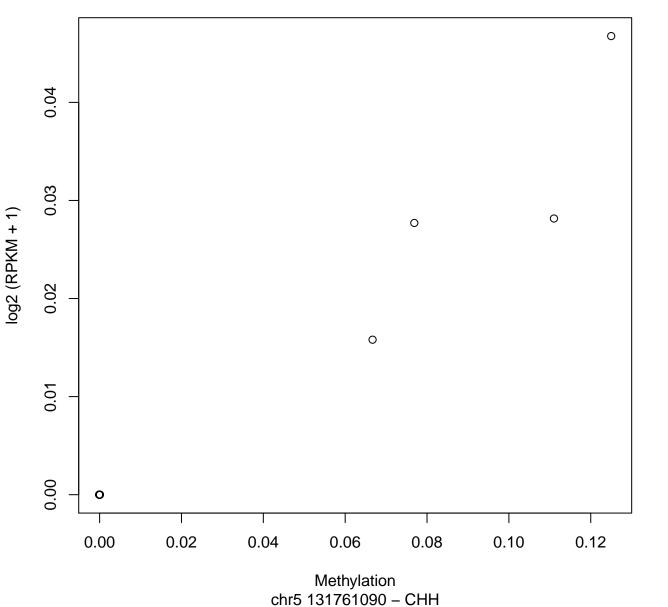
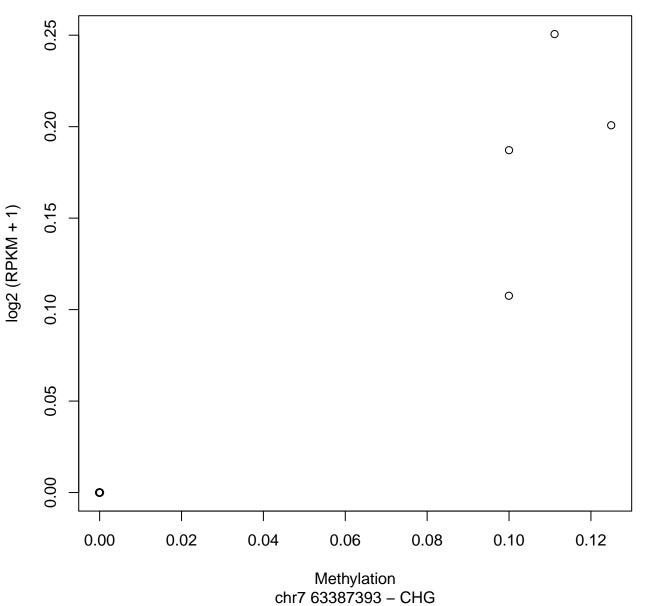
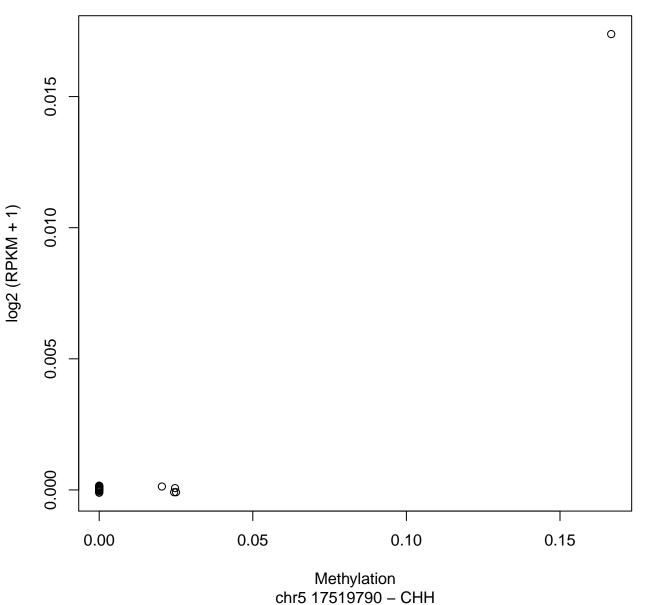
### exon ENSE00001758766.1\_1 FDR 2.94e-11 AC116366.5



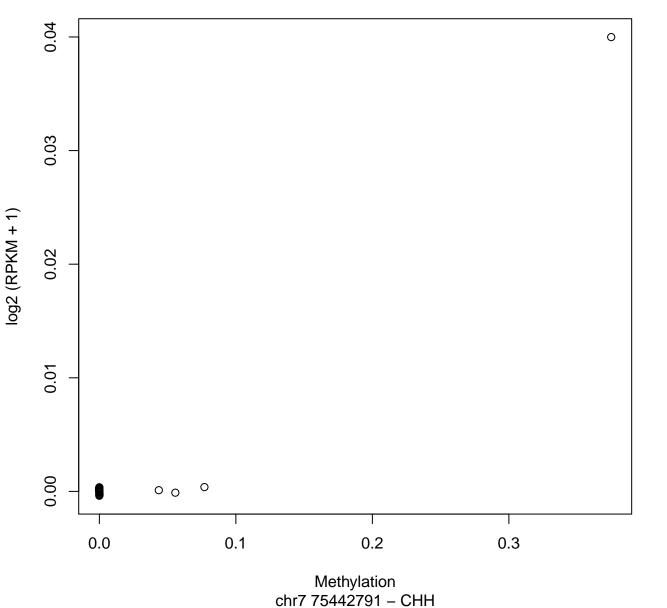
#### exon ENSE00002702765.1 FDR 2.49e-09 AC092634.2



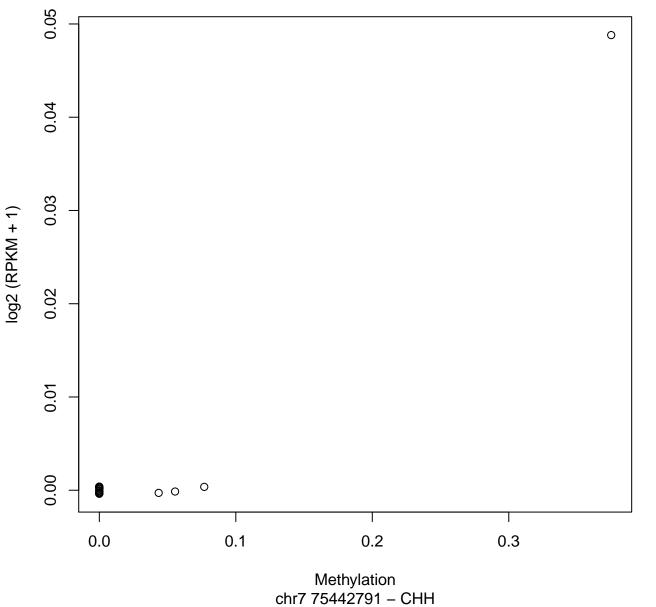
### exon ENSE00002087060.1\_1 FDR 3.5e-09 RP11-321E2.2



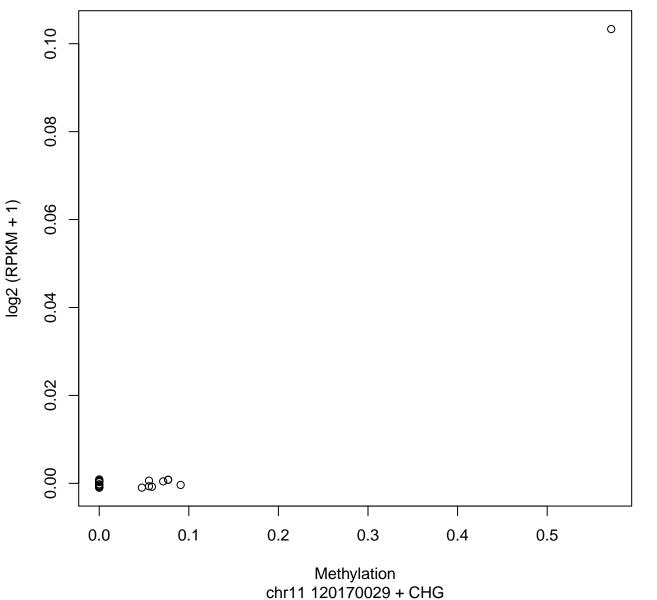
### exon ENSE00001661780.1\_1 FDR 3.91e-09 CCL24



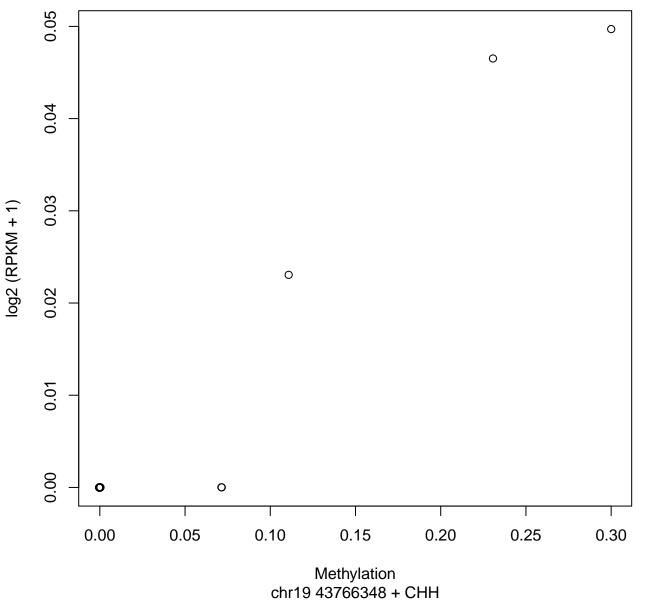
### exon ENSE00000696985.3\_1 FDR 3.91e-09 CCL24



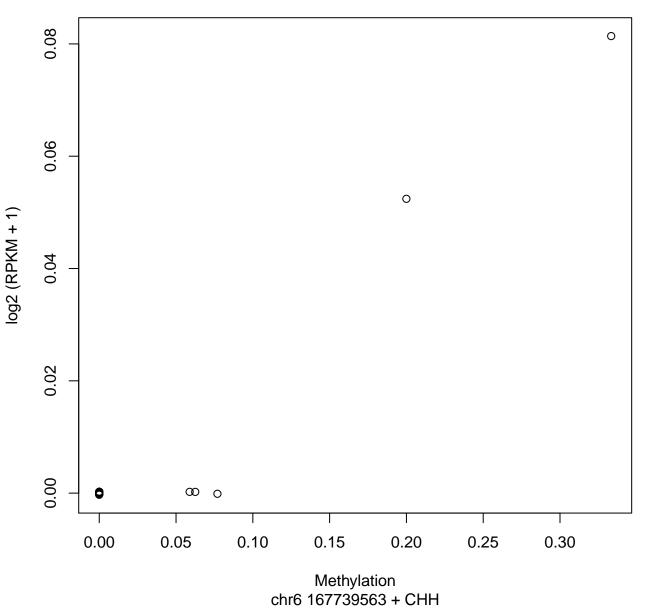
# exon ENSE00003696147.1\_1 FDR 1.16e-08 POU2F3



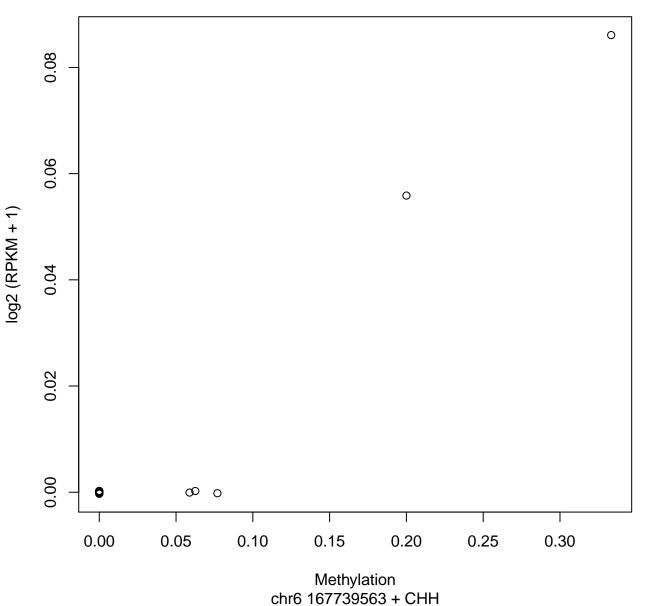
# exon ENSE00002503378.1\_1 FDR 1.58e-08 PSG9



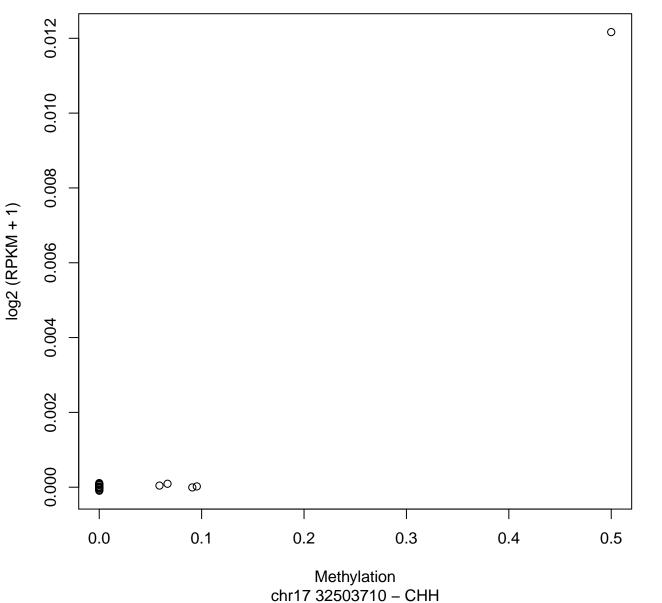
### exon ENSE00002063271.1\_1 FDR 2.5e-08 TTLL2



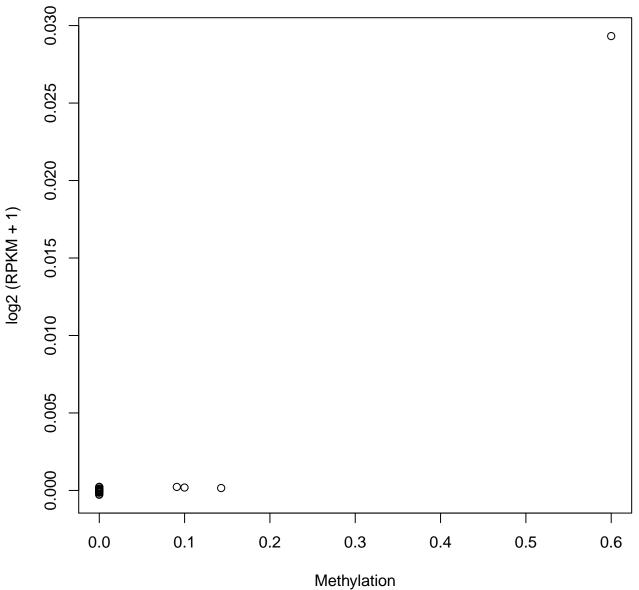
### exon ENSE00001411481.3\_1 FDR 2.5e-08 TTLL2



#### exon ENSE00002724203.1\_1 FDR 2.95e-08 RP11-215E13.2

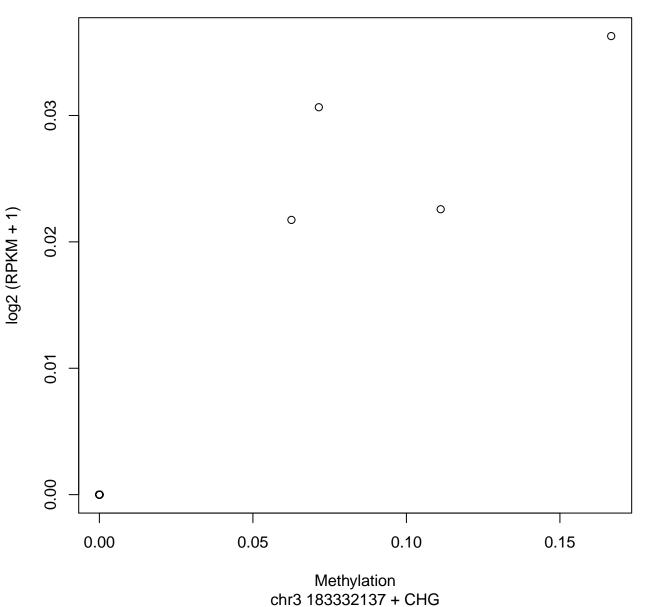


# exon ENSE00001870912.1\_1 FDR 7.94e-08 RP11-572C15.5

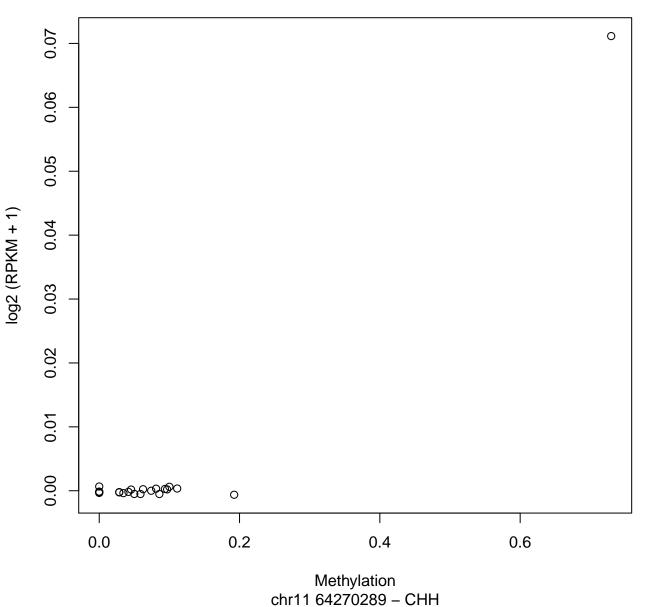


chr3 112461186 - CHH

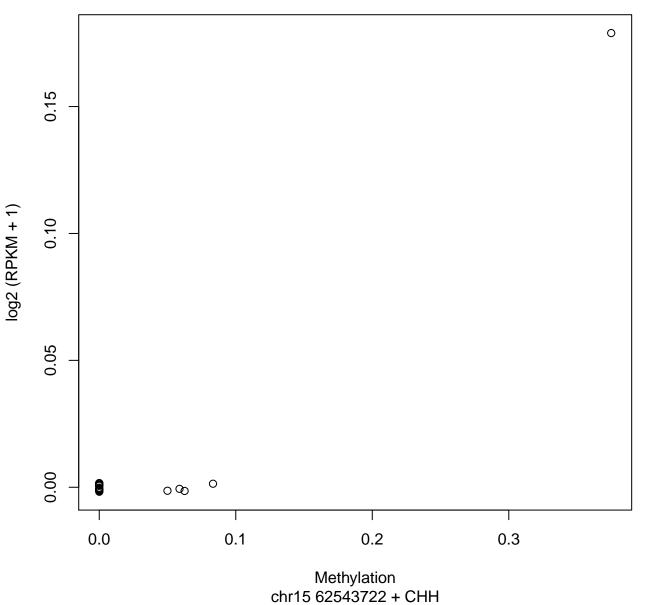
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#### exon ENSE00001638742.1\_1 FDR 1.21e-07 AP005273.1



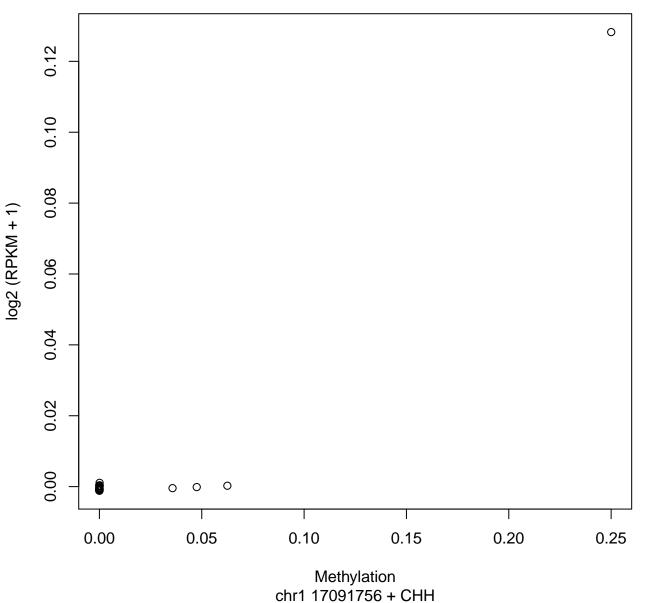
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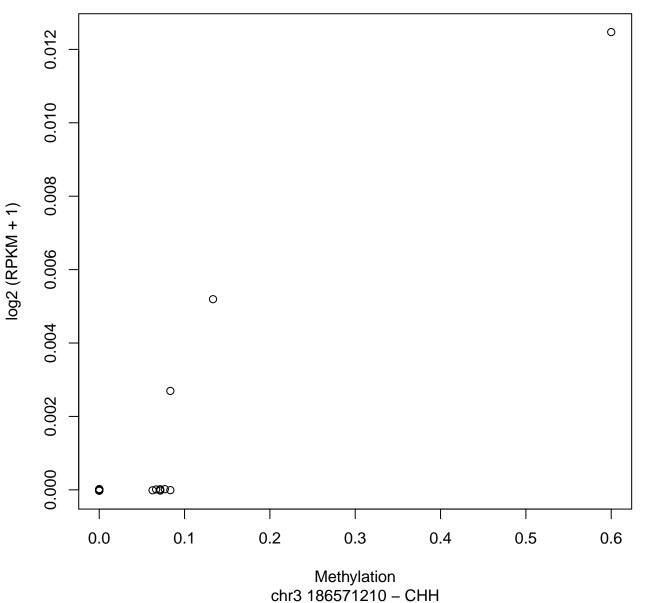
#### exon ENSE00001661971.1\_1 FDR 1.4e-07 AC008937.3 0.030 0 0.025 0.020 log2 (RPKM + 1) 0.015 0.010 0.005 0.000 0 0 0 0.0 0.2 0.4 0.6

Methylation chr5 56195584 + CHH

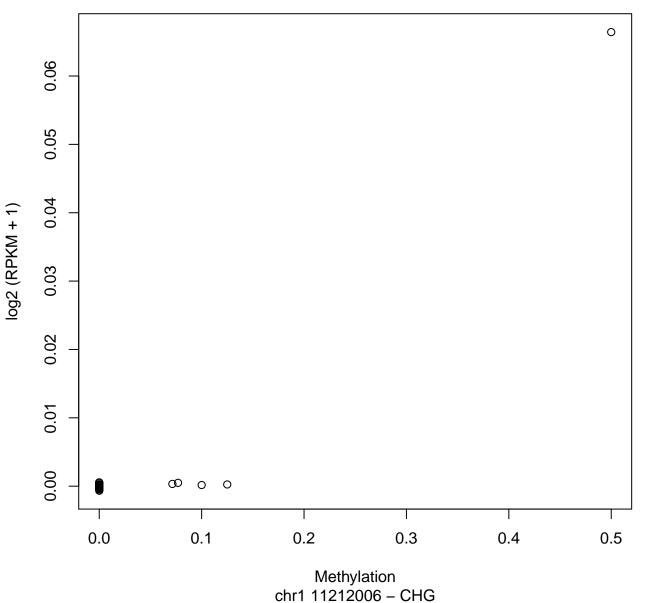
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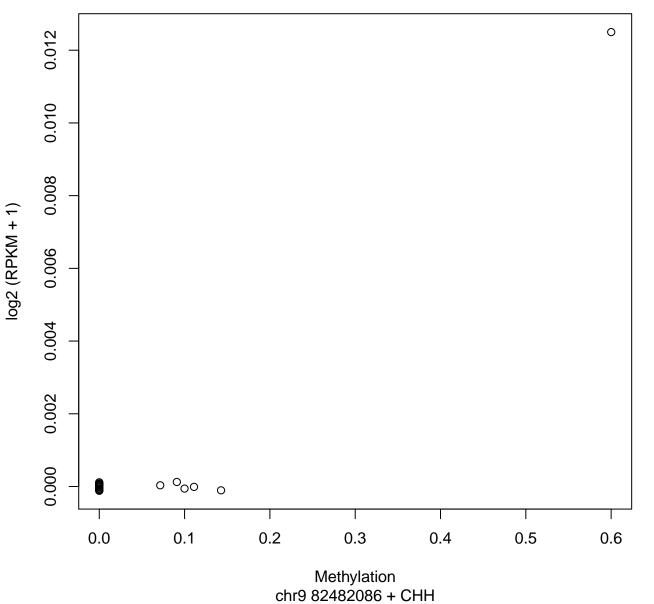
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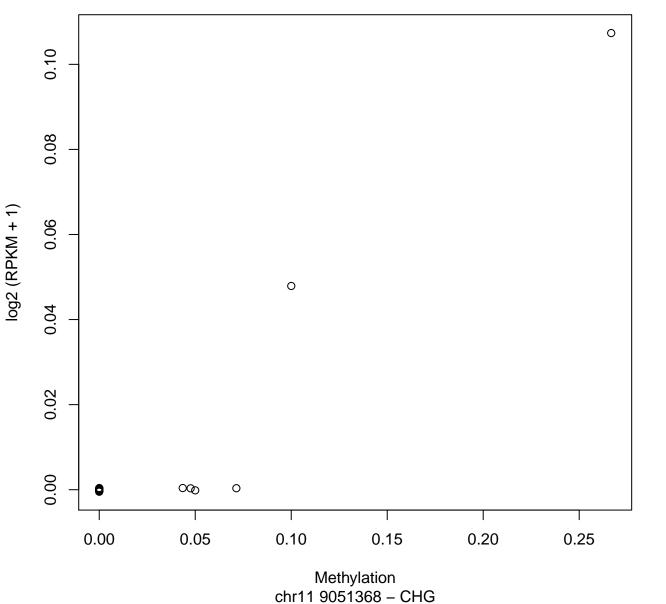
#### exon ENSE00002089554.1 FDR 8.73e-07 RNU6-537P



#### exon ENSE00003452155.1\_1 FDR 1e-06 NPAP1P6



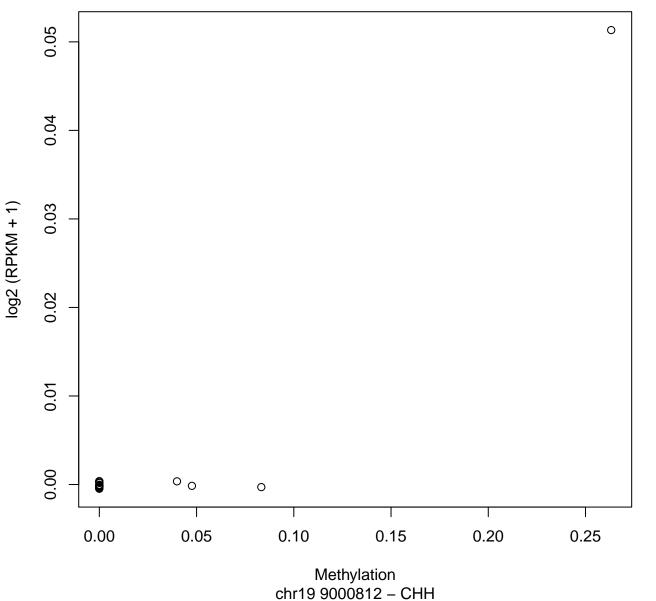
### exon ENSE00002124792.1\_1 FDR 1.94e-06 RP11-467K18.2



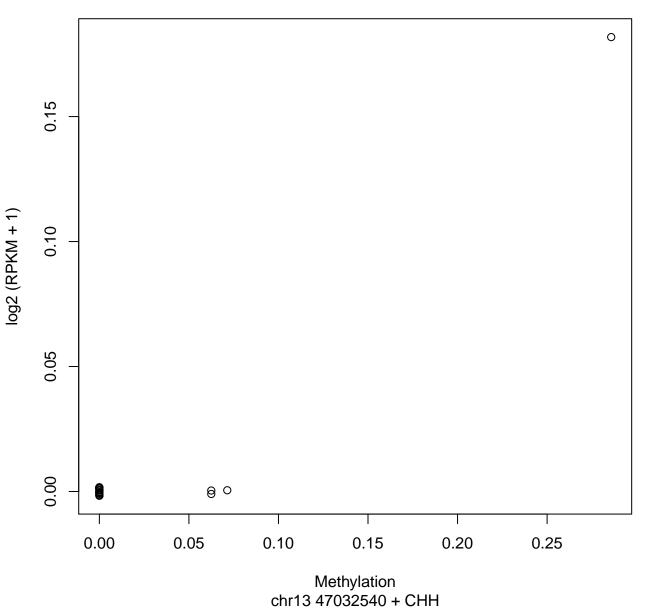
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chr17 26010905 - CHH

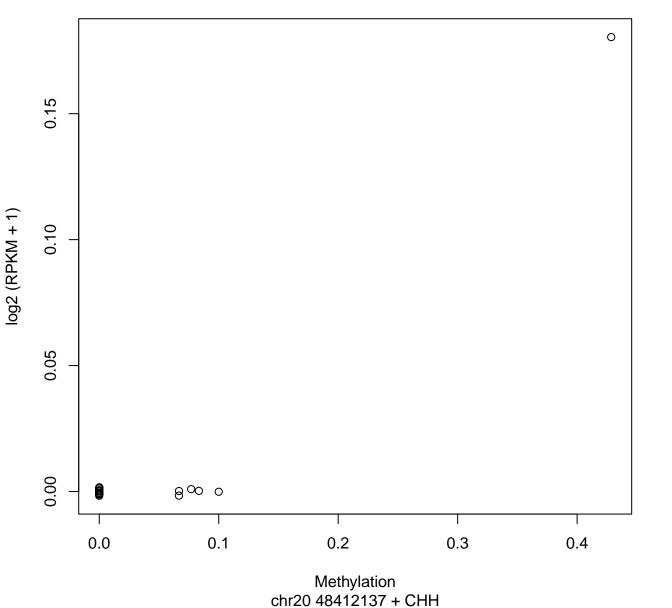
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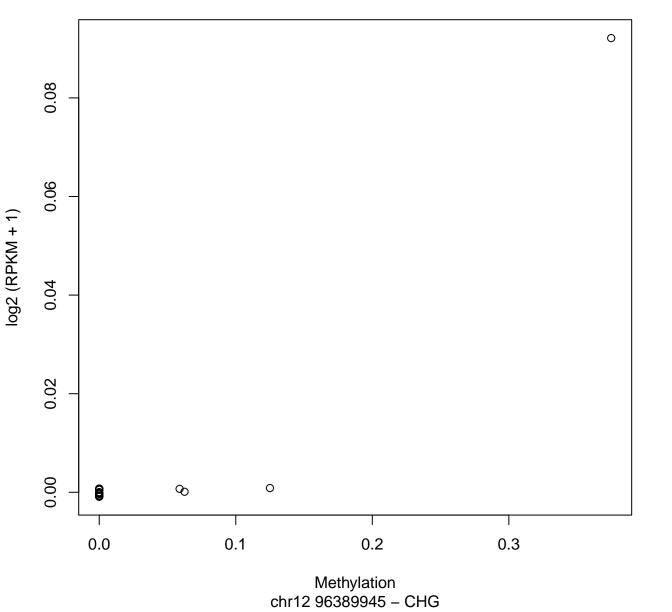
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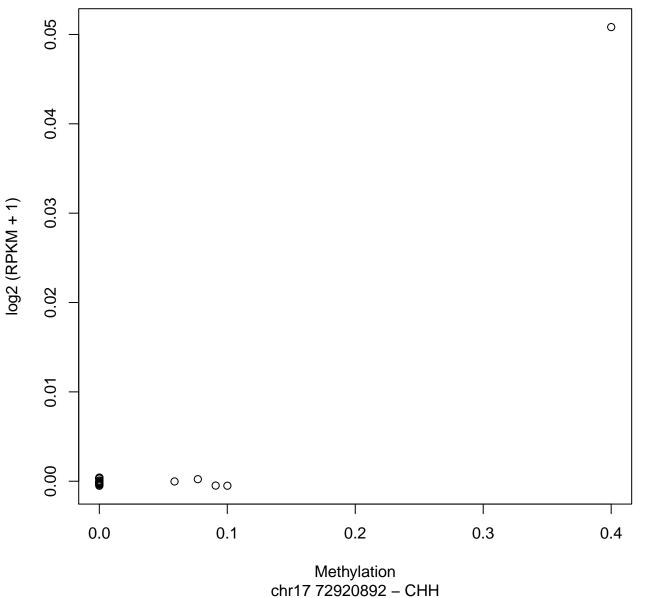
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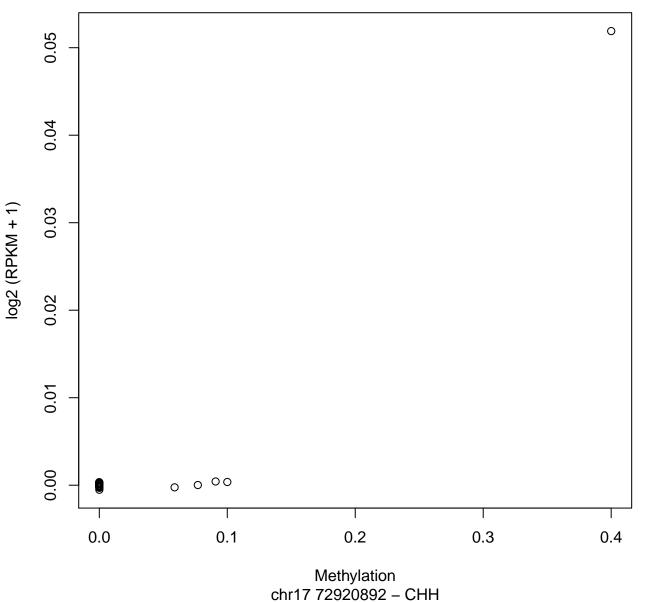
#### exon ENSE00001332536.2\_1 FDR 3.49e-06 HAL



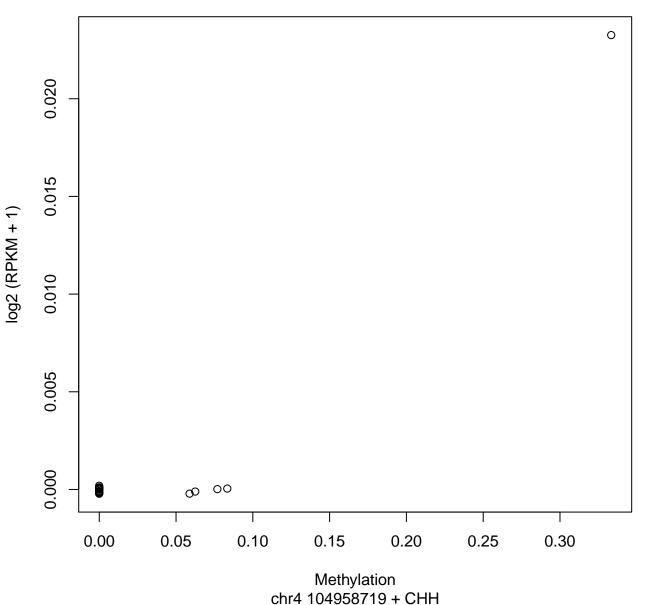
#### exon ENSE00003492138.1\_1 FDR 3.55e-06 OTOP2



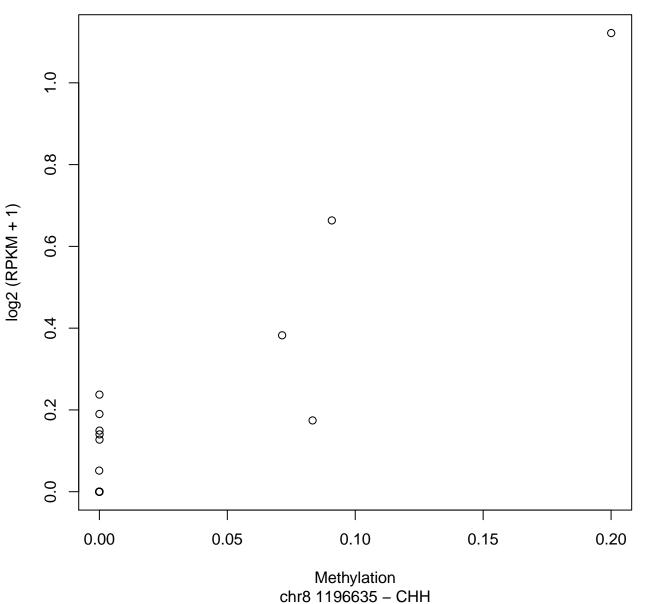
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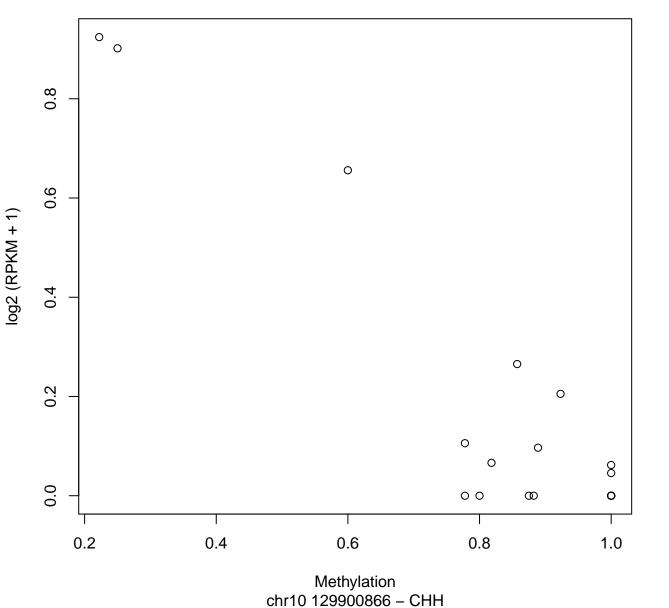
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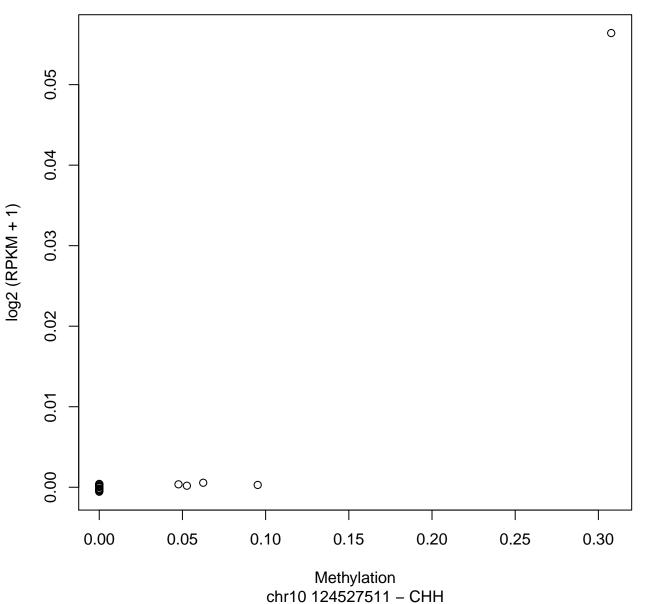
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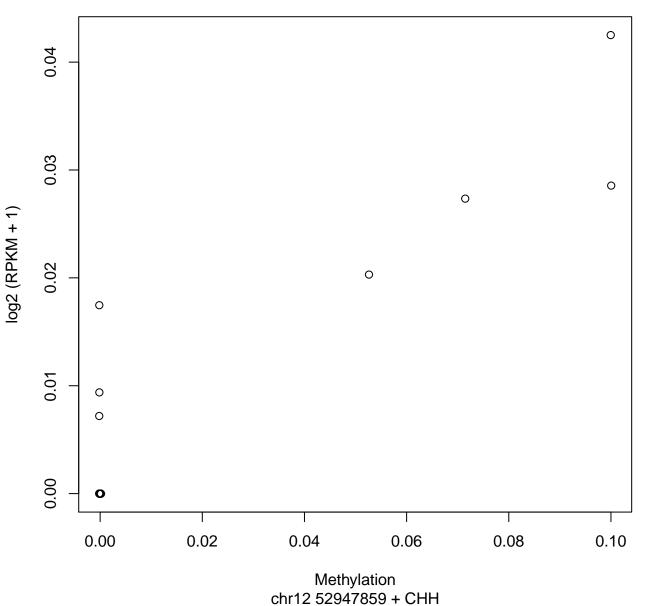
#### exon ENSE00001883513.1\_1 FDR 7.57e-06 MKI67



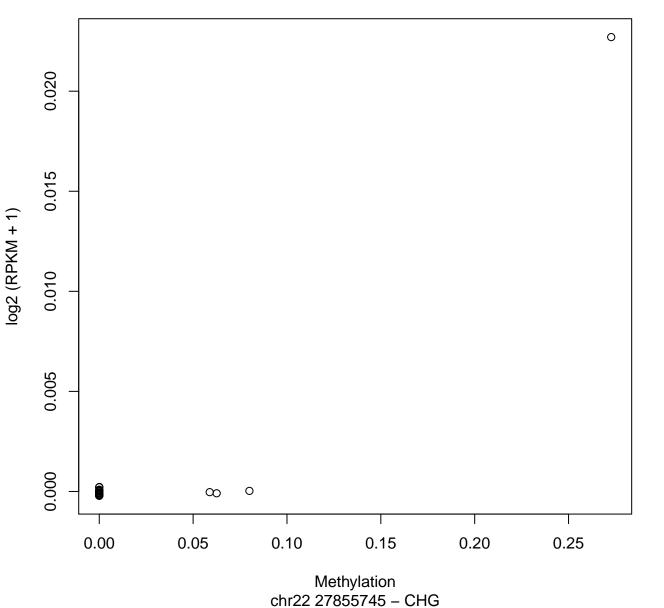
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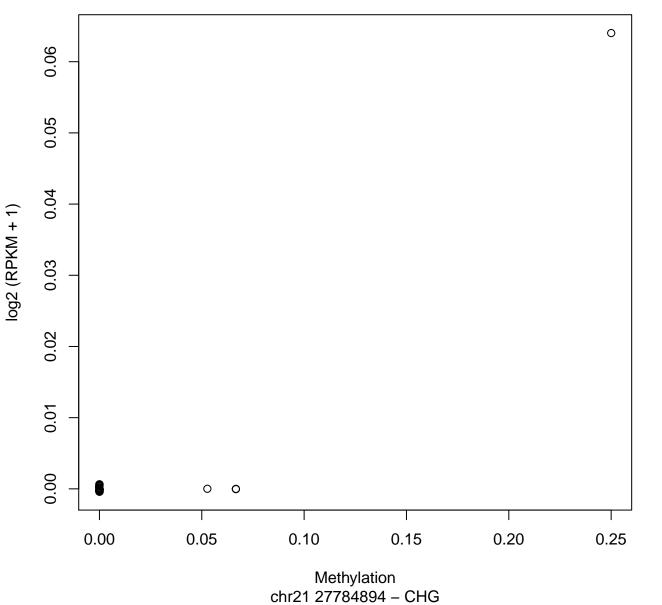
### exon ENSE00001060904.6\_1 FDR 8.68e-06 KRT71



#### exon ENSE00001621270.1\_1 FDR 8.89e-06 RP11-46E17.6



#### exon ENSE00001675911.1\_1 FDR 9.38e-06 AP001596.6

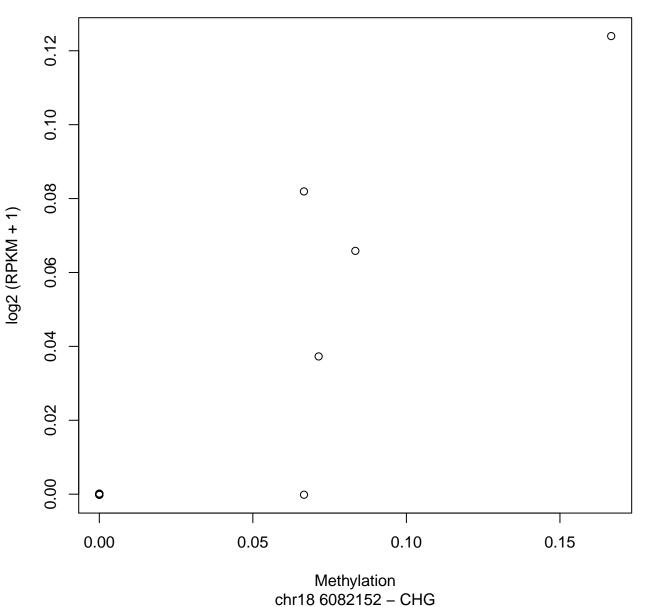


#### RP5-859D4.3 0.035 0 0.030 0.025 0.020 log2 (RPKM + 1) 0.015 0.010 0.005 0.000 တတ 0 0.0 0.1 0.2 0.3 0.4 Methylation

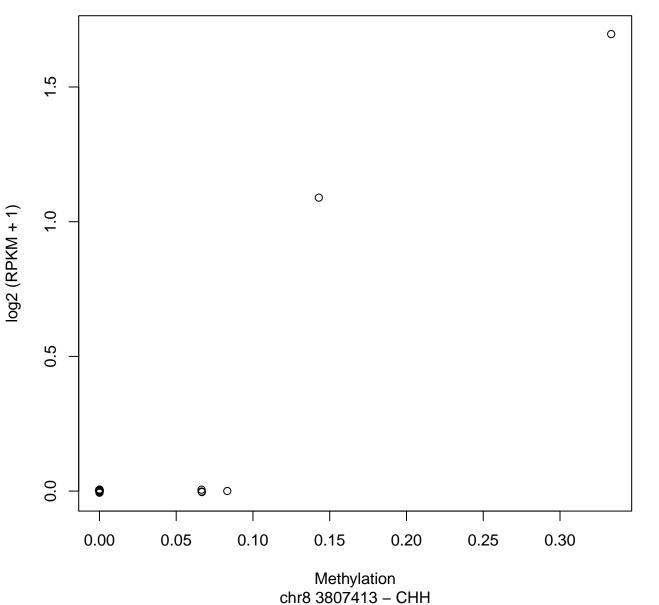
chr20 6711600 - CHH

exon ENSE00001594390.1\_1 FDR 1.11e-05

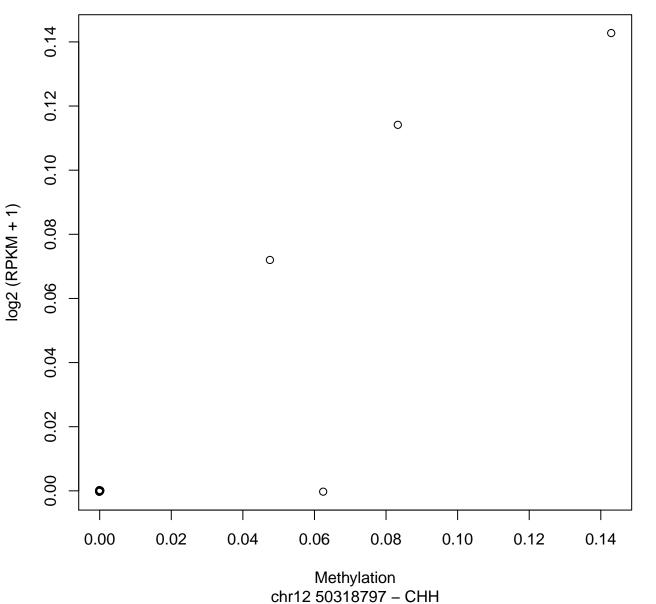
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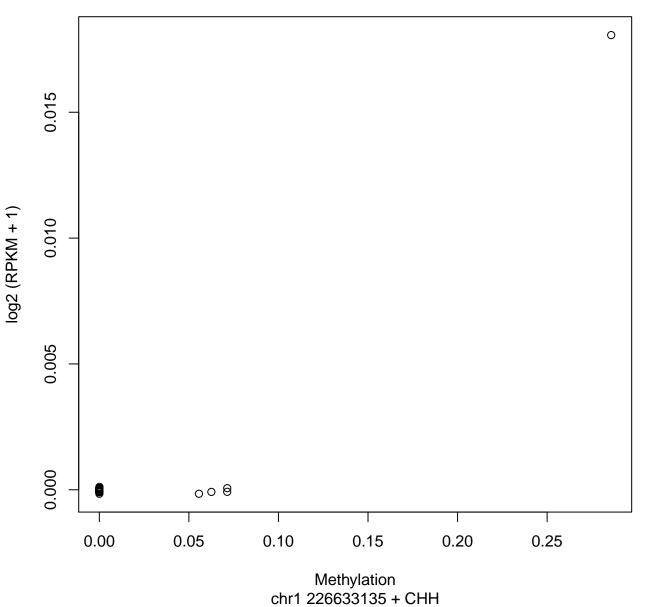
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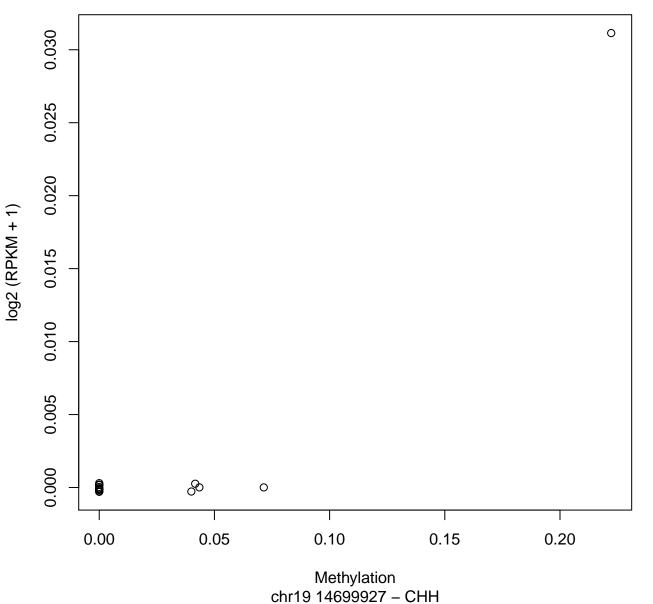
## exon ENSE00002368058.1\_1 FDR 1.61e-05 RP11-70F11.8



#### exon ENSE00001591168.1 FDR 1.8e-05 RN7SKP165



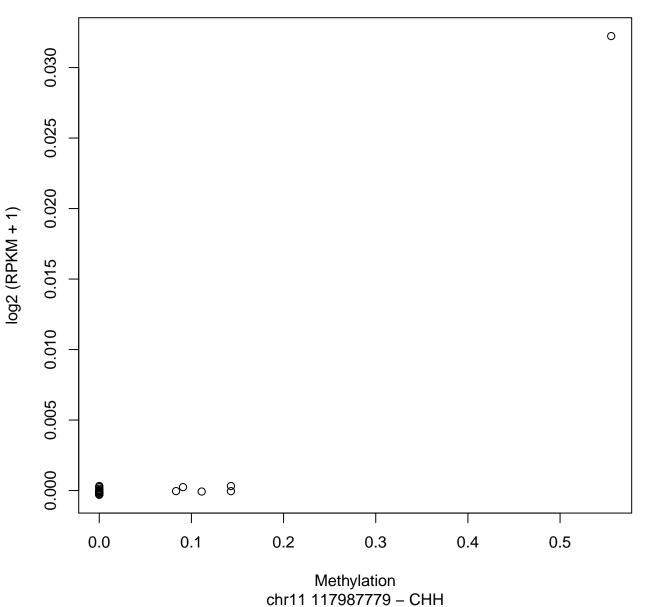
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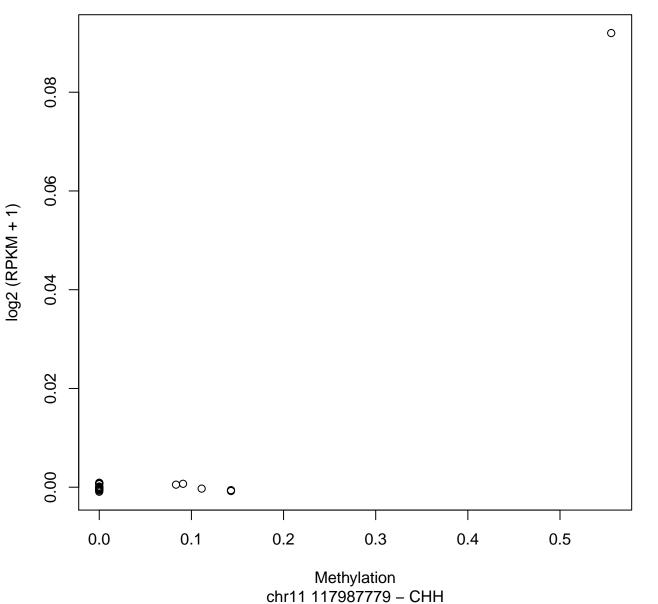
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Methylation chr19 46259725 – CHG

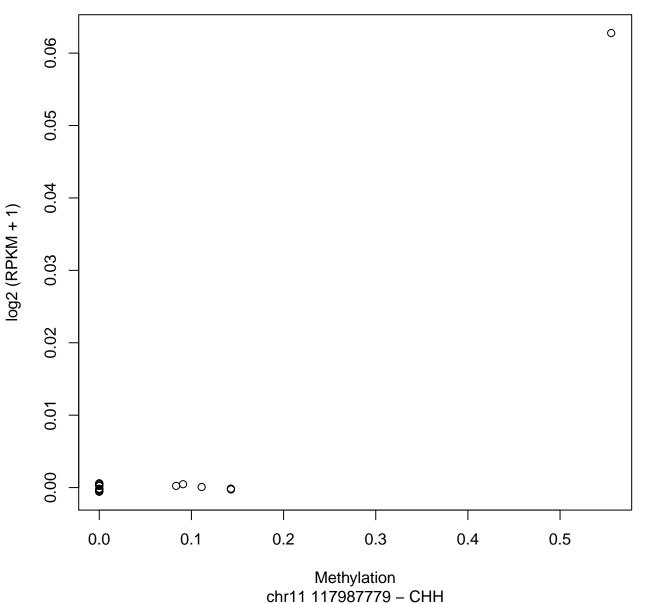
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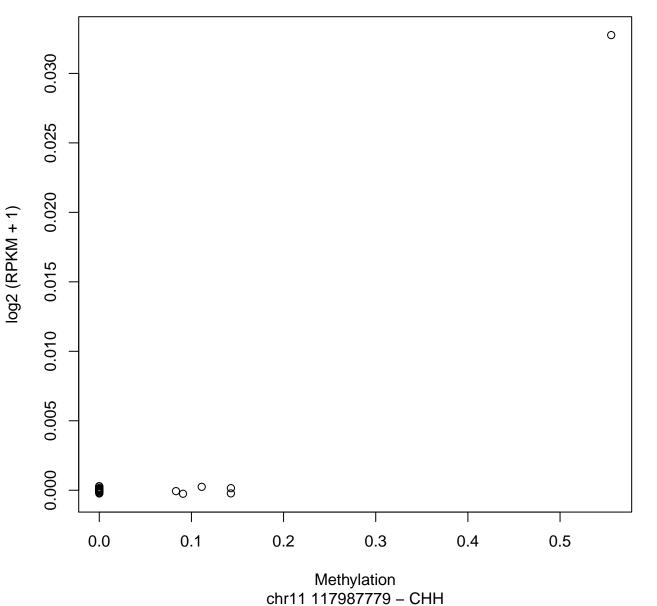
## exon ENSE00002120153.1\_1 FDR 1.99e-05 TMPRSS4



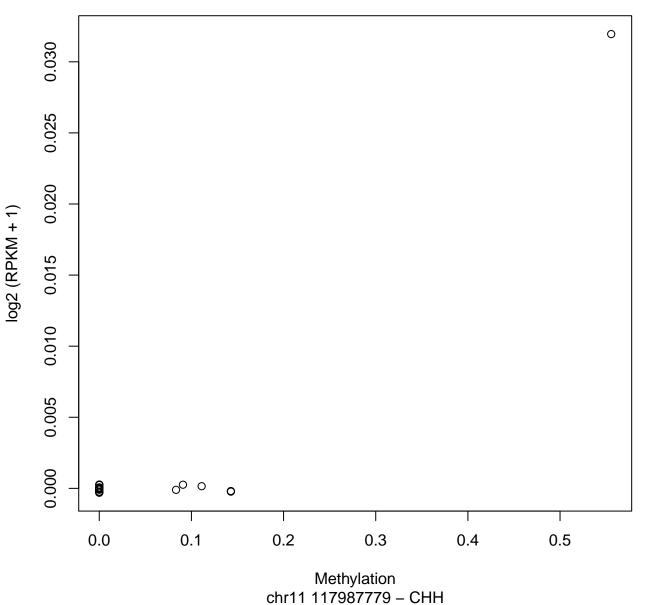
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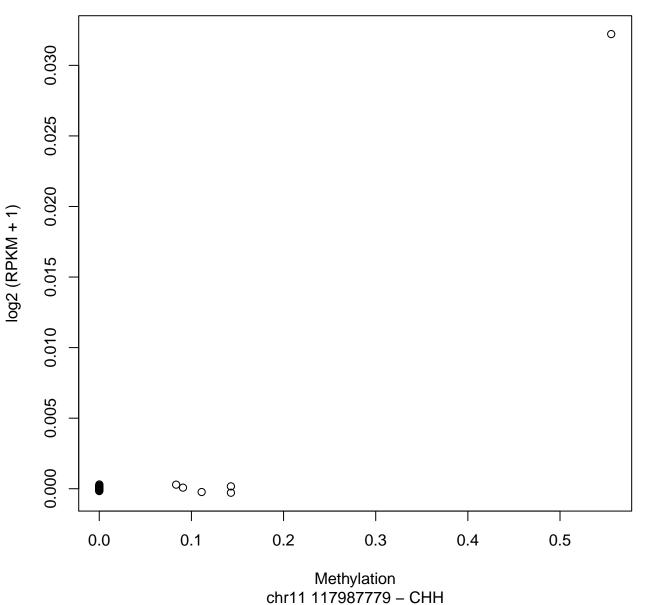
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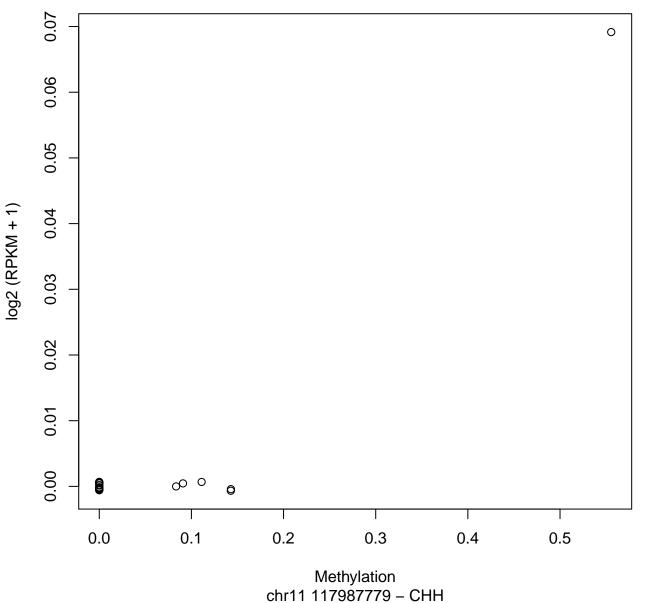
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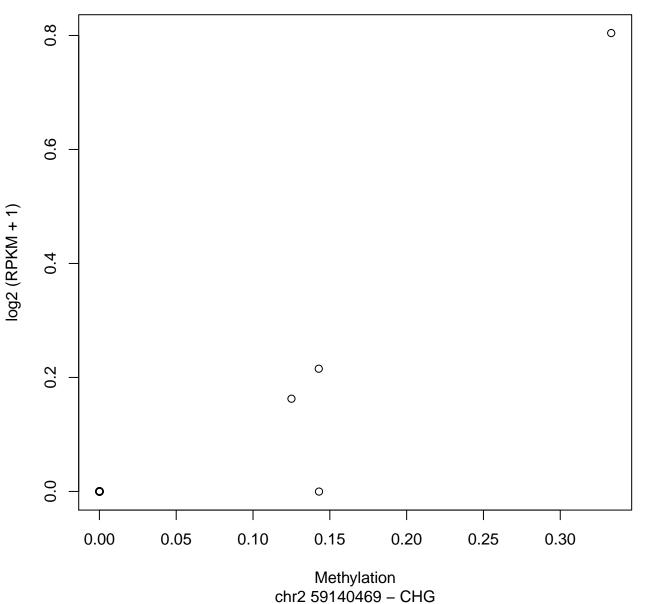
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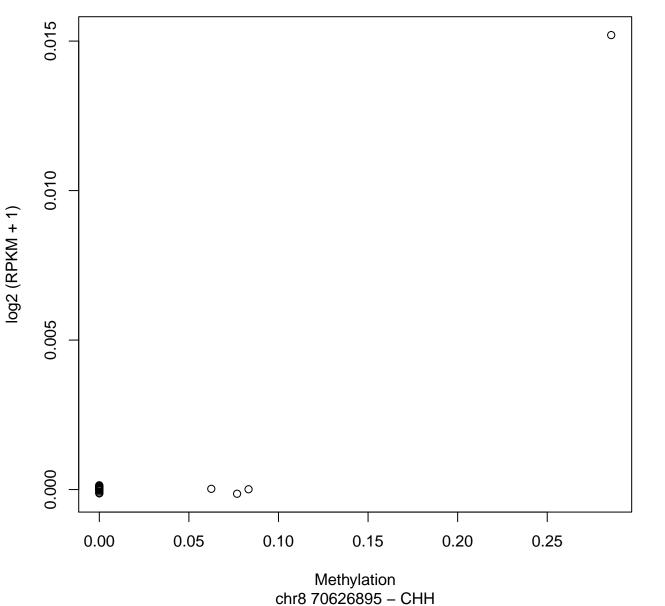
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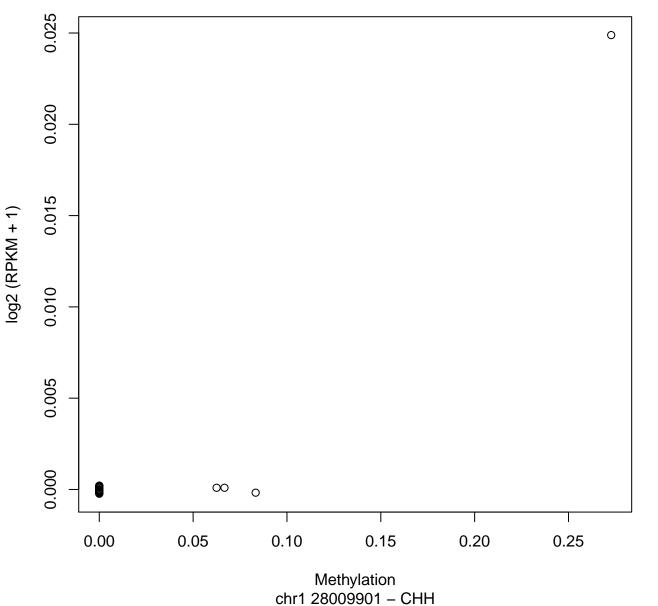
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#### exon ENSE00002173303.1\_1 FDR 2.06e-05 RP11-102F4.2



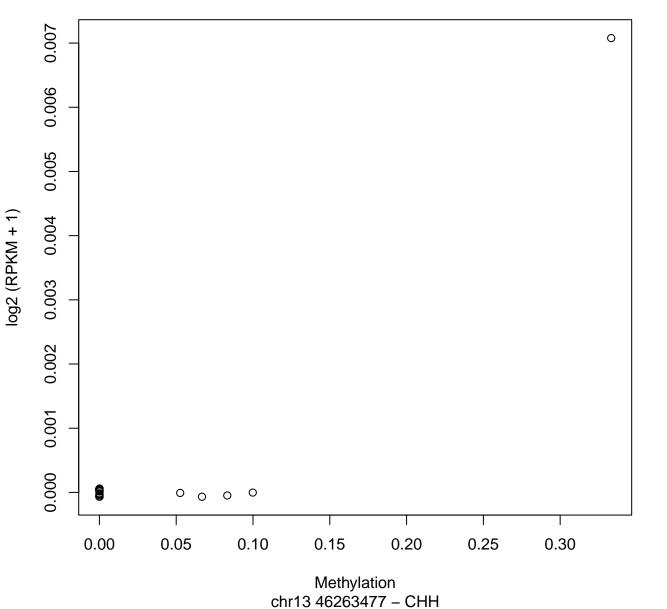
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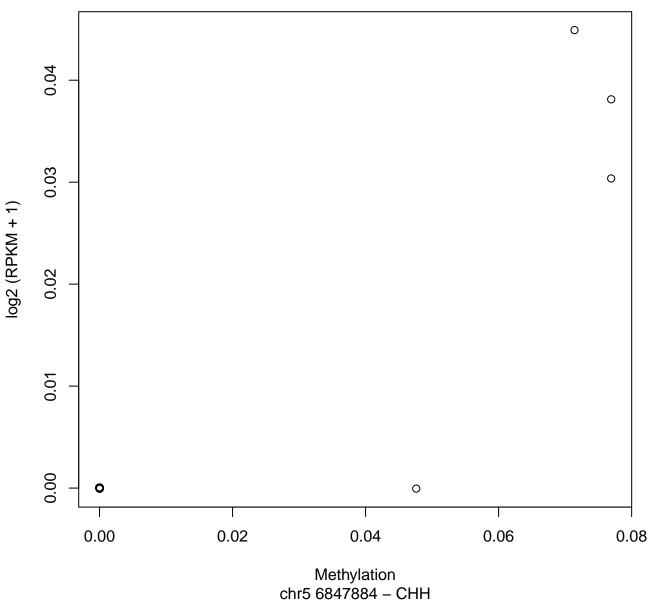
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chr17 25748370 - CHG

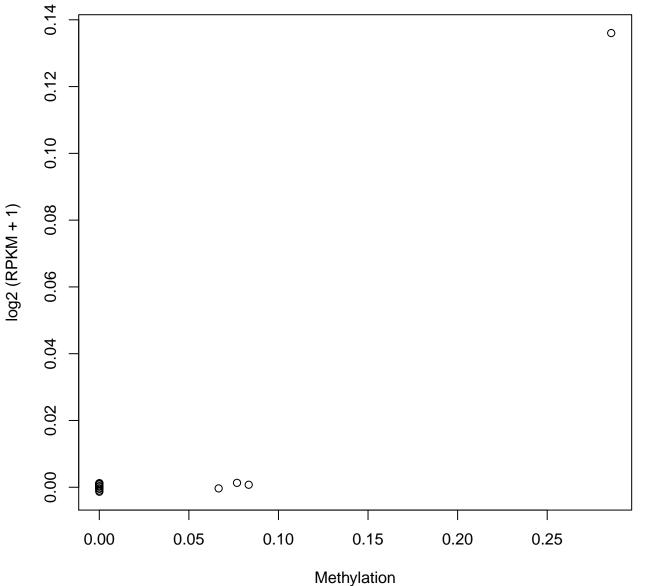
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## exon ENSE00001438136.1 FDR 2.58e-05 RN7SKP79

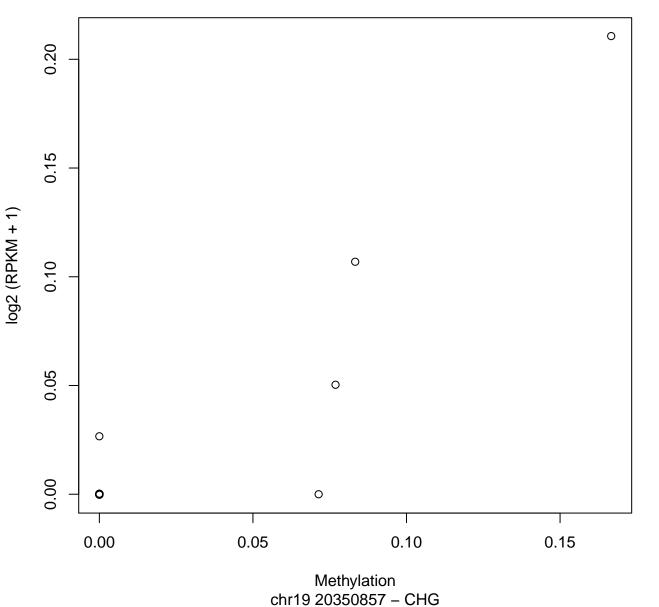


# exon ENSE00002272282.1\_1 FDR 2.63e-05 MUC16

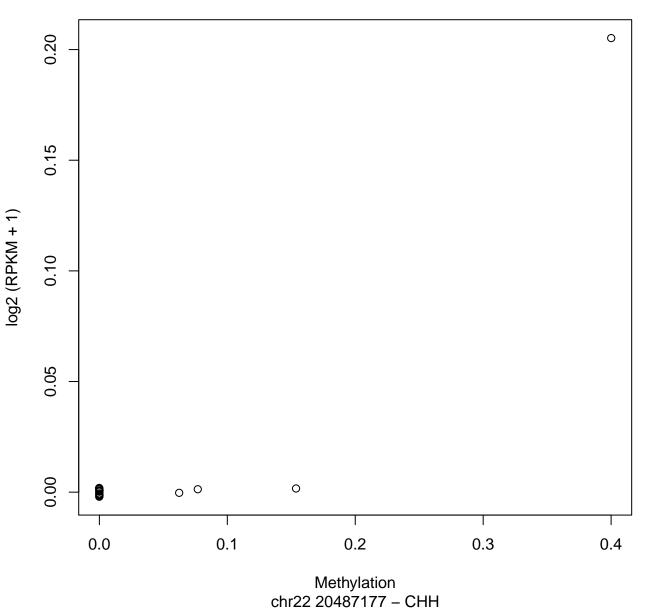


chr19 9005193 + CHG

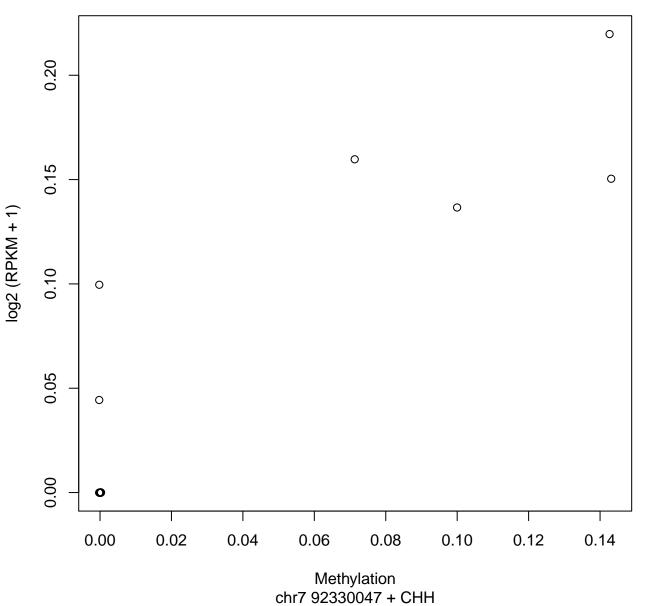
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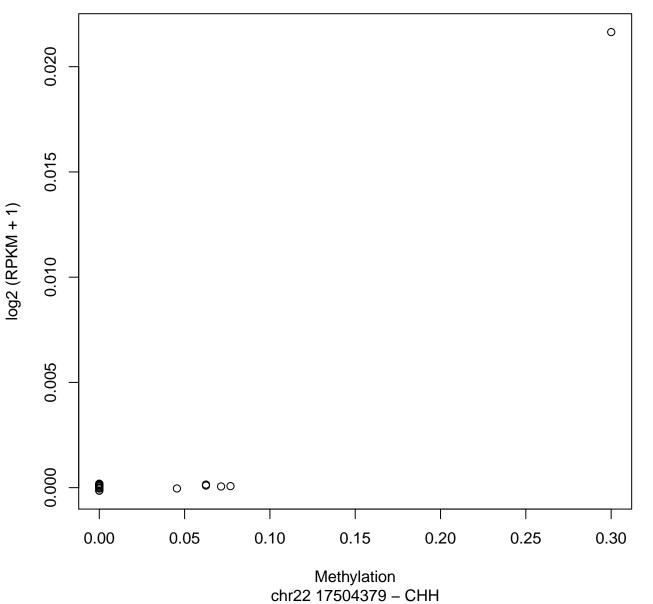
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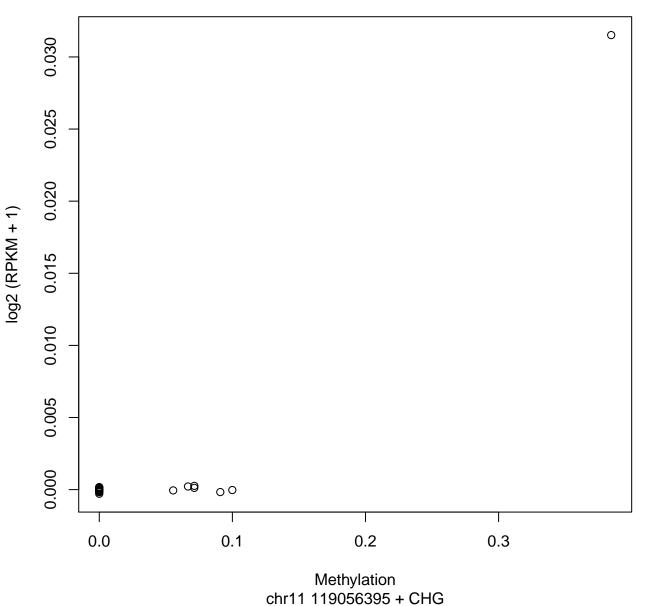
### exon ENSE00001809019.1 FDR 2.73e-05 RNU6-10P



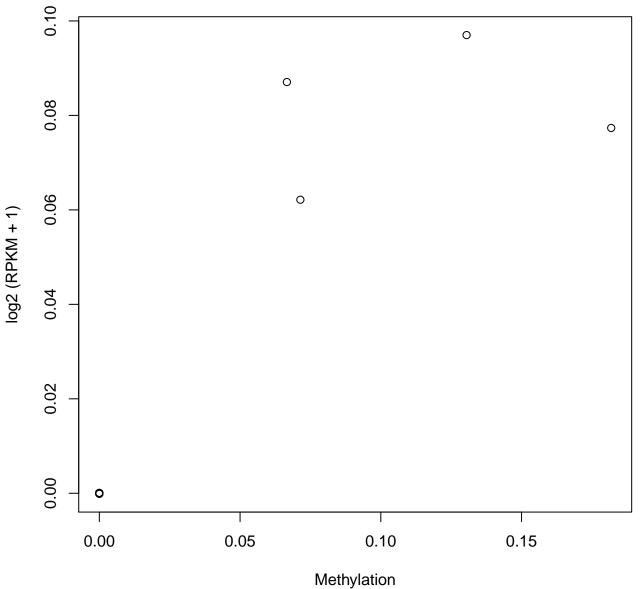
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# exon ENSE00002178197.1\_1 FDR 2.81e-05 PDZD3

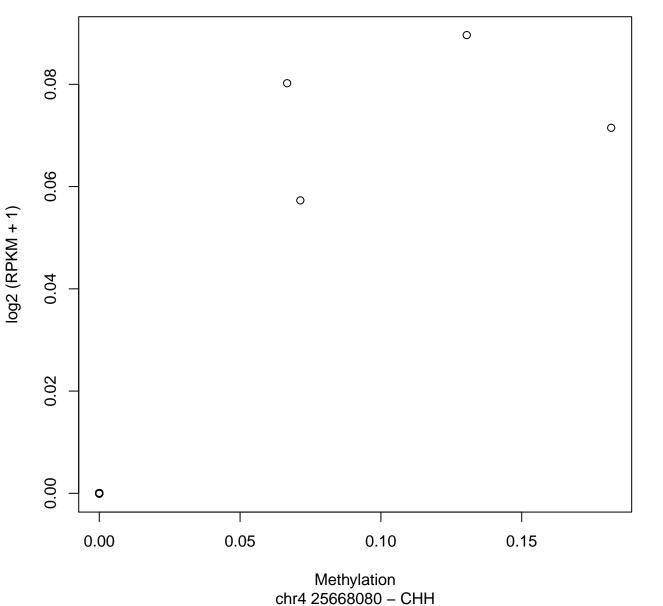


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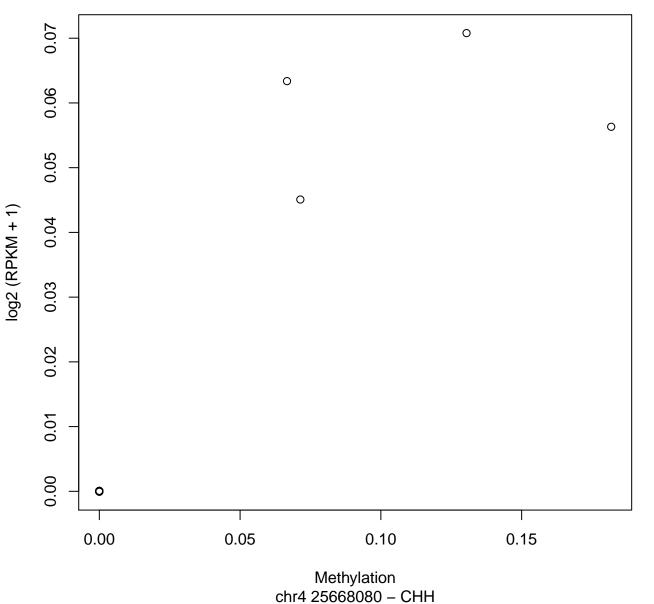


chr4 25668080 - CHH

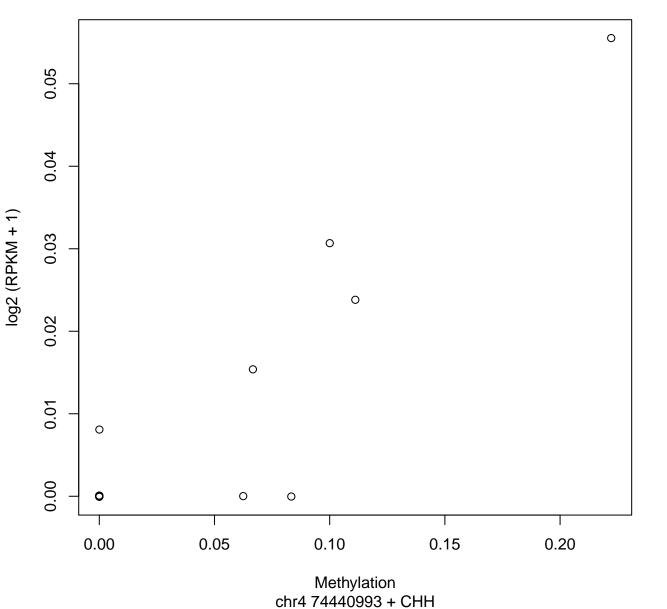
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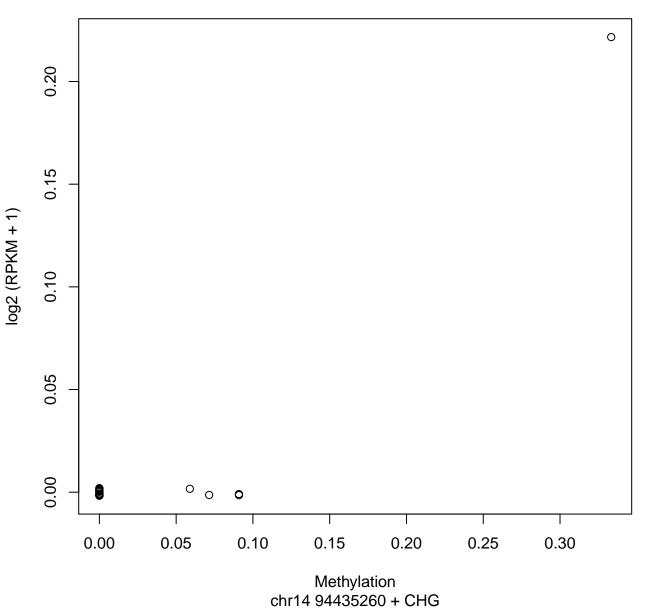
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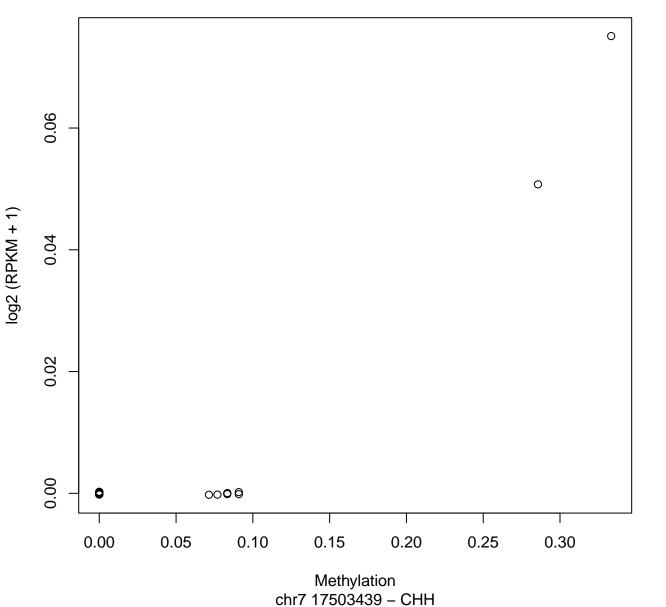
# exon ENSE00001902229.1\_1 FDR 2.94e-05 RASSF6



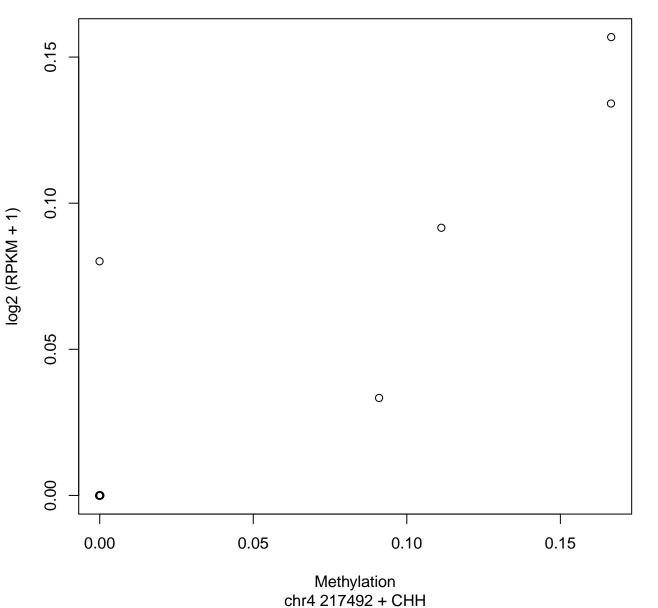
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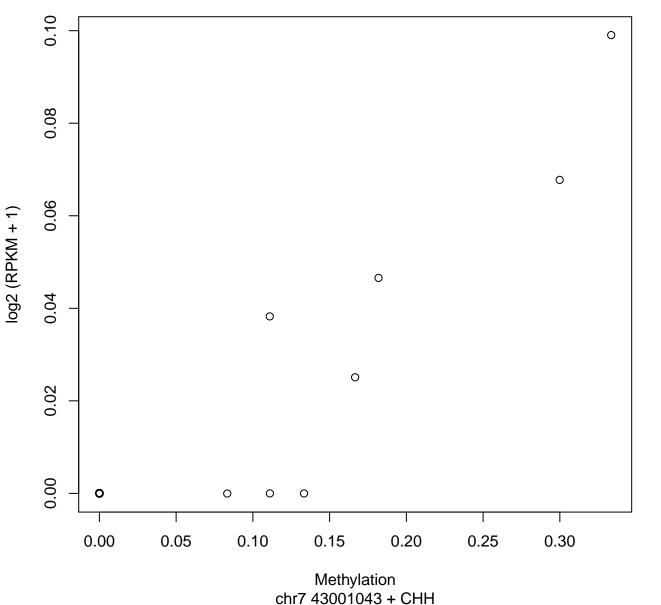
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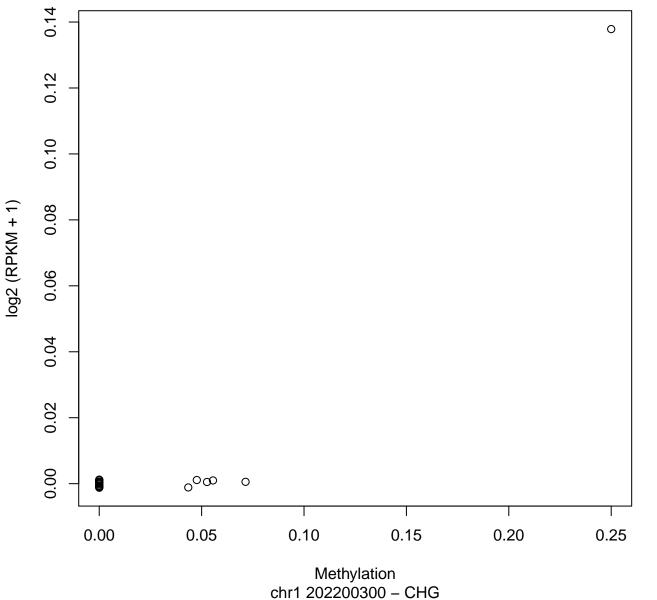
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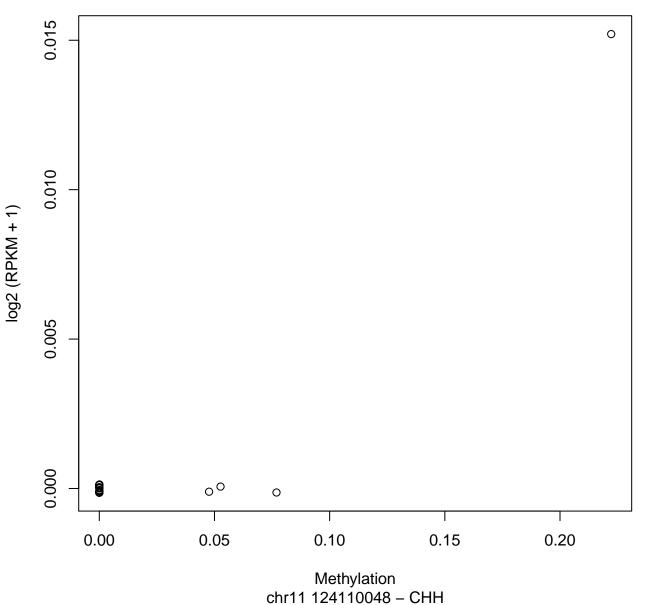
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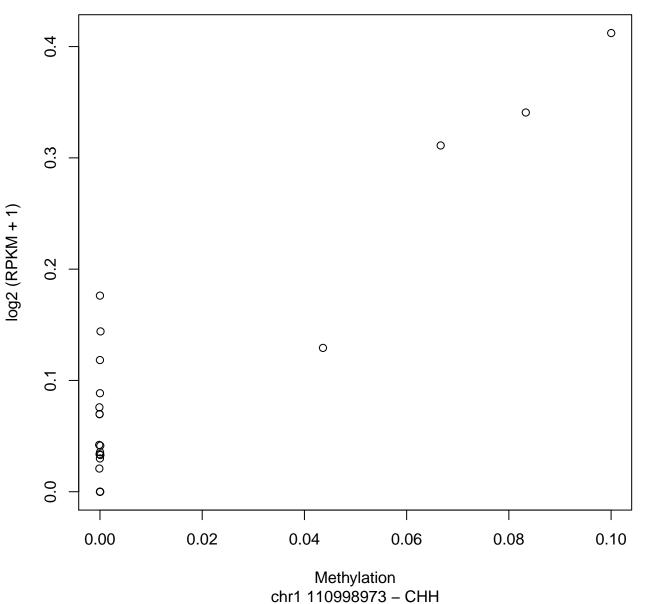
# exon ENSE00001807530.1 FDR 3.89e-05 snoU13



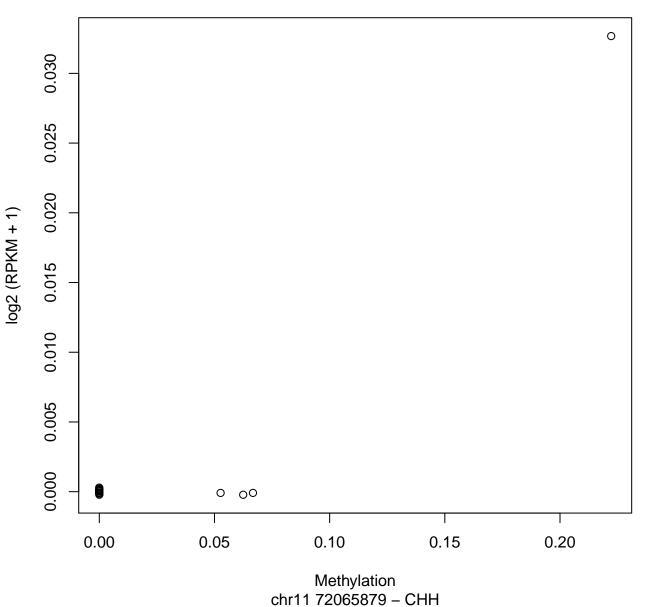
#### exon ENSE00003753782.1\_1 FDR 3.91e-05 OR8G7P



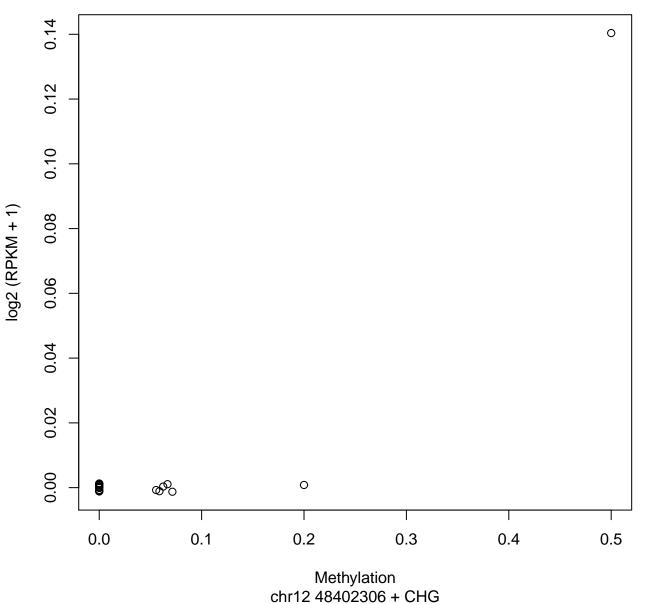
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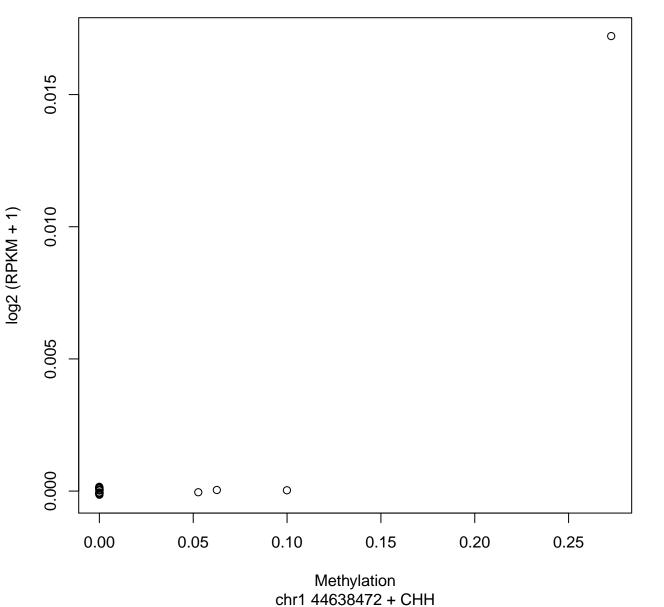
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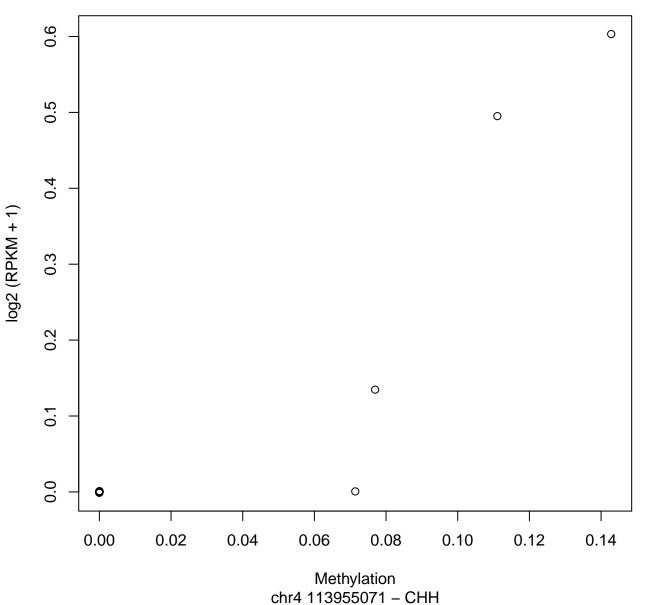
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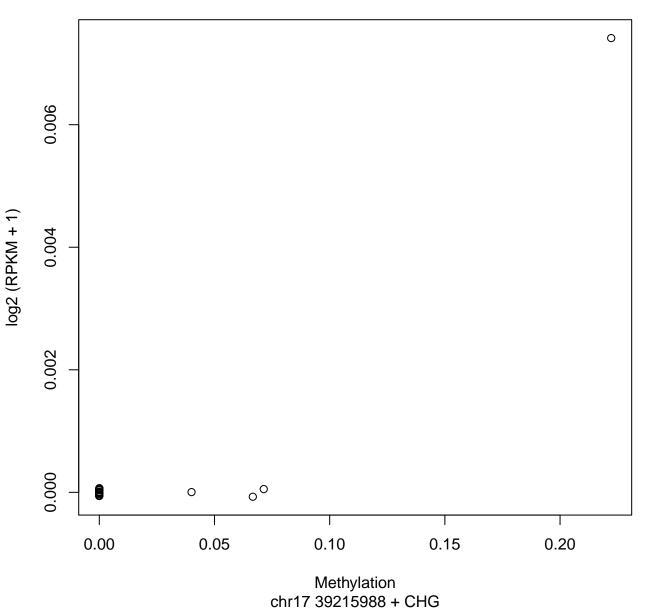
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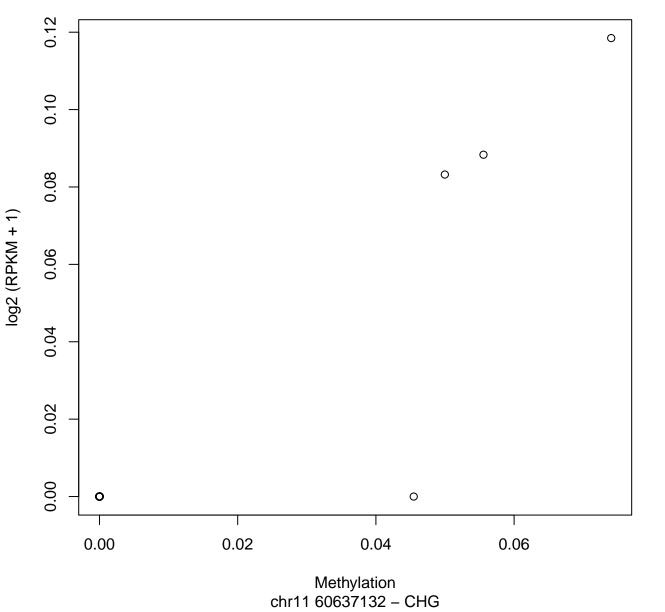
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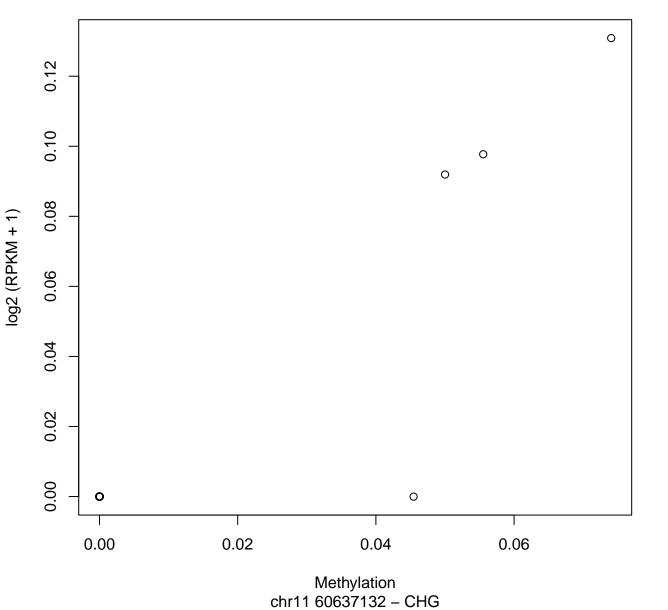
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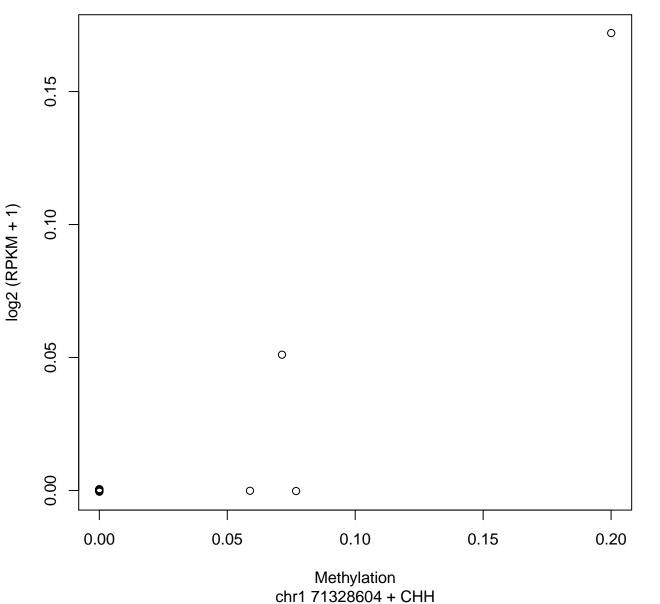
### exon ENSE00000990965.1\_1 FDR 4.54e-05 ZP1



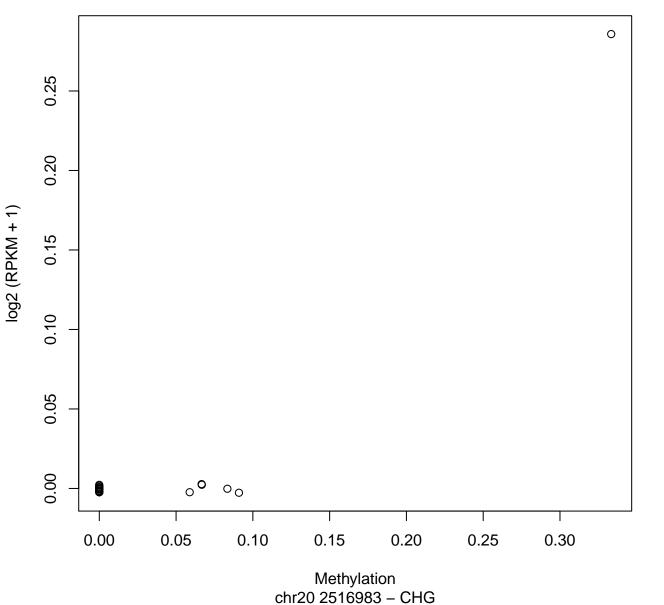
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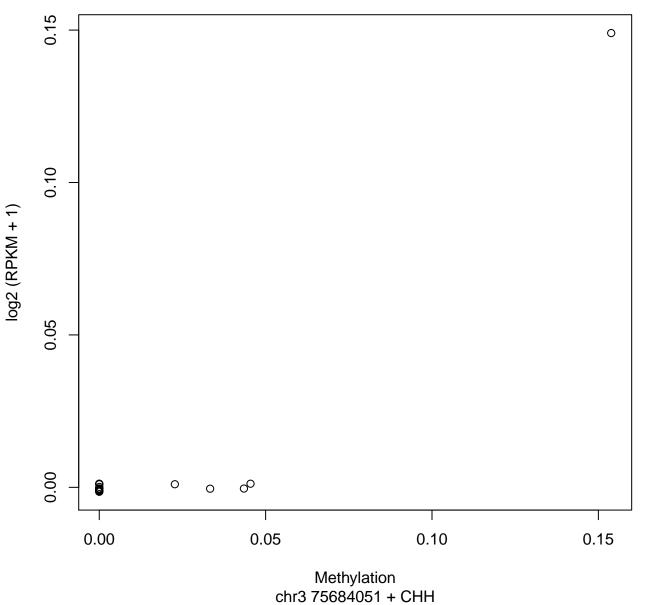
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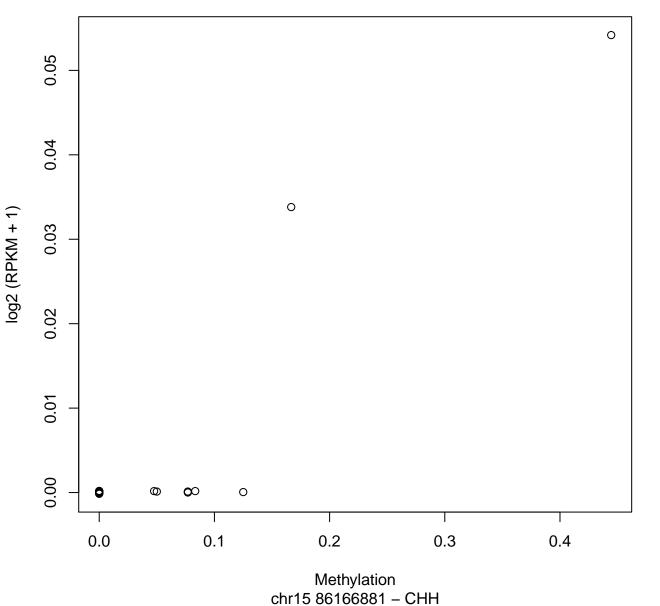
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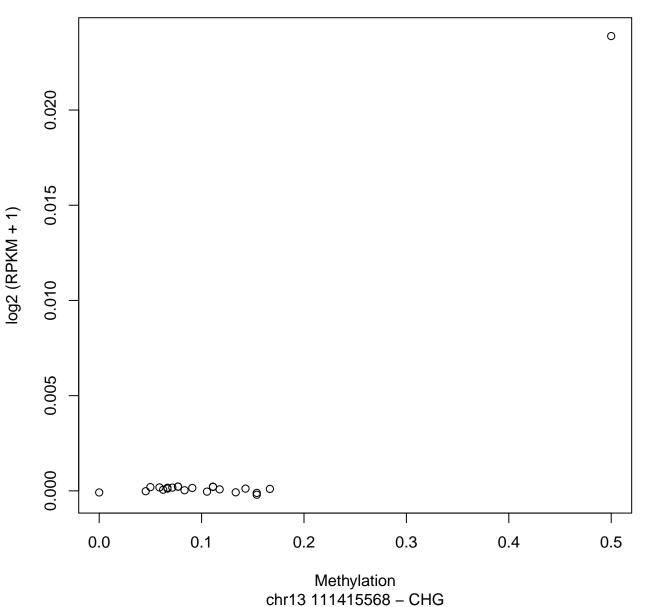
# exon ENSE00003780099.1\_1 FDR 4.71e-05 CLUHP10



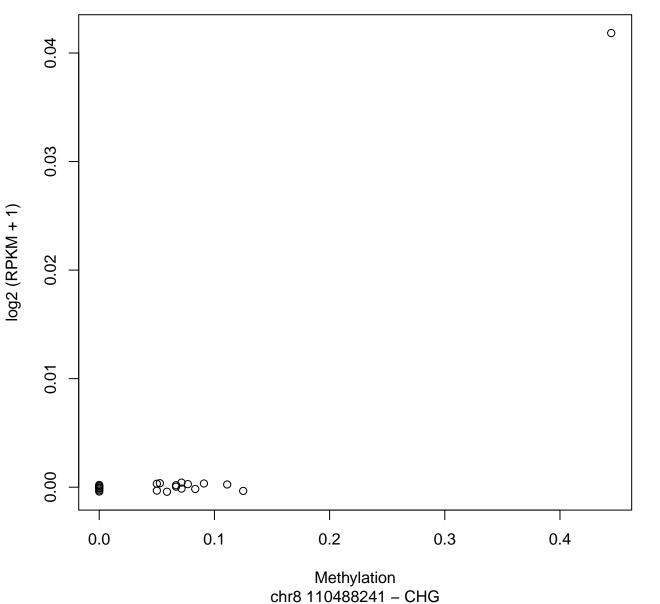
#### exon ENSE00002538902.1\_1 FDR 5.03e-05 RP11-815J21.3



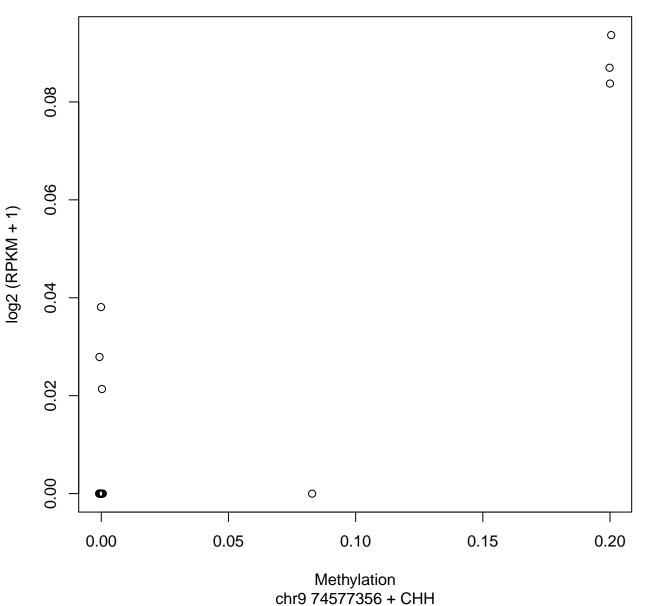
#### exon ENSE00001699357.1\_1 FDR 5.18e-05 RPL21P107



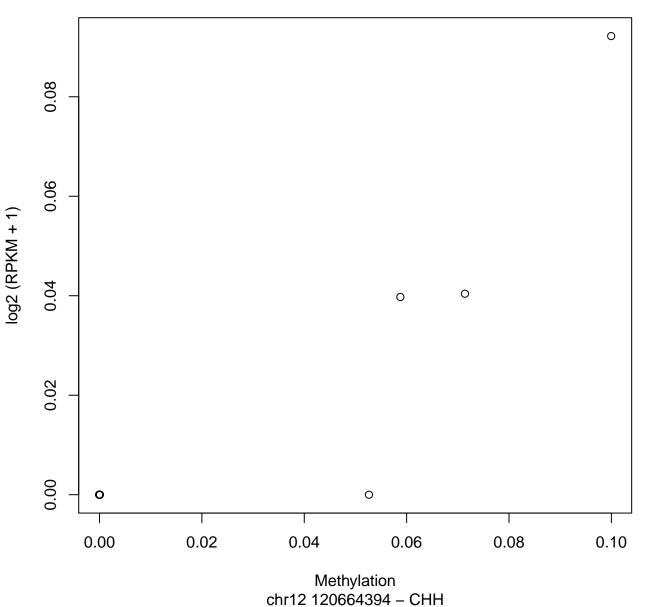
# exon ENSE00001477397.1\_1 FDR 5.3e-05 PKHD1L1



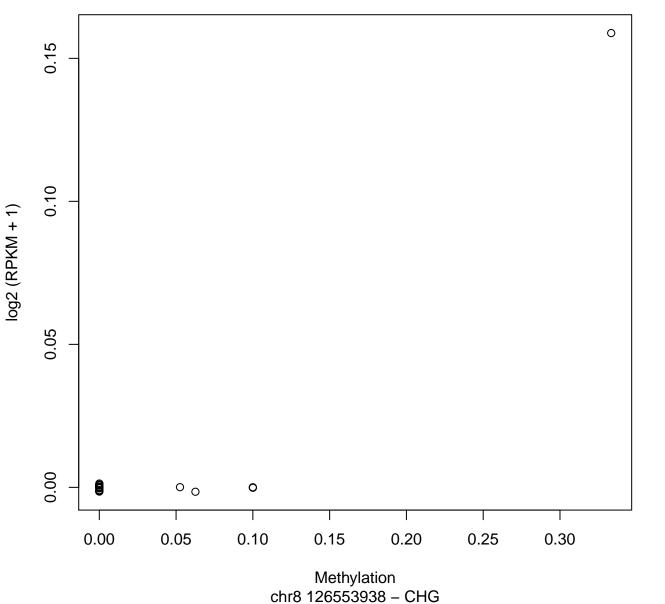
#### exon ENSE00001425316.1 FDR 5.35e-05 AL138751.1



### exon ENSE00002328773.1\_1 FDR 5.38e-05 PXN



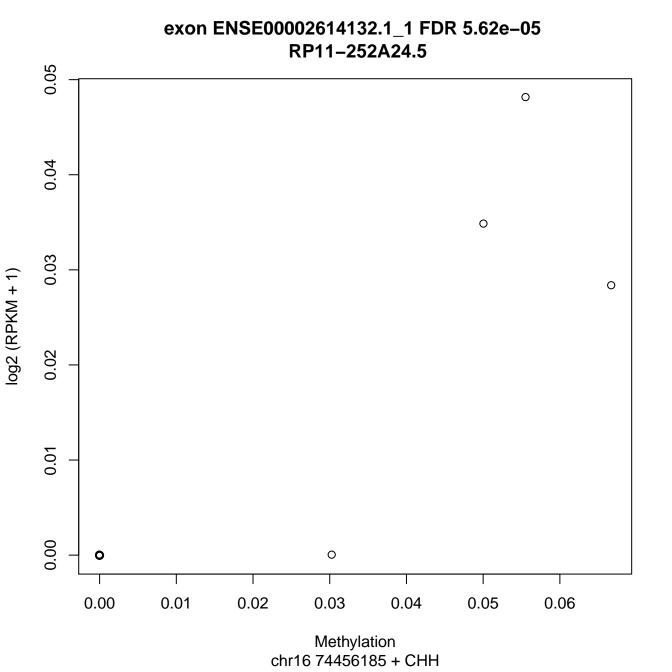
#### exon ENSE00002125460.1\_1 FDR 5.48e-05 RP11-136O12.2



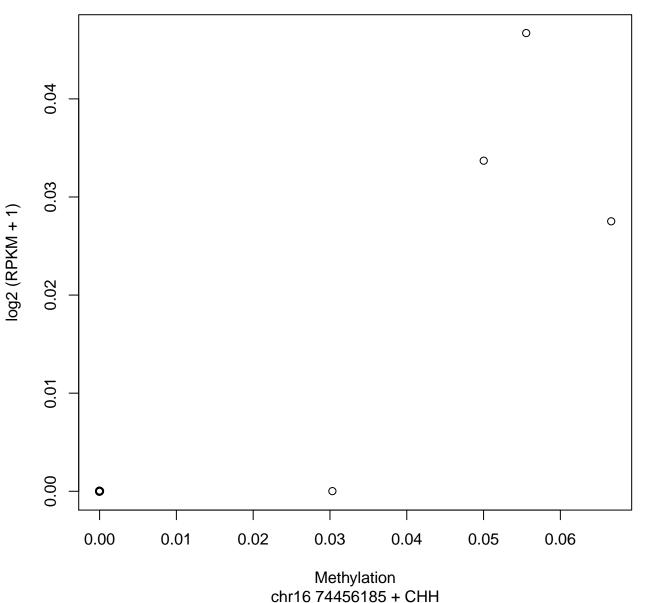
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chr3 137744061 + CHG

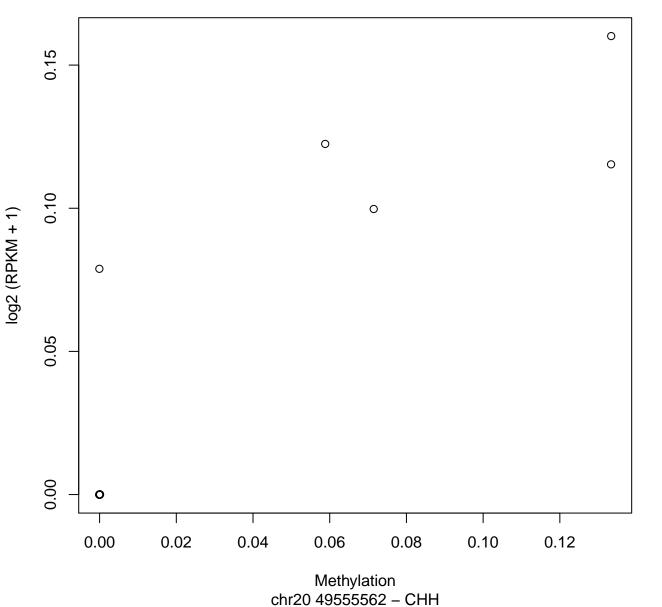
exon ENSE00000778672.1\_1 FDR 5.6e-05



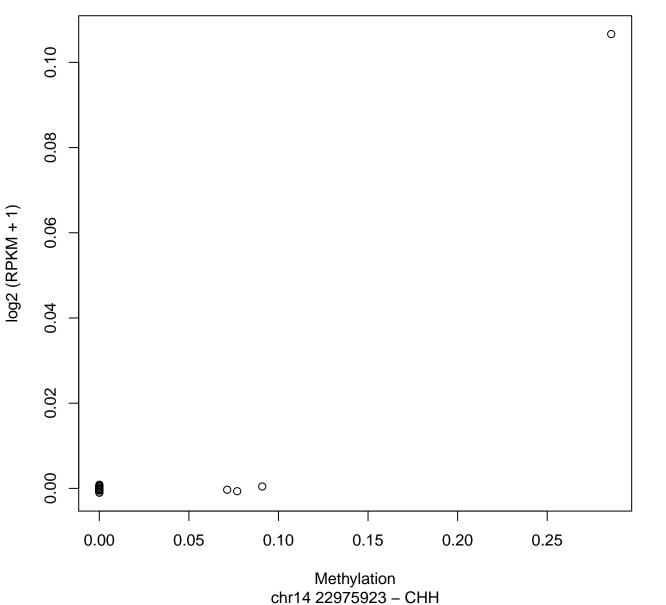
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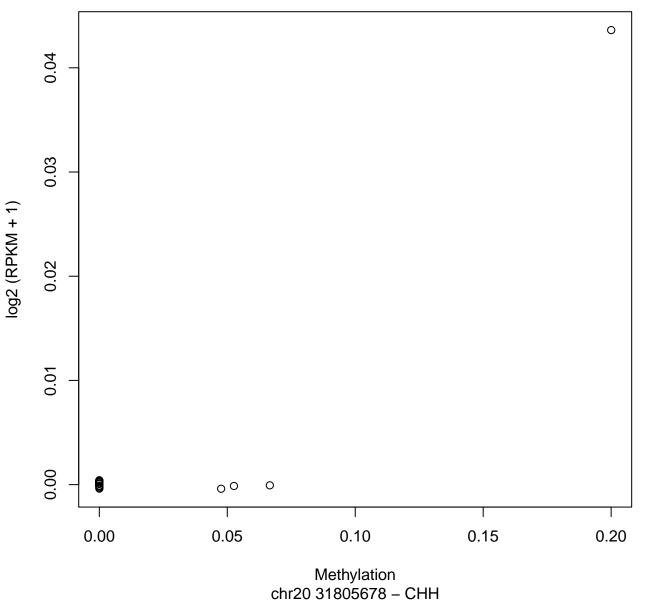
#### exon ENSE00002717942.1 FDR 6.02e-05 AL034553.1



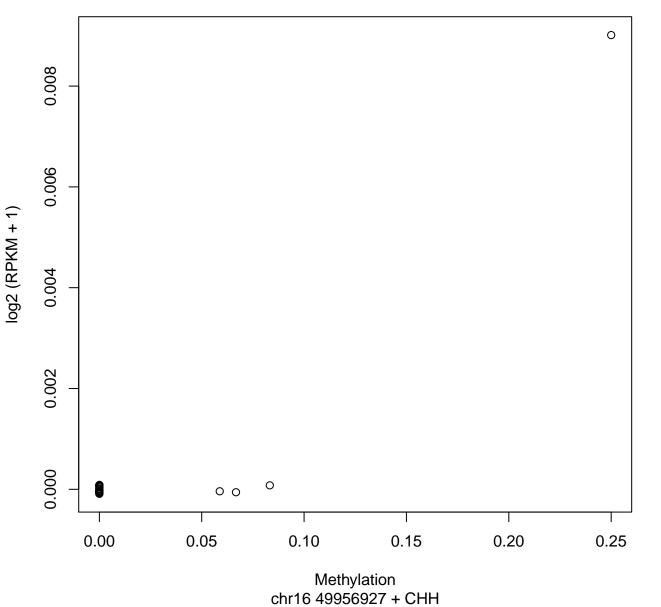
# exon ENSE00001508131.1\_1 FDR 6.16e-05 TRAJ35



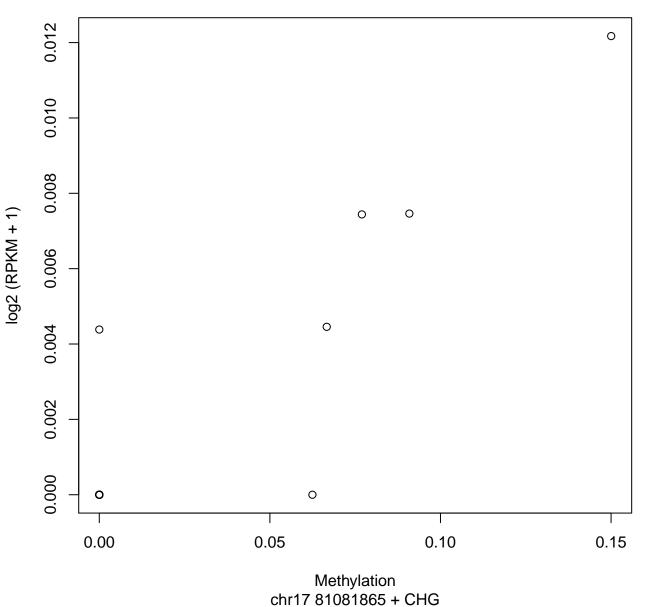
#### exon ENSE00001753340.1\_1 FDR 6.4e-05 RP11-49G10.3



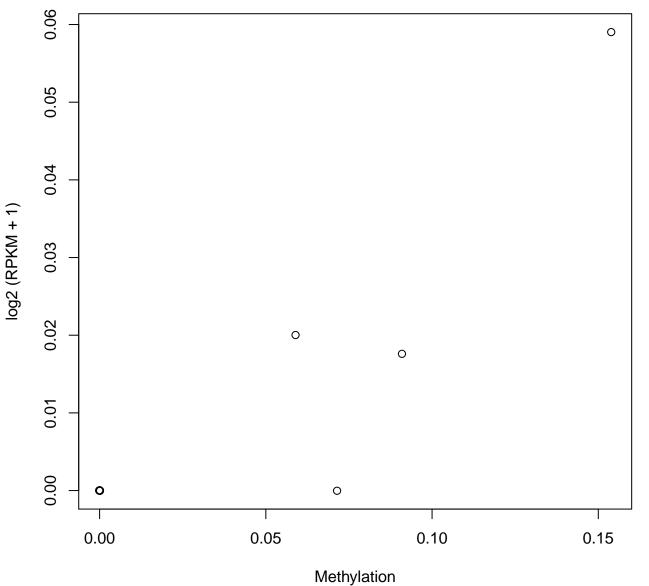
## exon ENSE00002579587.1\_1 FDR 6.52e-05 RP11-305A4.3



### exon ENSE00003757671.1\_1 FDR 6.59e-05 AC139099.7

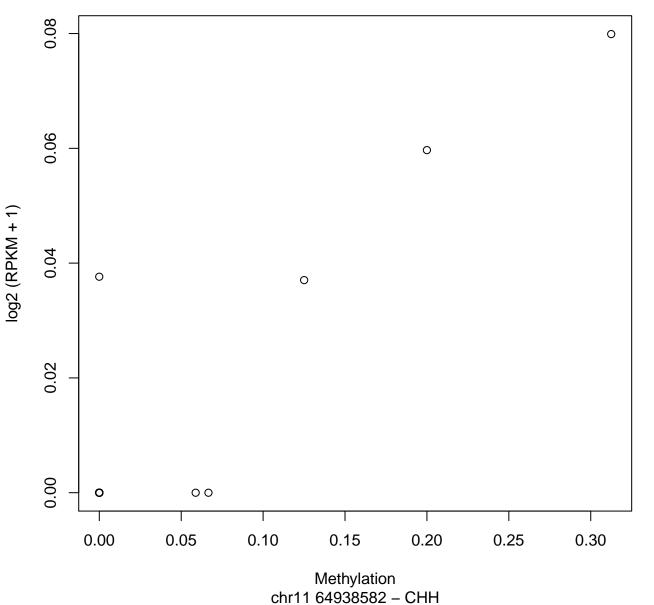


#### exon ENSE00002553941.1\_1 FDR 6.66e-05 RP11-19J5.2

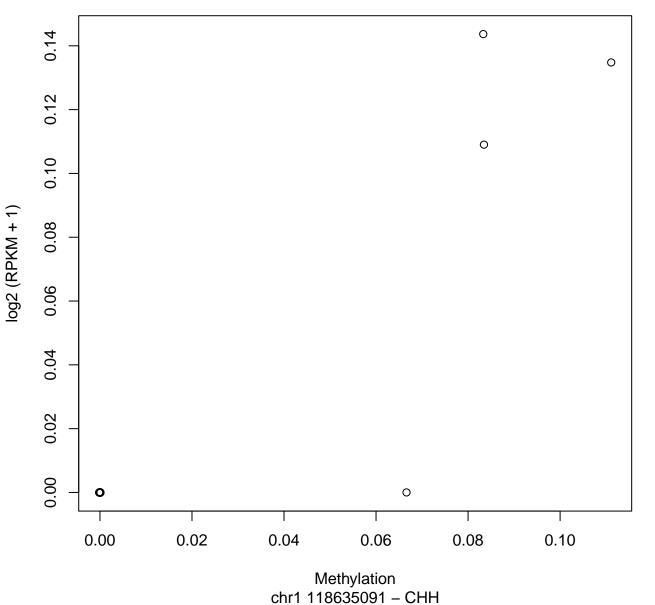


chr15 70113504 + CHH

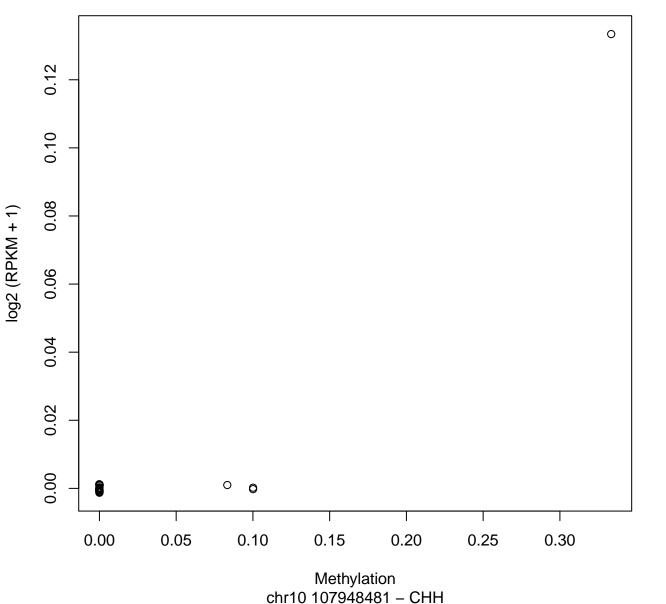
# exon ENSE00001473046.1\_1 FDR 6.77e-05 SPDYC



#### exon ENSE00001249619.3\_1 FDR 7.16e-05 SPAG17



#### exon ENSE00001761258.1\_1 FDR 7.53e-05 RP11-298H24.1



## exon ENSE00003544517.1\_1 FDR 7.72e-05 PSG6

