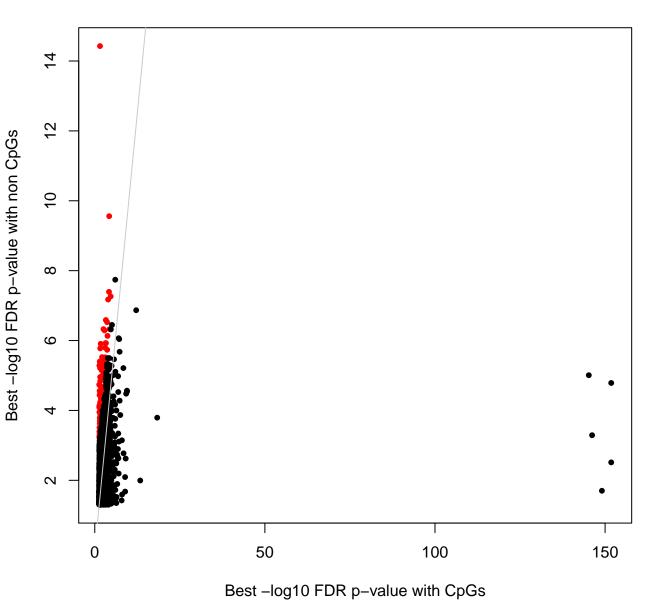
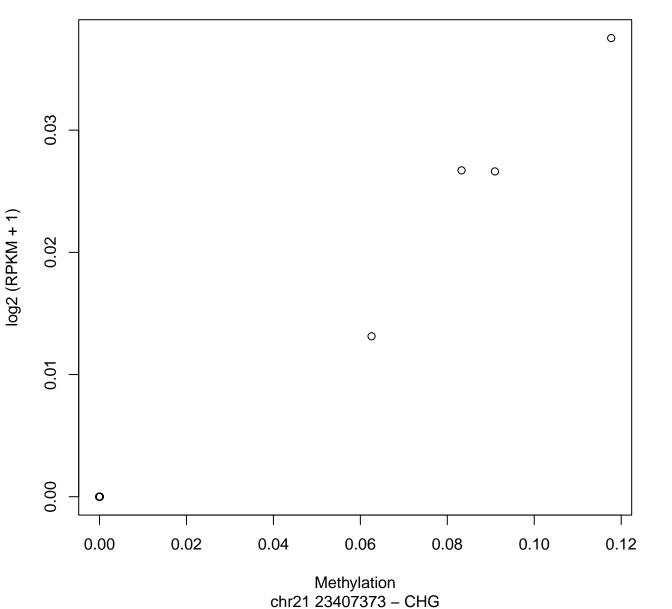
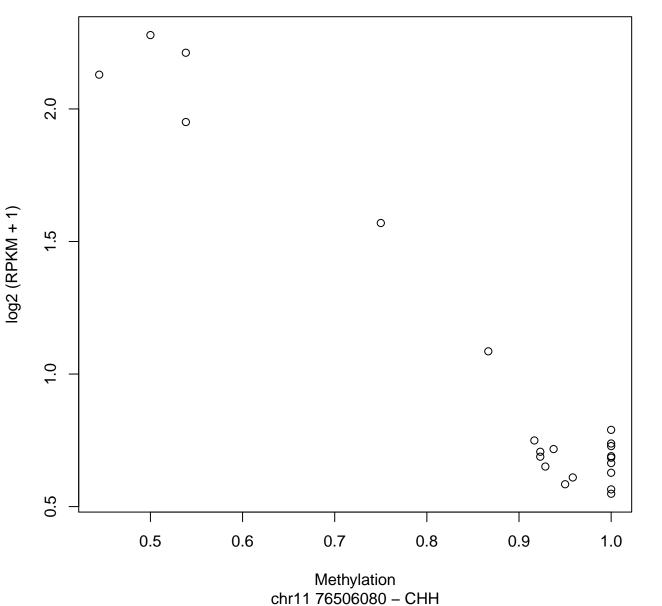
## 3811 common gene meQTLs



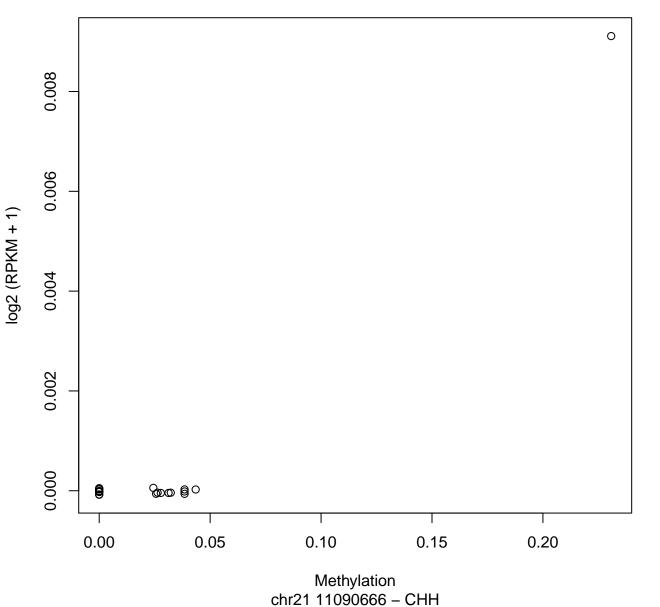
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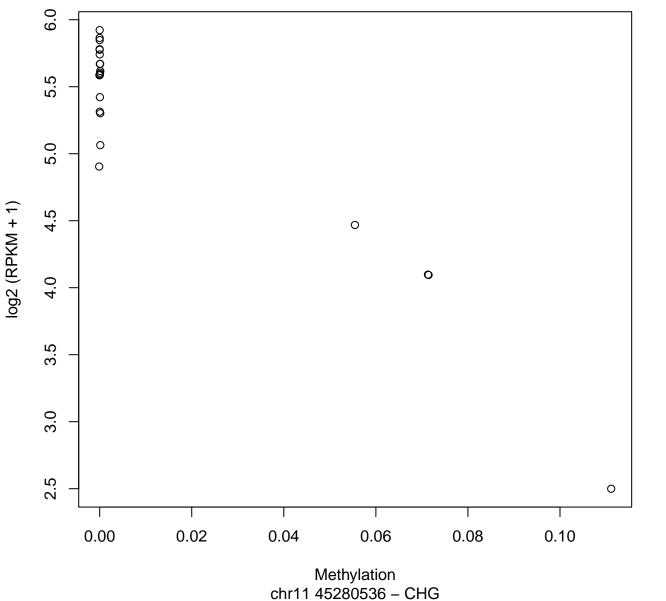
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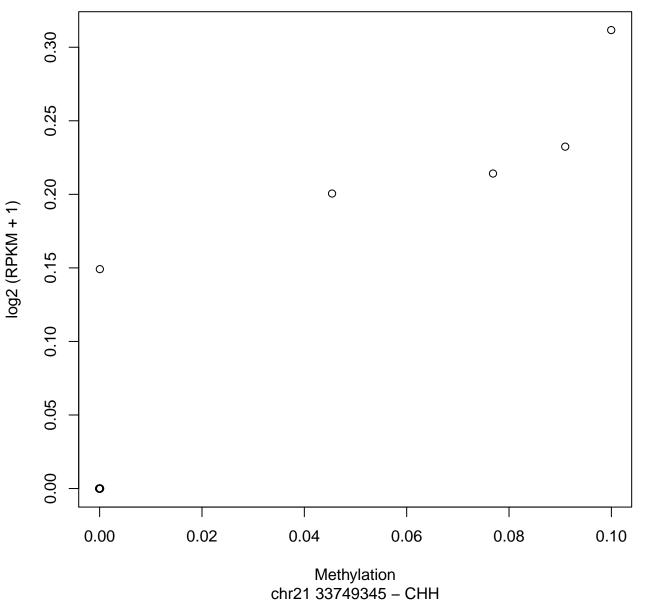
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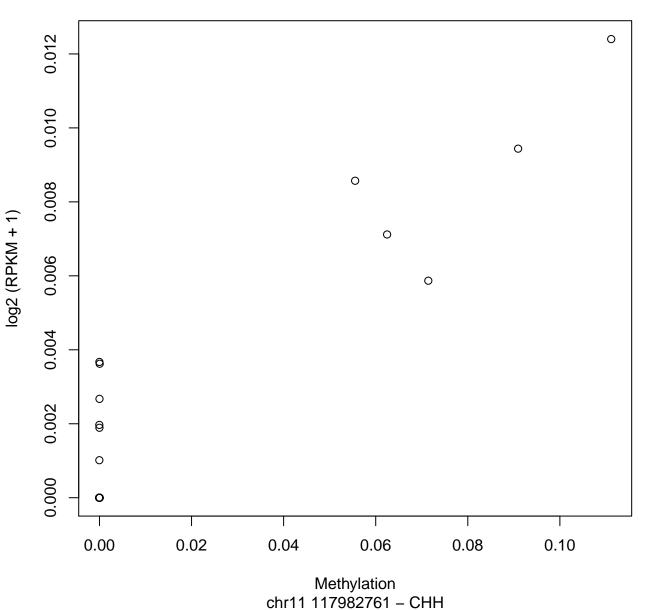
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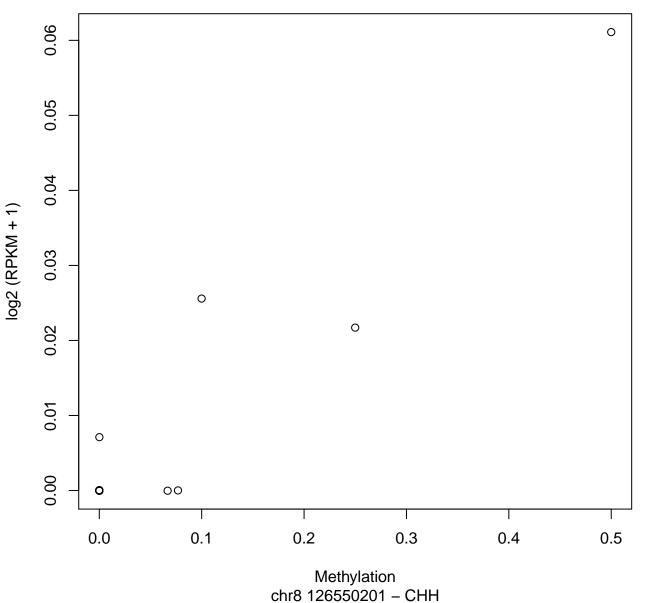
# gene ENSG00000200792.1 FDR 4e-06 SNORA80



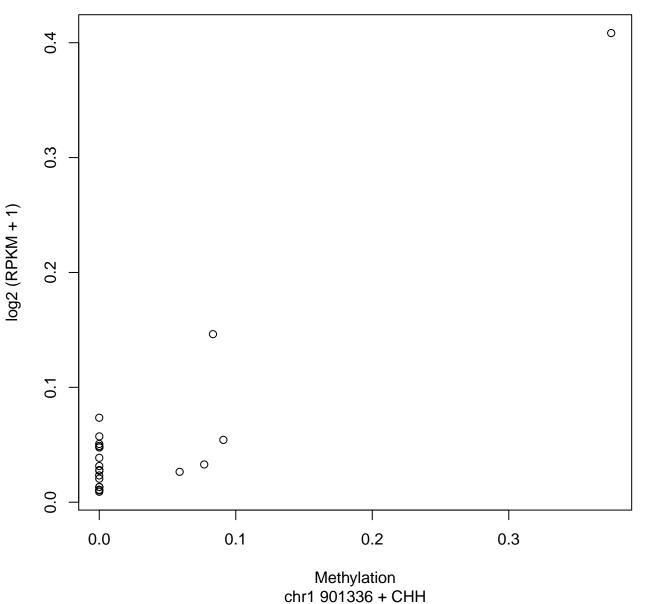
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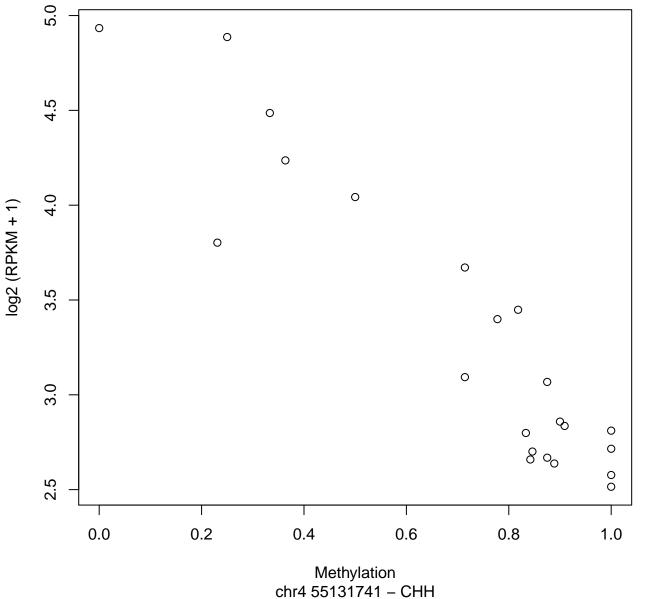
## gene ENSG00000253111.1\_2 FDR 4.71e-07 RP11-136O12.2



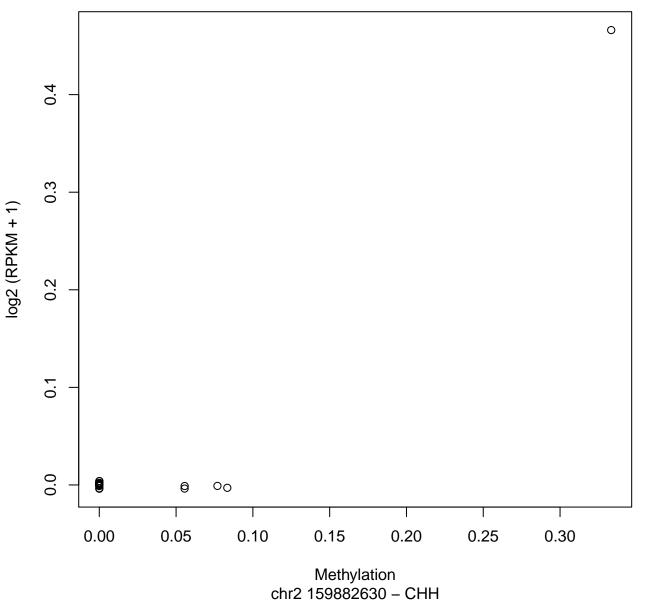
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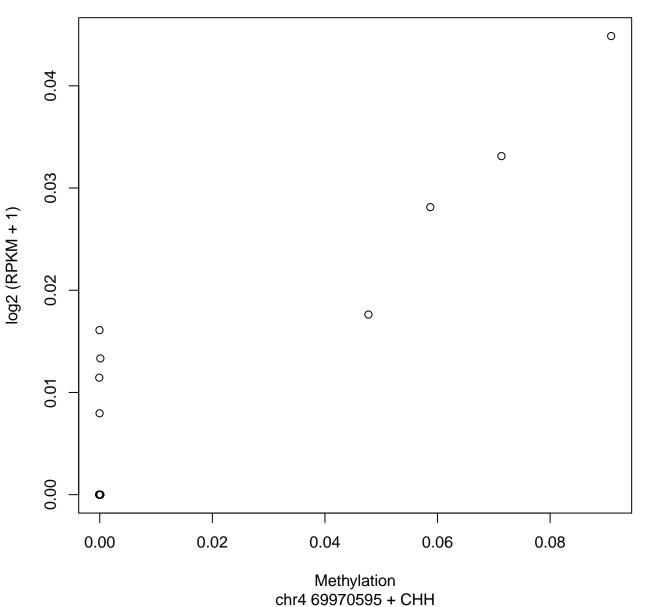
# gene ENSG00000134853.11\_2 FDR 3.99e-06 PDGFRA



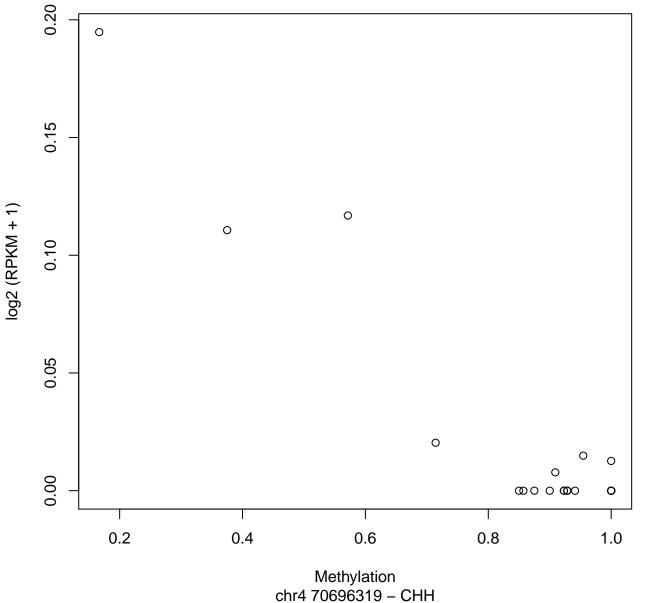
# gene ENSG00000202029.1 FDR 6.13e-06 RNU6-580P



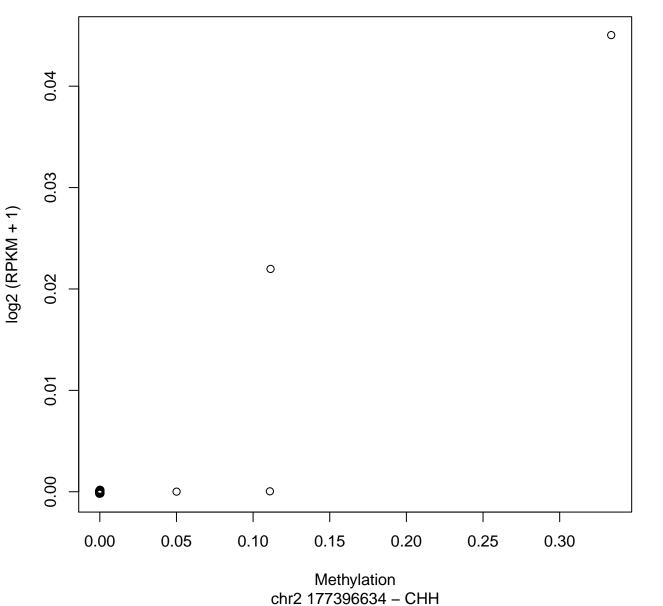
## gene ENSG00000171234.13\_2 FDR 1.83e-05 UGT2B7



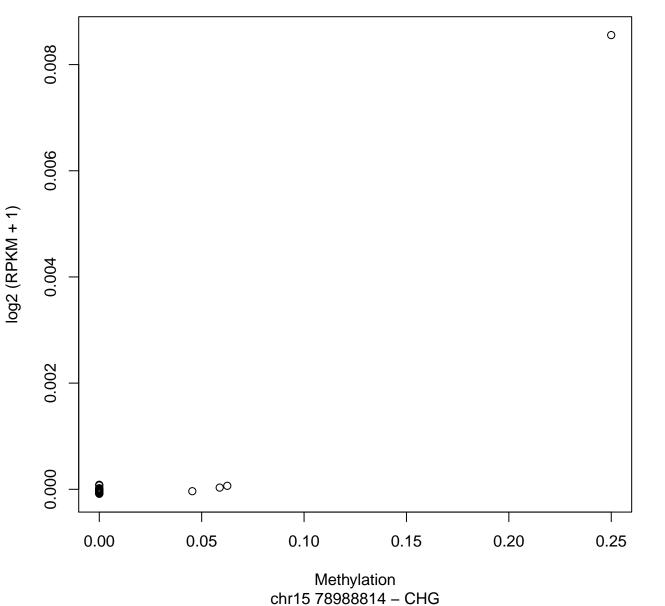
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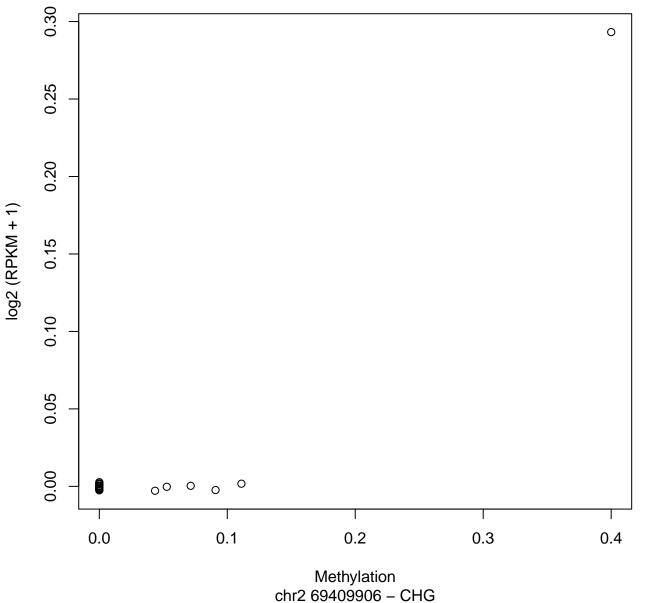
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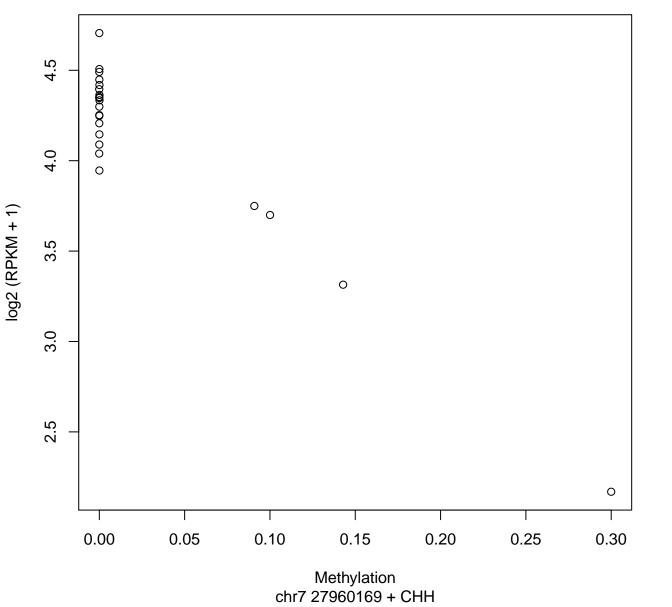
### gene ENSG00000261303.5\_1 FDR 2.98e-06 RP11-160C18.2



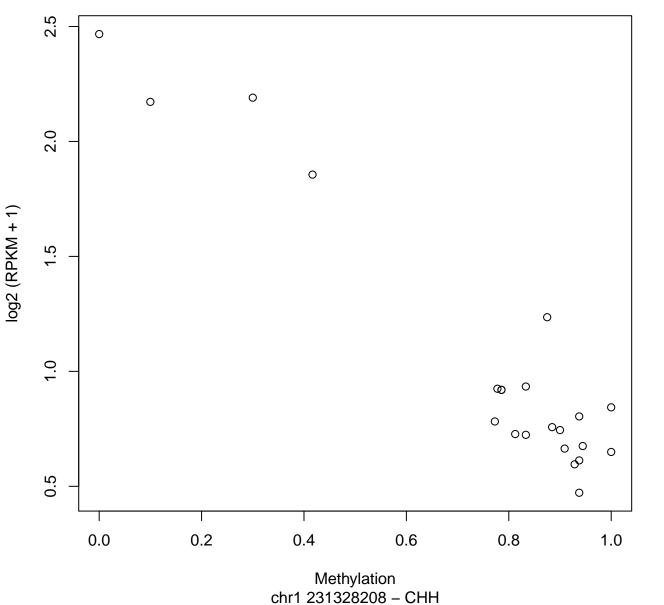
# gene ENSG00000199460.2 FDR 1.11e-05 RNU6-1216P



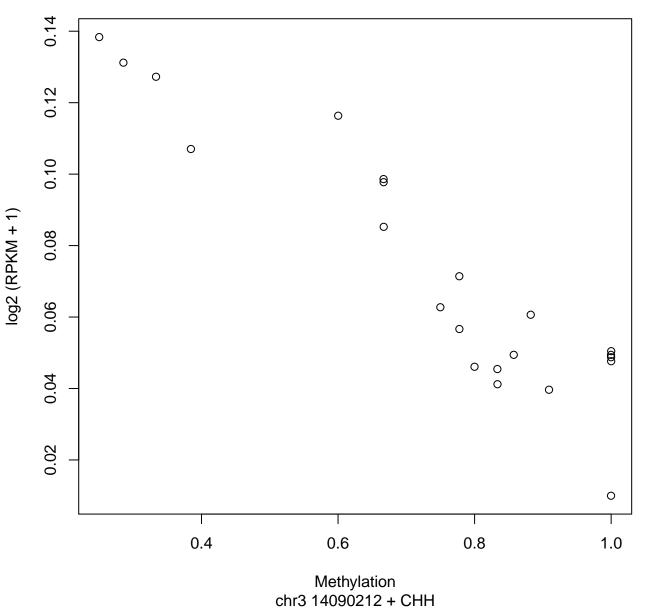
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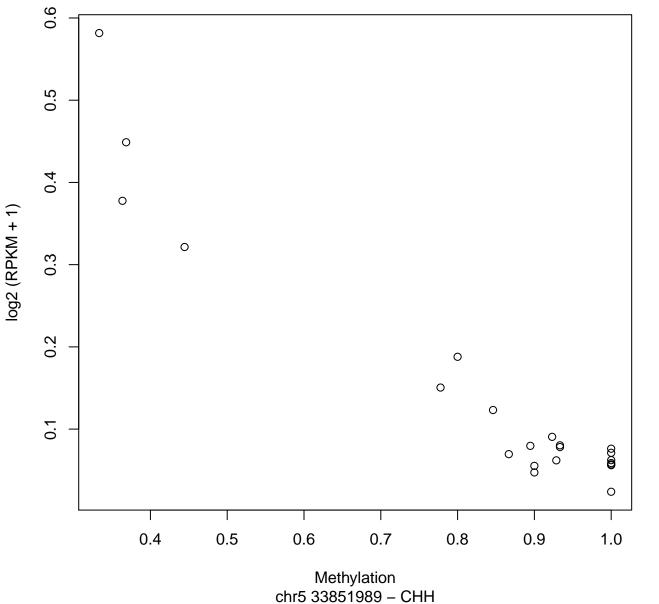
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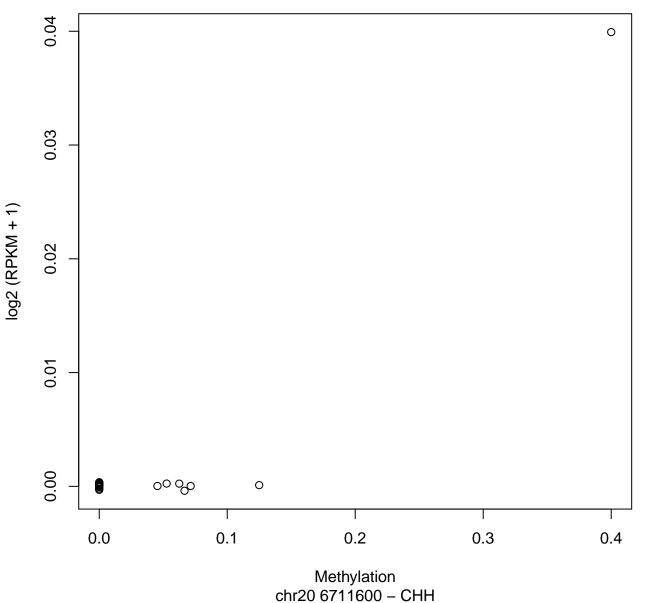
### gene ENSG00000180438.15\_2 FDR 1.45e-05 TPRXL



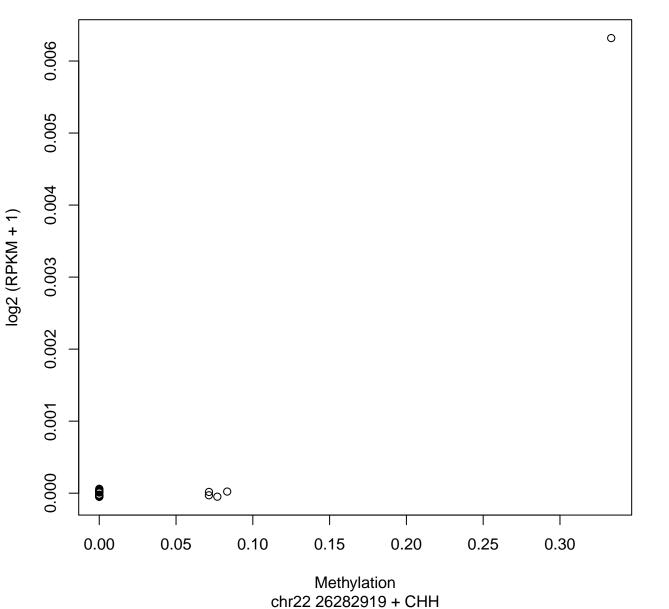
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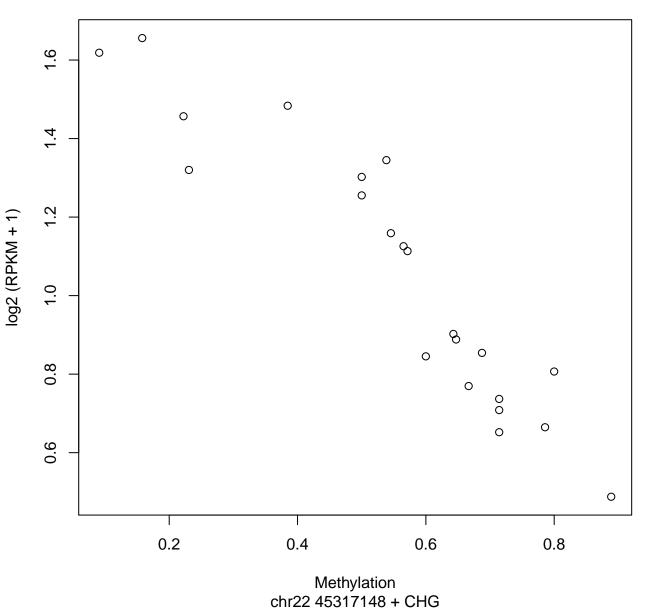
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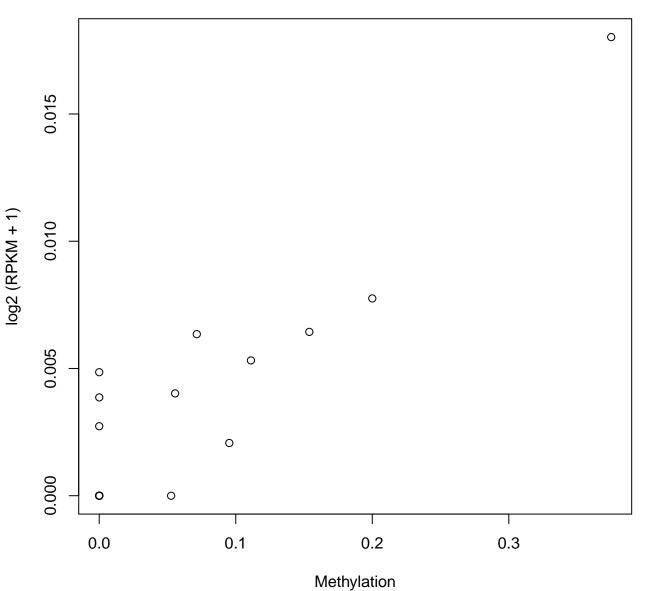
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#### gene ENSG00000056487.15\_2 FDR 7.14e-06 PHF21B

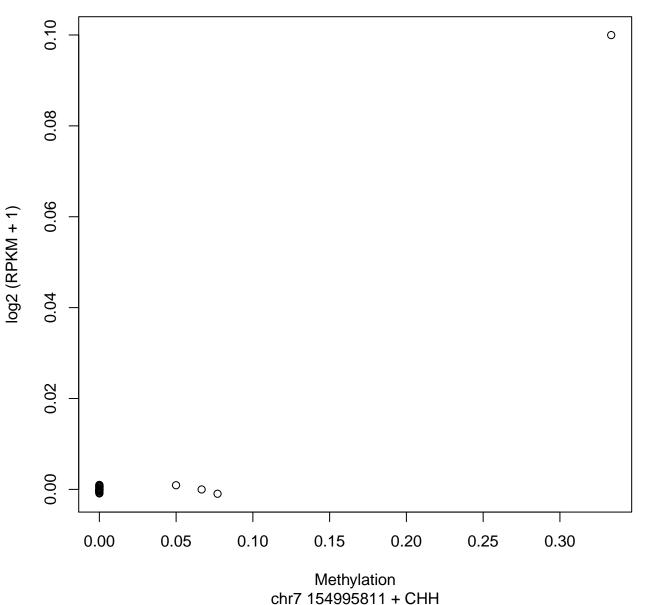


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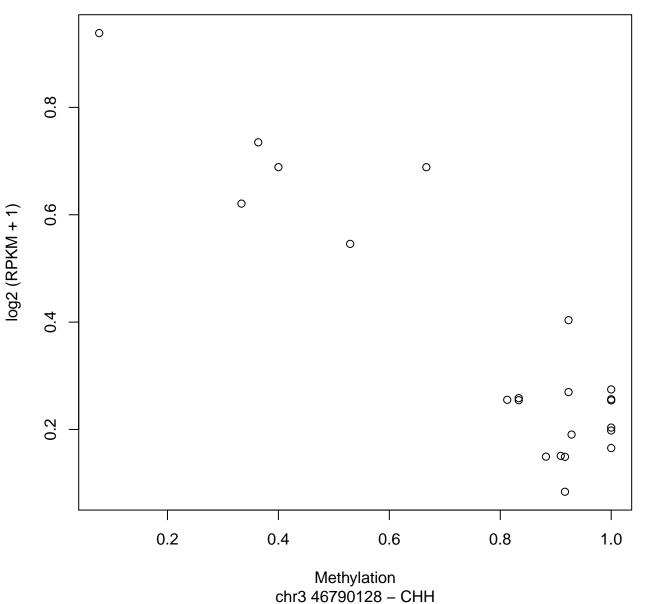


chr3 128865650 + CHG

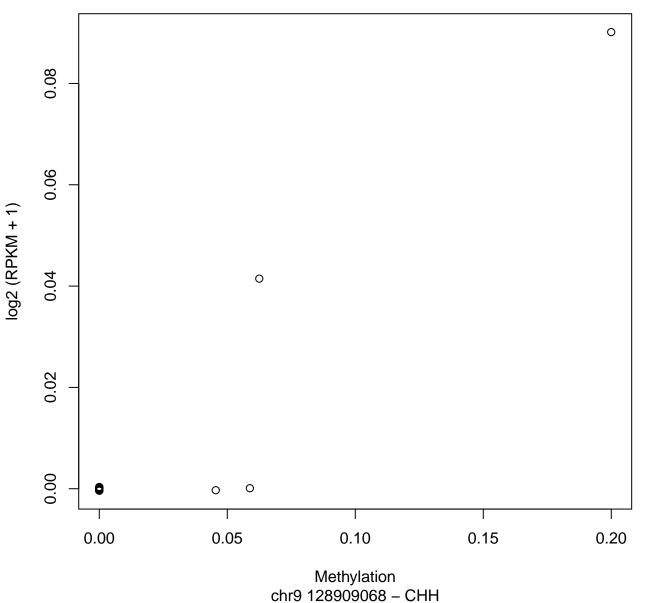
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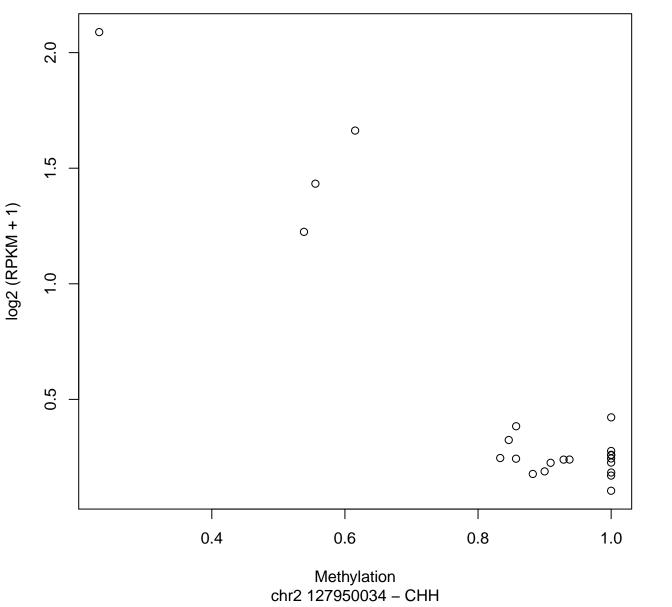
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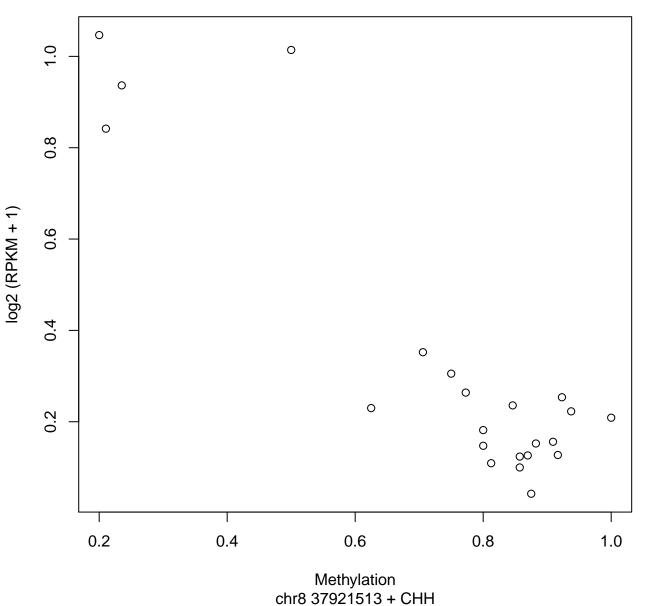
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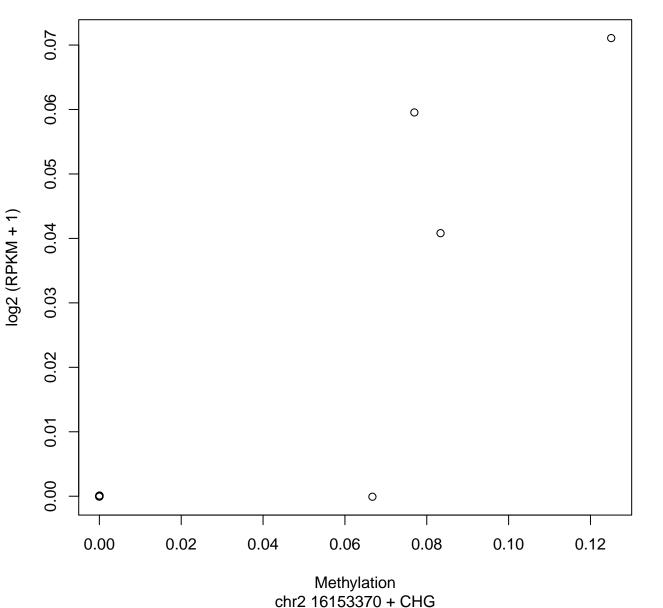
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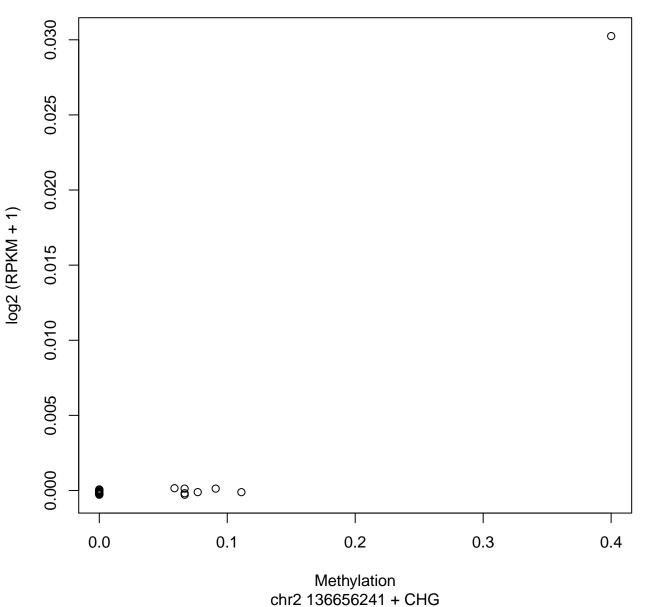
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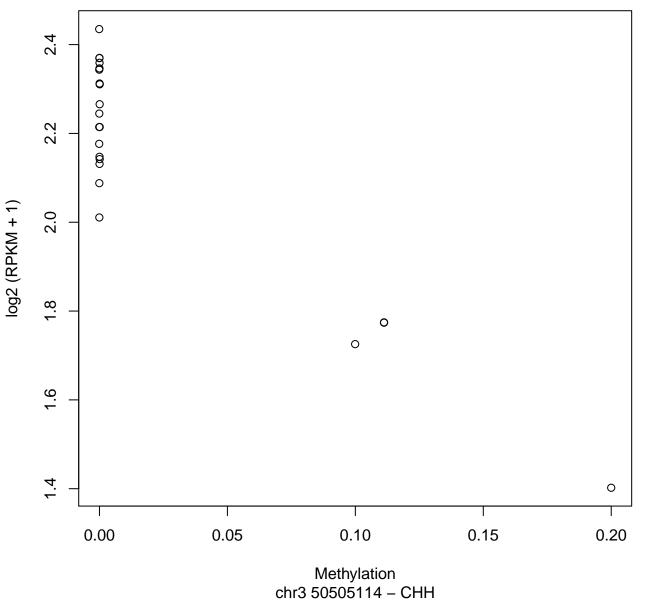
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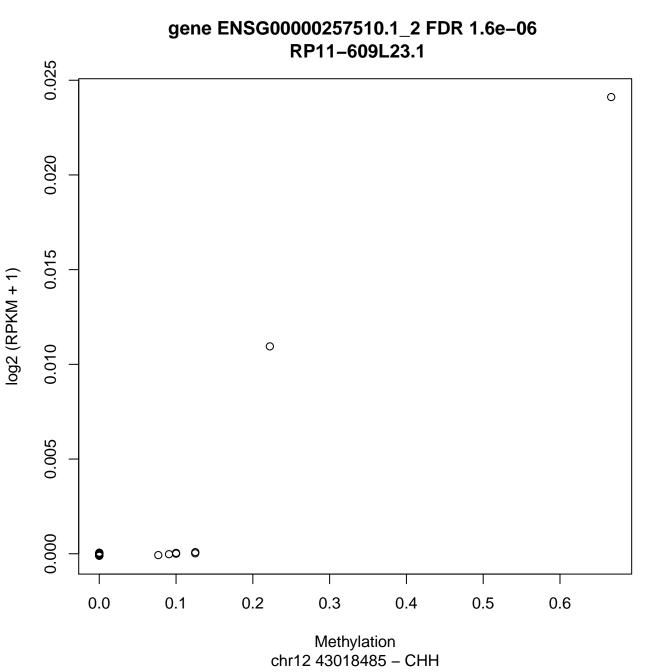


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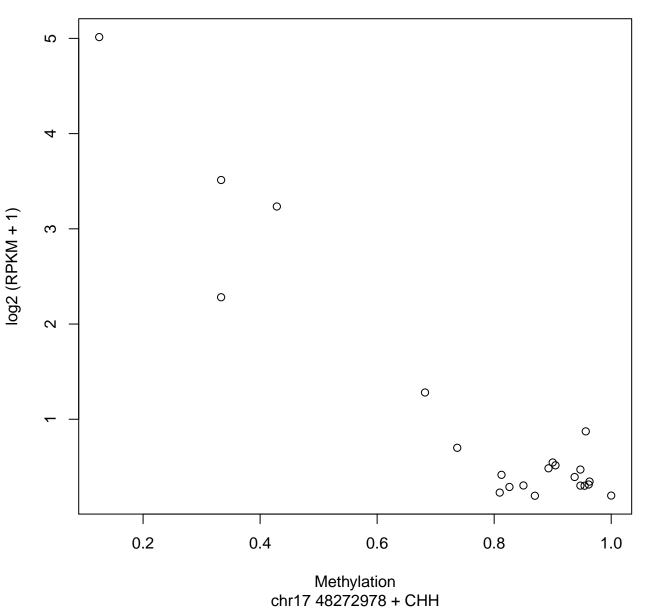


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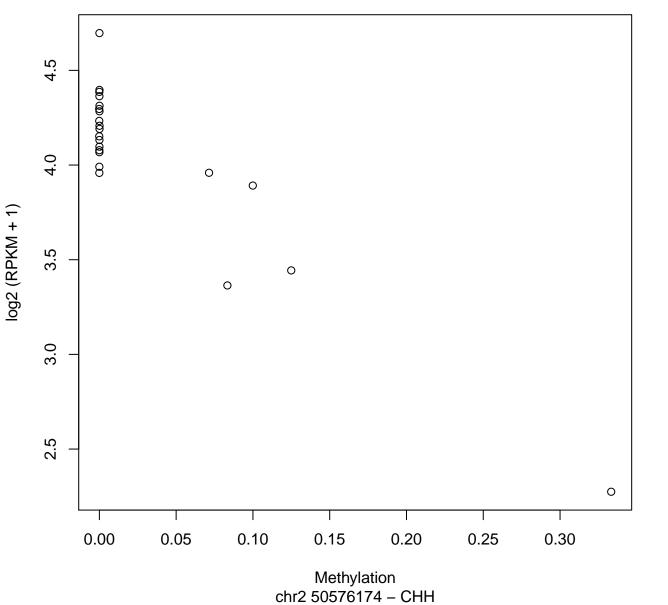




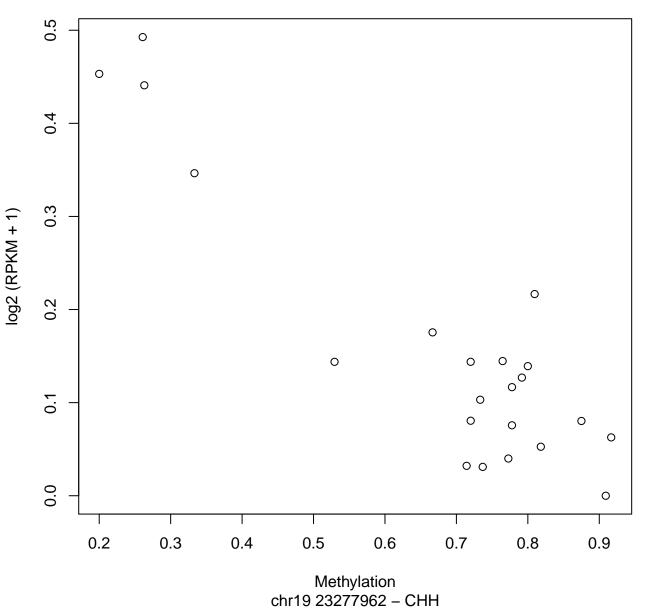
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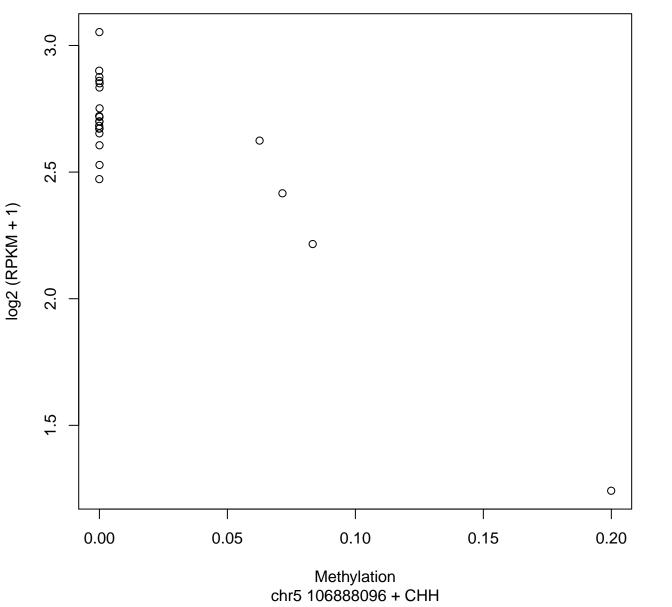
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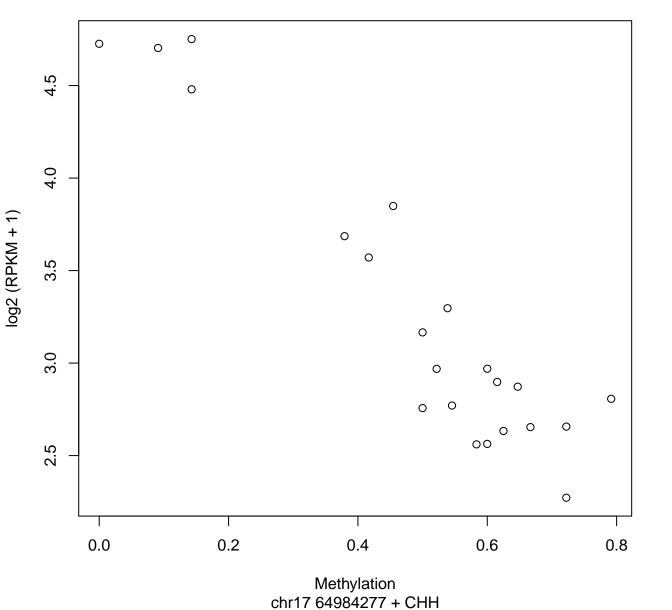
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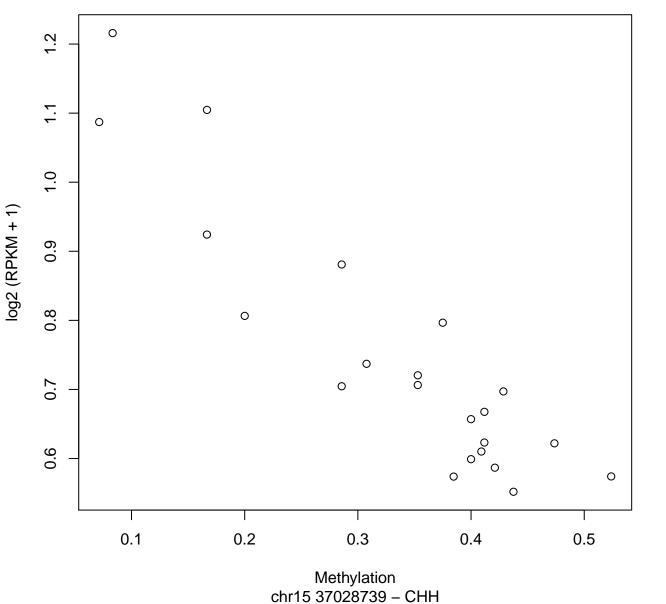
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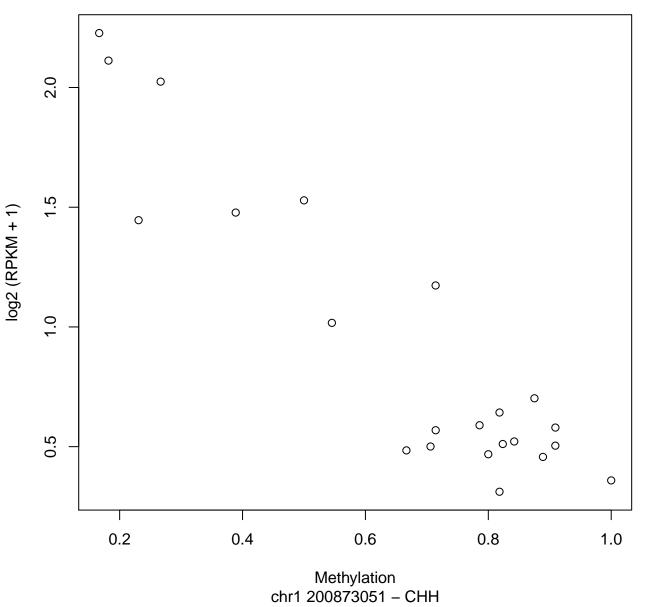
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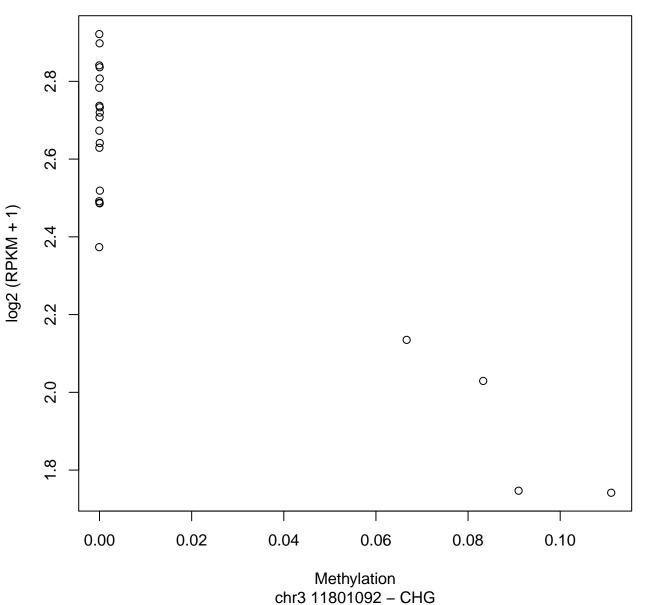
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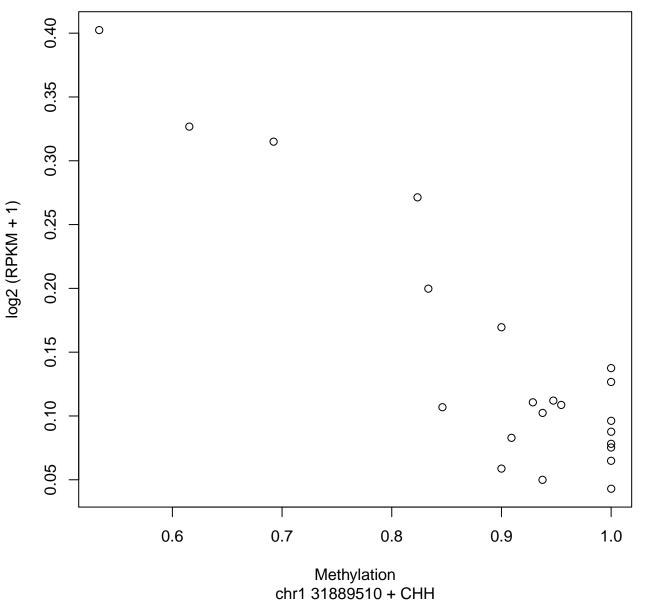
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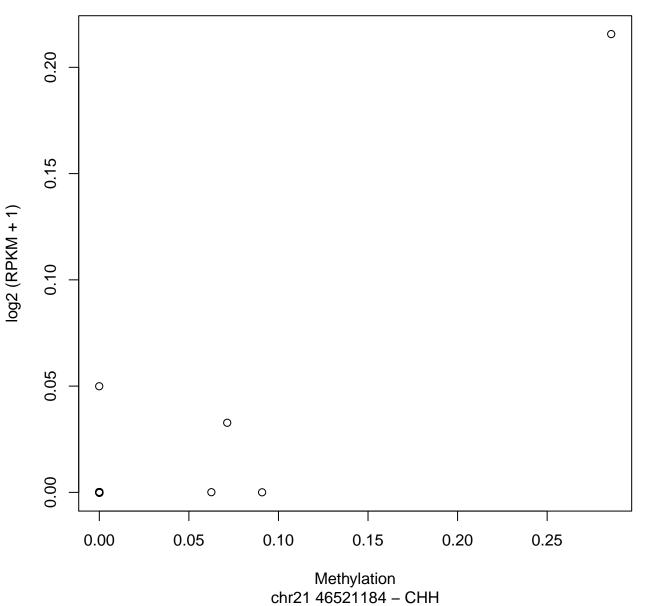
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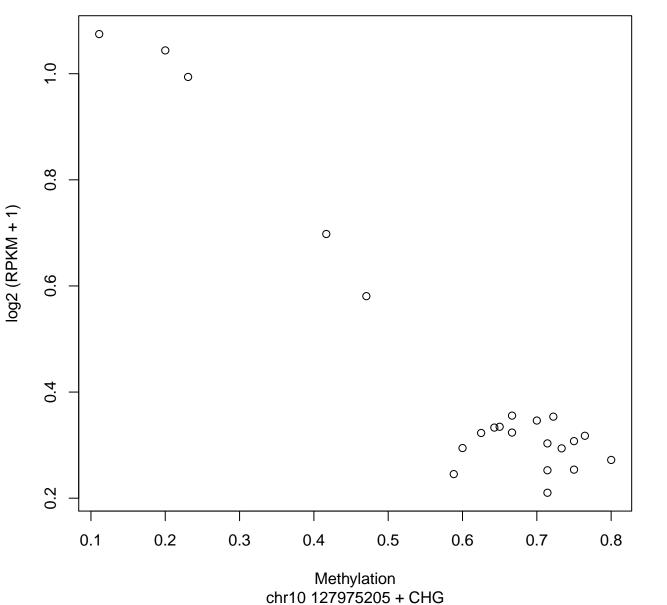
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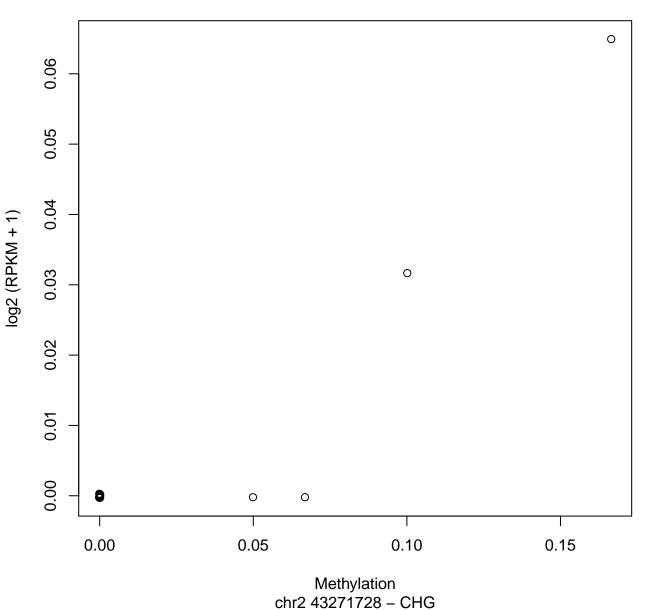
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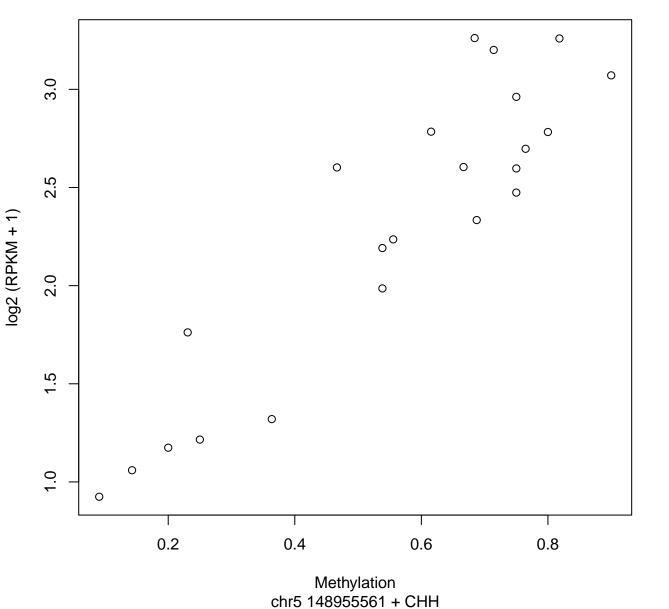
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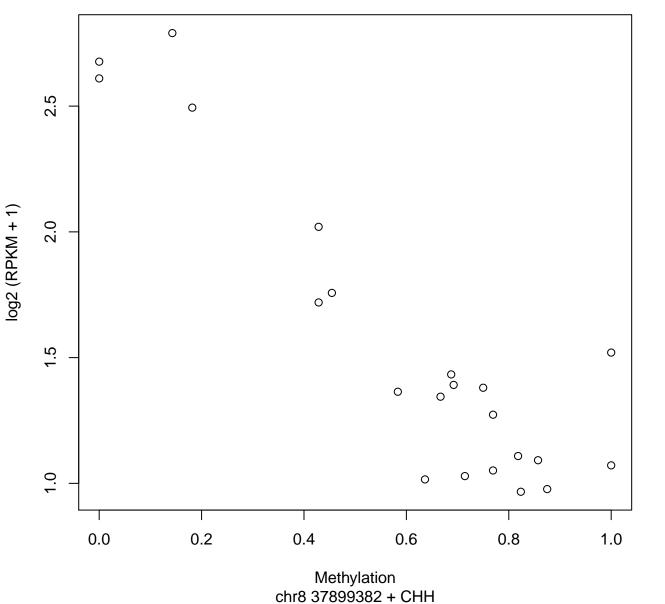
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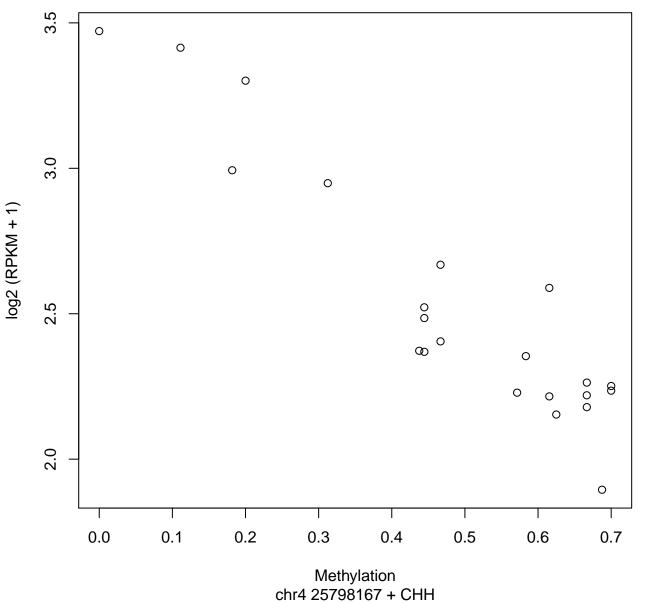
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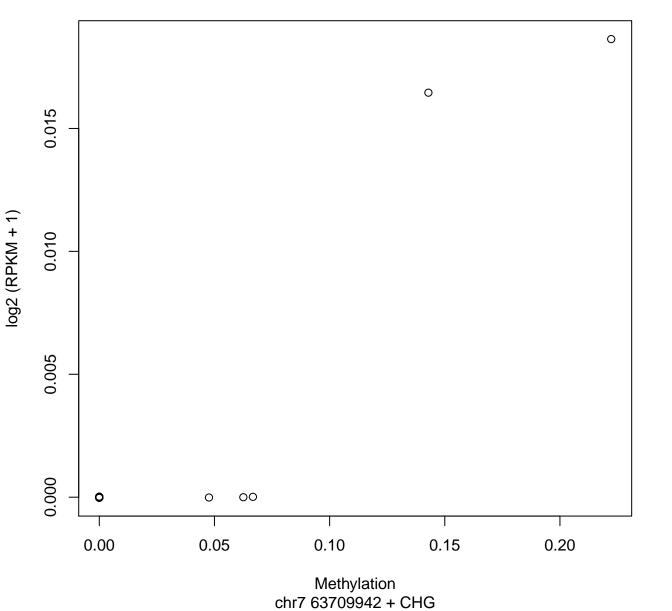
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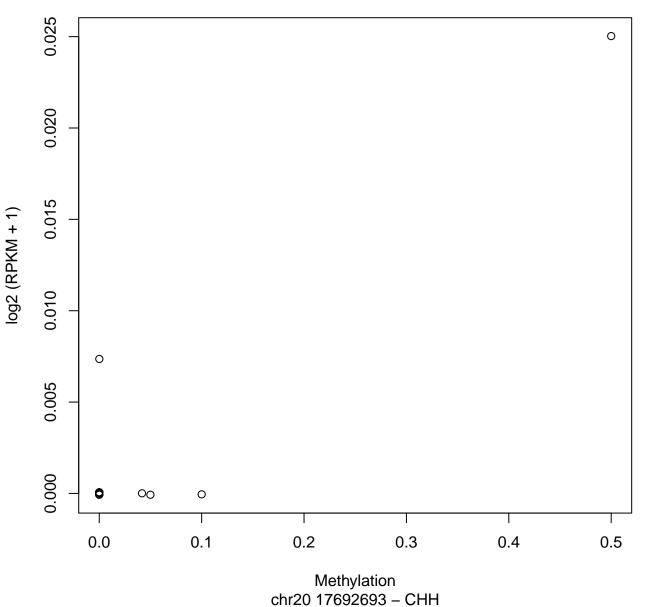
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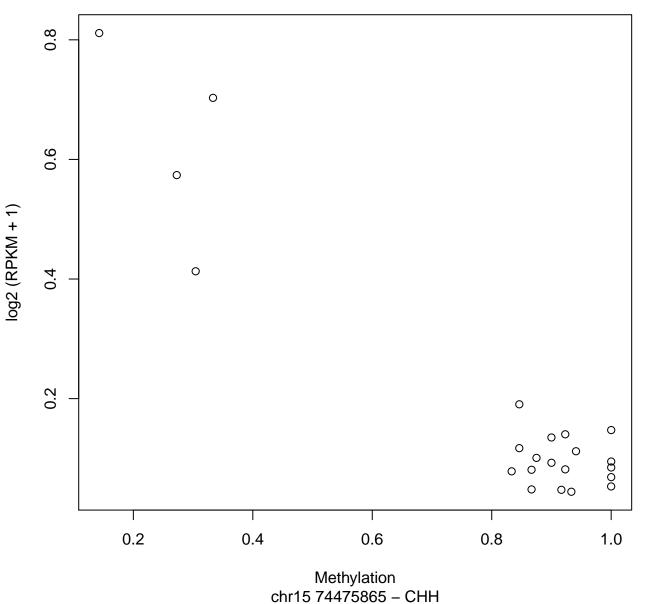
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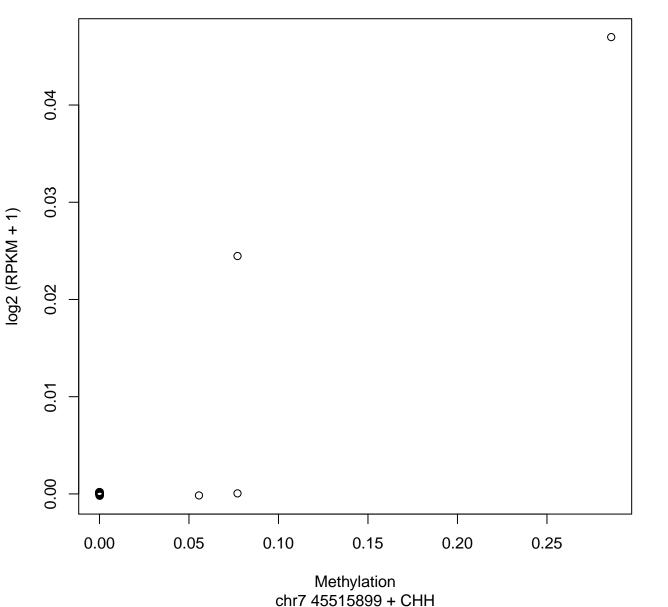
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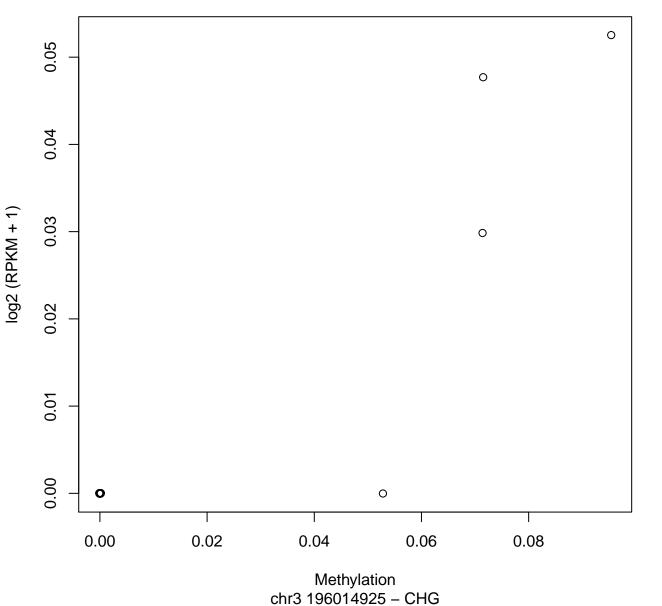
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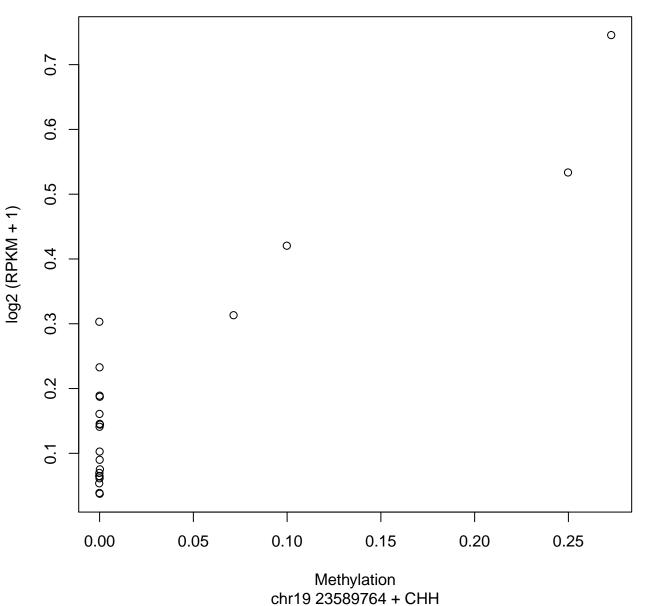
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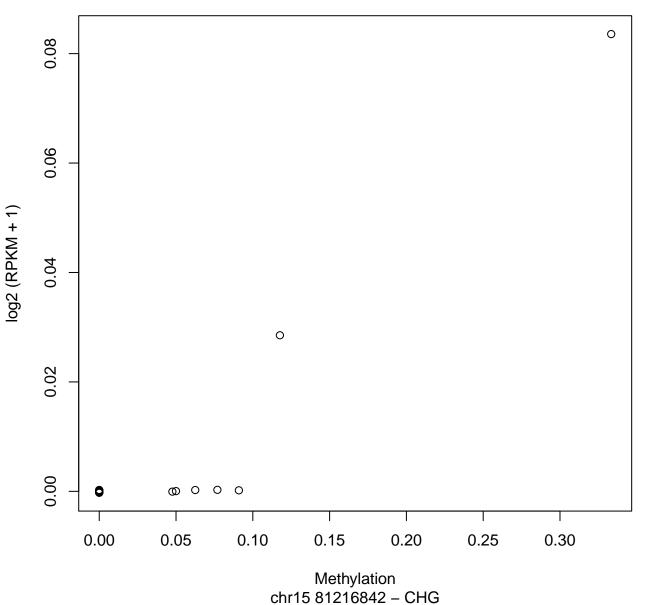
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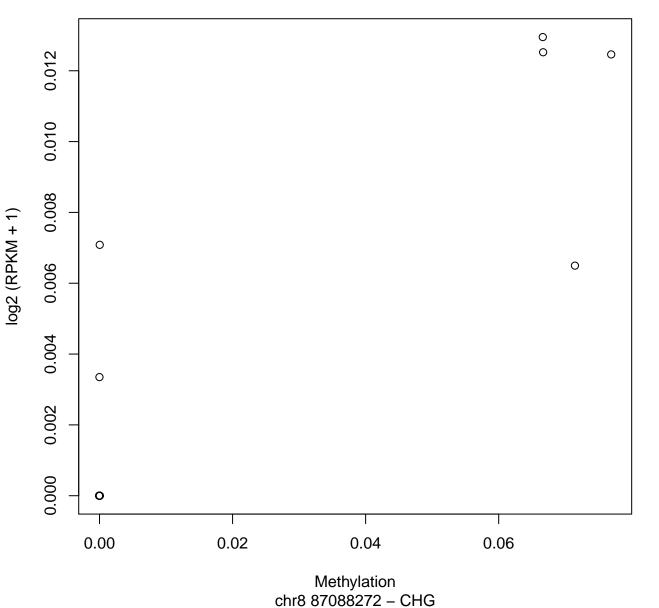
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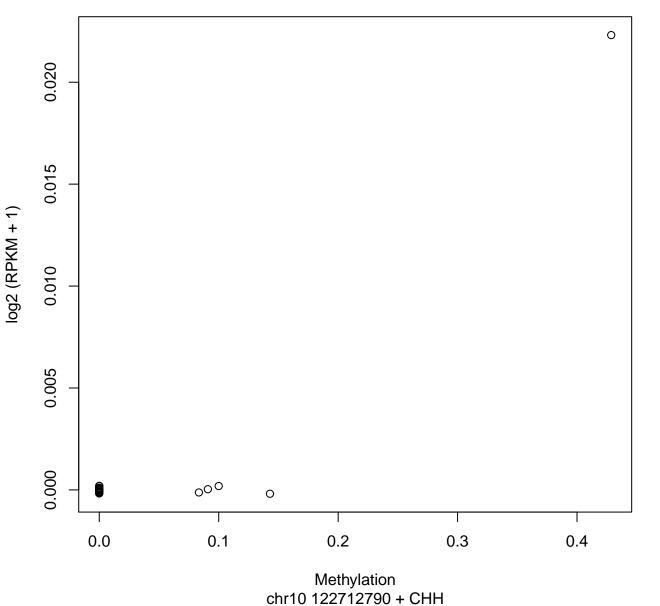
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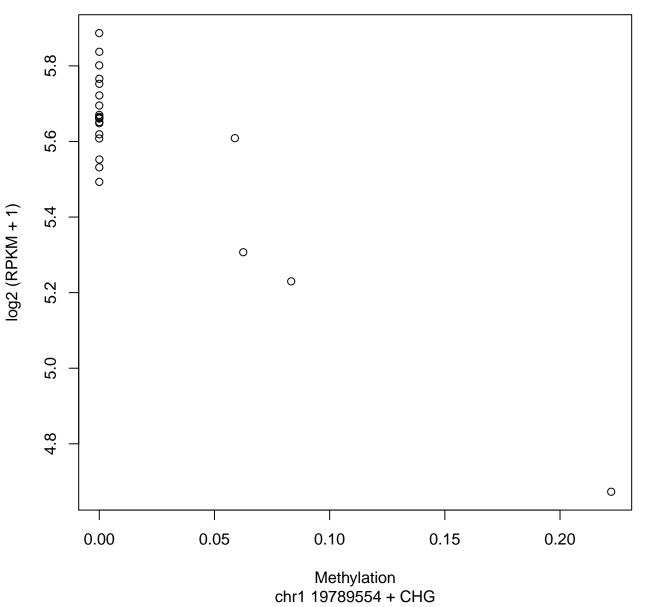
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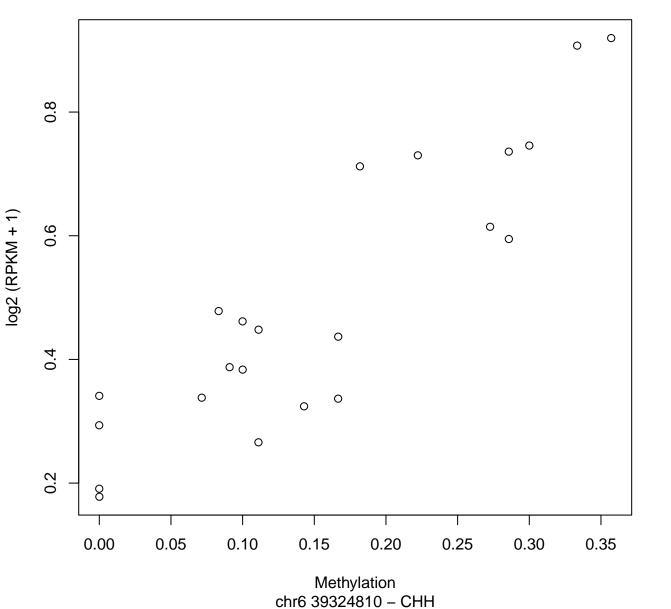
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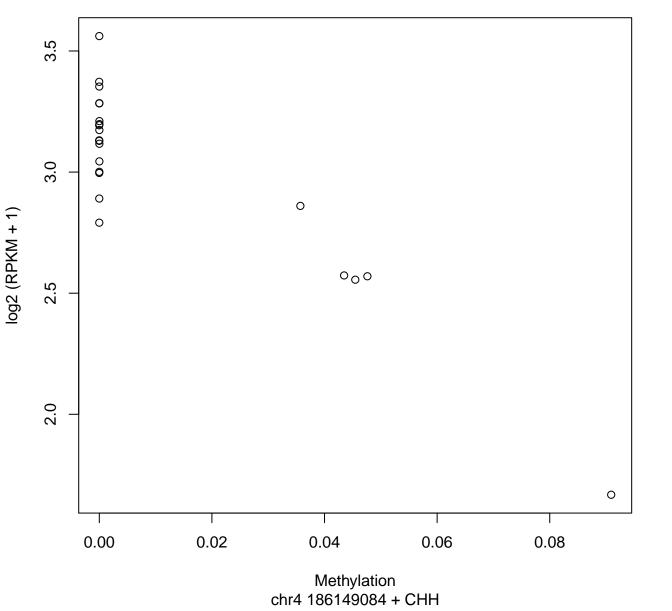
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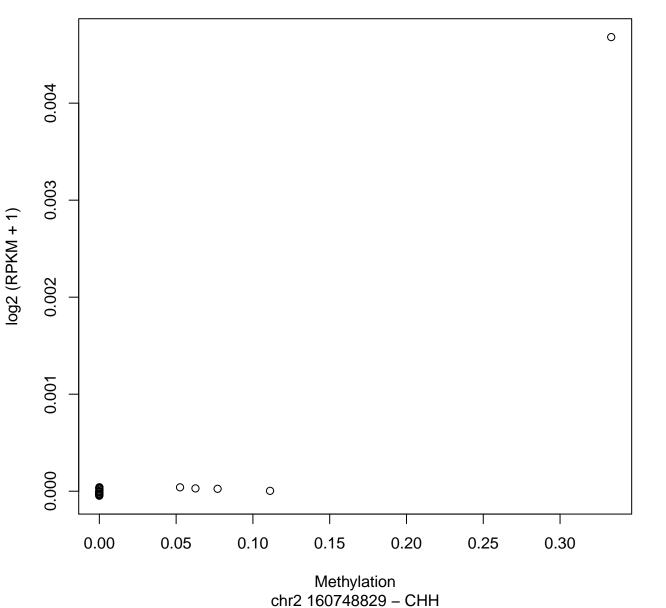
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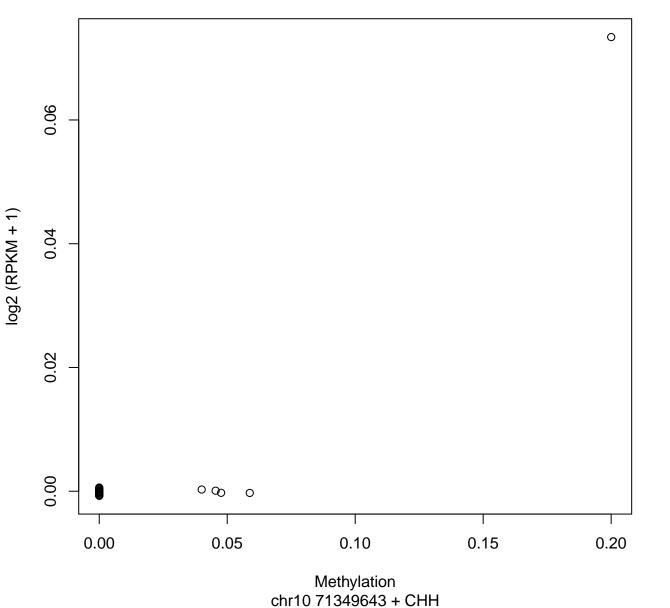
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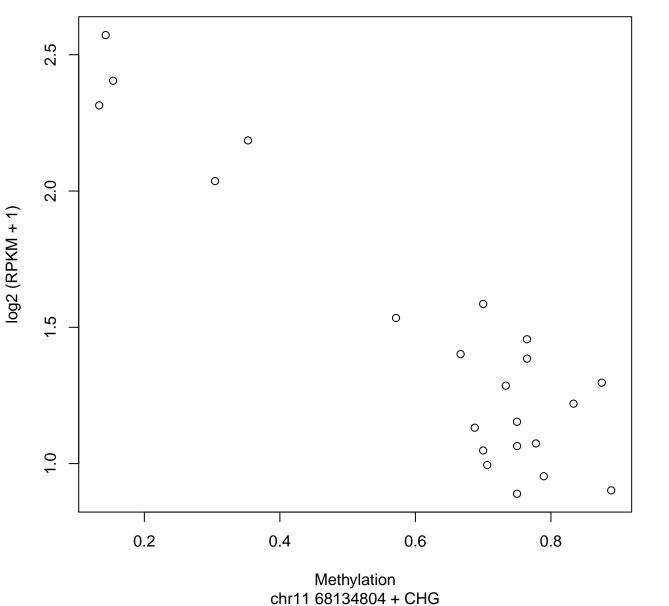
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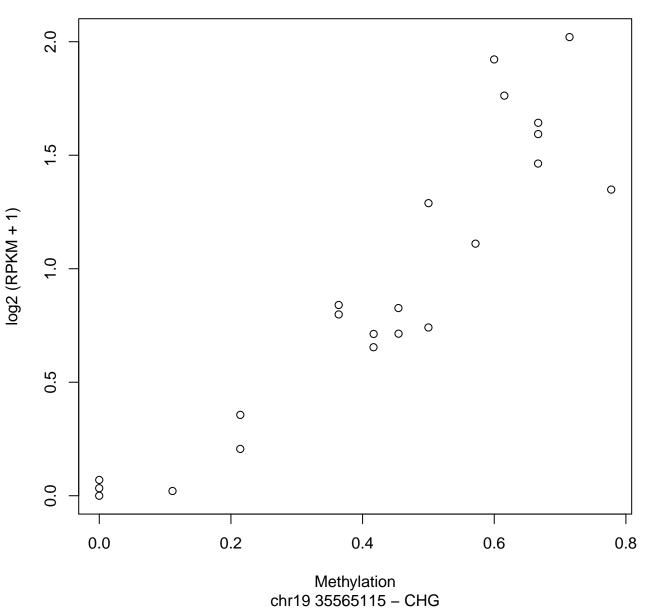
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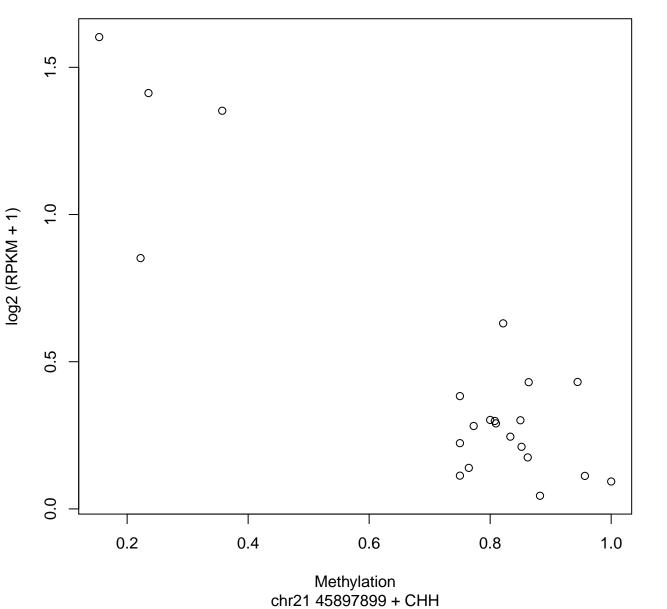
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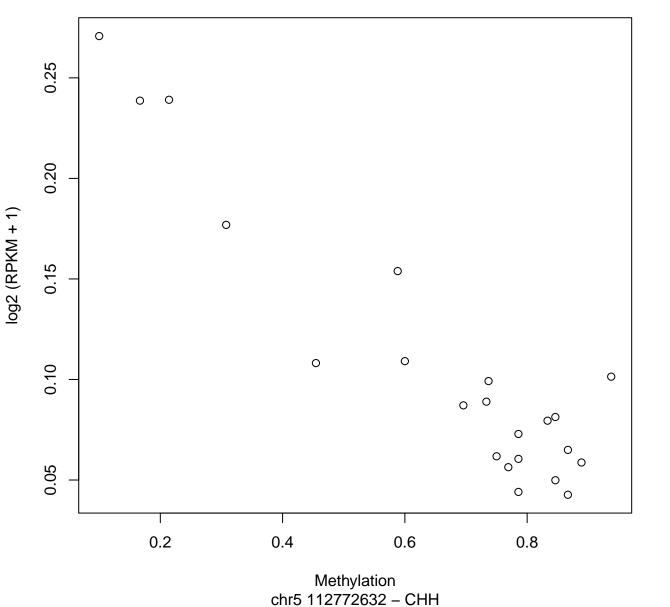
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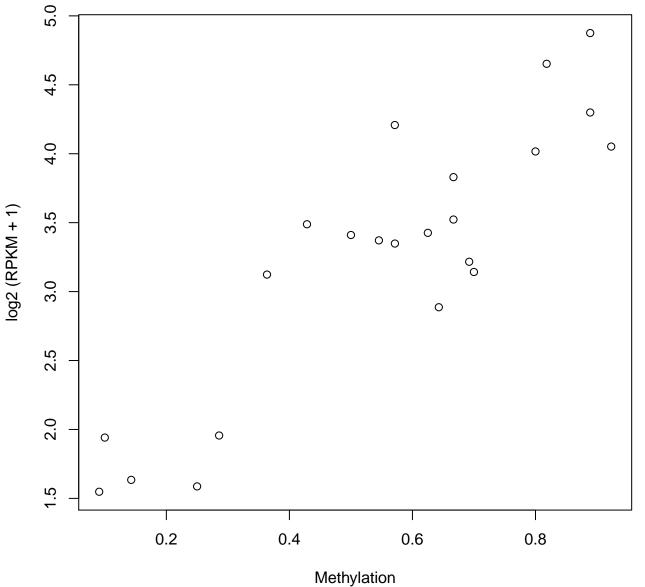
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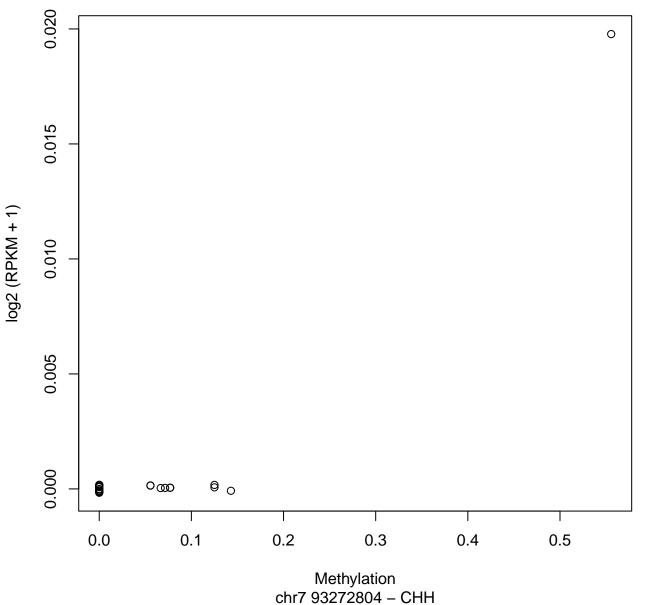


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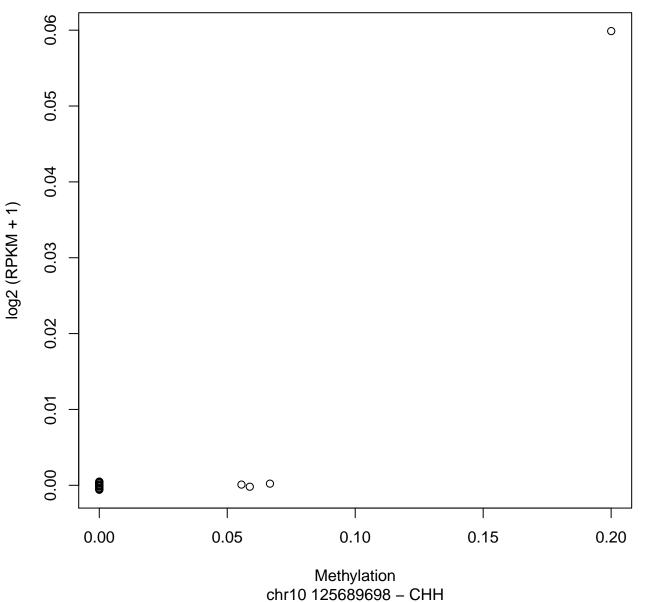


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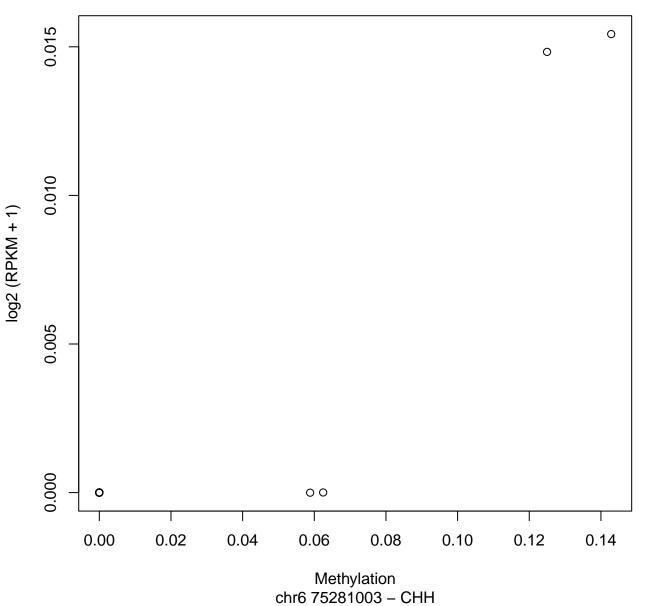
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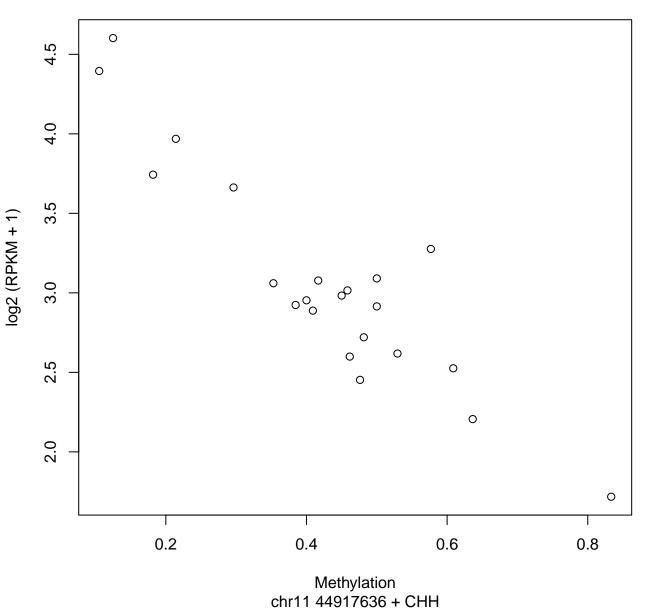
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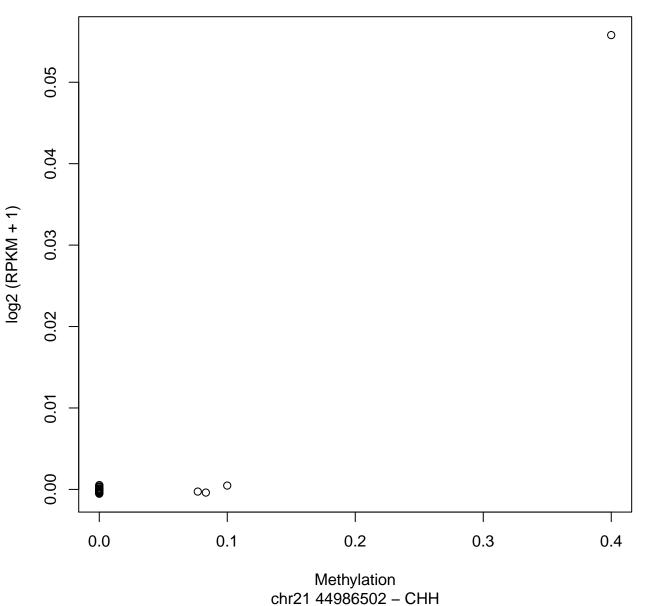
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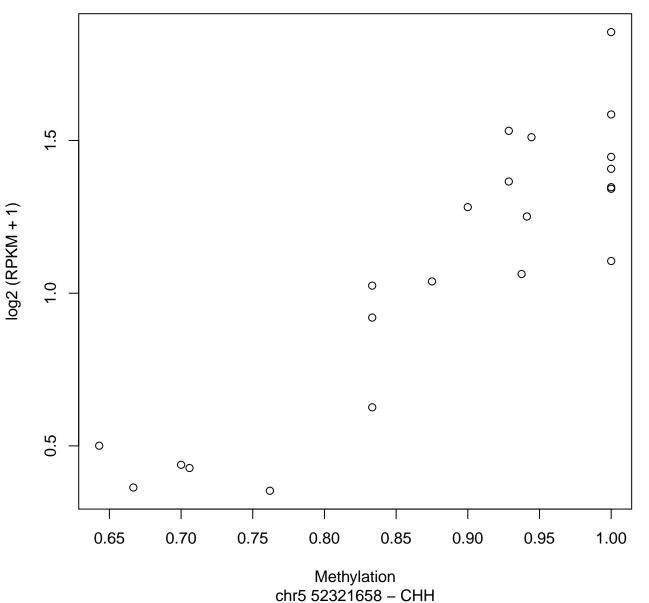
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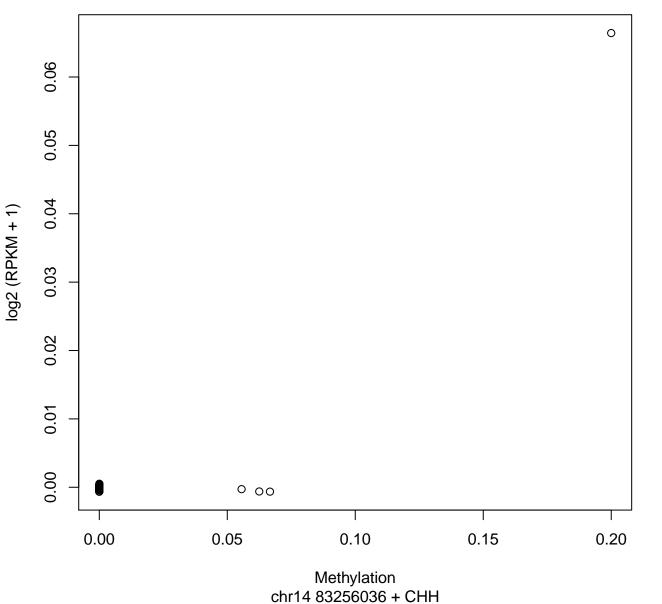
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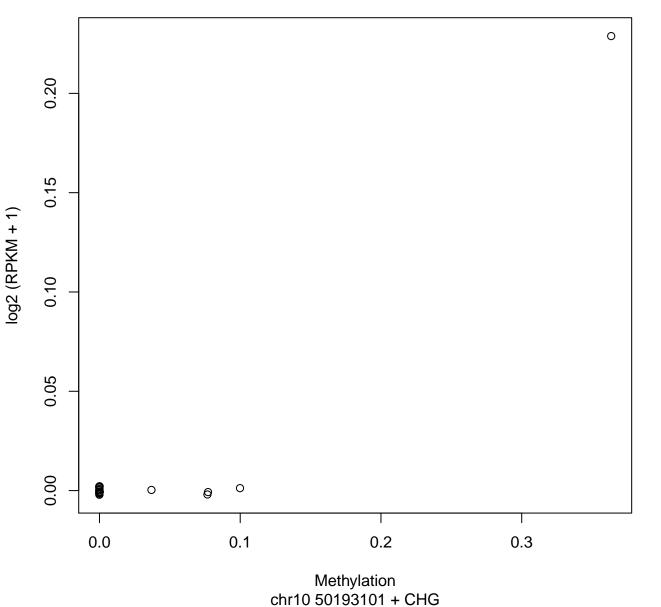
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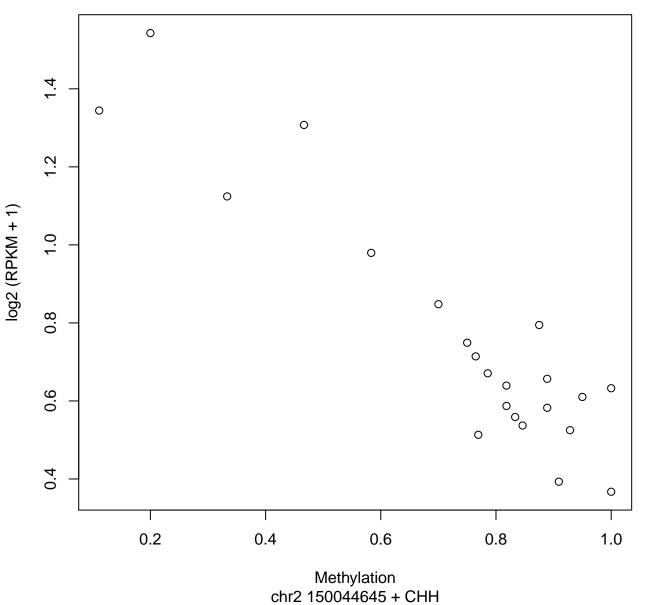
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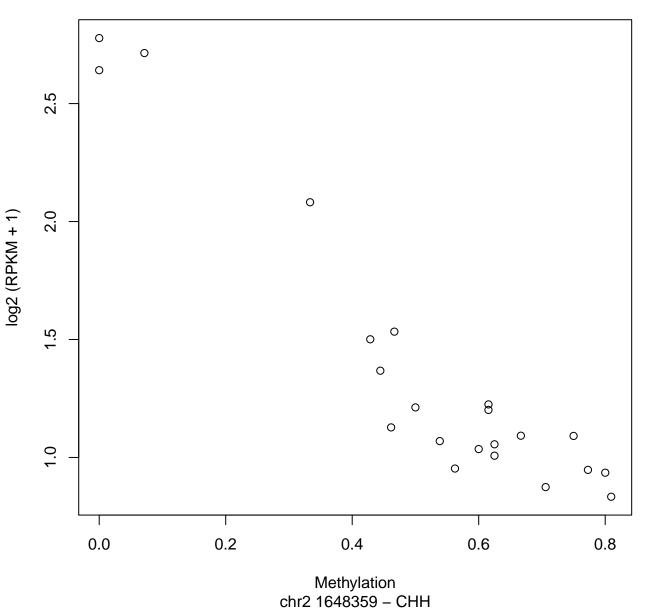
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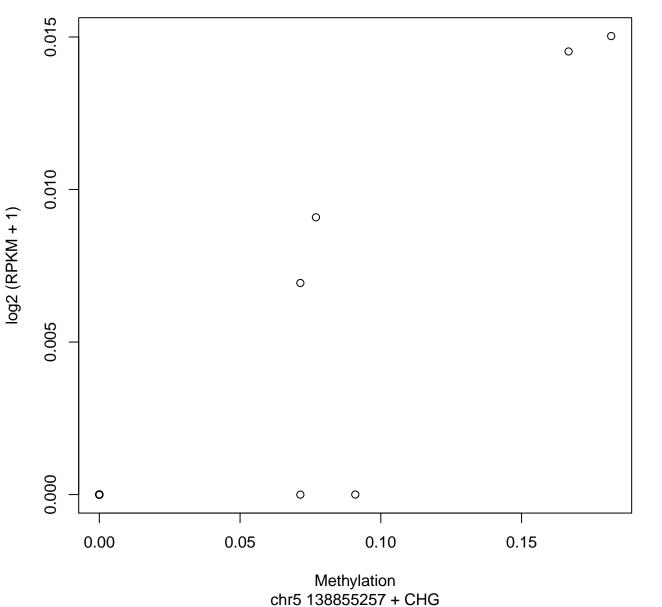
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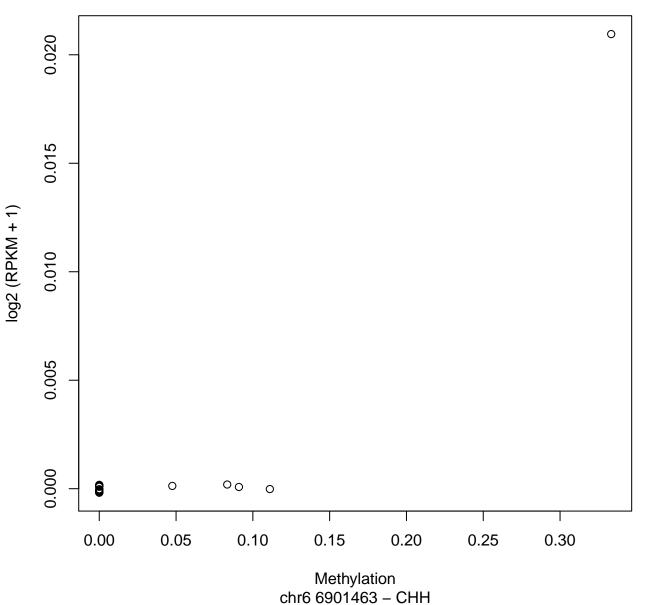
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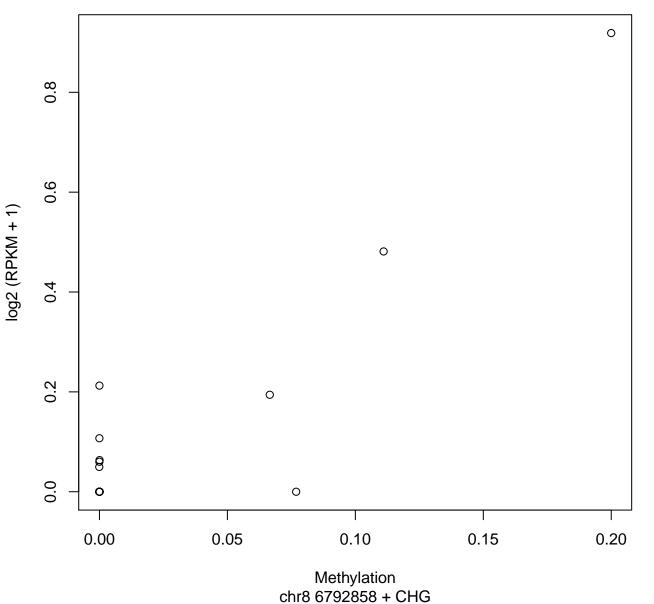
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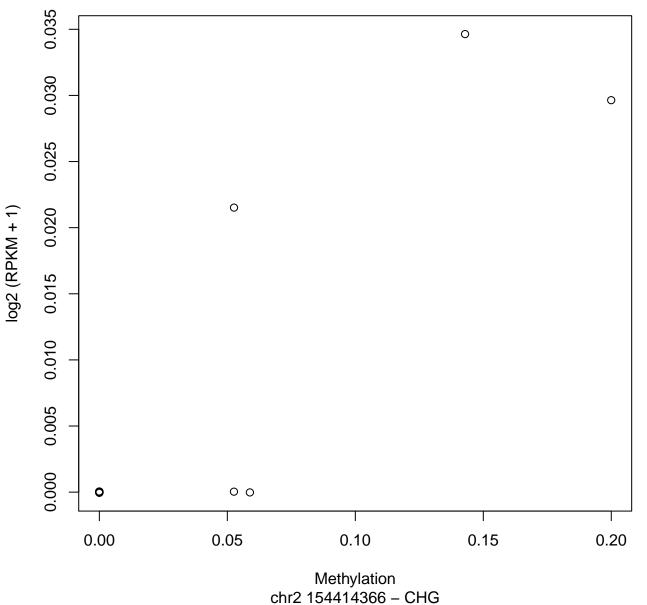
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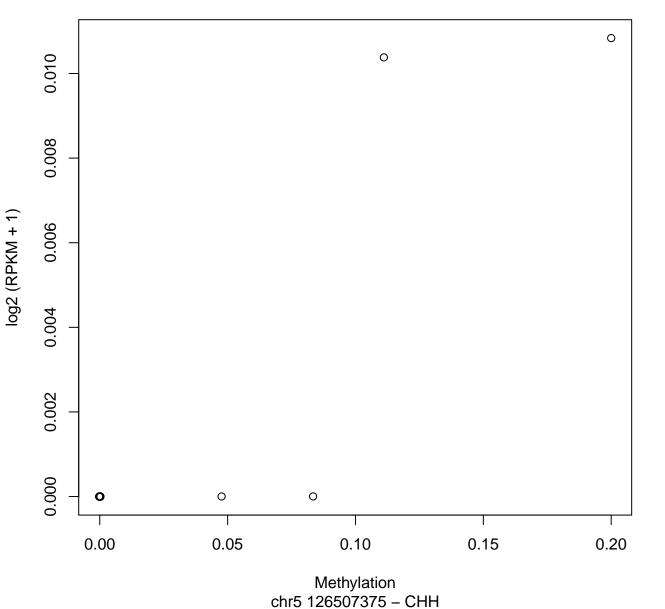
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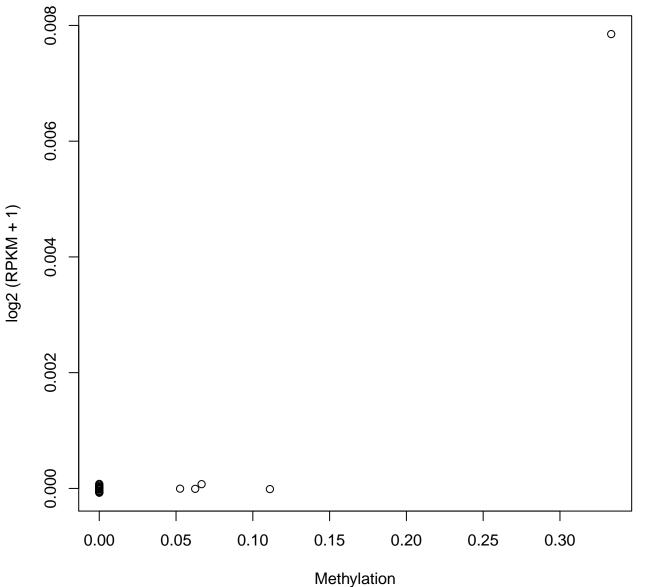
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#### gene ENSG00000250080.1\_1 FDR 0.000463 CTB-88F18.2

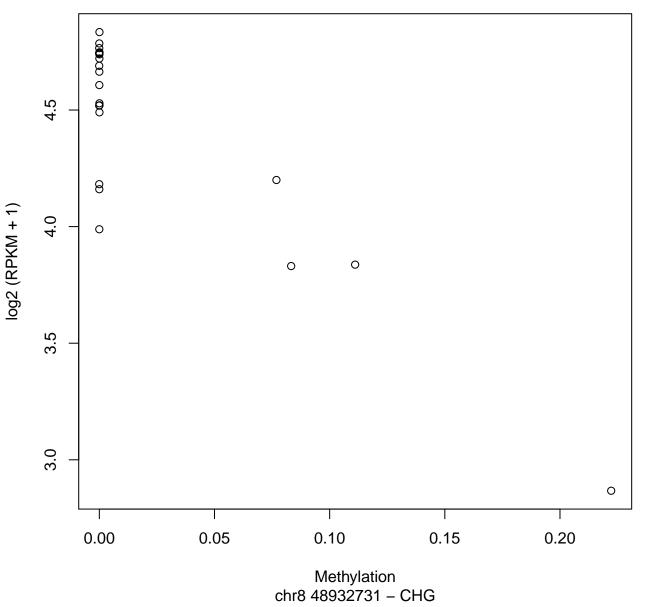


### gene ENSG00000280086.1\_1 FDR 3.57e-05 RP11-544P1.3

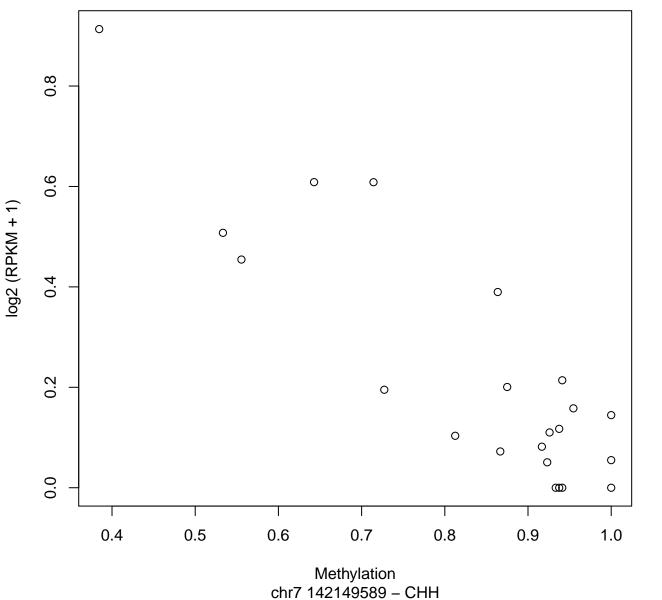


chr10 11438063 + CHG

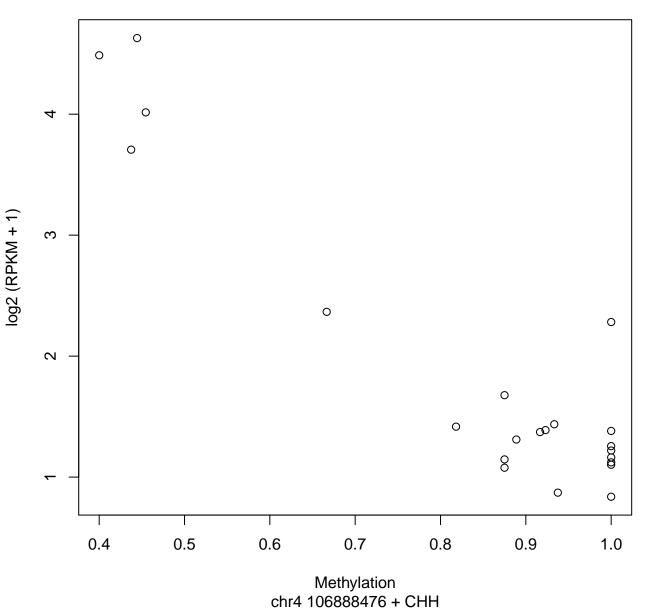
### gene ENSG00000169139.11\_1 FDR 0.000575 UBE2V2



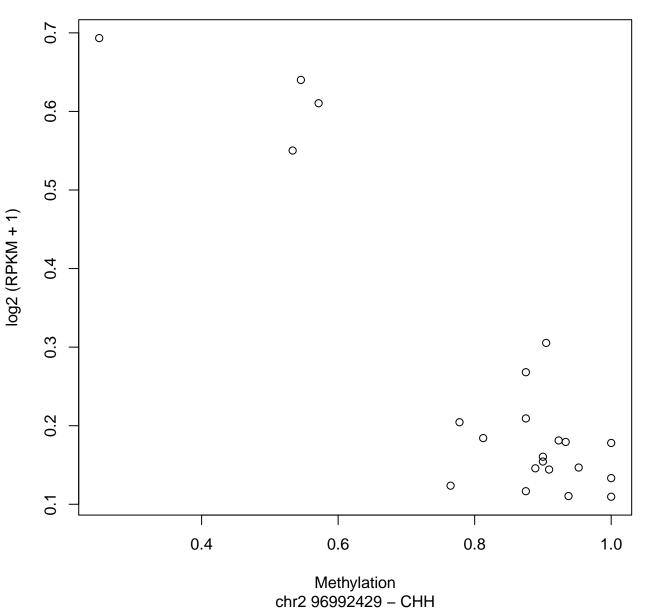
### gene ENSG00000211725.3\_1 FDR 0.000502 TRBV5-5



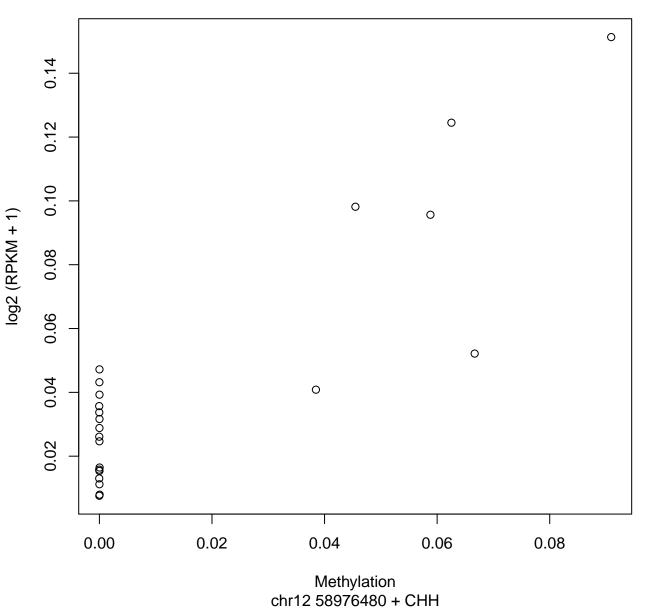
#### gene ENSG00000168743.12\_1 FDR 4.81e-06 NPNT



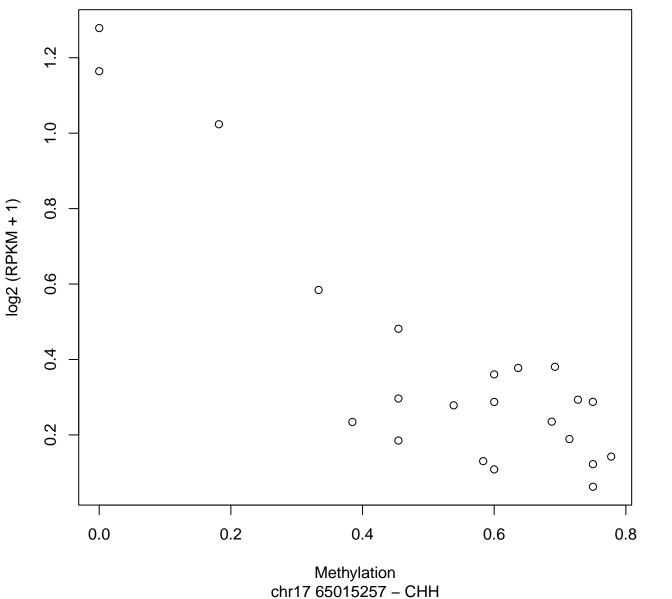
# gene ENSG00000198885.9\_2 FDR 8.74e-05 ITPRIPL1



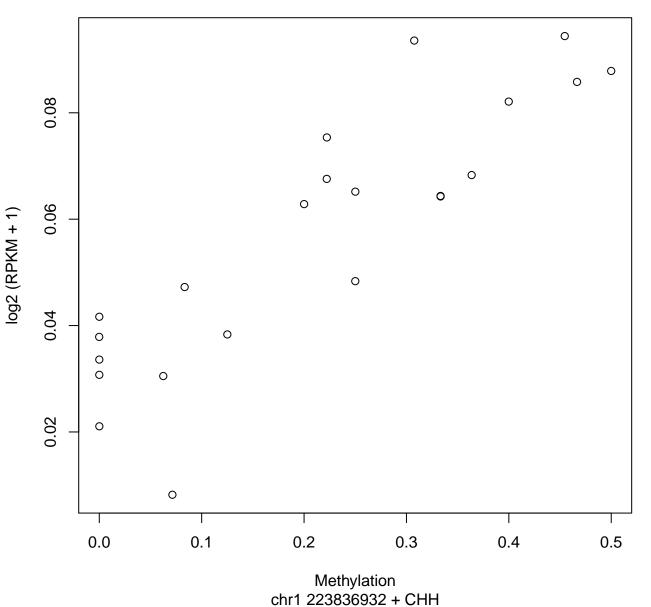
# gene ENSG00000257259.1\_1 FDR 0.000597 RP11-767I20.1



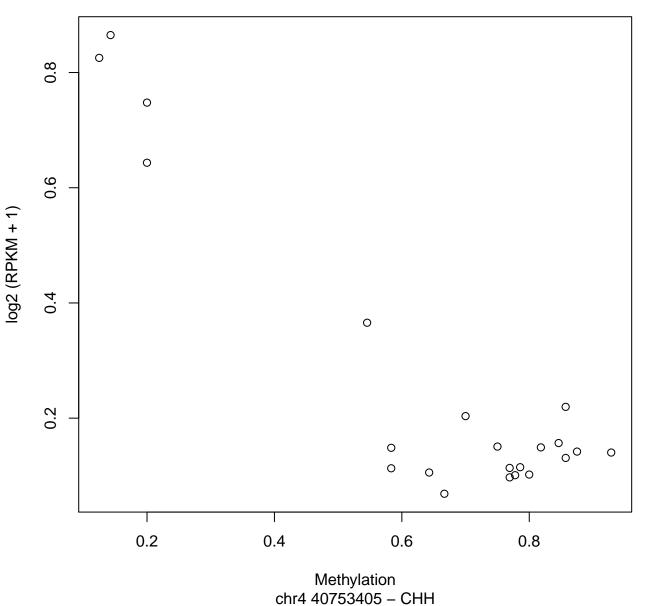
### gene ENSG00000264491.1\_1 FDR 0.000518 RP11-349A8.3



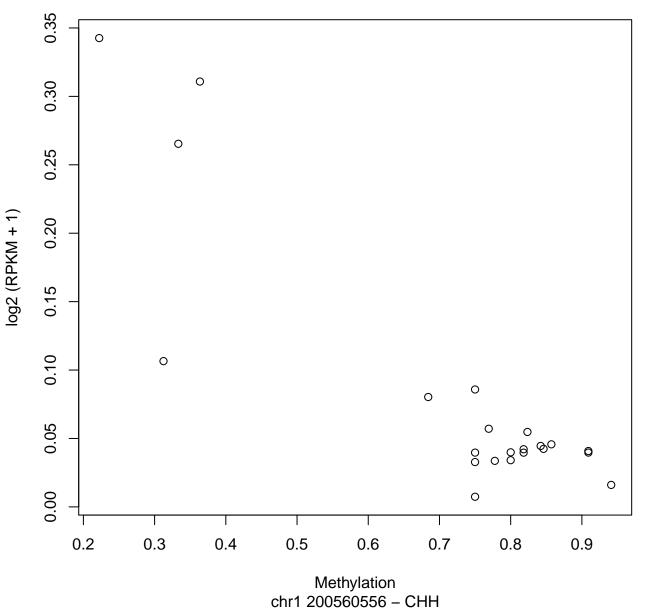
### gene ENSG00000203697.11\_2 FDR 0.000581 CAPN8



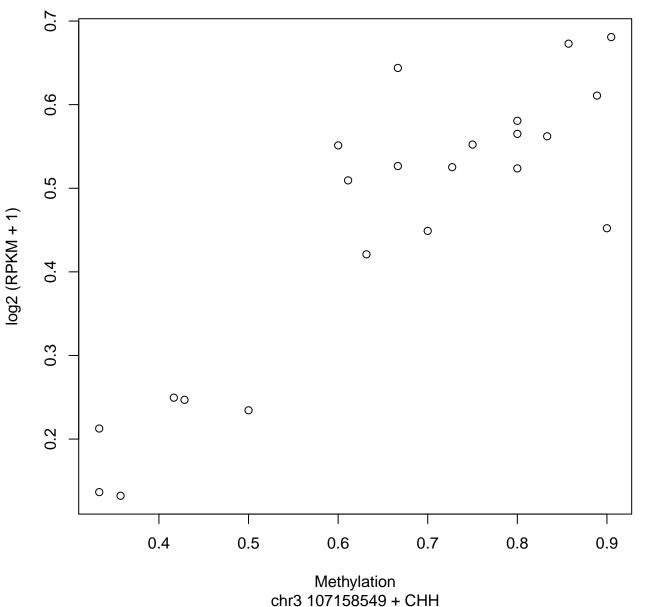
# gene ENSG00000179299.16\_2 FDR 3.7e-05 NSUN7



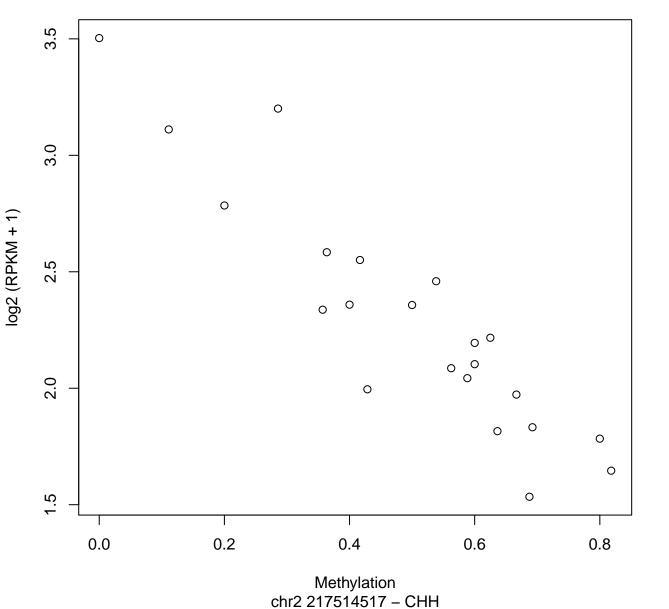
### gene ENSG00000118193.11\_1 FDR 0.00053 KIF14



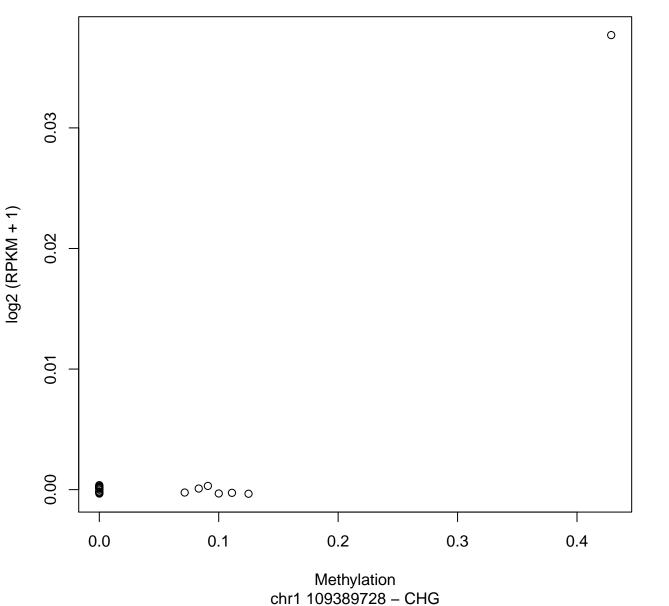
#### gene ENSG00000273125.1\_1 FDR 0.000328 RP11-115H18.1



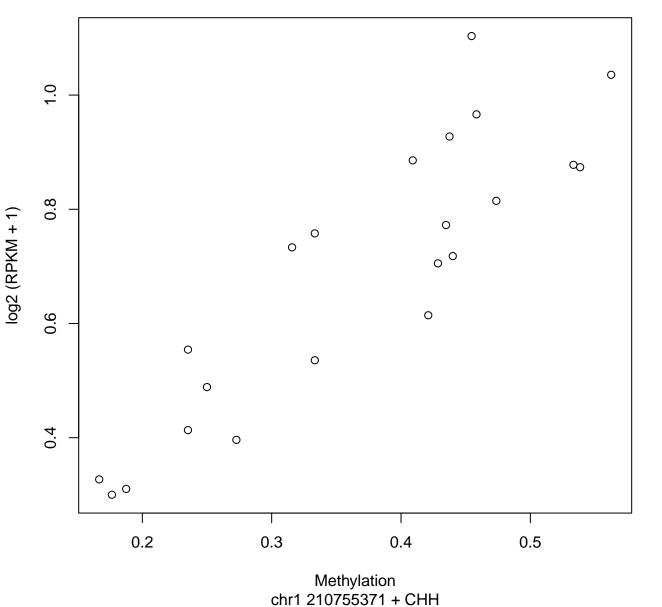
# gene ENSG00000115457.9\_2 FDR 2.05e-05 IGFBP2



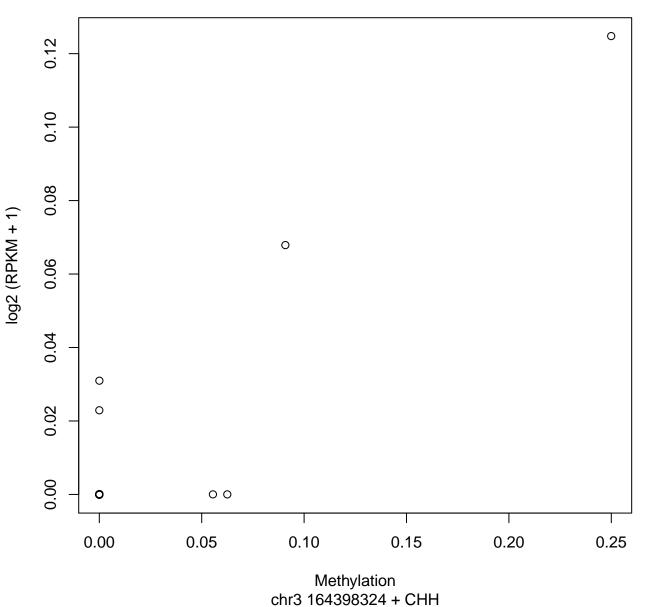
### gene ENSG00000274068.1\_2 FDR 0.000349 RP11-475E11.9



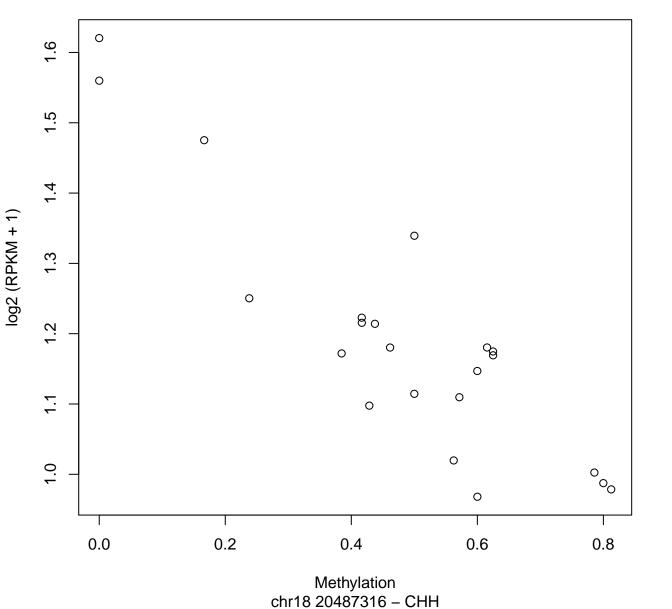
### gene ENSG00000054392.12\_2 FDR 0.000312 HHAT



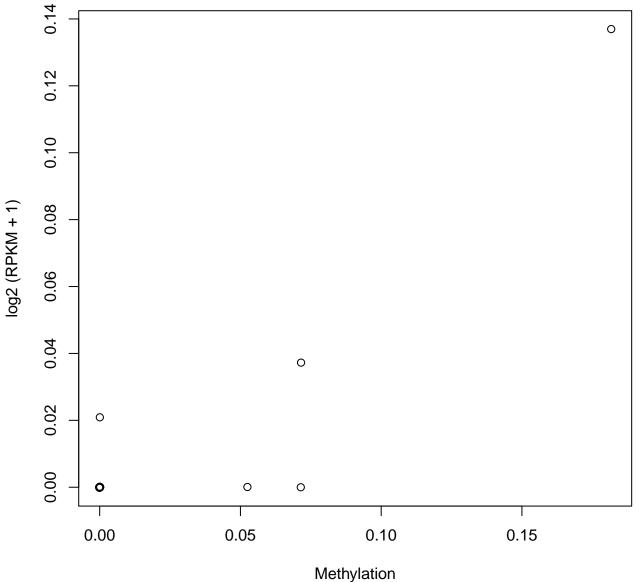
### gene ENSG00000241636.1\_1 FDR 0.000145 LINC01323



### gene ENSG00000101773.17\_2 FDR 0.000289 RBBP8

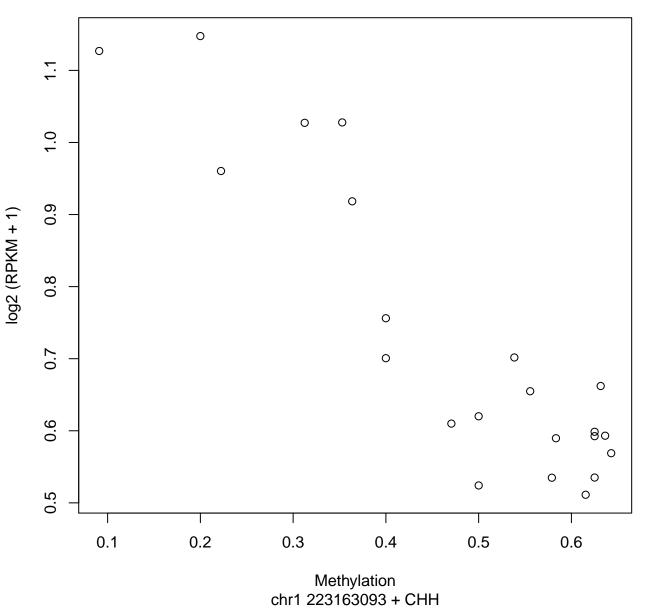


## gene ENSG00000240919.1\_1 FDR 0.000328 RP11-400K9.1



chr8 54446962 + CHH

# gene ENSG00000154309.8\_2 FDR 5.69e-05 DISP1



### gene ENSG00000260192.1\_1 FDR 0.000637 RP11-756H20.1

