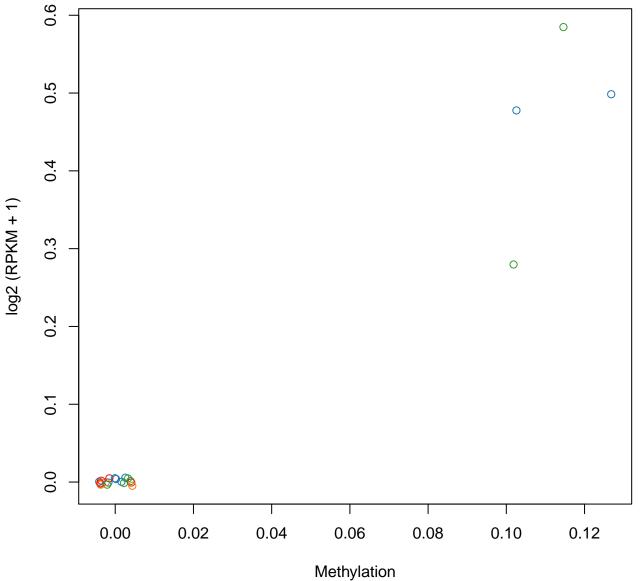
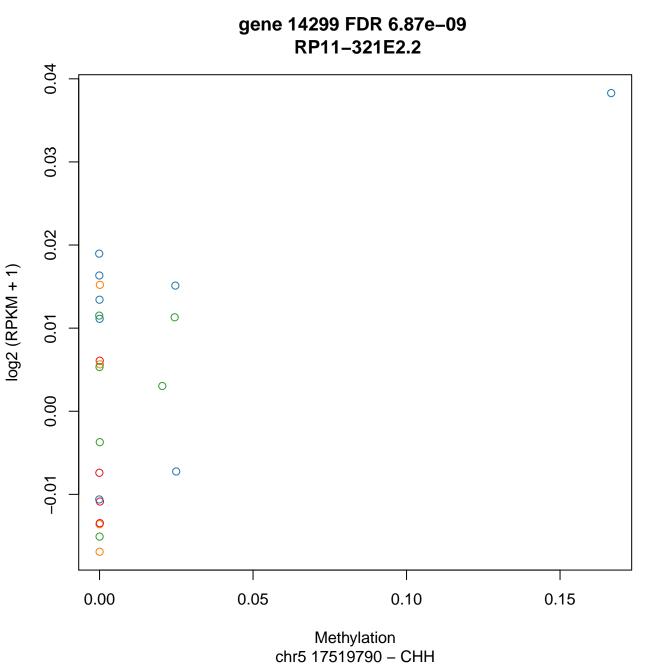
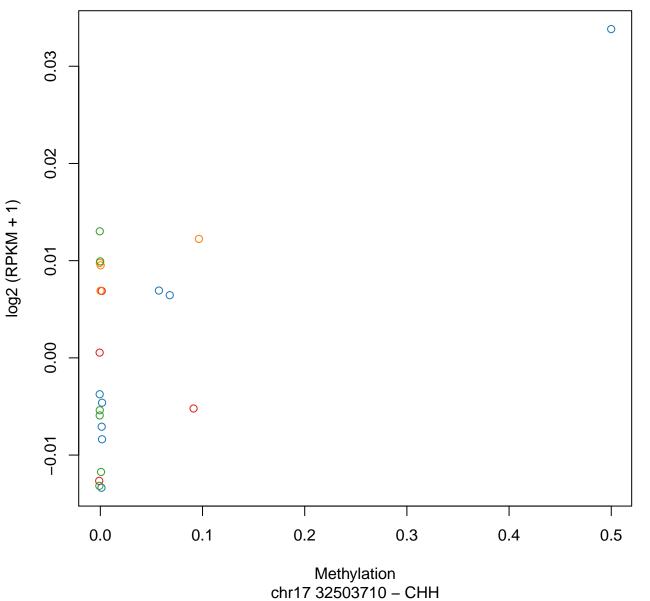
gene 17145 FDR 4.25e-10 AC092634.2



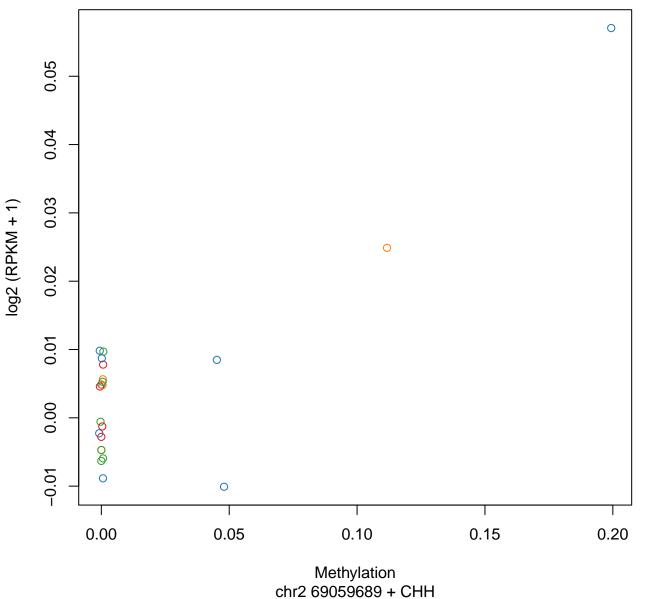
chr7 63387393 - CHG



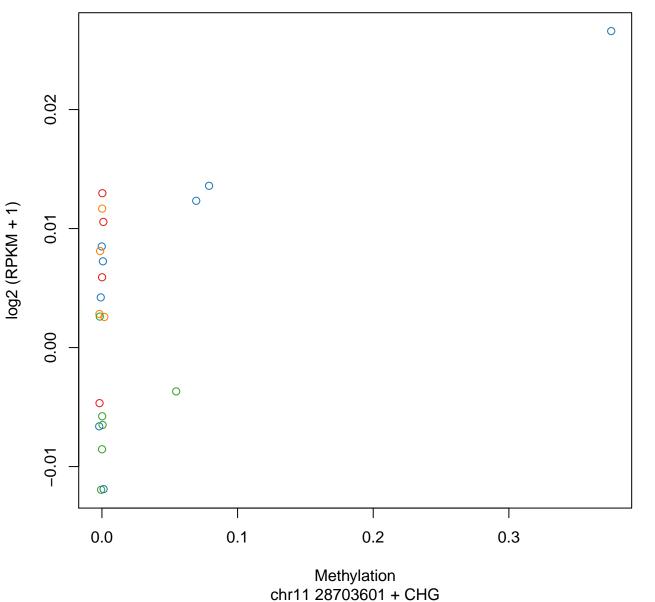
gene 17142 FDR 5.8e-08 RP11-215E13.2



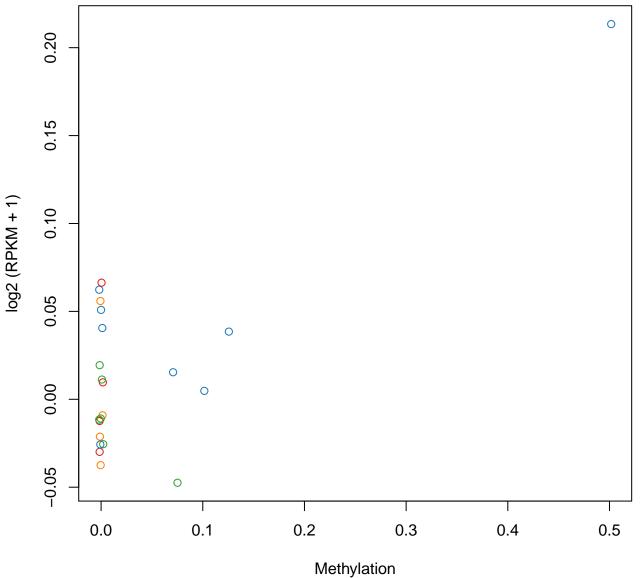
gene 11623 FDR 6.88e-08 AC097495.3



gene 14326 FDR 8.06e-08 RP11-960D24.1

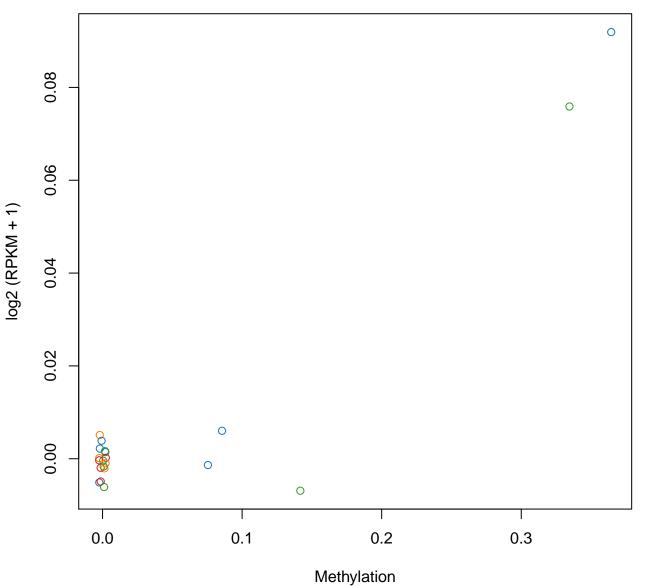


gene 14973 FDR 1.66e-06 RNU6-537P



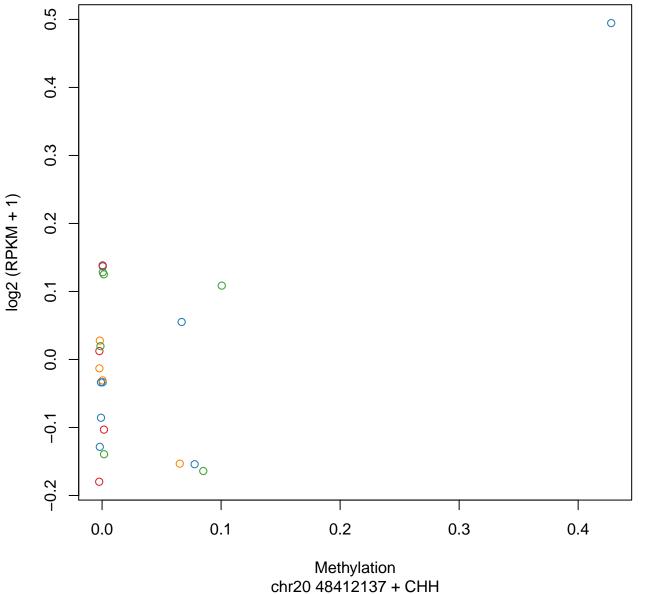
chr1 11212006 - CHG

gene 18791 FDR 3.19e-06 RP3-335E1.1

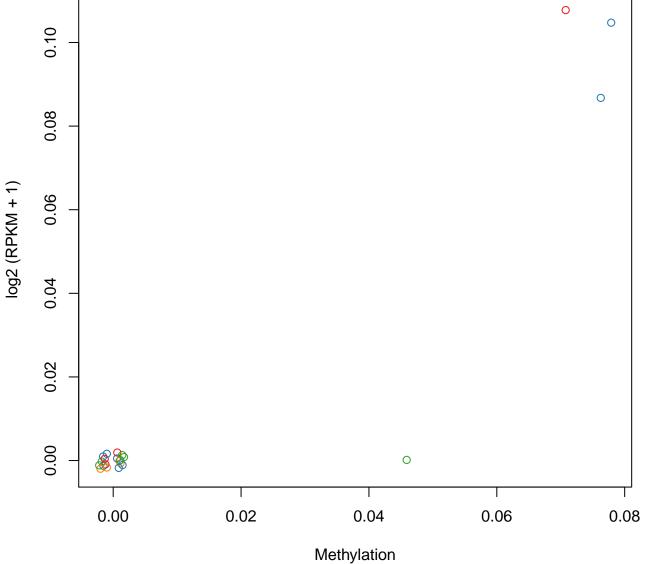


chr6 106905416 - CHH

gene 14917 FDR 4.4e-06 RNU6-919P

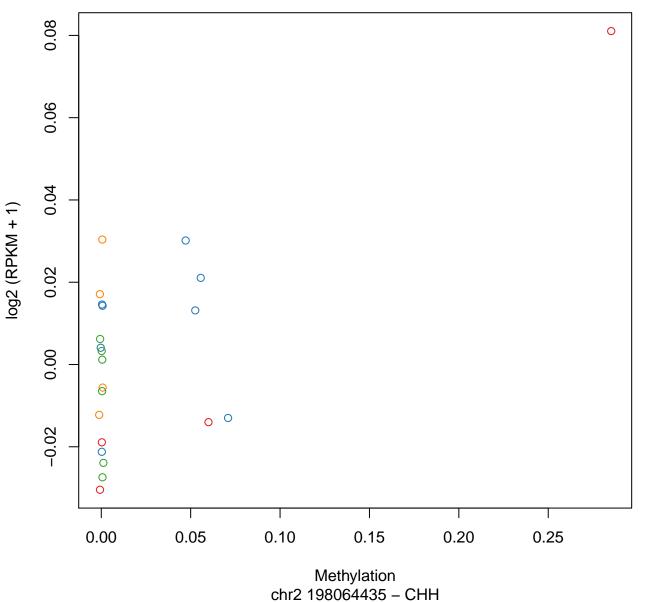


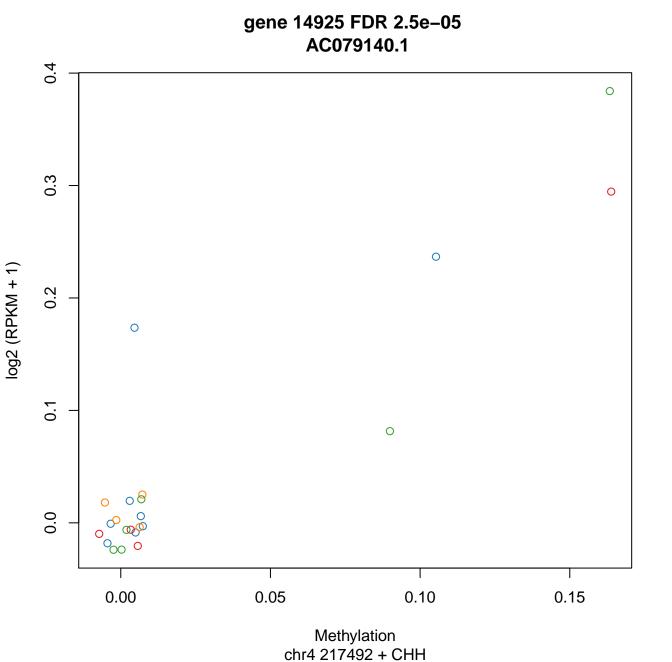




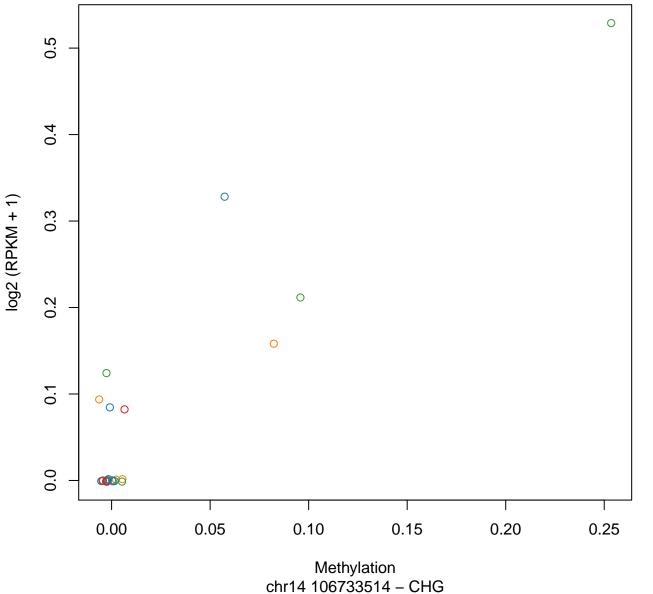
chr5 6847884 - CHH

gene 12203 FDR 1.7e-05 AC013264.2

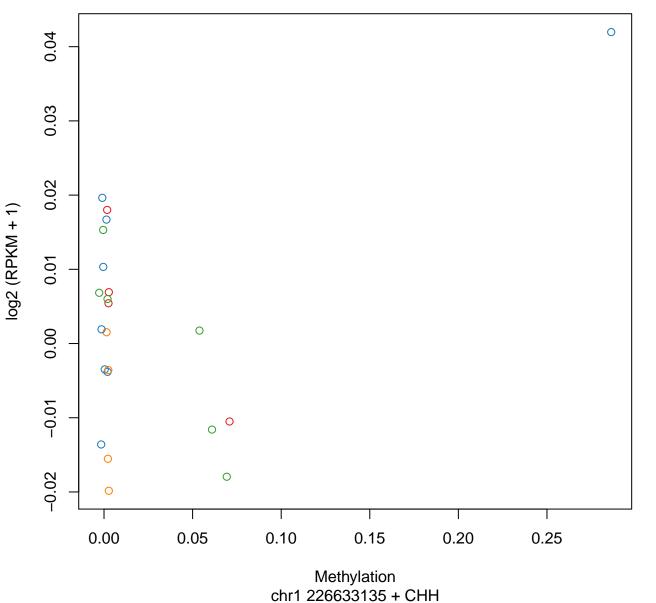




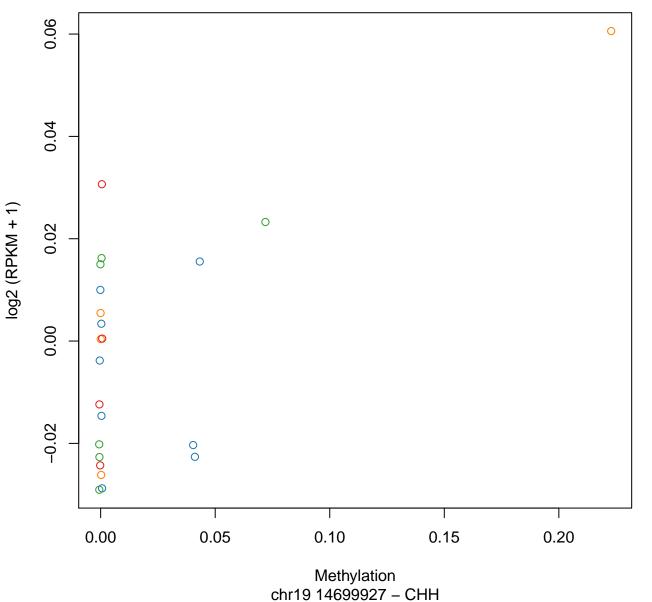
gene 10098 FDR 3.1e-05 IGHV1-24



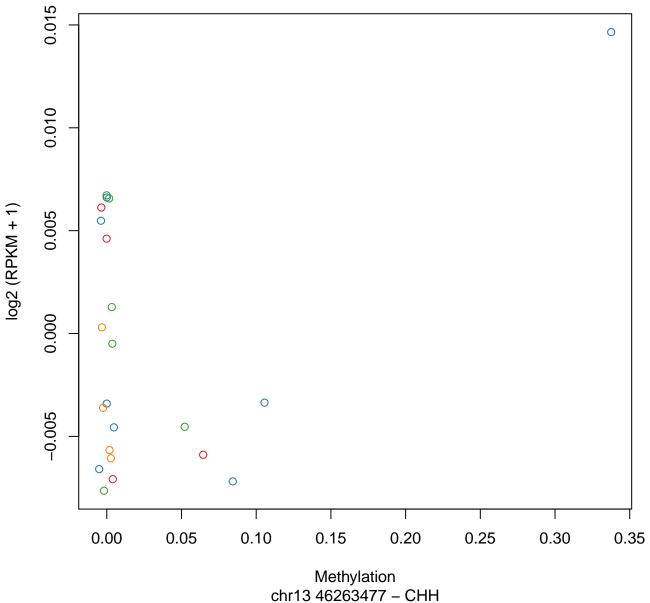
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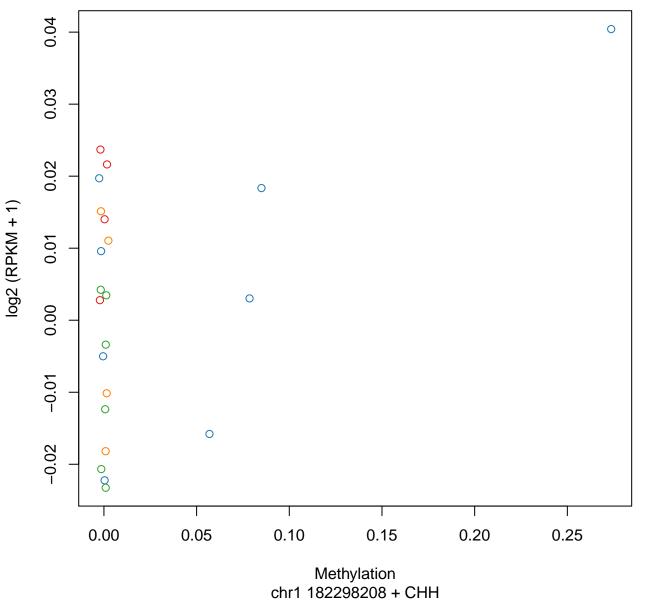
gene 13871 FDR 3.21e-05 RN7SL842P



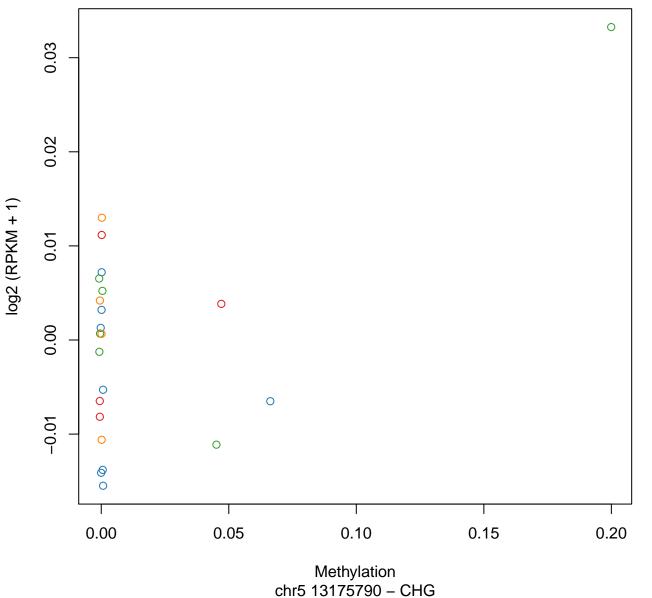
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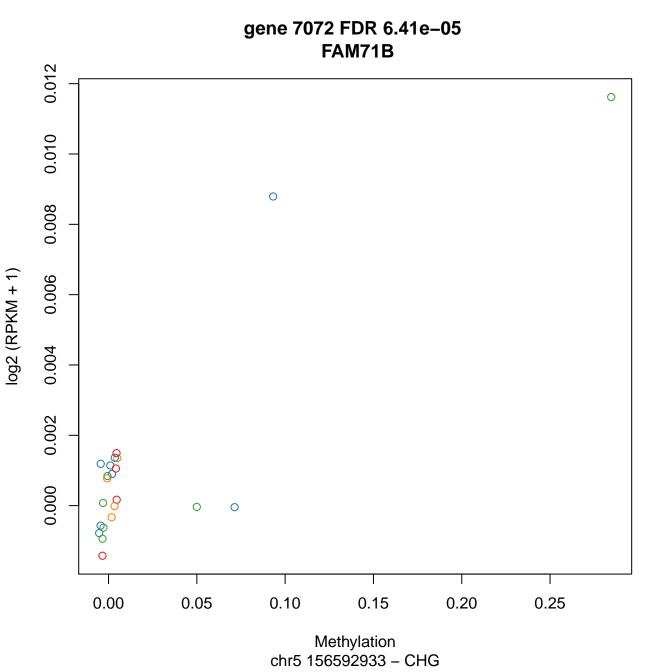


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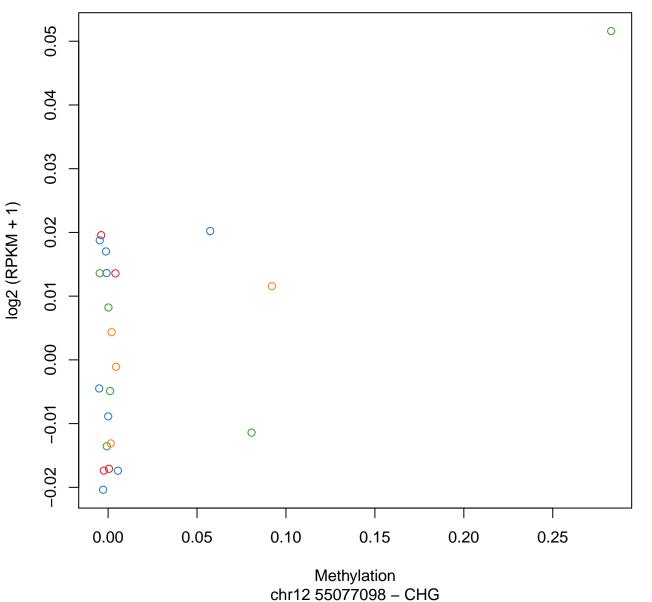


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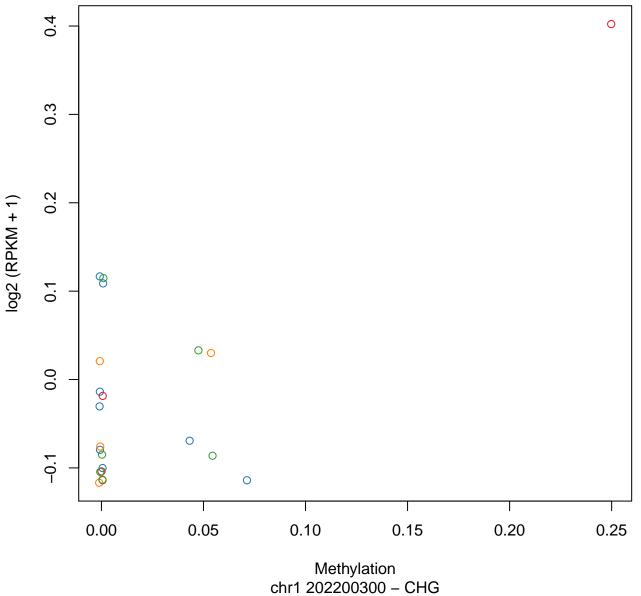




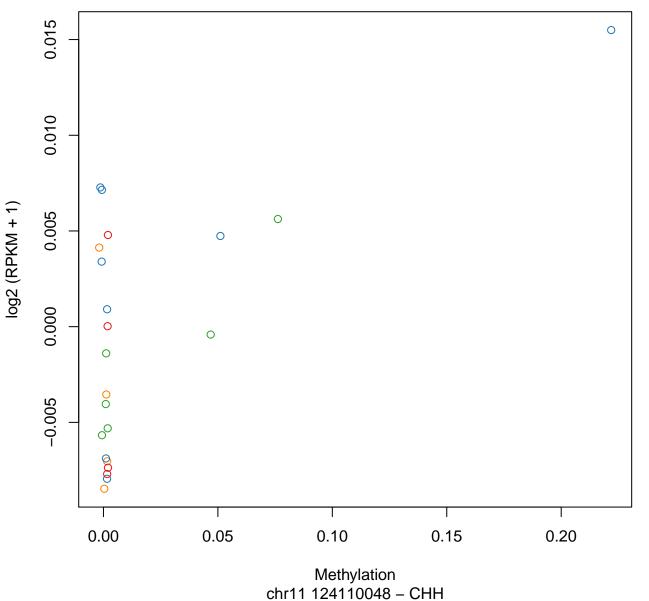
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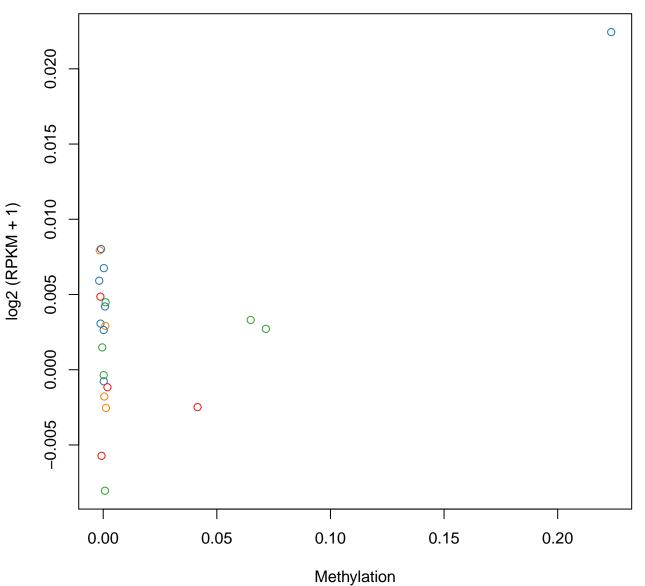
gene 13517 FDR 6.77e-05 snoU13



gene 18316 FDR 6.81e-05 OR8G7P

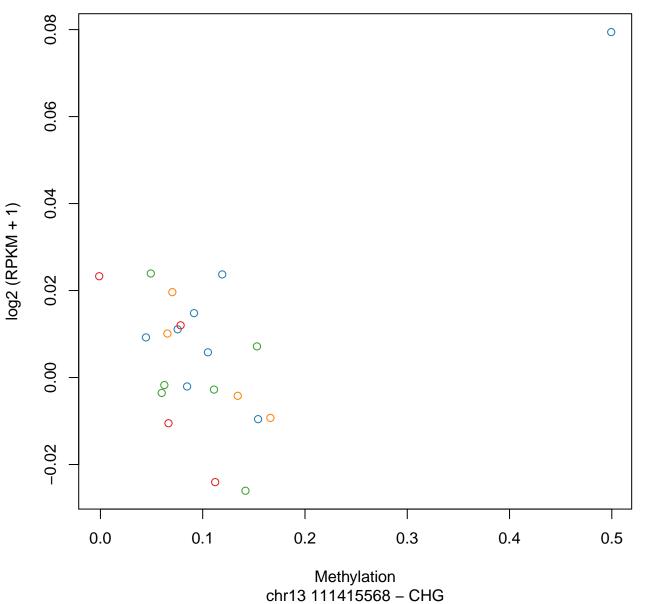


gene 10148 FDR 7.8e-05 KRTAP2-3

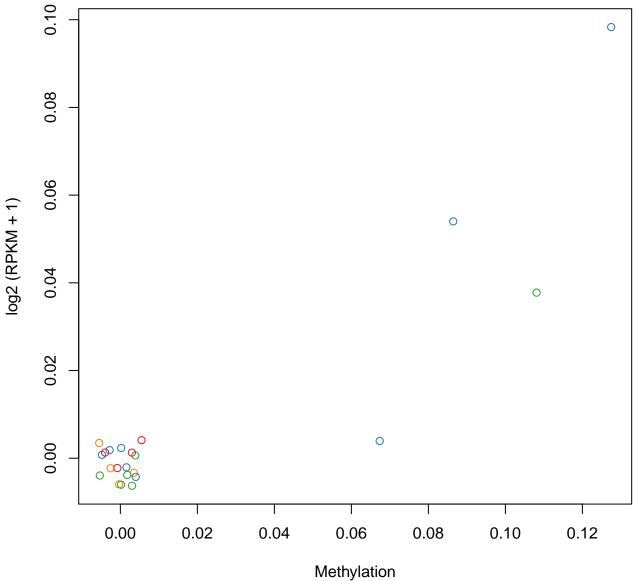


chr17 39215988 + CHG

gene 11093 FDR 8.93e-05 RPL21P107

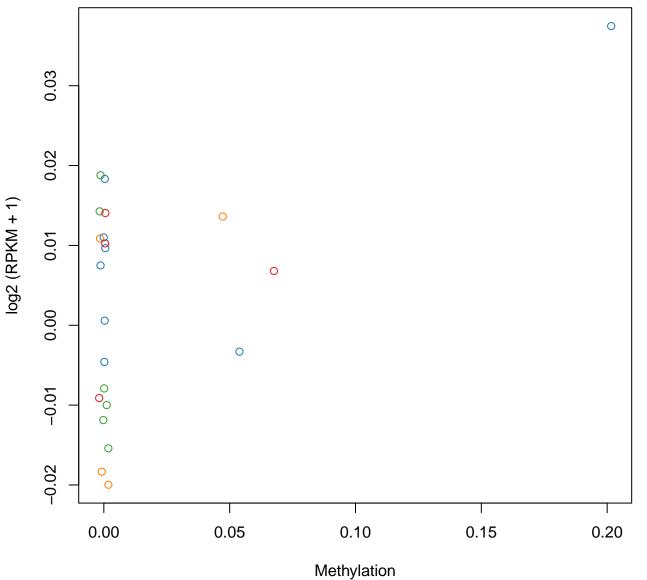


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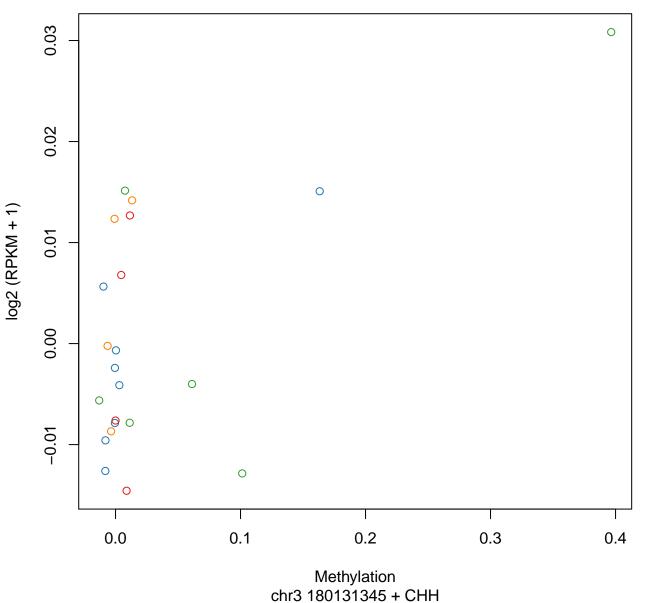
chr1 248662644 - CHH

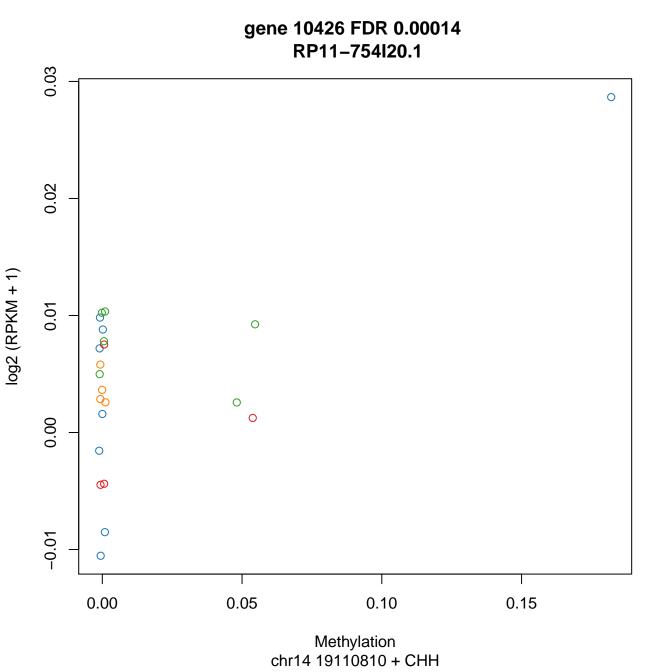
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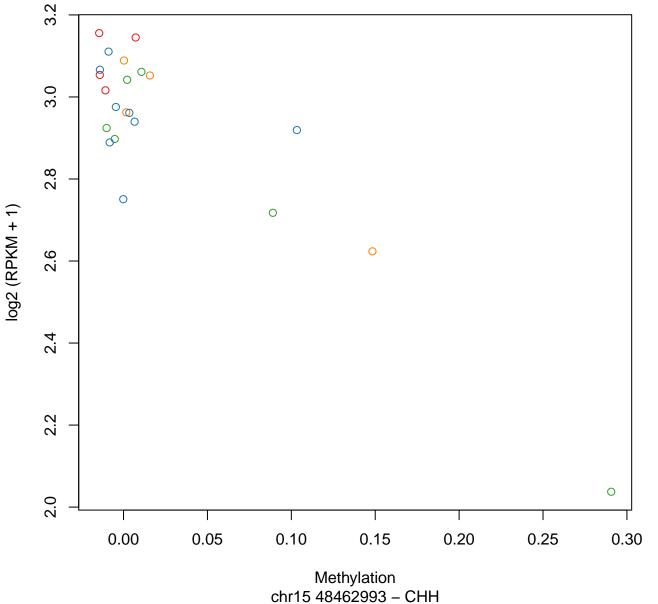
chr20 31805678 - CHH

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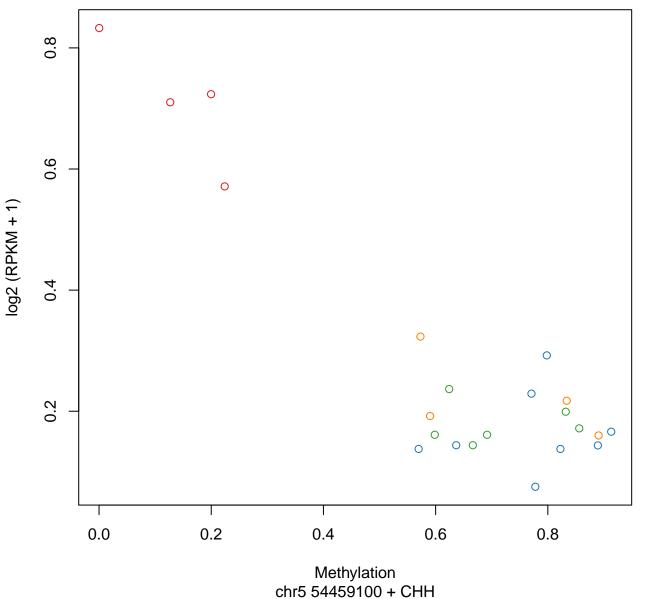




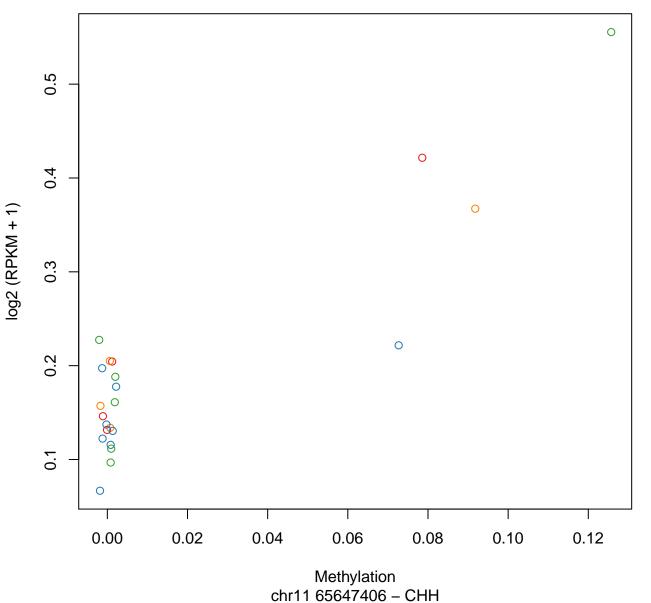
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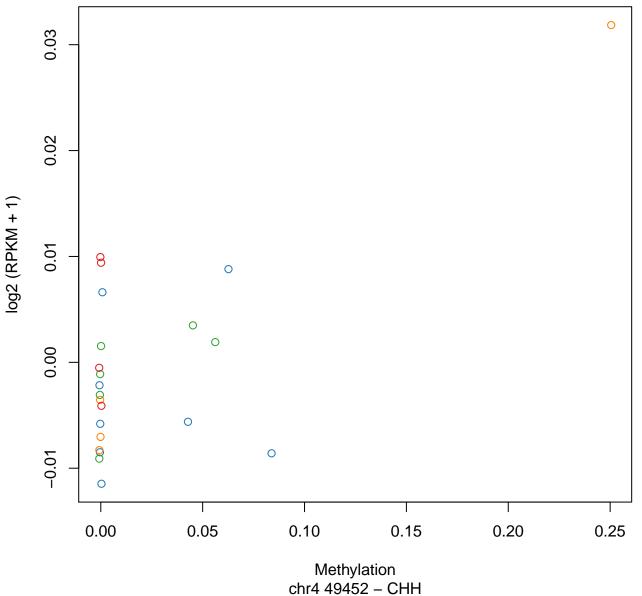
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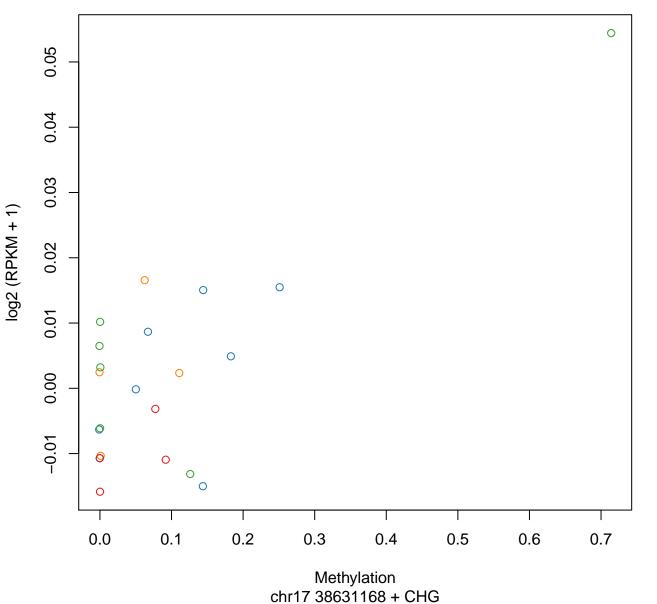
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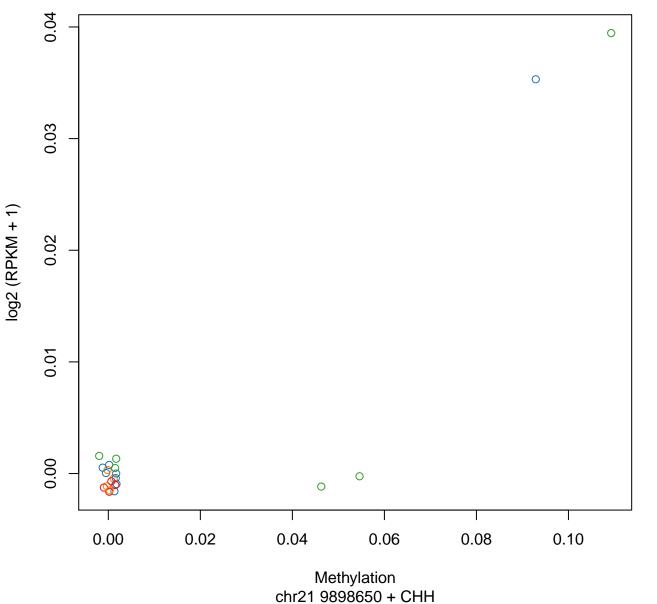
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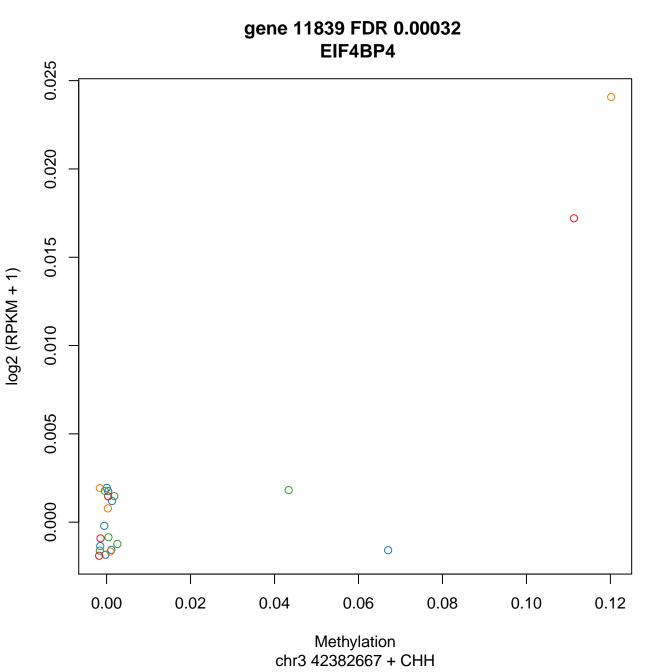


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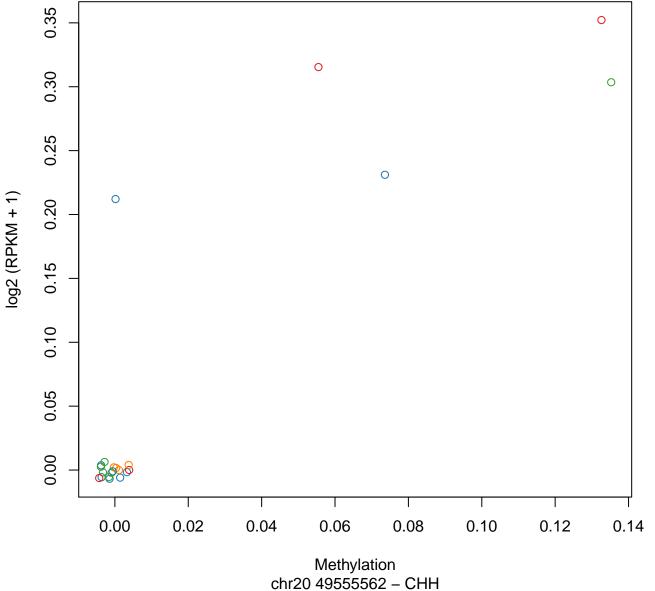


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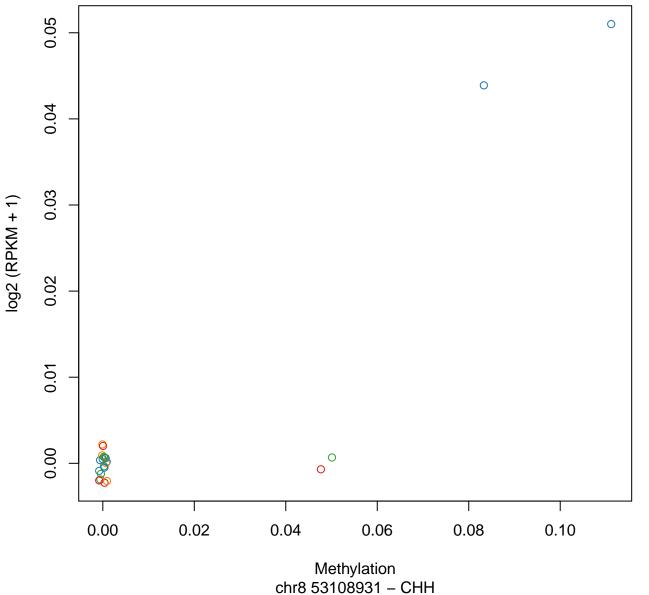




gene 17238 FDR 0.000344 AL034553.1



gene 15087 FDR 0.000399 RP11-26M5.2



gene 10543 FDR 0.000438 **B3GALNT2P1** 0.015 0 0.010 0 00 0 0.005 log2 (RPKM + 1) 0 0 0.000 0 0 0 -0.0050 0 0 0 8 0

0.2

0.0

0.1

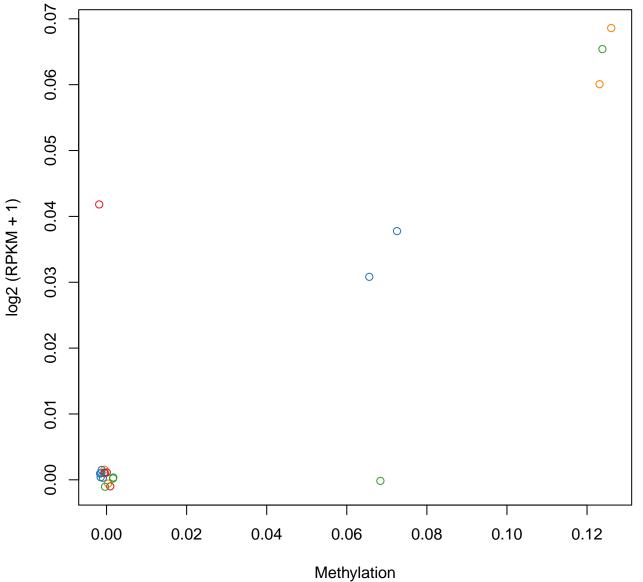
Methylation chr6 130123352 + CHH

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0.4

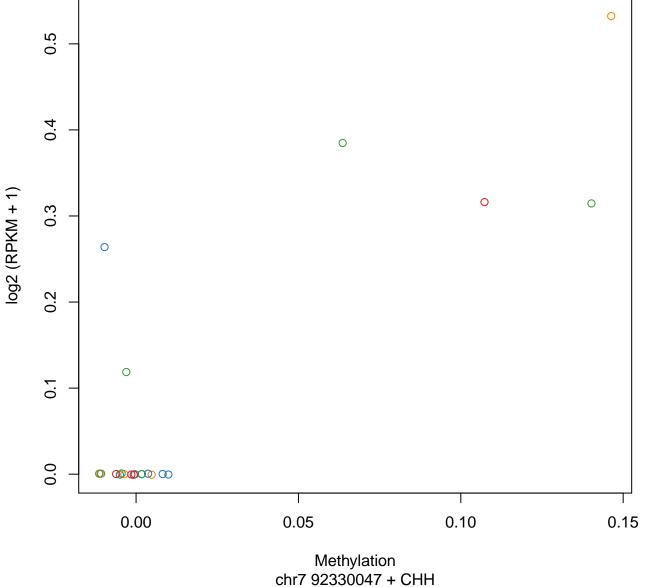
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gene 16256 FDR 0.000484 RP11-62C7.2

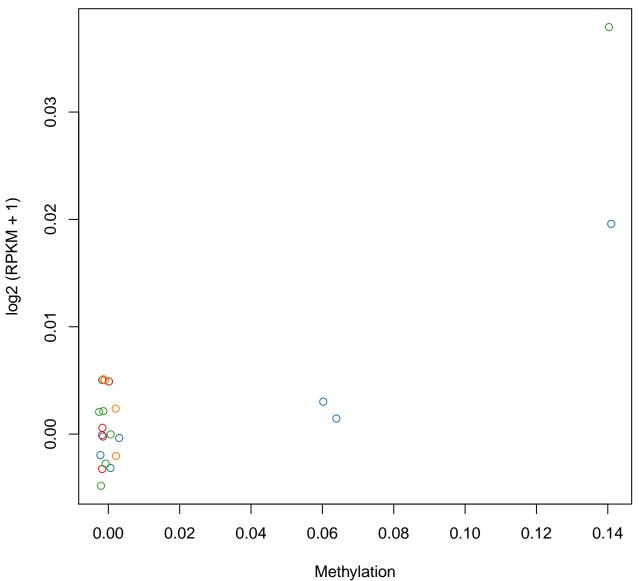


chr15 39311119 - CHH

gene 9933 FDR 0.000498 RNU6-10P

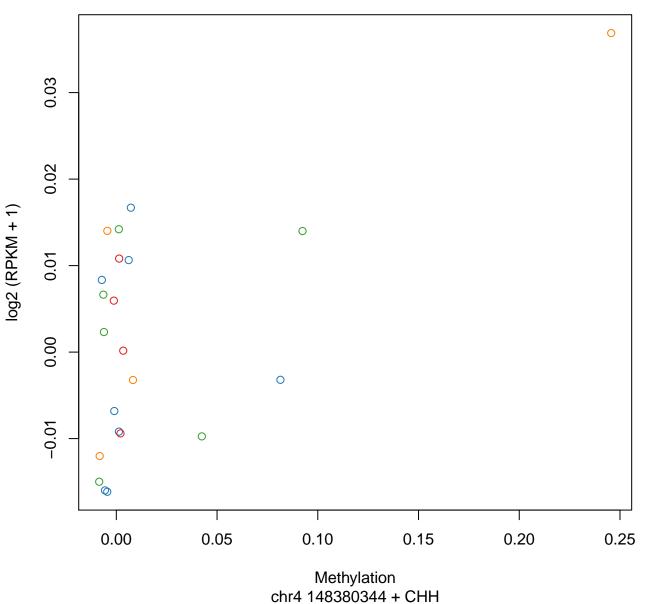


gene 13100 FDR 0.000509 RP5-1050E16.1

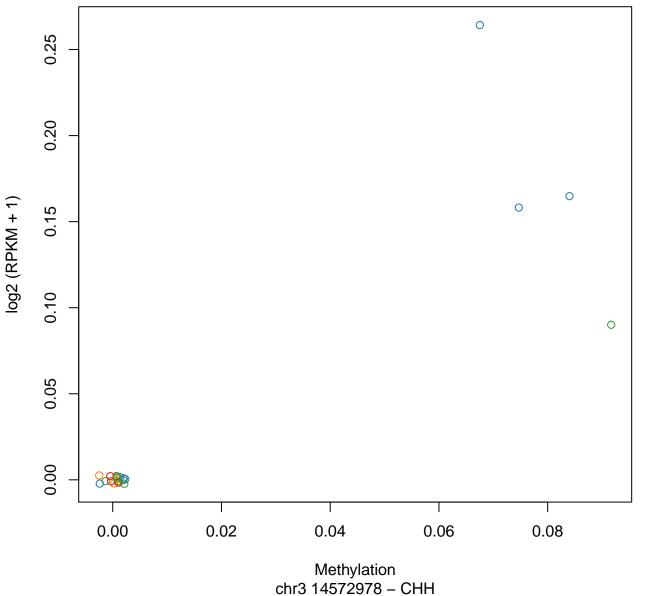


chr9 92503052 - CHH

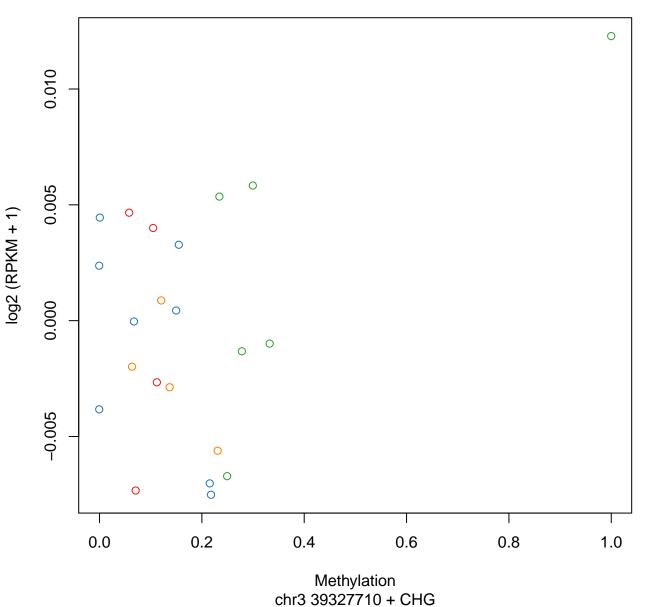
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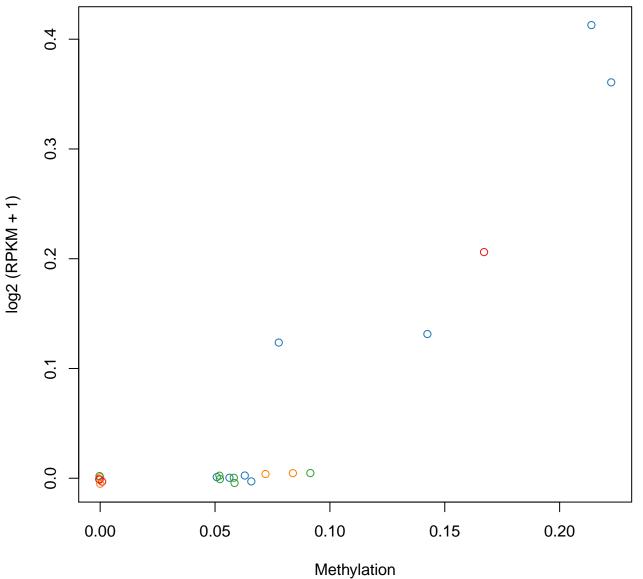
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gene 17823 FDR 0.000565 RP11-331G2.6

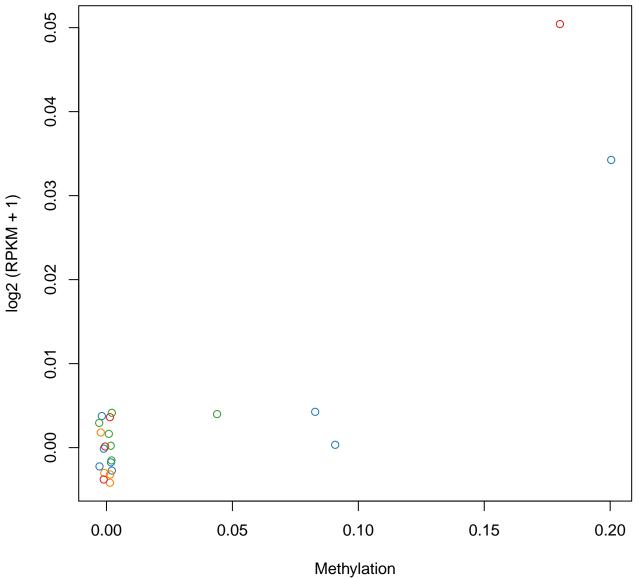


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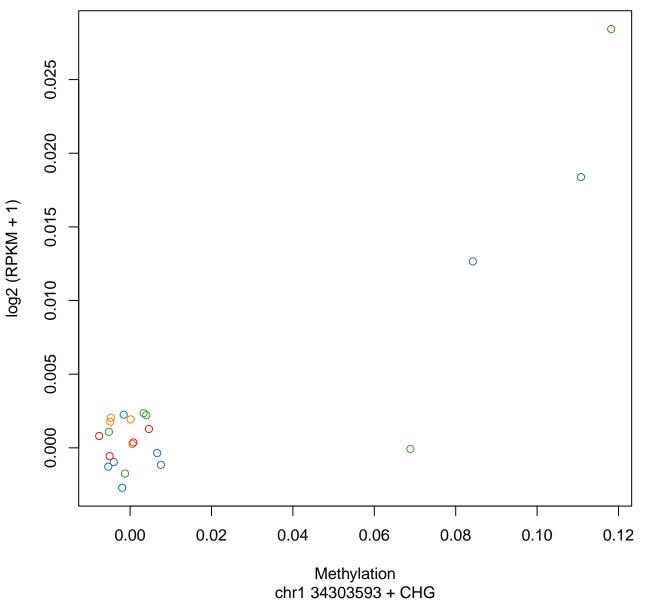
chr11 36410712 - CHH

gene 10213 FDR 0.000584 AC091654.7

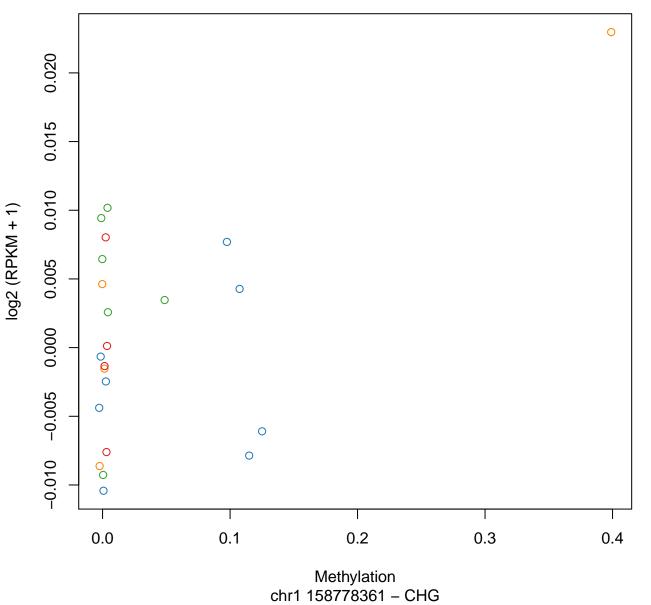


chr7 98083606 - CHH

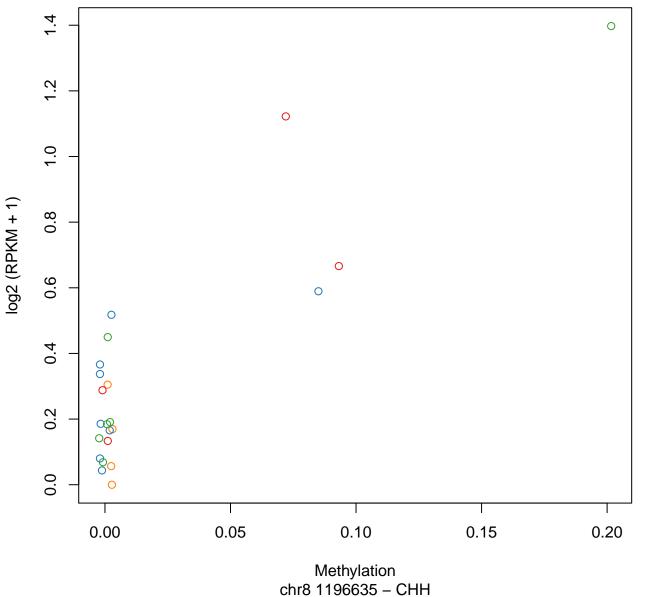
gene 11392 FDR 0.000653 HSPD1P14



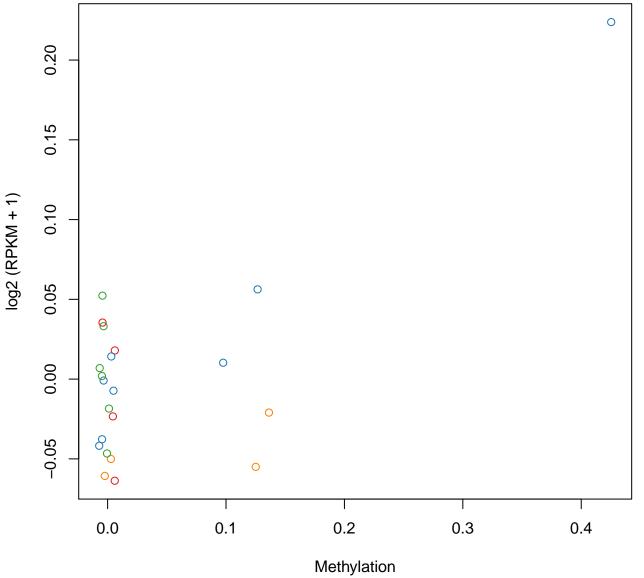
gene 7986 FDR 0.000668 OR10AA1P



gene 15250 FDR 0.000716 CTD-2281E23.3

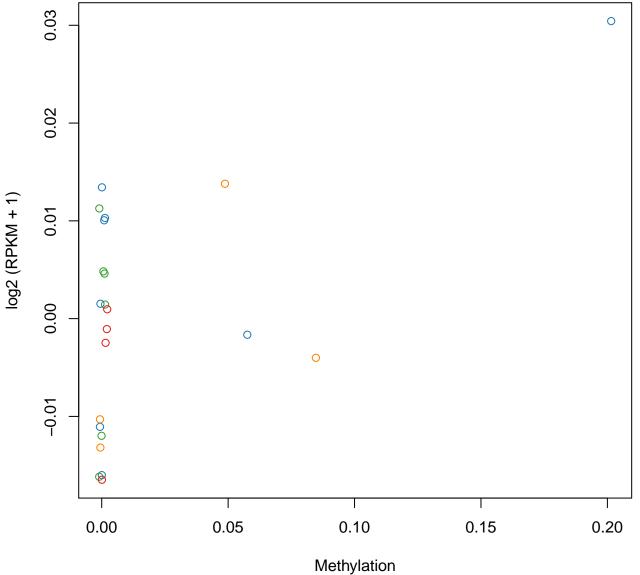


gene 9448 FDR 0.000749 RNU6-992P



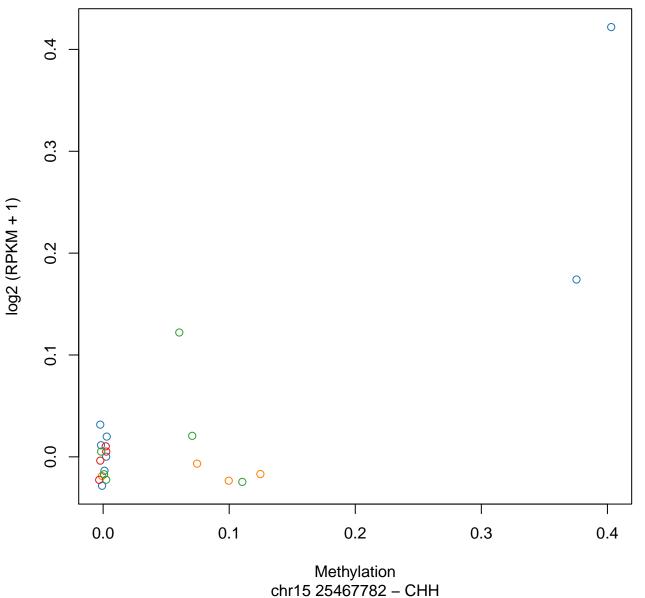
chr21 37437941 - CHH

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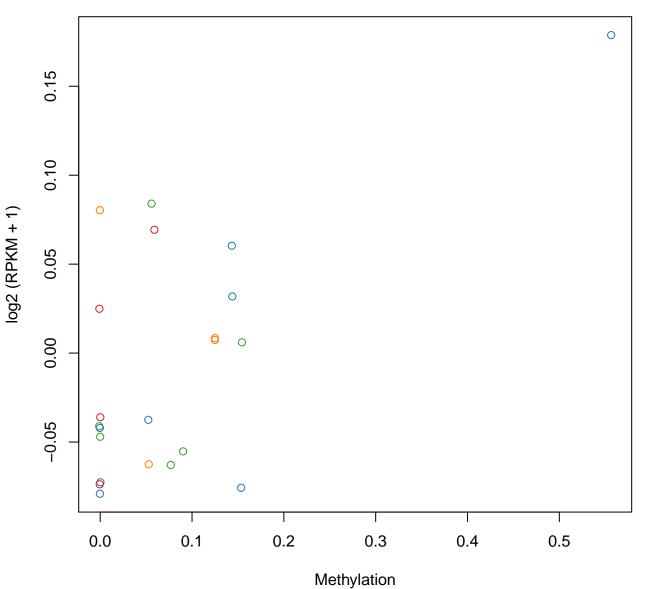


chr5 2738423 - CHG

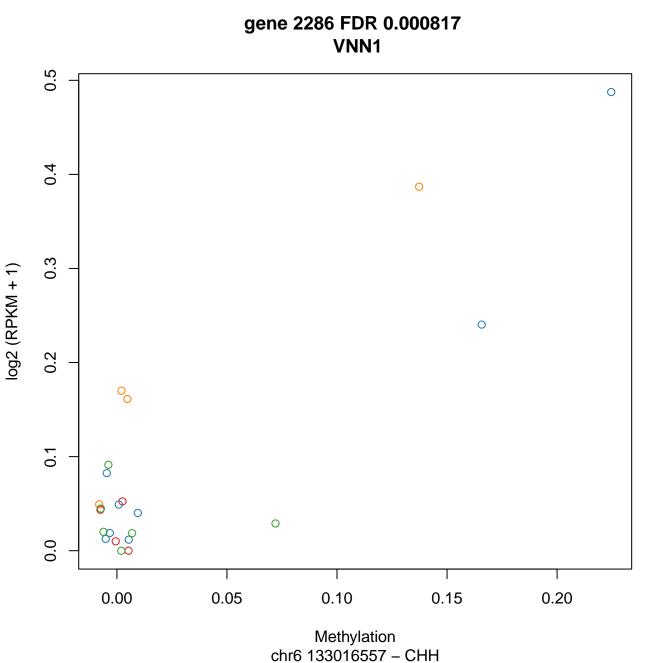
gene 9420 FDR 0.000756 SNORD115-29



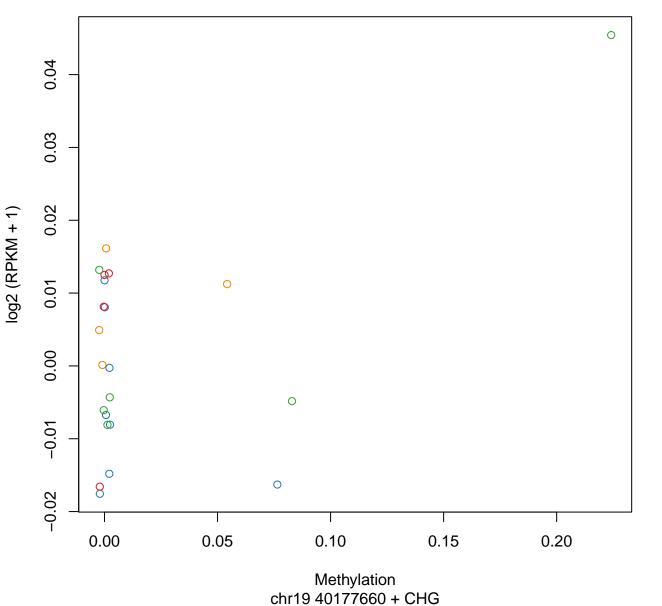
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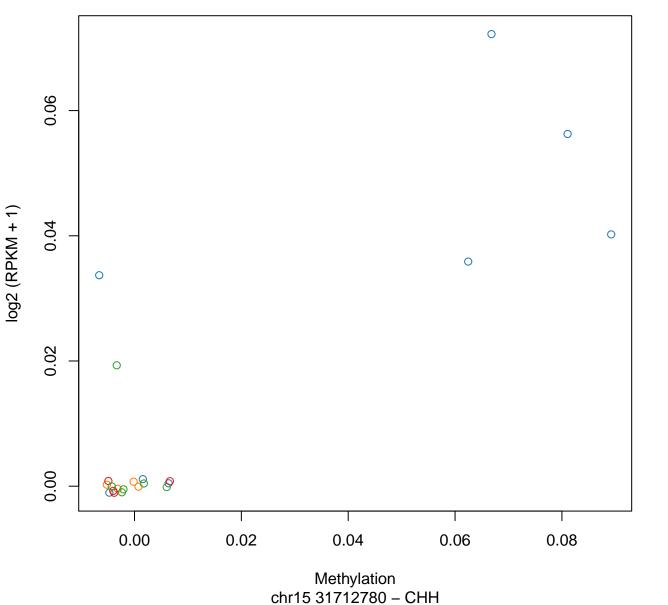
chr6 67068737 + CHG

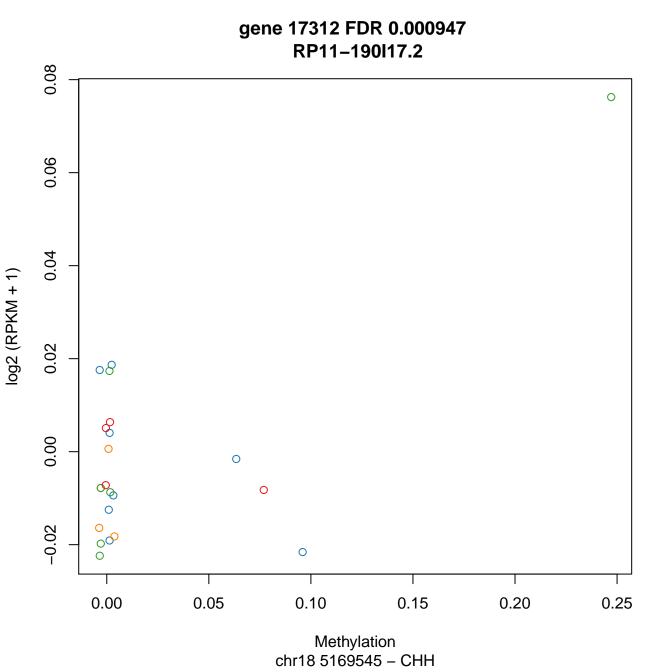


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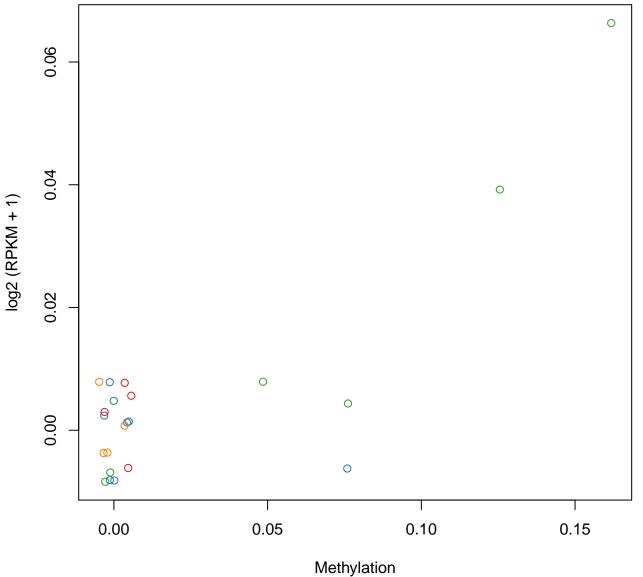


gene 16225 FDR 9e-04 UBE2CP4

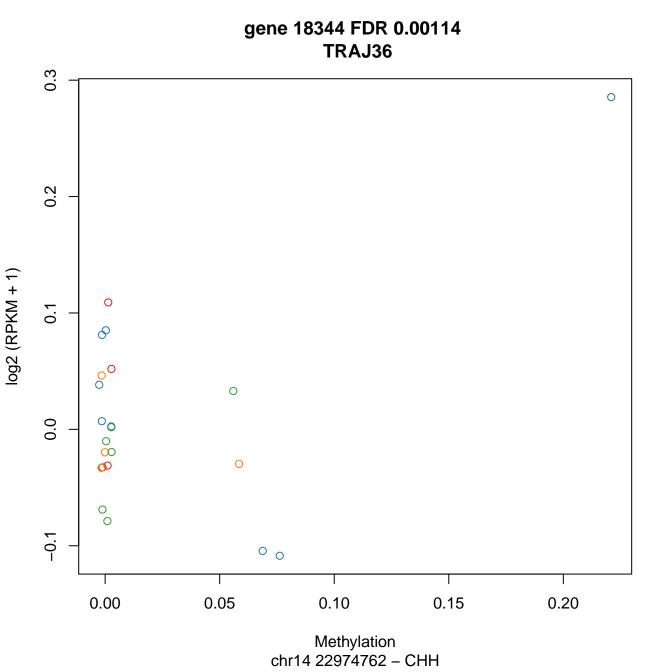


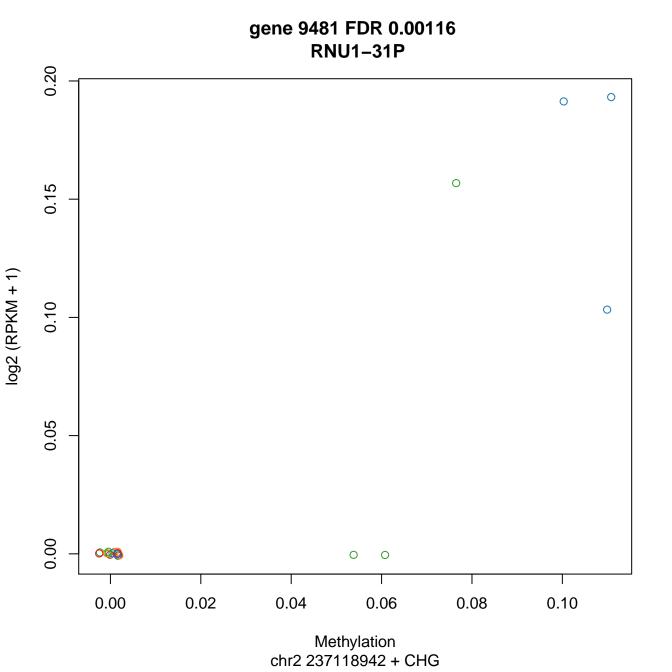


gene 13307 FDR 0.000983 RP3-449H6.1

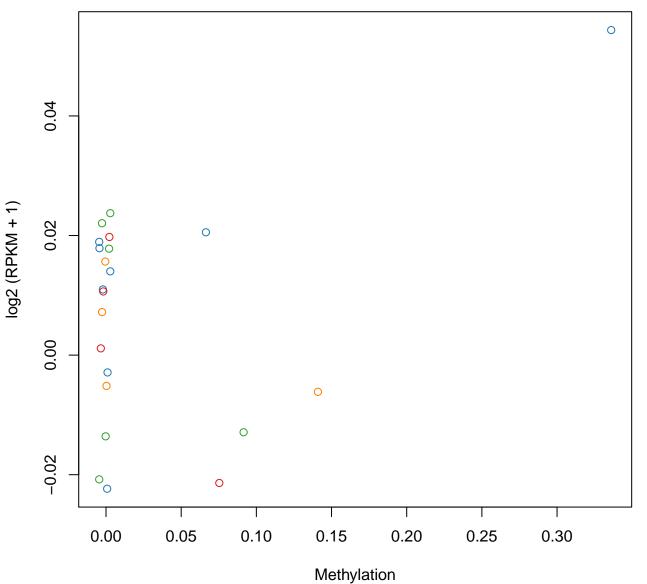


chr6 44495234 + CHG



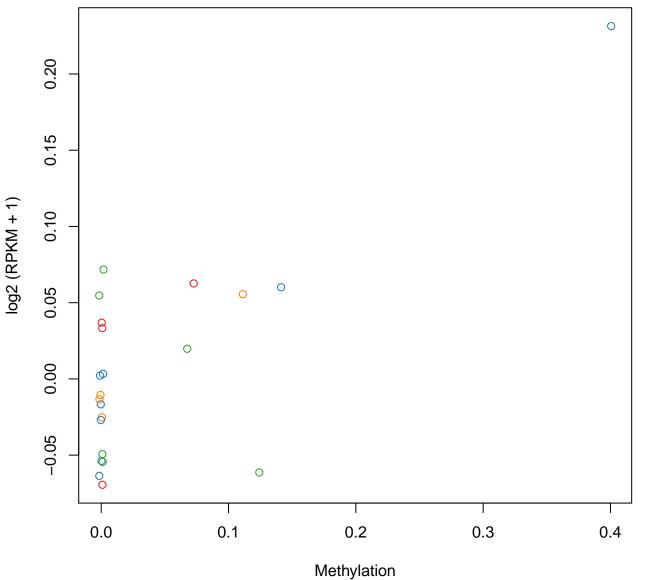


gene 17394 FDR 0.00117 RN7SL7P



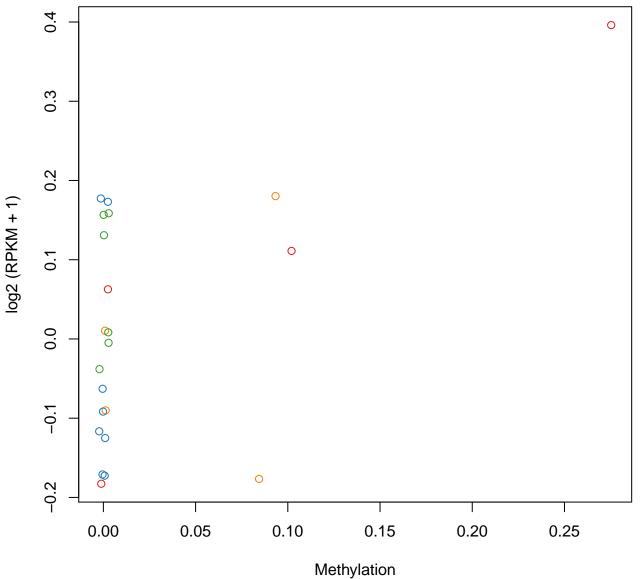
chr7 92601464 + CHG

gene 17212 FDR 0.00118 MIR4684



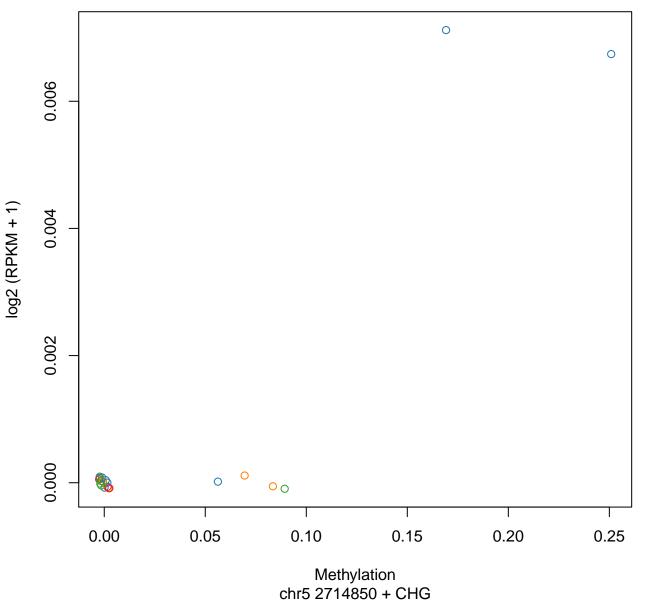
chr1 23045109 - CHG

gene 13464 FDR 0.00118 RNU7-38P

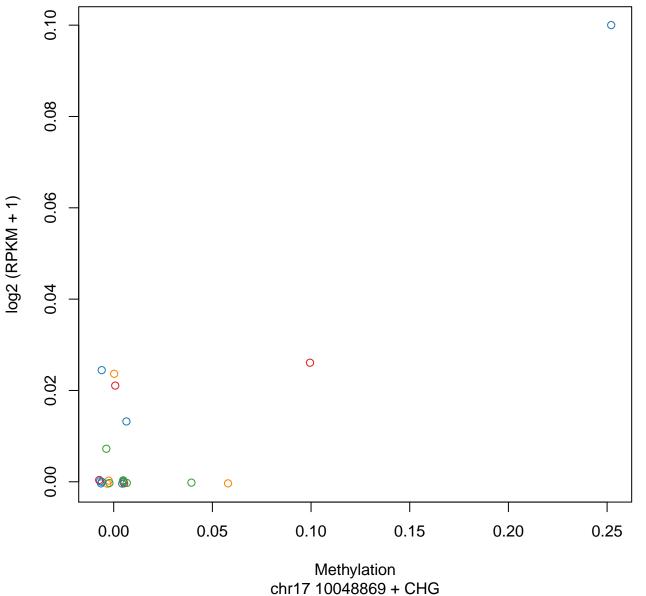


chr10 73603512 + CHH

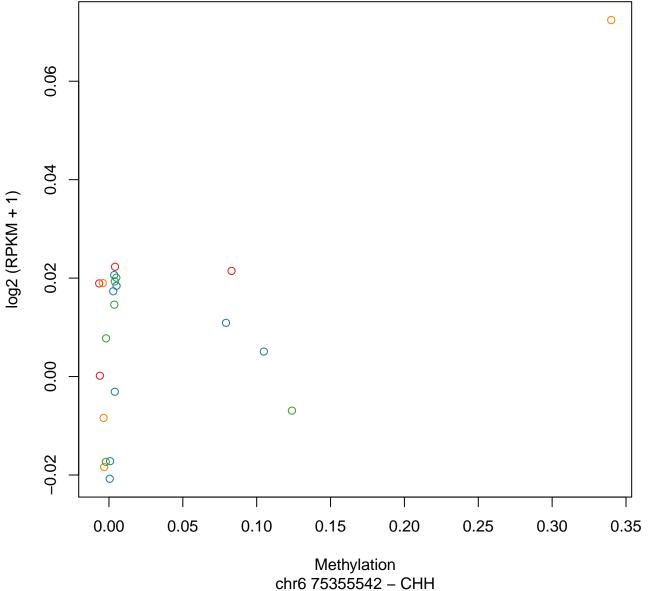
gene 18762 FDR 0.00125 LSINCT5



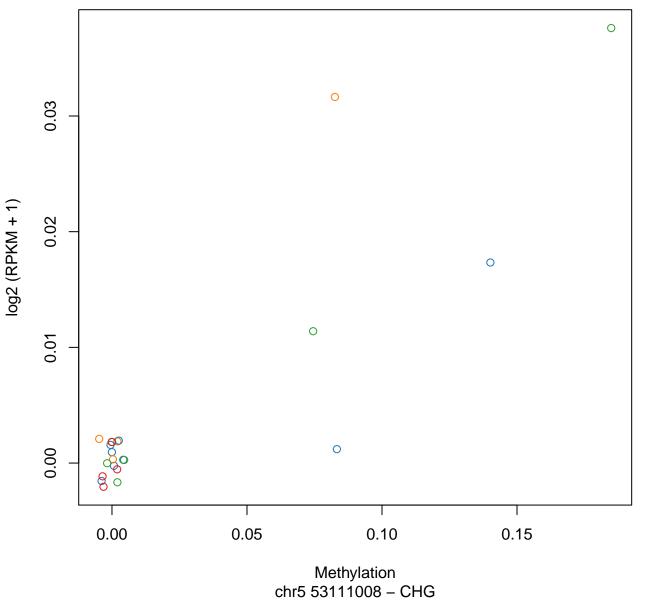
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gene 10973 FDR 0.00128 RP11-554D15.4

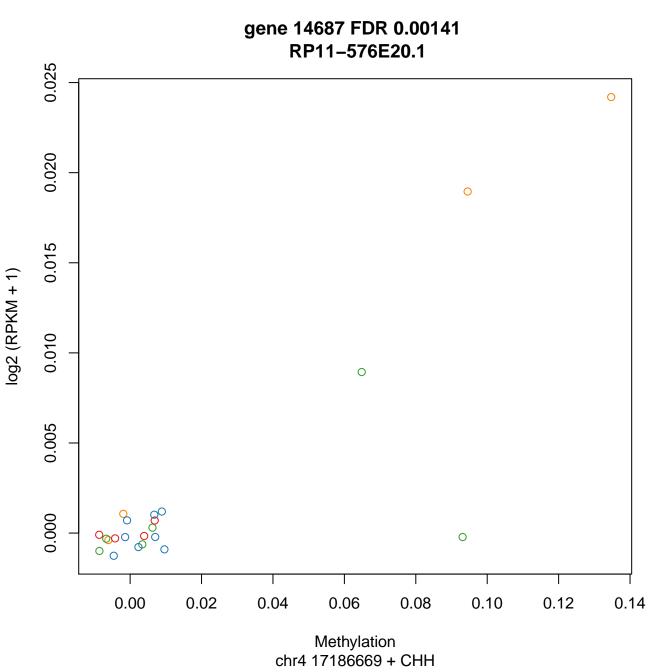


gene 14612 FDR 0.00129 CTD-2081C10.1

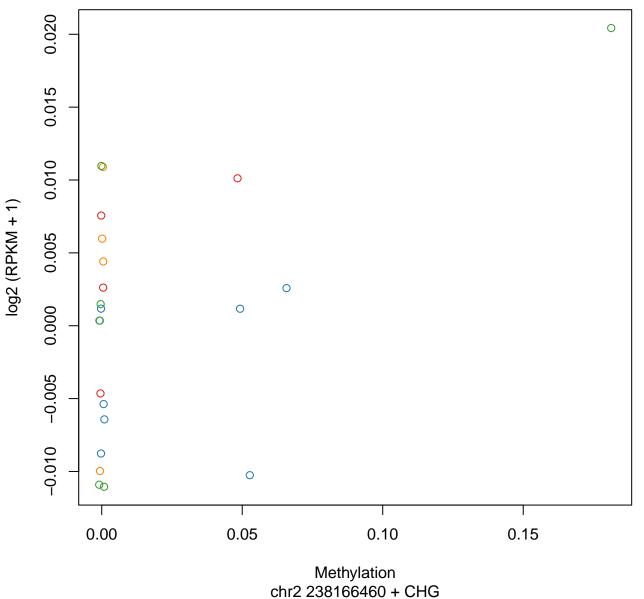


gene 14762 FDR 0.00139 RP11-1057B8.2 0.025 0 0.020 0.015 log2 (RPKM + 1) 0.010 000 0.005 0 0 0 0.000 0 00 0 -0.0050 0 0.00 0.05 0.10 0.15 0.20 0.25 0.30 0.35 Methylation

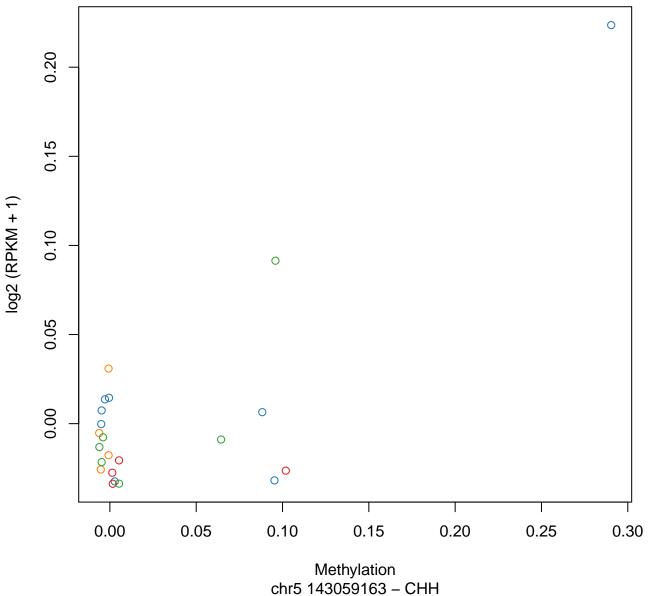
chr8 135869807 + CHG



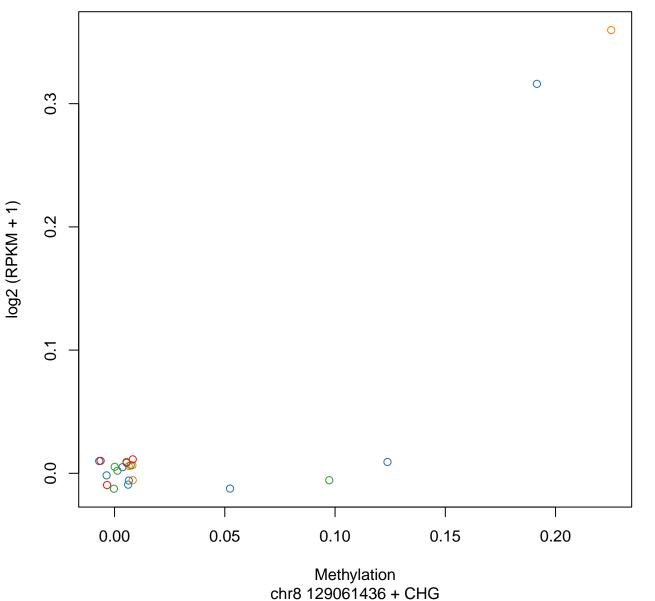
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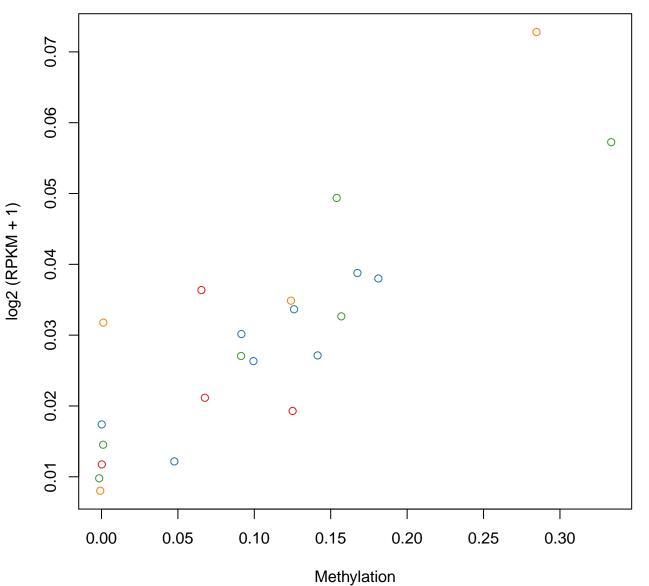
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gene 10600 FDR 0.00147 MIR1207

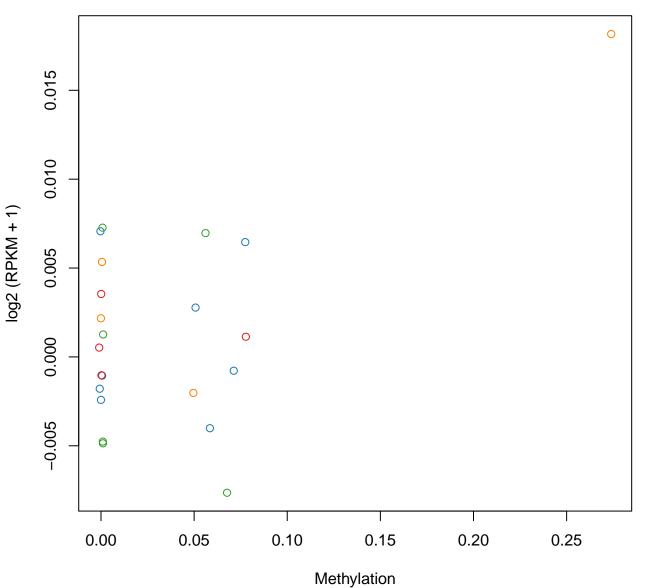


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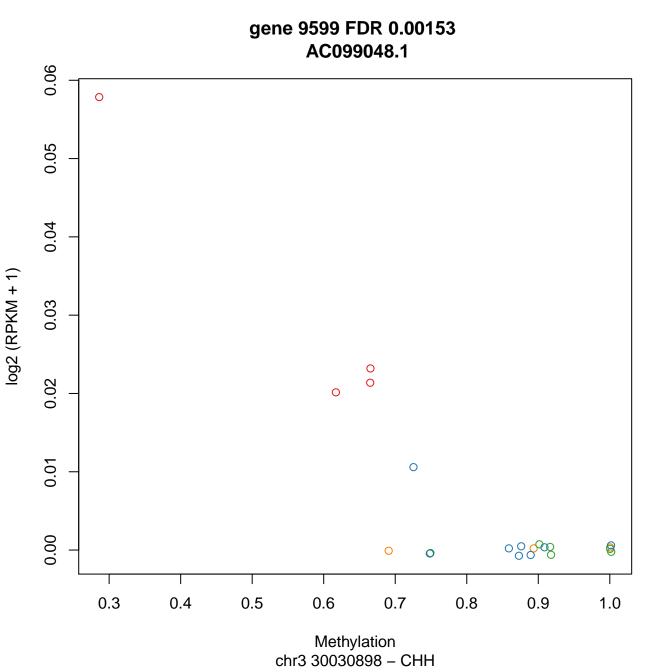


chr1 207265808 - CHG

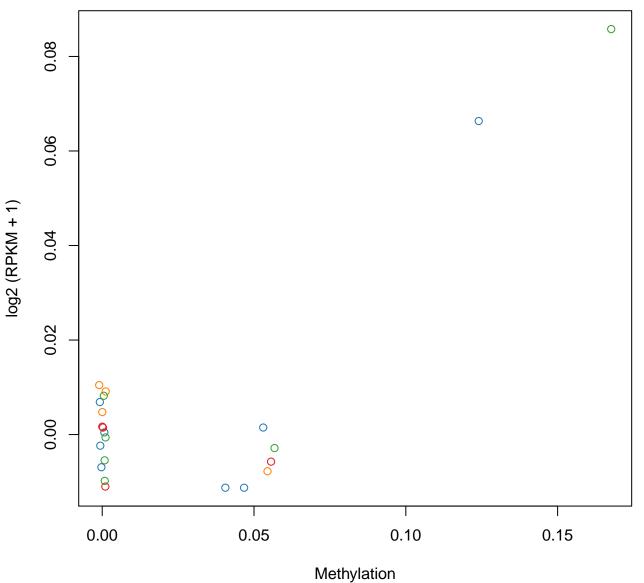
gene 11384 FDR 0.00151 HAUS4P1



chr1 170338816 - CHG

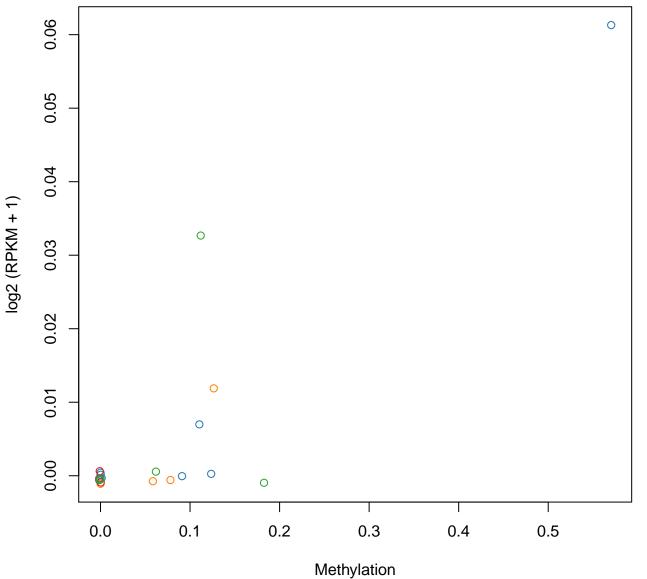


gene 13792 FDR 0.00162 RP11-237P21.1



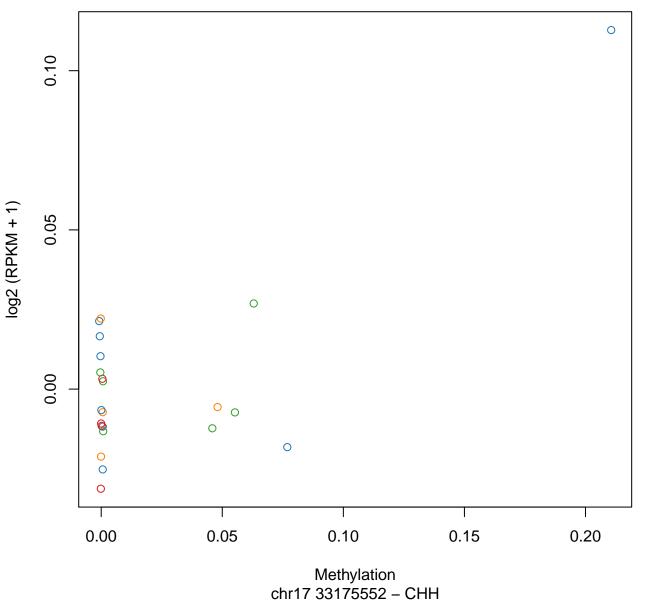
chr3 135644209 - CHH

gene 18833 FDR 0.00166 RP11-1223D19.6

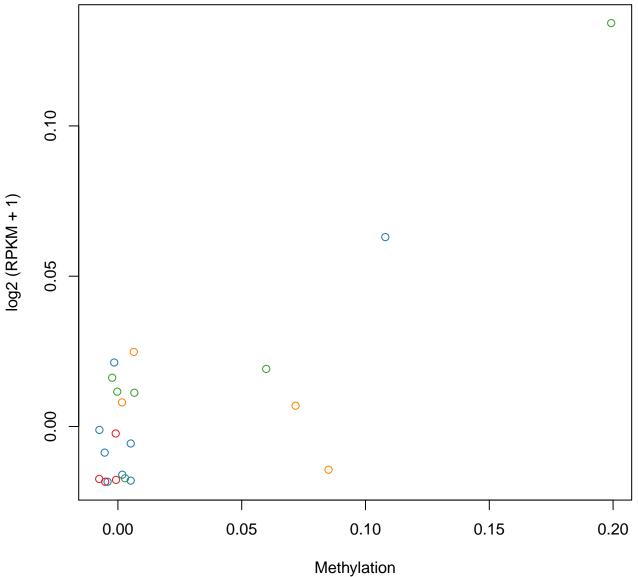


chr2 111092312 - CHH

gene 17086 FDR 0.00172 RP11-642M2.1

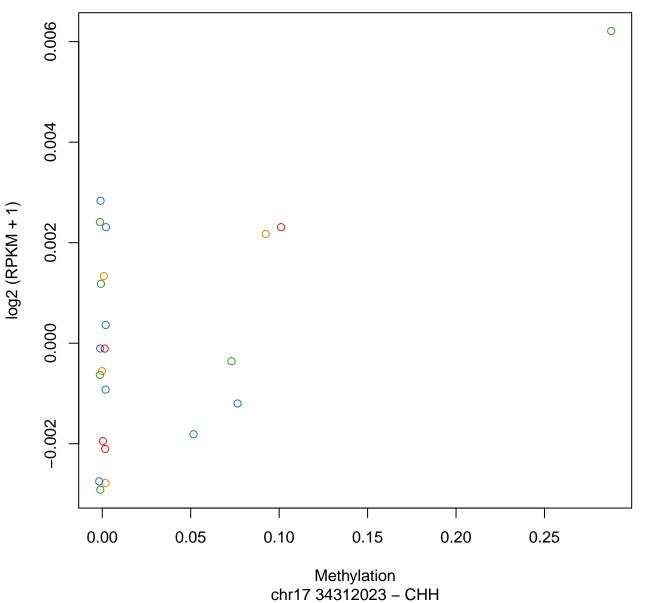


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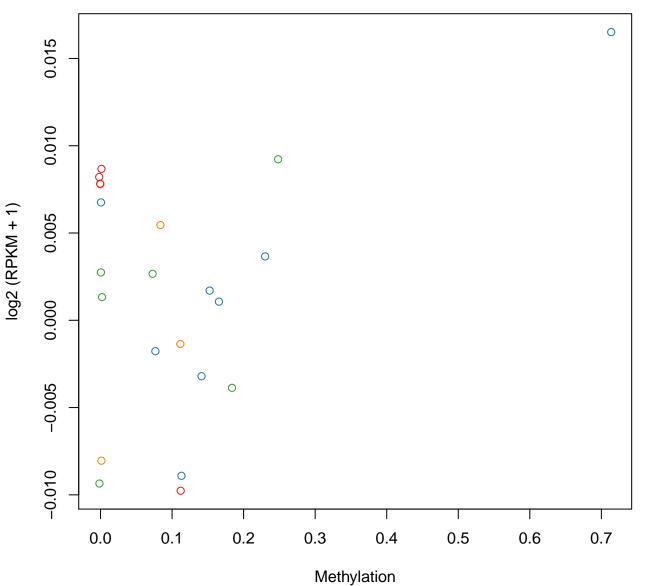


chr10 112864051 - CHH

gene 18260 FDR 0.00183 CTB-186H2.3

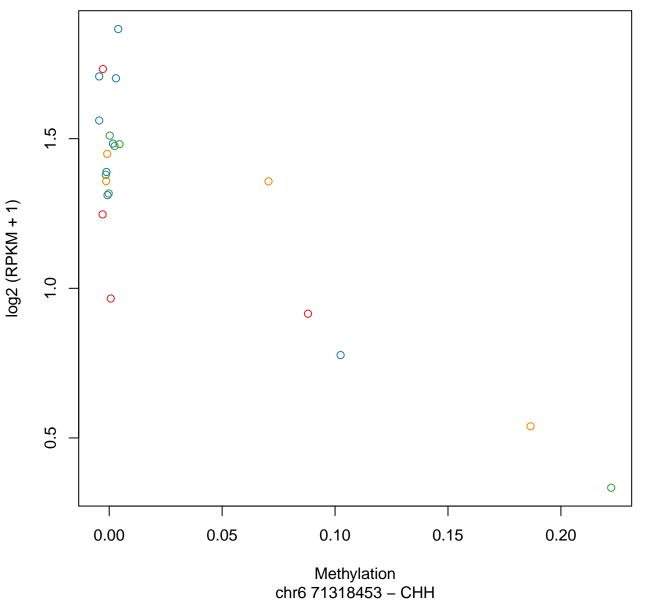


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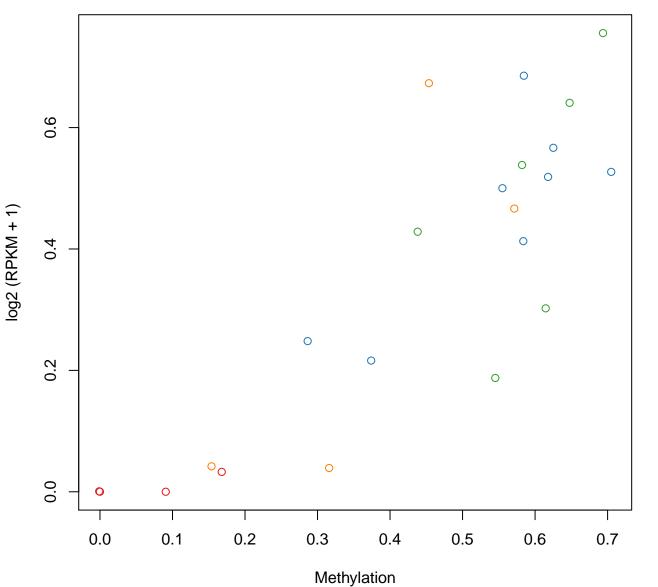


chr1 234844292 + CHH

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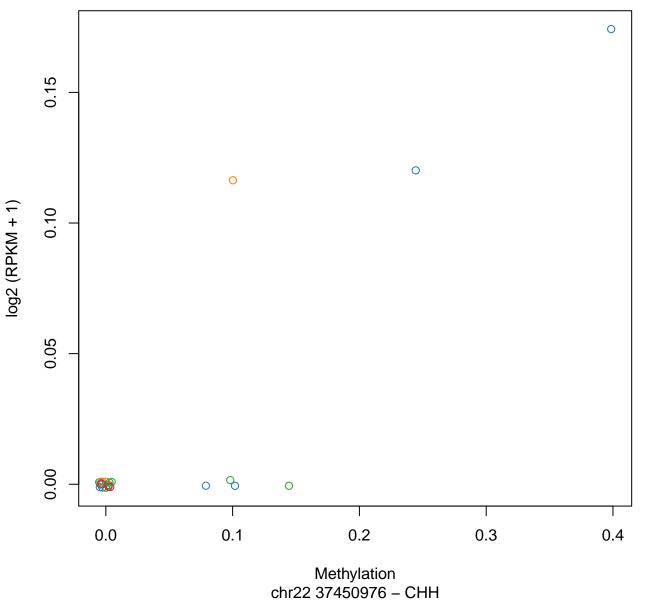


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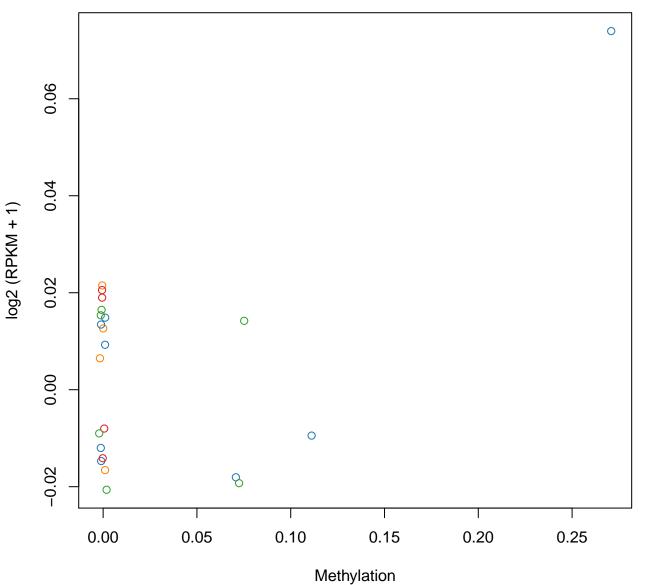


chr1 191115863 - CHH

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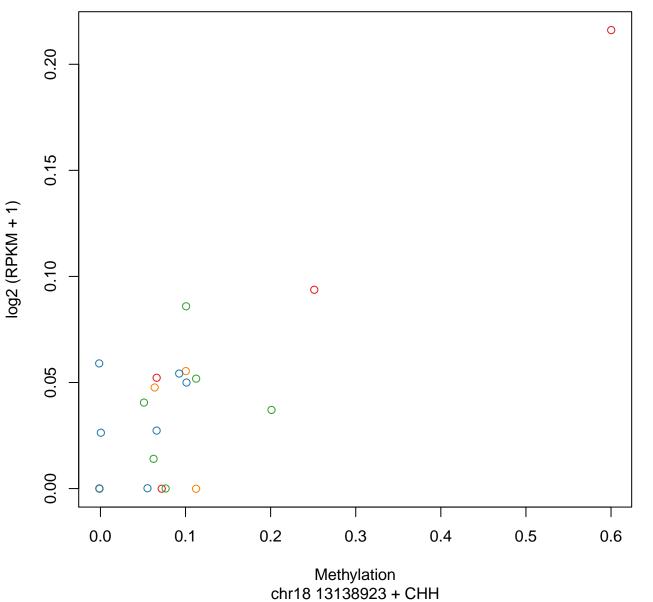


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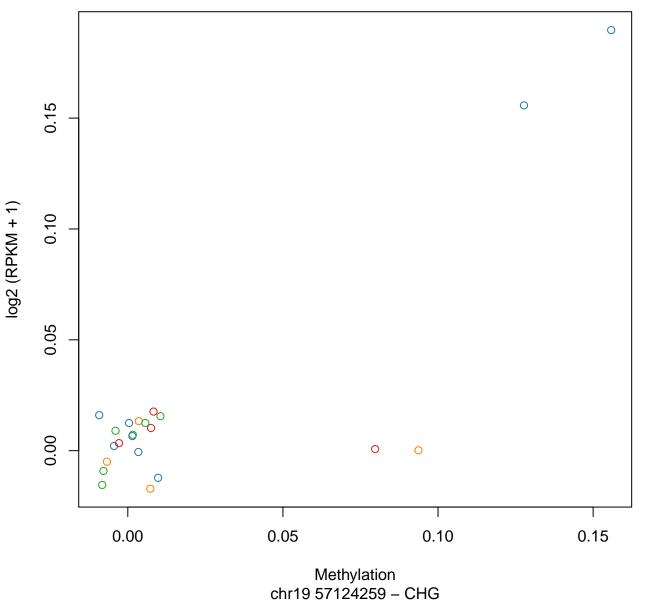


chr2 112432812 + CHH

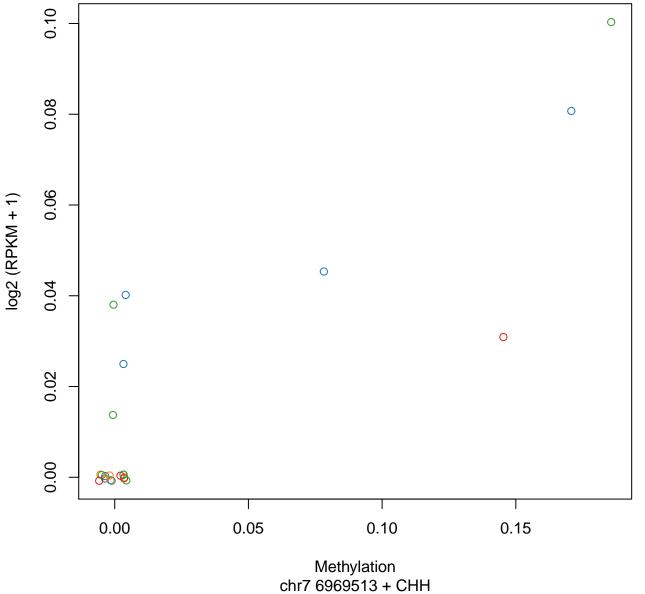
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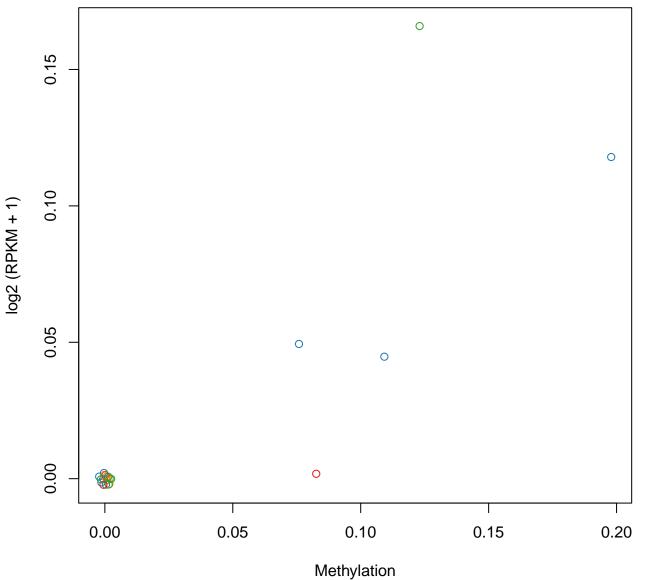
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gene 11381 FDR 0.0022 ALG1L5P

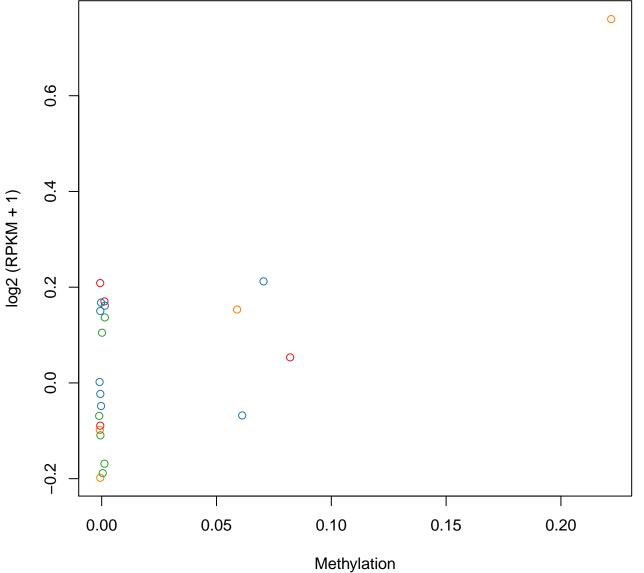


gene 11438 FDR 0.00225 SPATA13-AS1



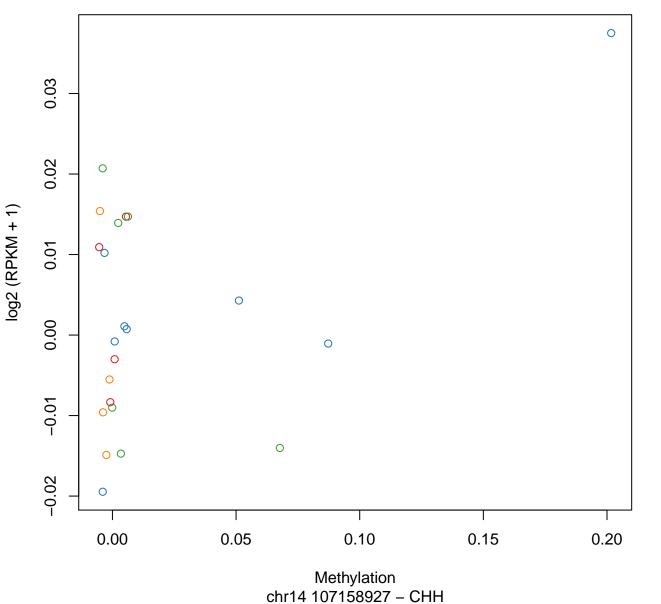
chr13 24828635 - CHH

gene 9545 FDR 0.00225 Y_RNA

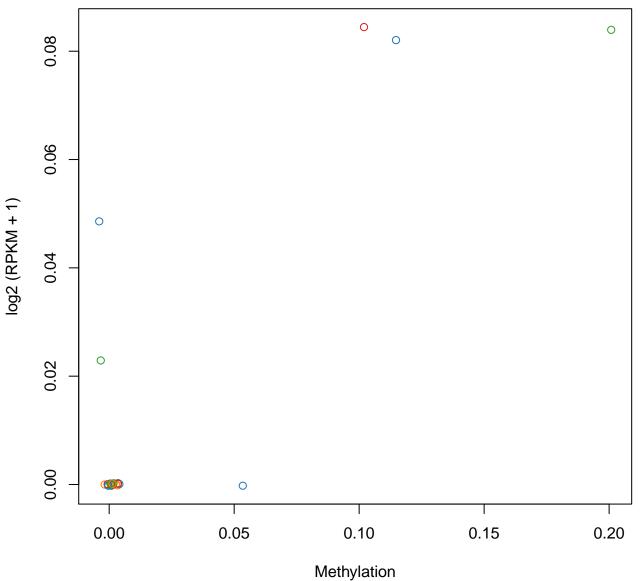


chr9 98864563 + CHH

gene 18789 FDR 0.00235 CH17-212P11.6

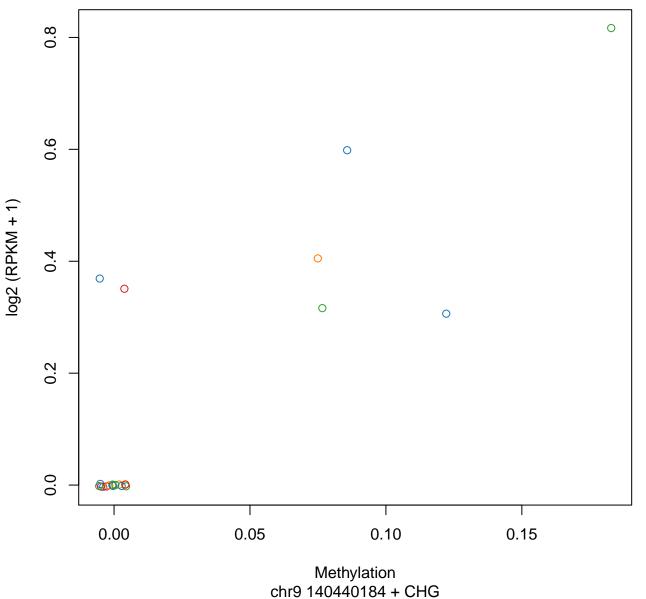


gene 15778 FDR 0.00235 RP11-7M8.2

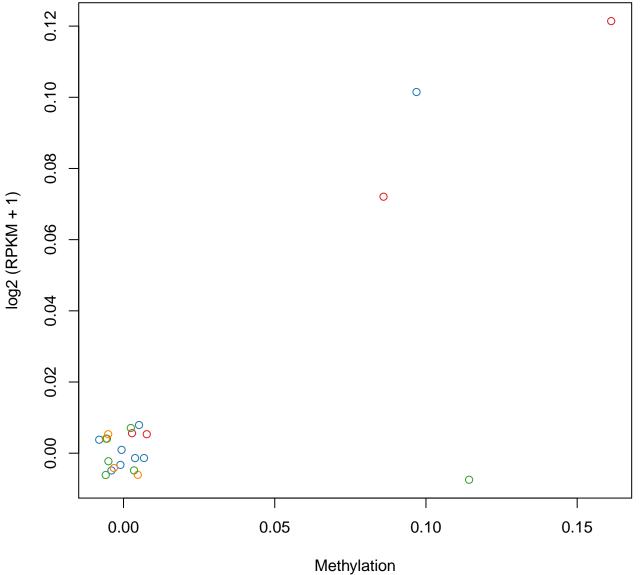


chr12 122295358 + CHG

gene 16983 FDR 0.00239 AL365502.1

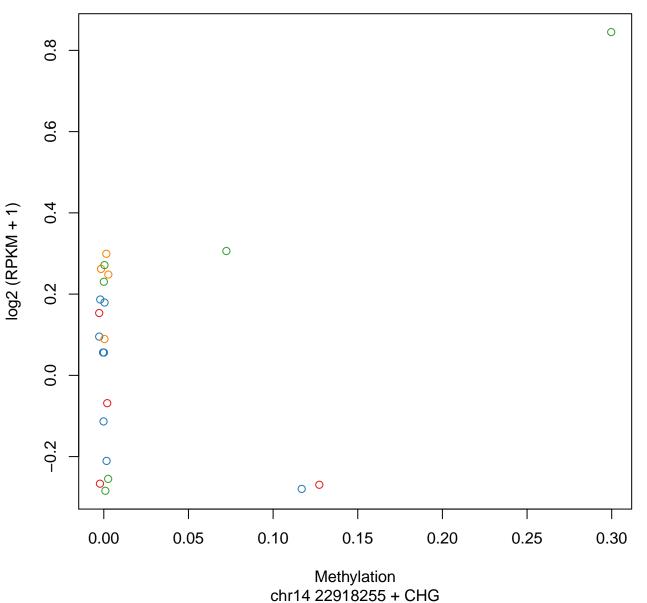


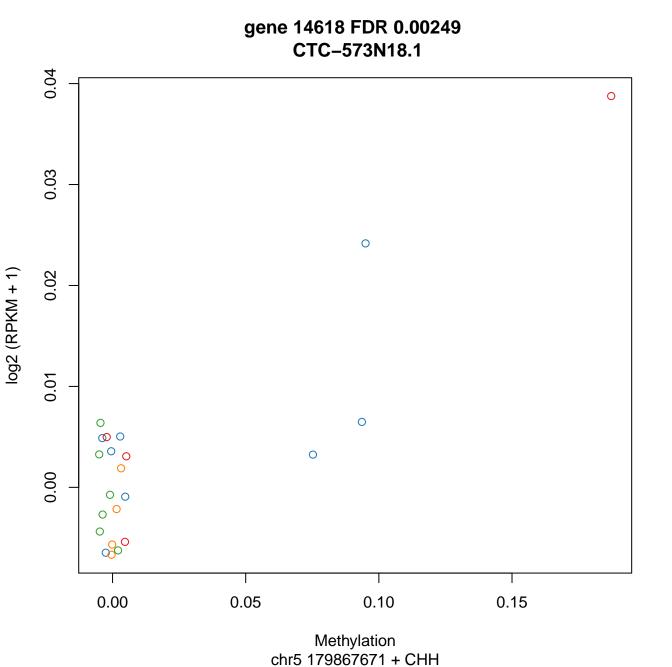
gene 14288 FDR 0.00241 RP11-287J9.1



chr5 65918555 + CHH

gene 11740 FDR 0.00247 TRDD3





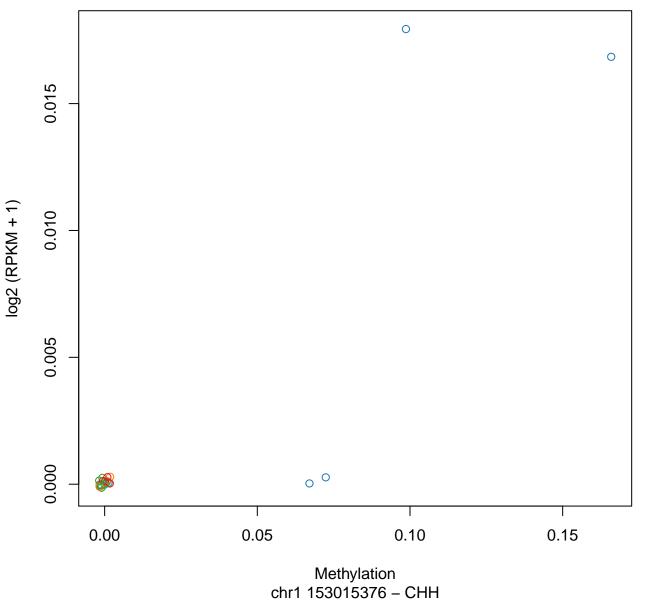
gene 10033 FDR 0.00252 AL121588.1 0.15 0 log2 (RPKM + 1) 0.05 00 0 00 0 0 0 8 0.00 0 0 0 8 -0.050 0.00 0.05 0.10 0.15 0.20 0.25 Methylation

chr20 37901118 + CHH

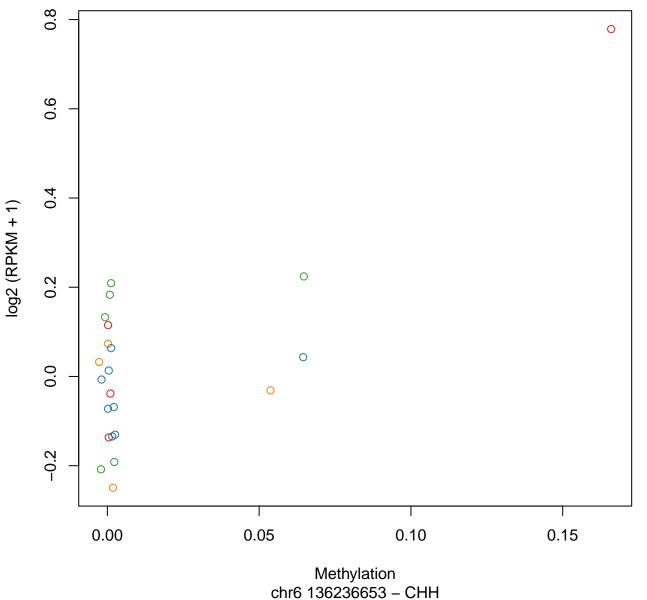
gene 15093 FDR 0.00259 RP11-107N7.1 0.10 0 0 0.08 90.0 log2 (RPKM + 1) 0 0.04 0 0 0 0.02 0 0.00 0 0.00 0.05 0.10 0.15

Methylation chr5 77242901 – CHH

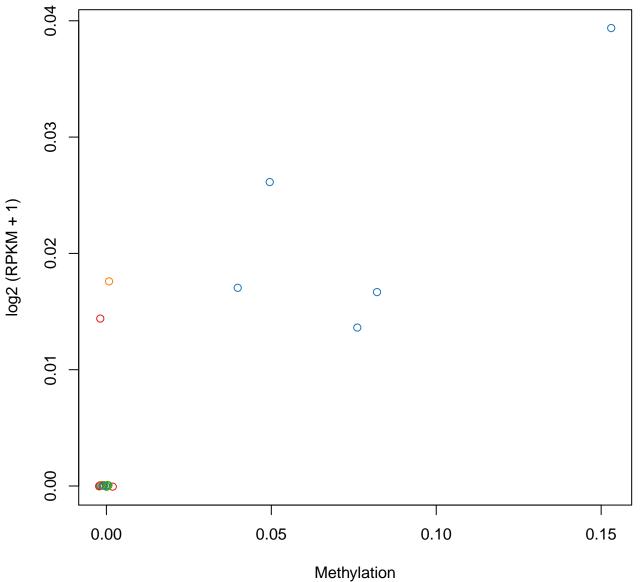
gene 6078 FDR 0.00264 SPRR2D



gene 13507 FDR 0.0027 AL360178.1



gene 16580 FDR 0.00275 LINC01227



chr16 80607366 + CHH