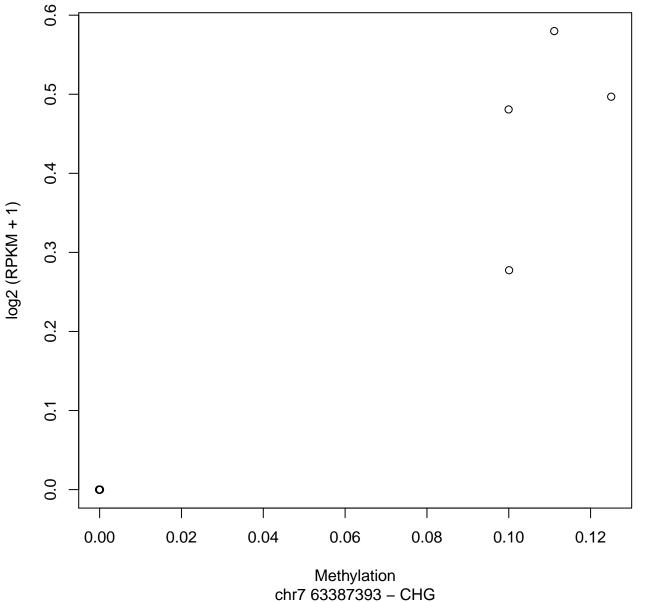
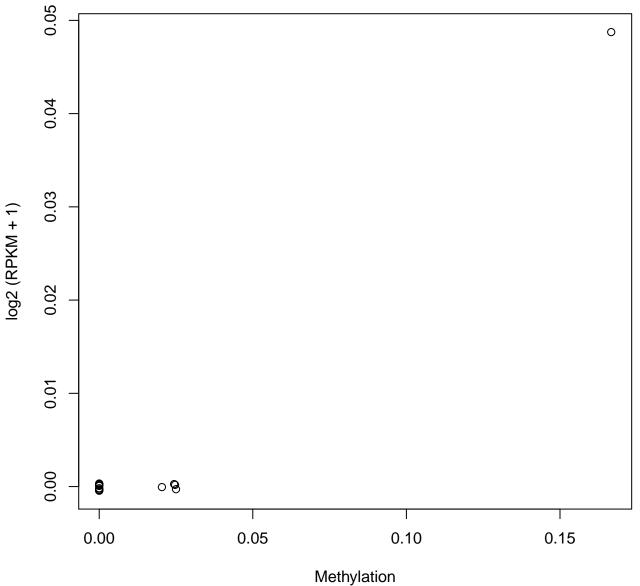
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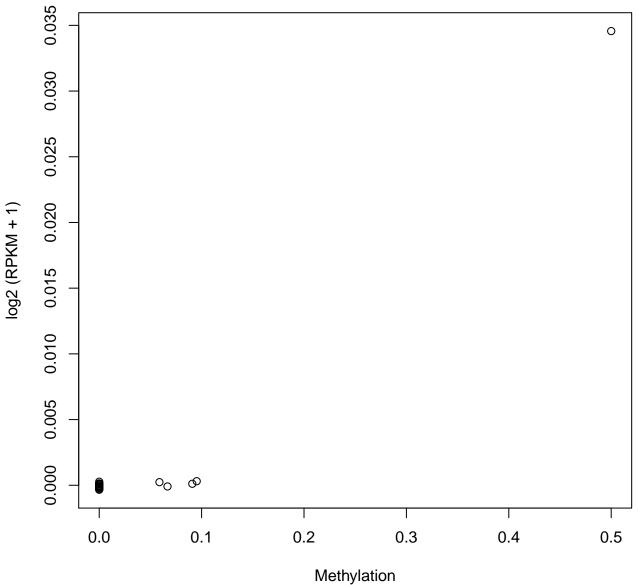


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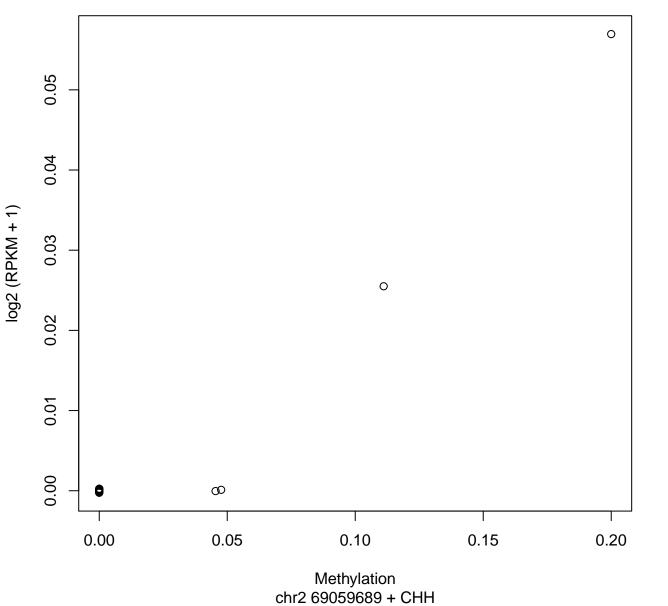
chr5 17519790 - CHH

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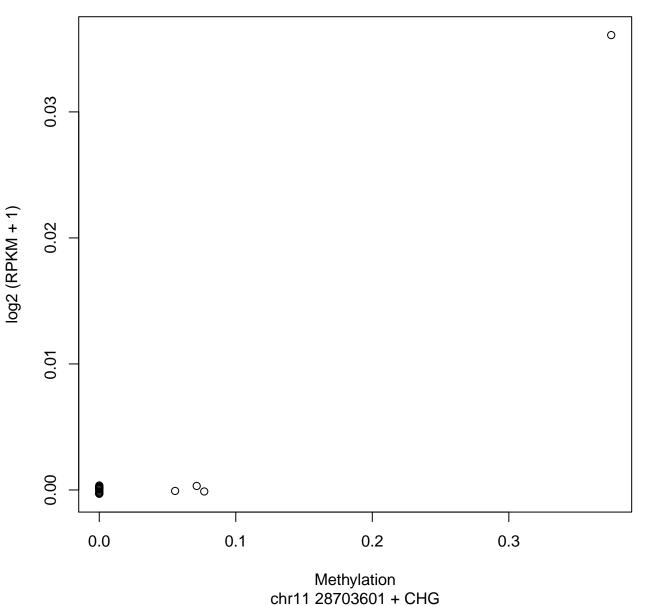


chr17 32503710 - CHH

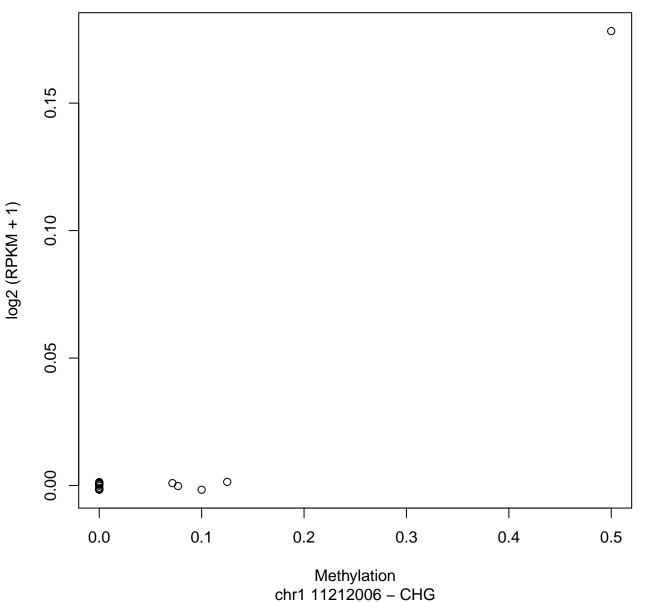
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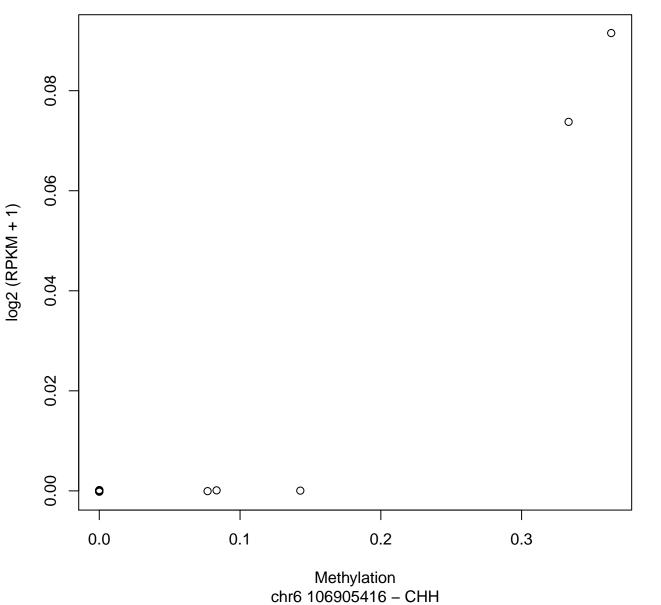
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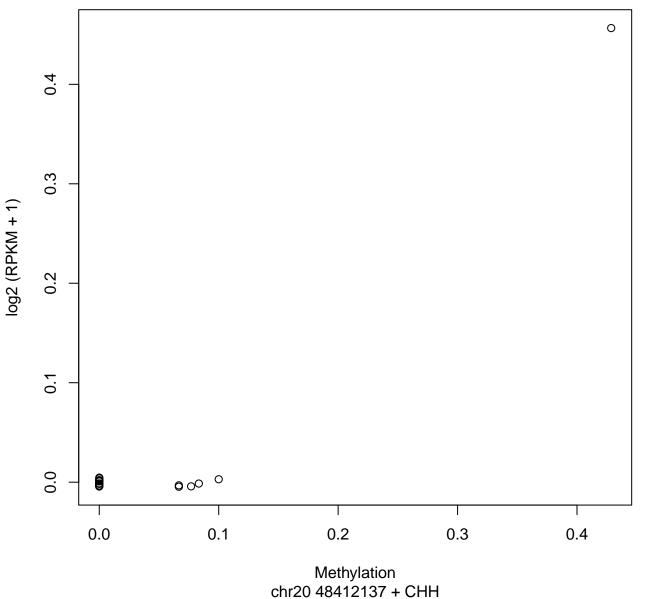
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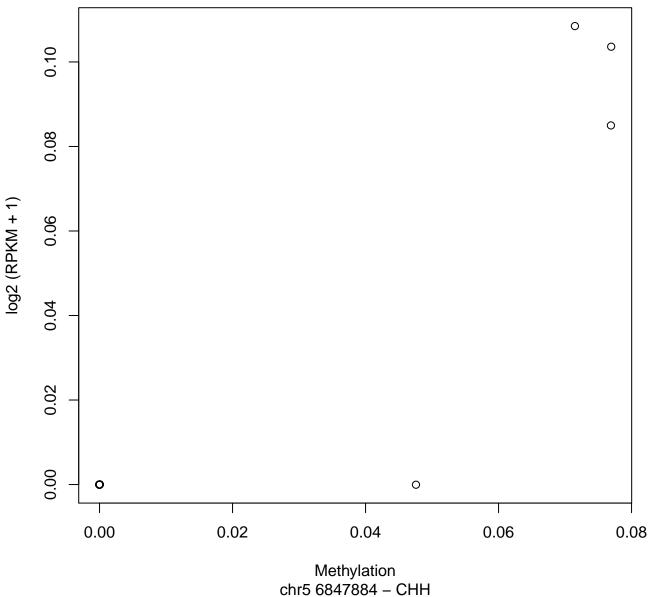
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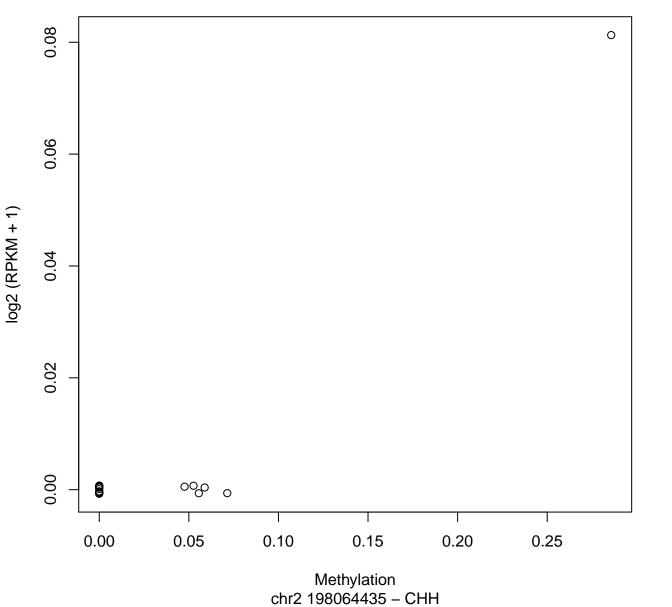
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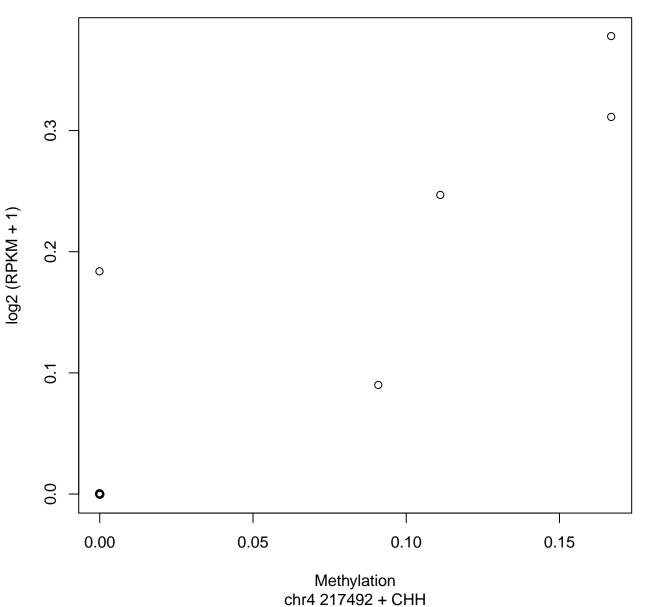
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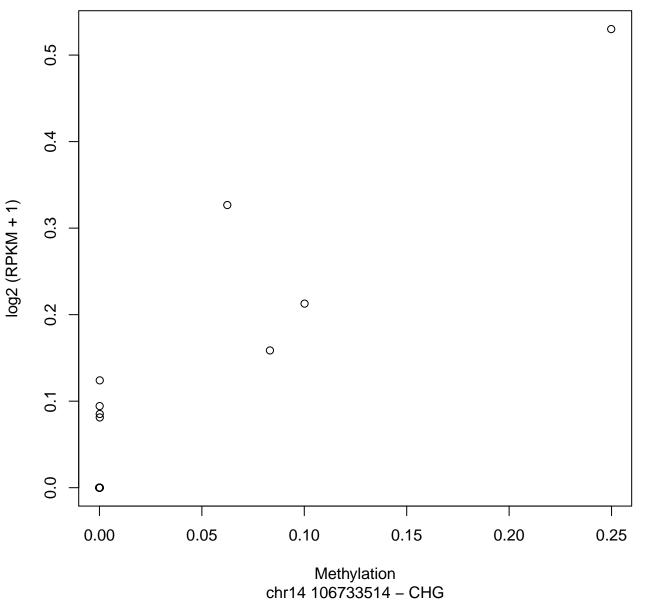
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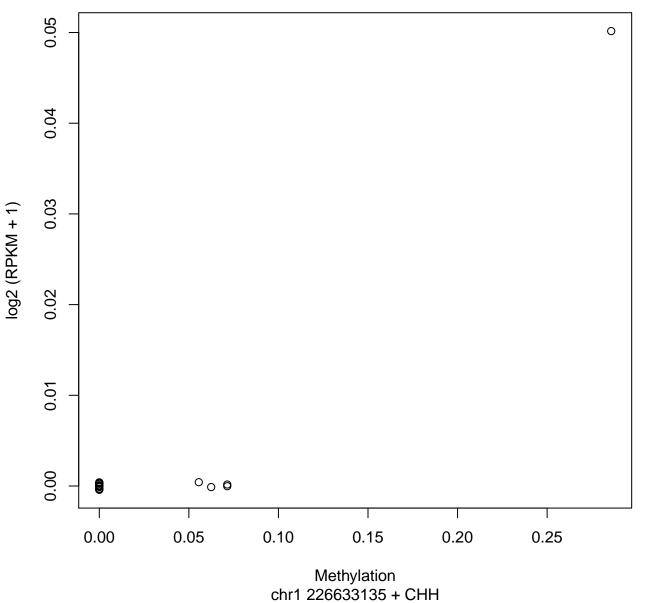
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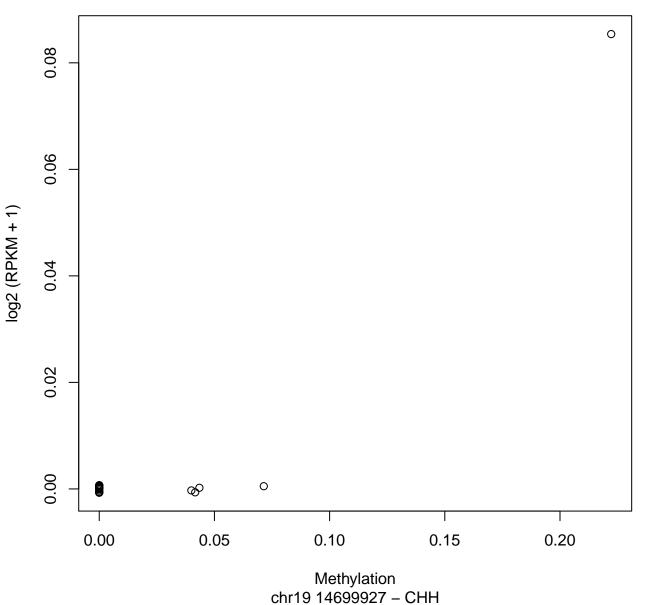
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gene ENSG00000223282.1 FDR 3.15e-05 RN7SKP165



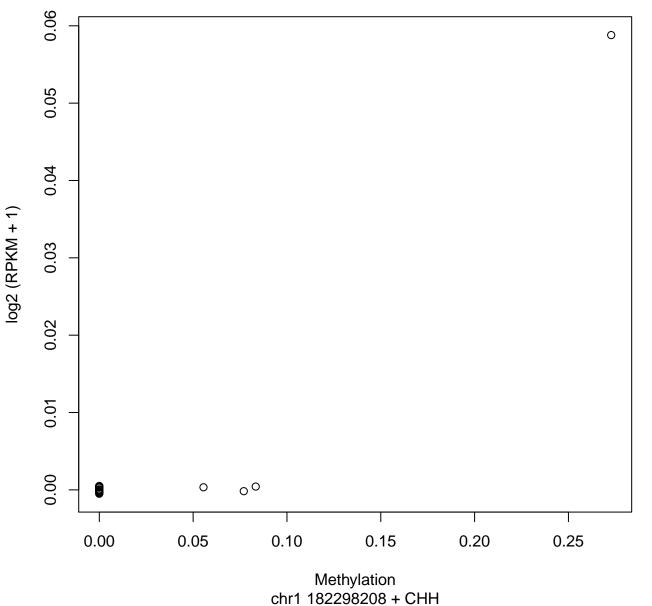
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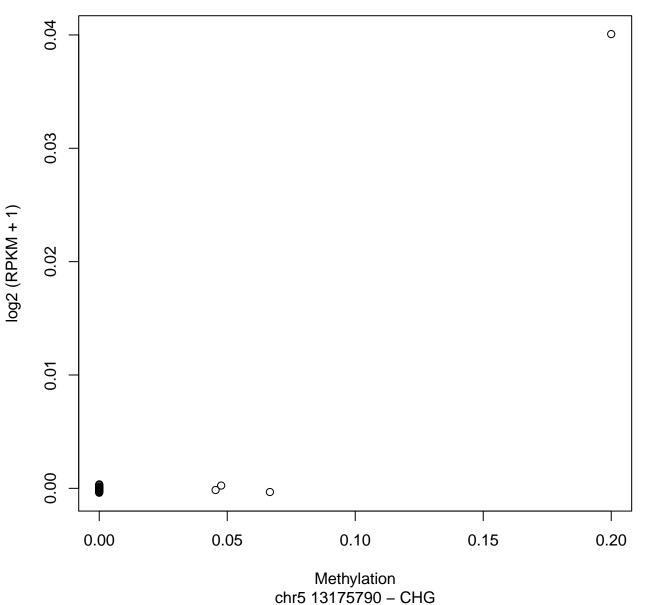
gene ENSG00000237876.1_1 FDR 4.29e-05 AKR1B1P4 0.020 0 0.015 log2 (RPKM + 1) 0.010 0.005 0.000 0 0 0.00 0.05 0.10 0.15 0.20 0.25 0.30 Methylation

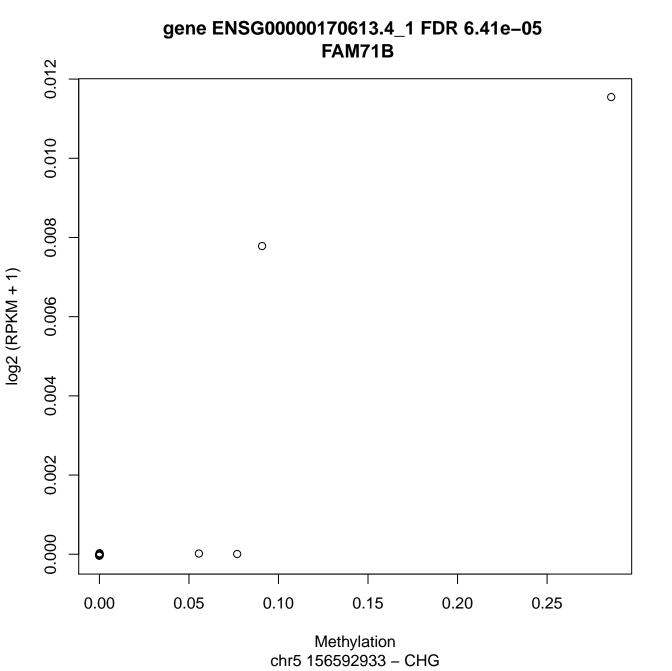
chr13 46263477 - CHH

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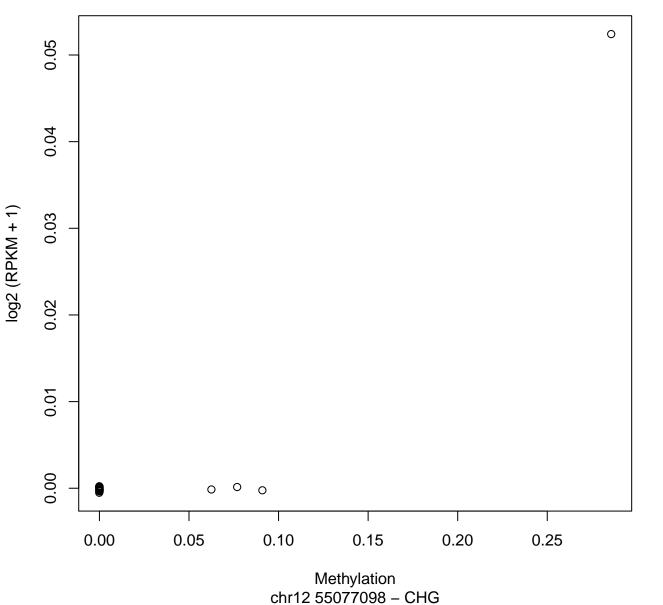


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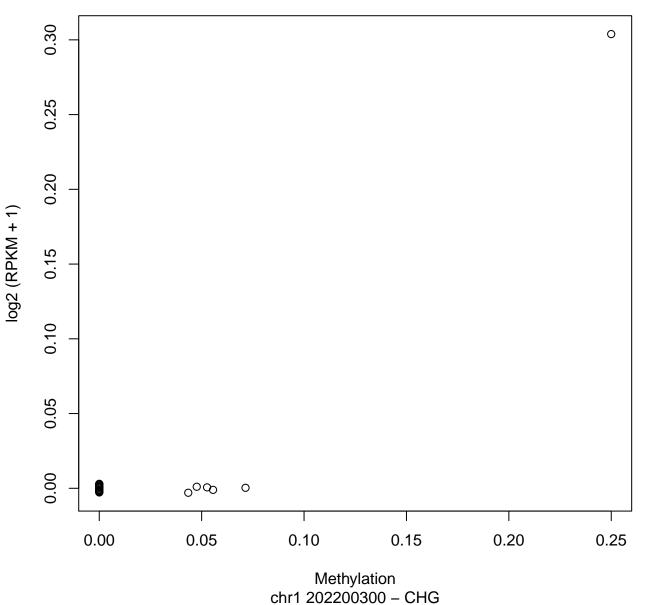




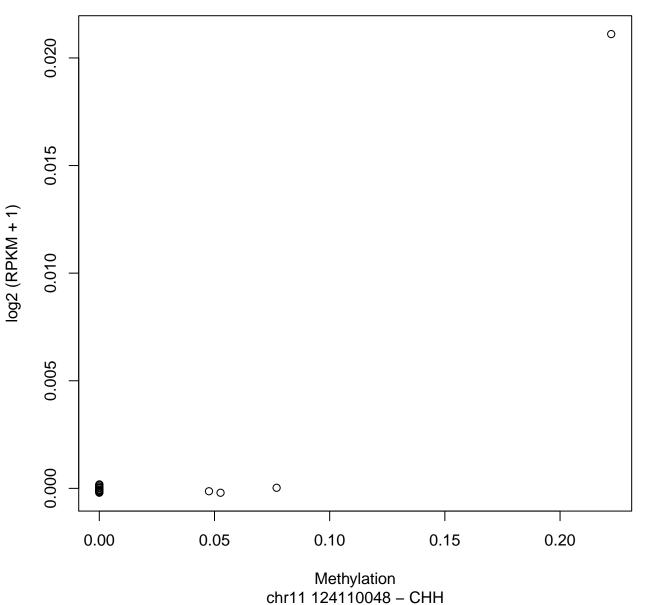
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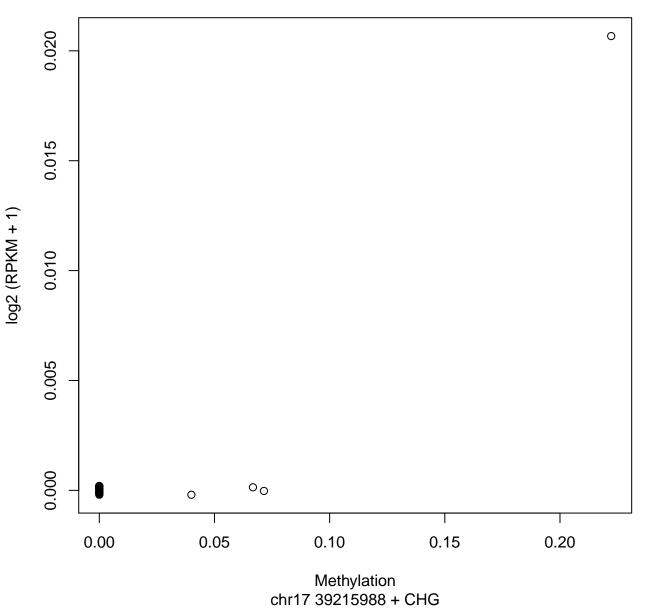
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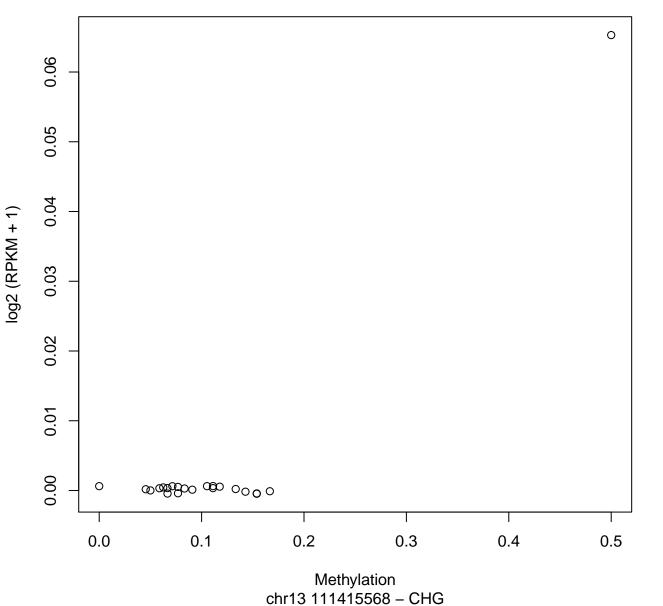
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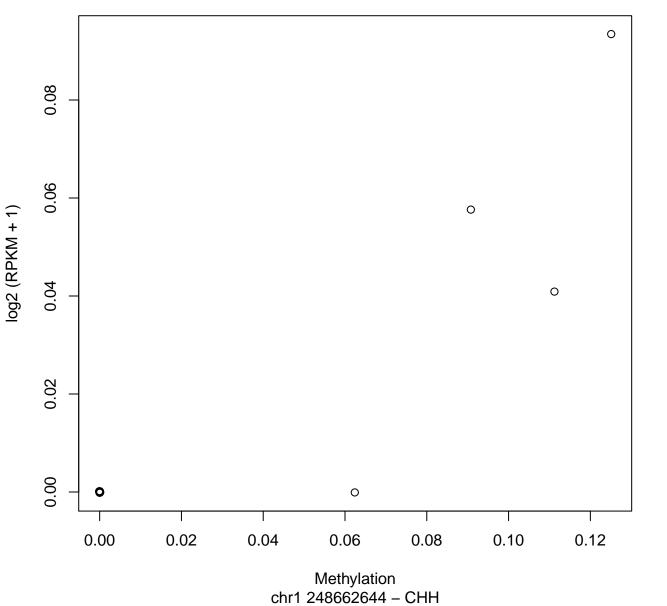
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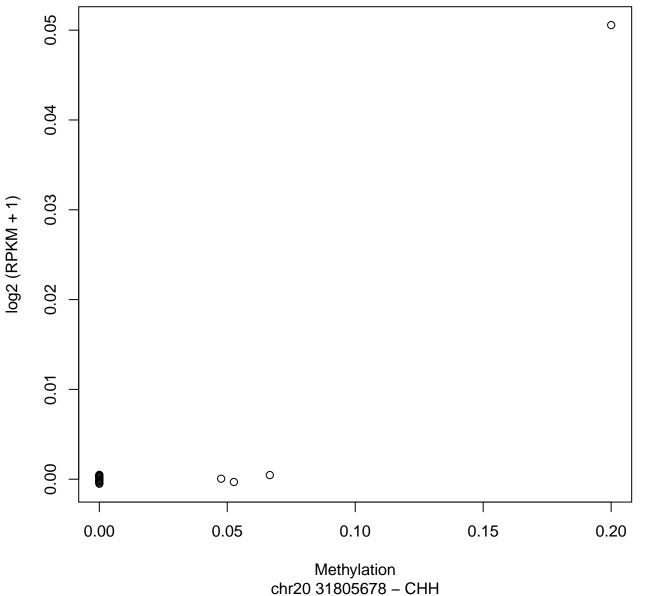
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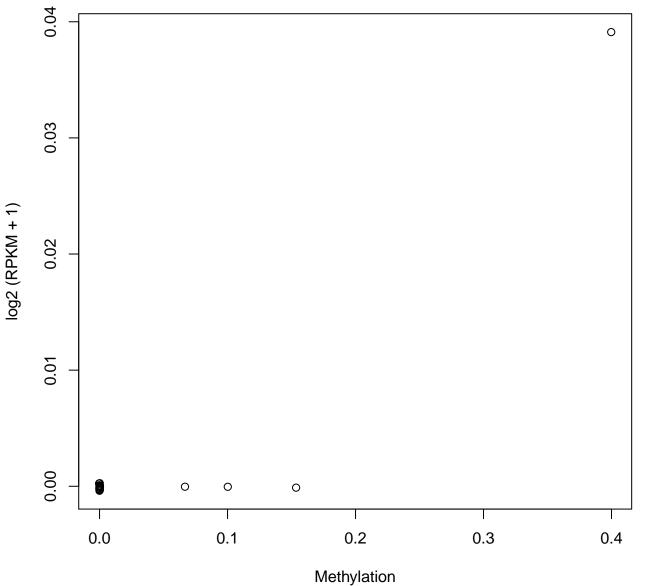
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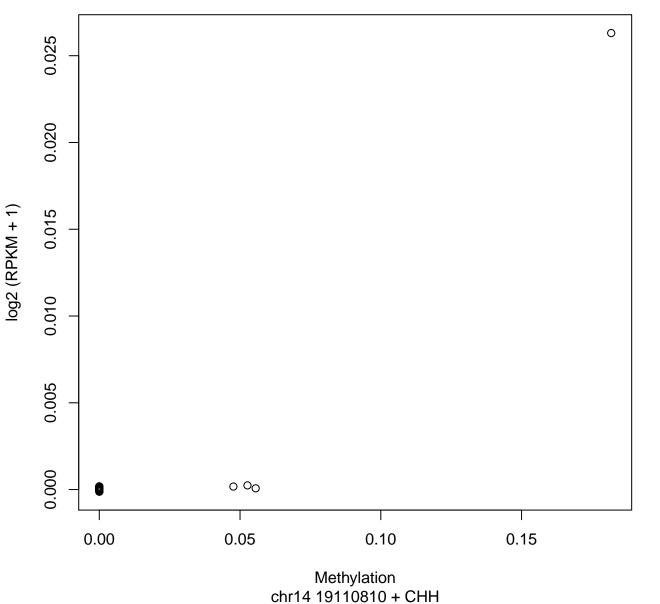


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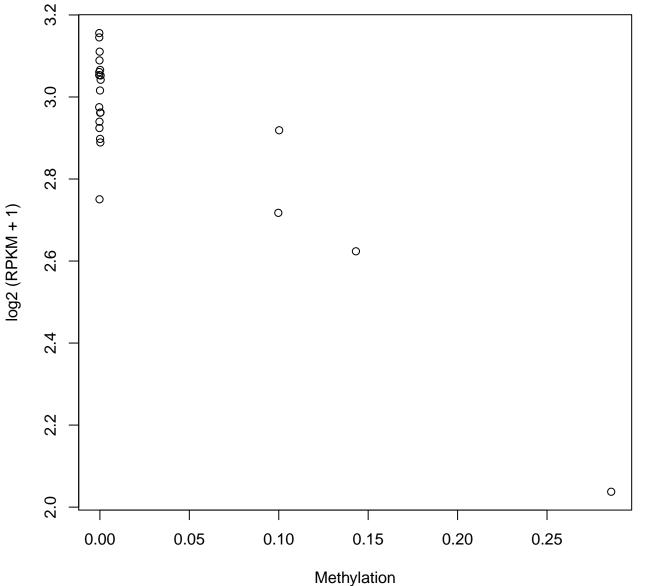


chr3 180131345 + CHH

gene ENSG00000215398.11_1 FDR 0.00014 RP11-754I20.1

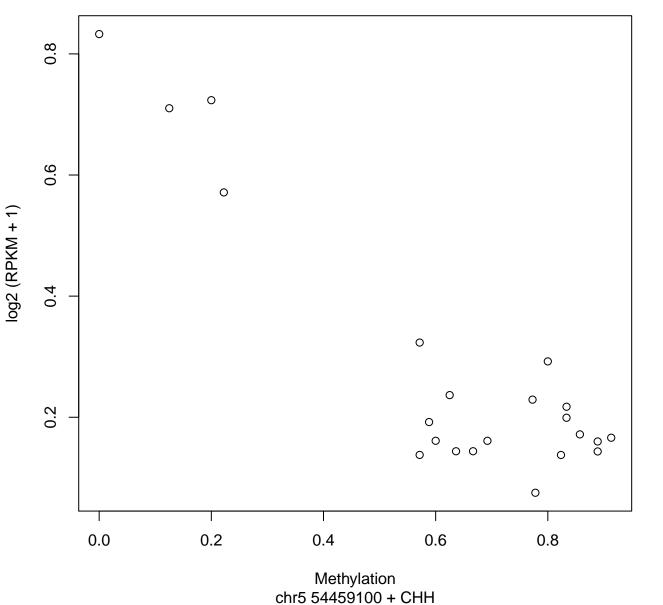


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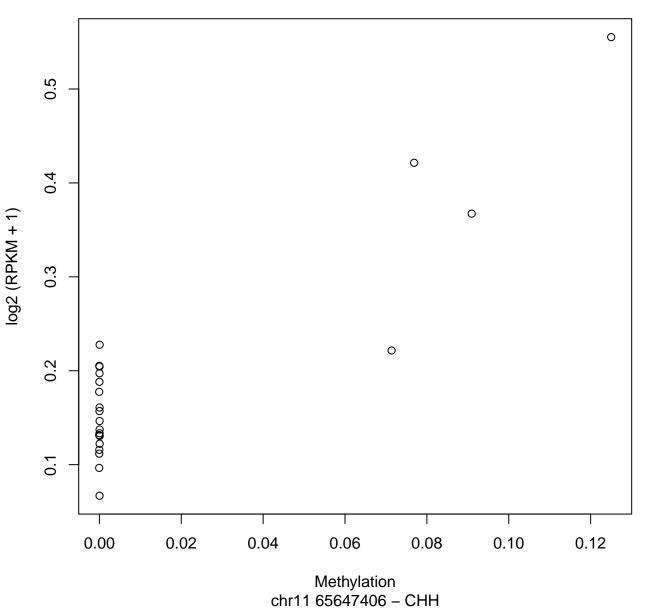


chr15 48462993 - CHH

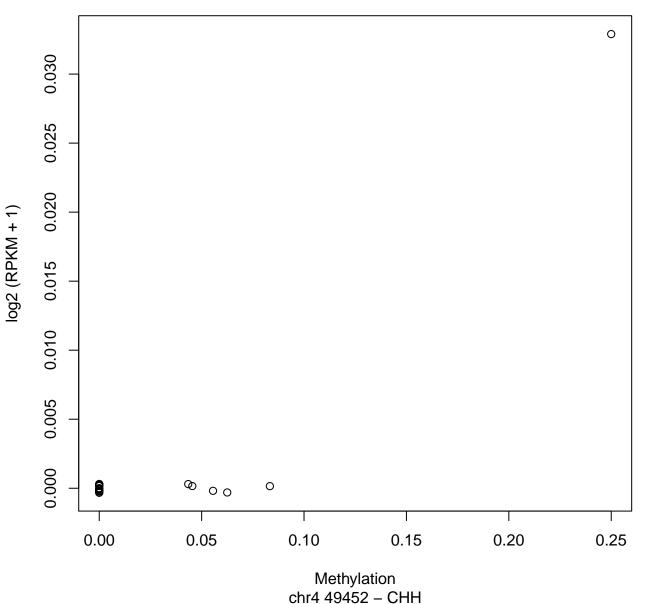
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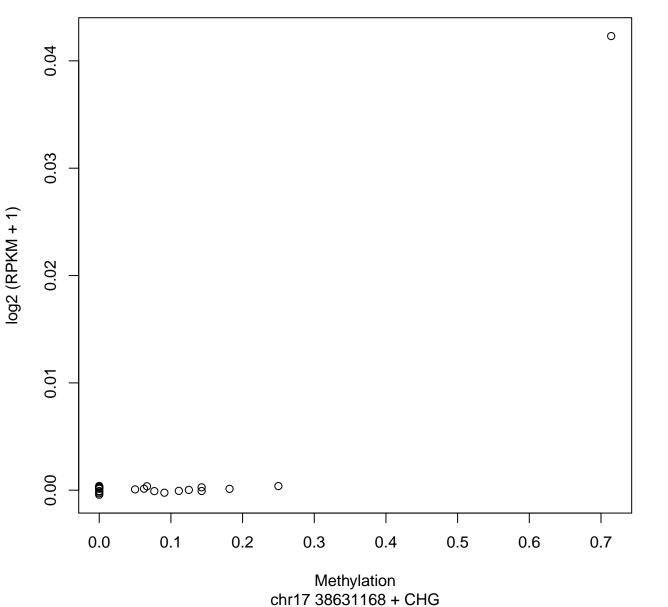
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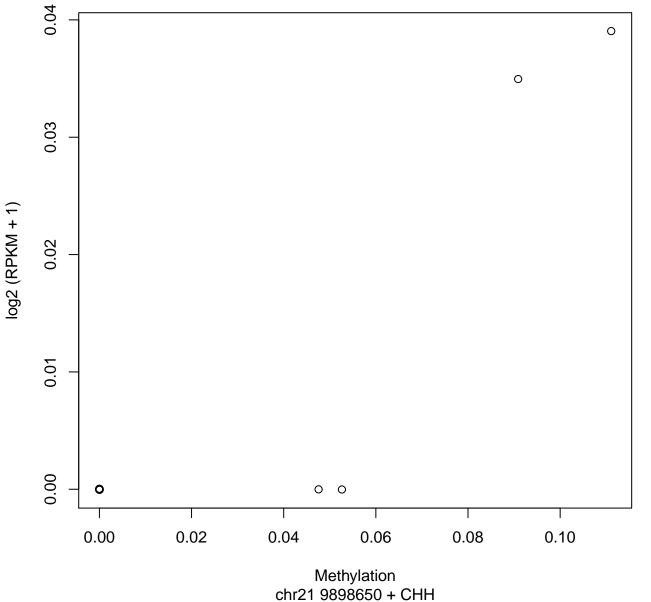
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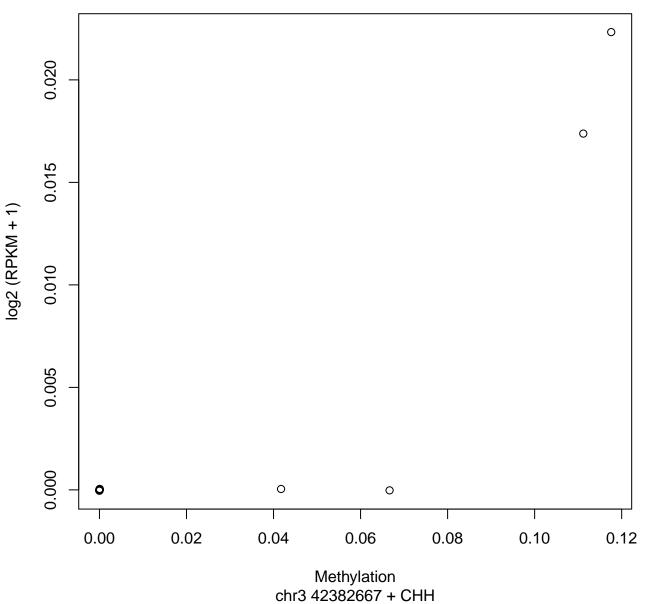
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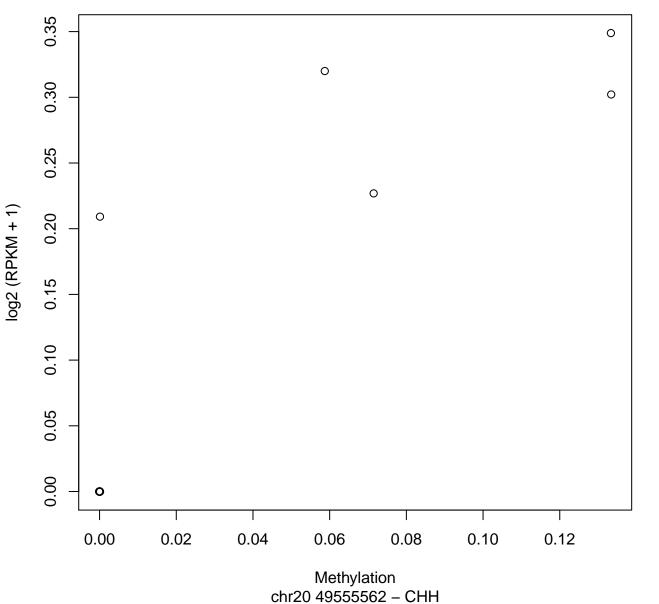
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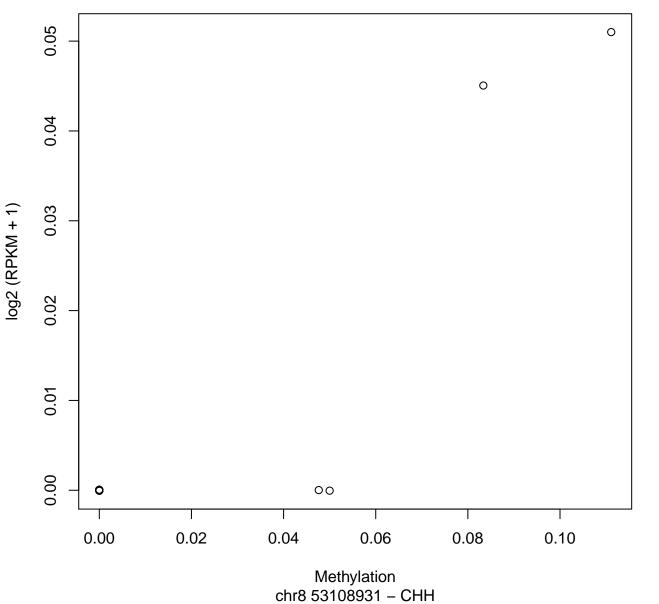
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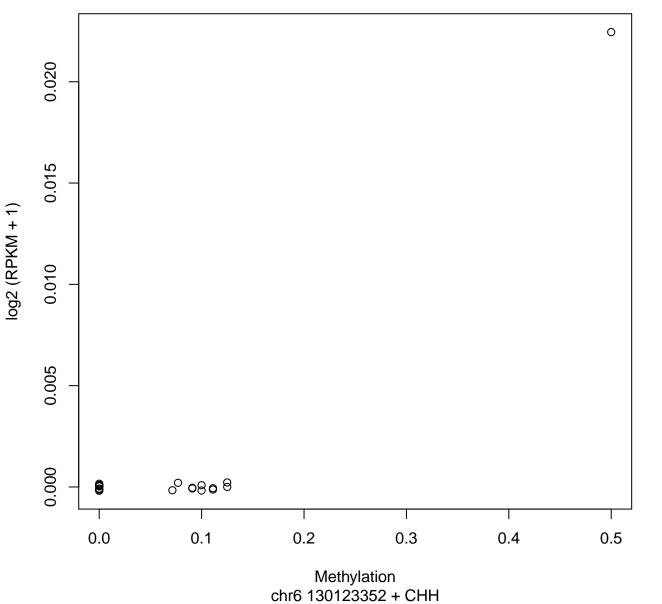
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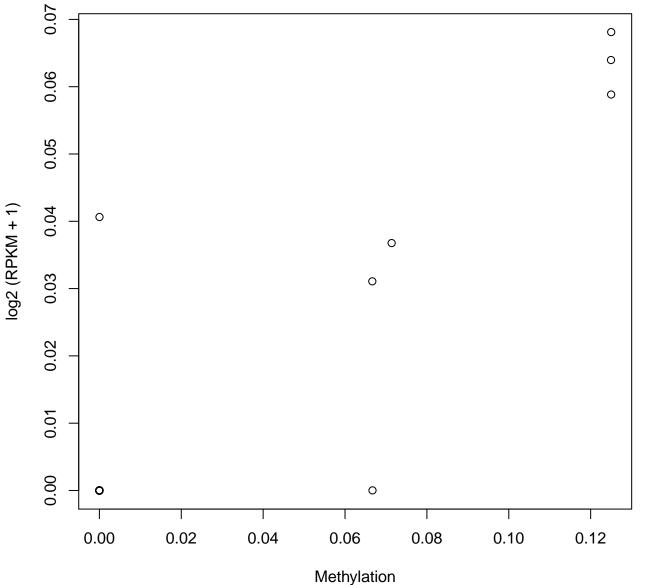
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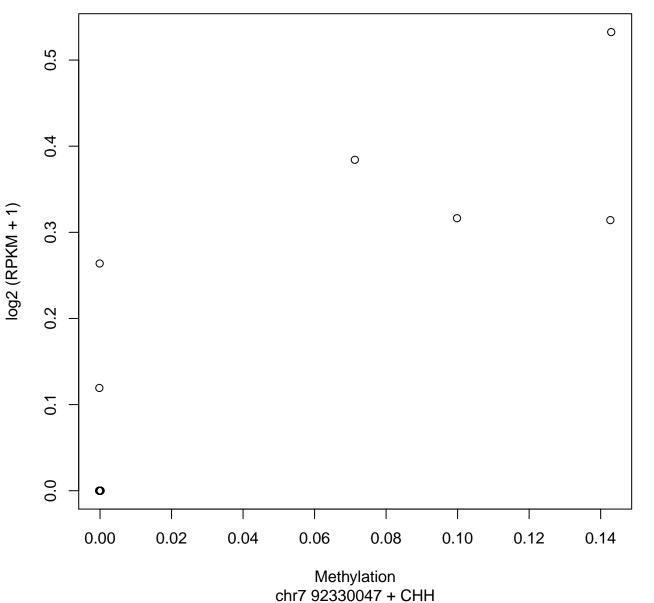


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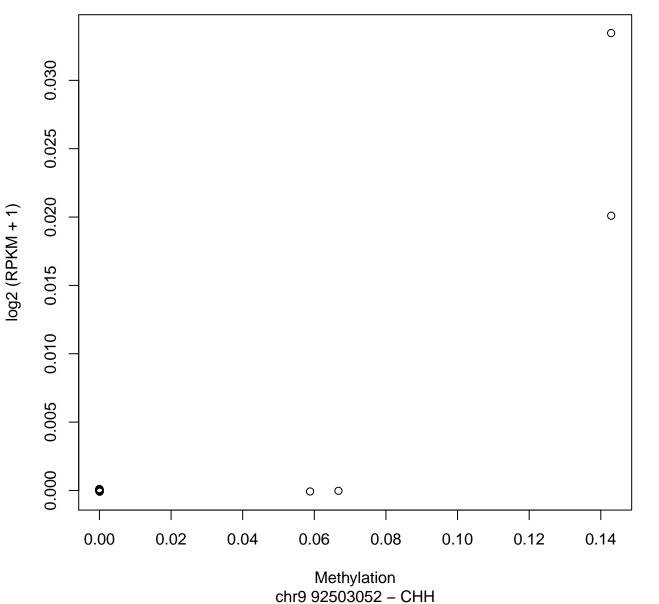


chr15 39311119 - CHH

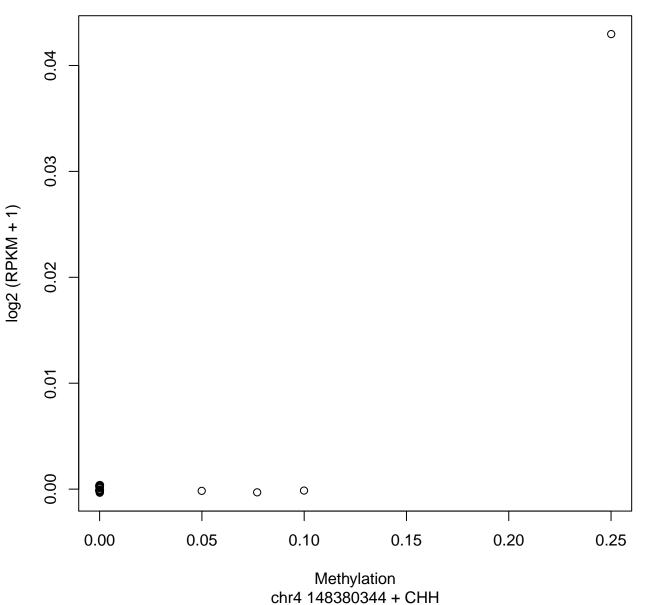
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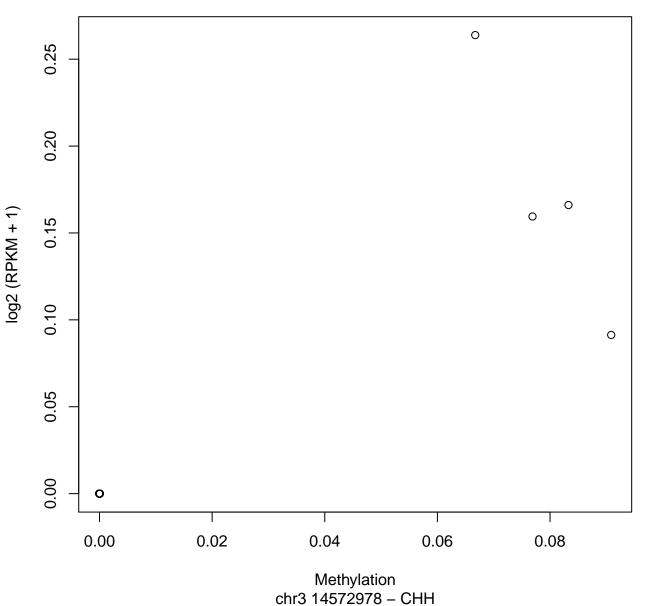
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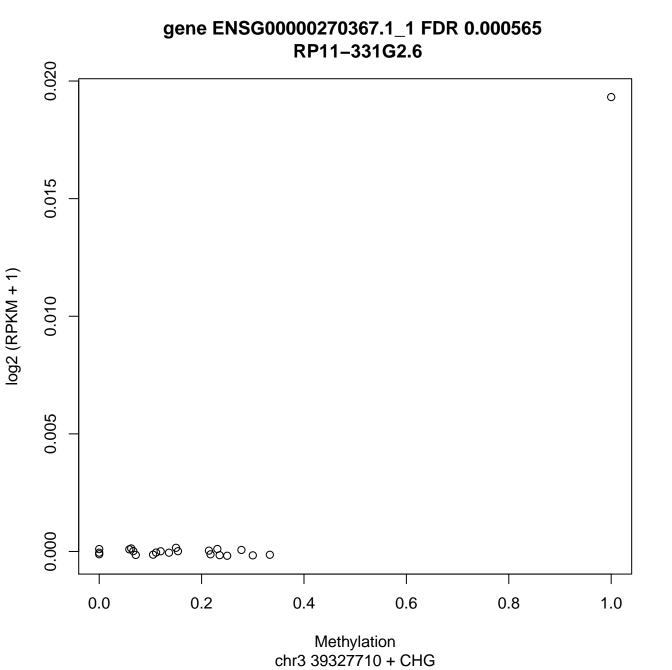


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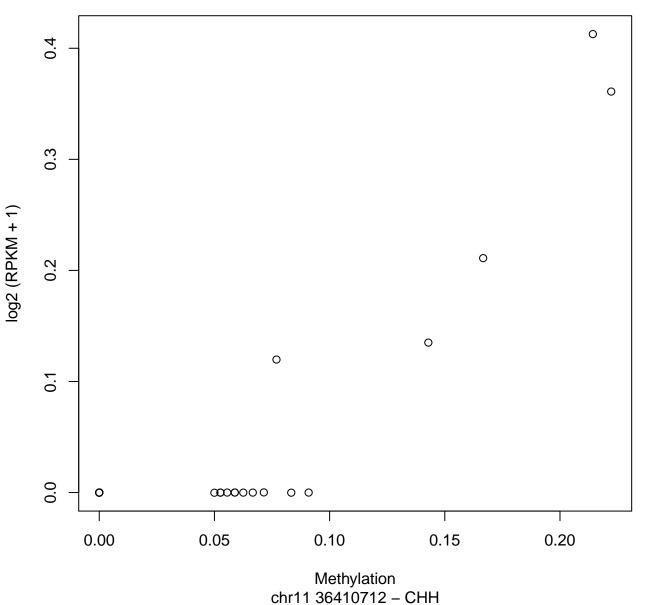


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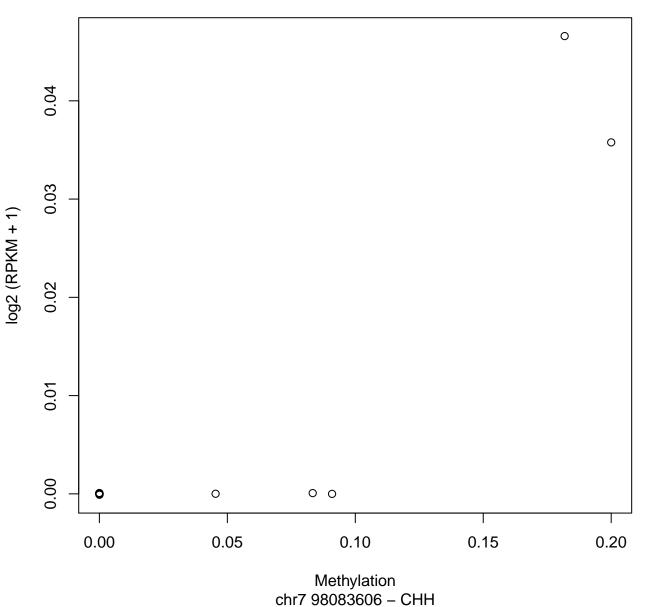




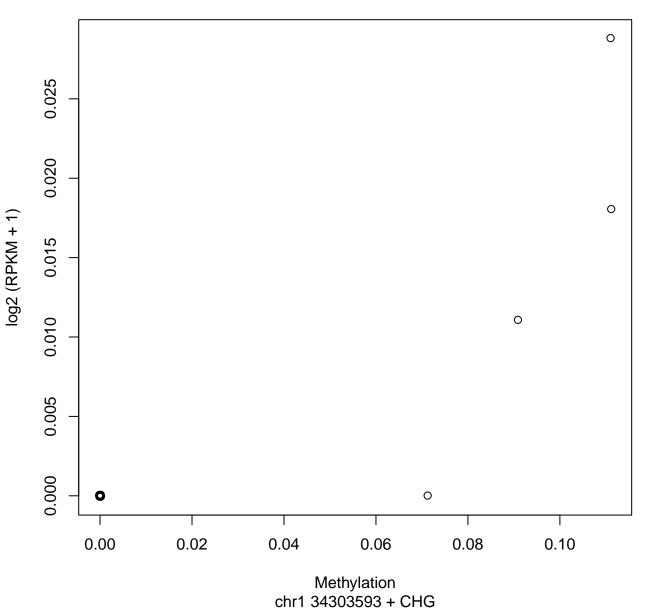
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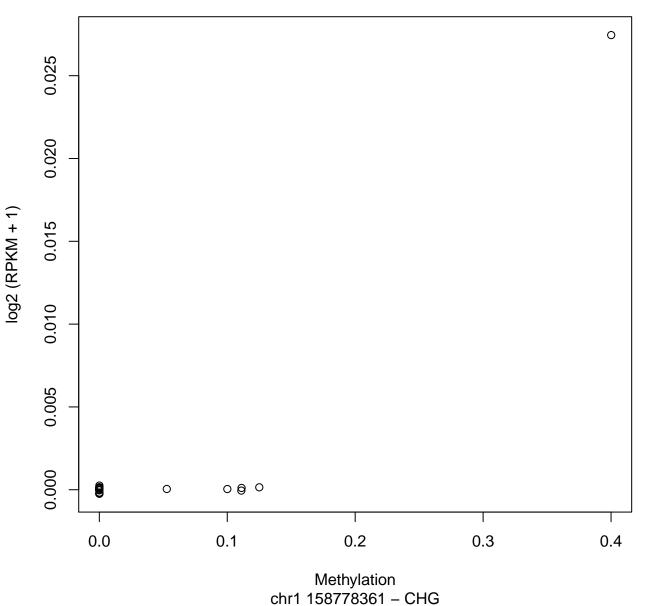
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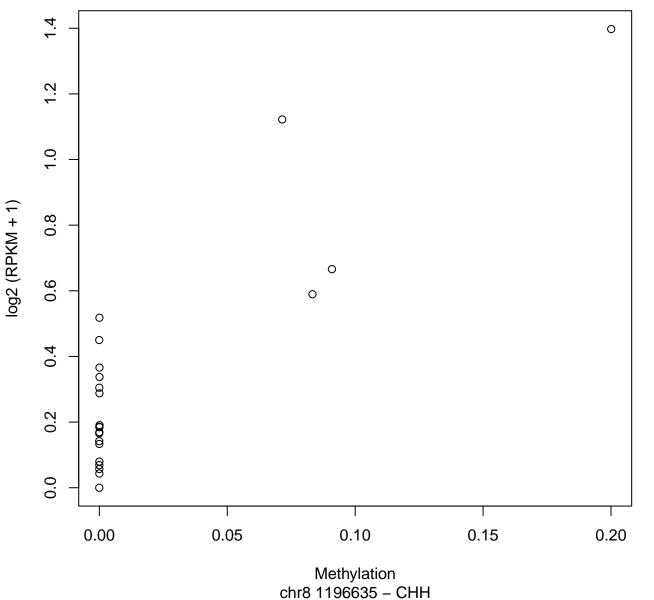
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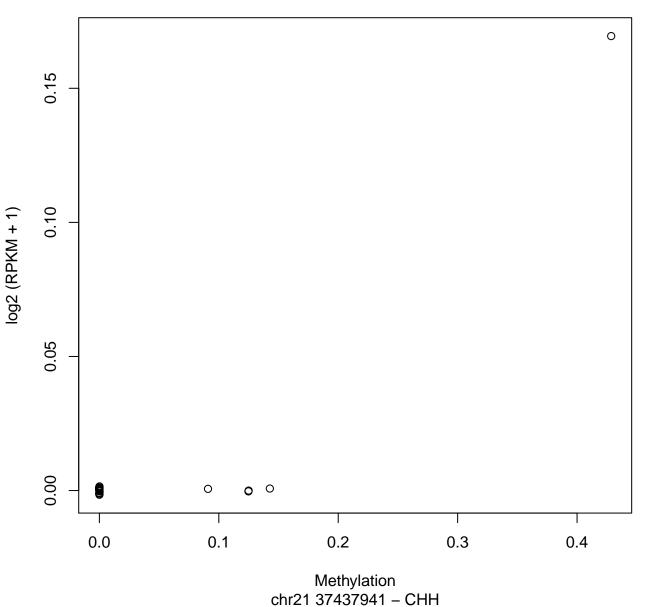
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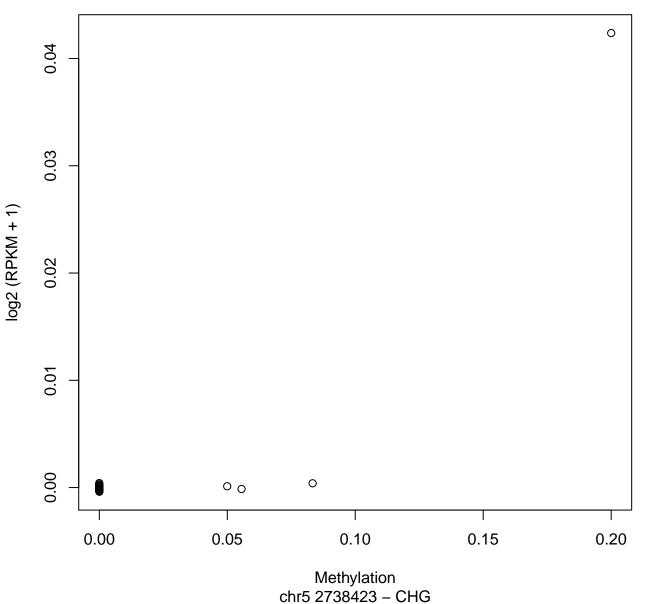
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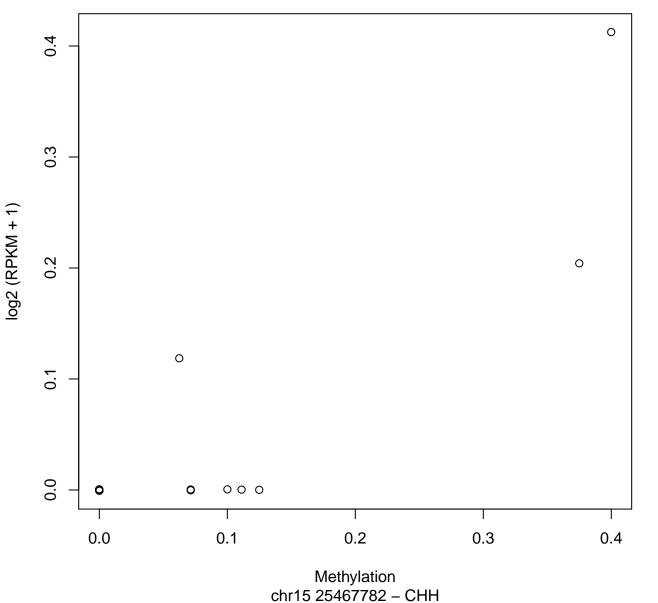
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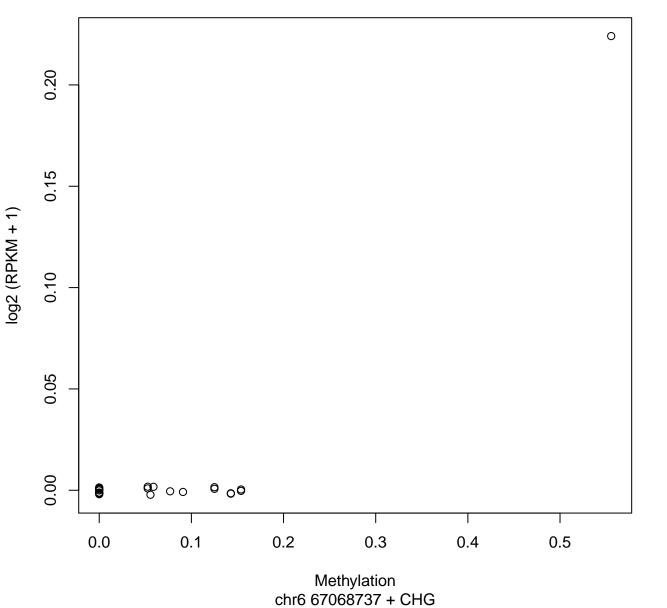
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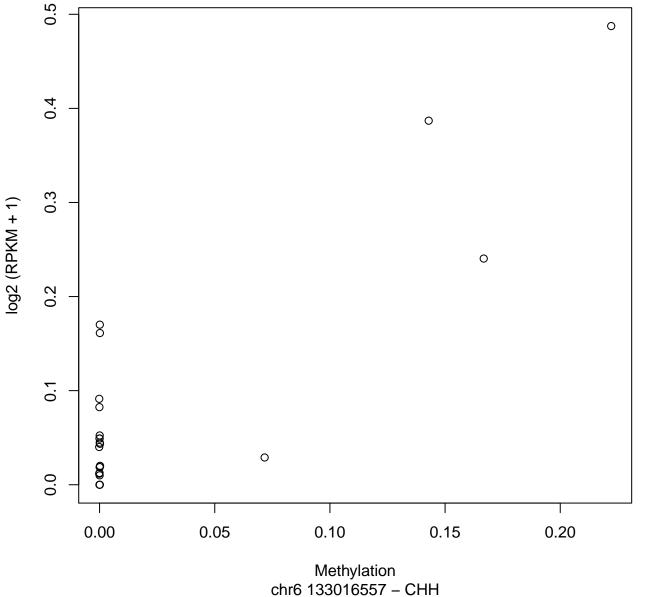
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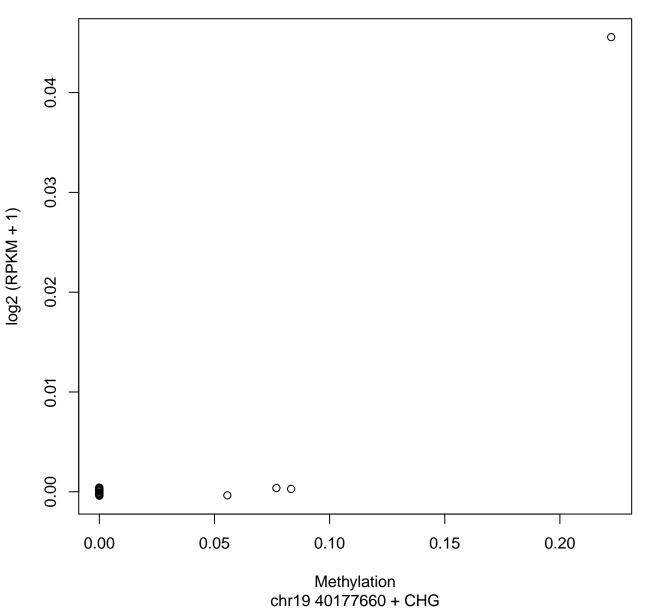
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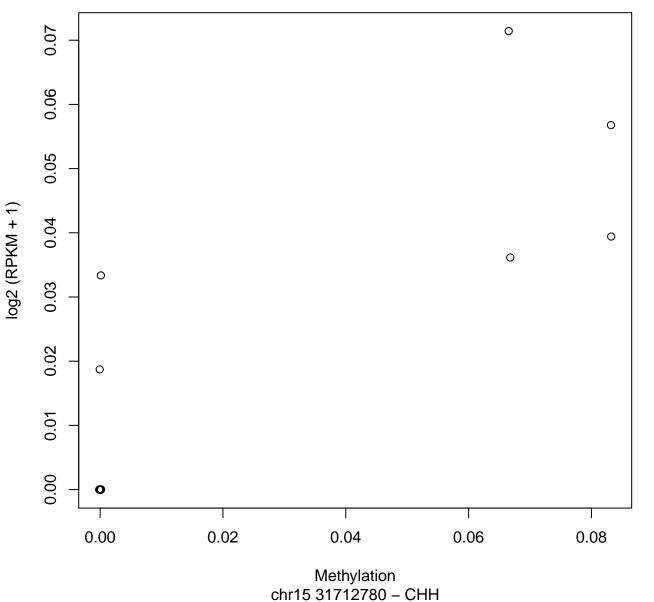
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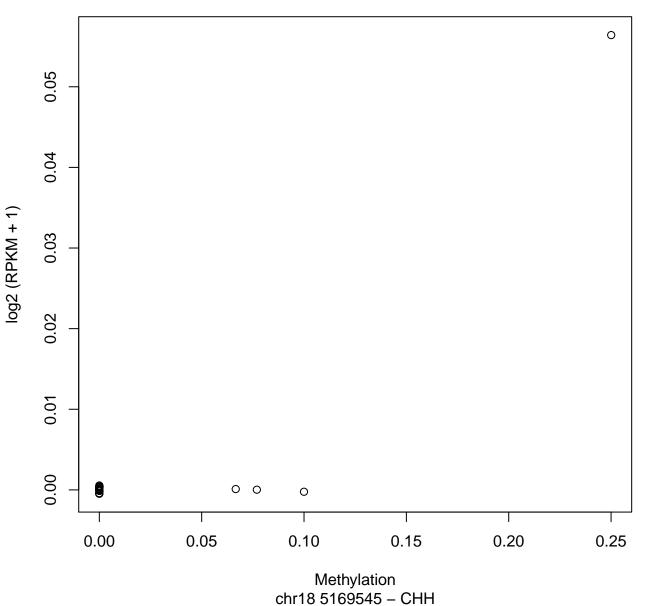
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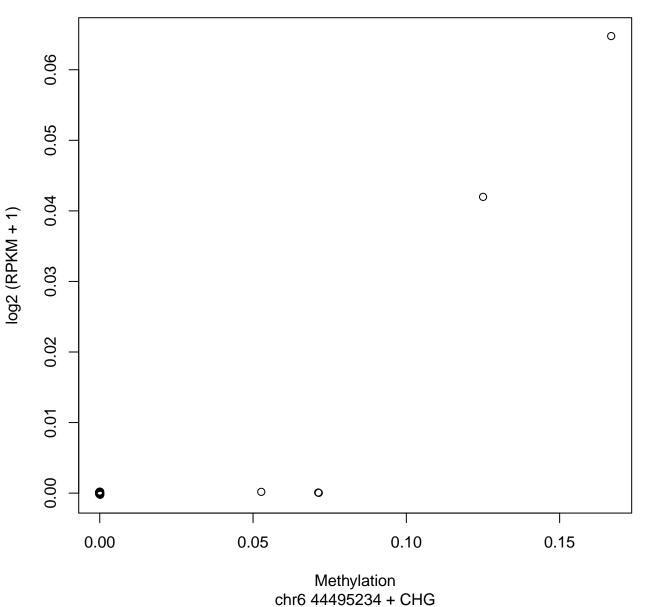
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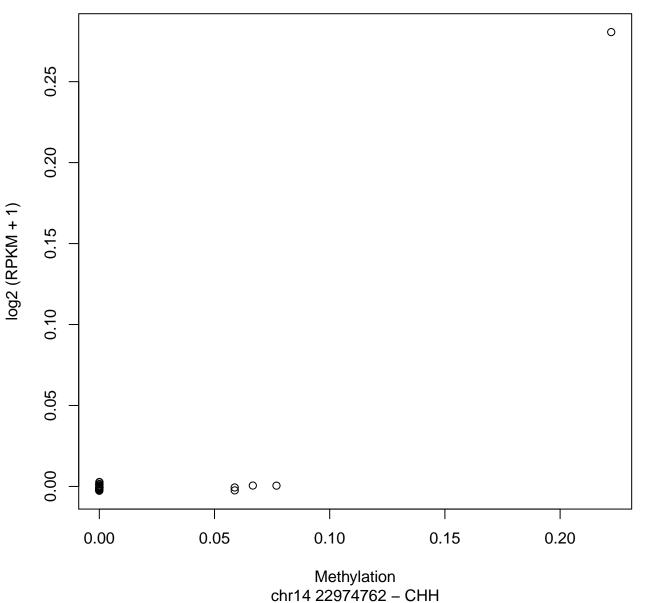
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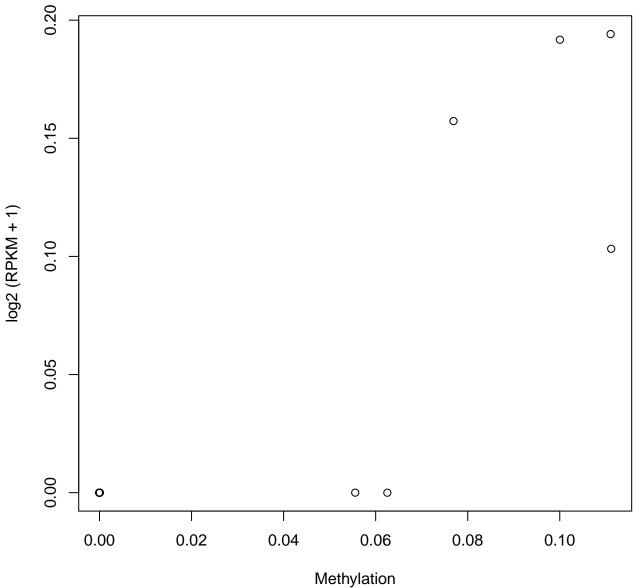
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gene ENSG00000276699.1_2 FDR 0.00114 TRAJ36

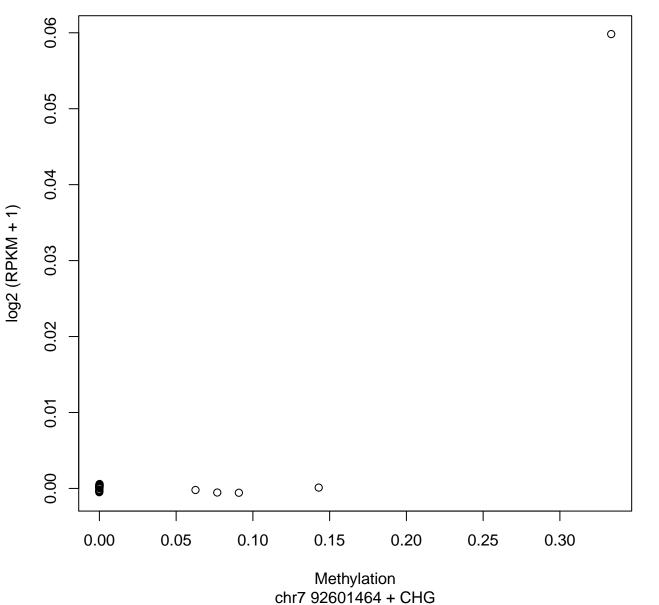


gene ENSG00000200745.1 FDR 0.00116 RNU1-31P

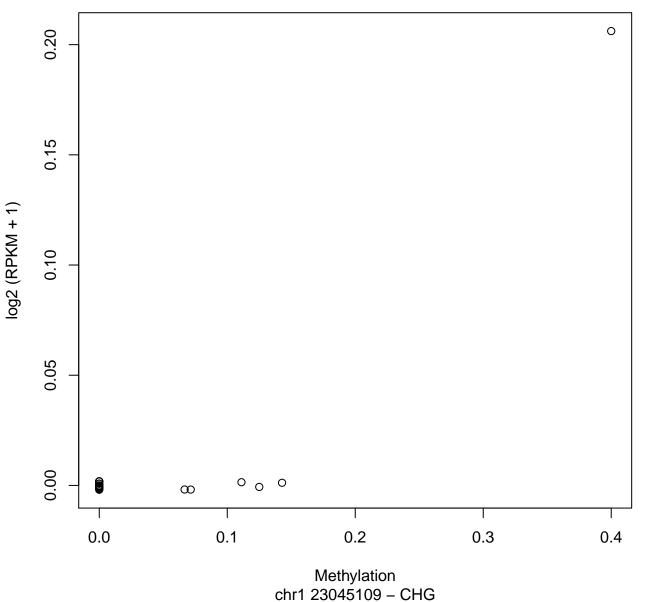


chr2 237118942 + CHG

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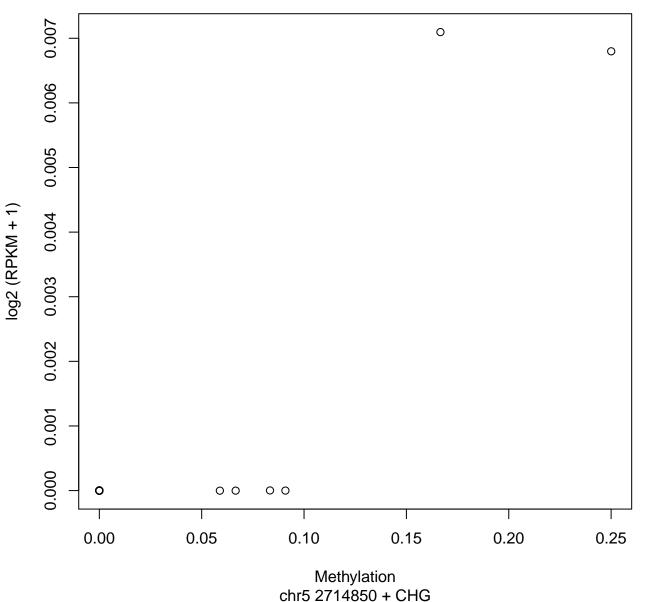
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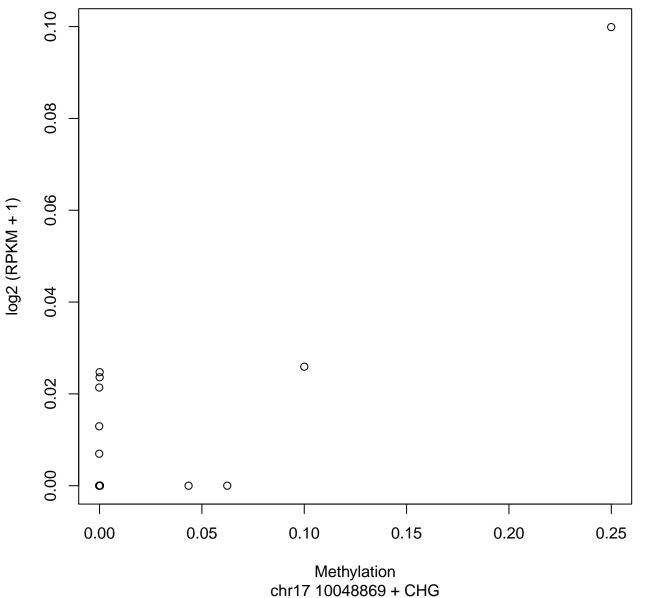
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chr10 73603512 + CHH

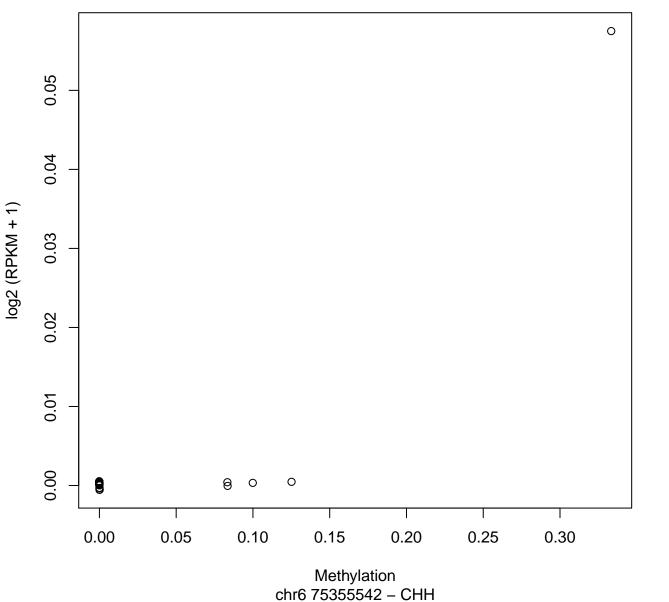
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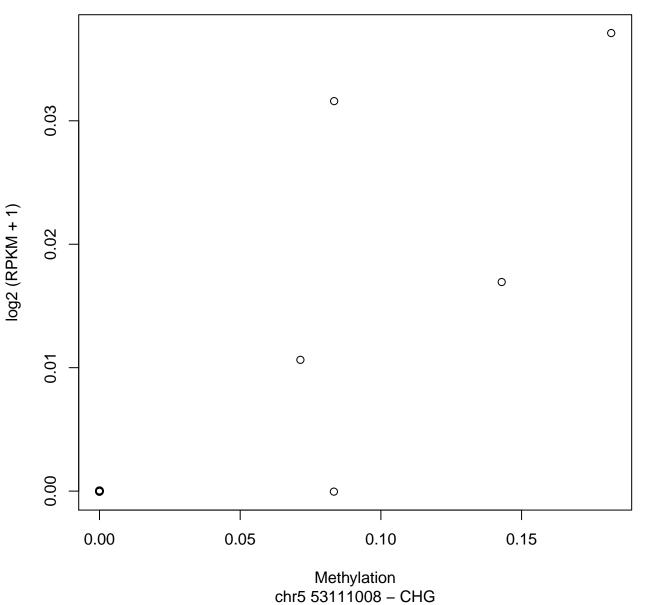
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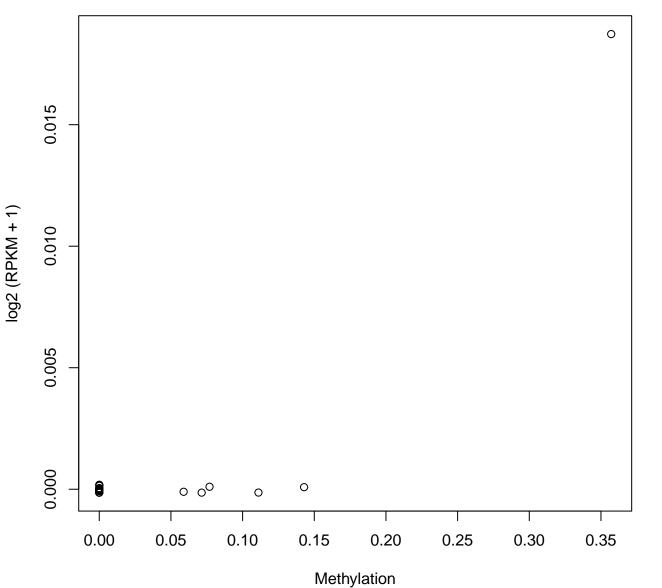
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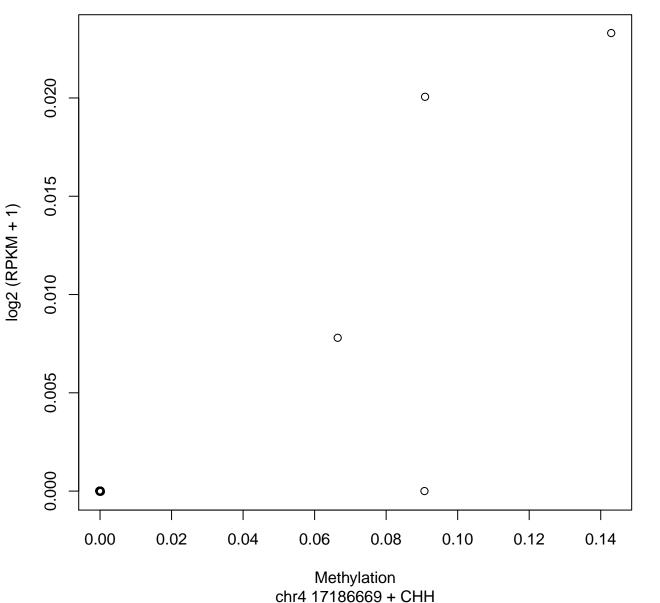


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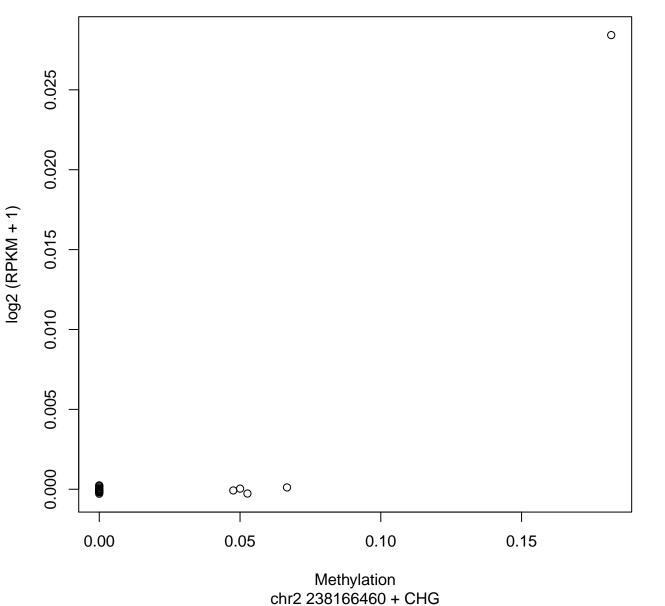


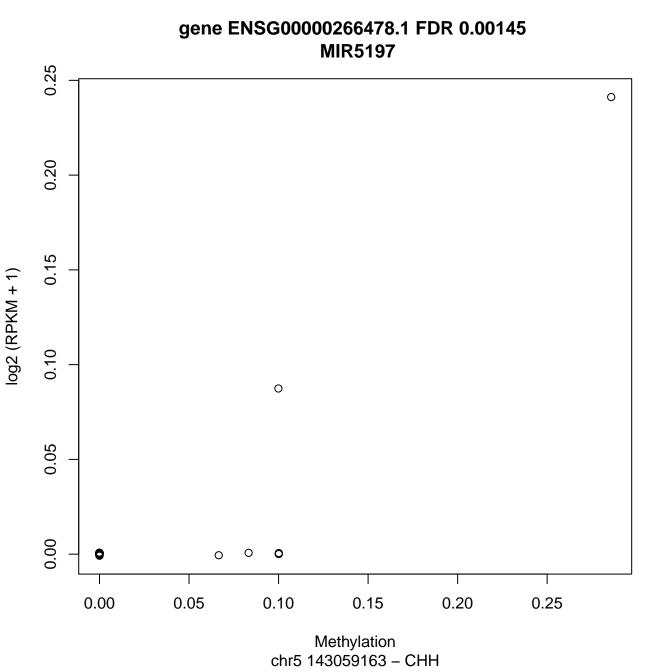
chr8 135869807 + CHG

gene ENSG00000250819.1_1 FDR 0.00141 RP11-576E20.1

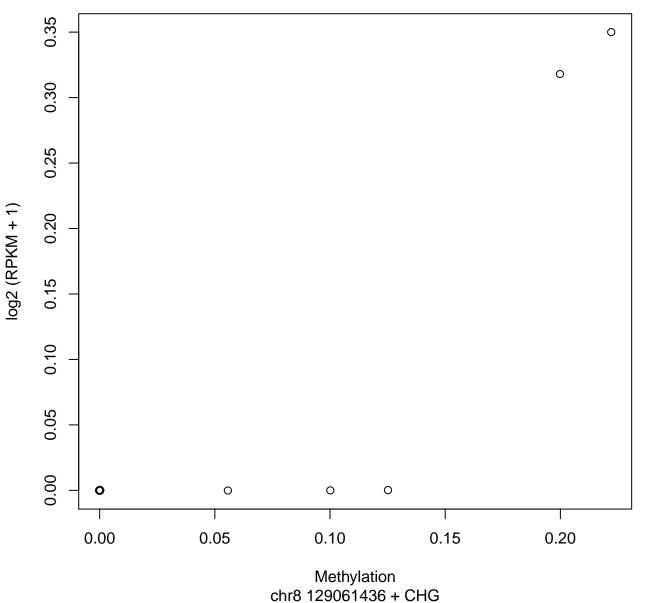


gene ENSG00000224132.2_1 FDR 0.00141 AC112715.2

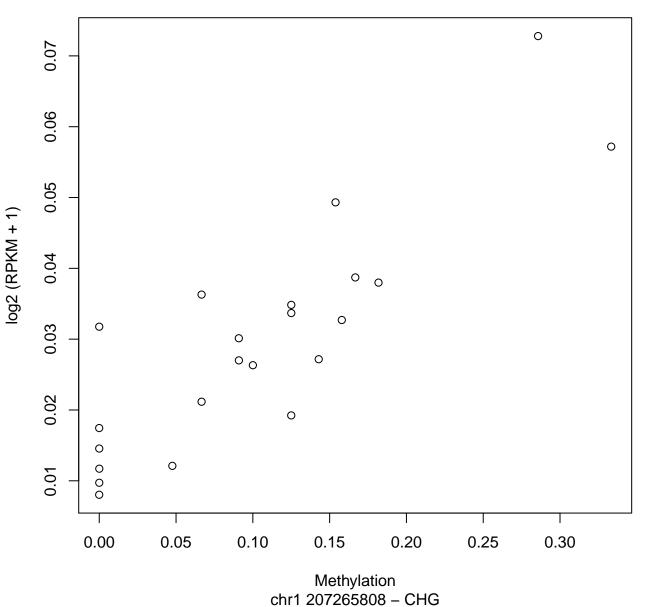




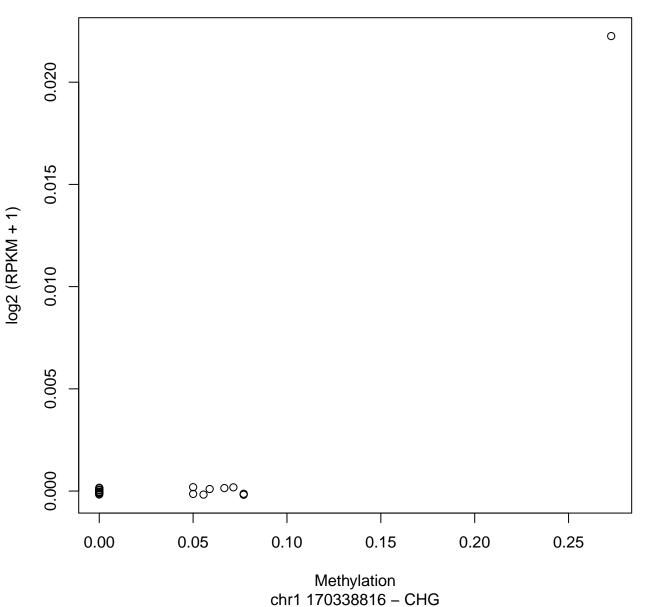
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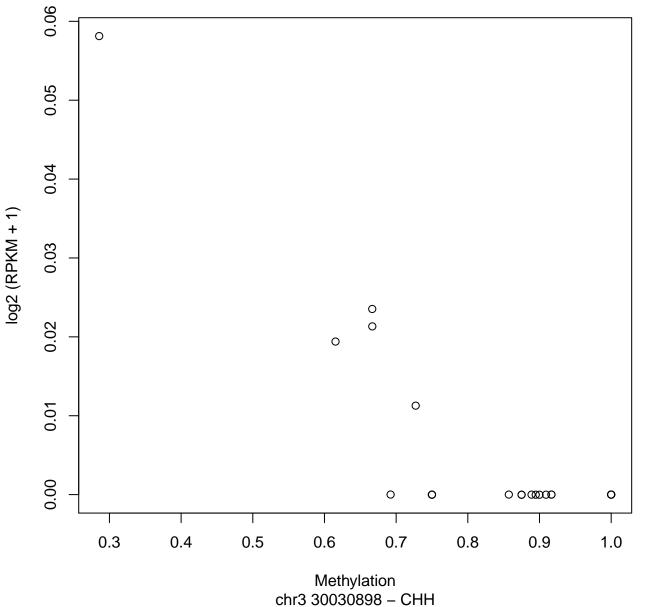
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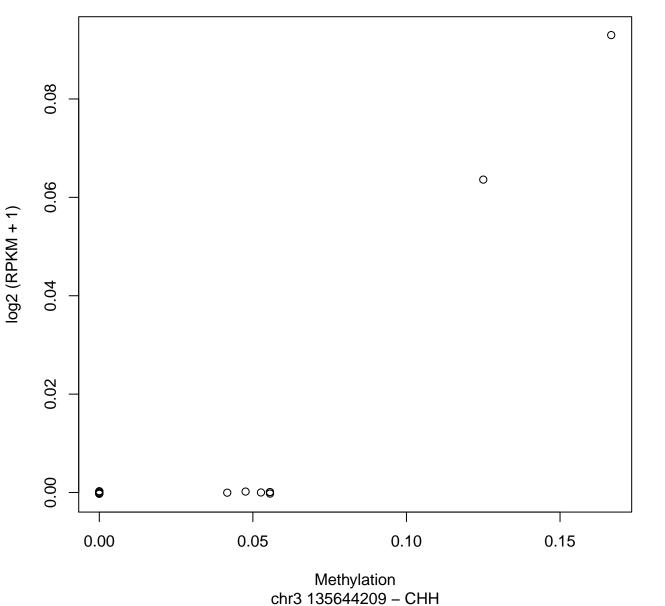
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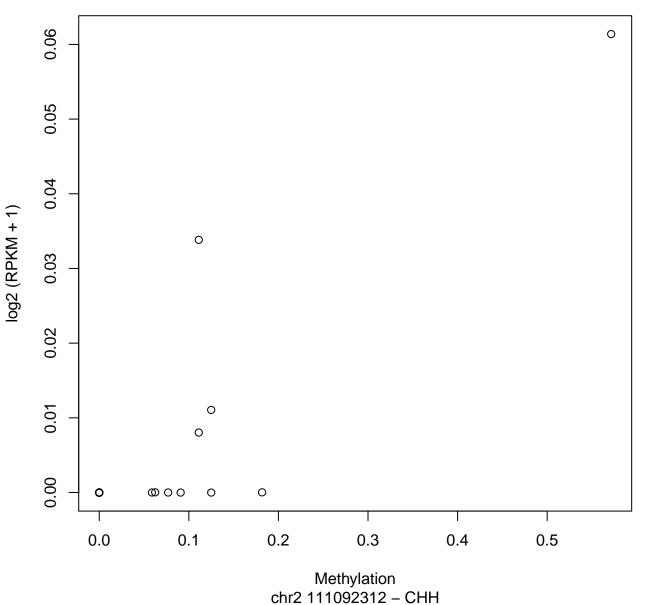
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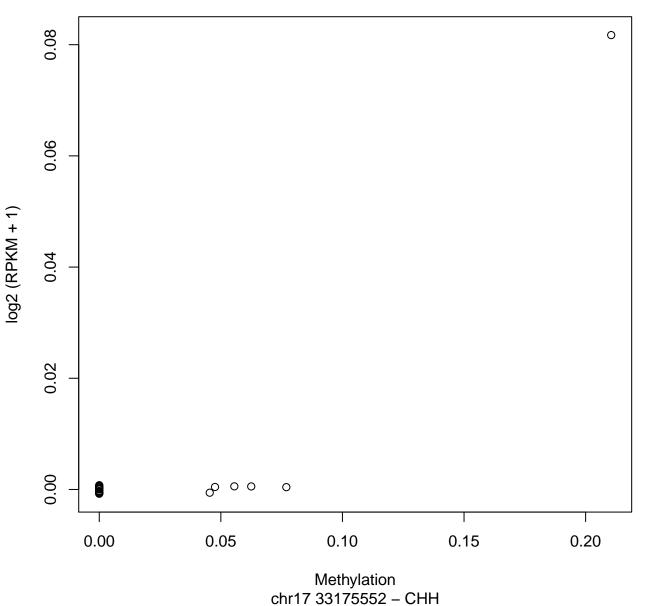
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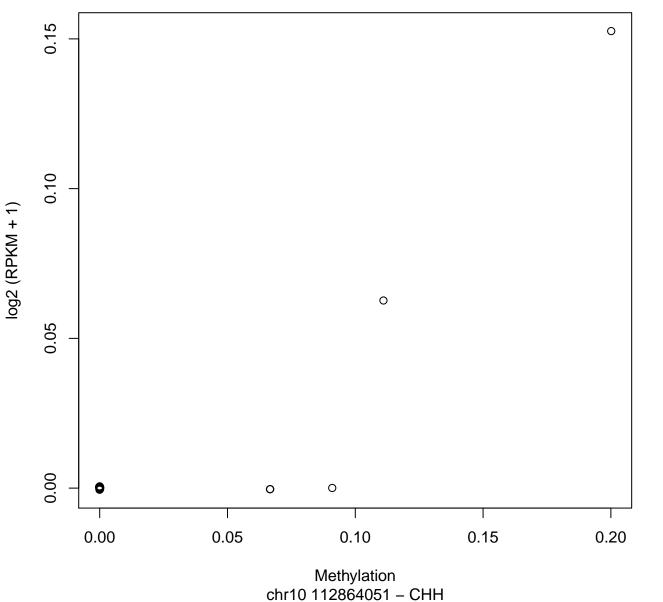
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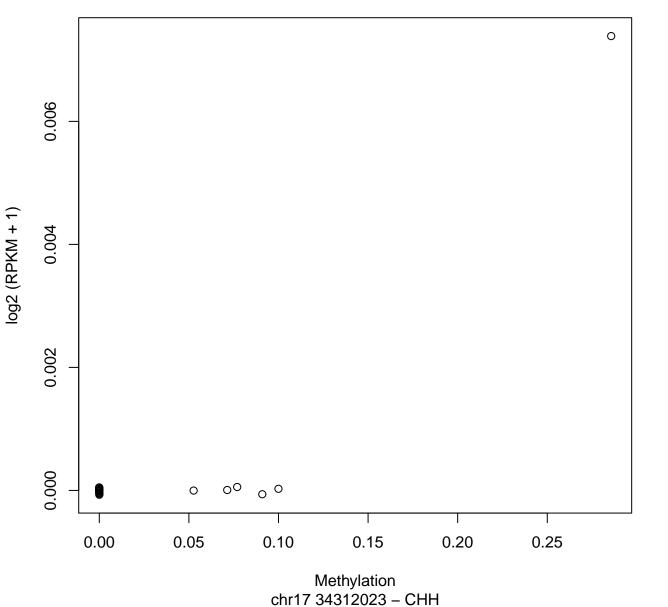
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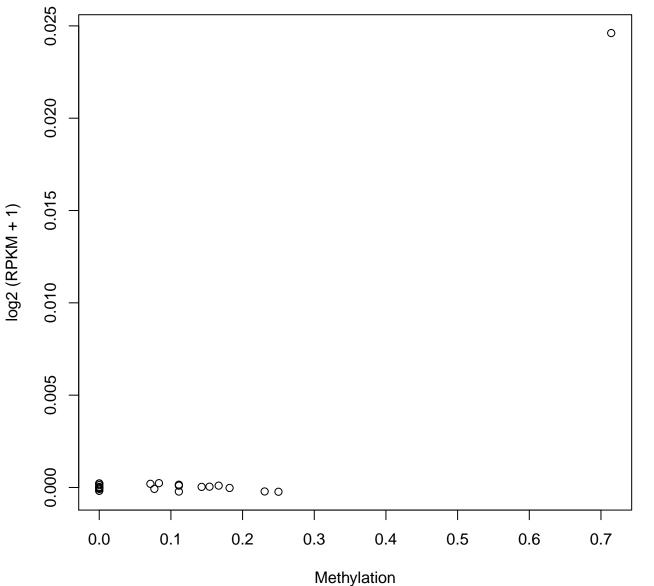
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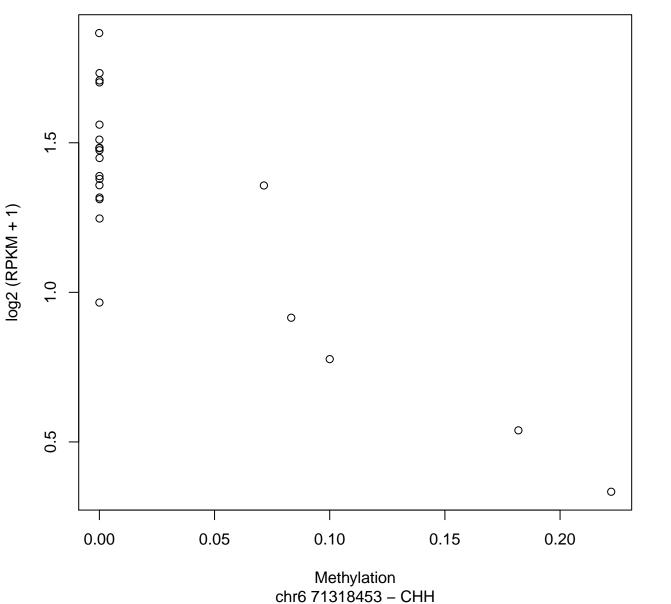


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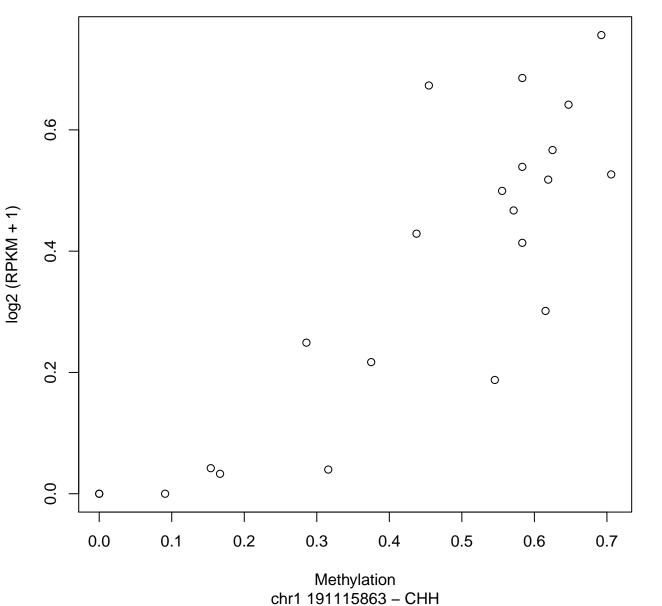


chr1 234844292 + CHH

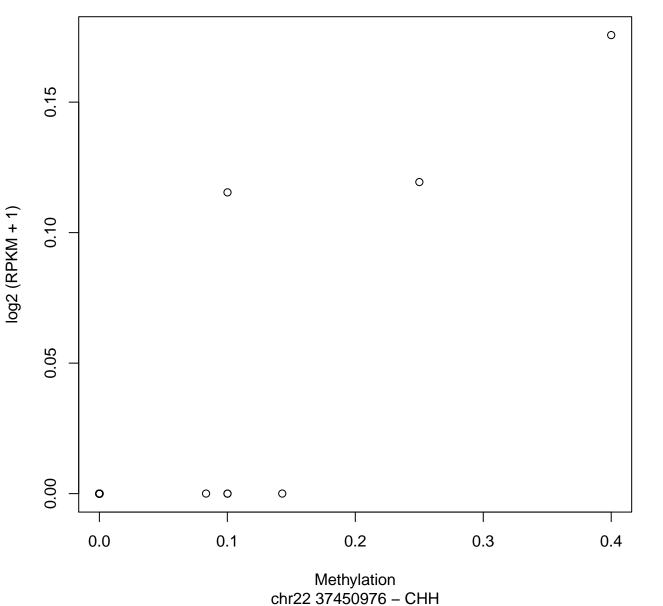
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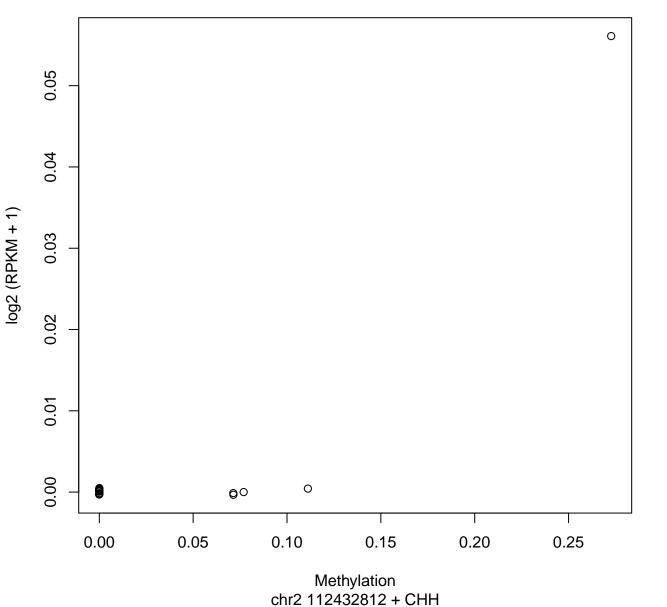
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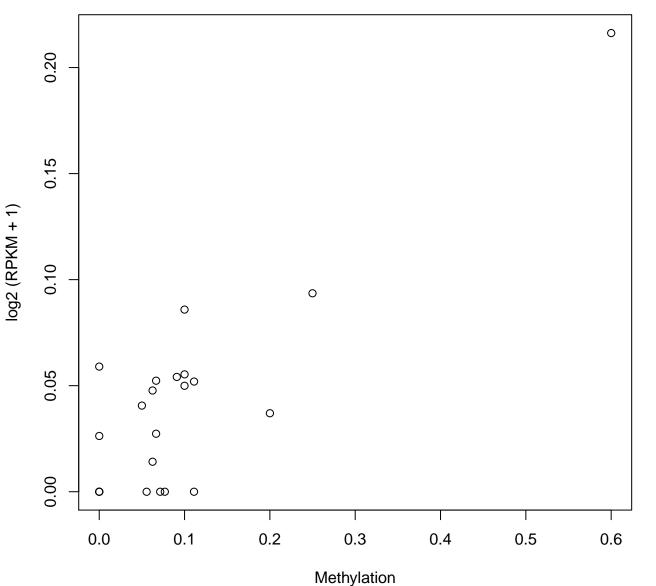
gene ENSG00000201078.1 FDR 0.00197 RN7SKP214



gene ENSG00000237583.1_1 FDR 0.00197 AC093166.1



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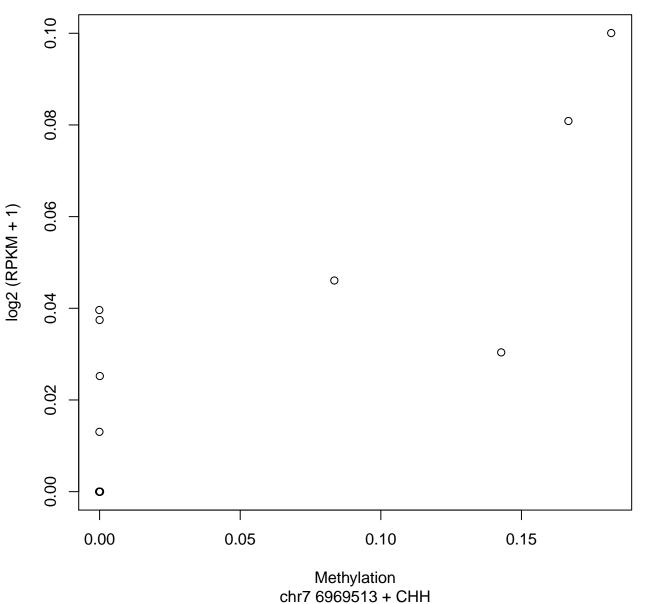


chr18 13138923 + CHH

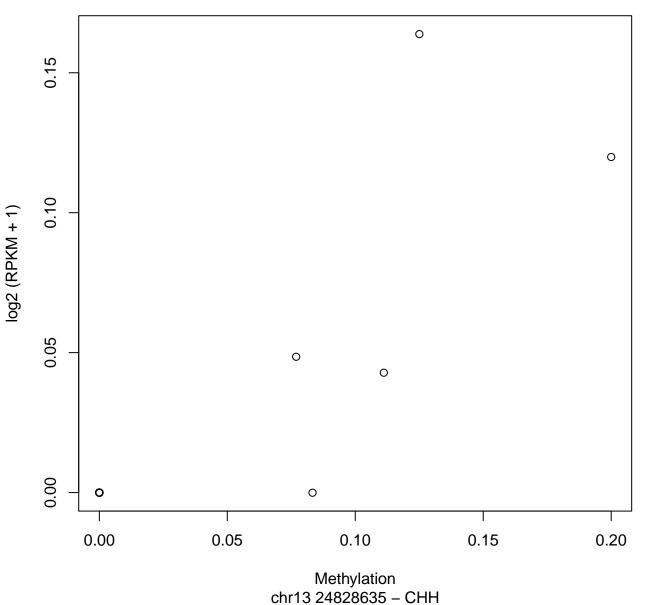
gene ENSG00000268036.1_1 FDR 0.00201 AC007228.8 0.20 0 0 log2 (RPKM + 1) 0.10 0.05 0.00 0 0 0.00 0.05 0.10 0.15

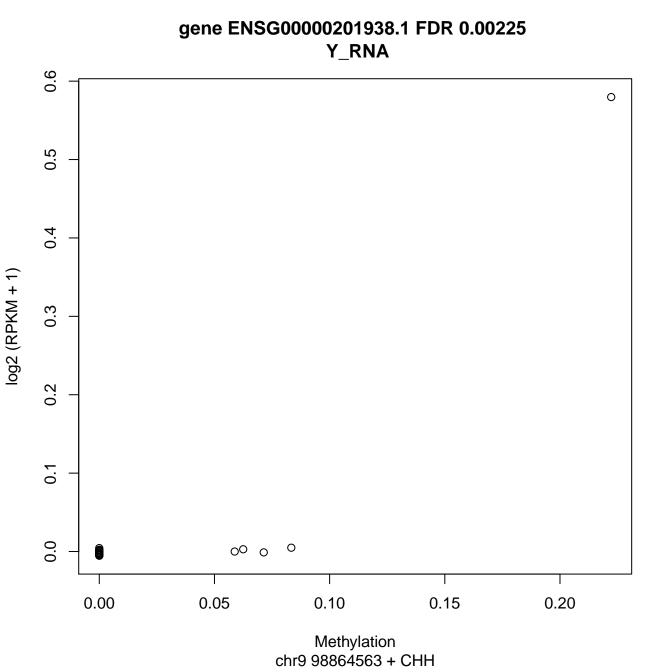
Methylation chr19 57124259 – CHG

gene ENSG00000226943.3_1 FDR 0.0022 ALG1L5P

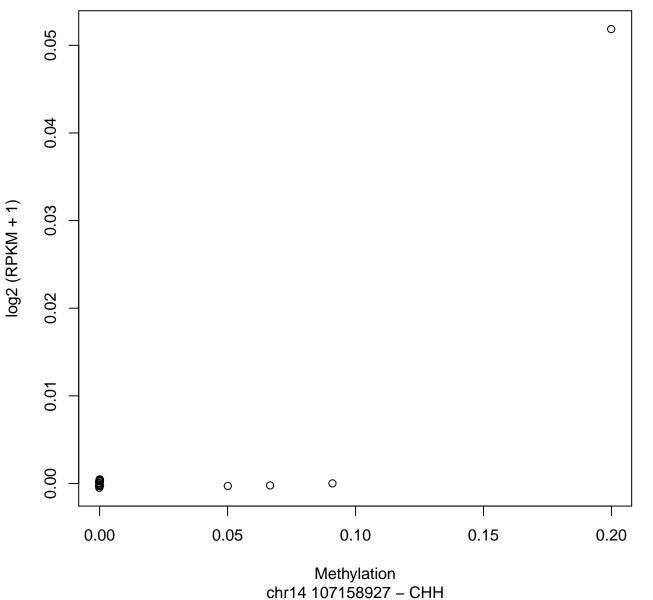


gene ENSG00000227213.1_1 FDR 0.00225 SPATA13-AS1

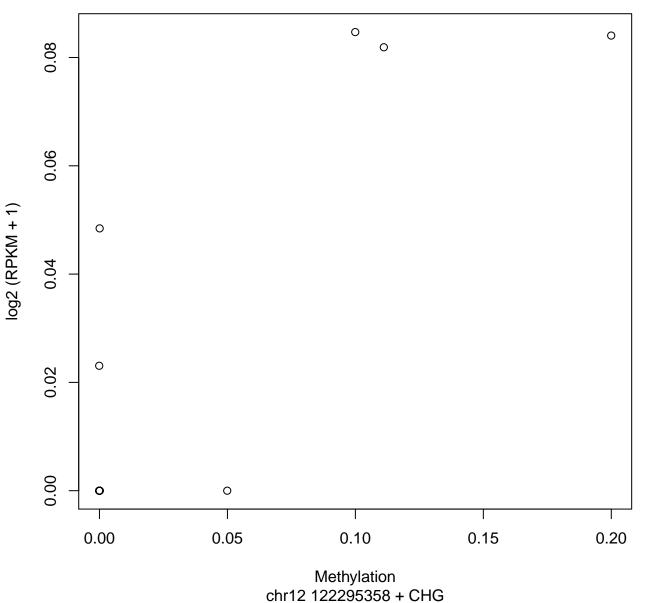




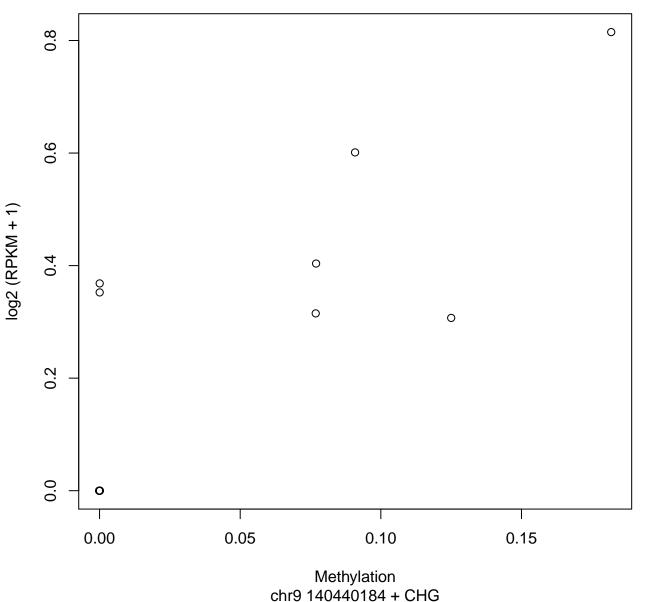
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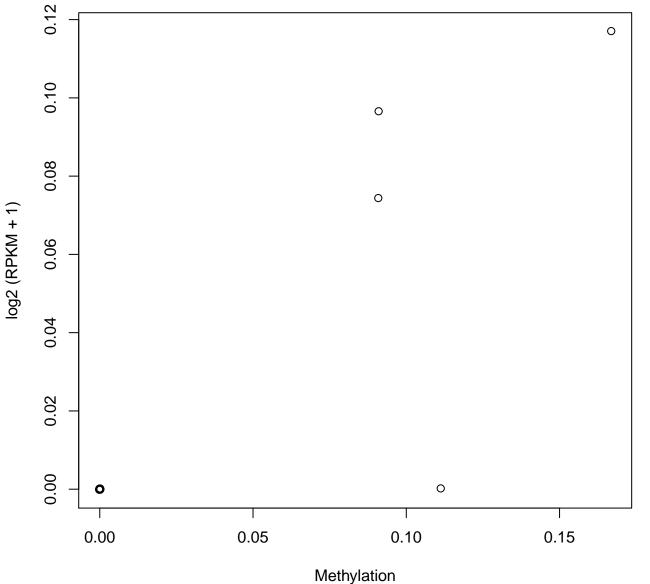
gene ENSG00000256811.1_1 FDR 0.00235 RP11-7M8.2



gene ENSG00000263933.1 FDR 0.00239 AL365502.1

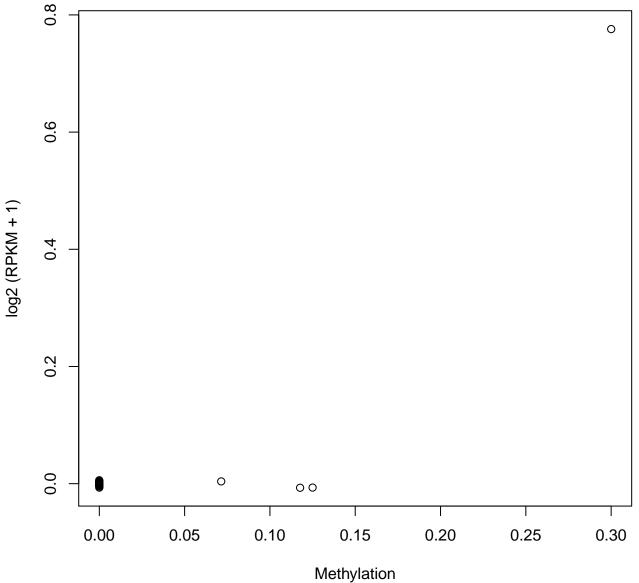


gene ENSG00000248803.1_1 FDR 0.00241 RP11-287J9.1



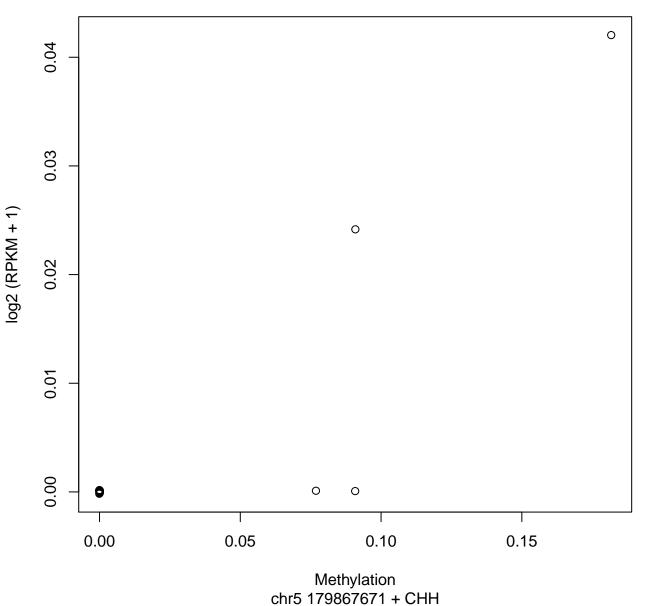
chr5 65918555 + CHH

gene ENSG00000228985.1_2 FDR 0.00247 TRDD3

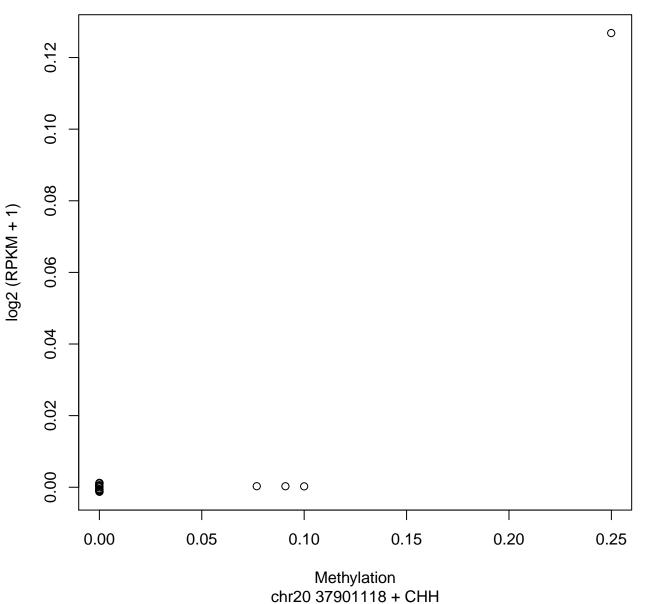


chr14 22918255 + CHG

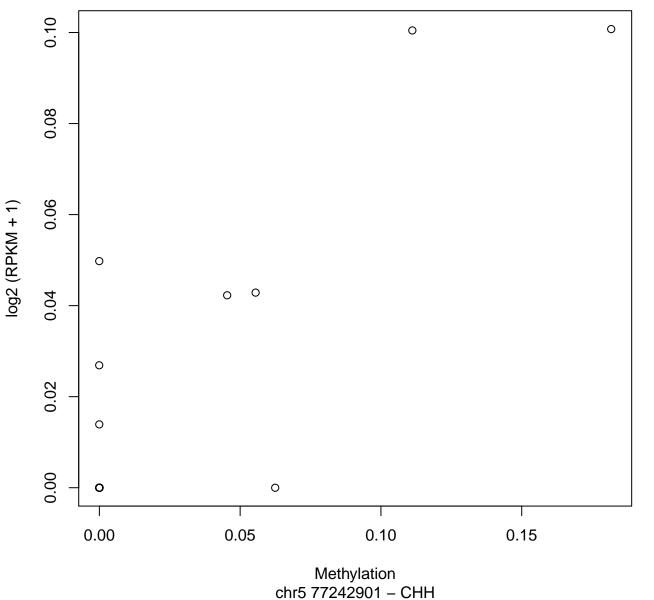
gene ENSG00000250509.1_1 FDR 0.00249 CTC-573N18.1



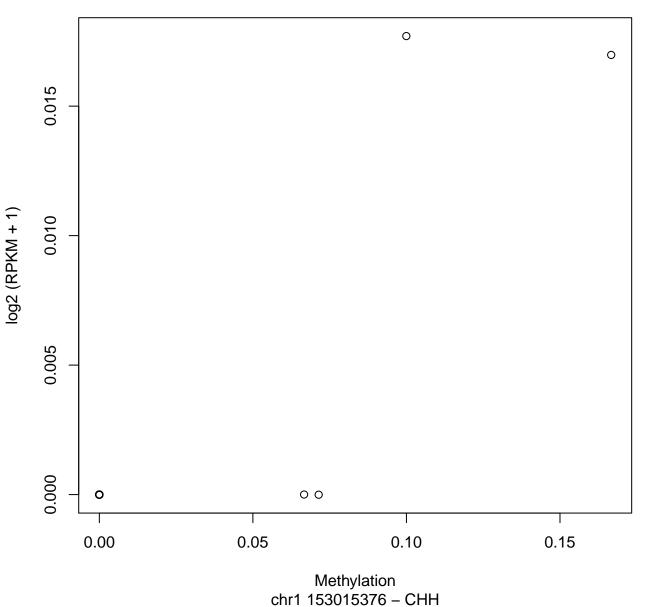
gene ENSG00000211534.2 FDR 0.00252 AL121588.1



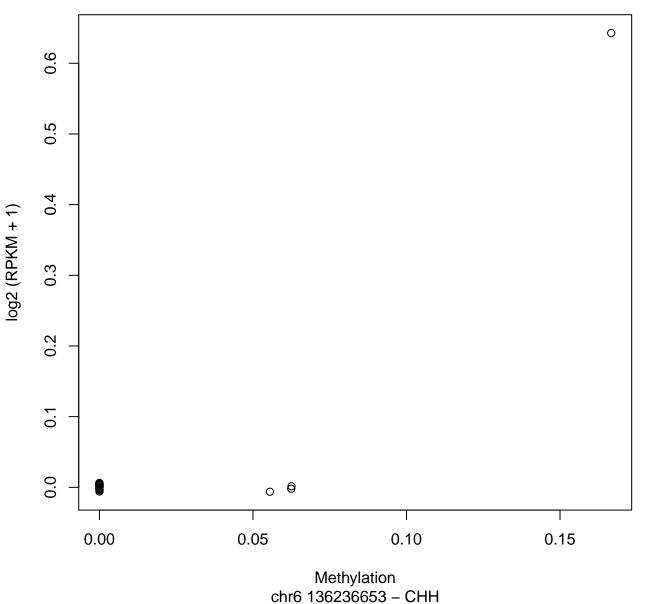
gene ENSG00000253572.2_1 FDR 0.00259 RP11-107N7.1



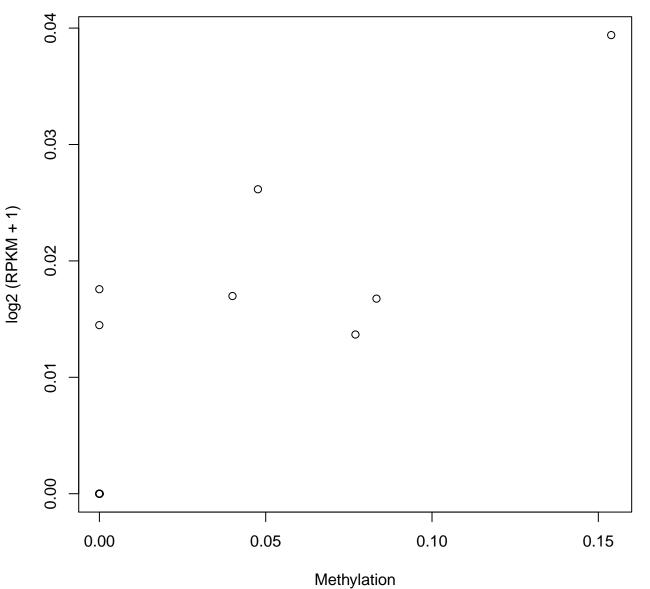
gene ENSG00000163216.6_1 FDR 0.00264 SPRR2D



gene ENSG00000238921.2 FDR 0.0027 AL360178.1



gene ENSG00000260737.5_1 FDR 0.00275 LINC01227



chr16 80607366 + CHH