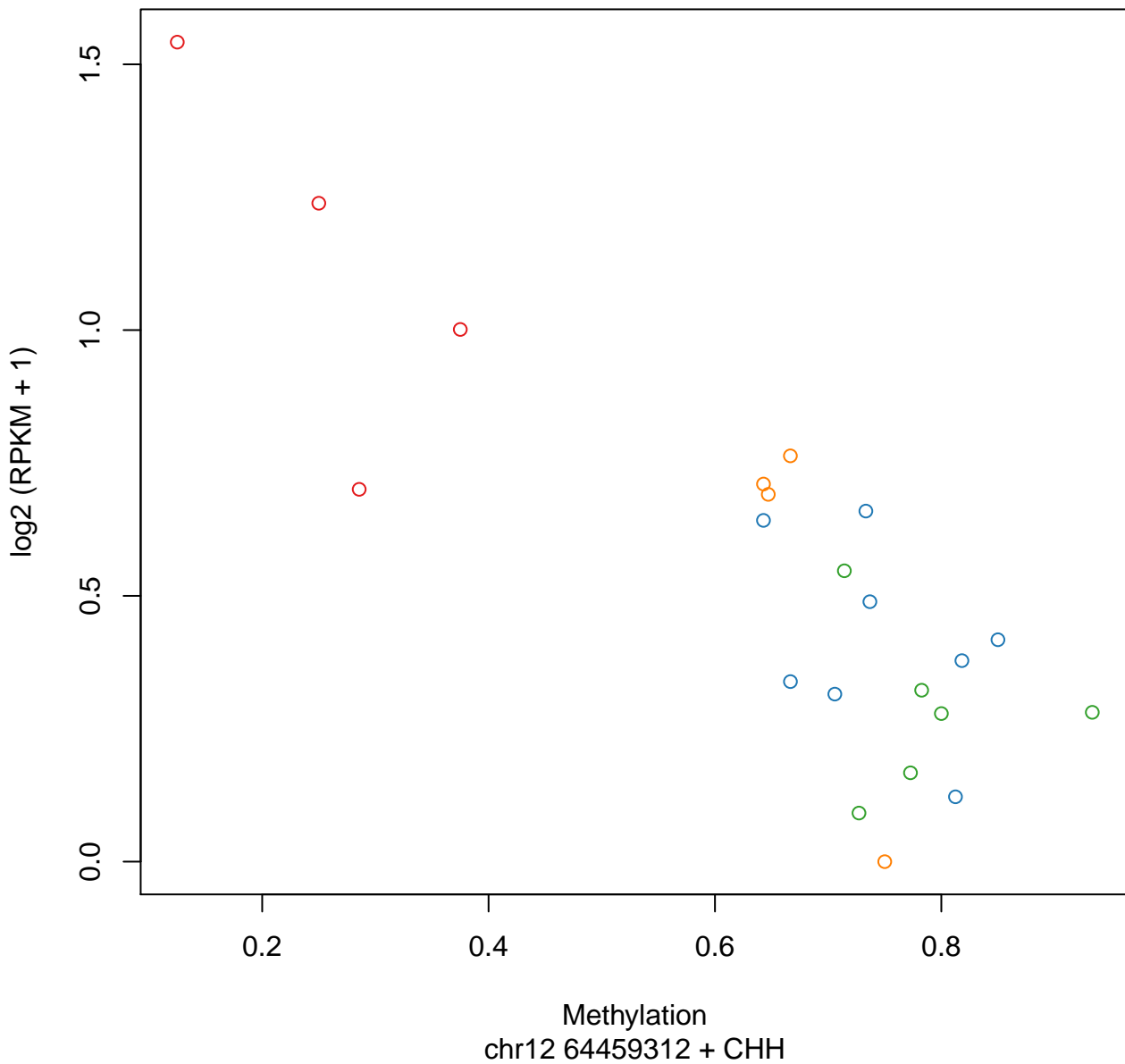


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TNC

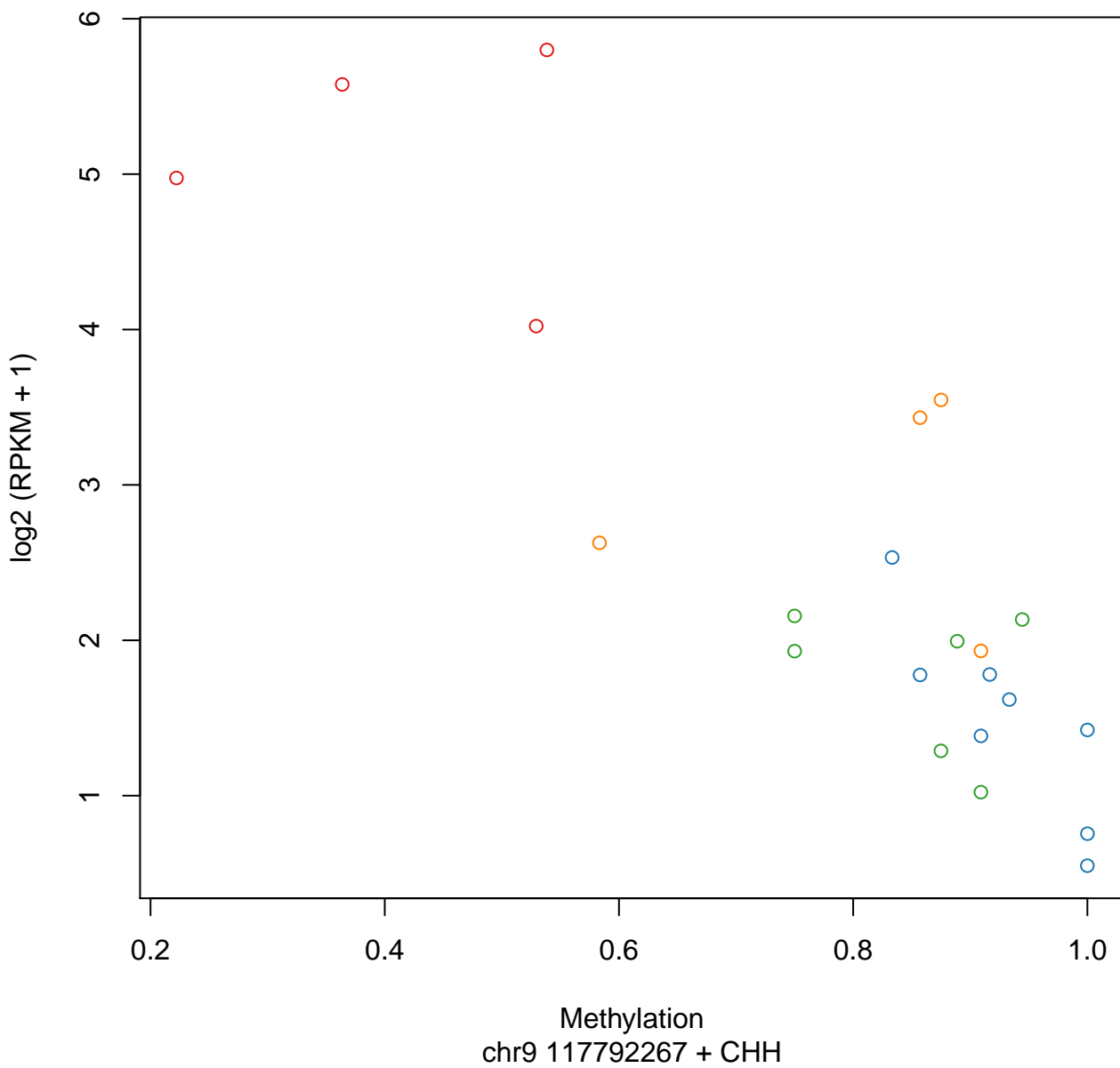


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SRGAP1



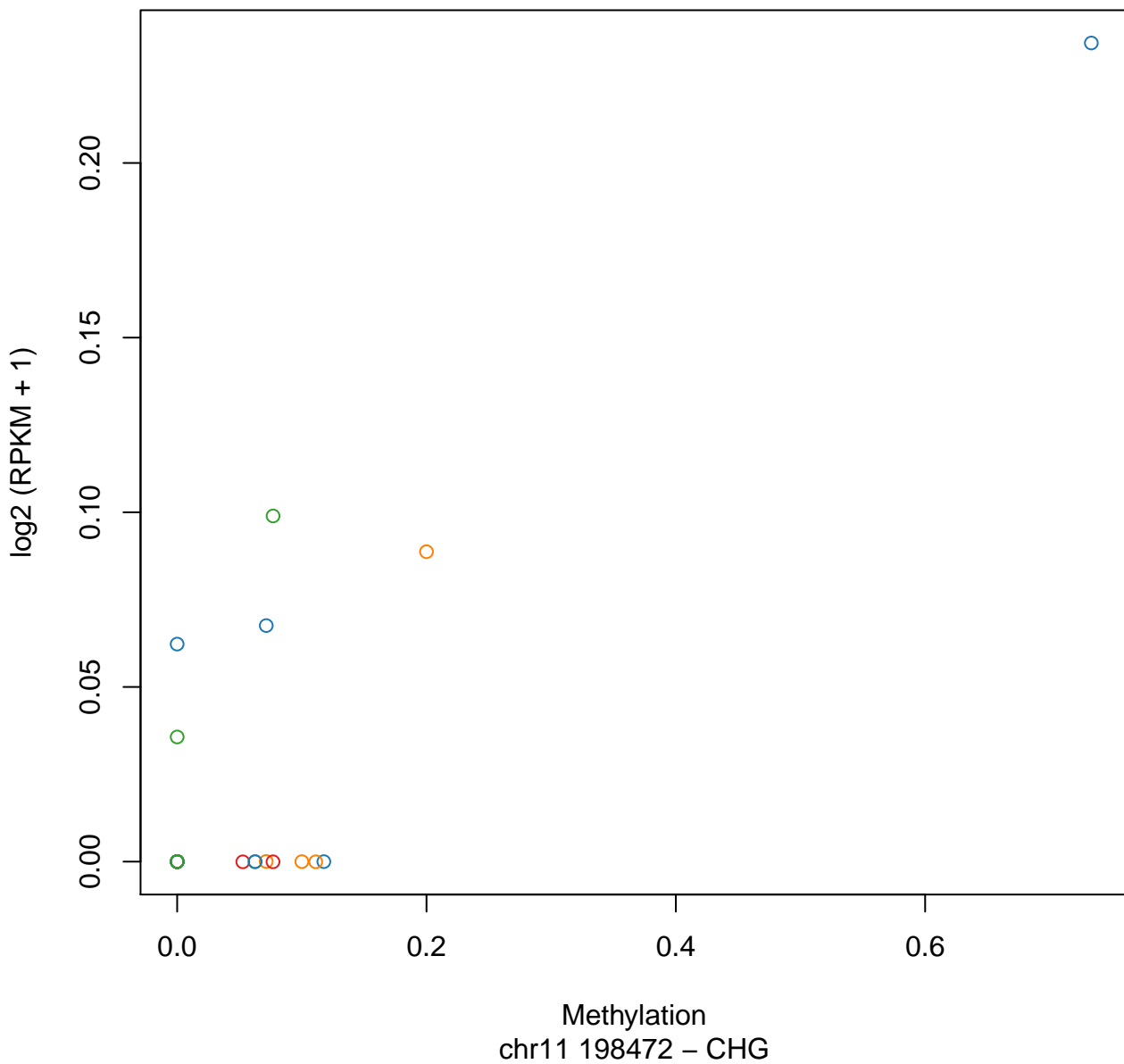
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TNC



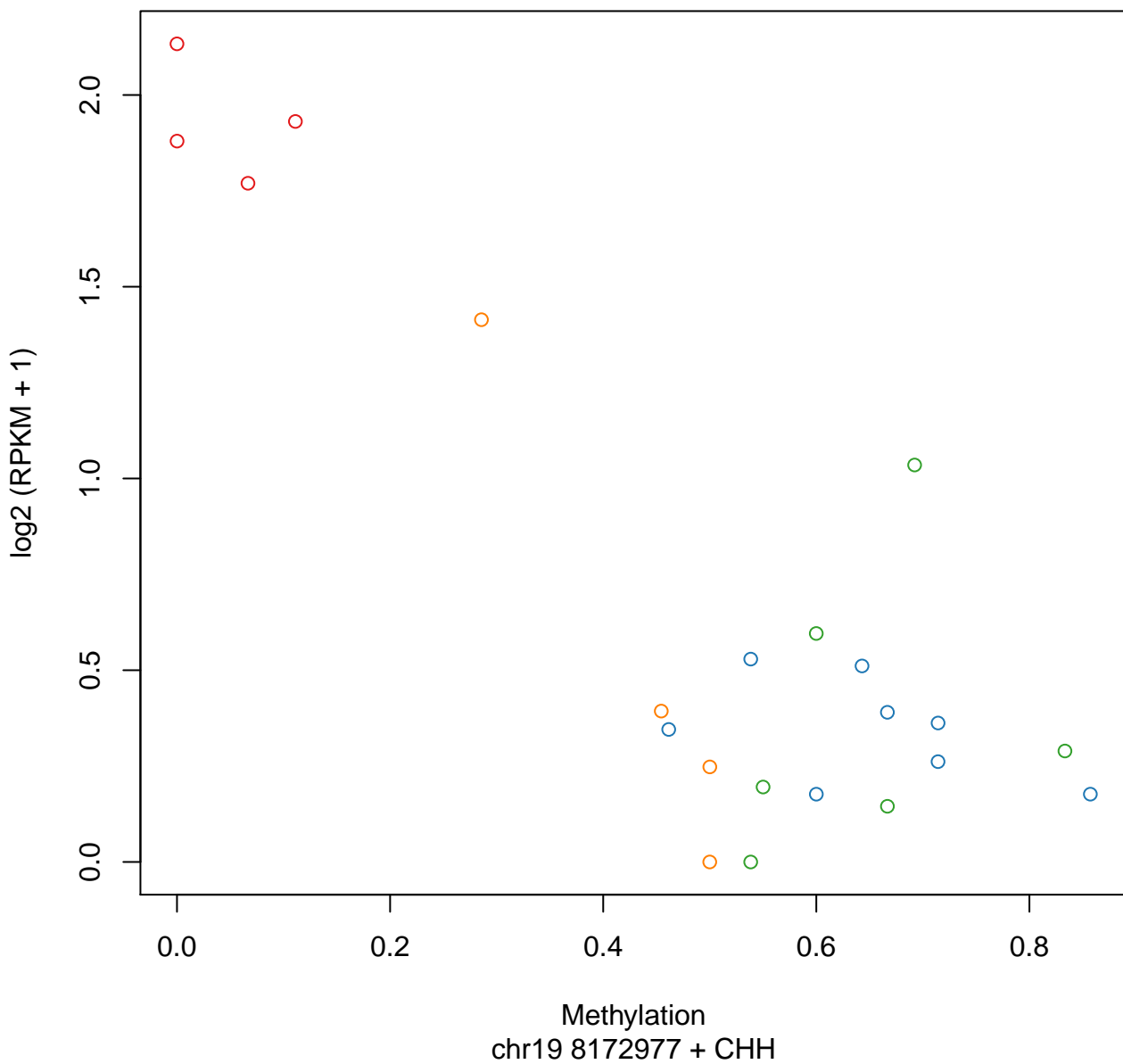
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ODF3



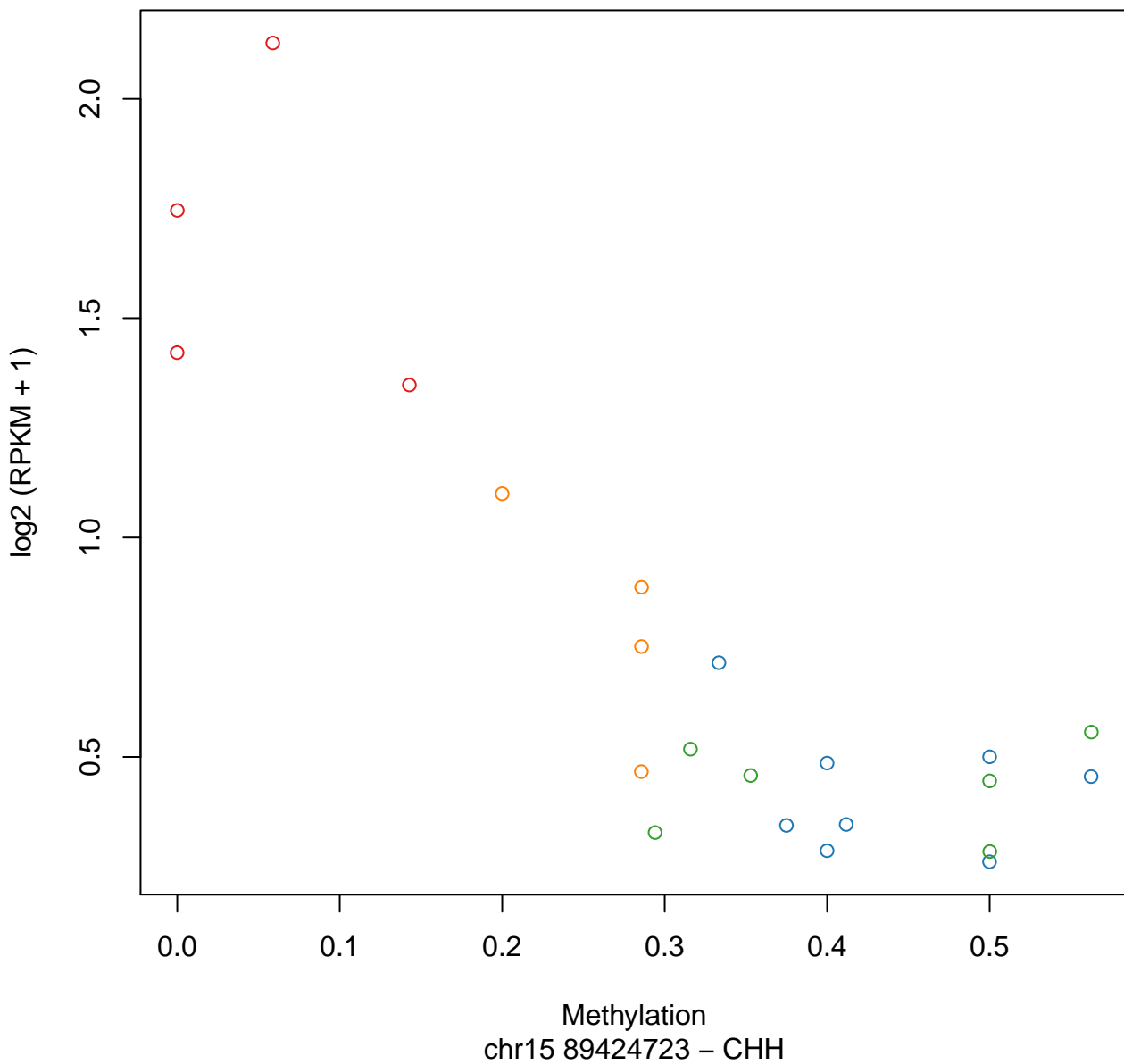
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FBN3

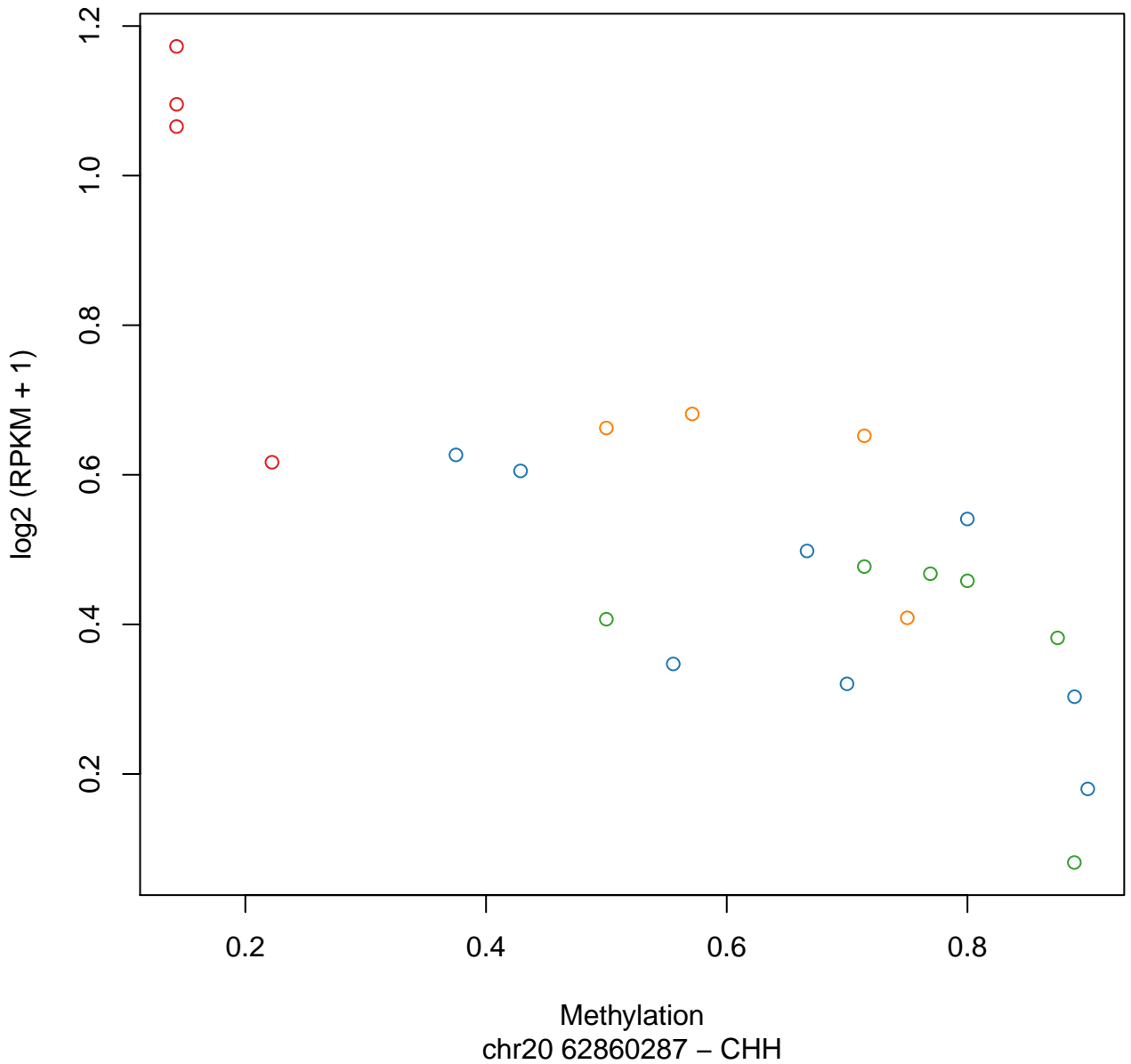


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HAPLN3

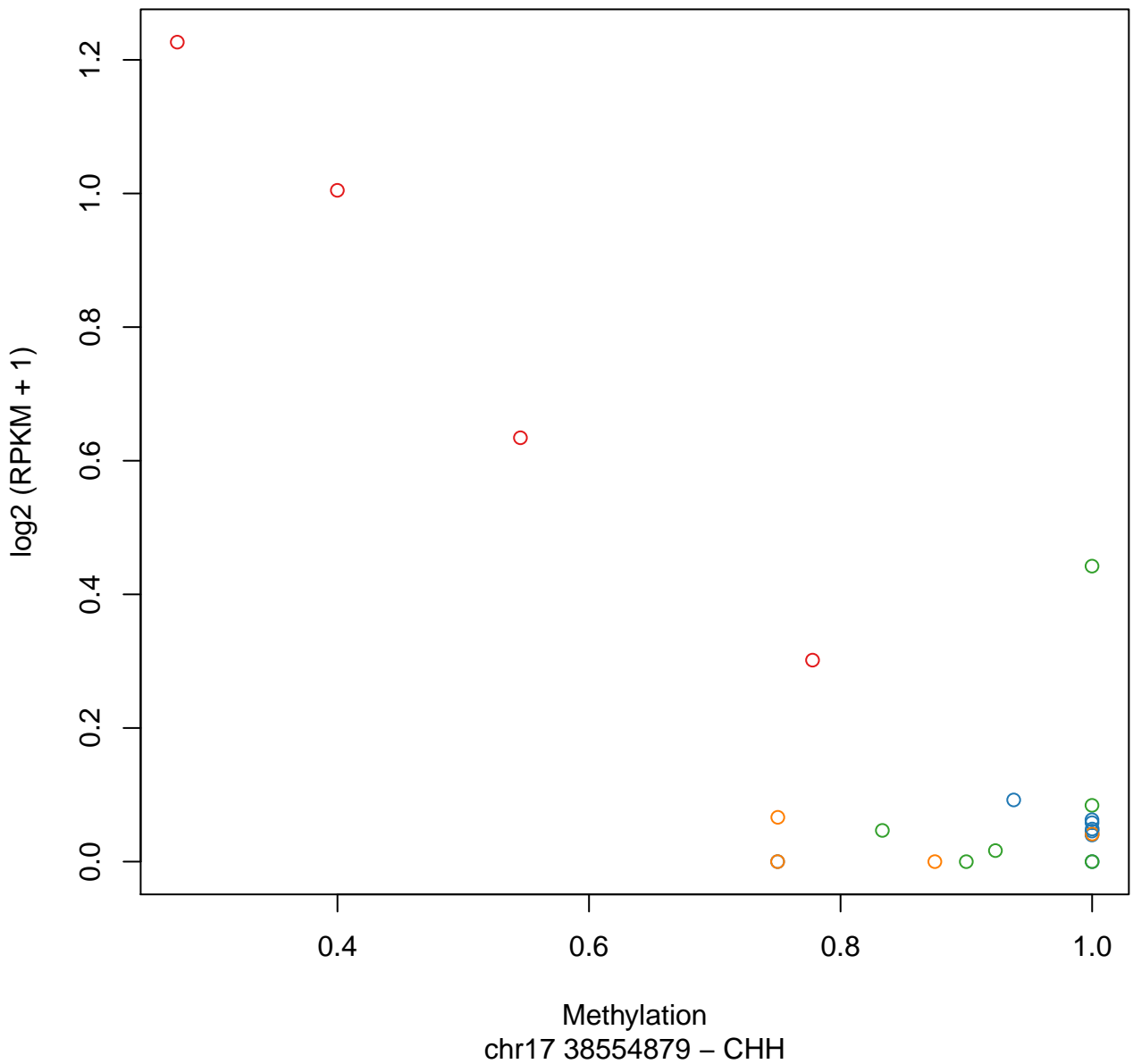


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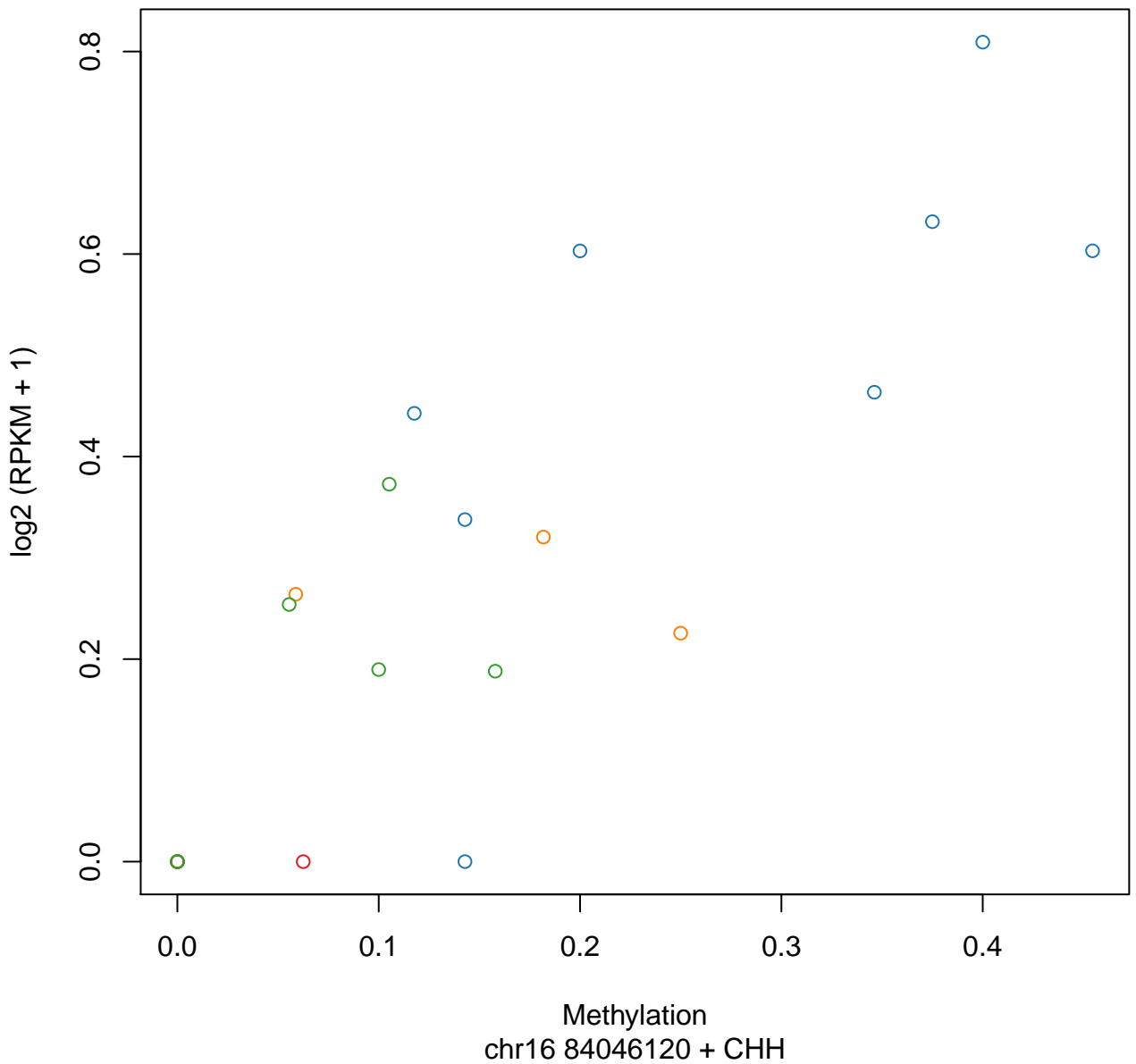


exon ENSE00002686808.1_1 FDR 0.00257

TOP2A



exon ENSE00003491917.1_1 FDR 0.00259
SLC38A8



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ASF1B

