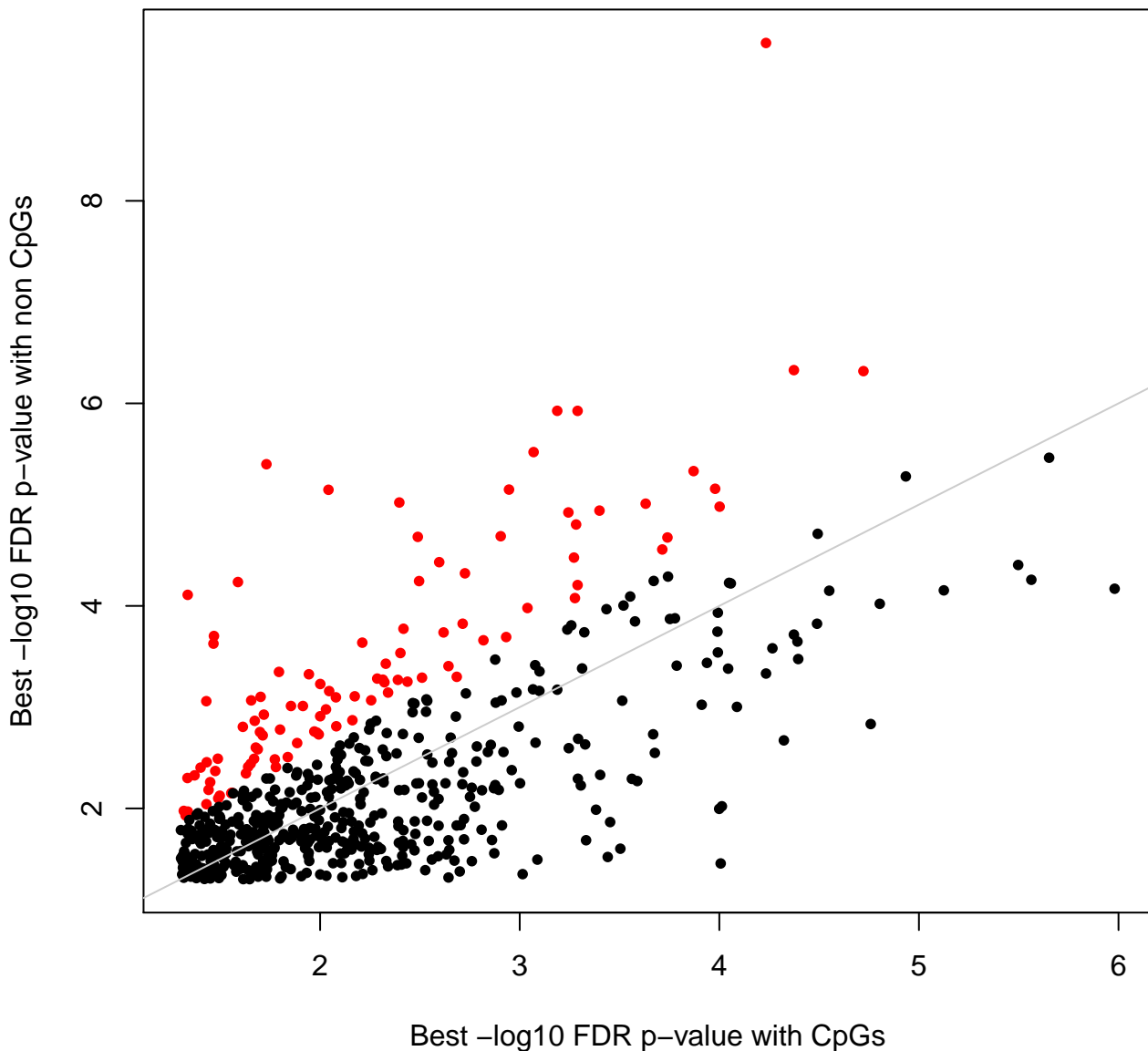
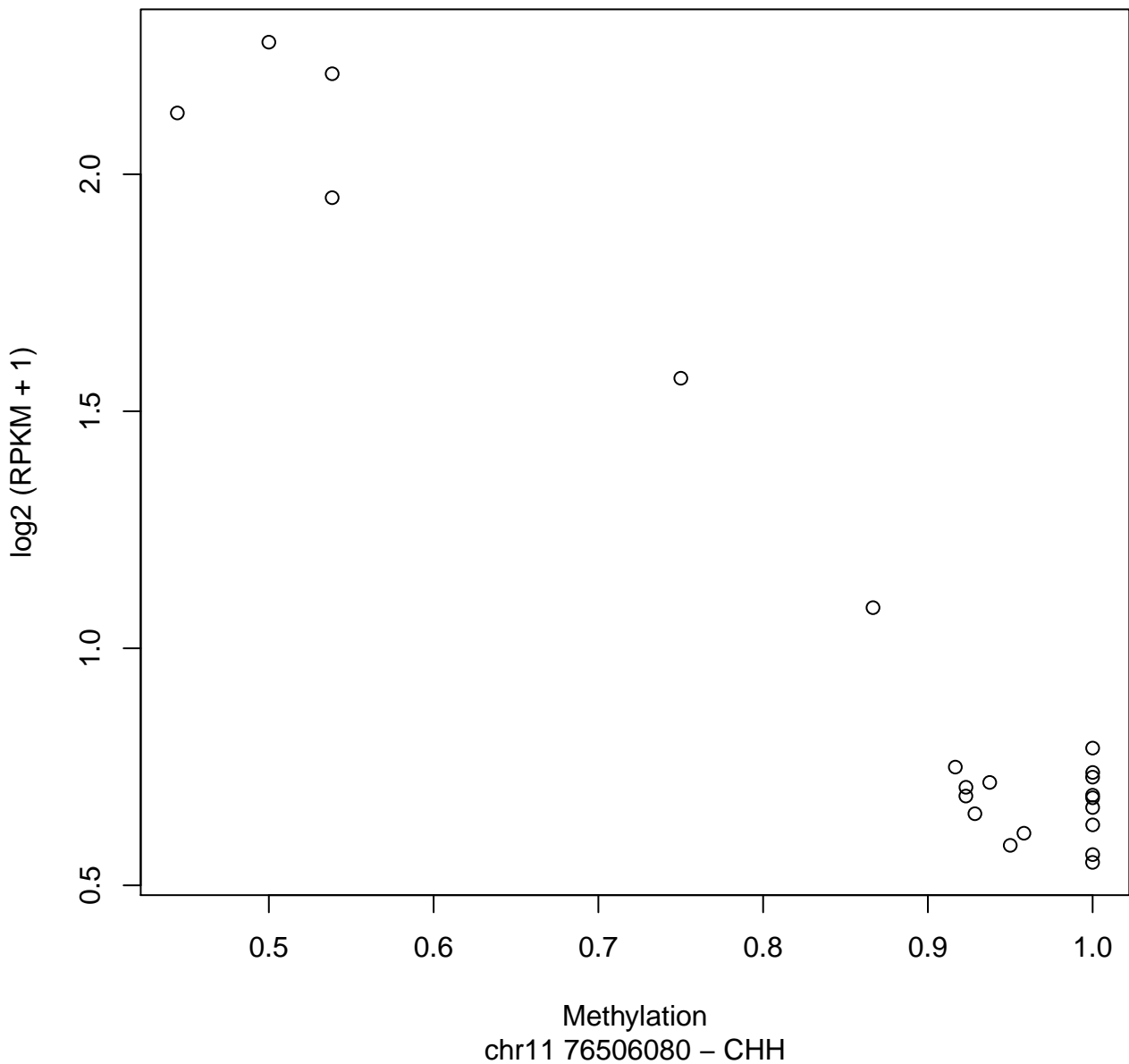


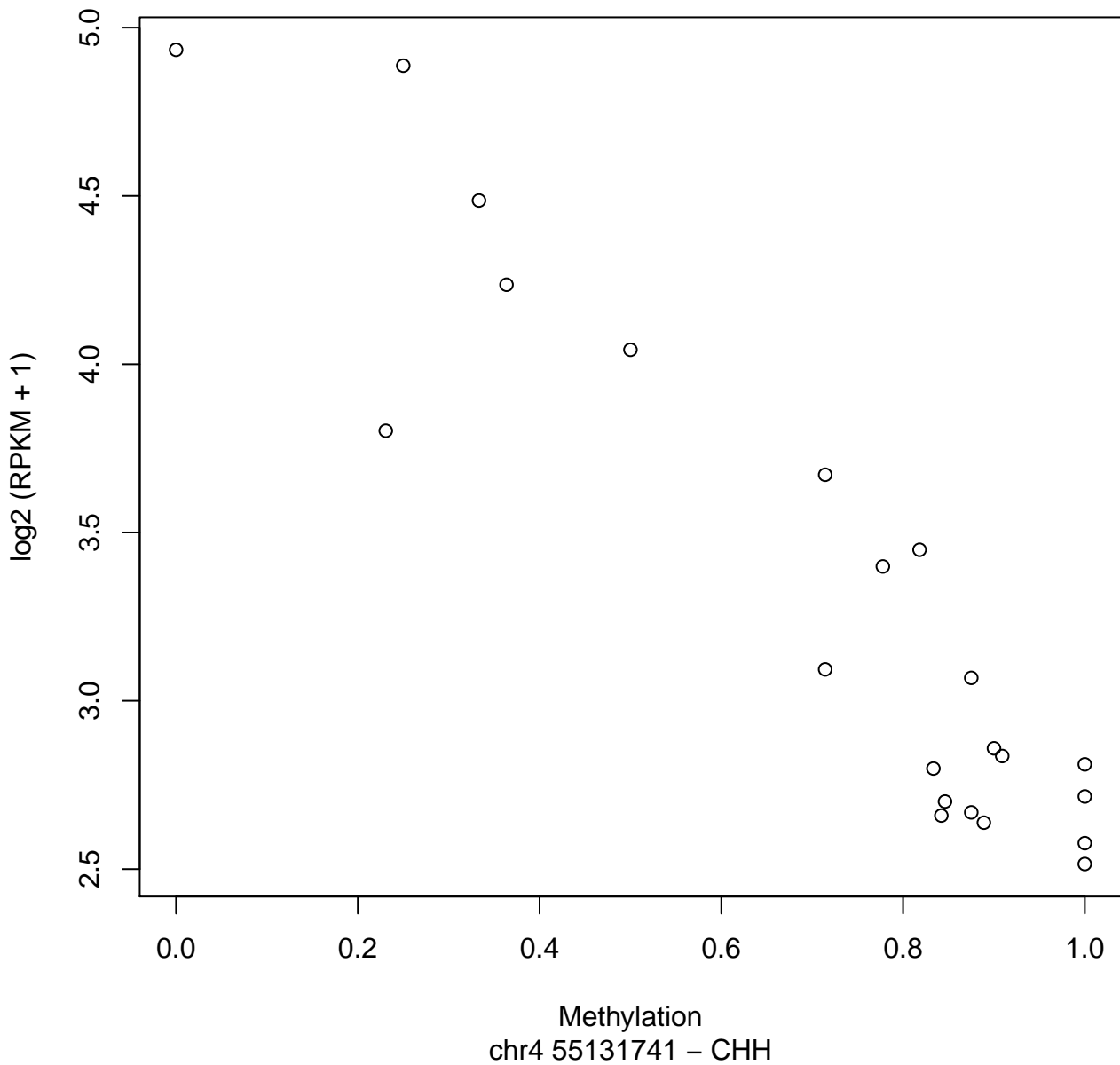
567 common gene meQTLs
Based on top 3473 gene s expressed in Neurons



gene ENSG00000182704.7_2 FDR 2.76e-10
TSKU

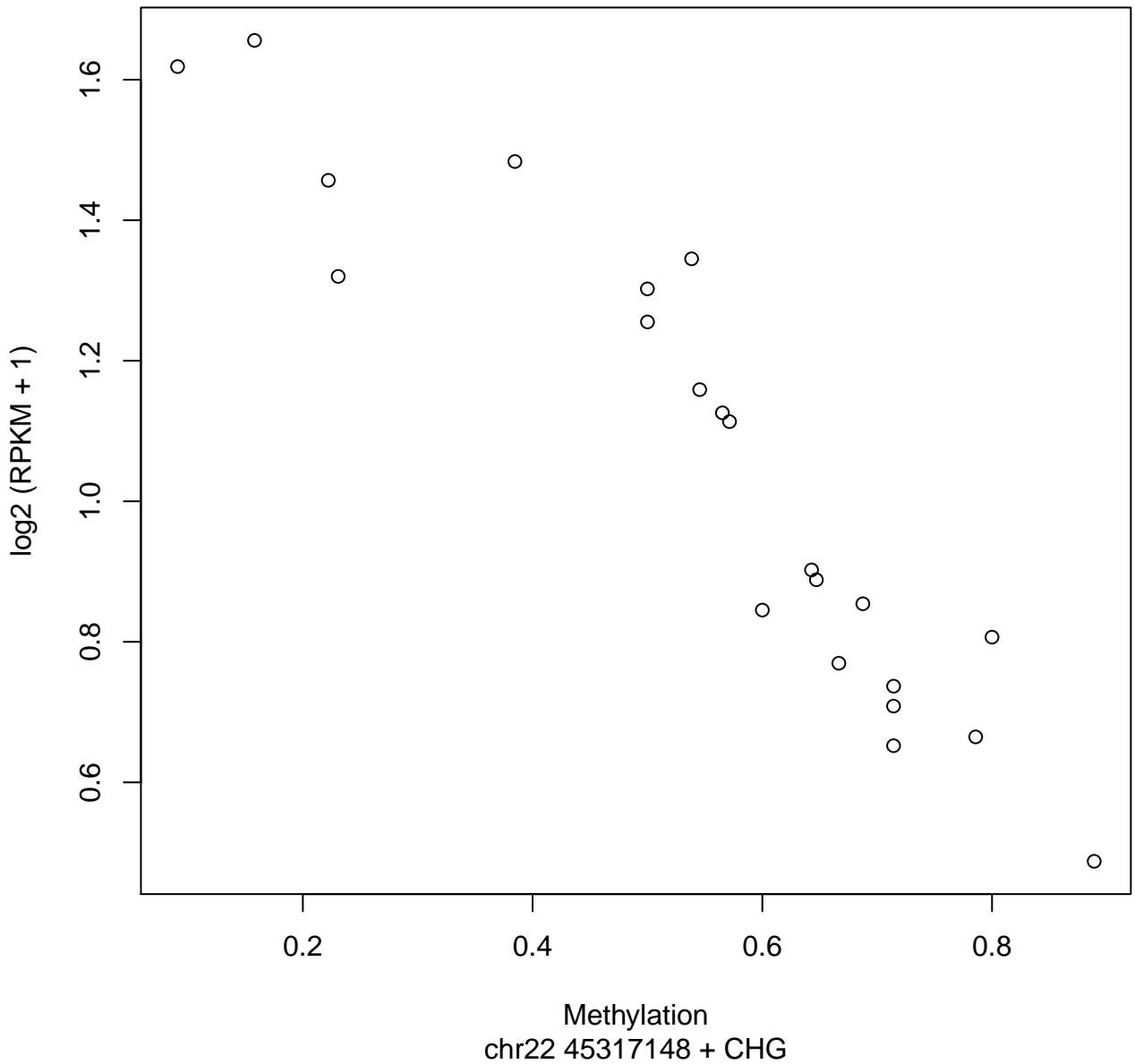


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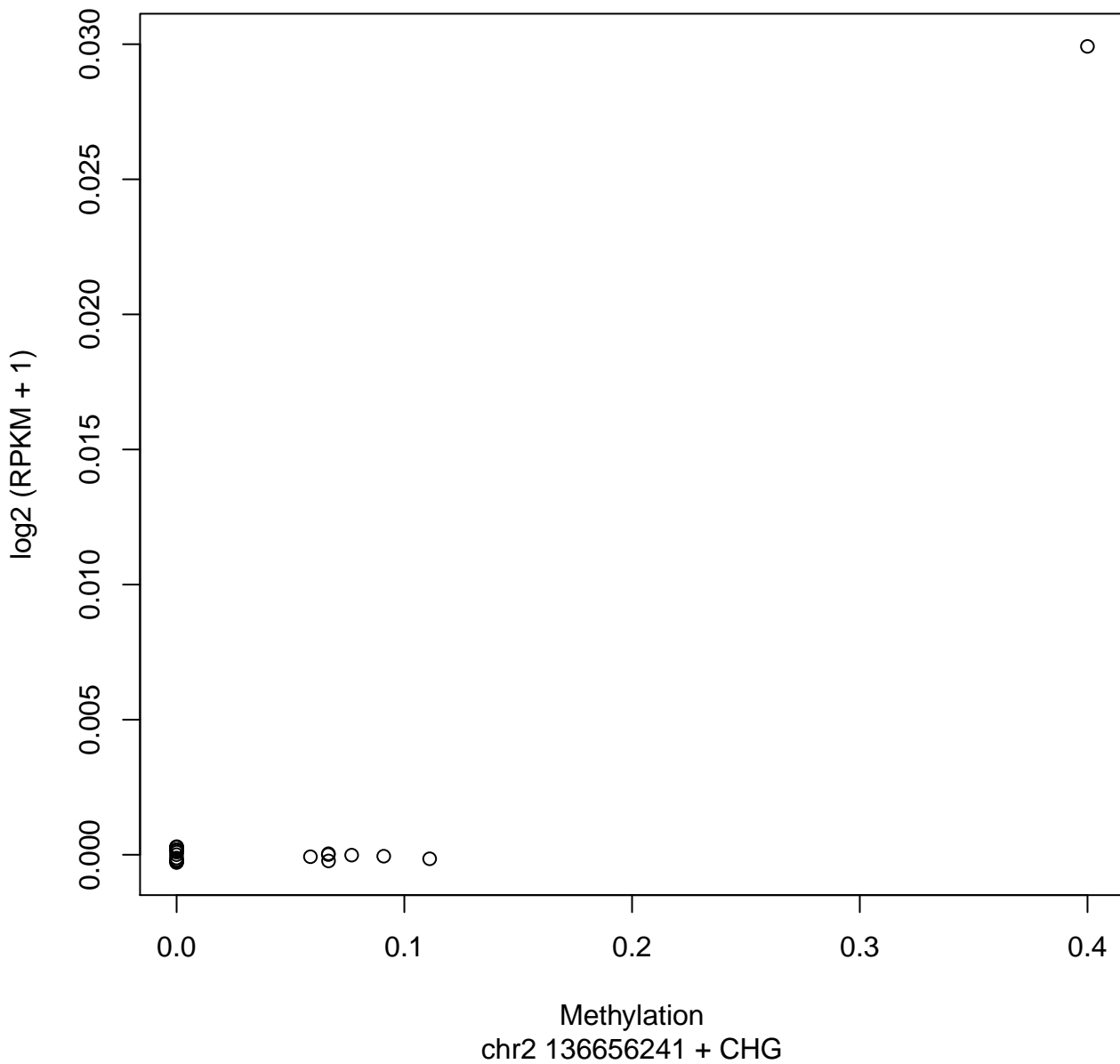


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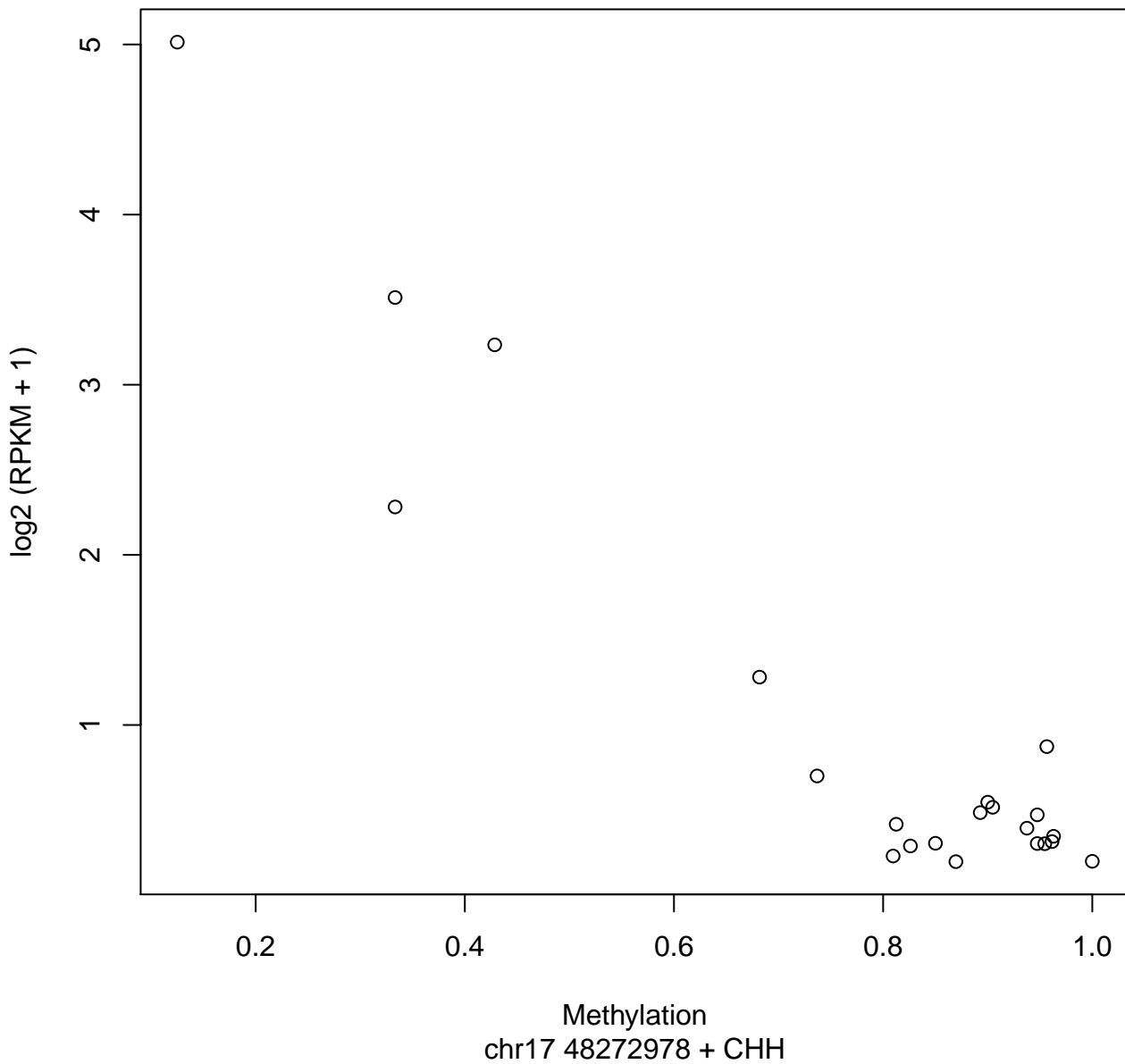
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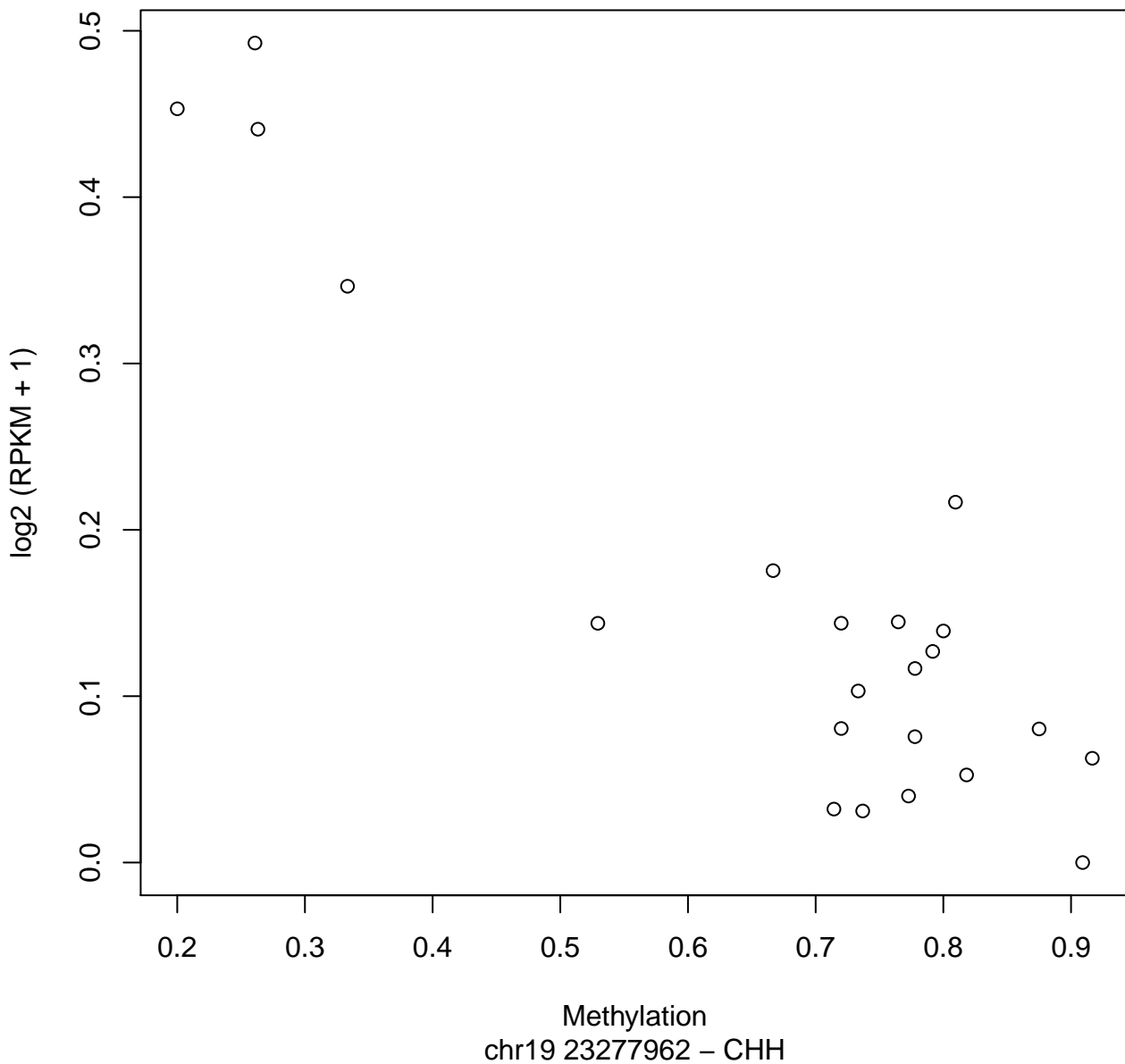
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AC011999.1



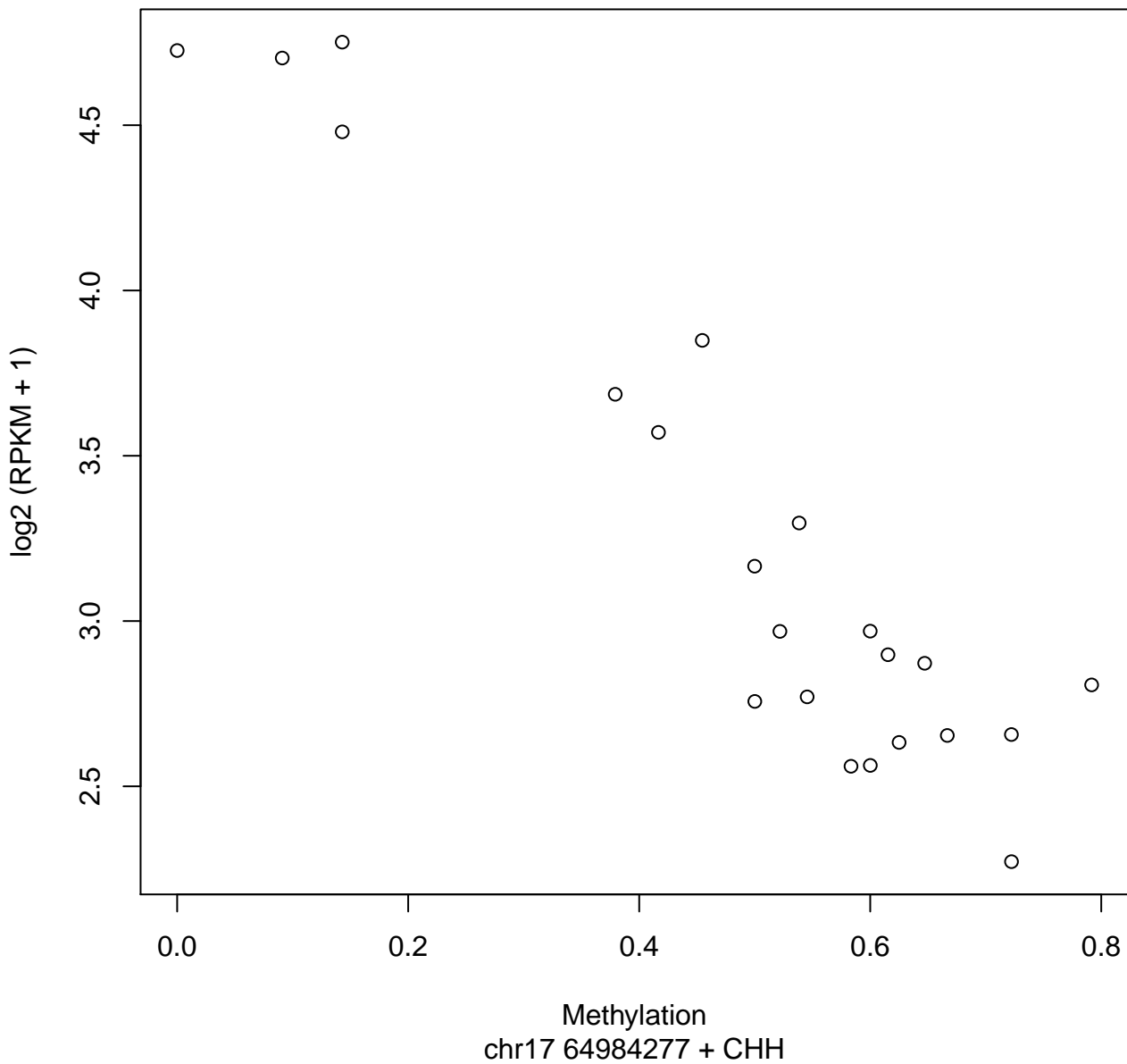
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COL1A1



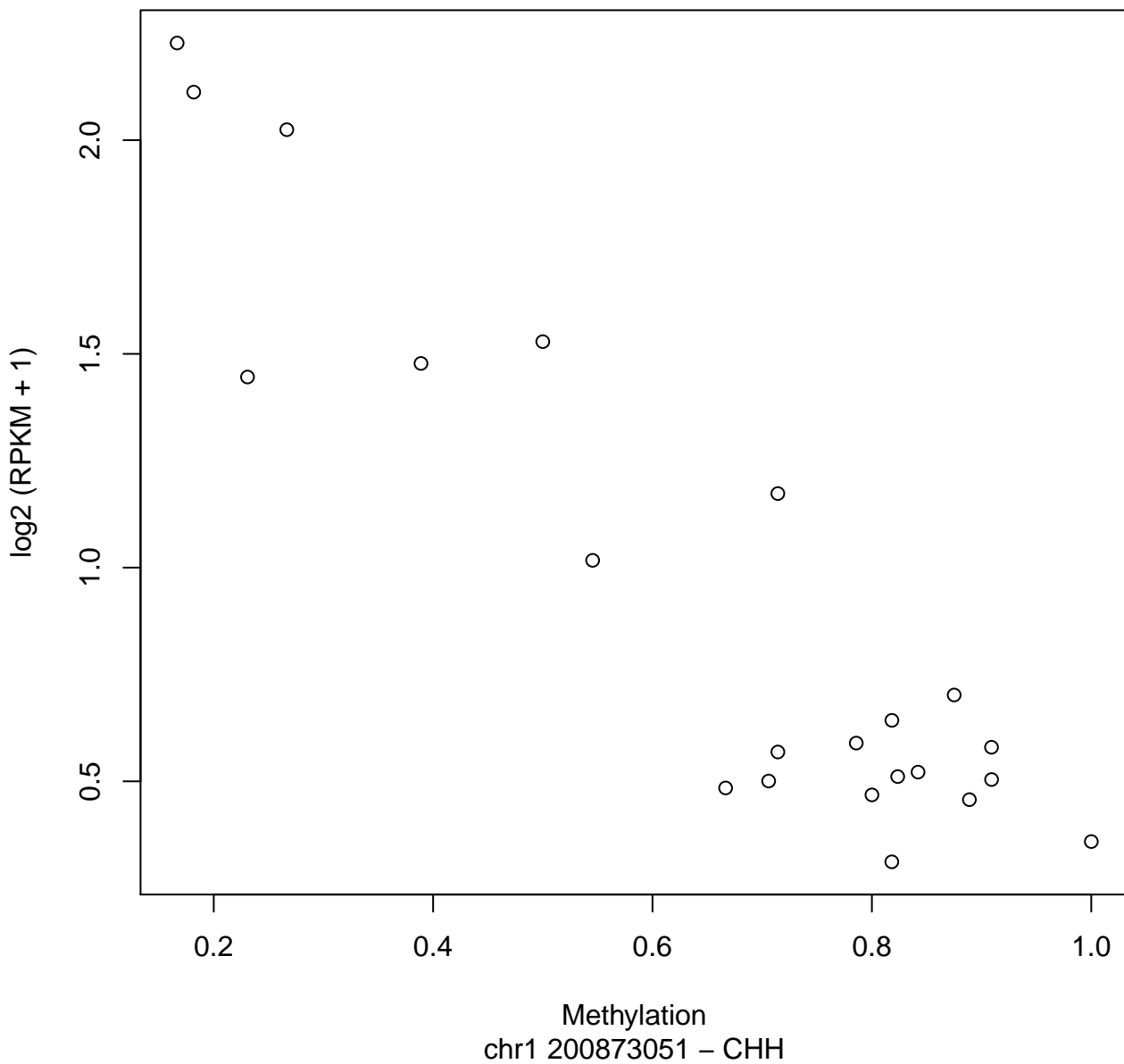
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CTD-2291D10.4



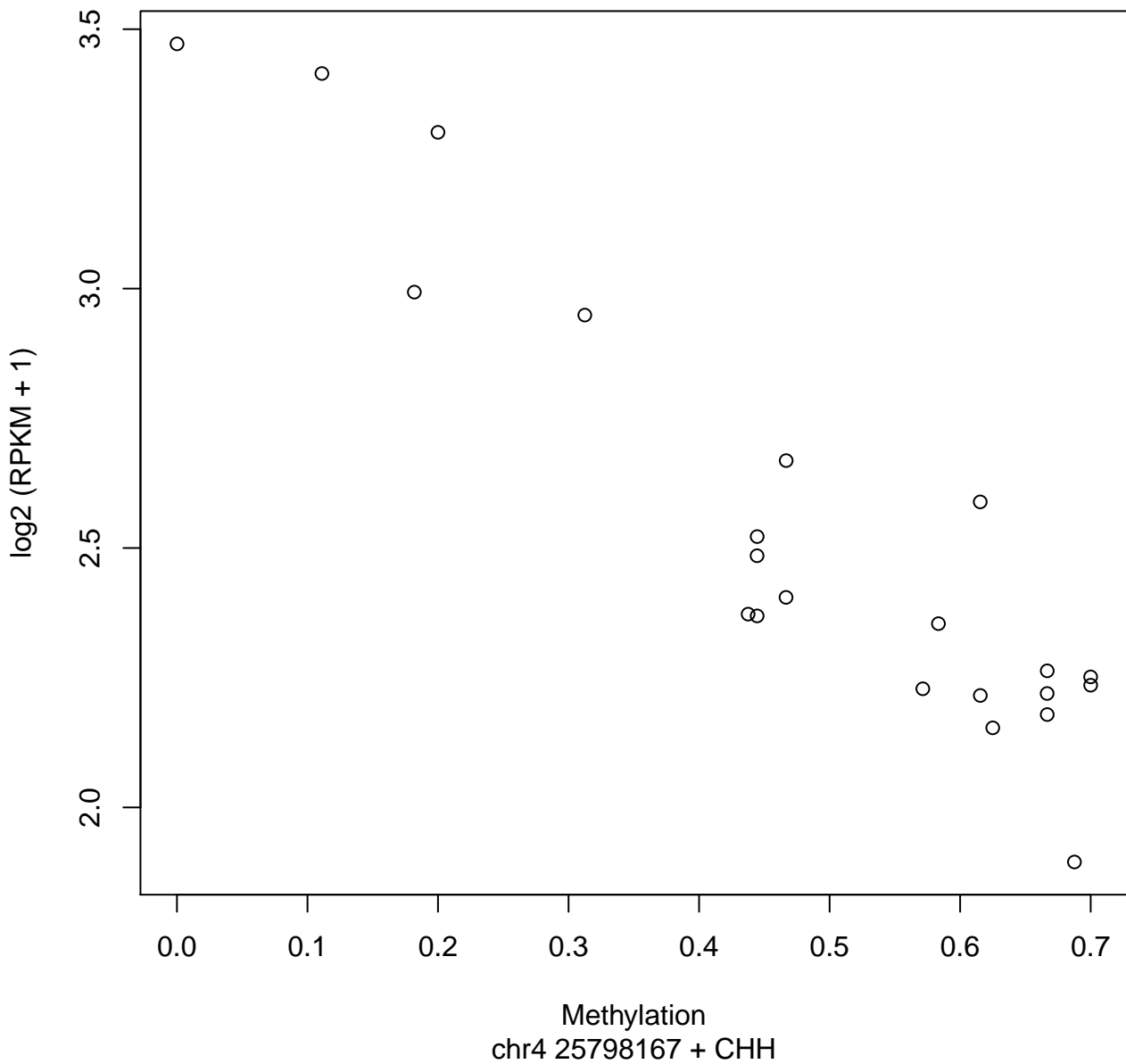
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gene ENSG00000163362.10_2 FDR 9.52e-06
C1orf106

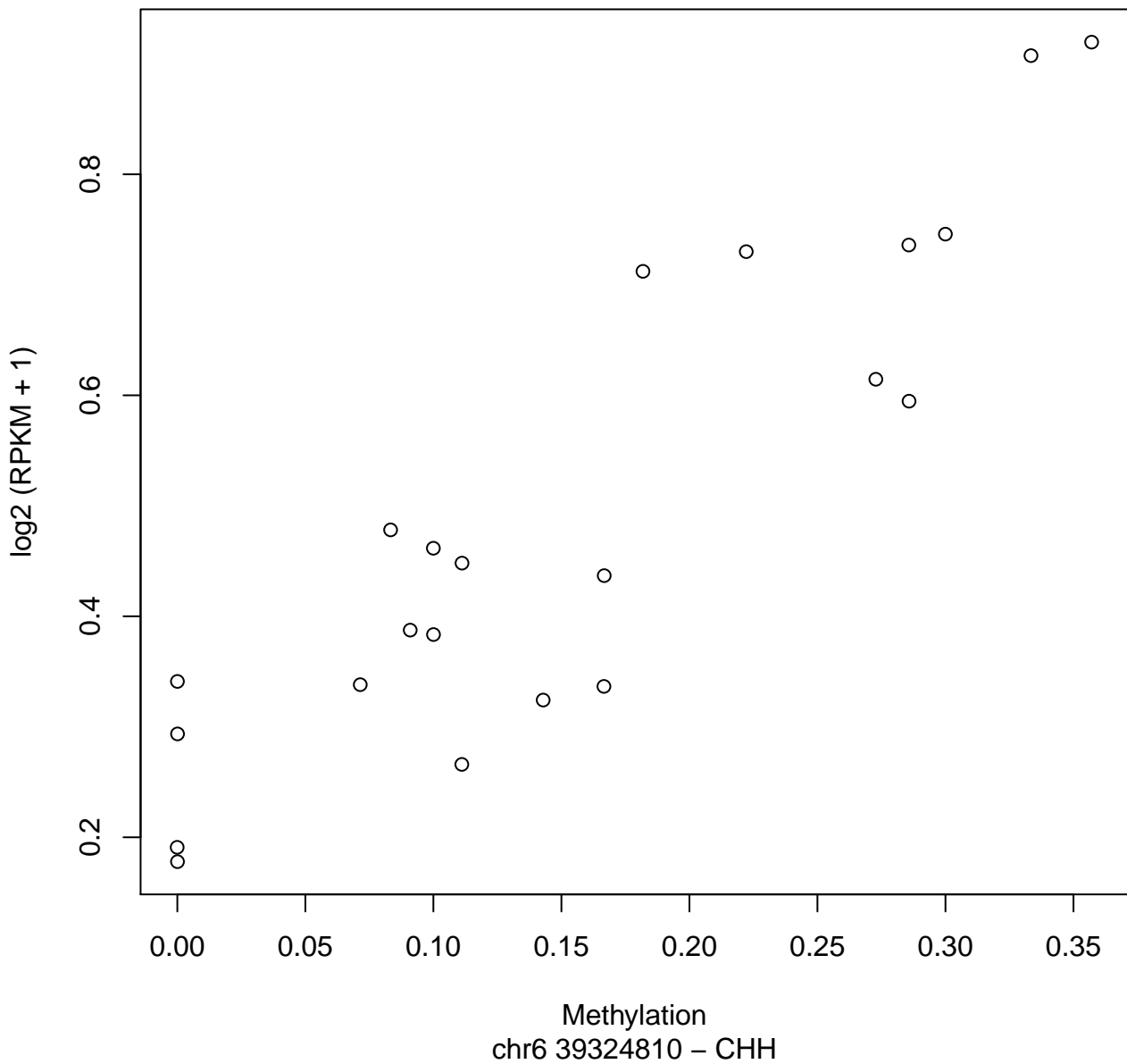


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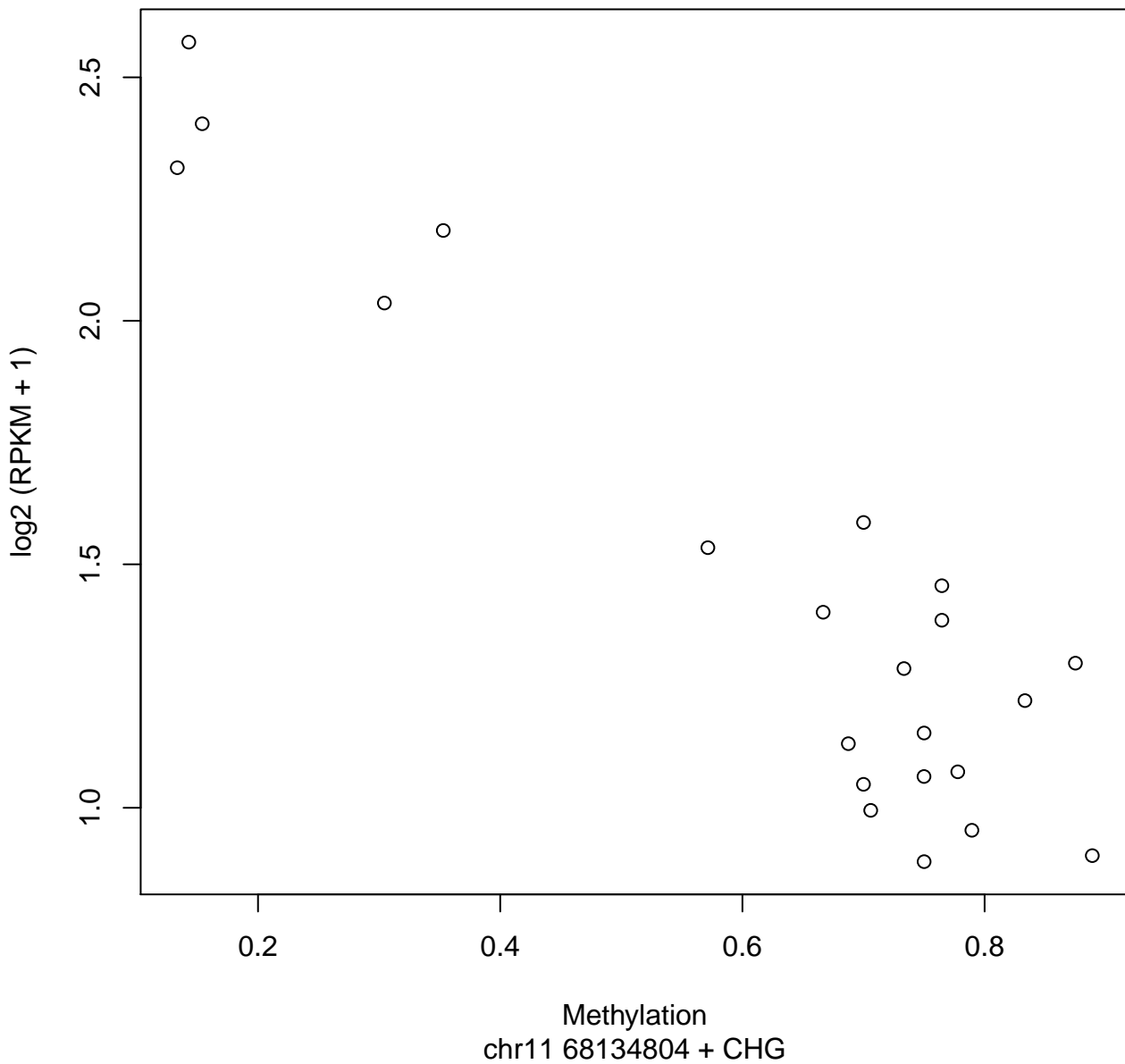


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KIF6

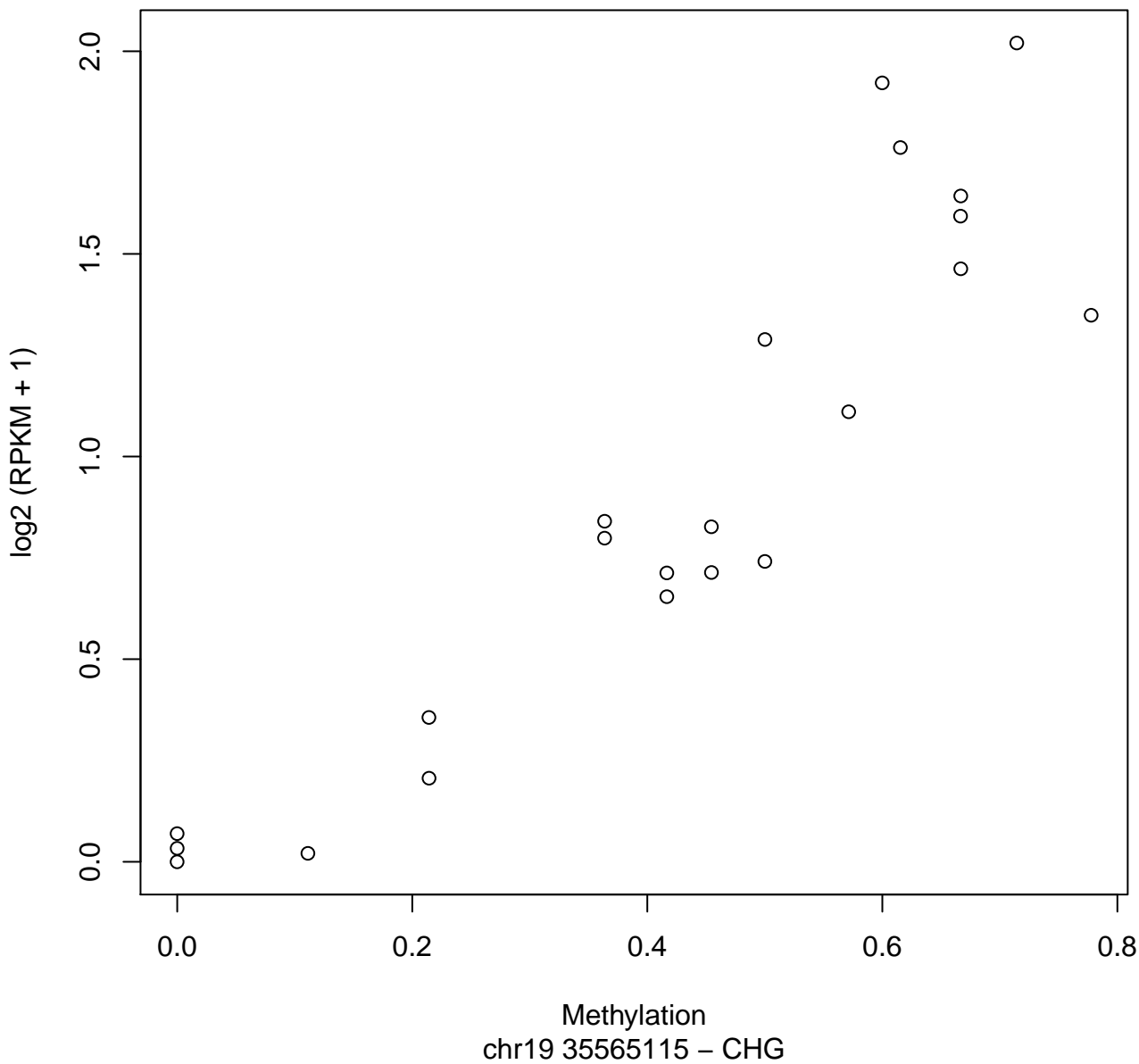


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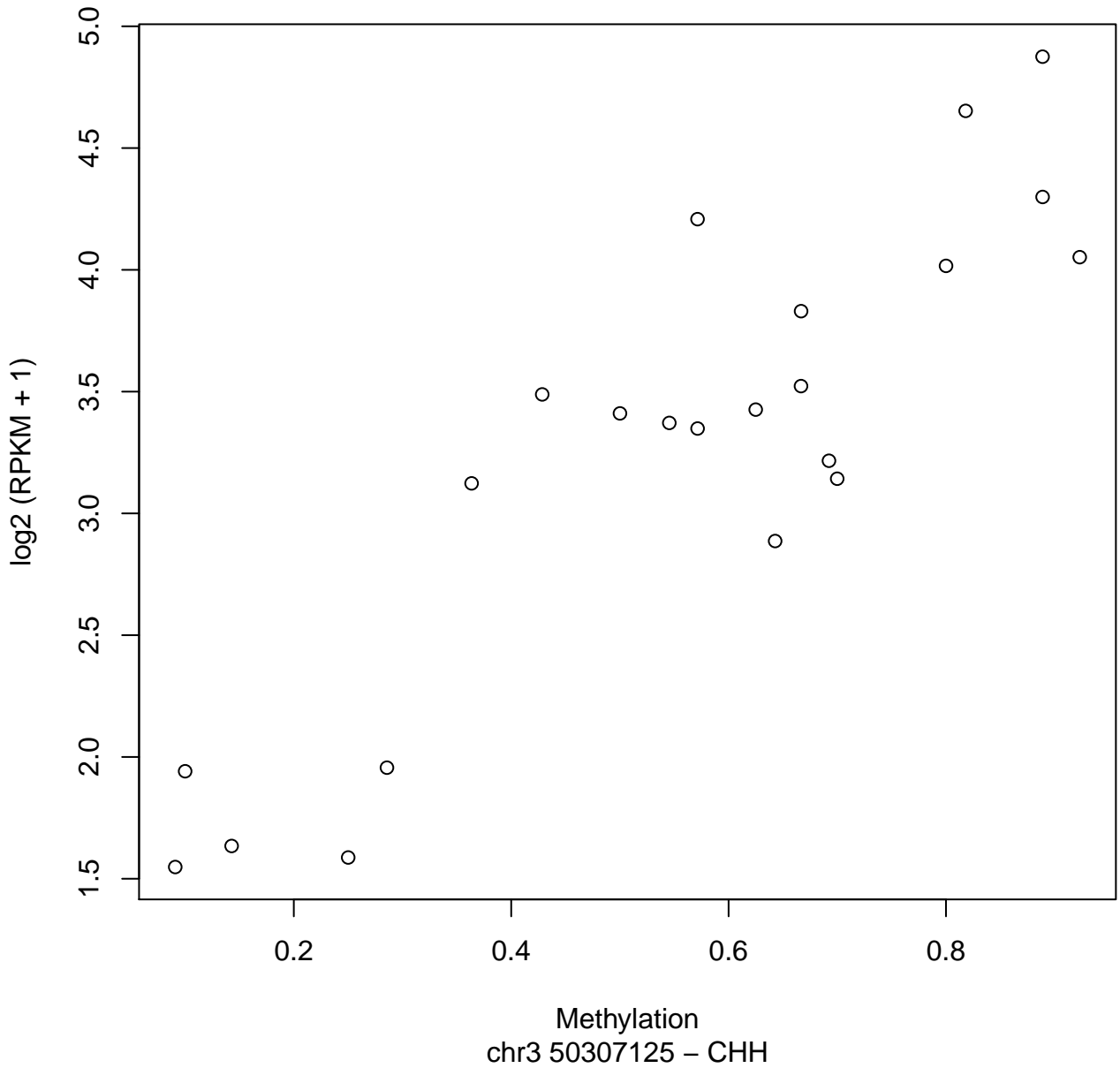
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HPN-AS1



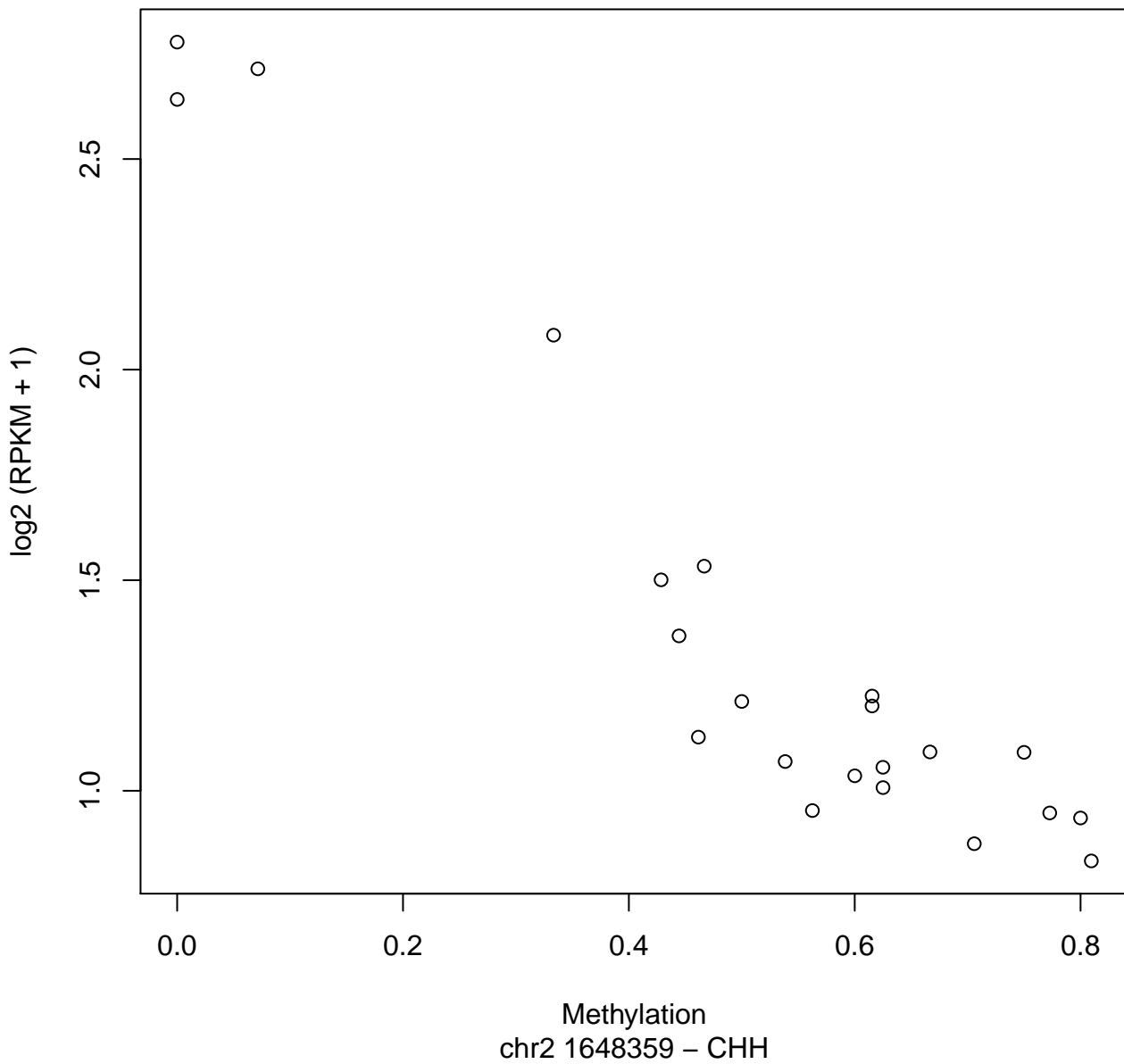
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SEMA3B

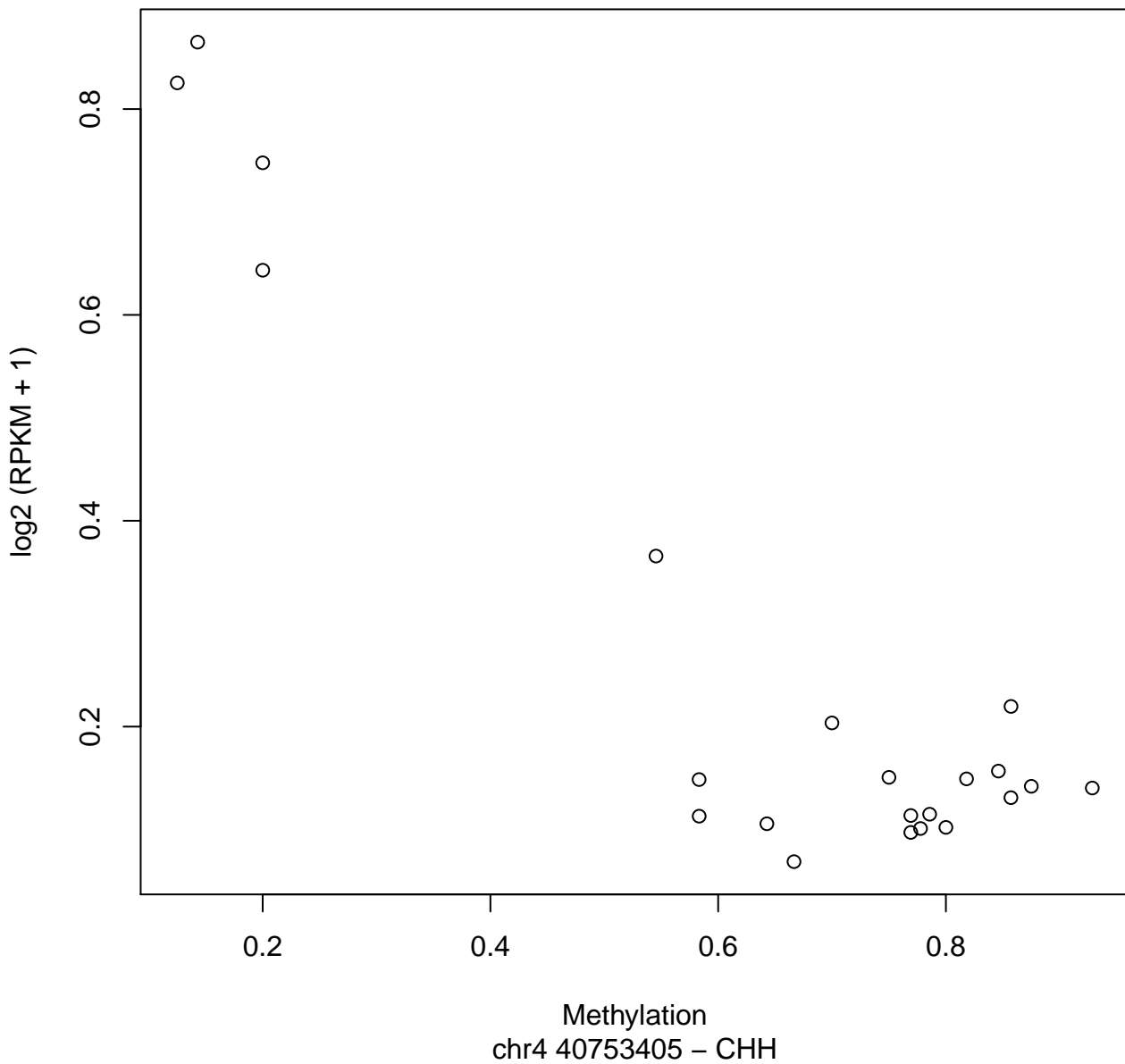


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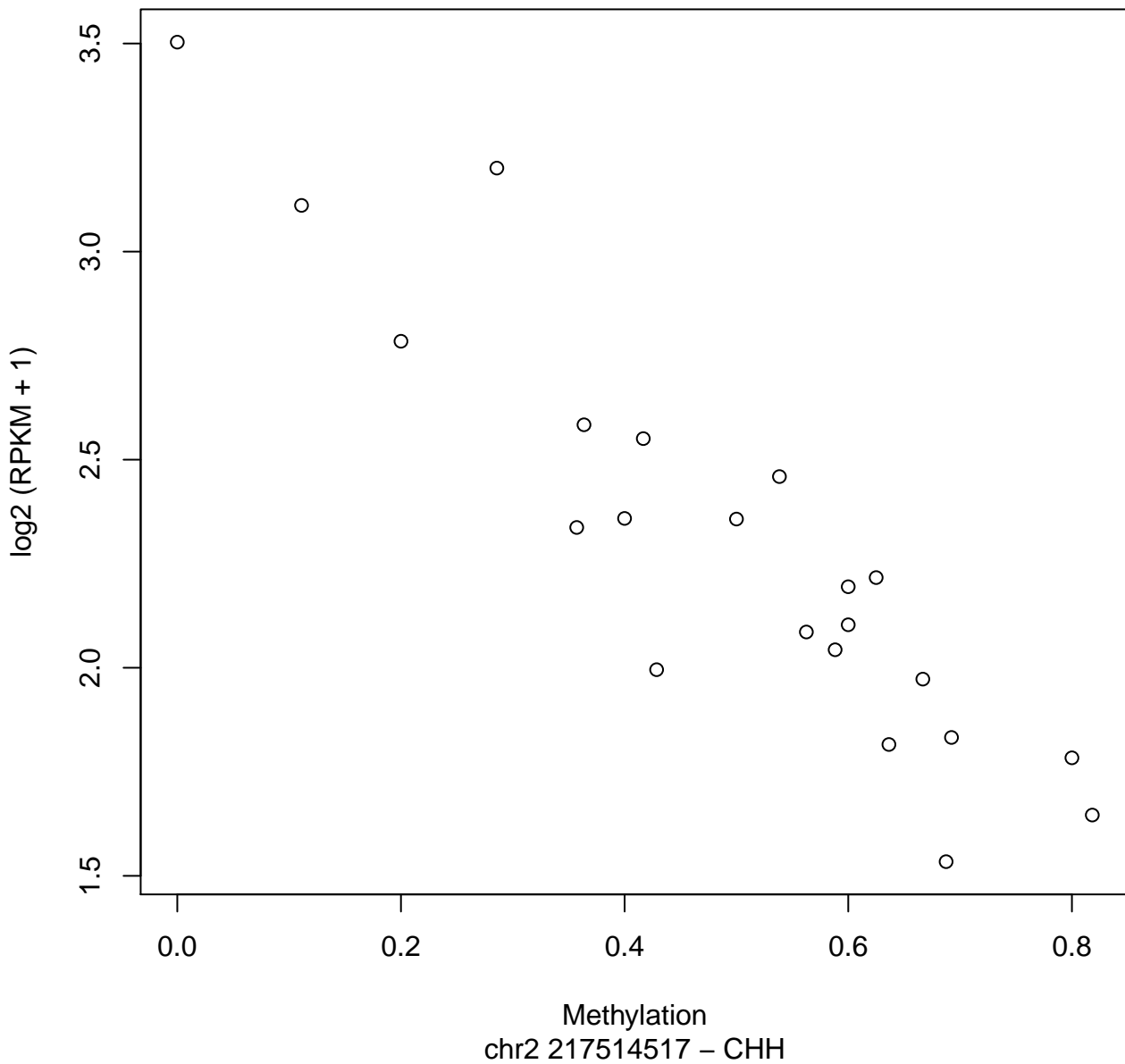
PXDN



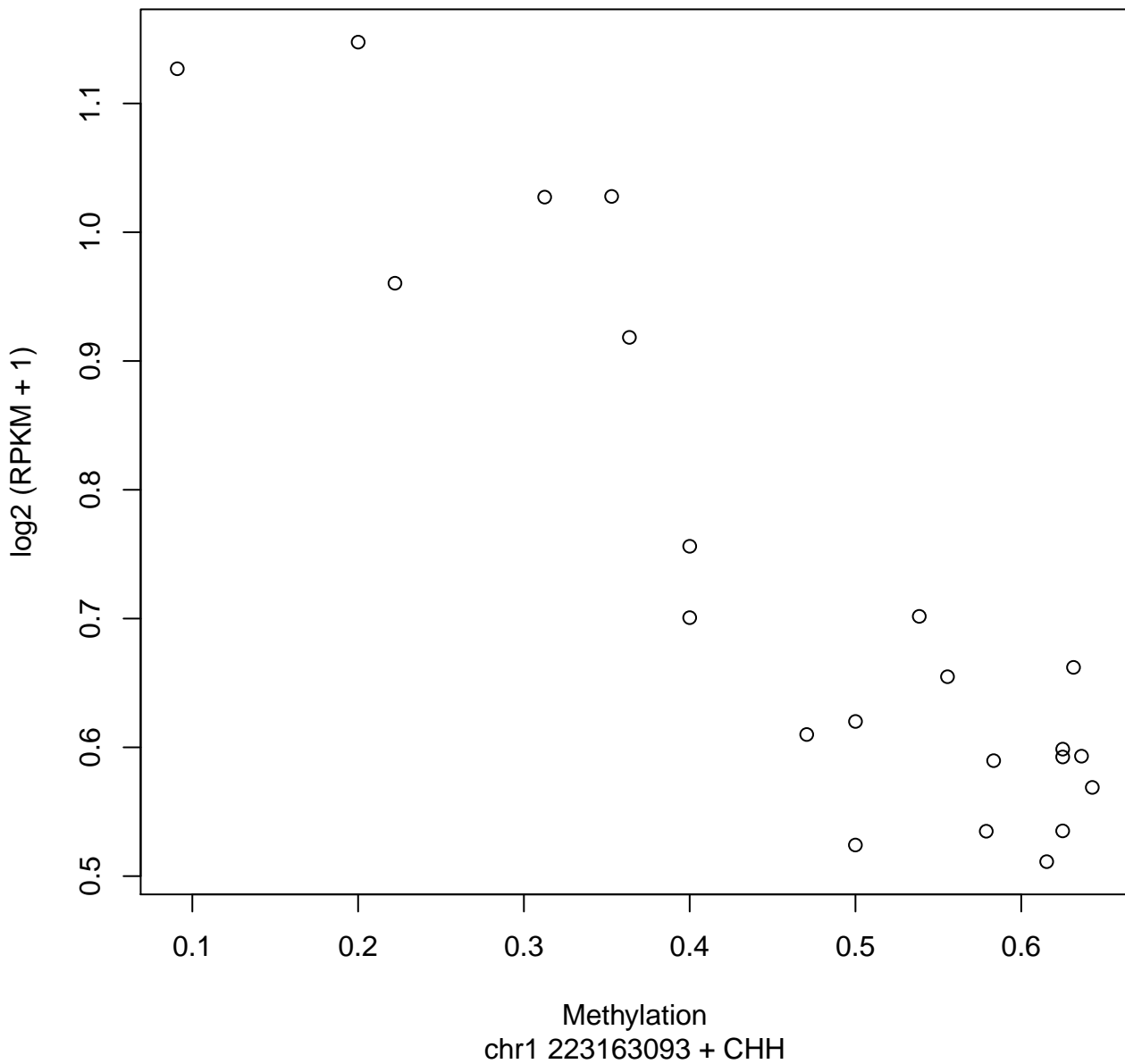
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gene ENSG00000115457.9_2 FDR 2.05e-05
IGFBP2

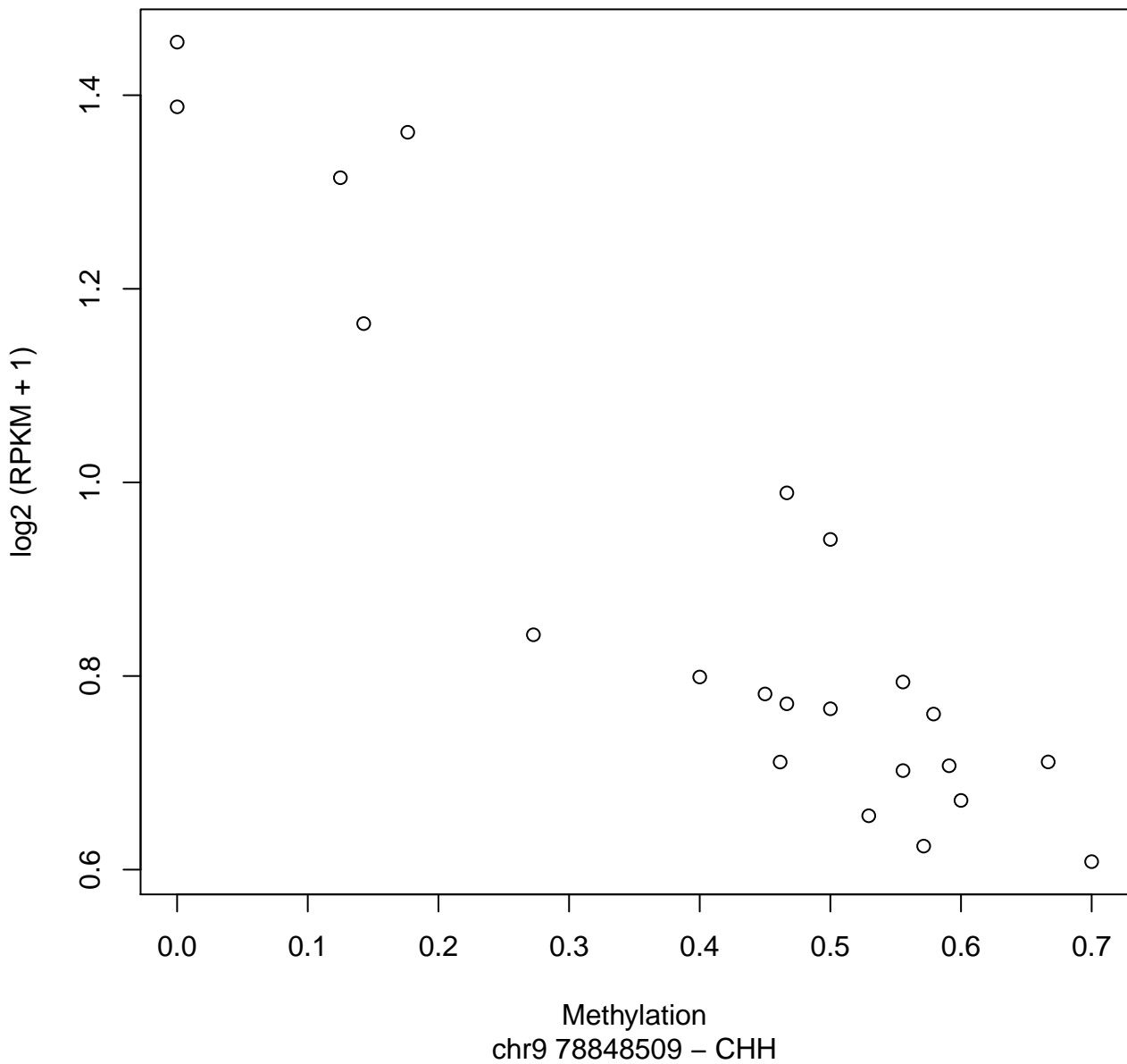


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DISP1



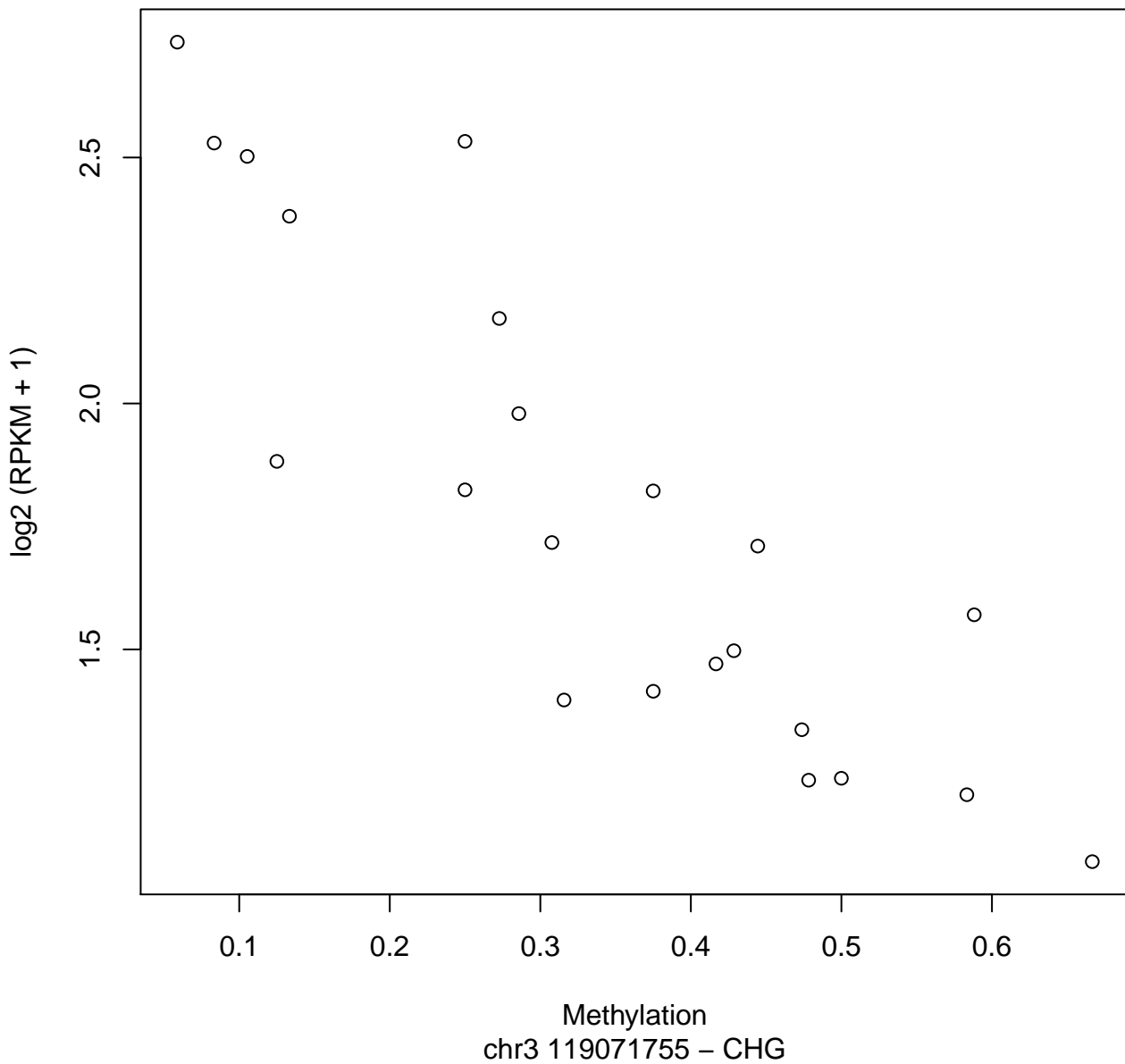
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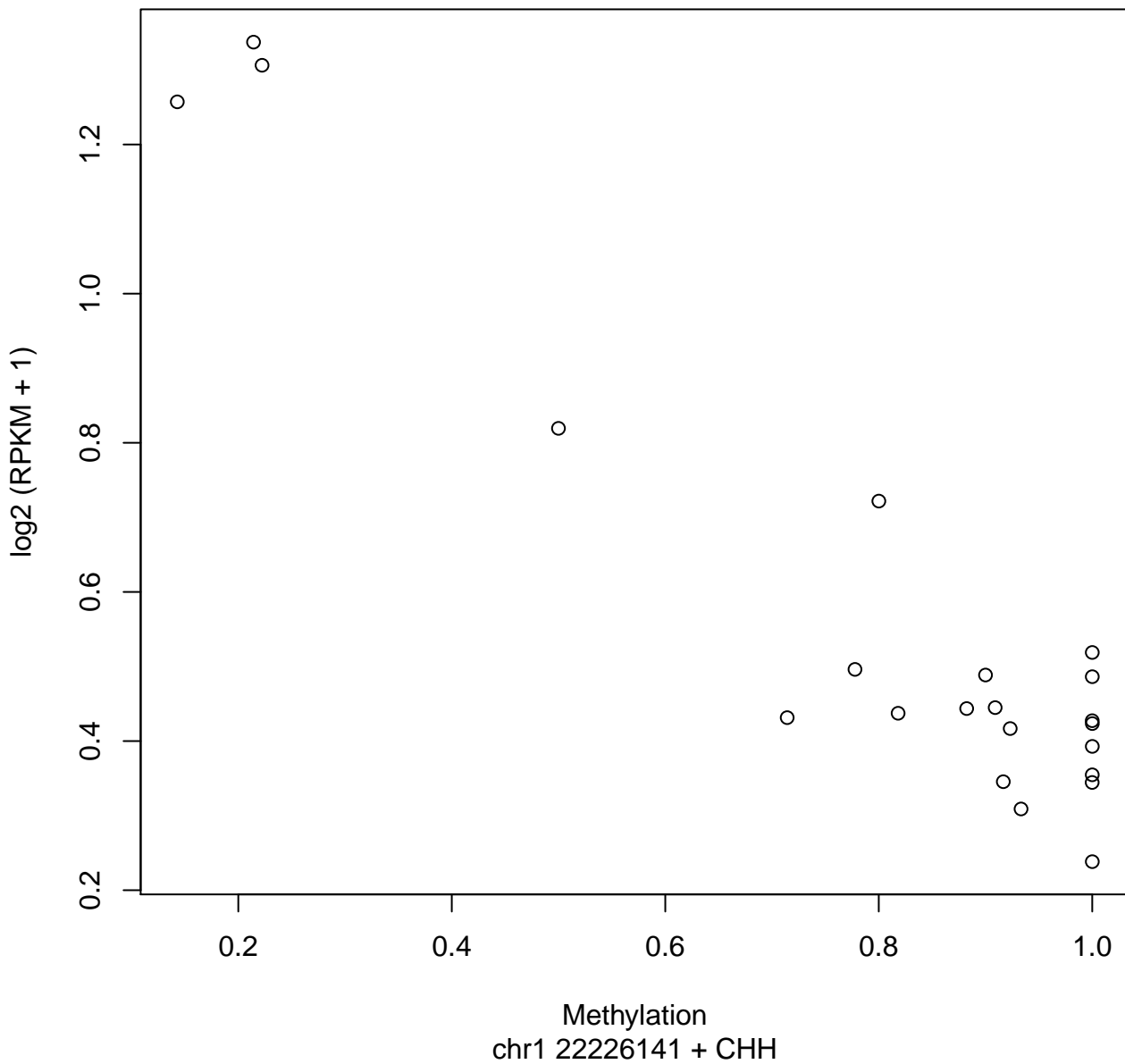


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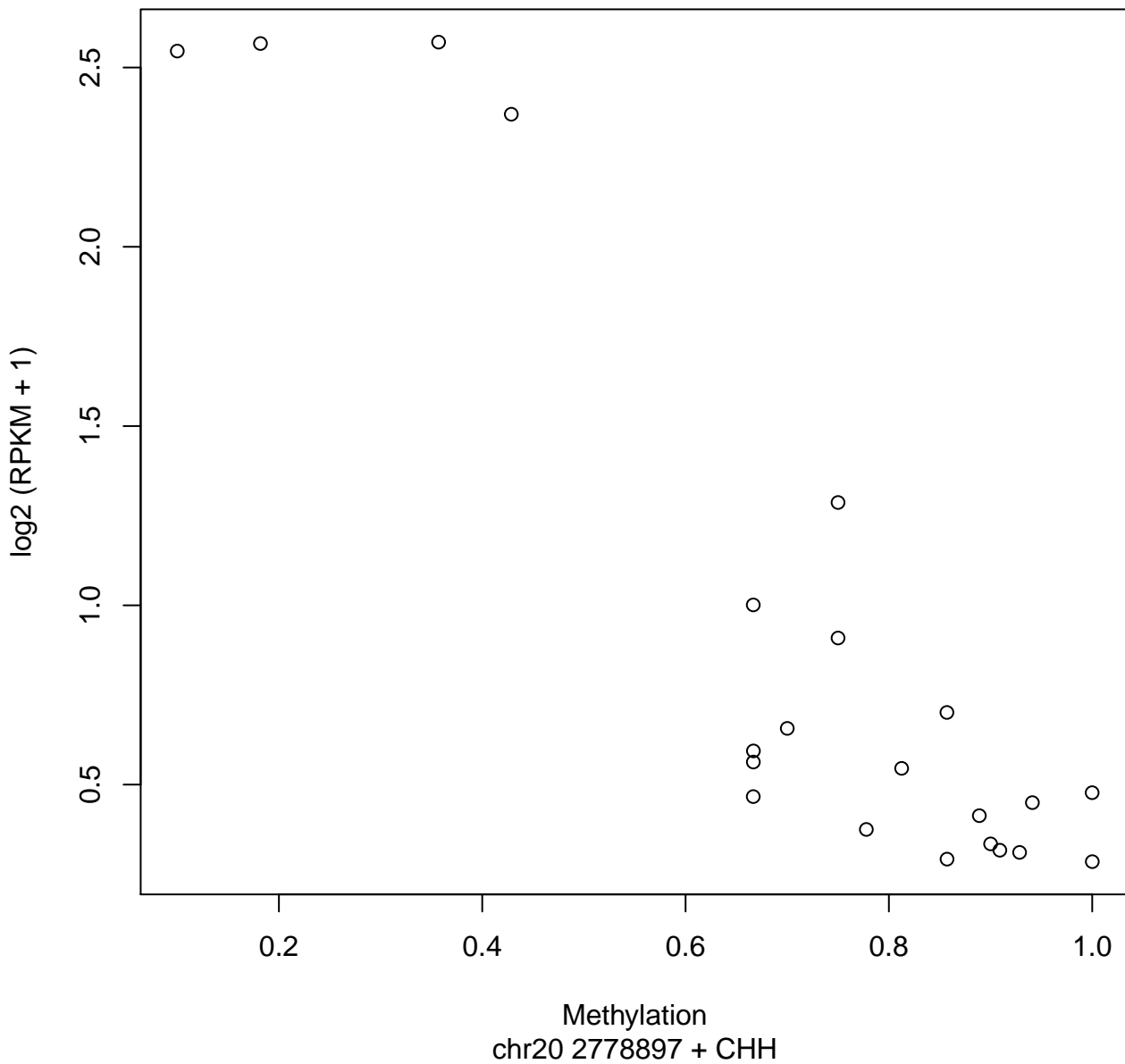
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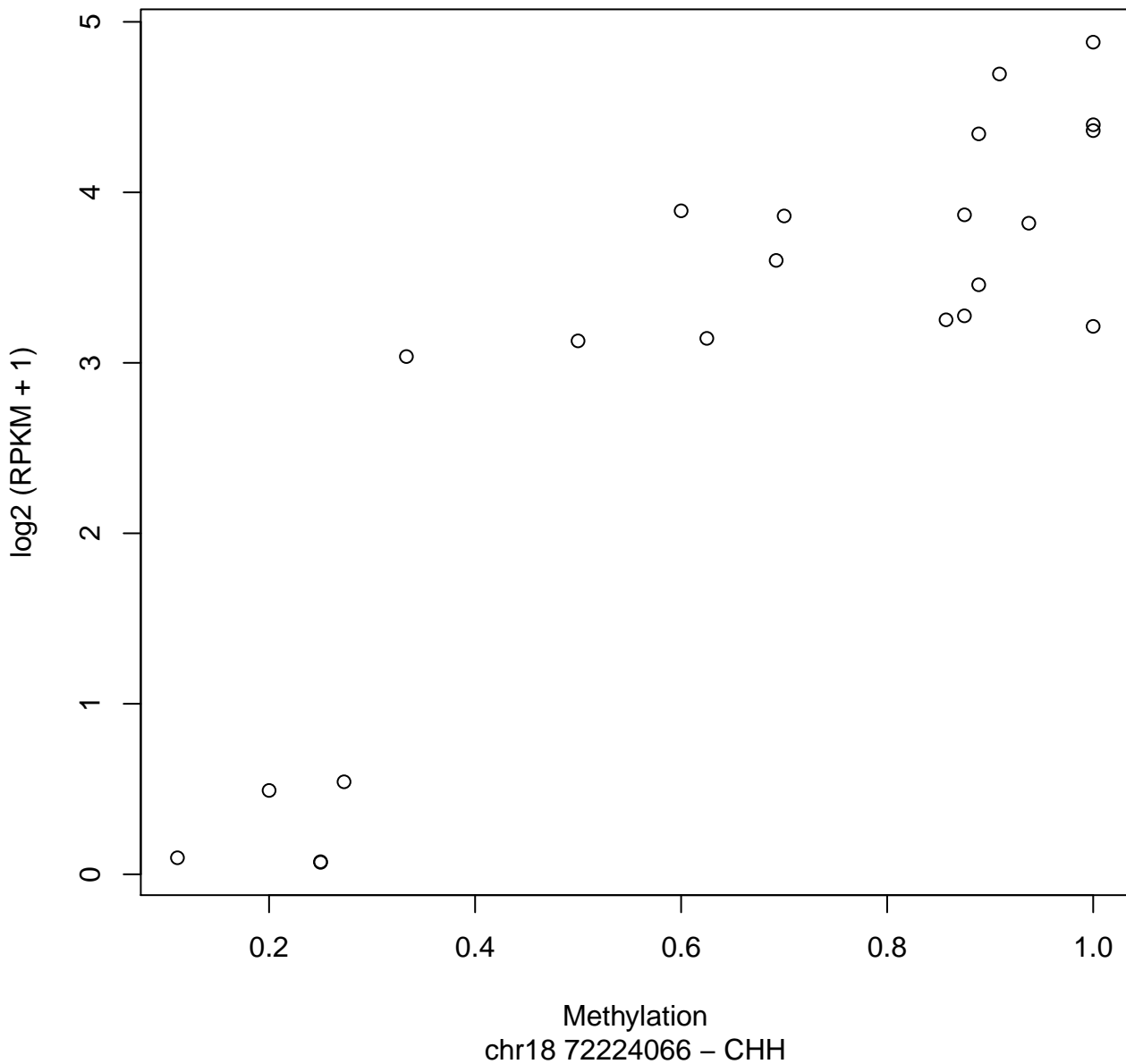


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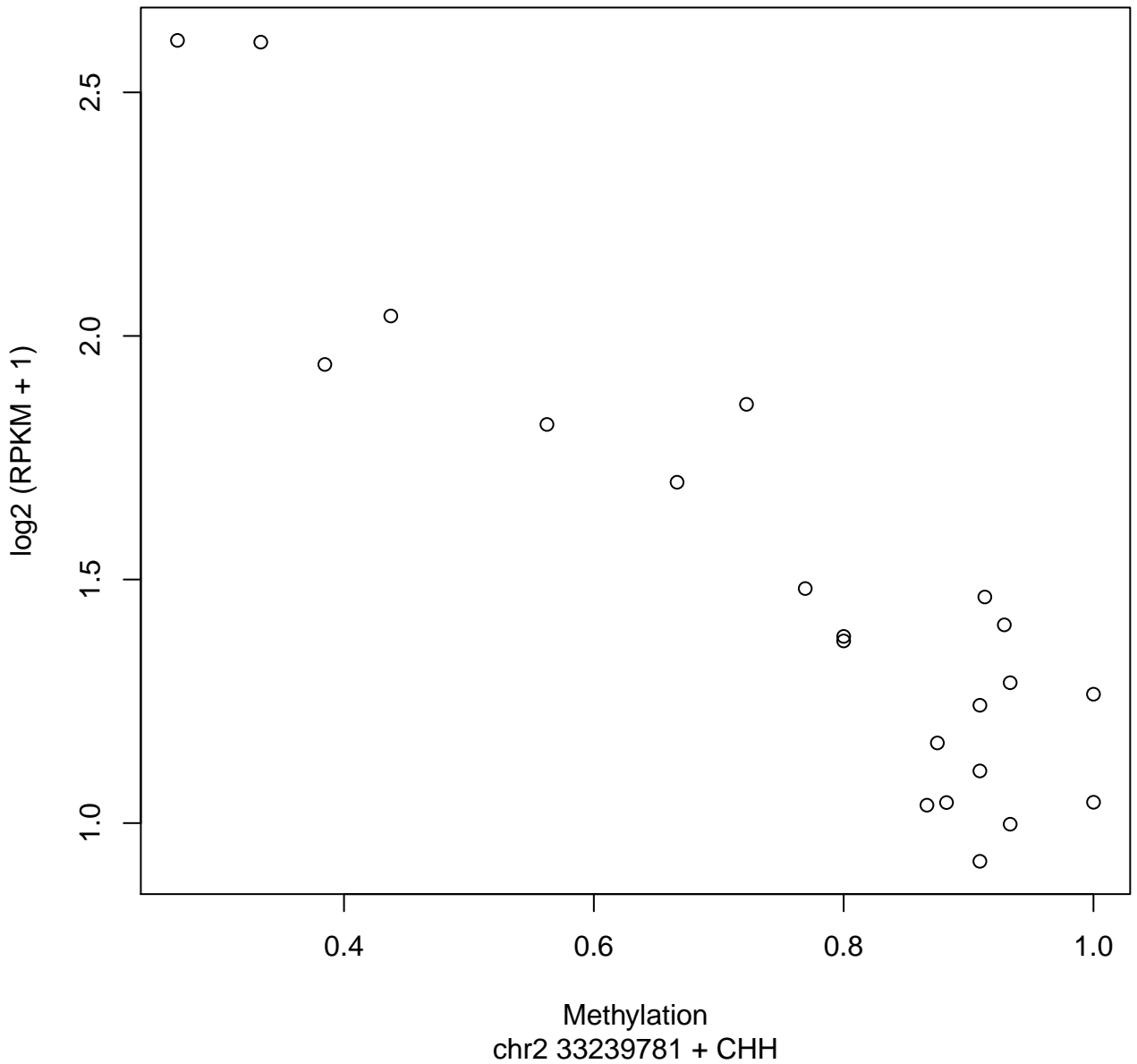
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CNDP1



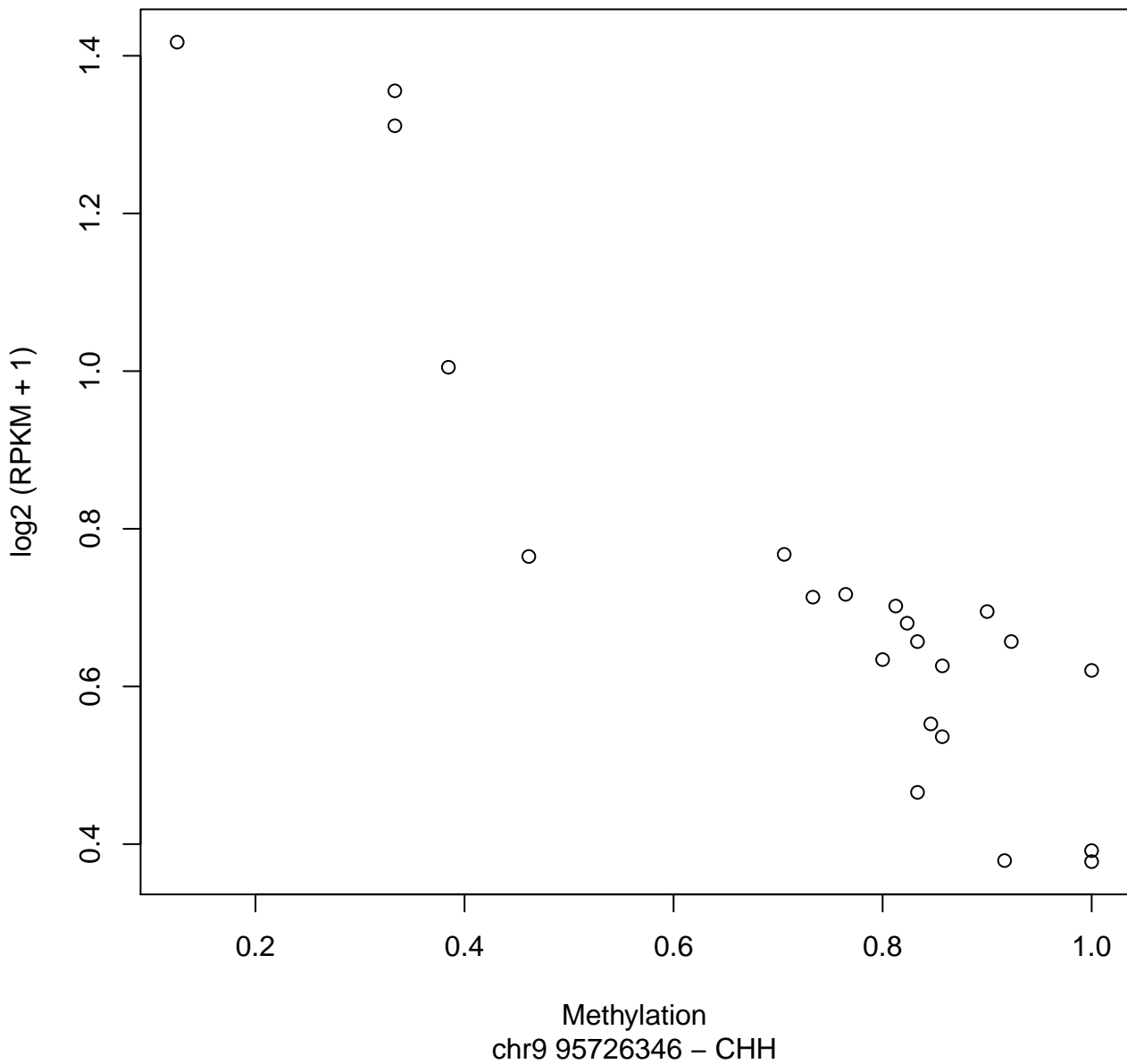
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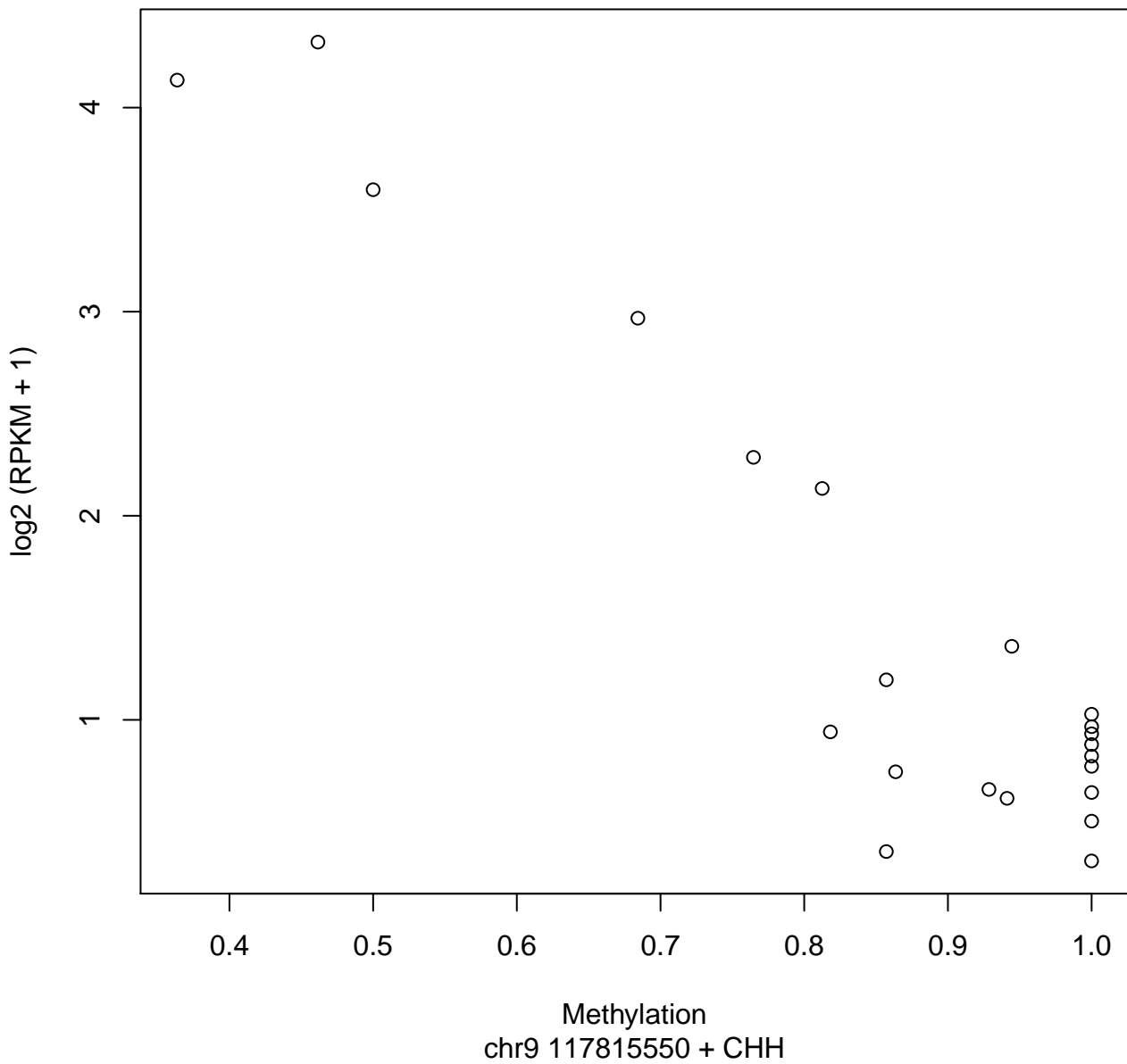
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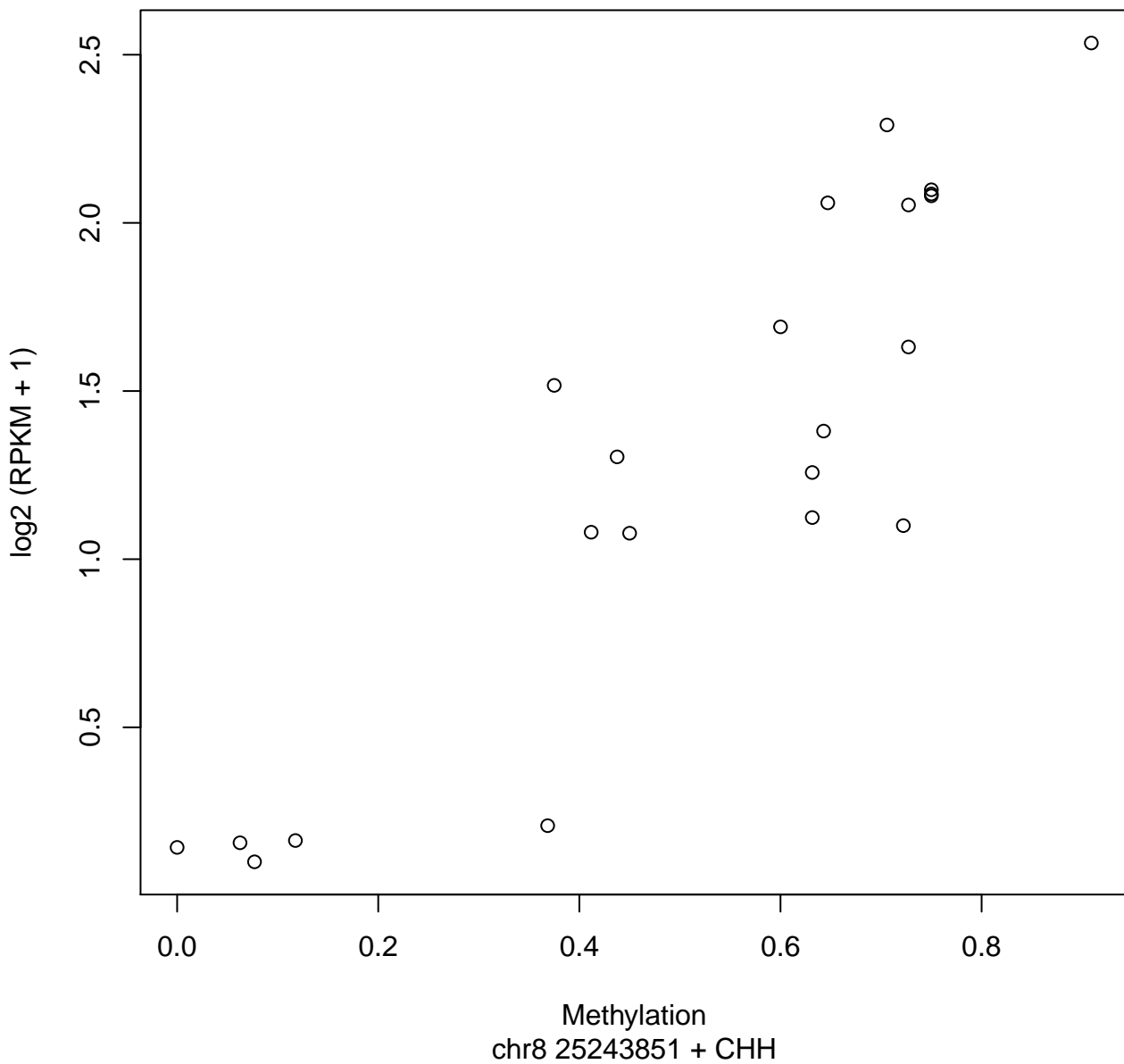
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TNC



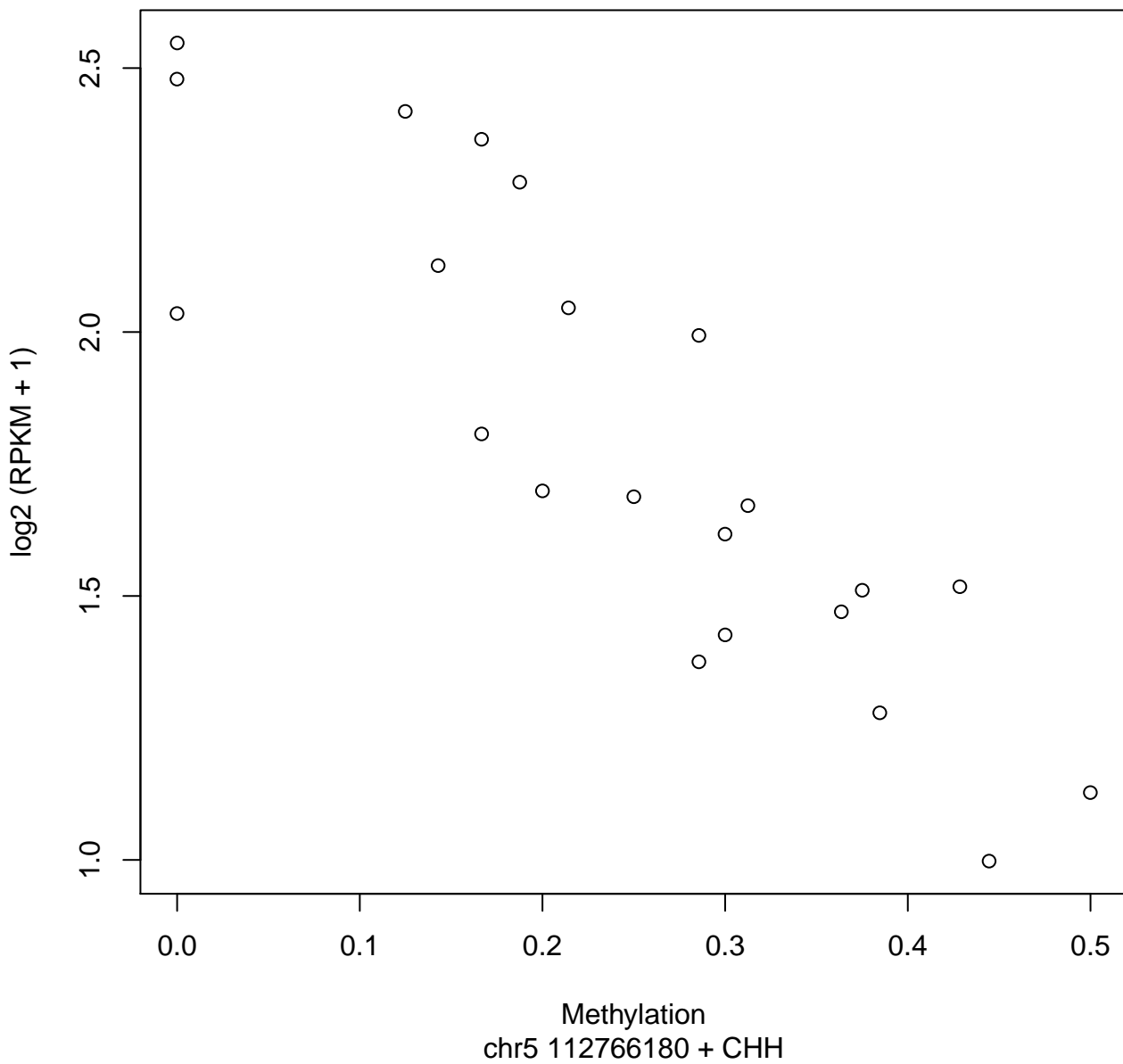
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DOCK5



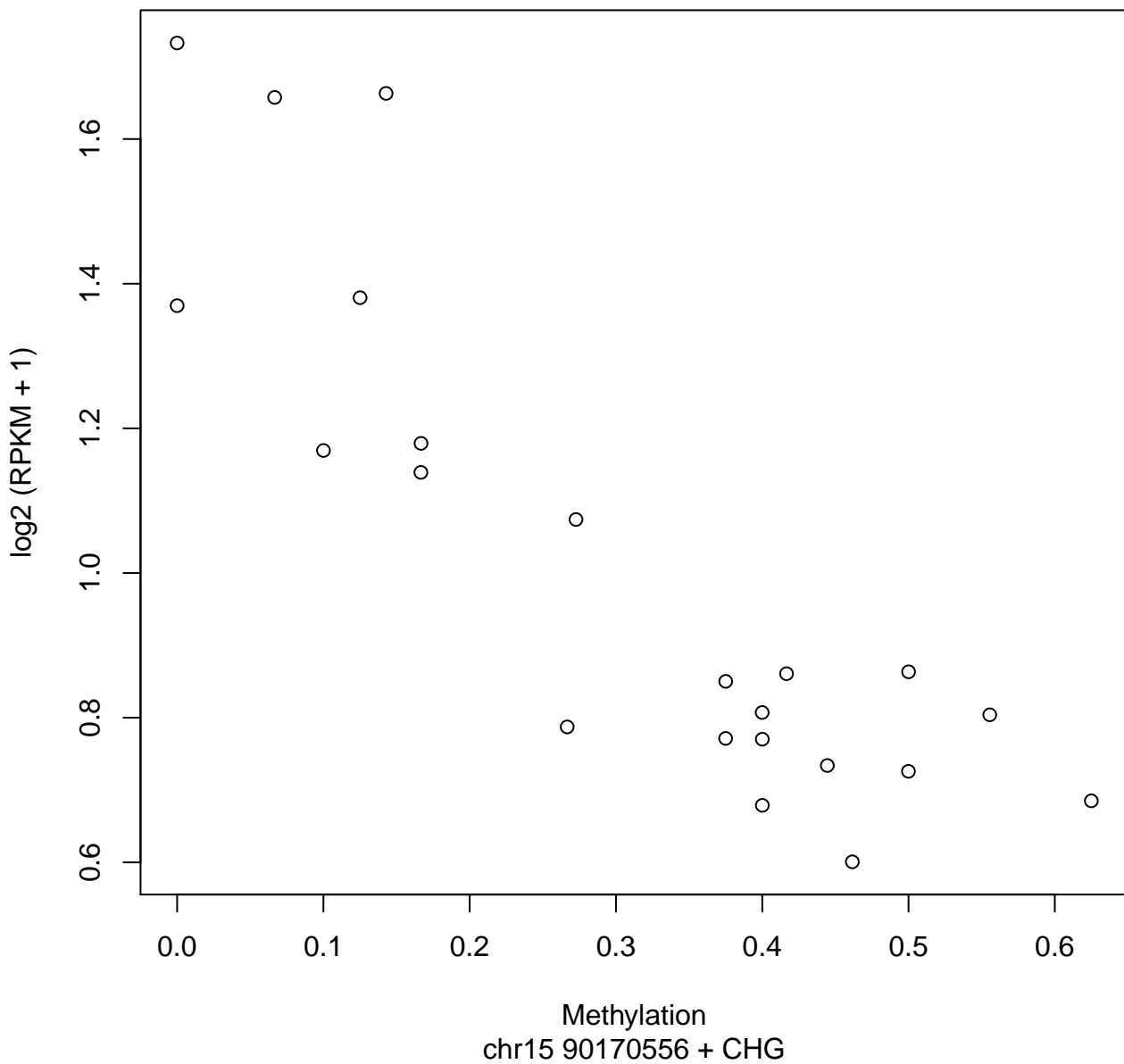
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MCC



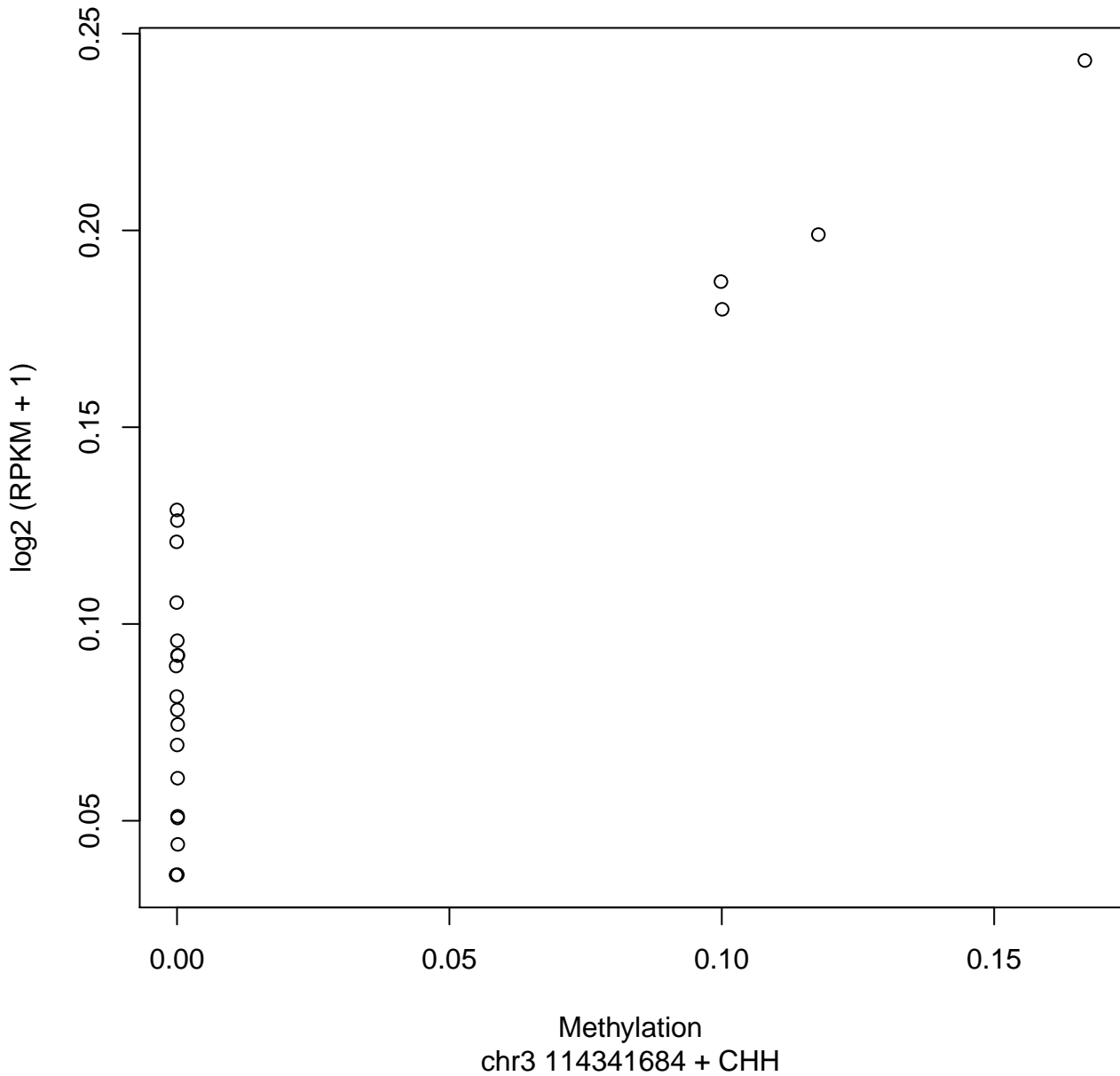
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KIF7

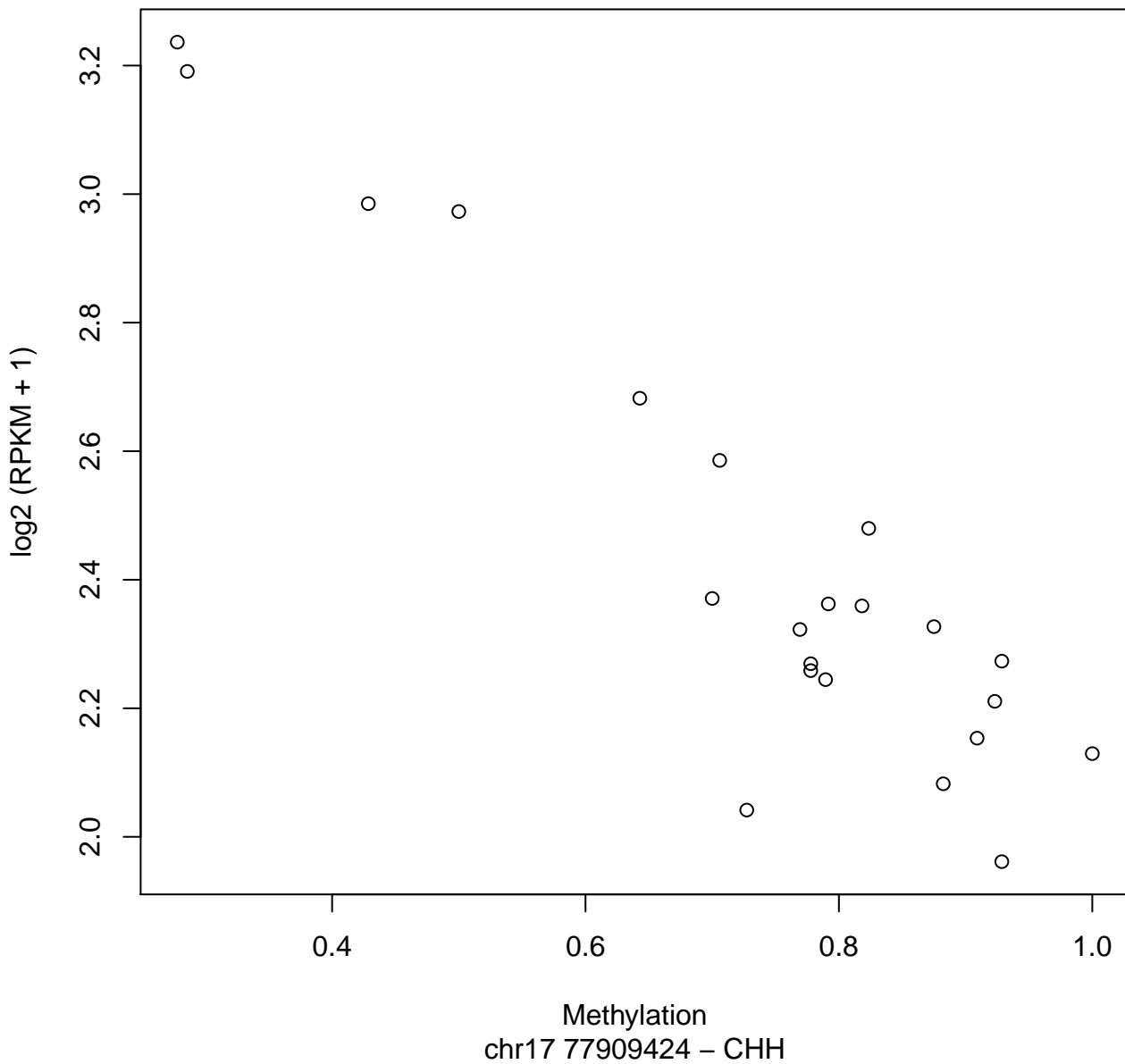


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ZBTB20

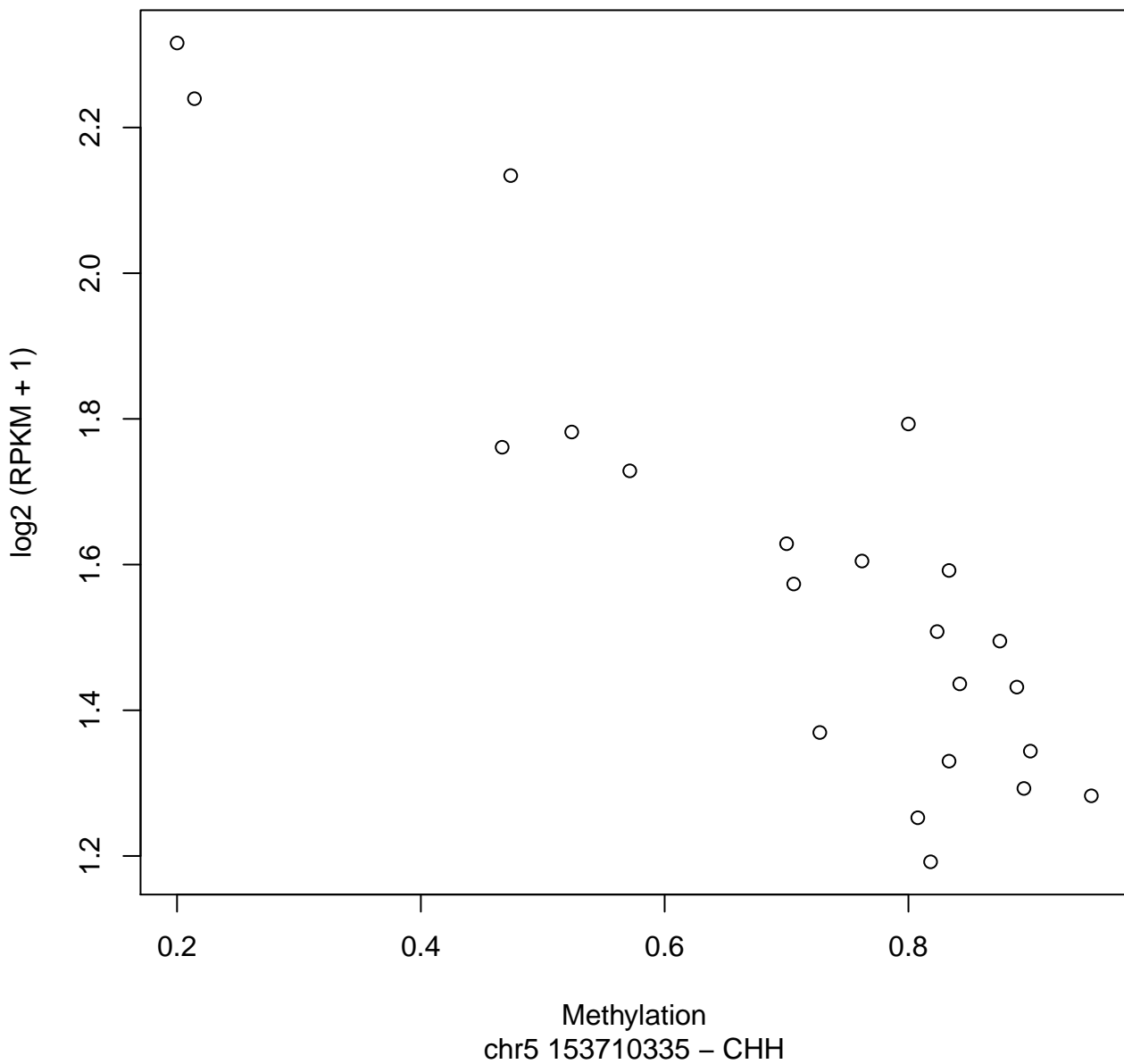


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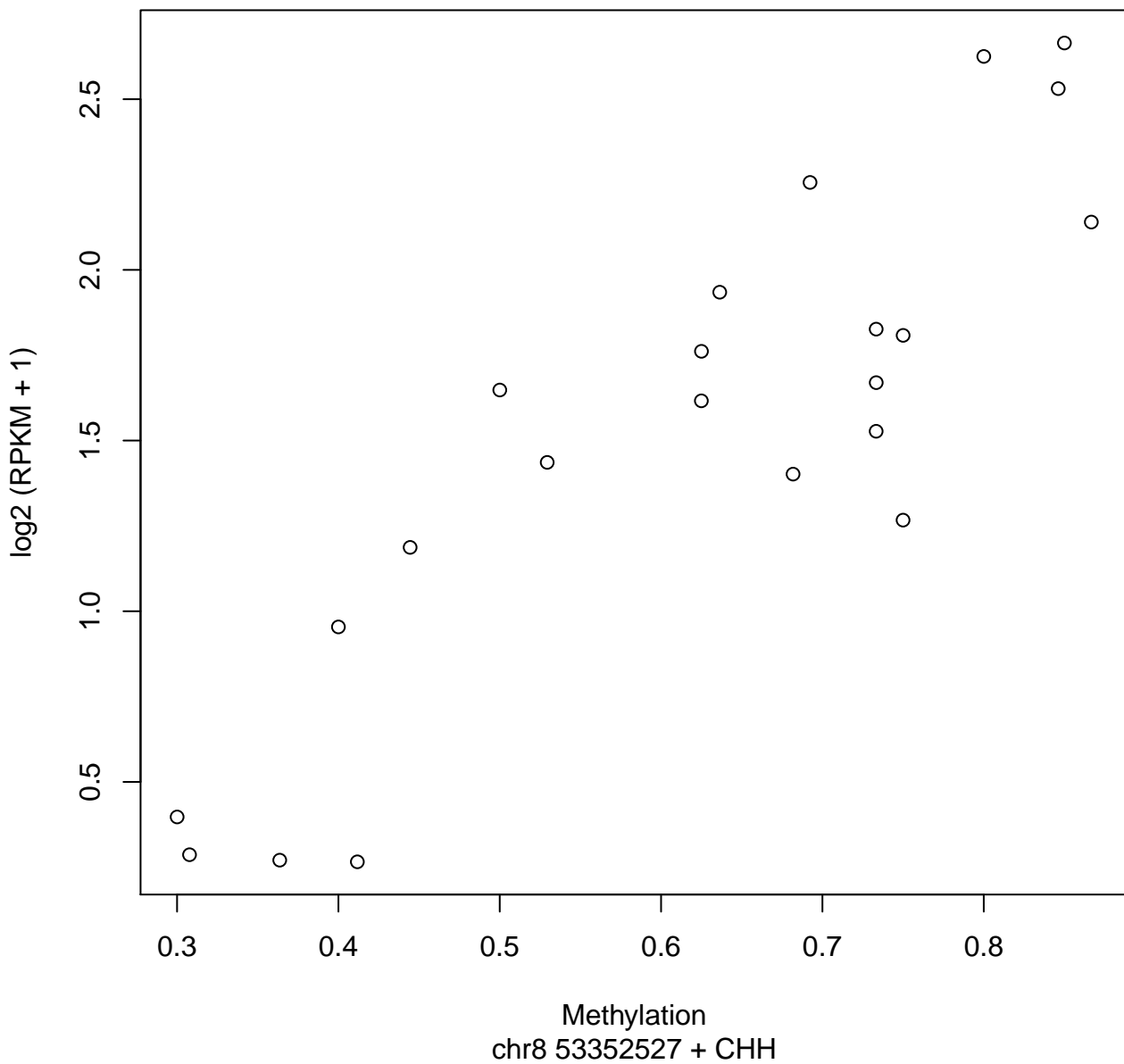
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GALNT10

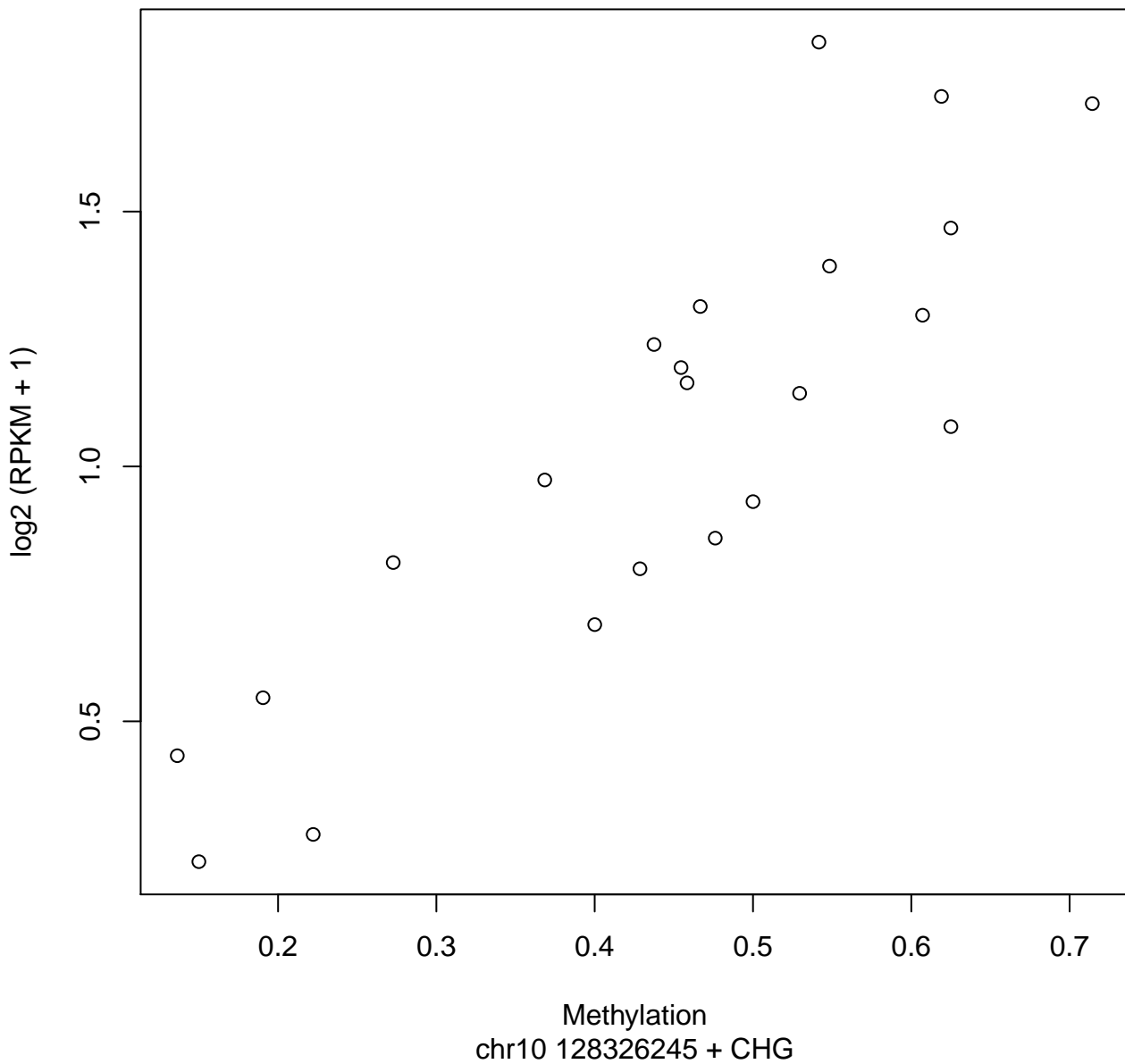


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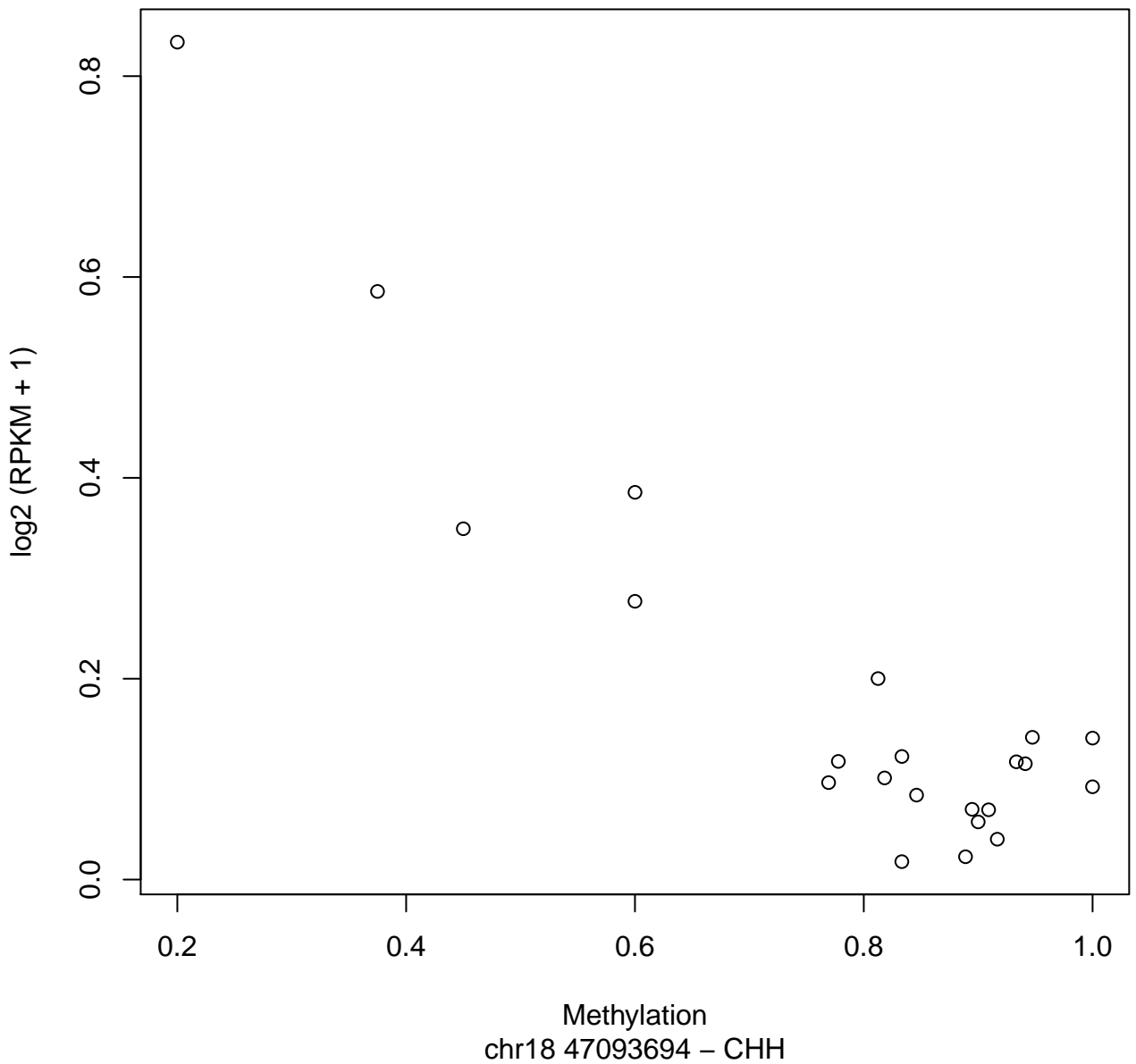
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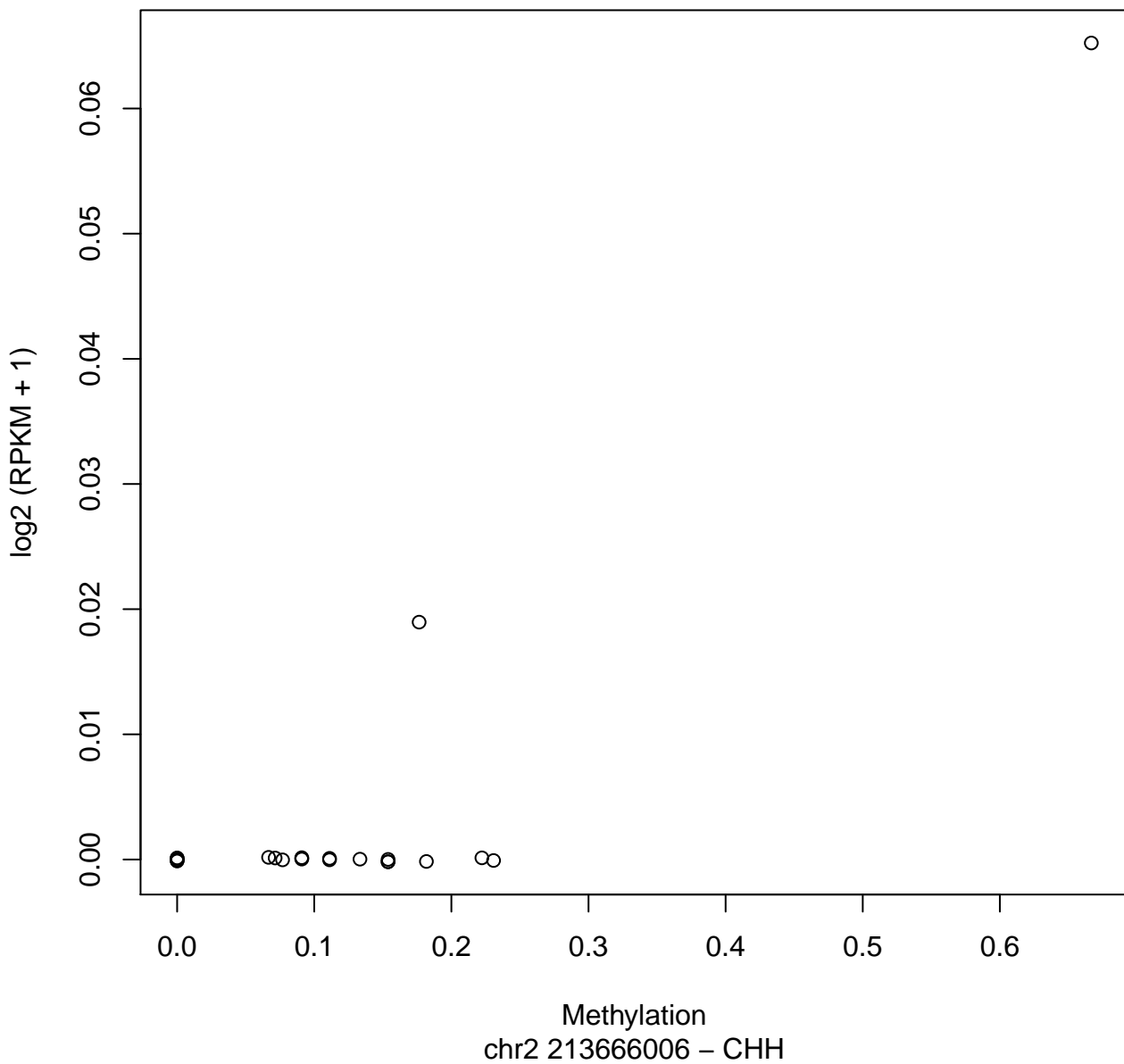
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C10orf90



gene ENSG00000101670.11_2 FDR 3.33e-05
LIPG

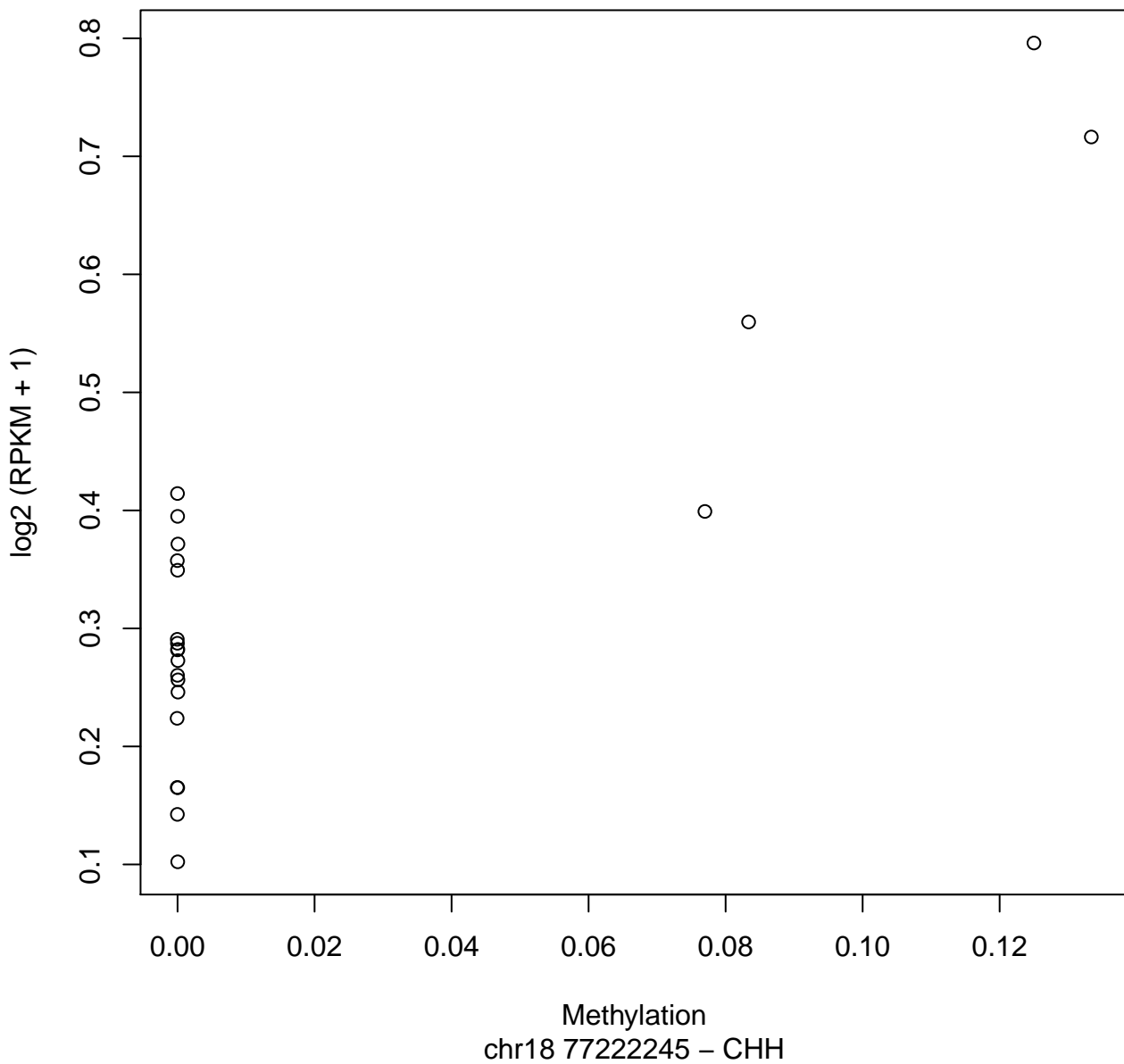


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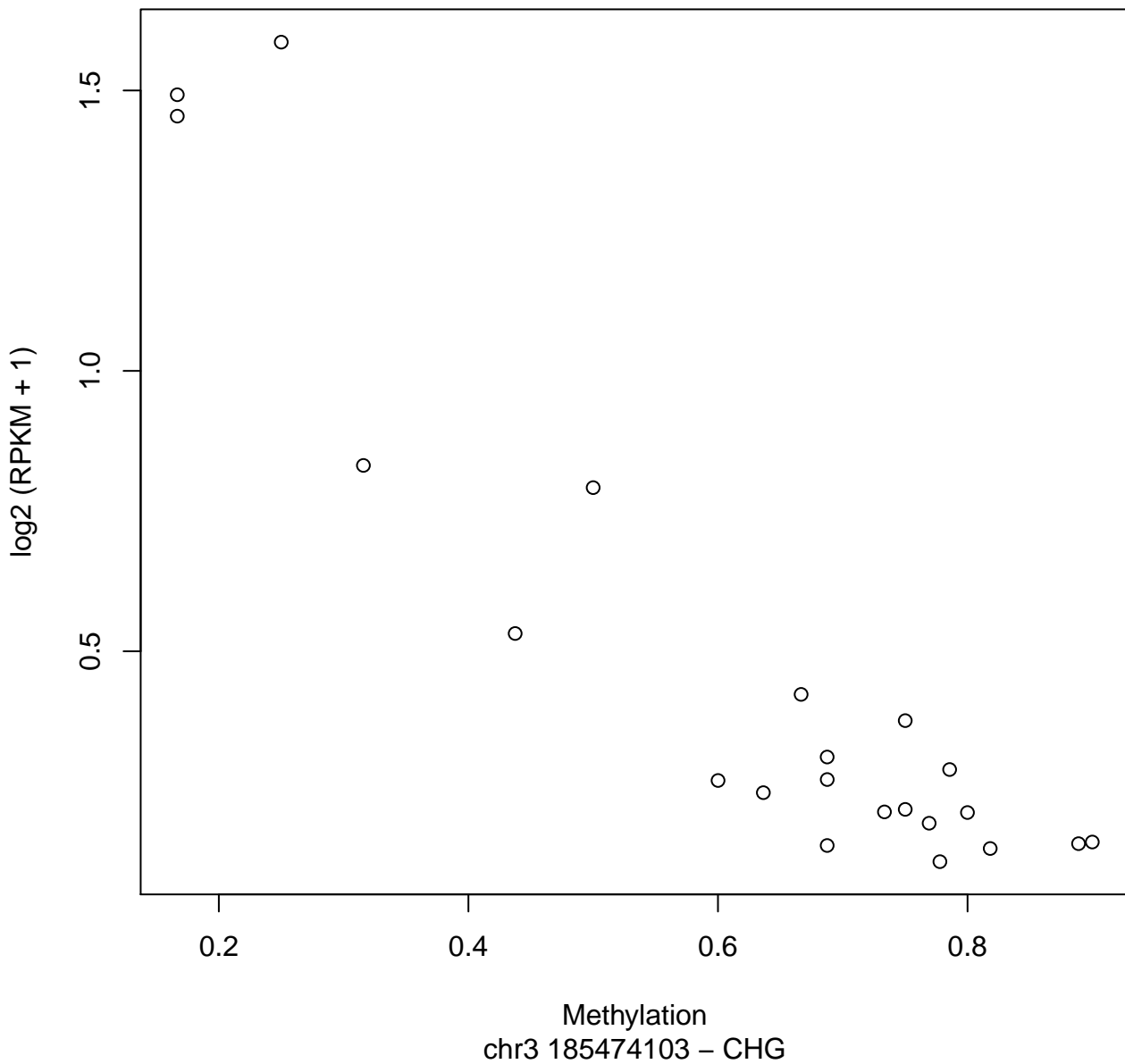


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NFATC1

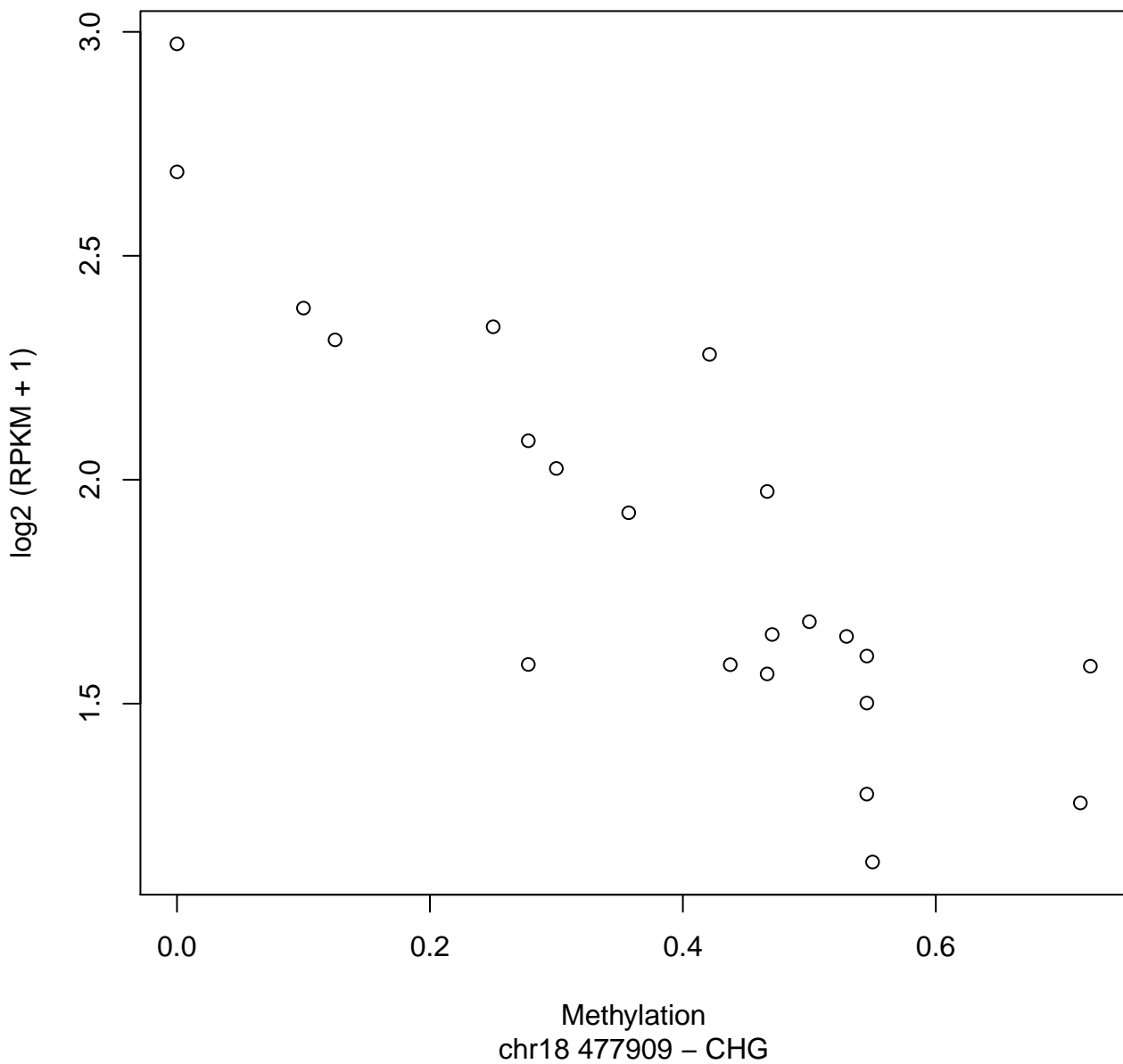


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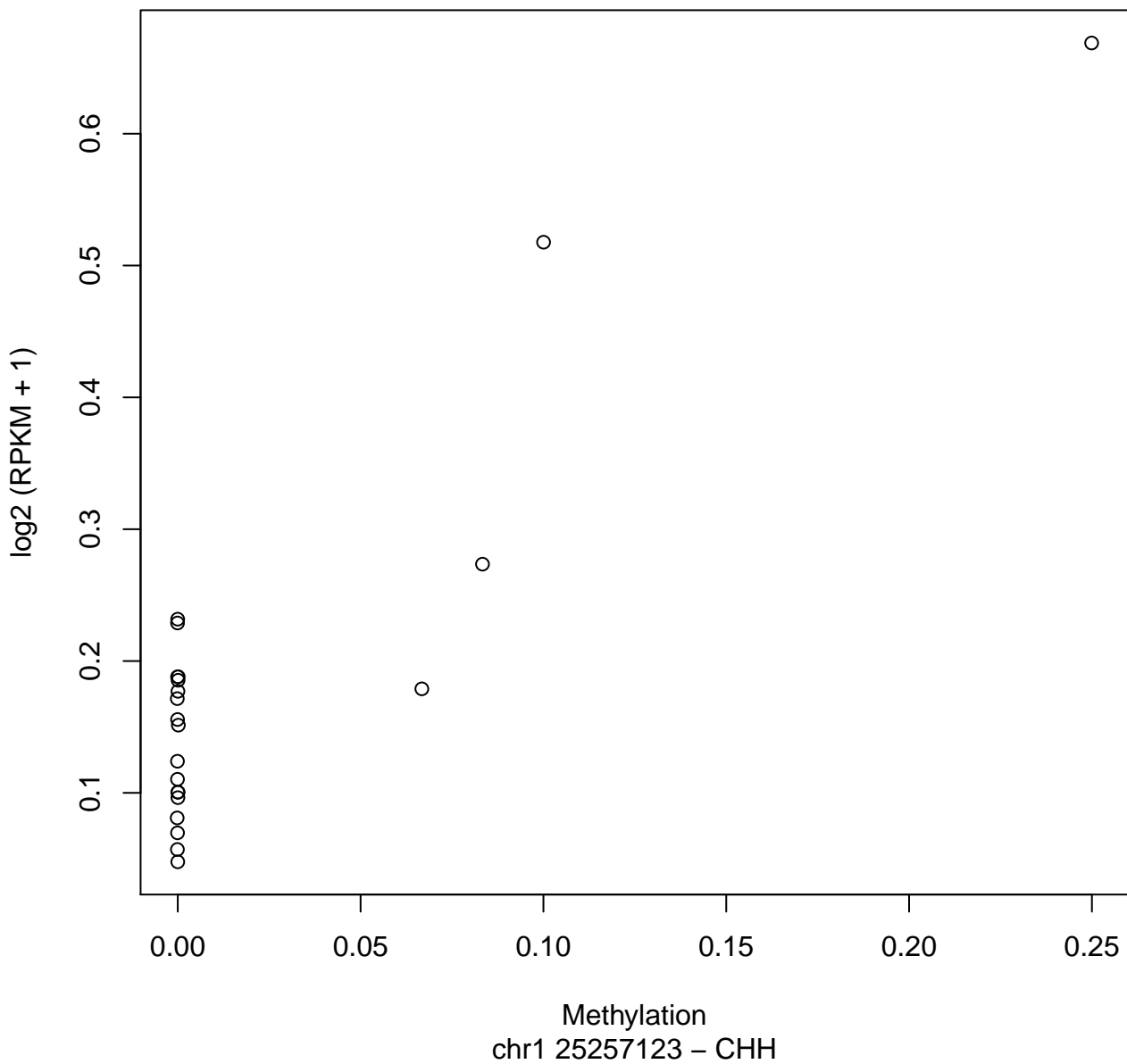
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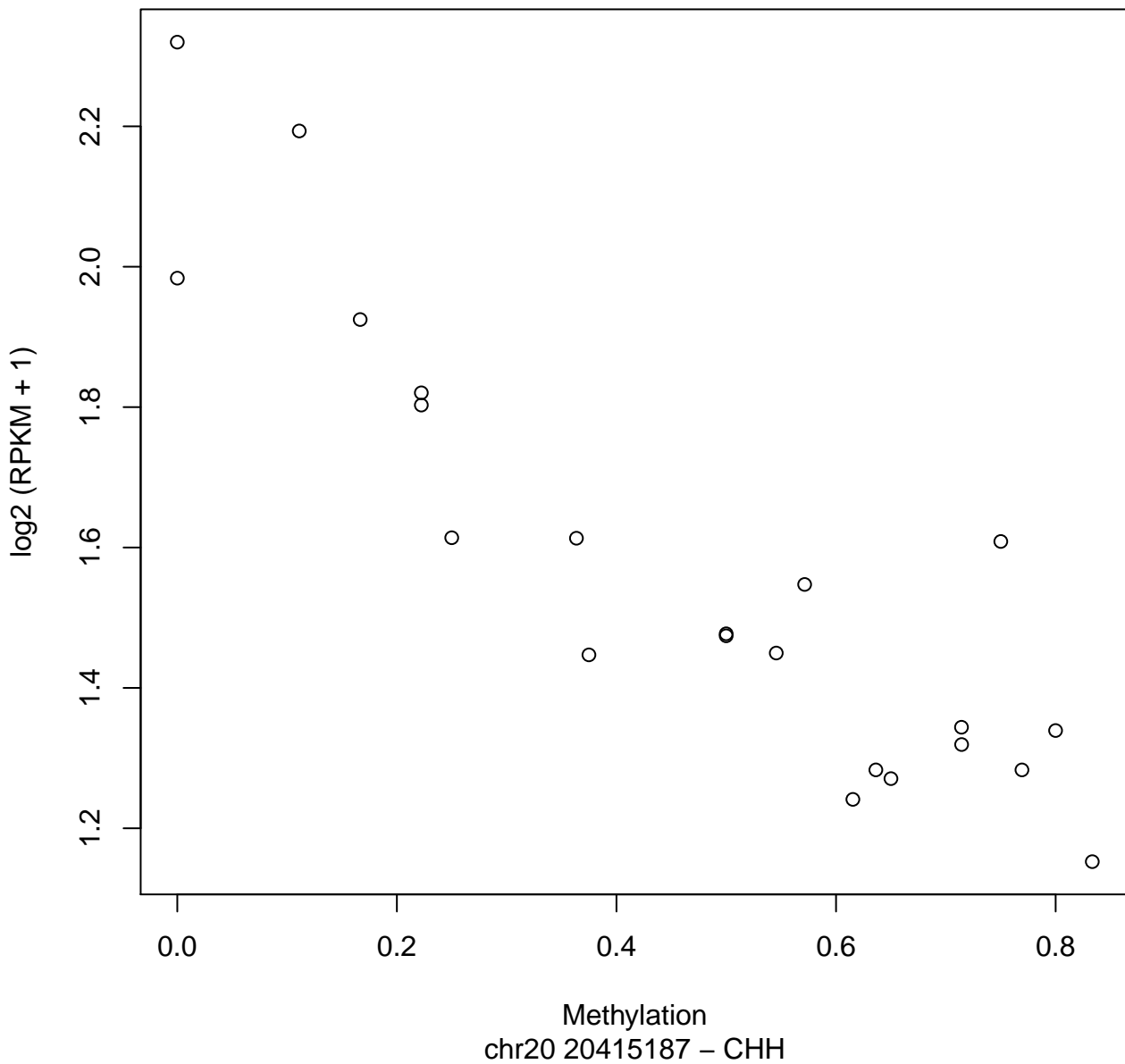


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RUNX3

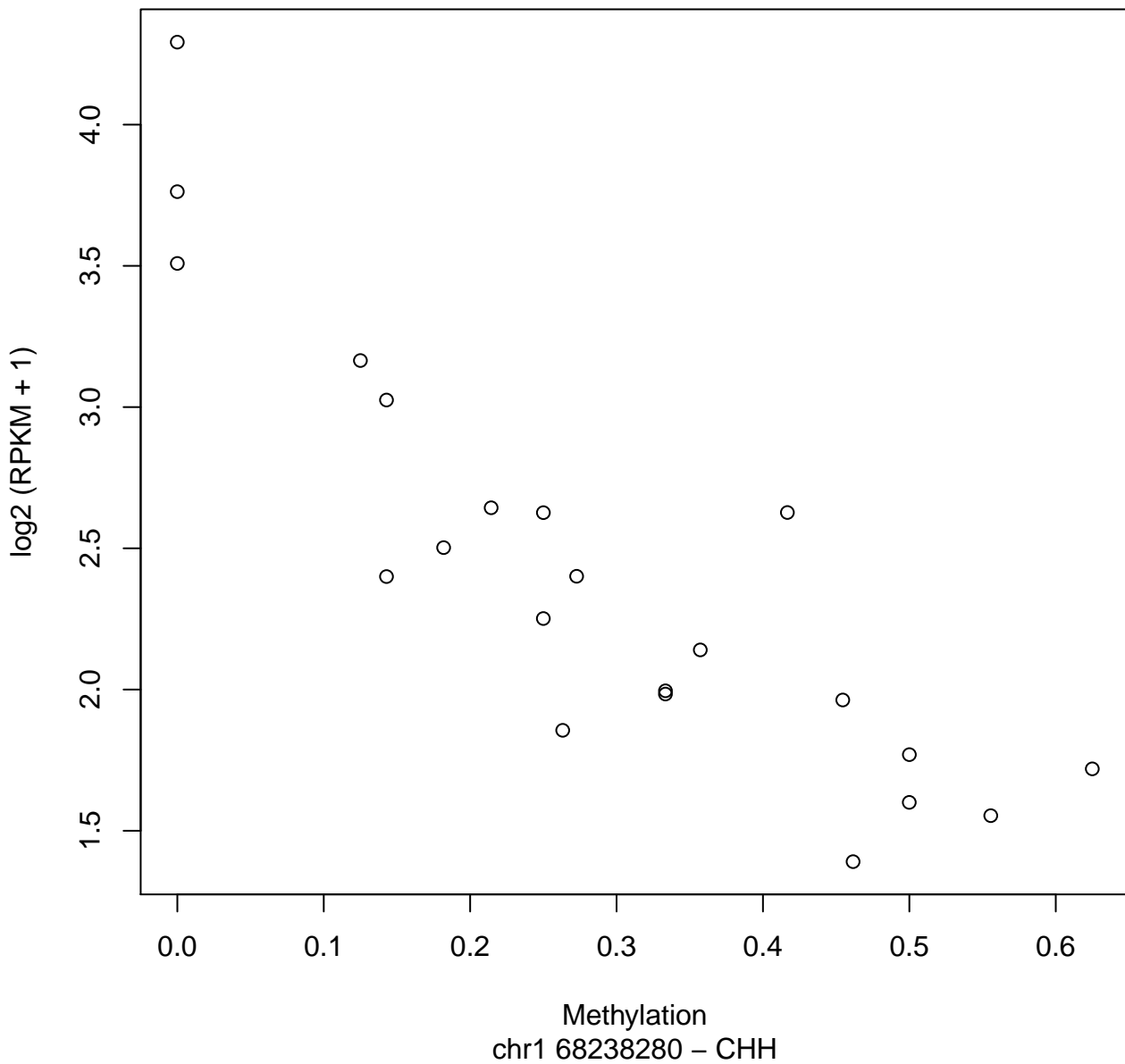


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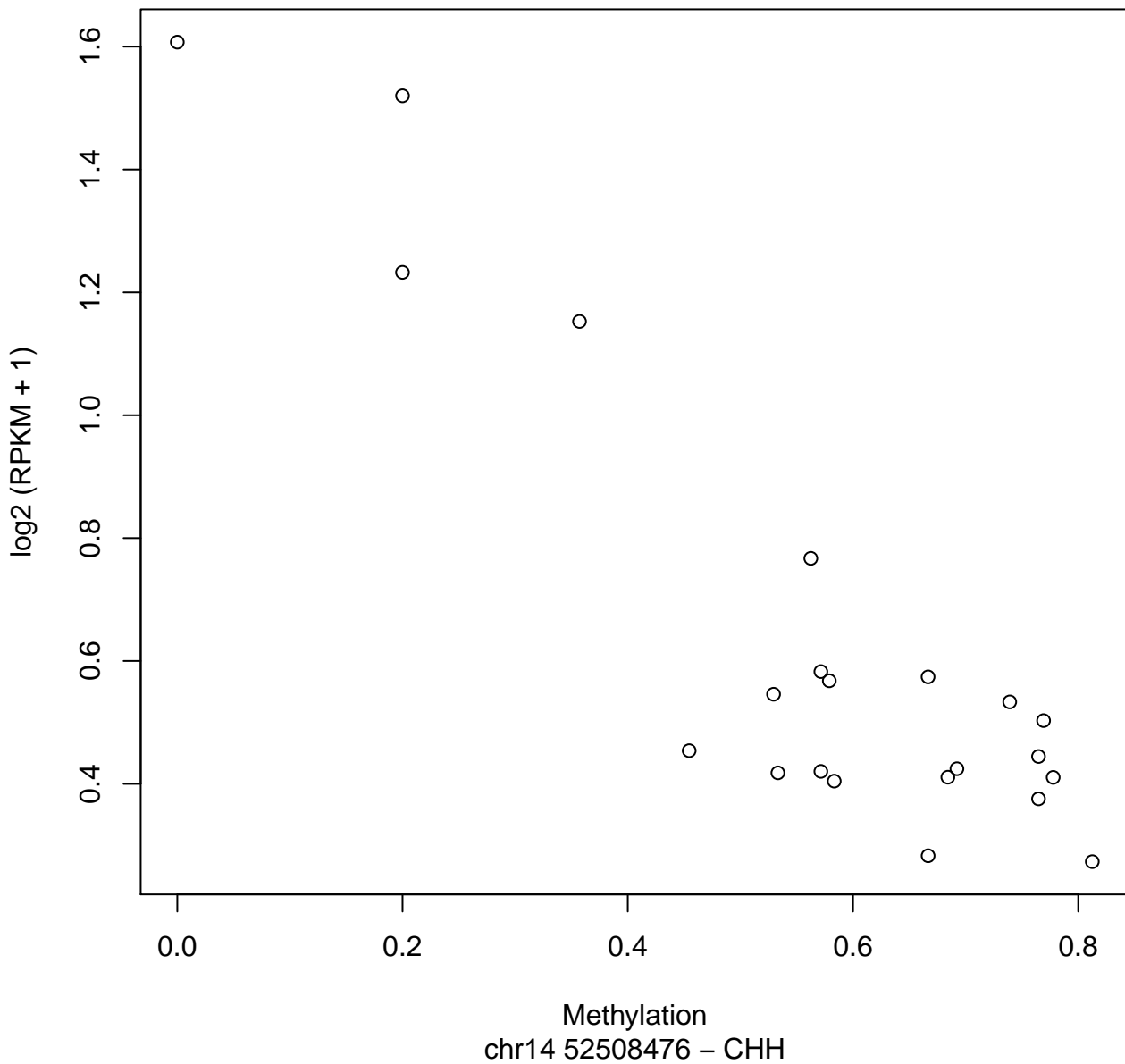


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GNG12

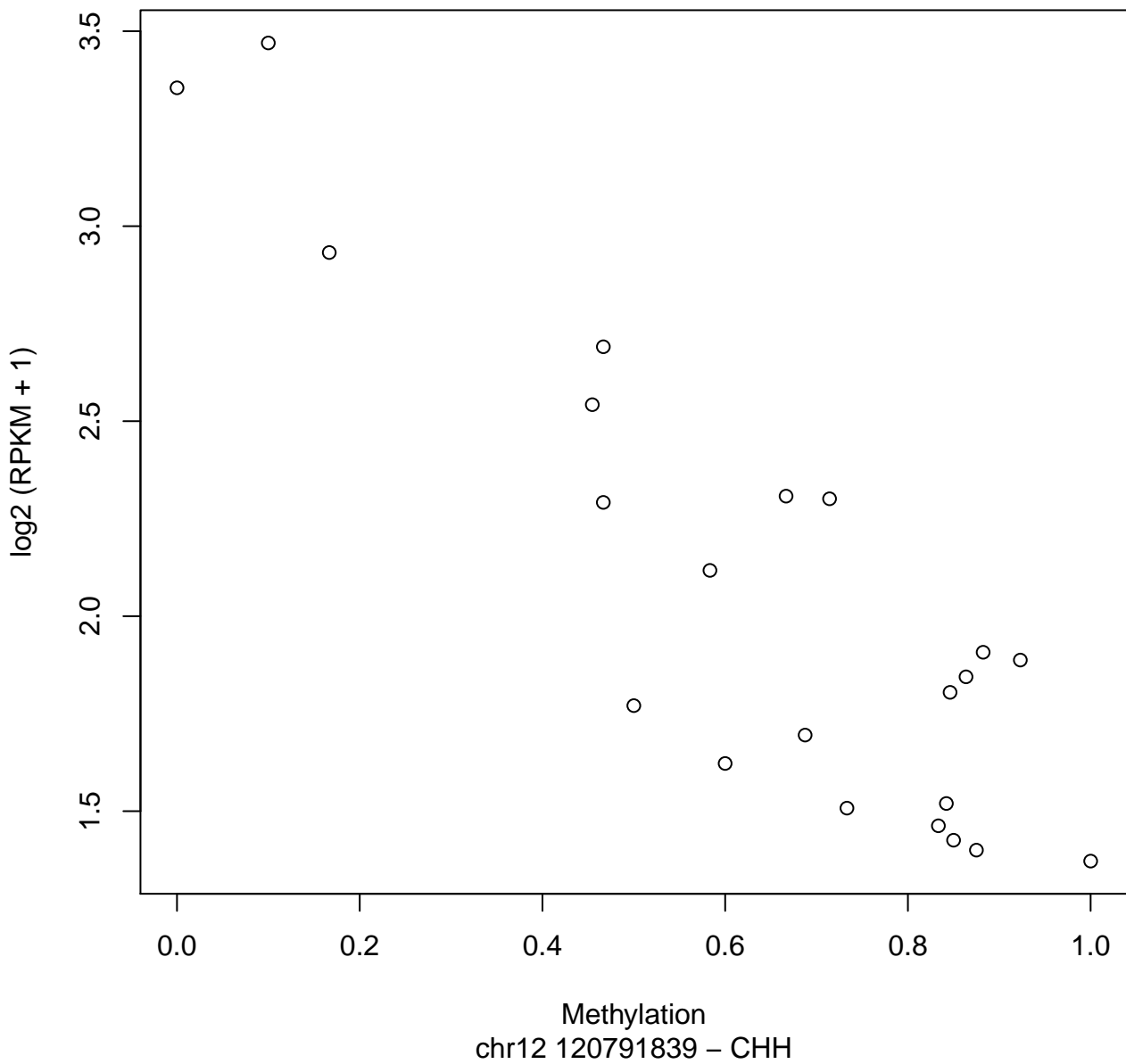


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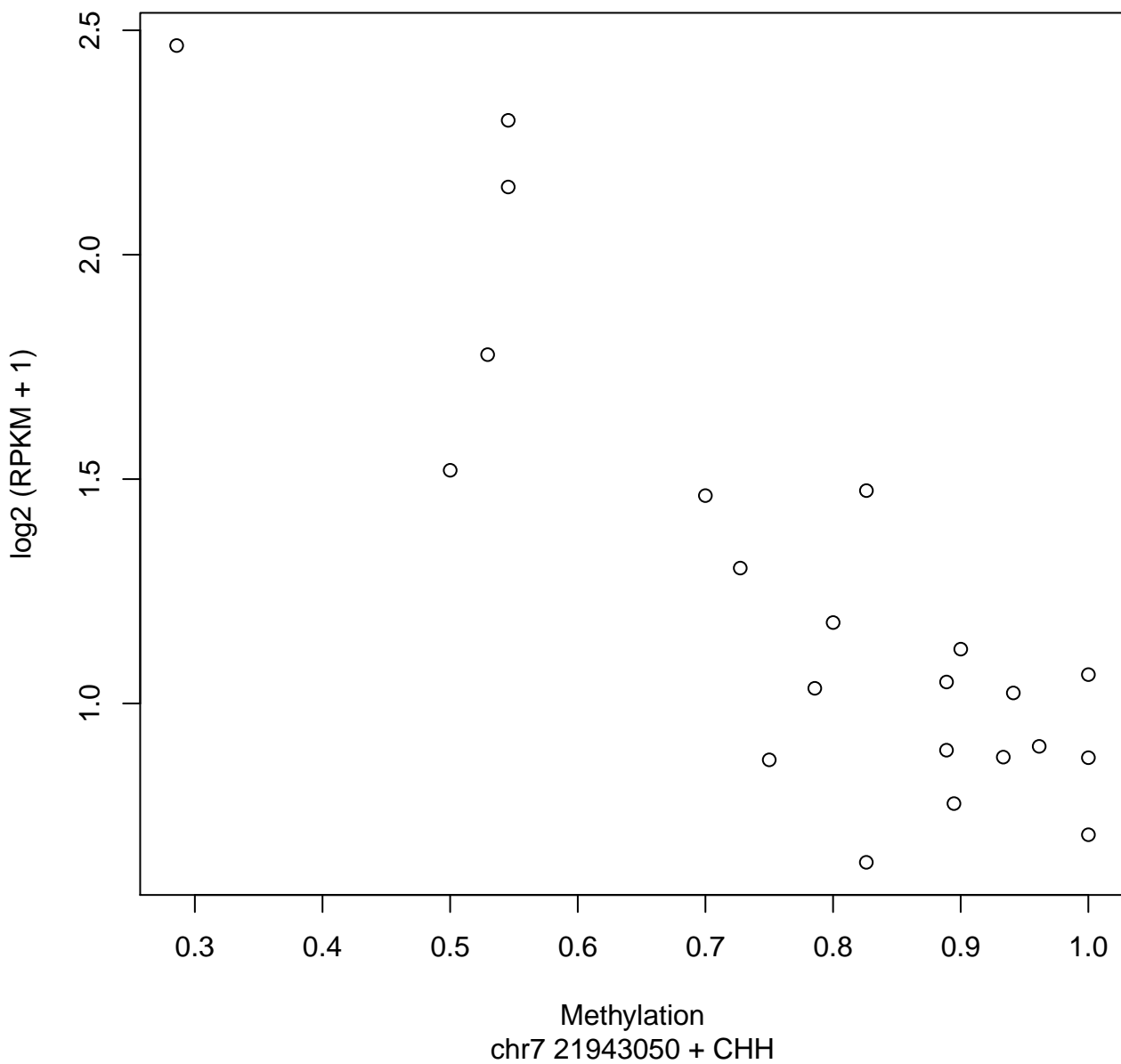
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MSI1



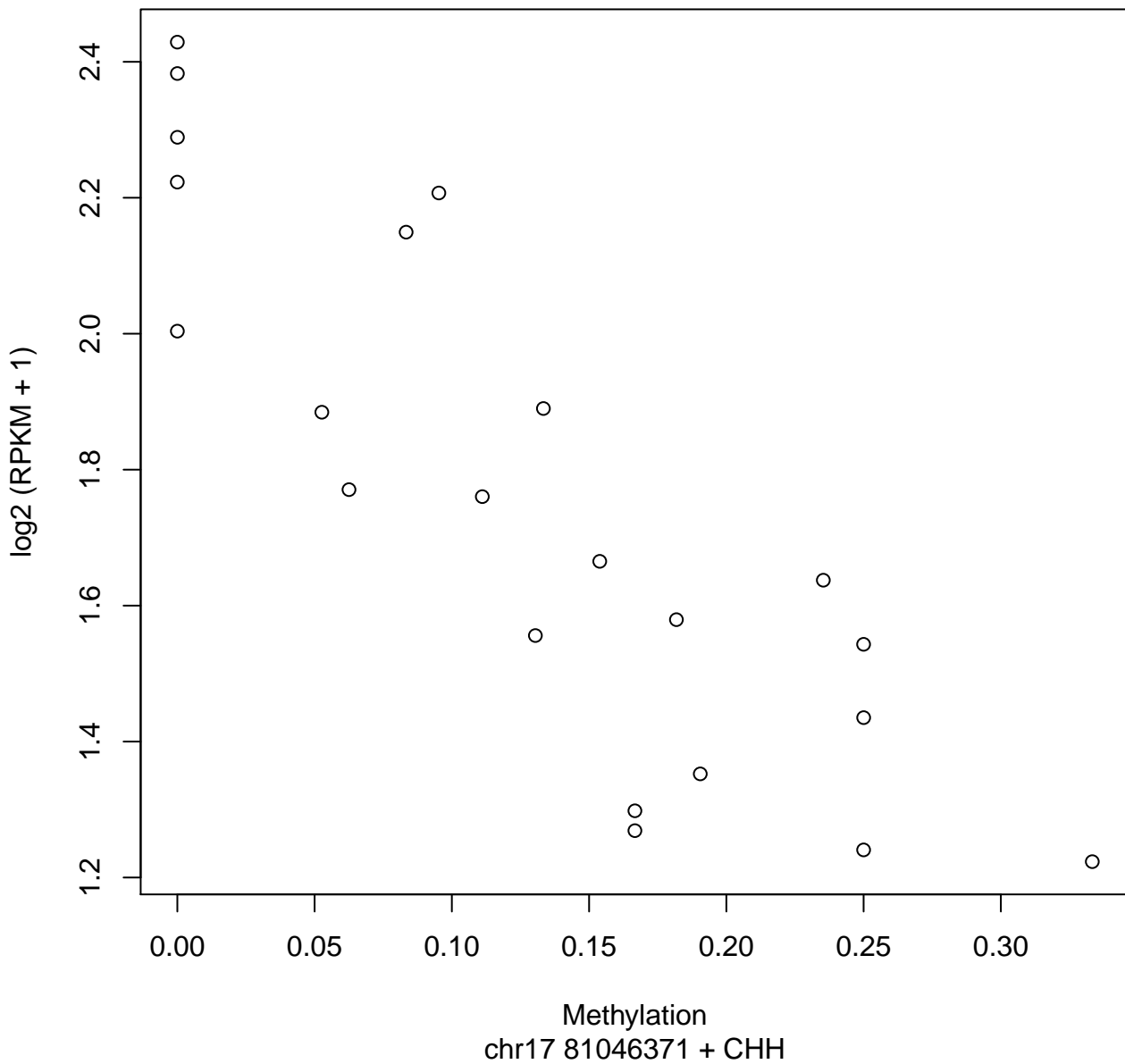
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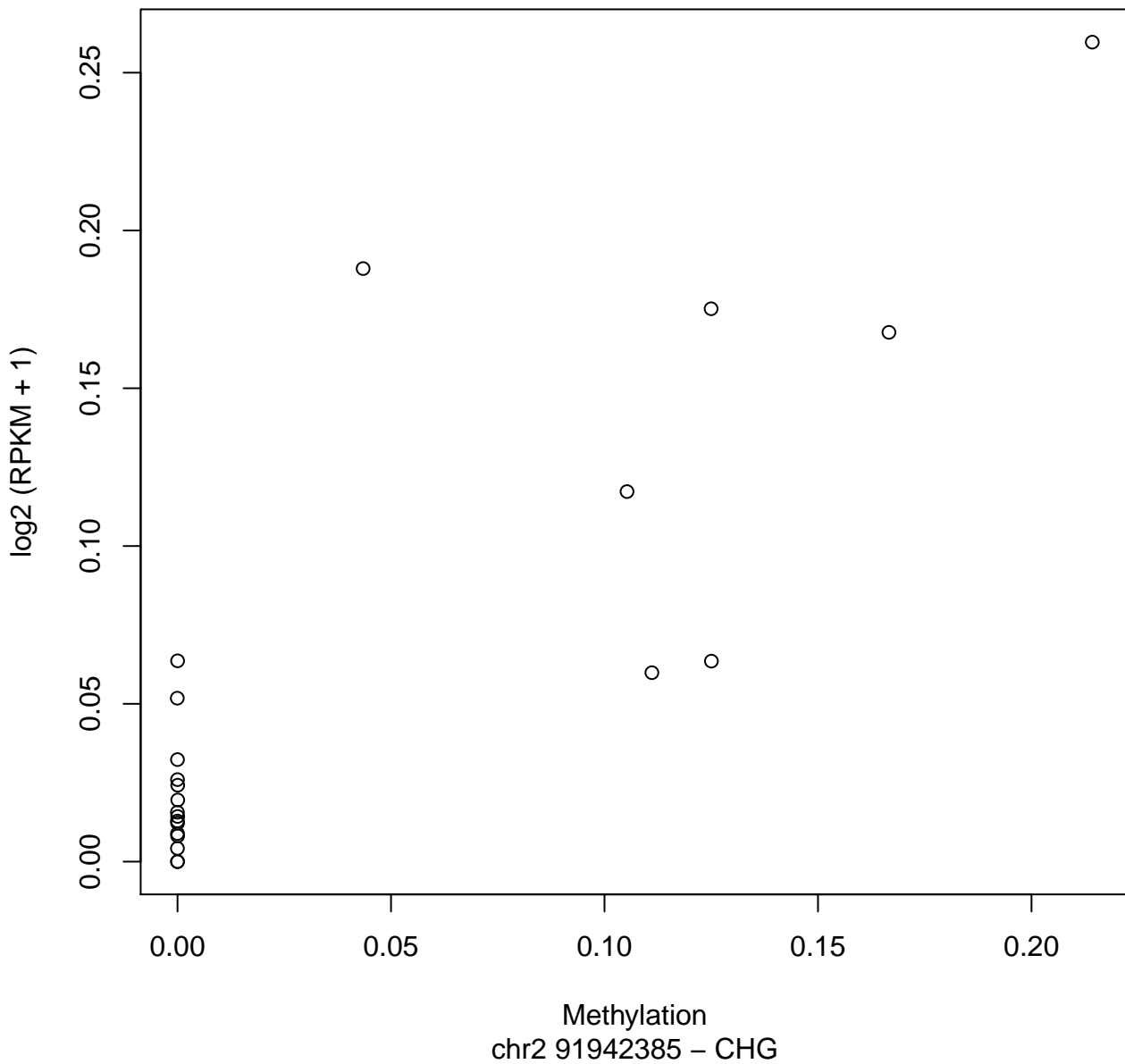


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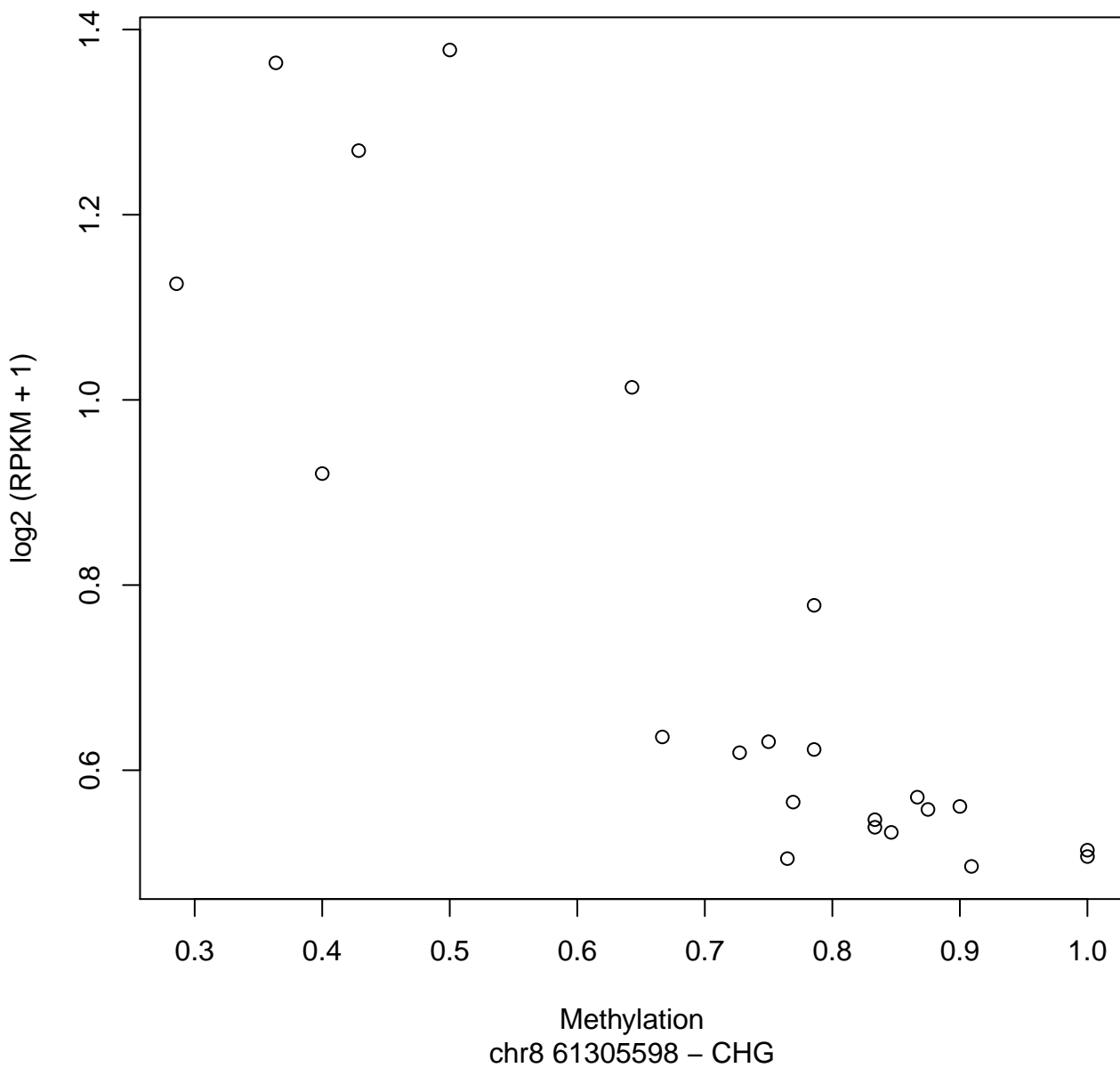
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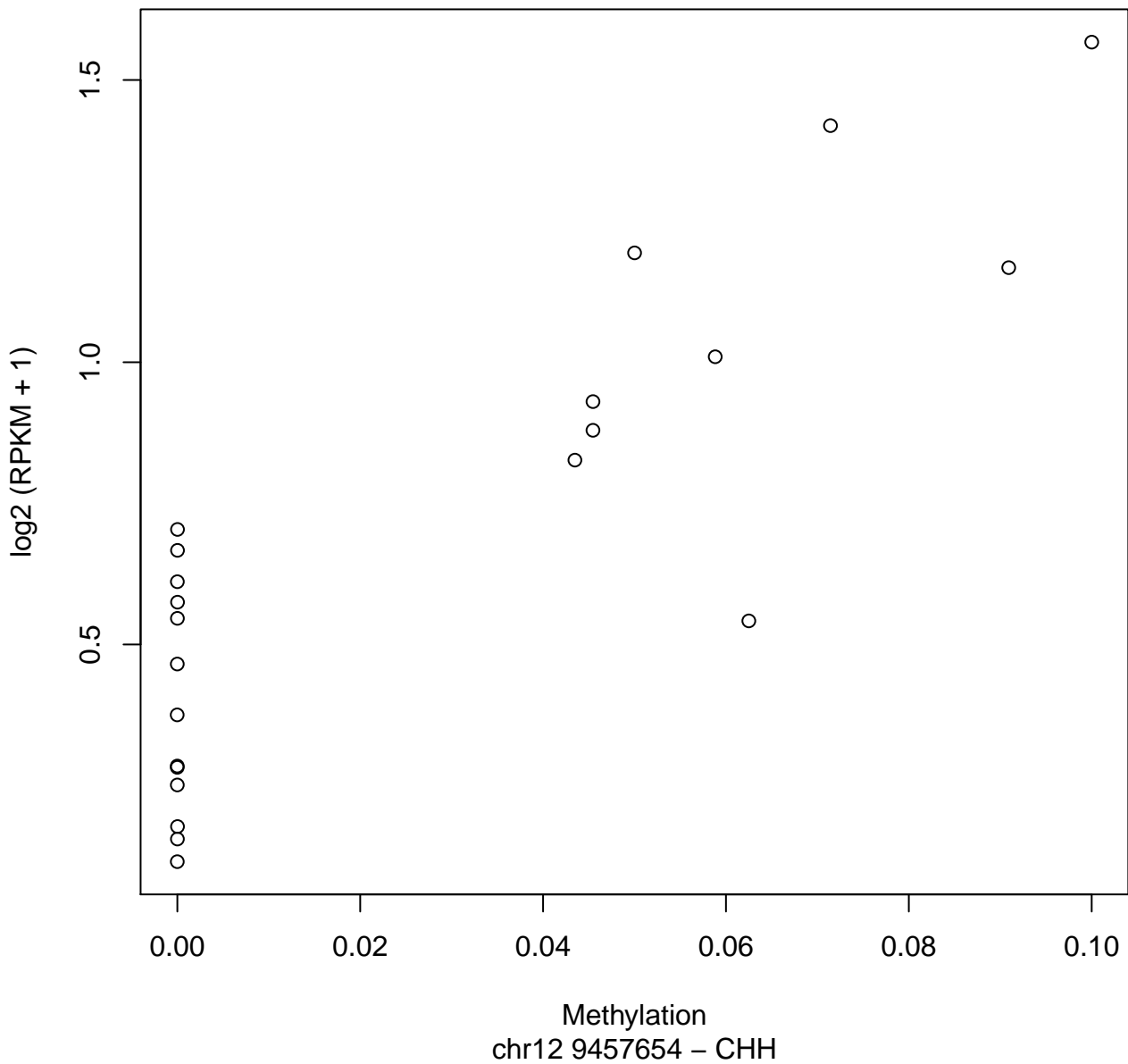
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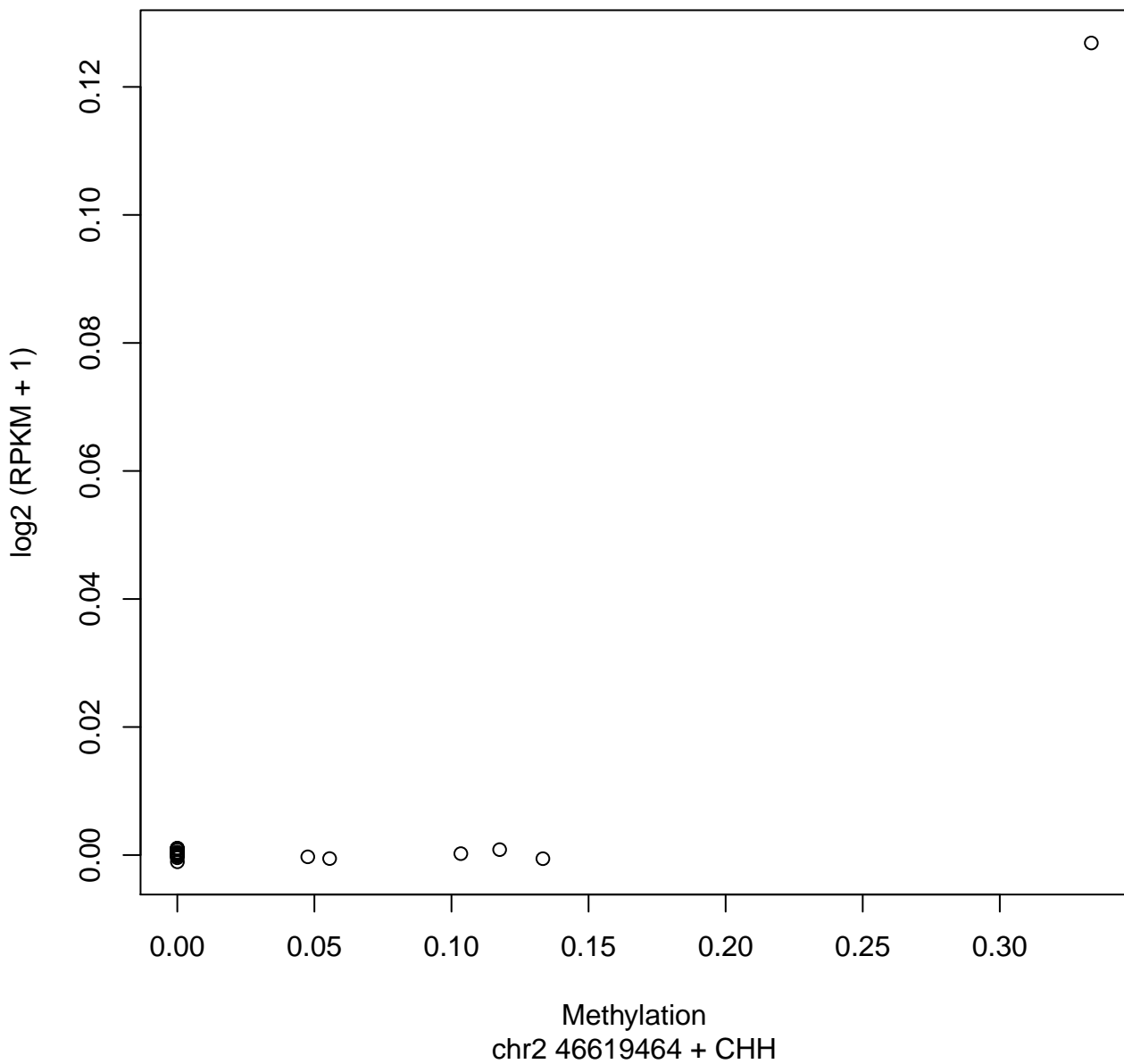
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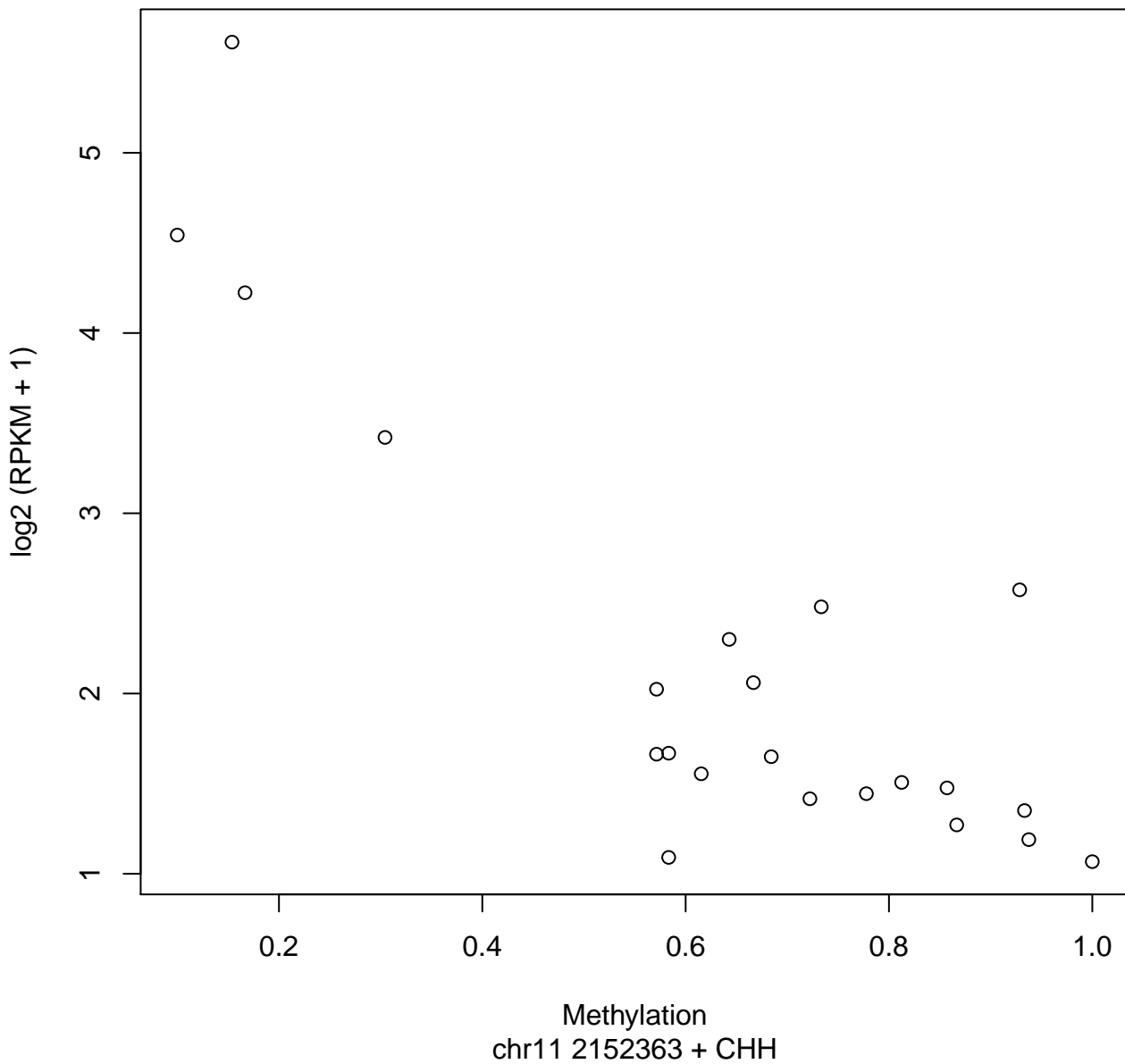


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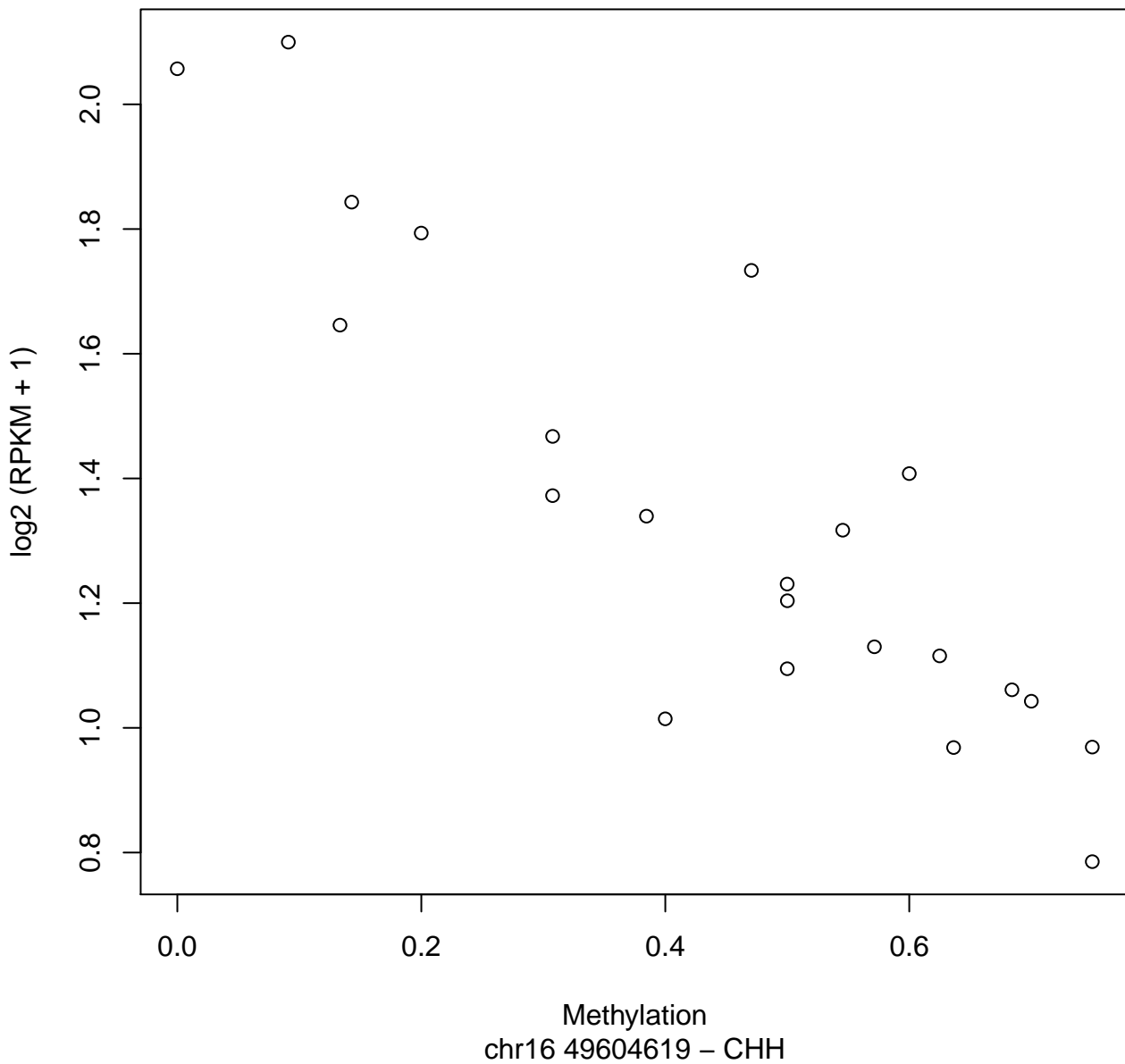


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IGF2

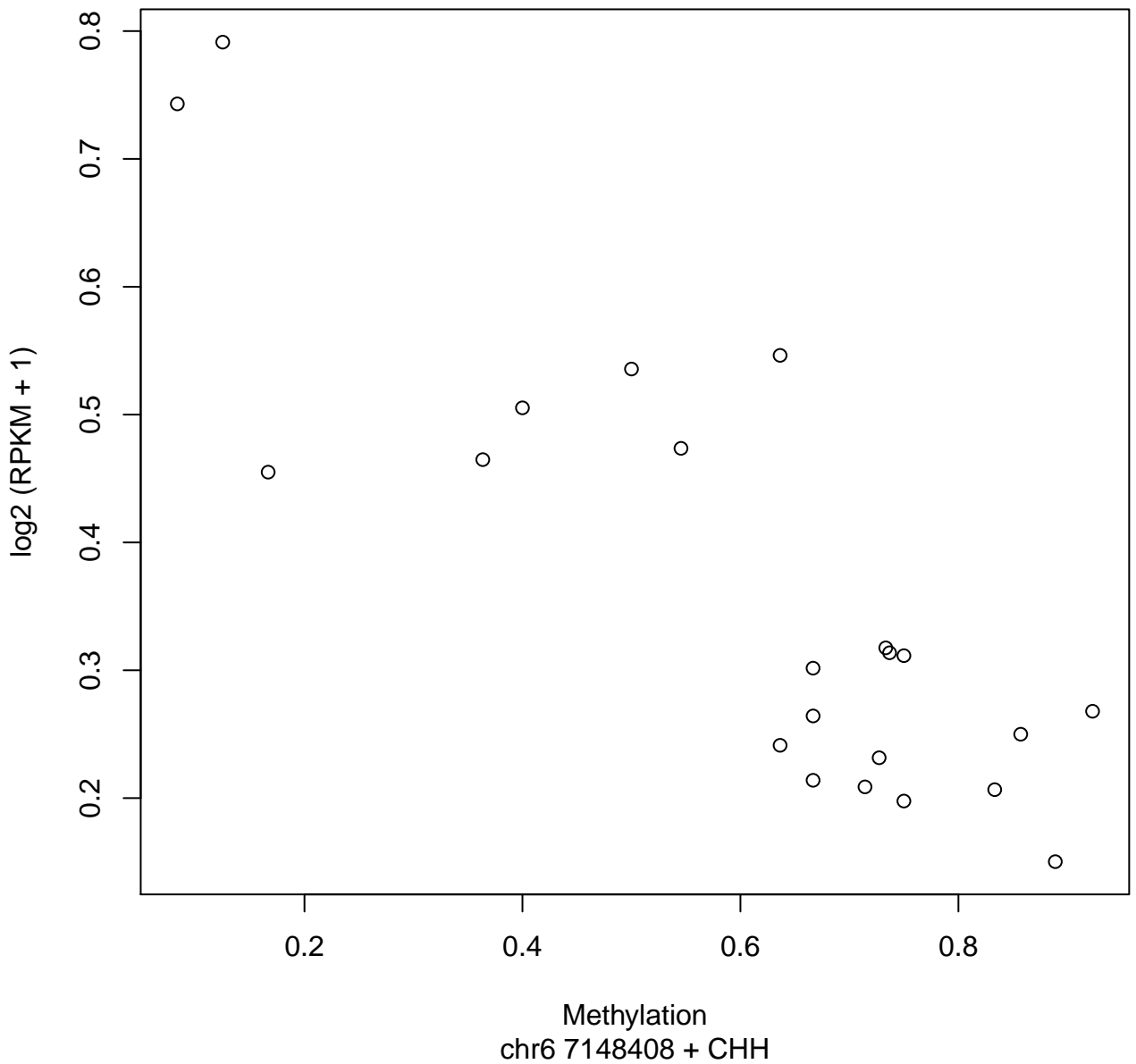


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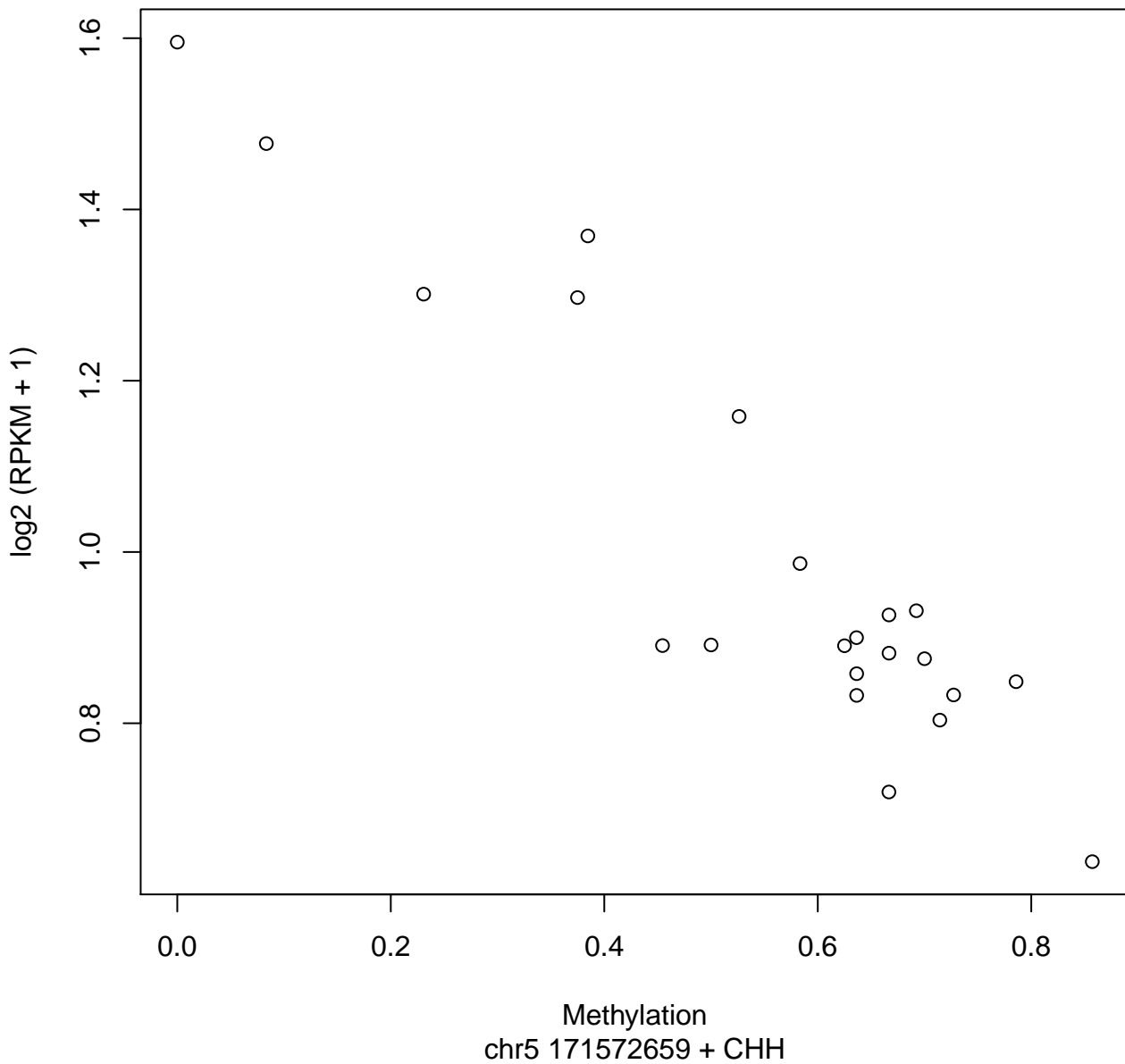


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RREB1

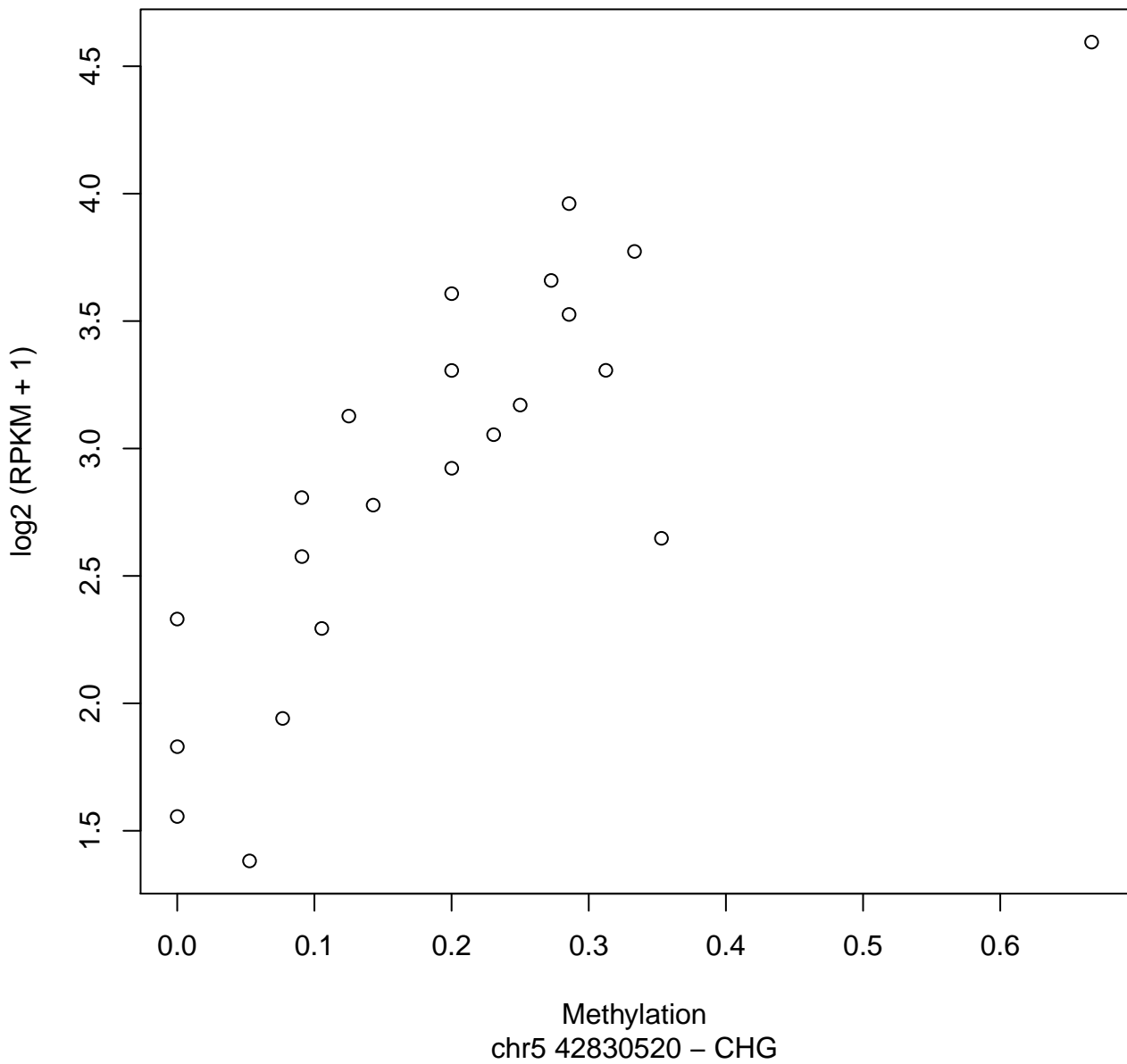


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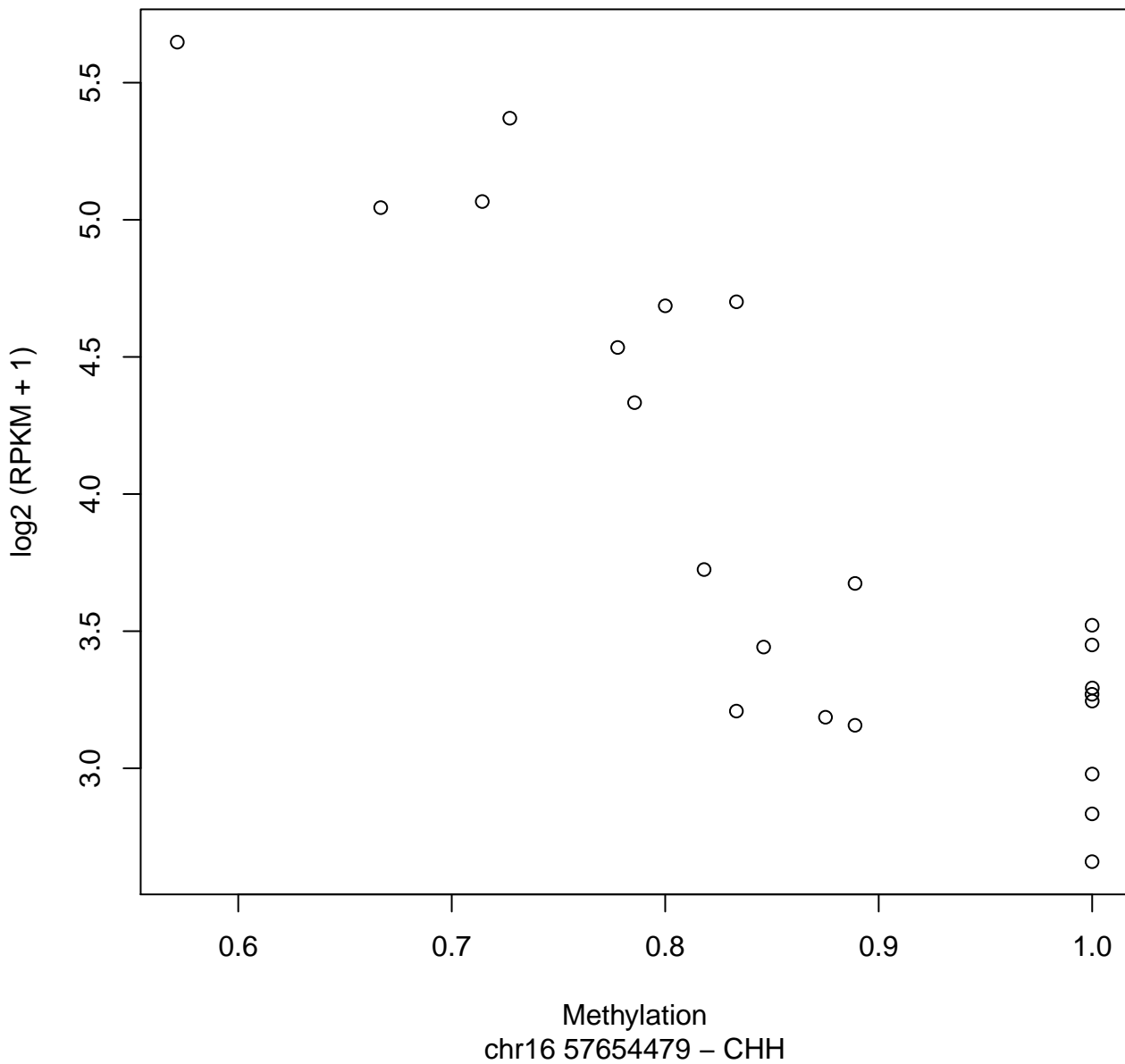


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SEPP1

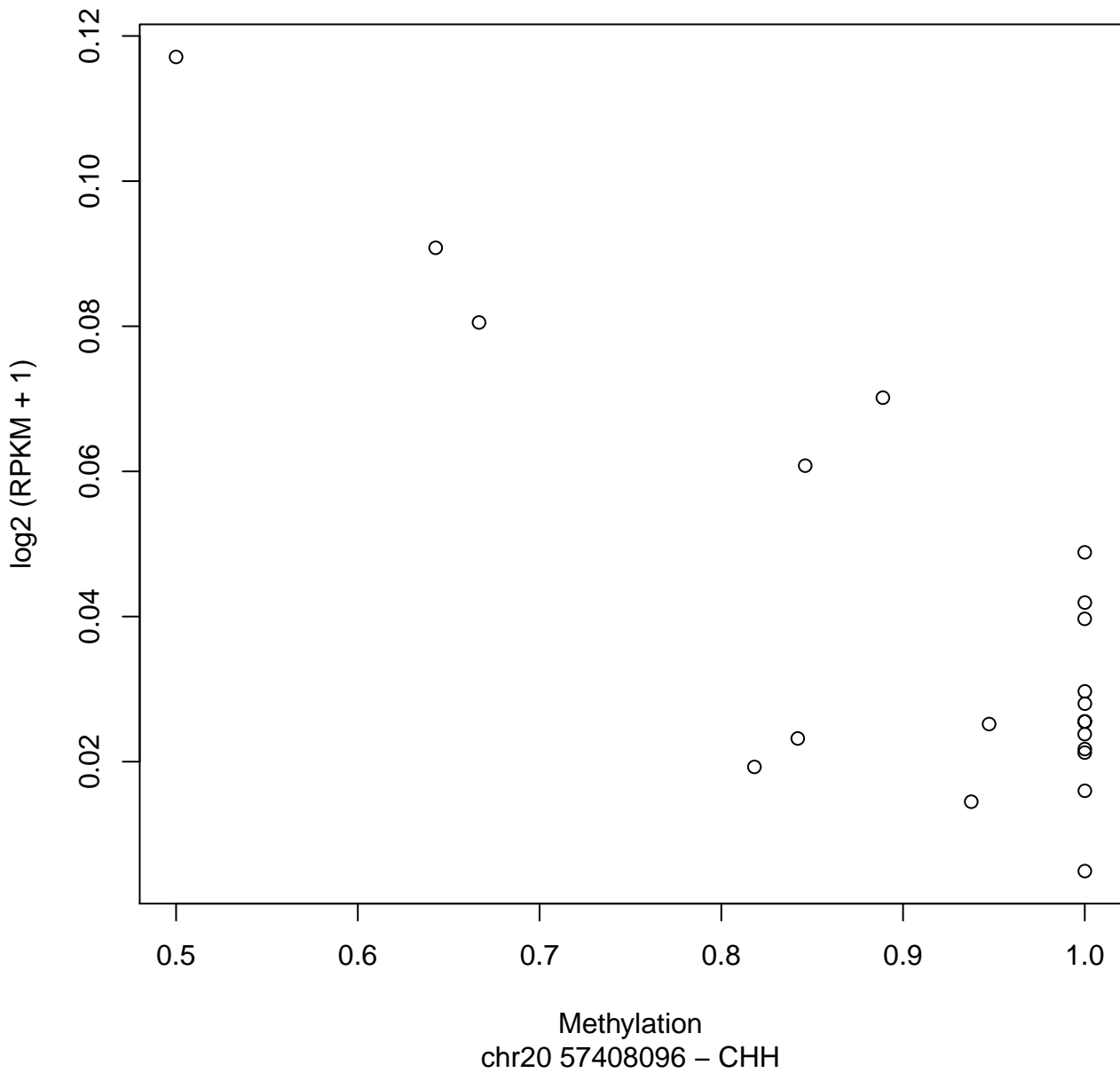


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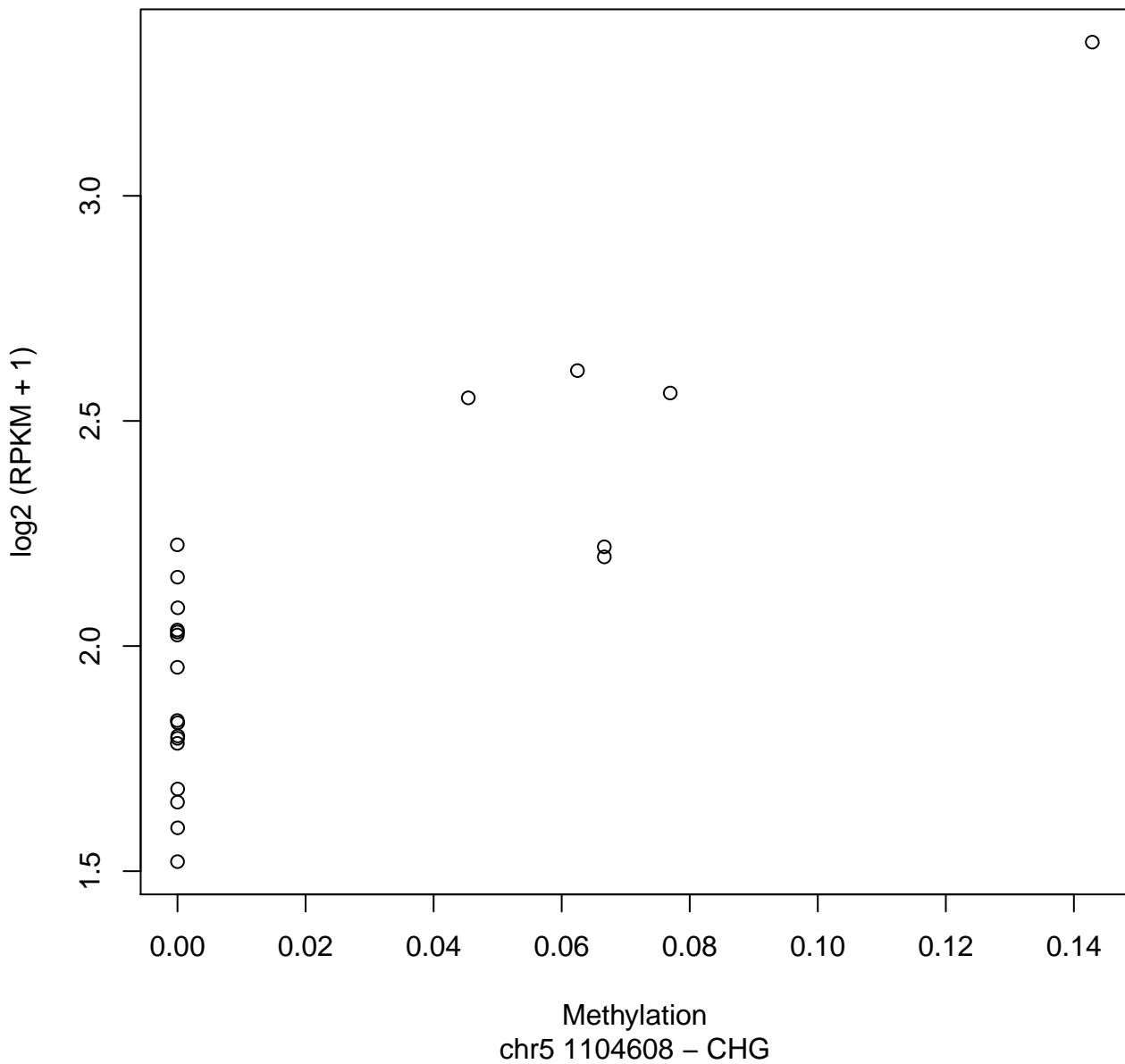


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GNAS-AS1

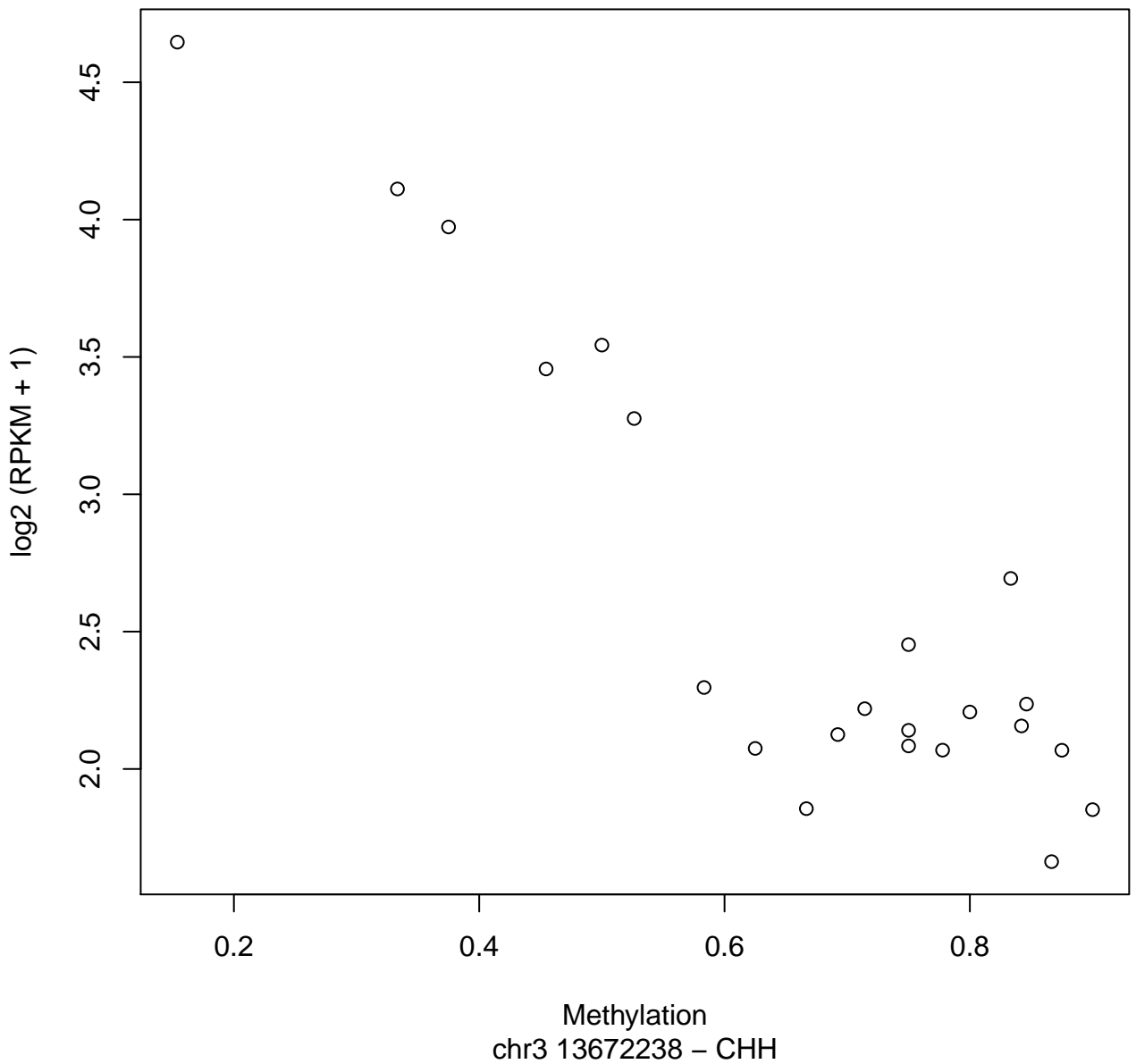


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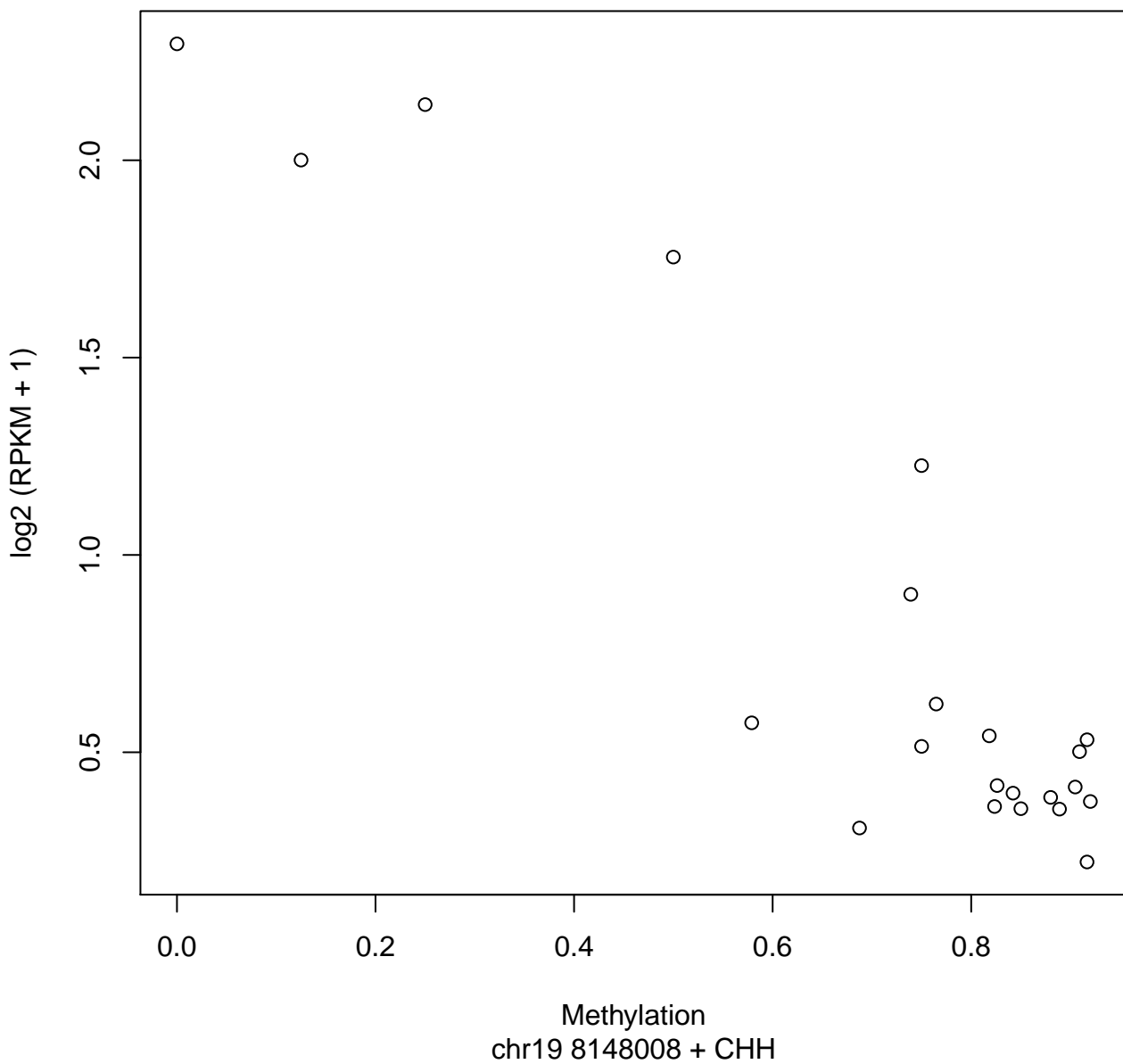


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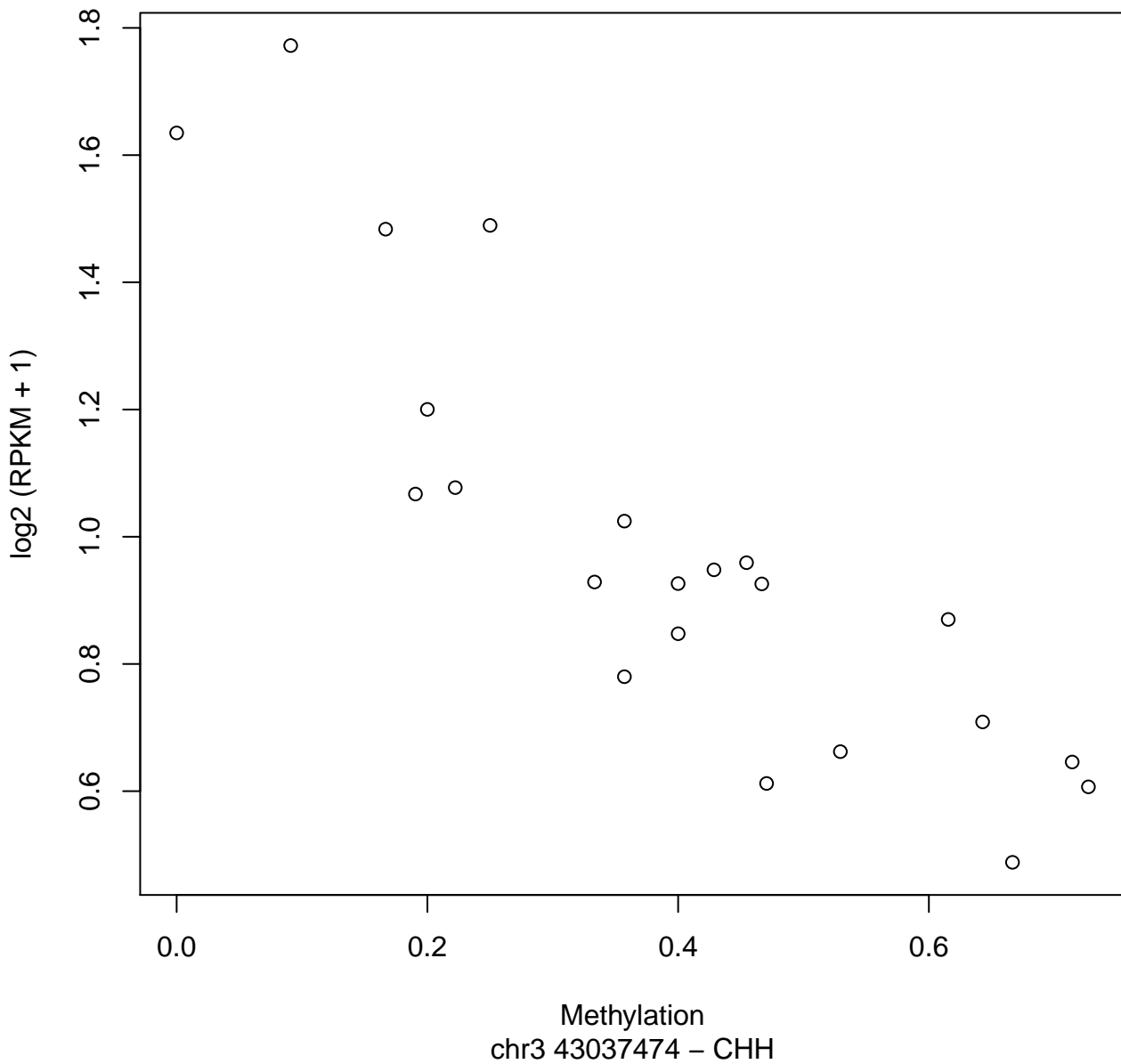


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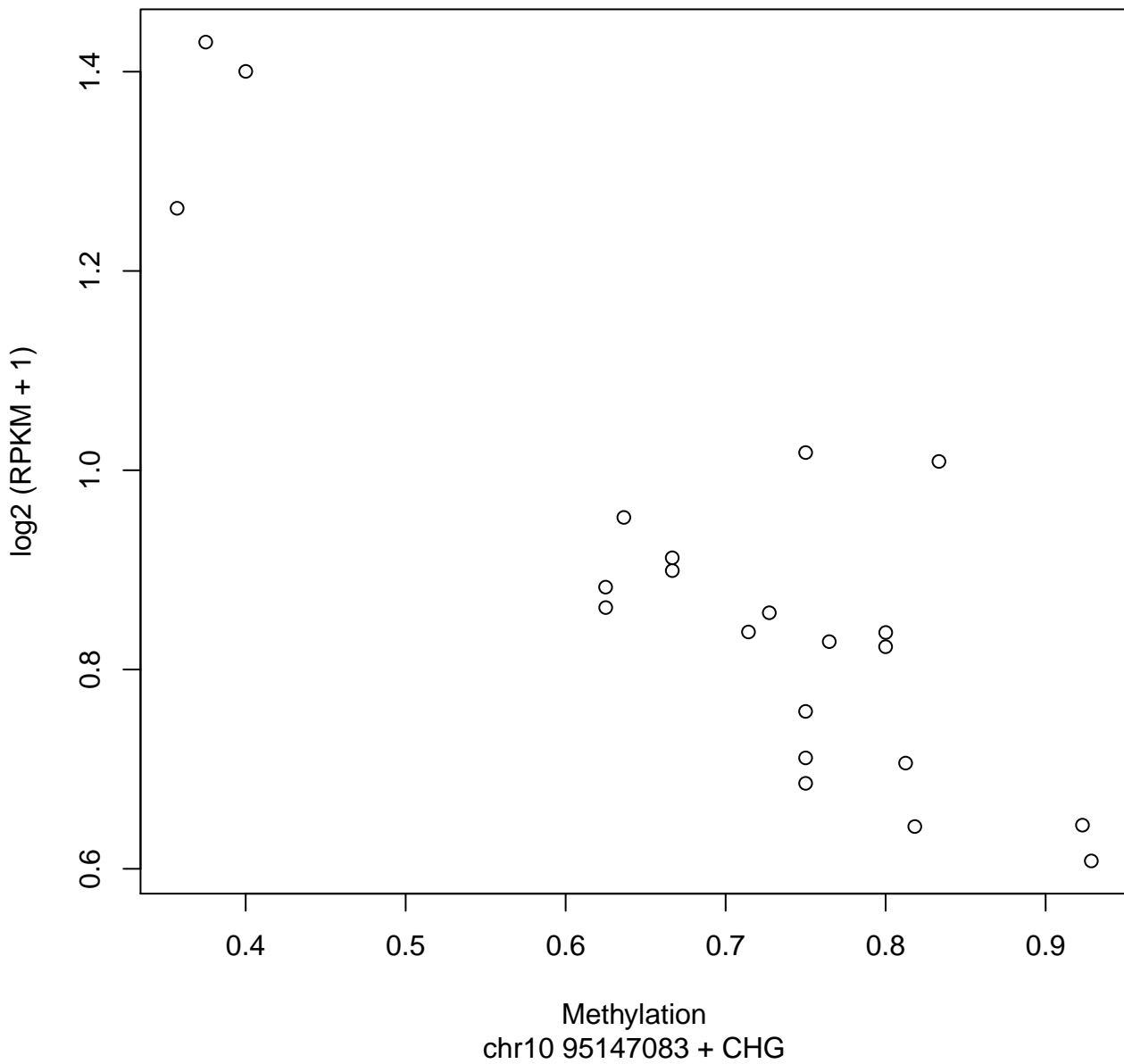
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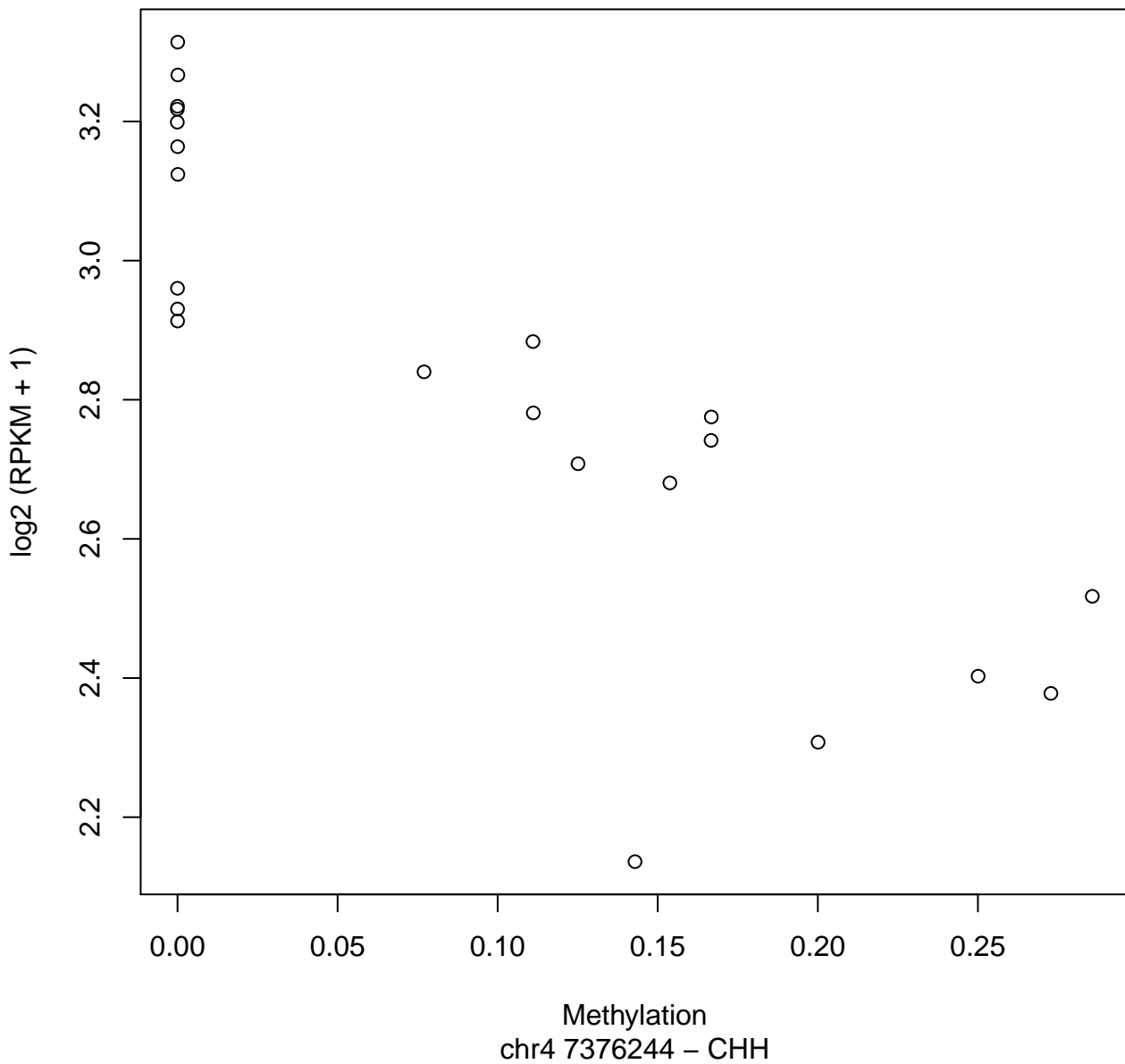
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MYOF



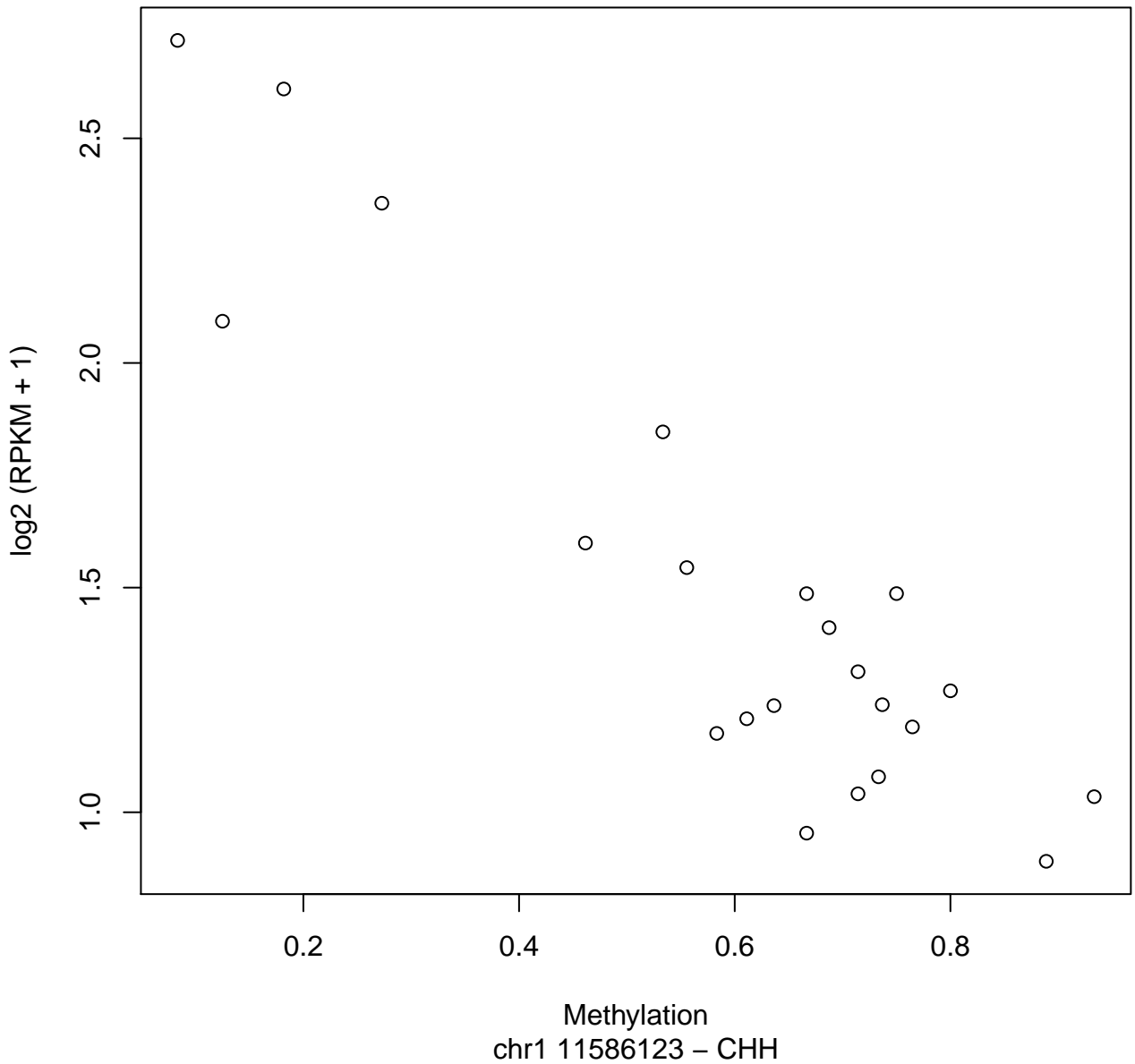
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SORCS2



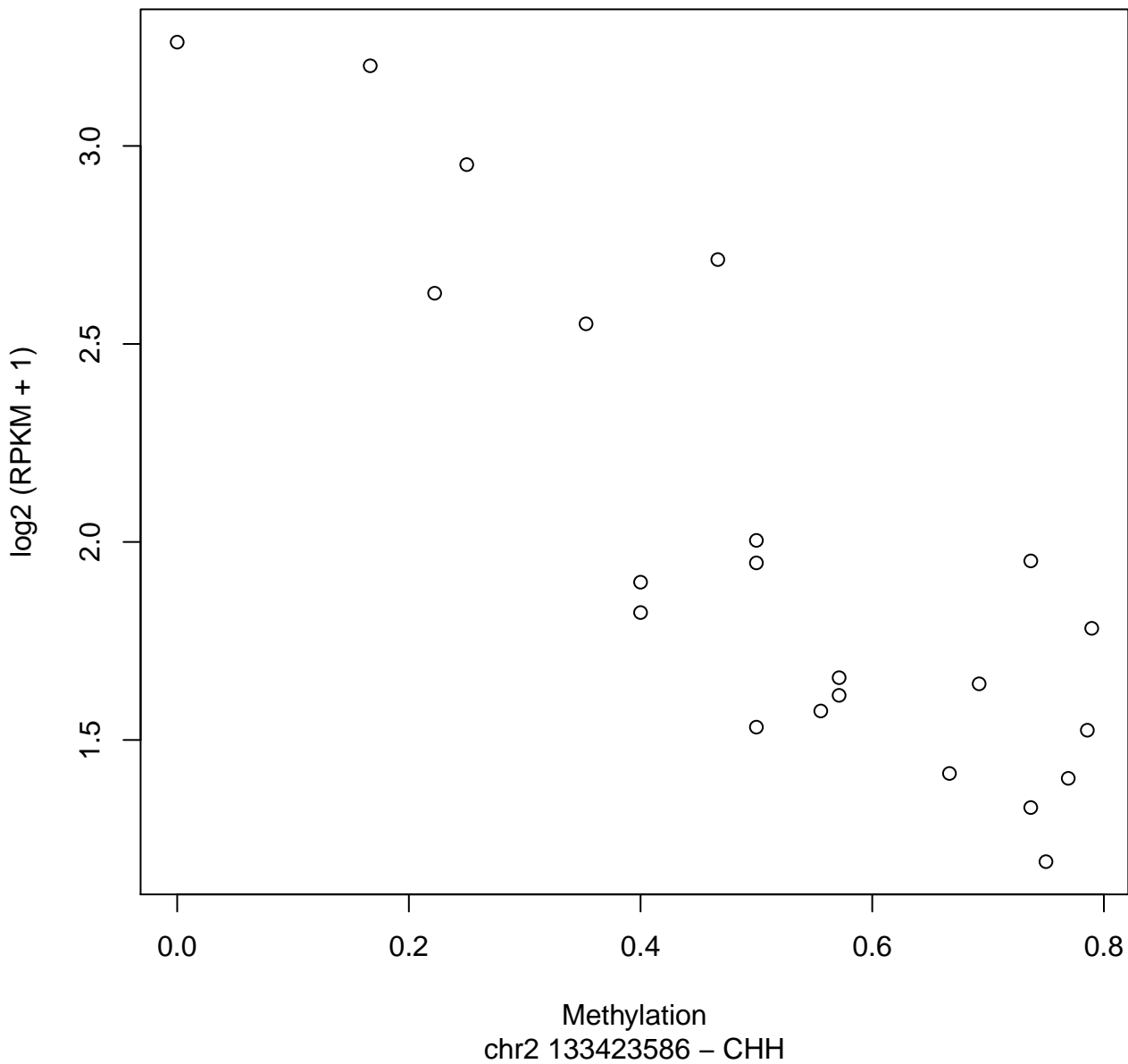
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DISP3



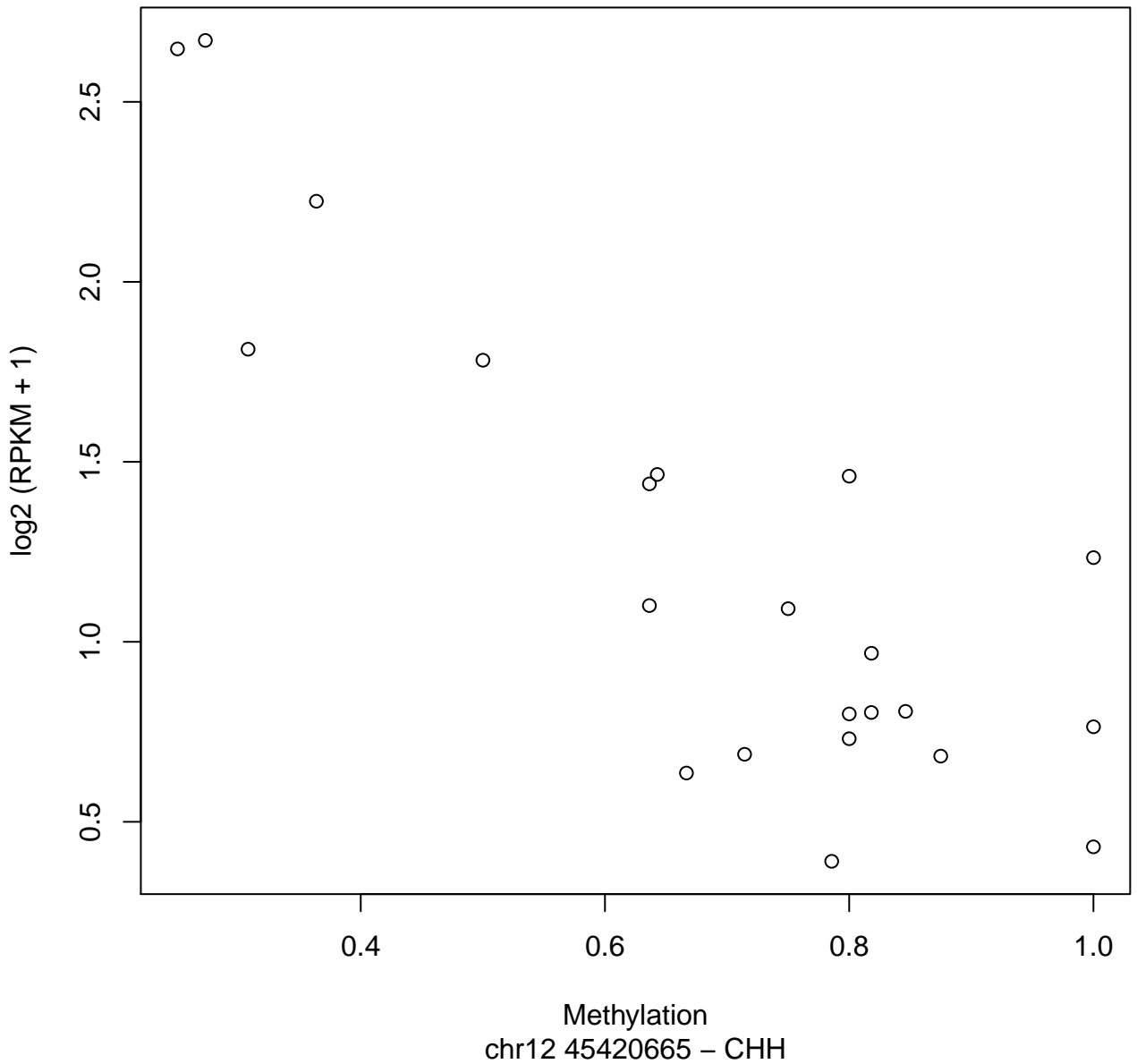
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LYPD1



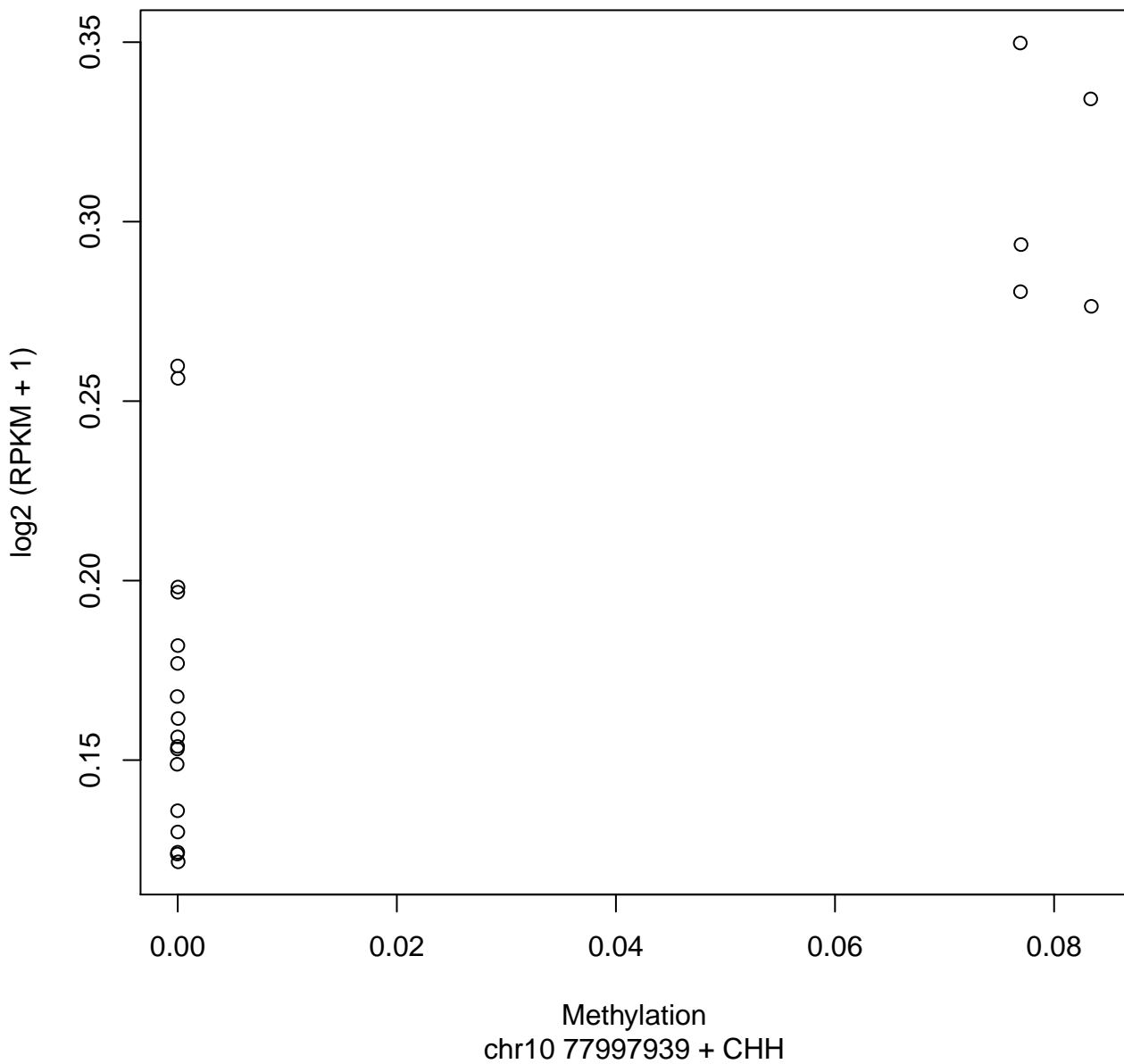
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DBX2



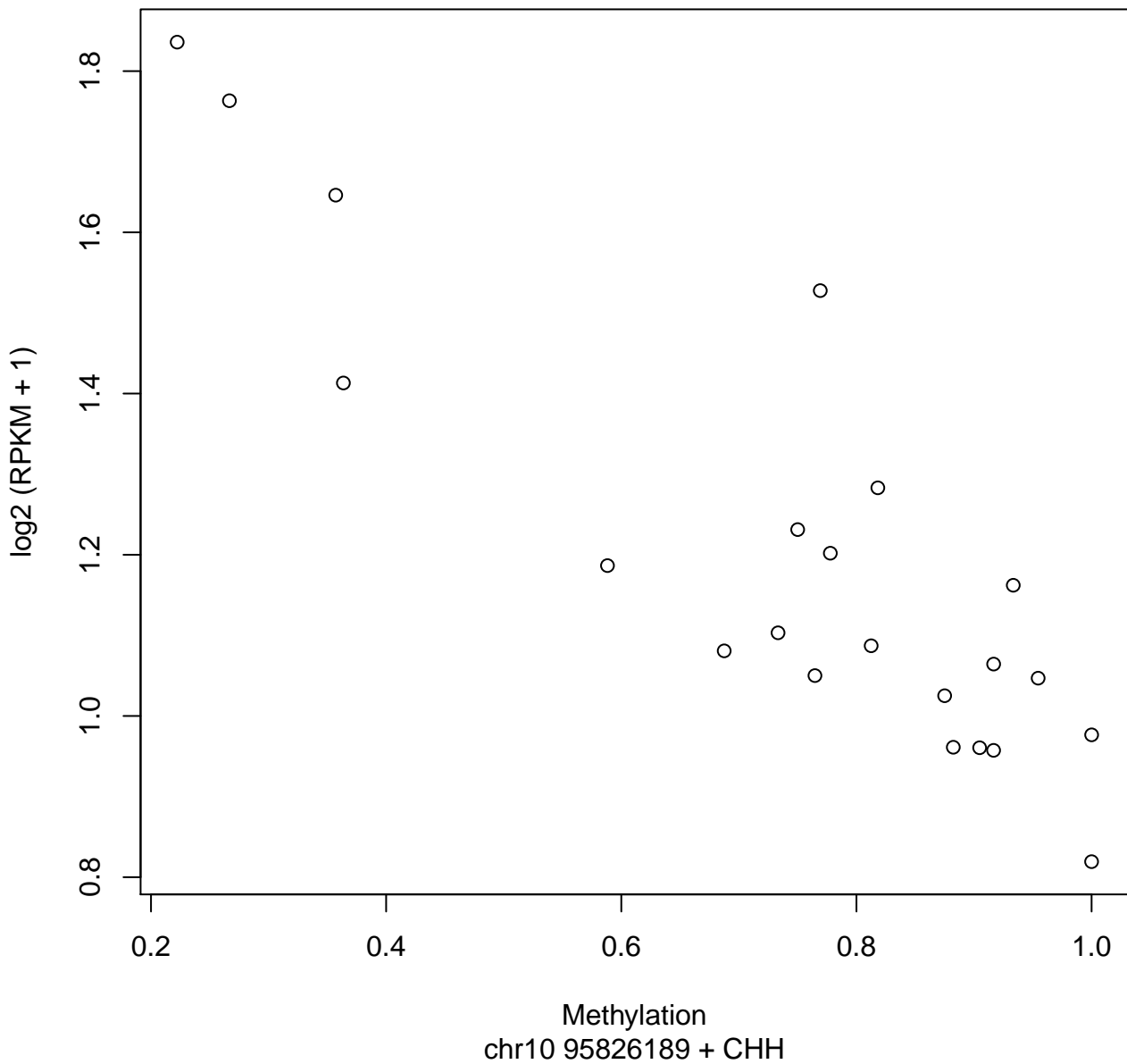
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C10orf11



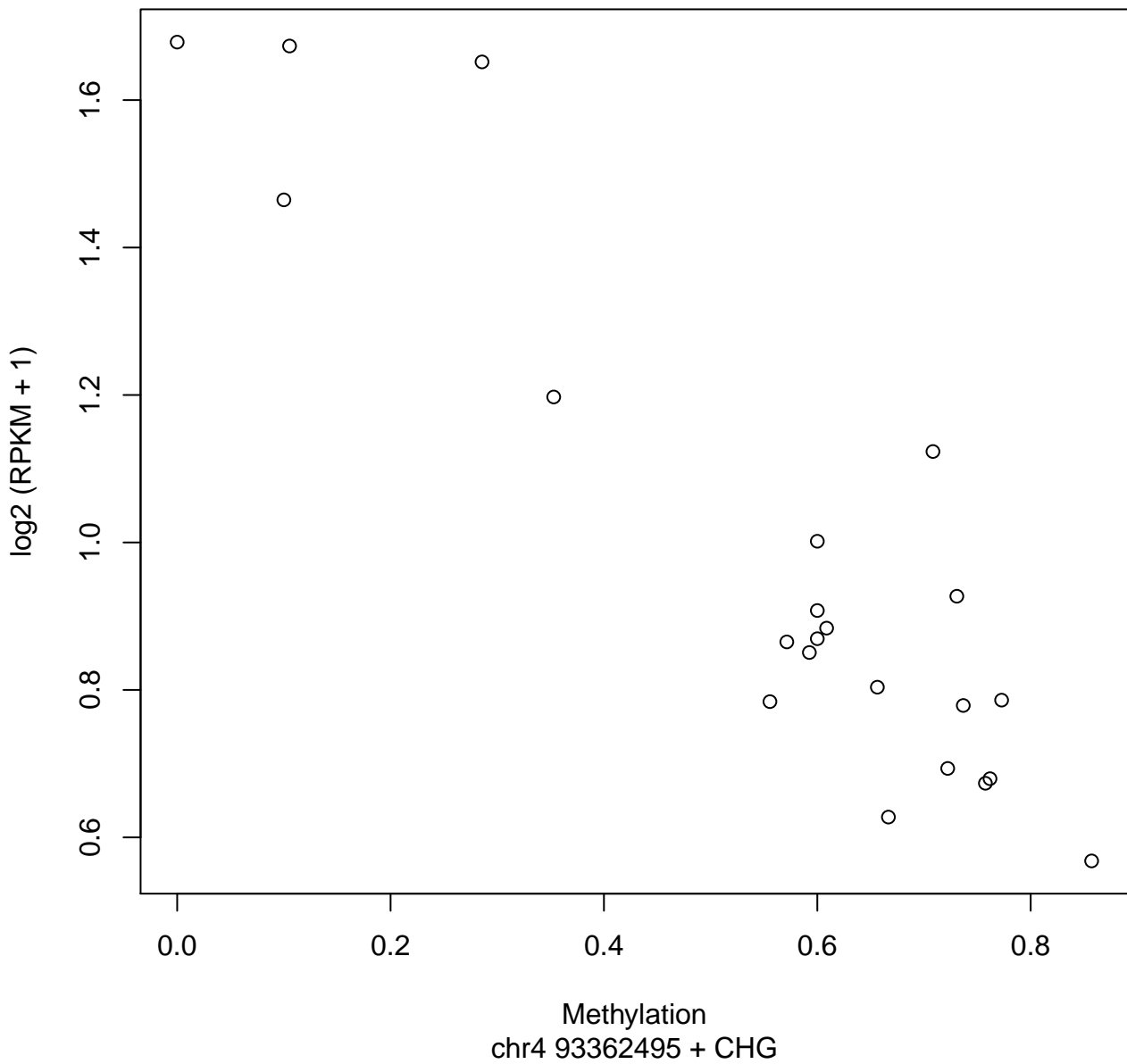
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PLCE1

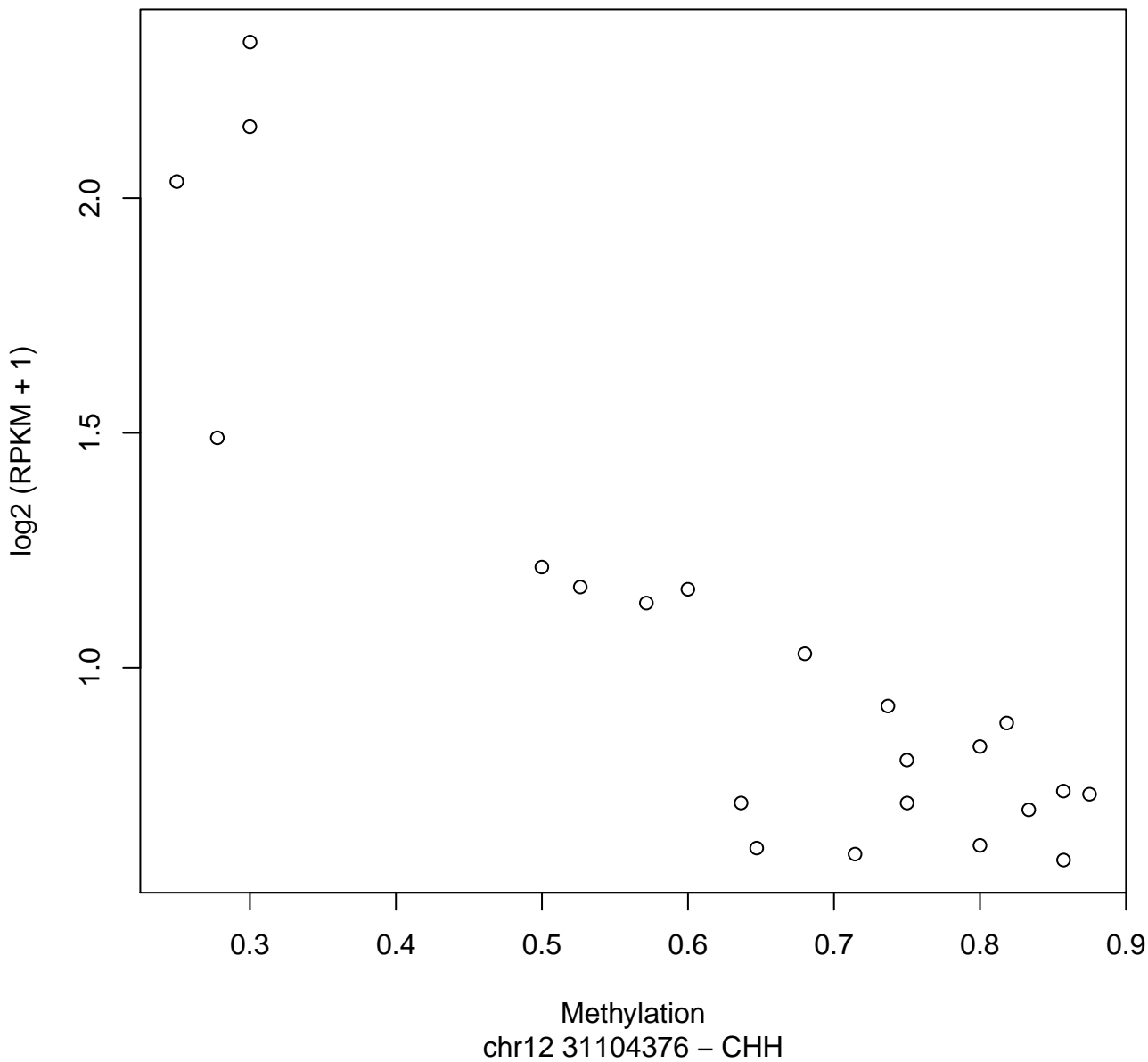


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GRID2

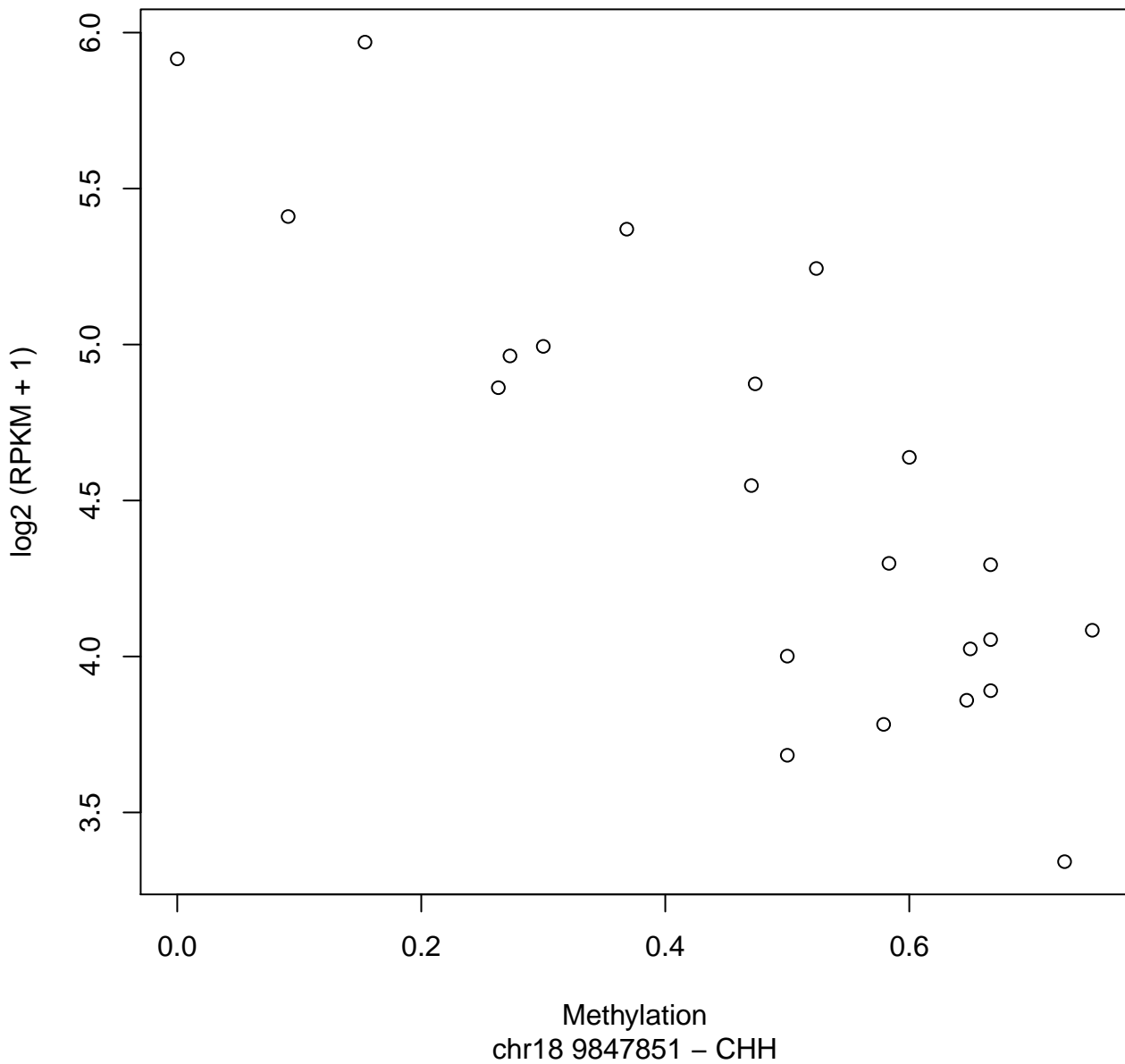


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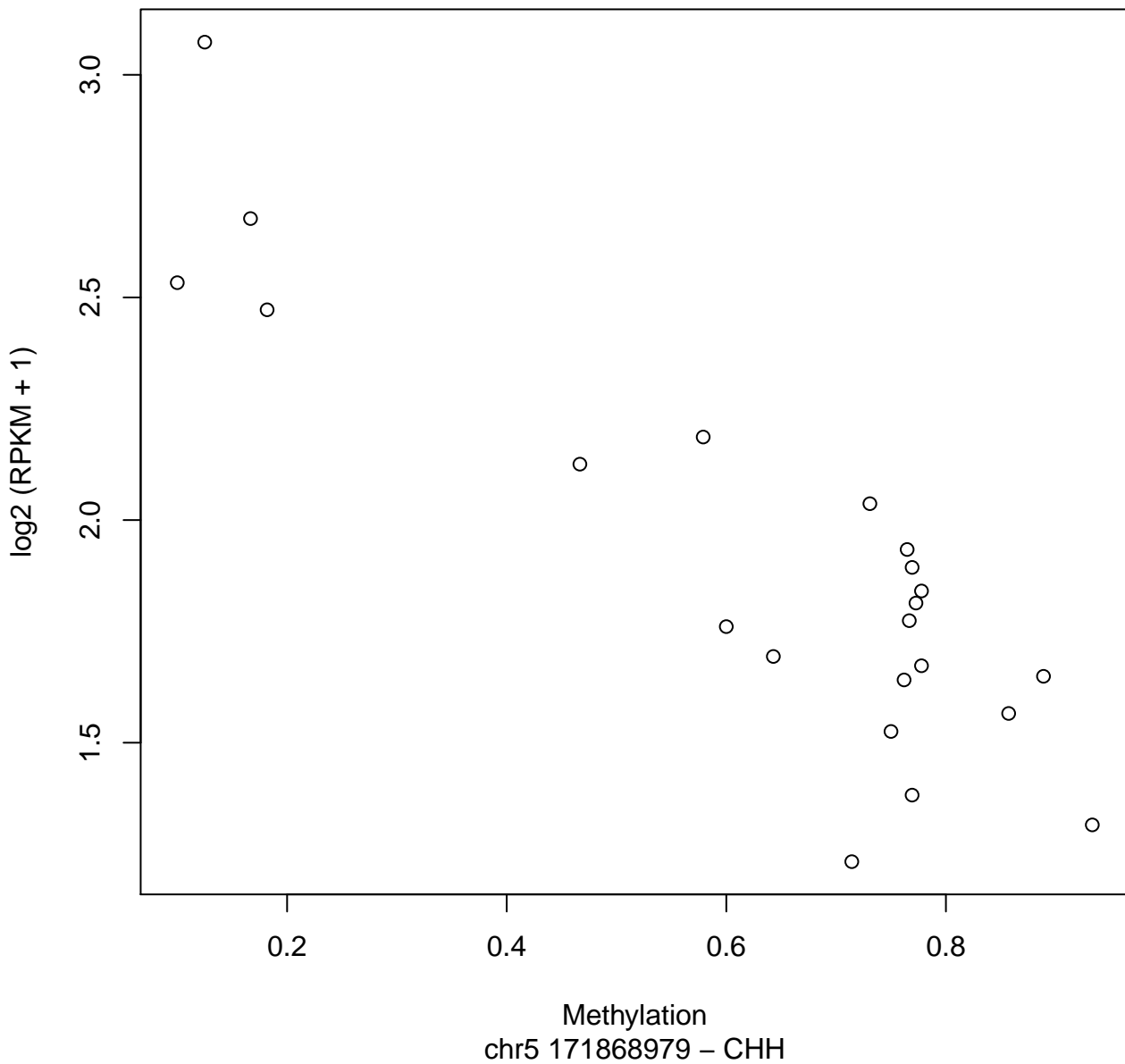
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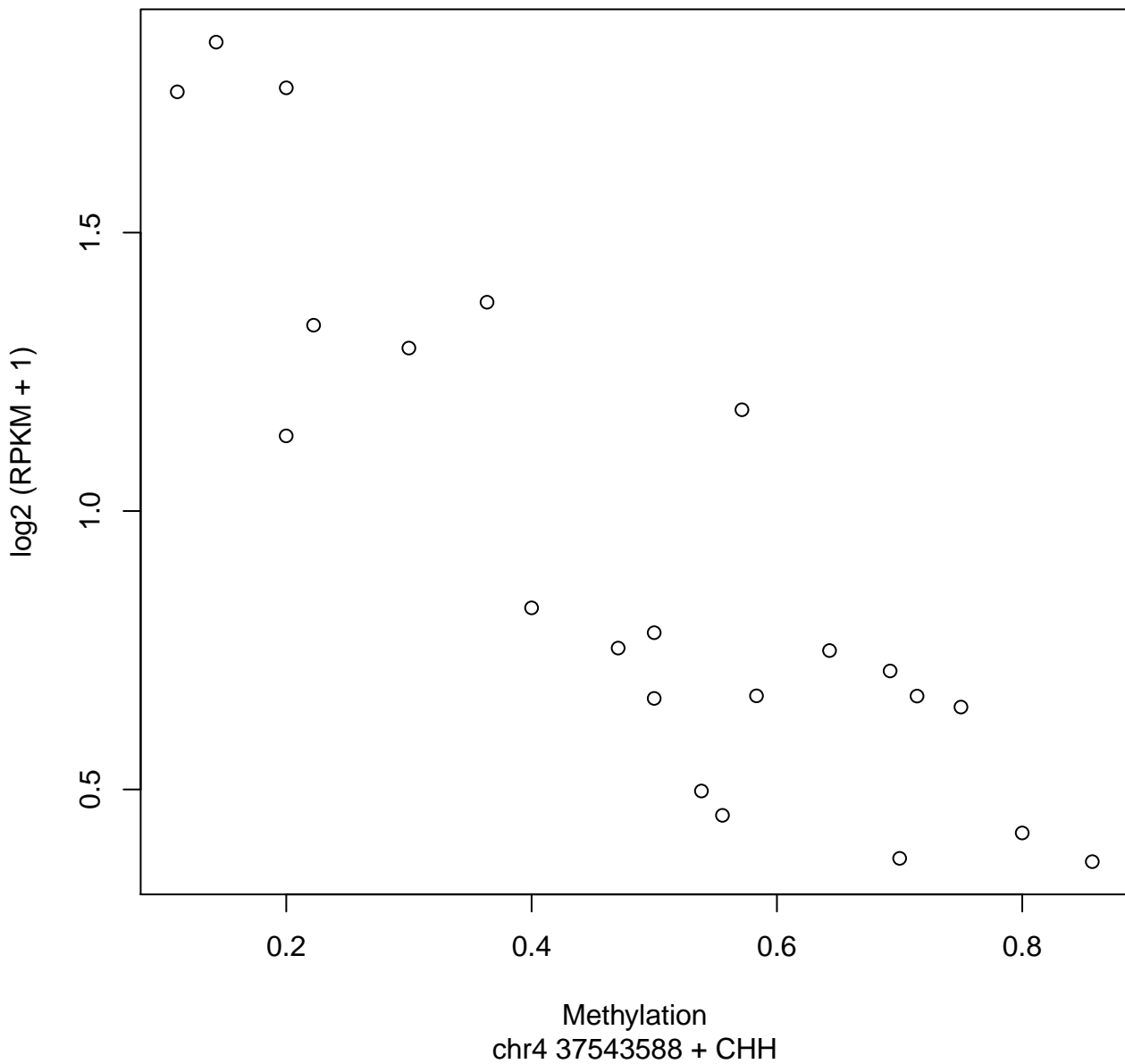
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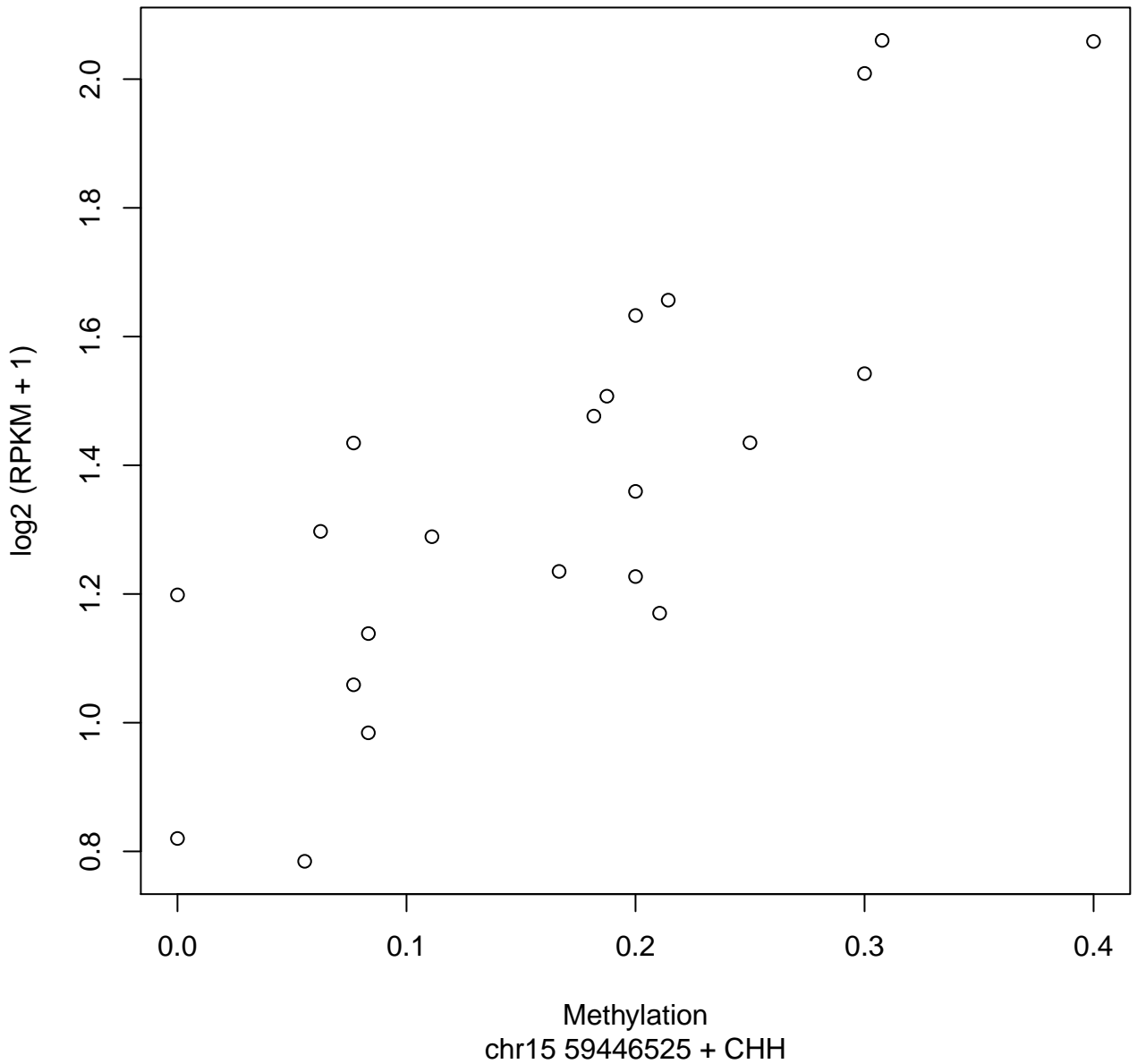
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C4orf19



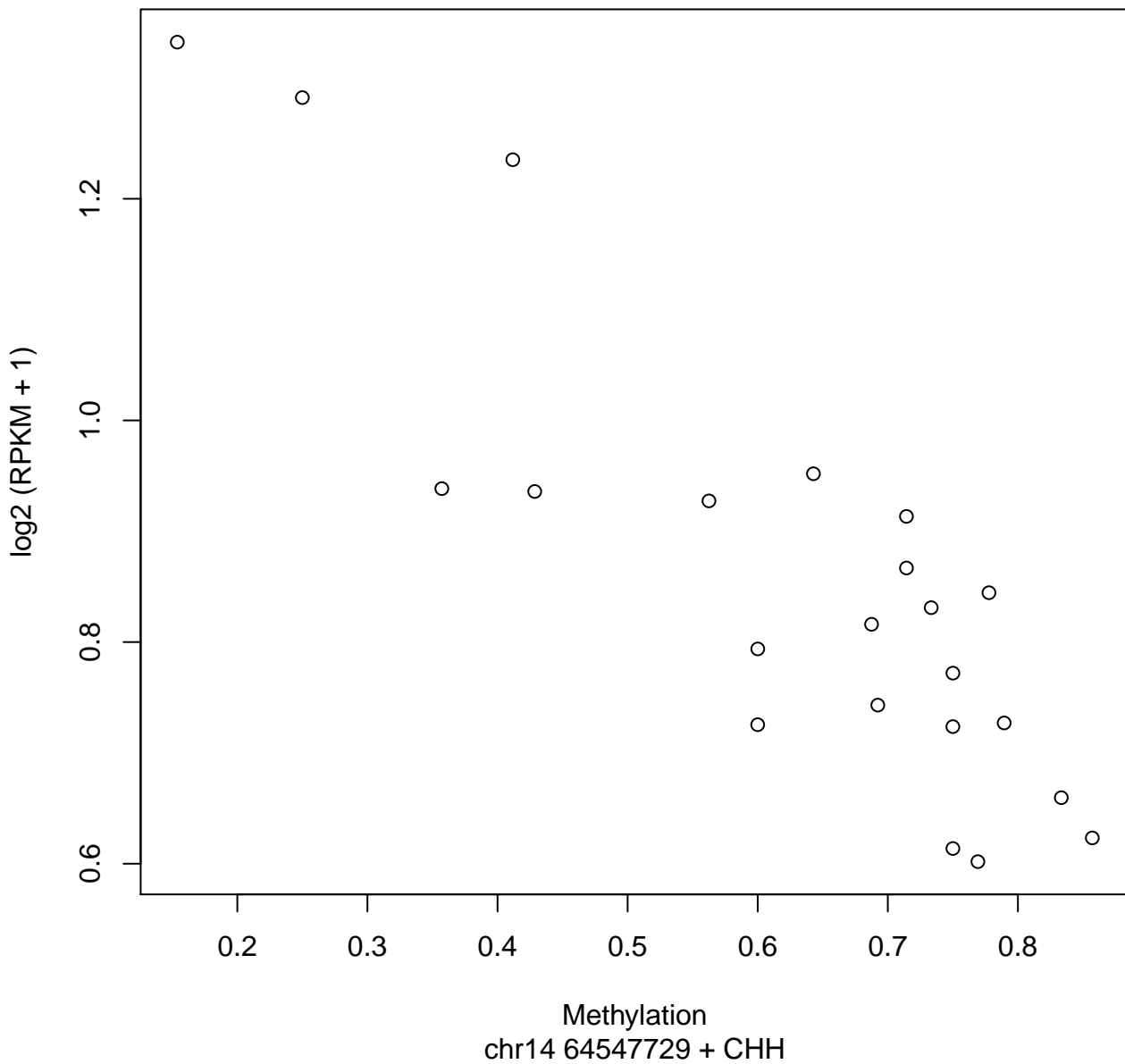
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MYO1E



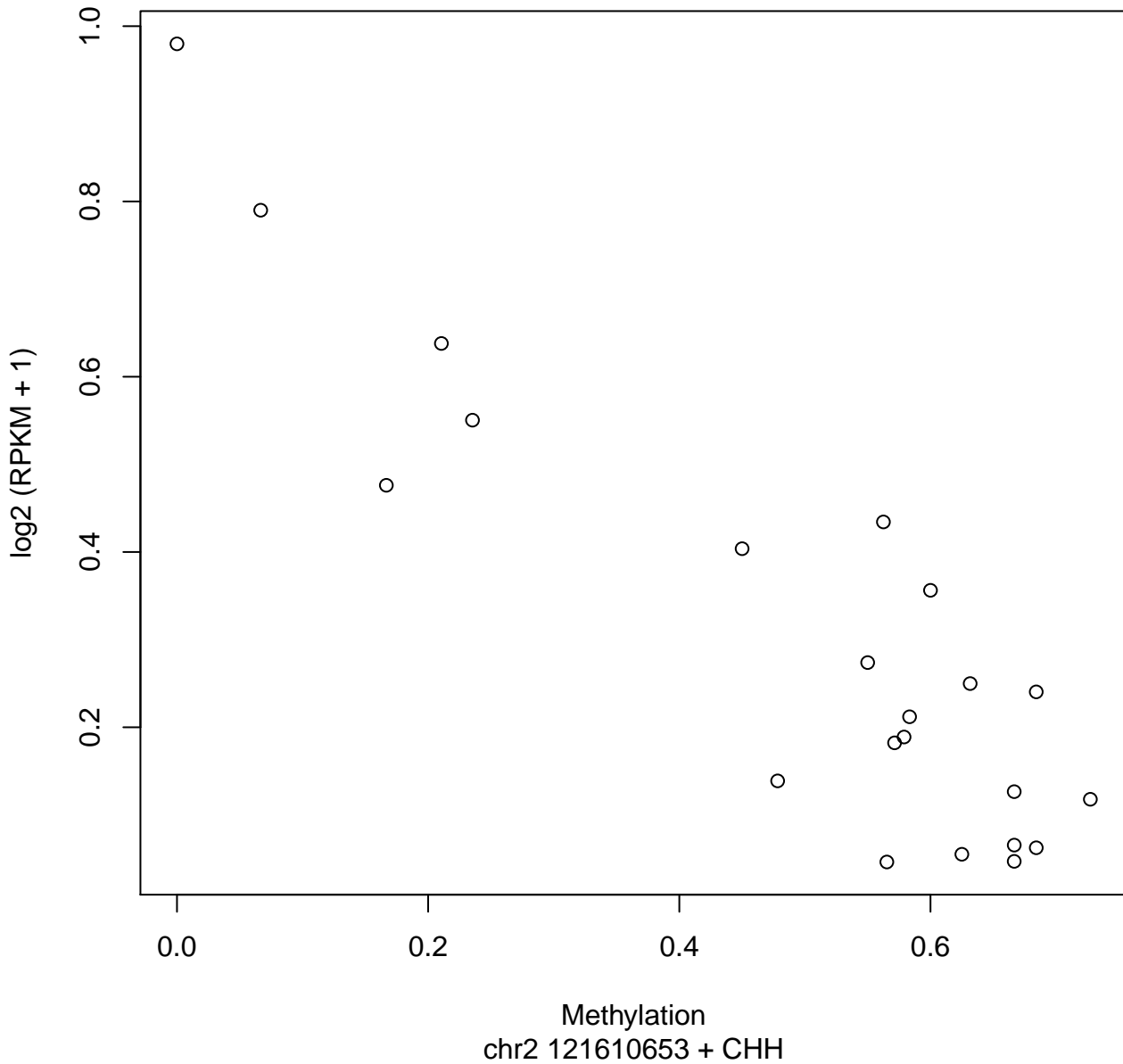
gene ENSG00000054654.15_2 FDR 0.000718

SYNE2

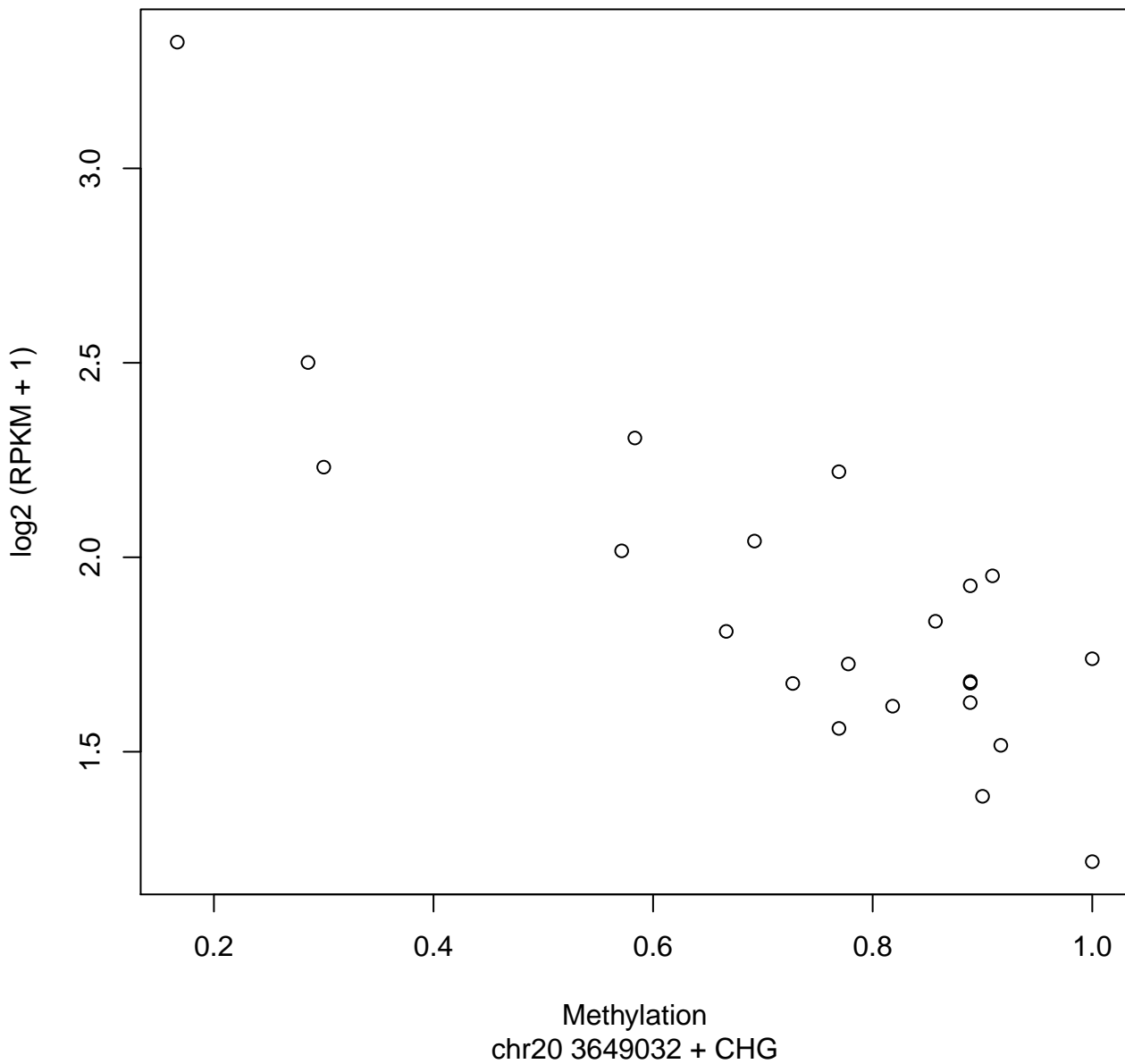


gene ENSG00000074047.21_2 FDR 8.37e-05

GLI2

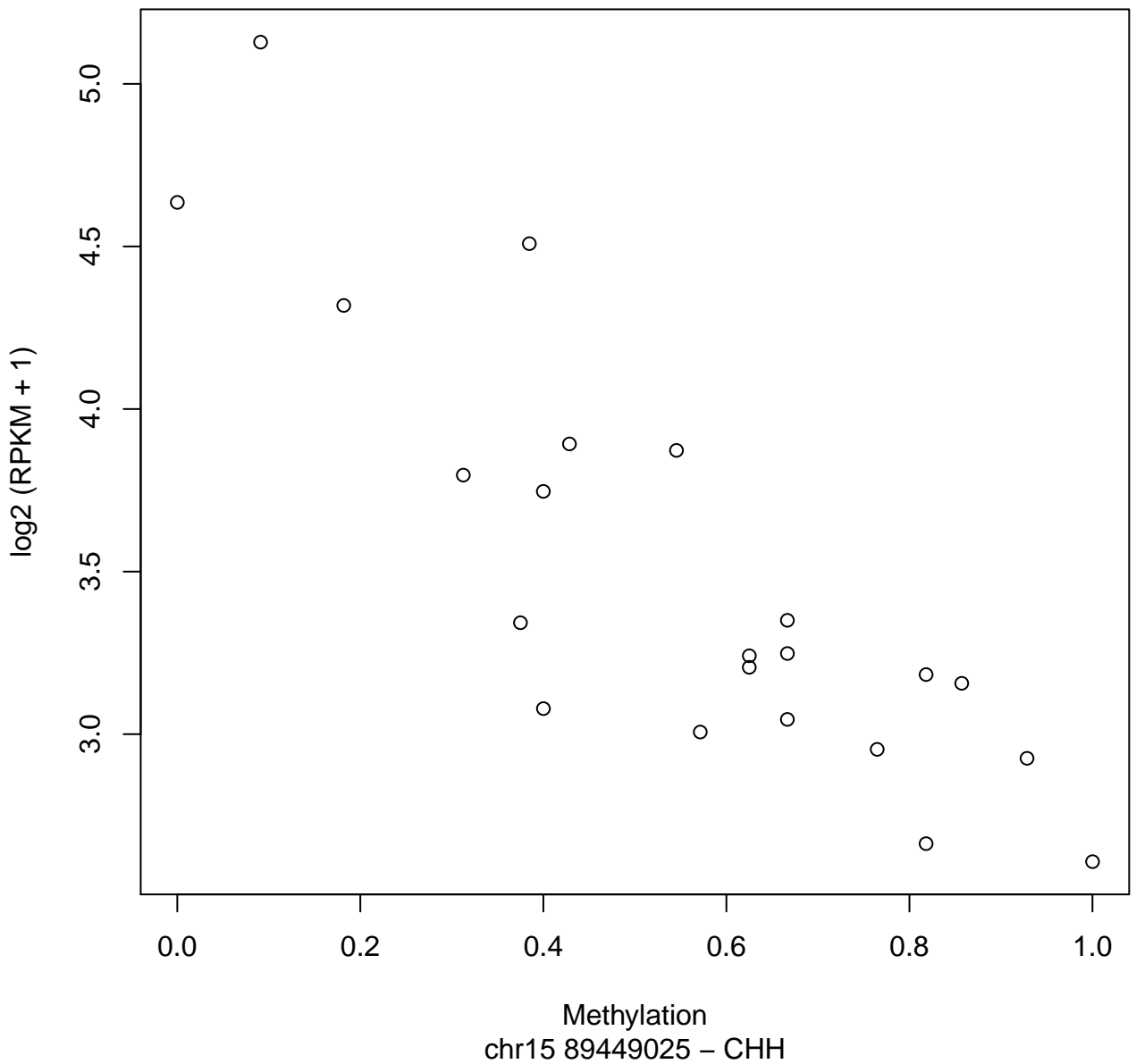


gene ENSG00000149451.17_1 FDR 0.00362
ADAM33



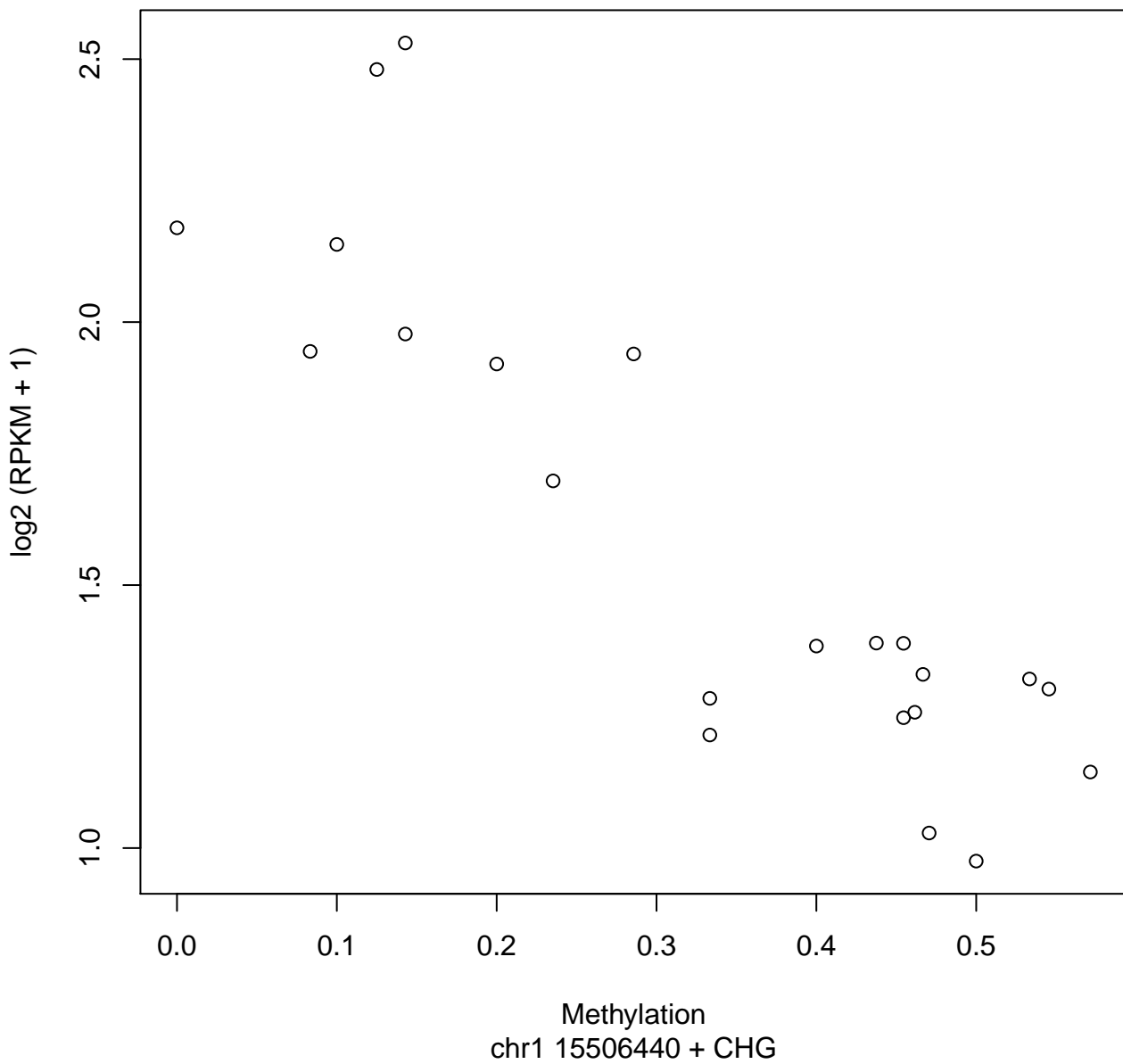
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MFGE8



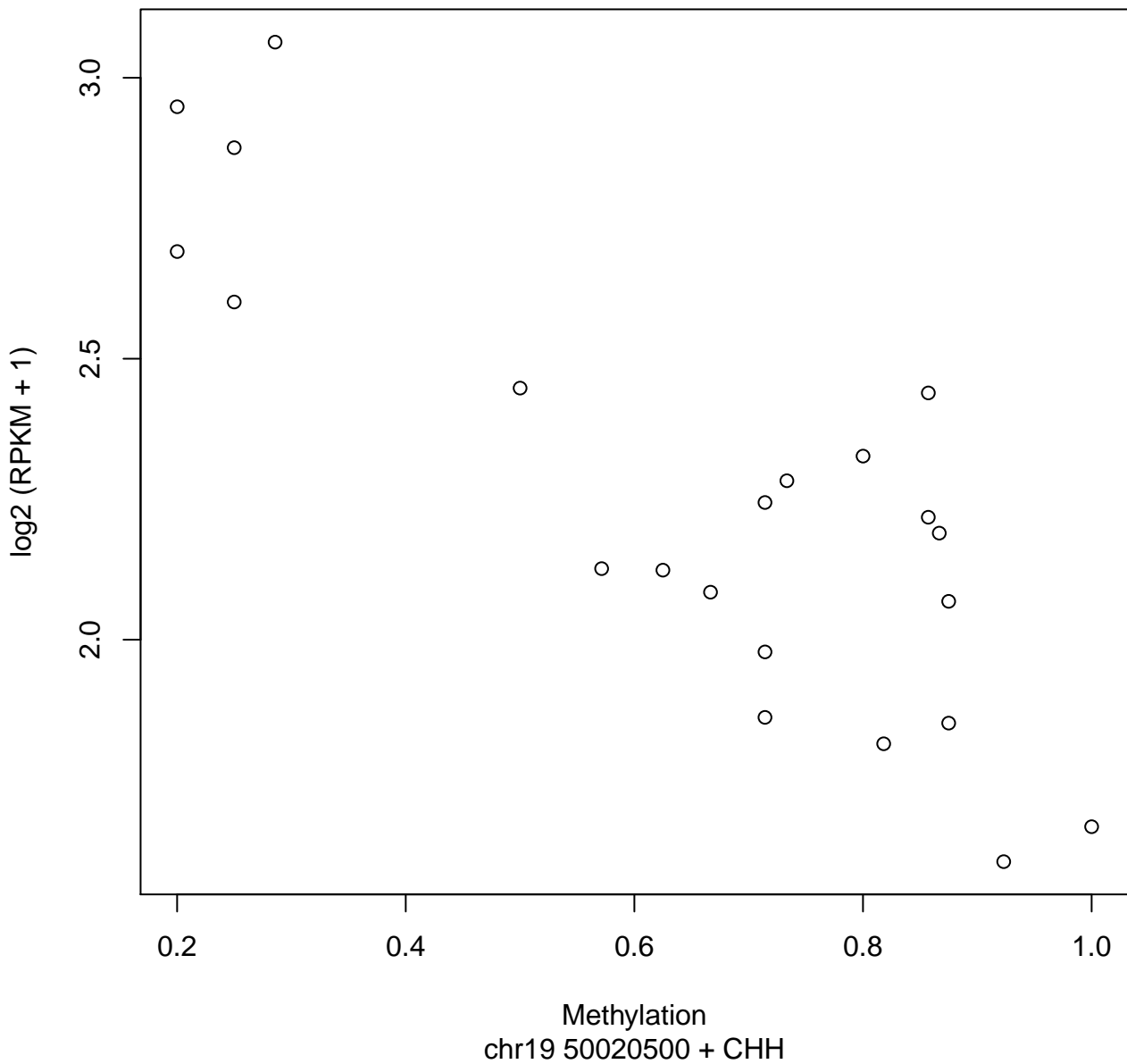
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TMEM51

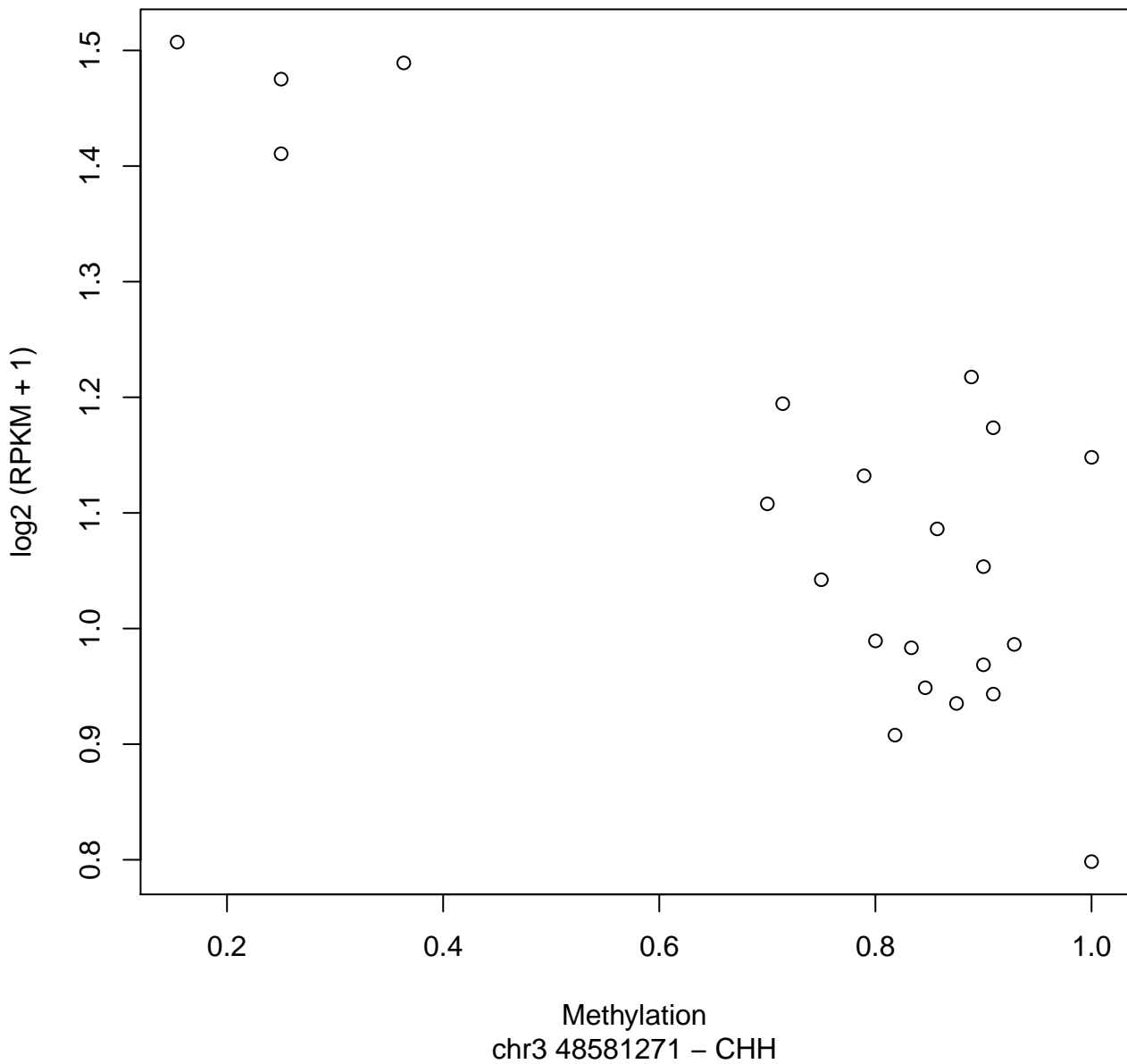


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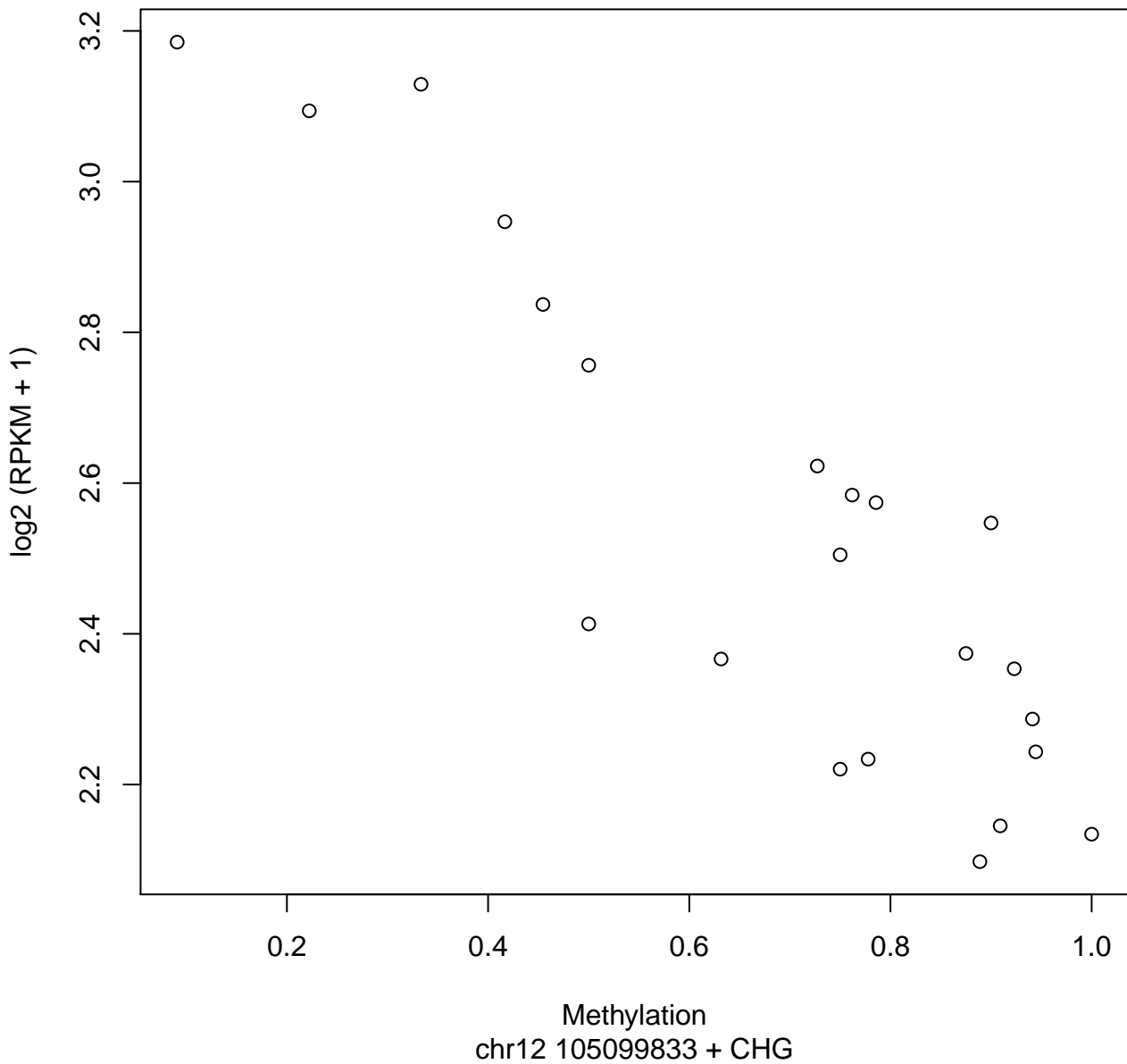
FCGRT



gene ENSG00000114268.11_2 FDR 0.00178
PFKFB4

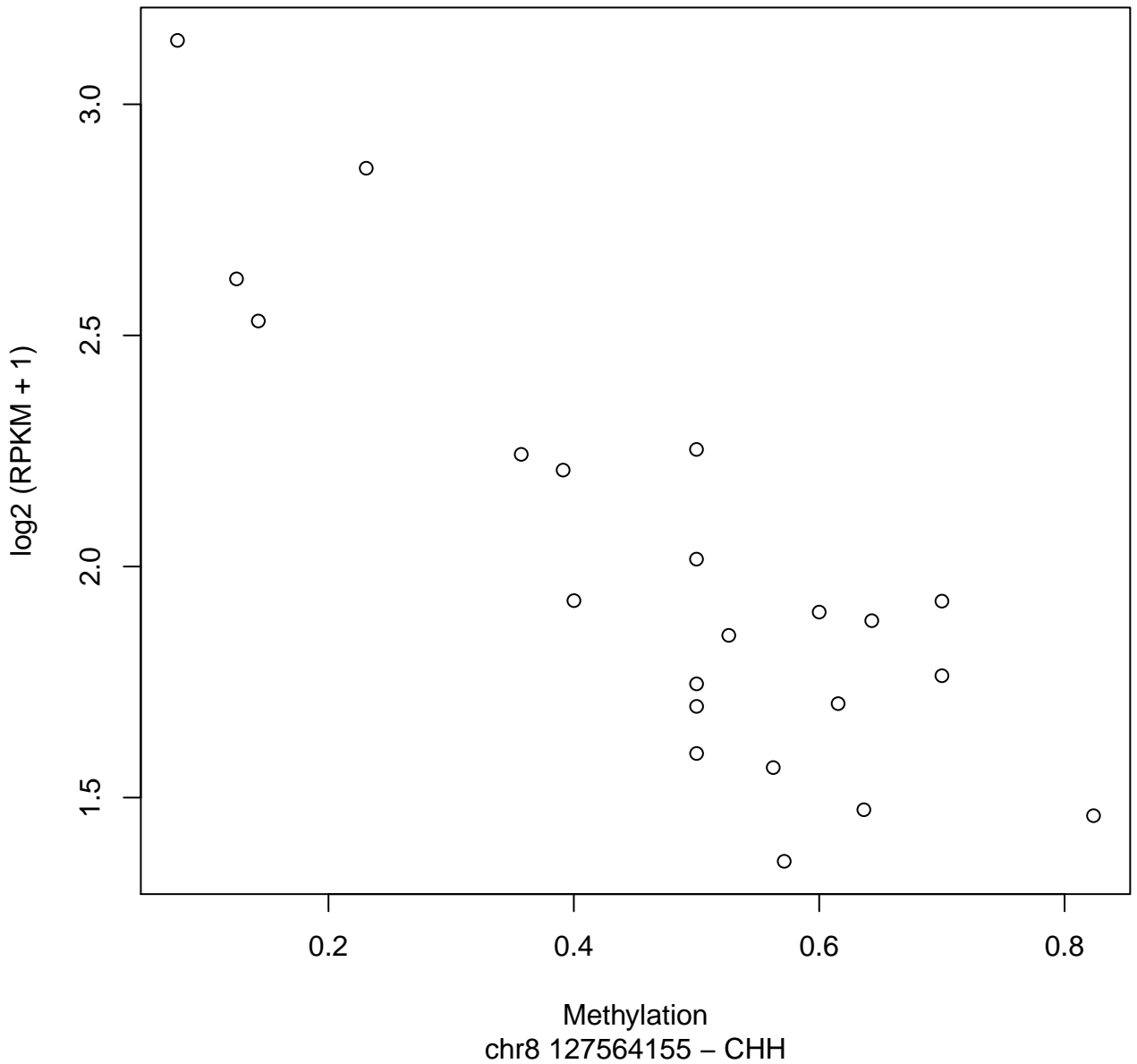


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CHST11



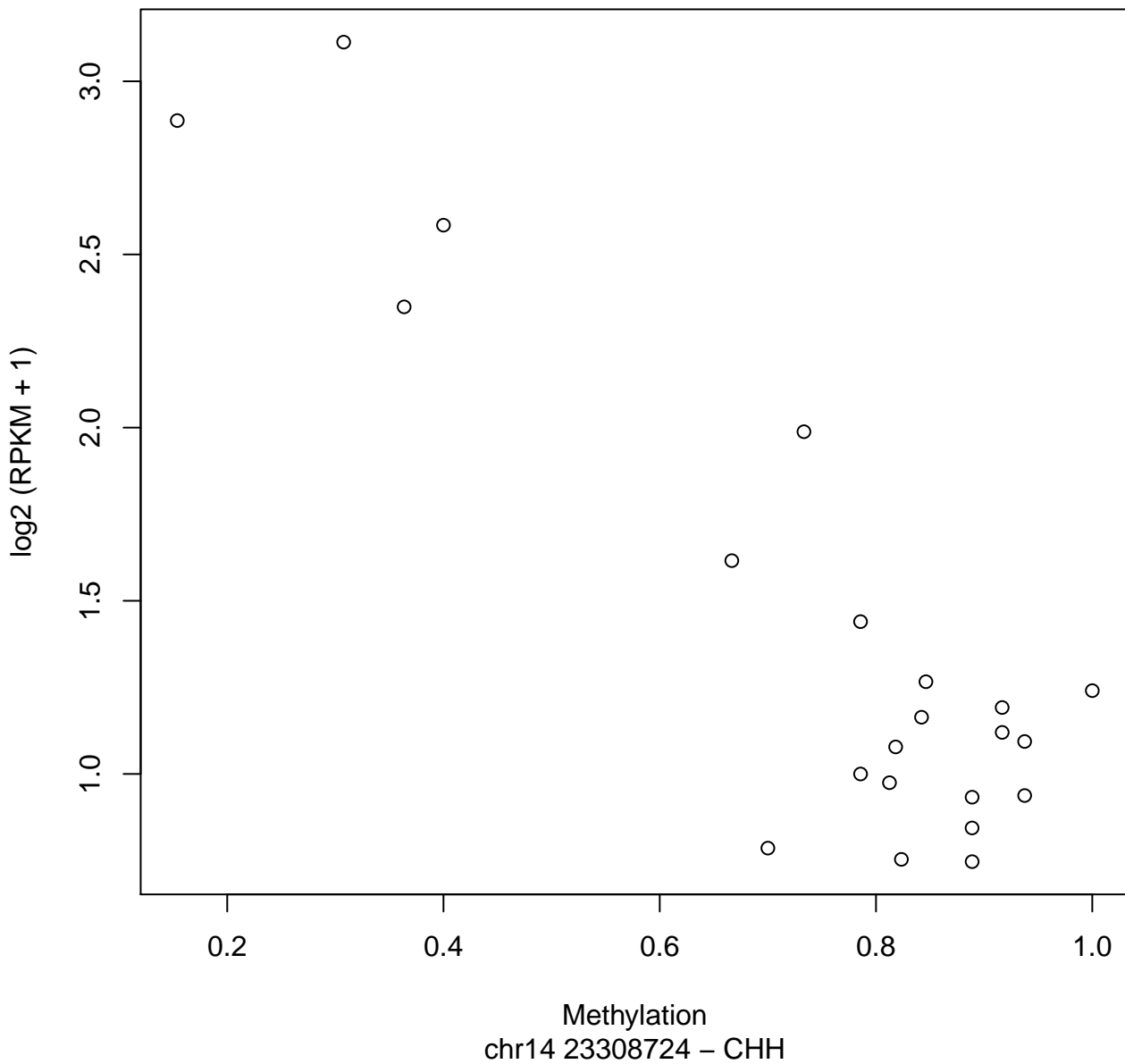
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FAM84B



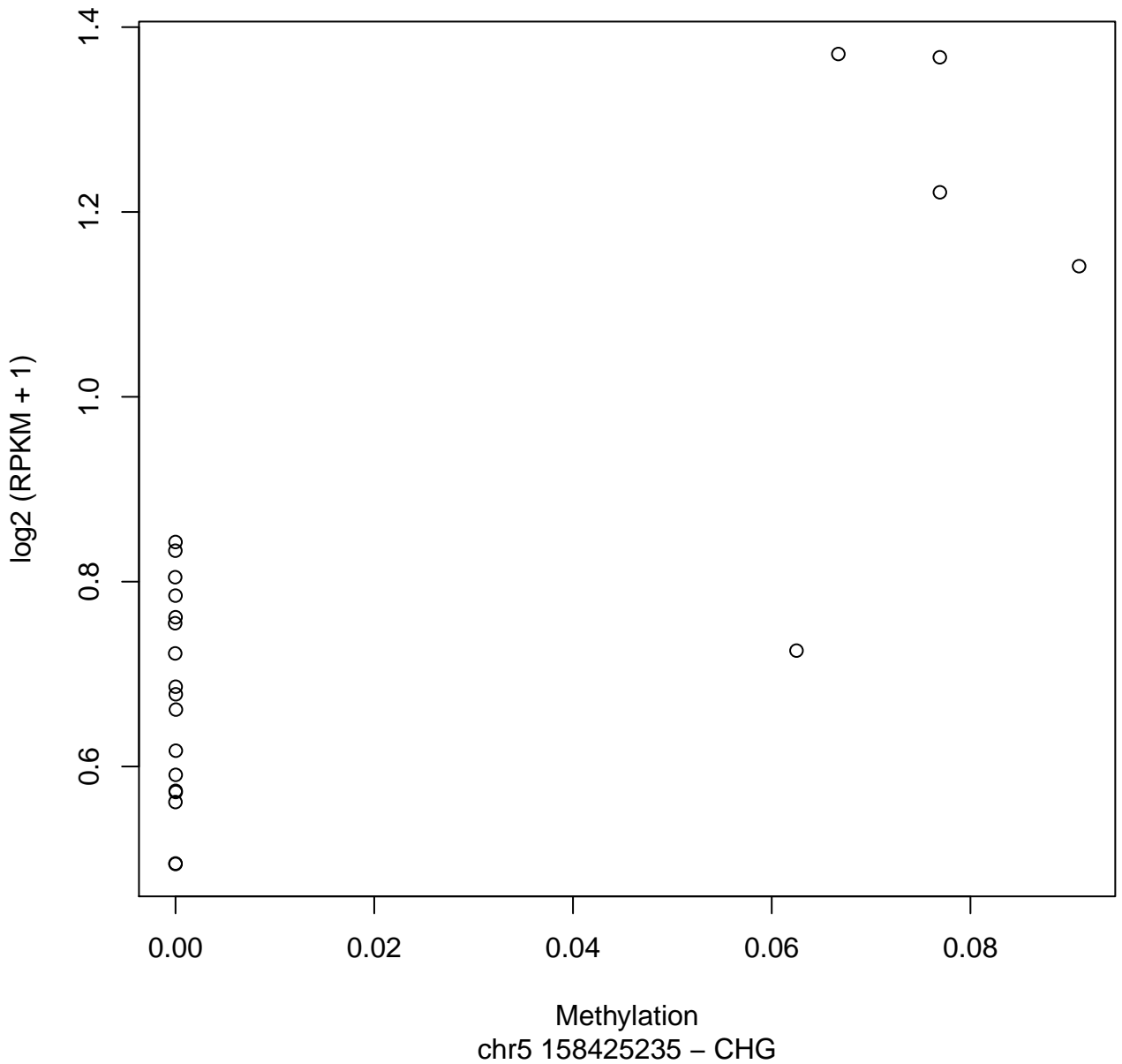
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MMP14



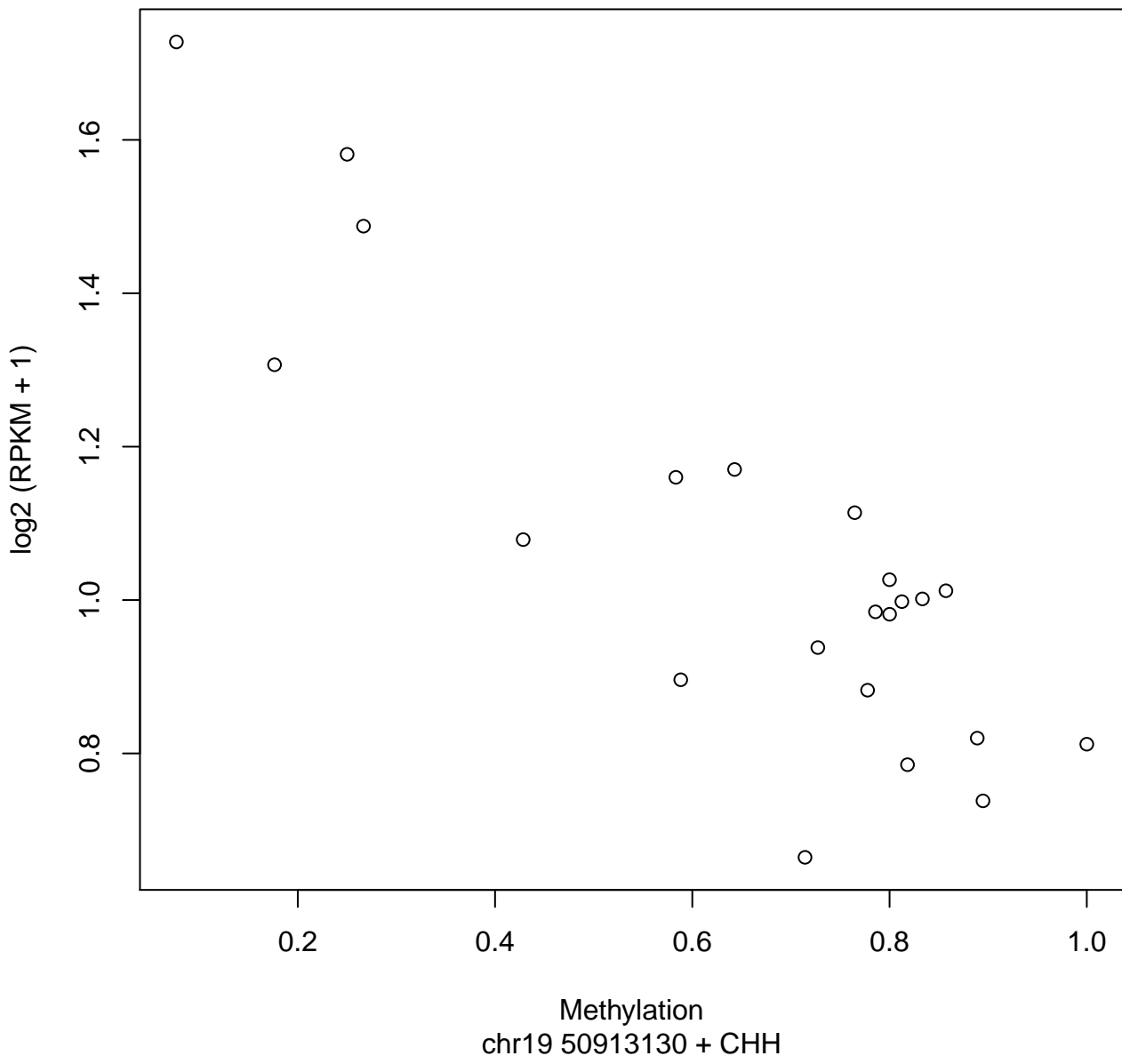
gene ENSG00000164330.16_1 FDR 0.00656

EBF1



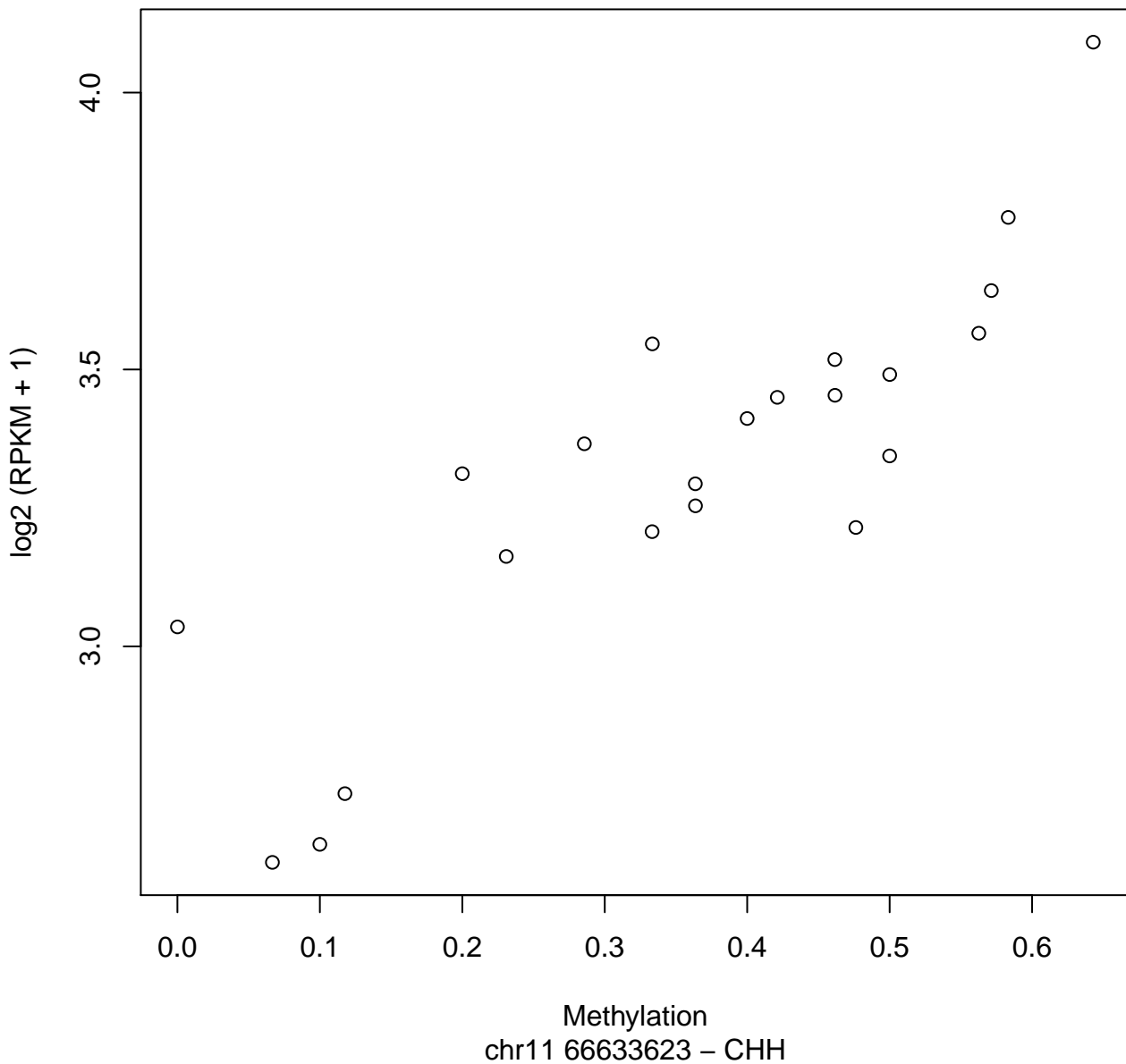
gene ENSG00000062822.12_2 FDR 0.00186

POLD1



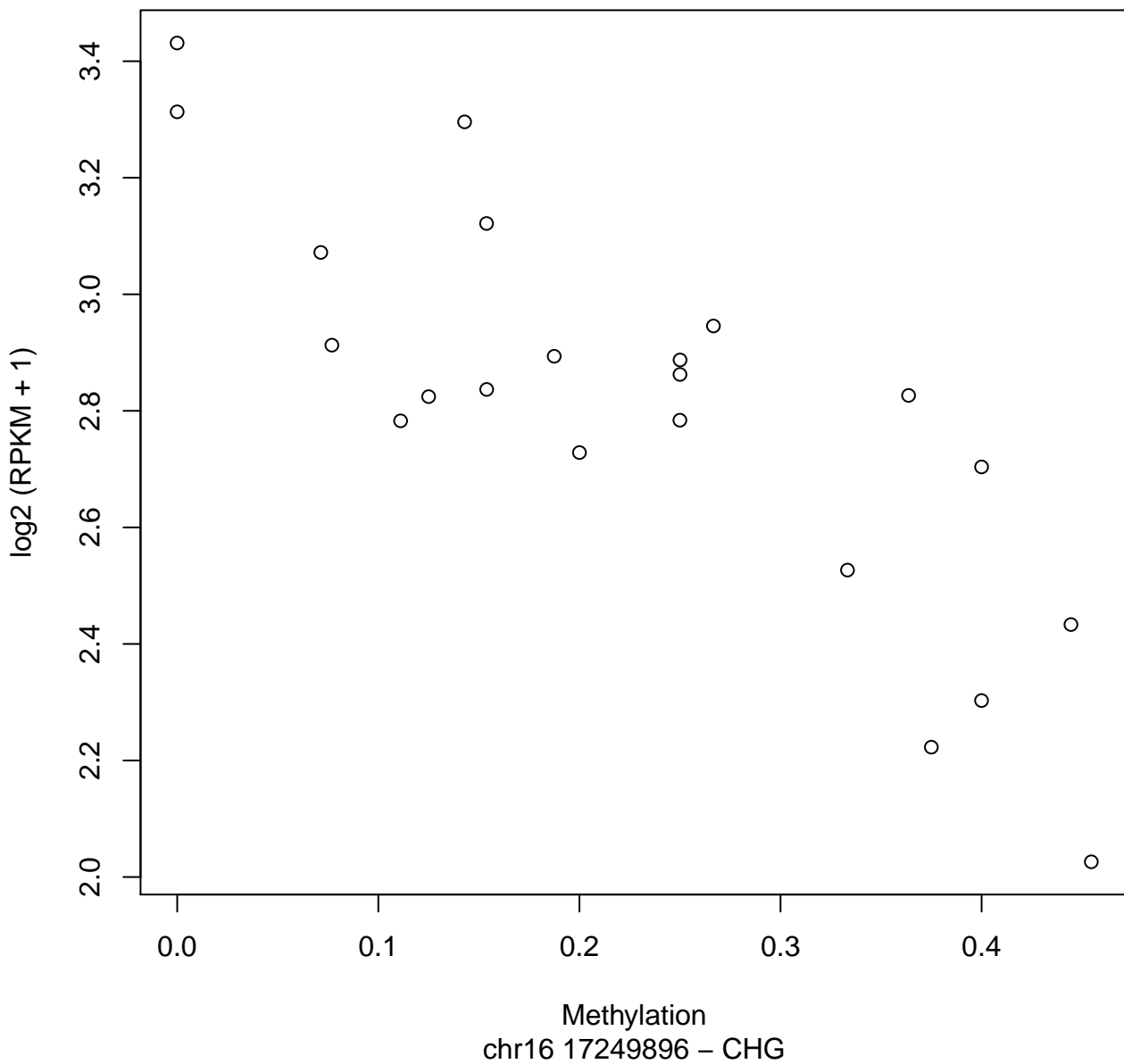
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PC



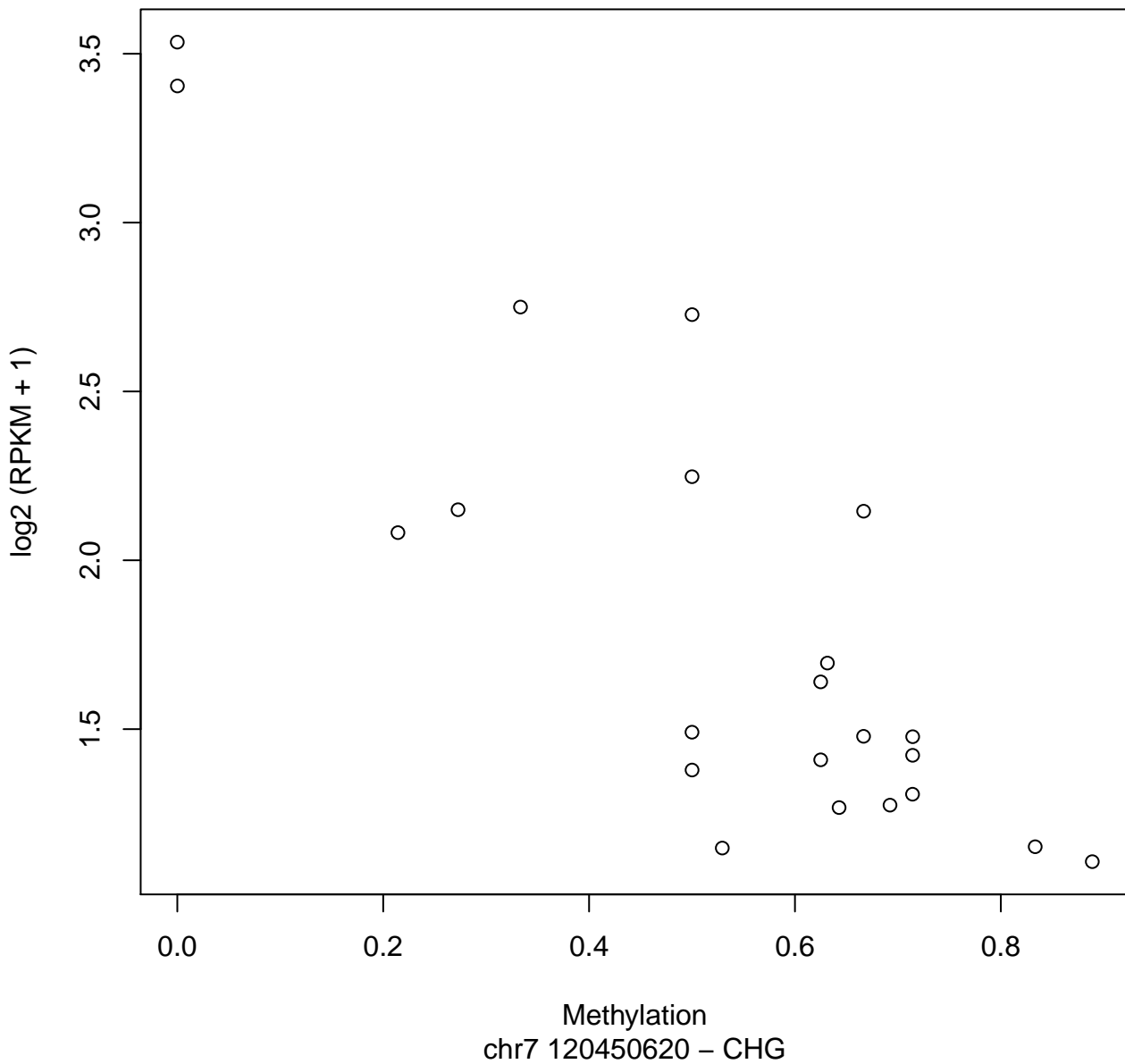
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XYLT1



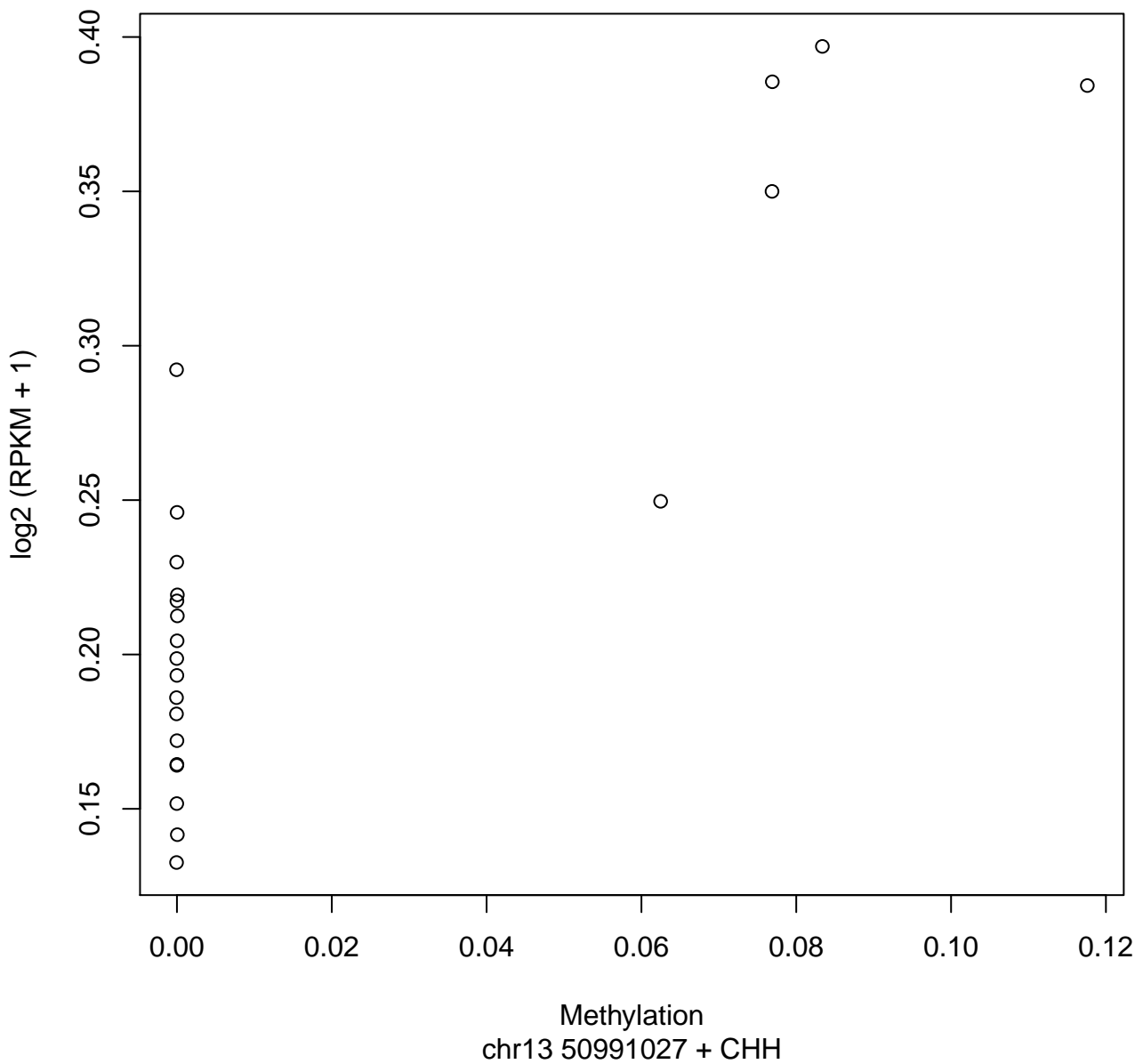
gene ENSG00000106025.8_1 FDR 0.00328

TSPAN12



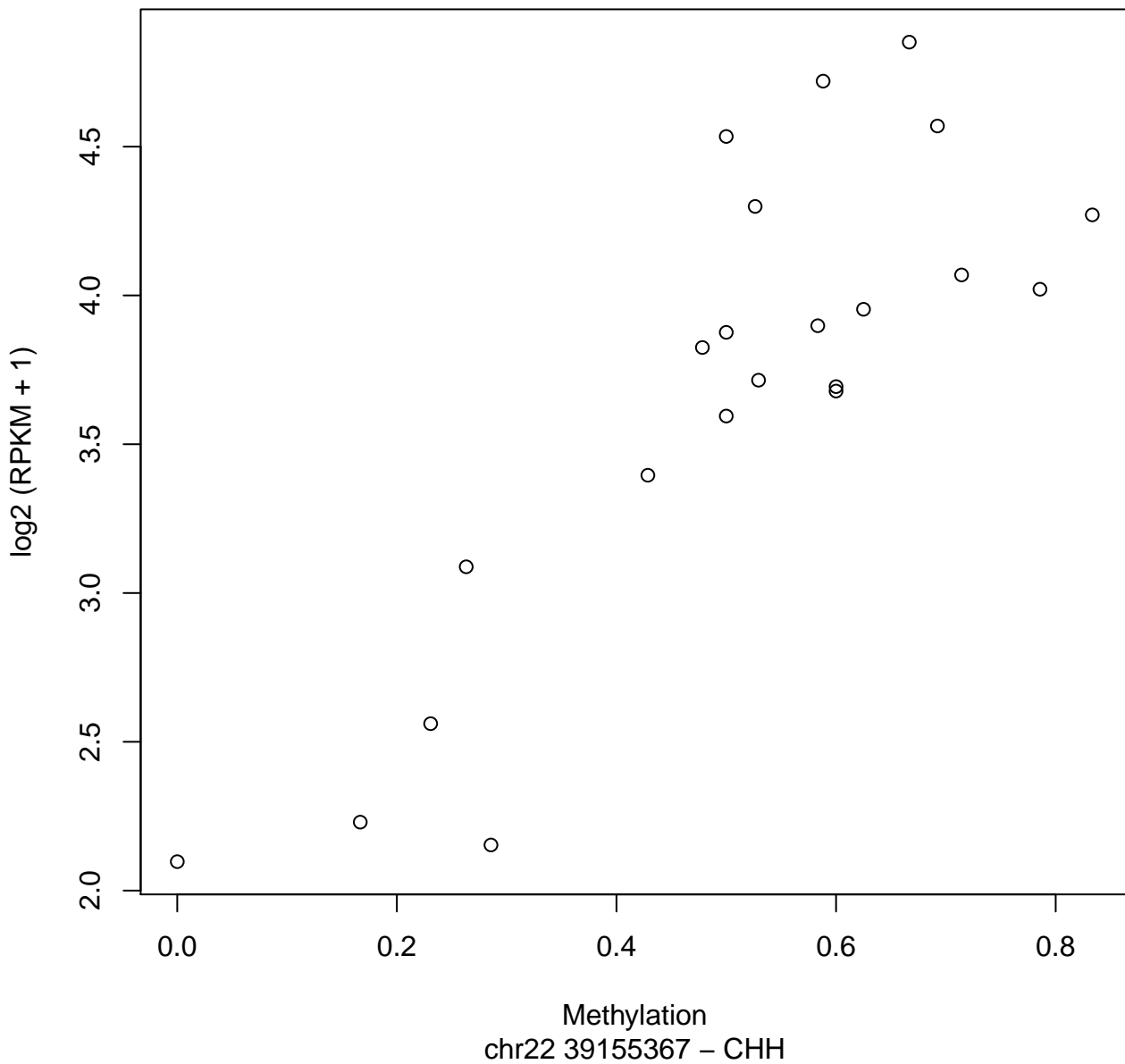
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DLEU1



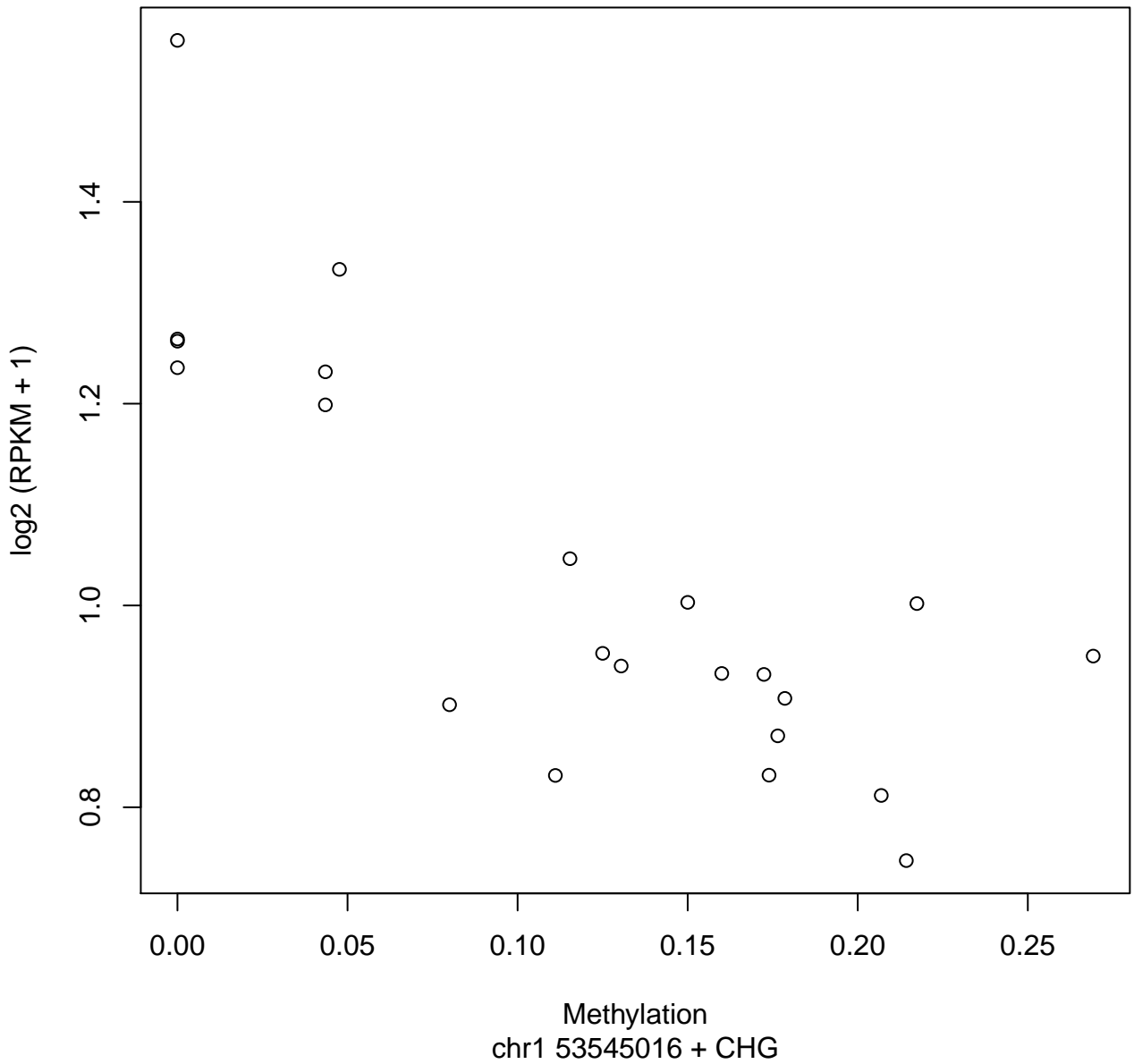
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SUN2



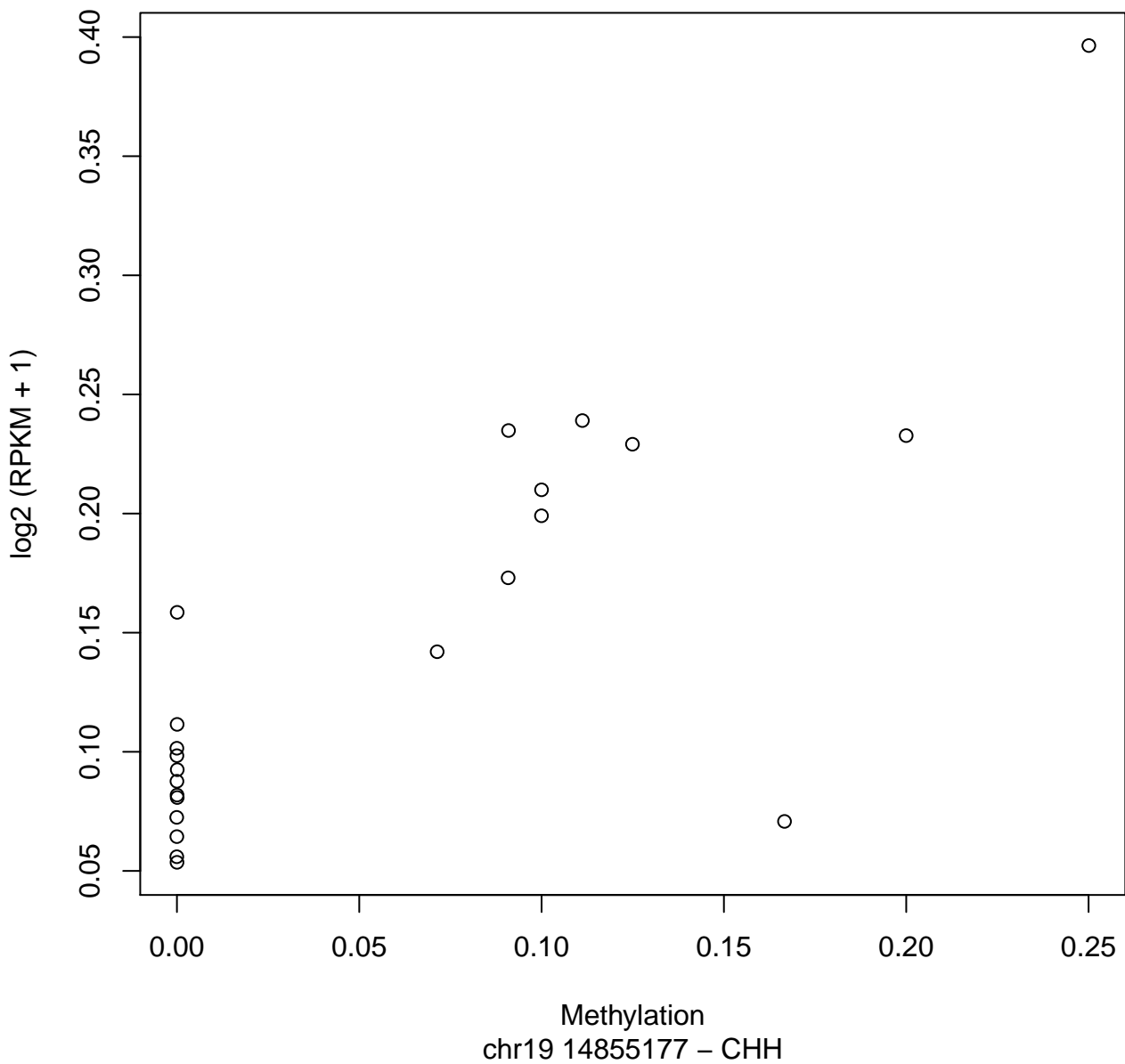
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PODN



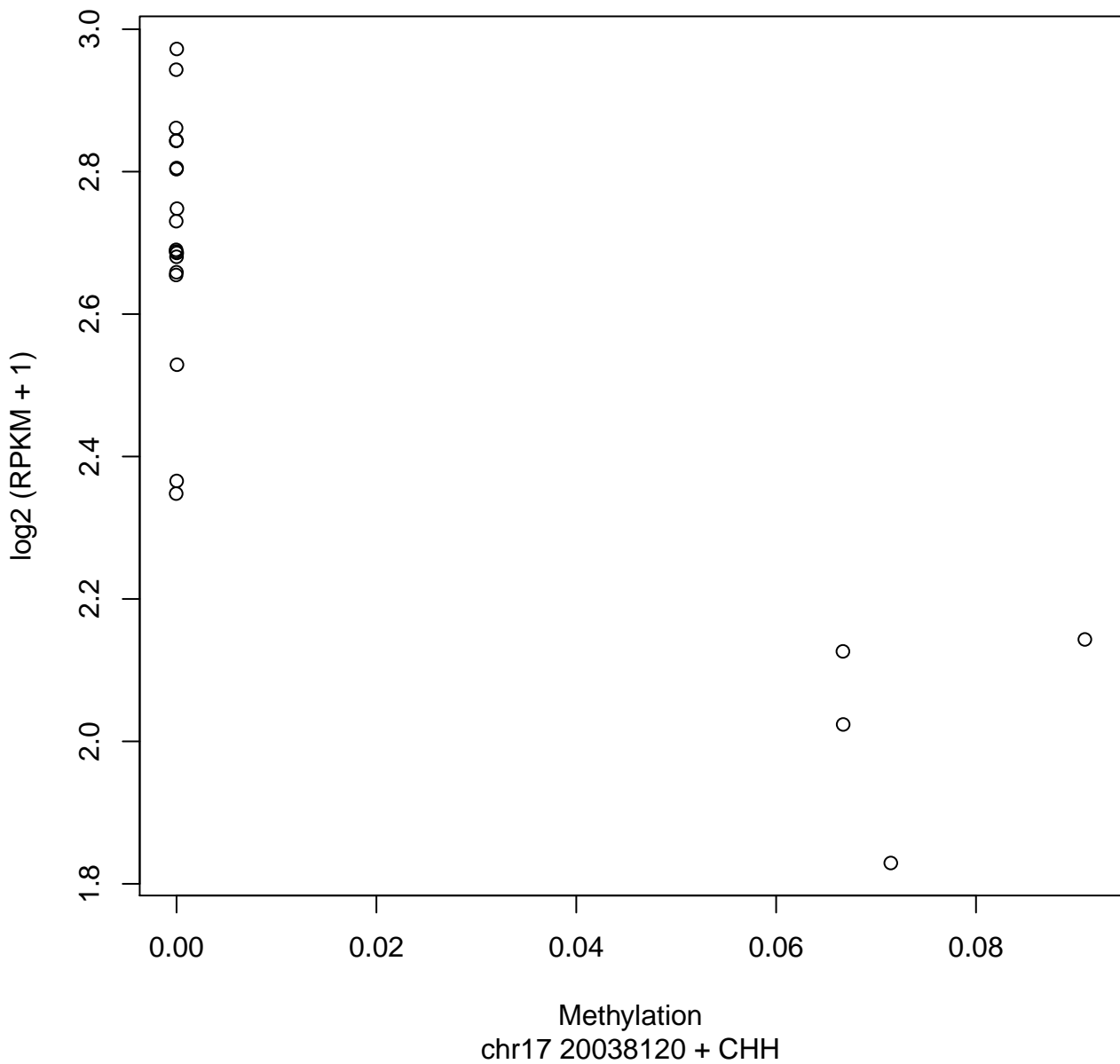
gene ENSG00000127507.17_2 FDR 0.0107

ADGRE2

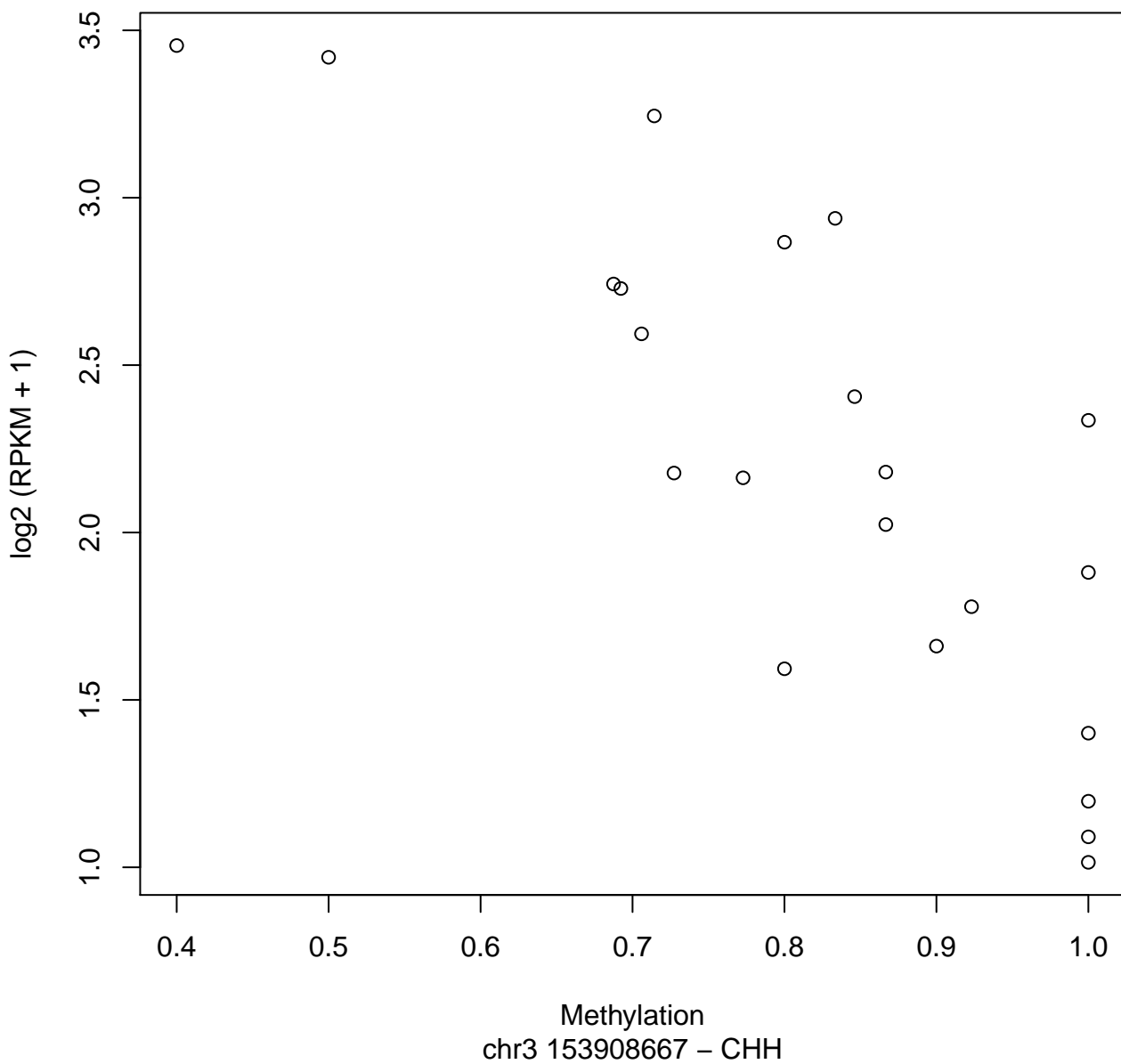


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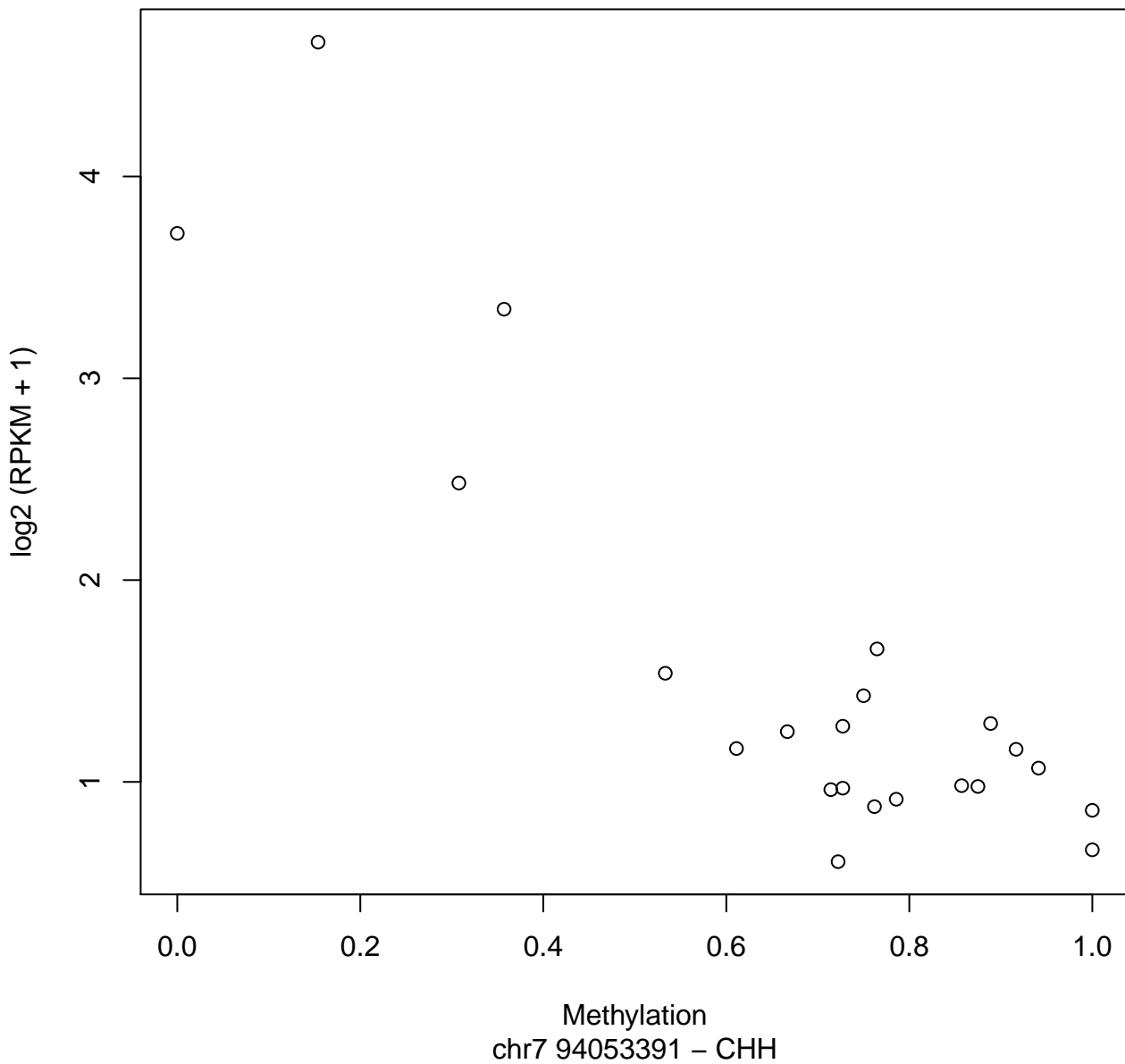
SPECC1



gene ENSG00000114790.12_1 FDR 0.00749
ARHGEF26

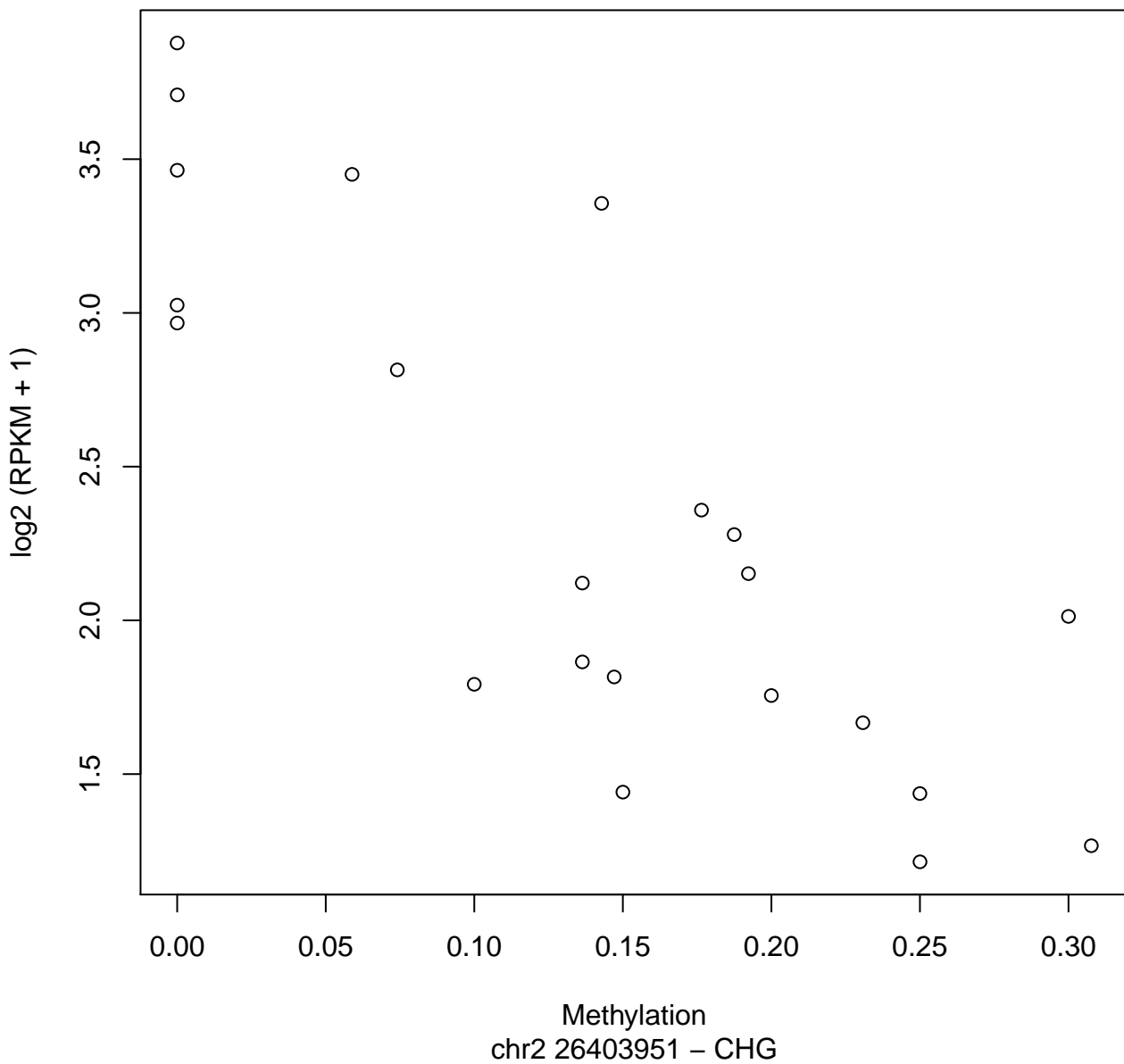


gene ENSG00000164692.17_1 FDR 0.000501
COL1A2

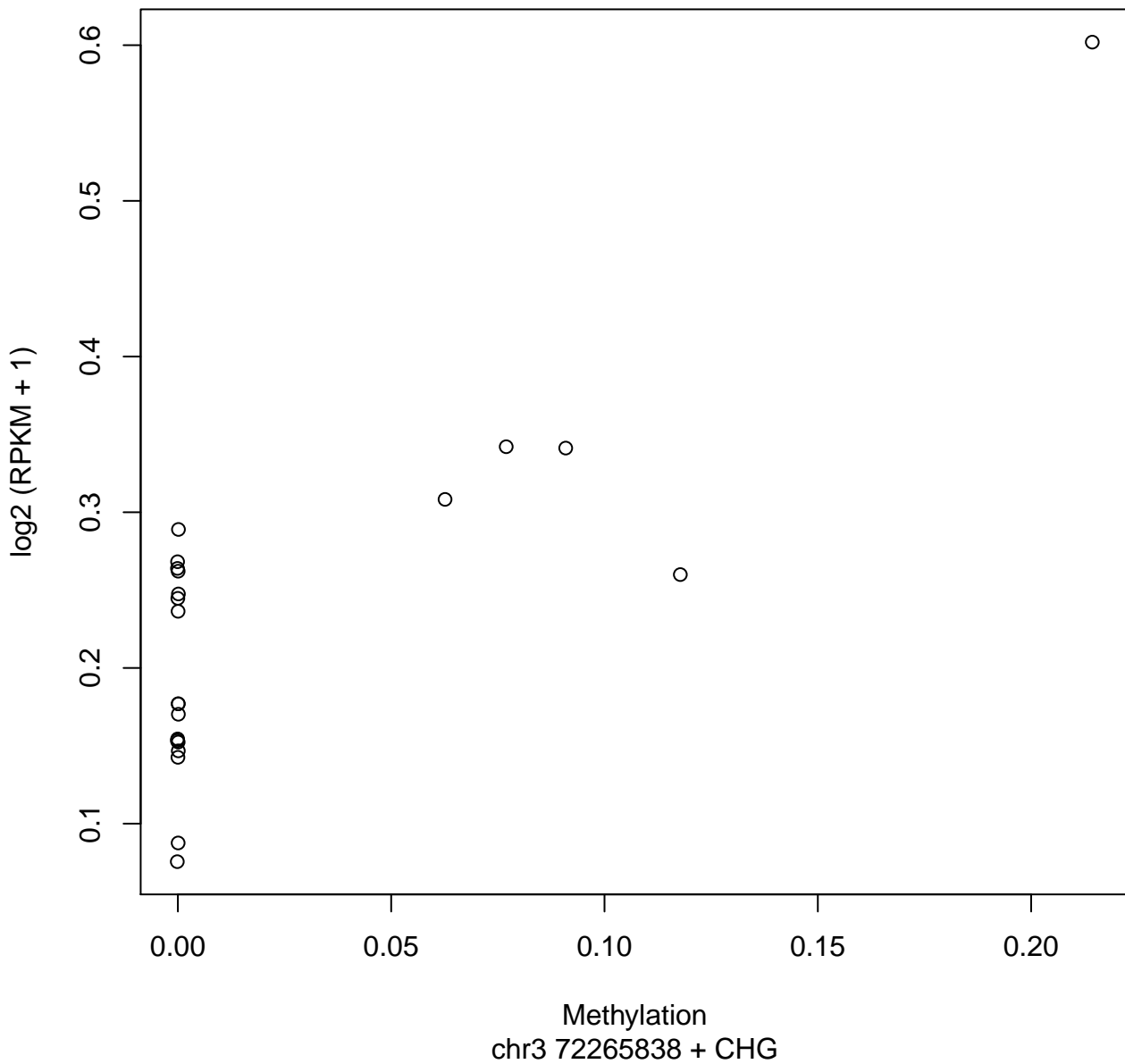


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GAREM2

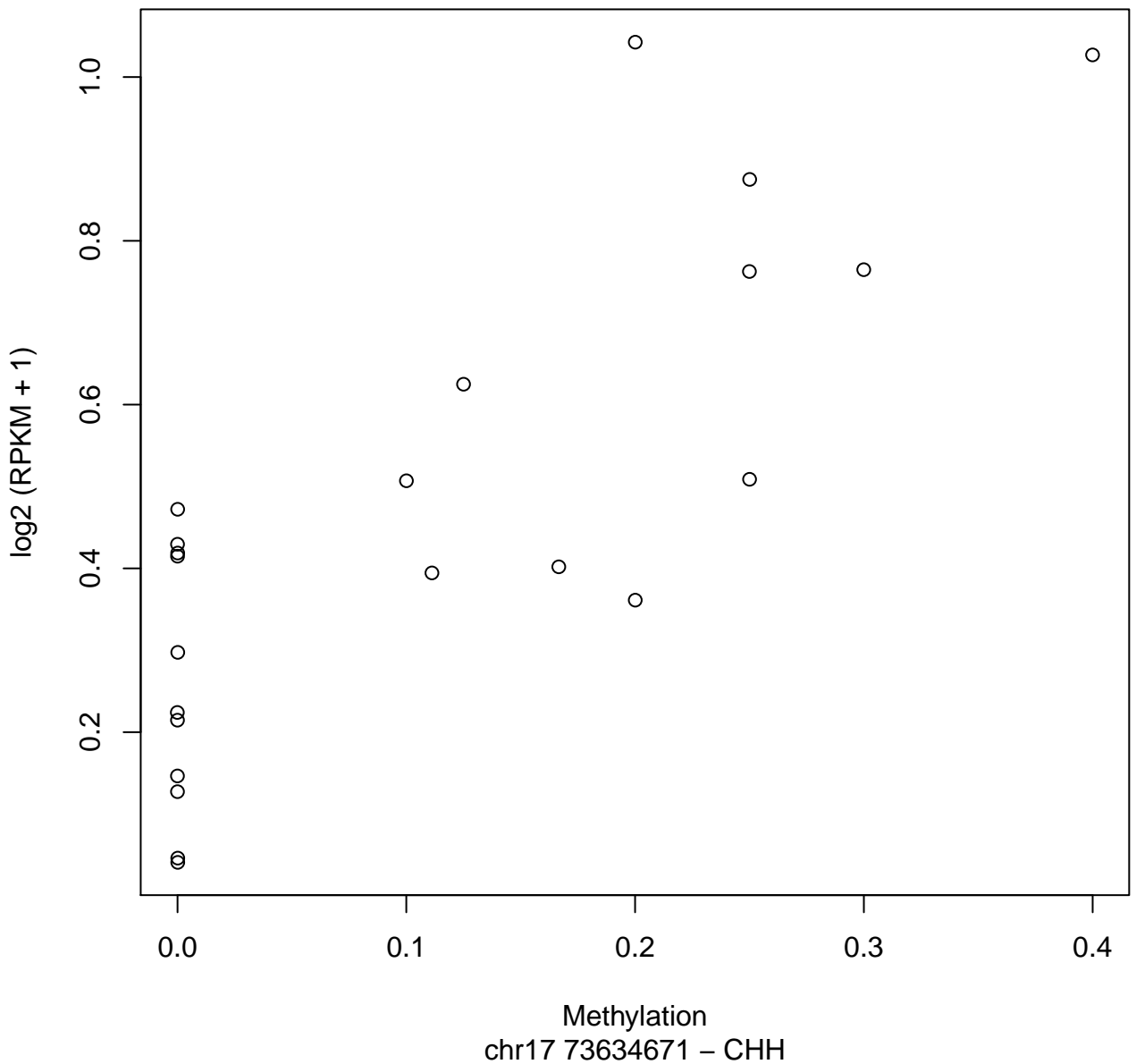


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LINC00877

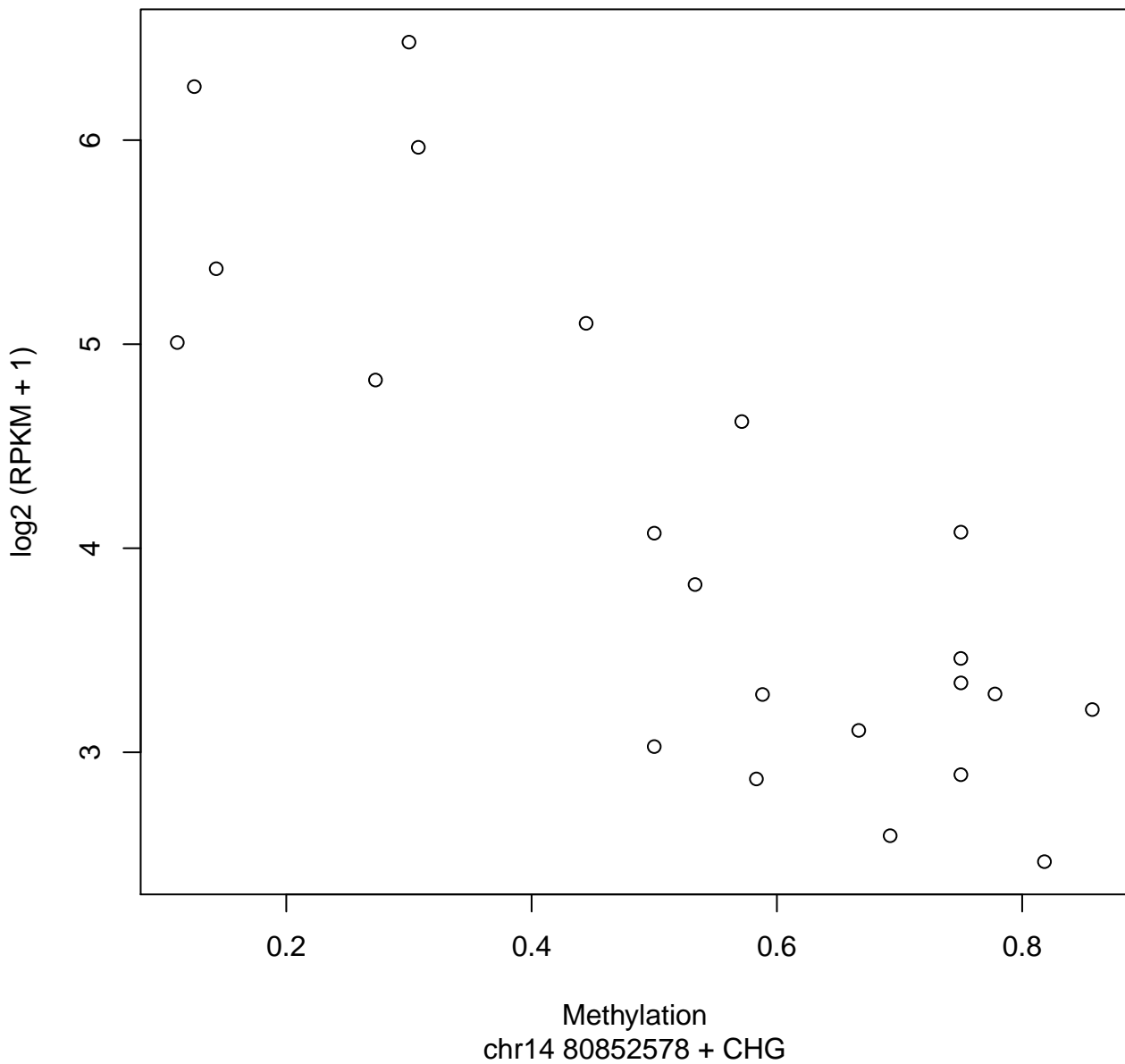


gene ENSG00000204323.5_1 FDR 0.0117

SMIM5



gene ENSG00000211448.11_2 FDR 0.007
DIO2



gene ENSG00000267432.5_2 FDR 0.00707

DNAH17-AS1

