exon ENSE00001418403.1_1 FDR 0.000154
NBEA

log2 (RPKM + 1)

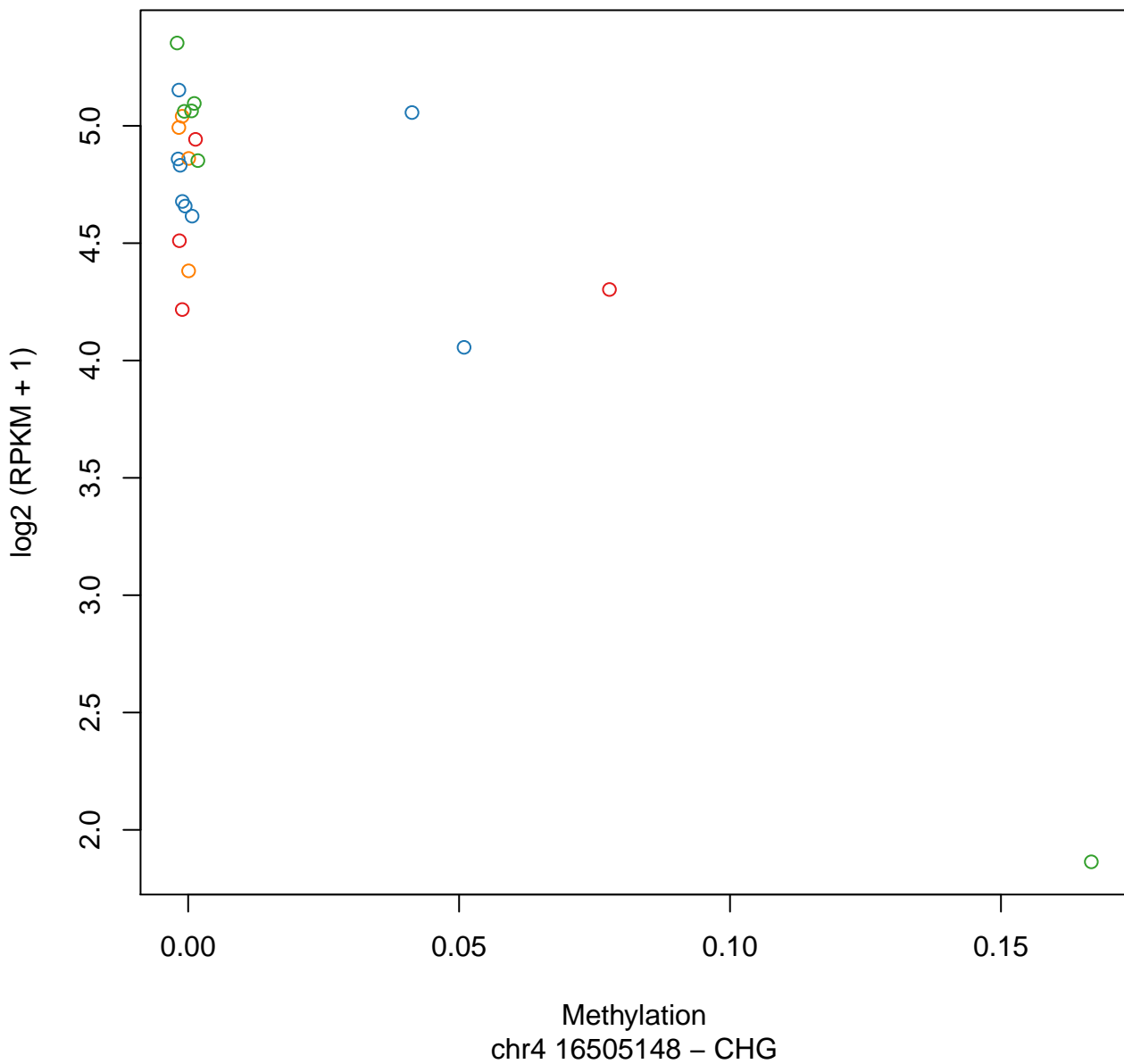Methylation
chr13 36247285 + CHG
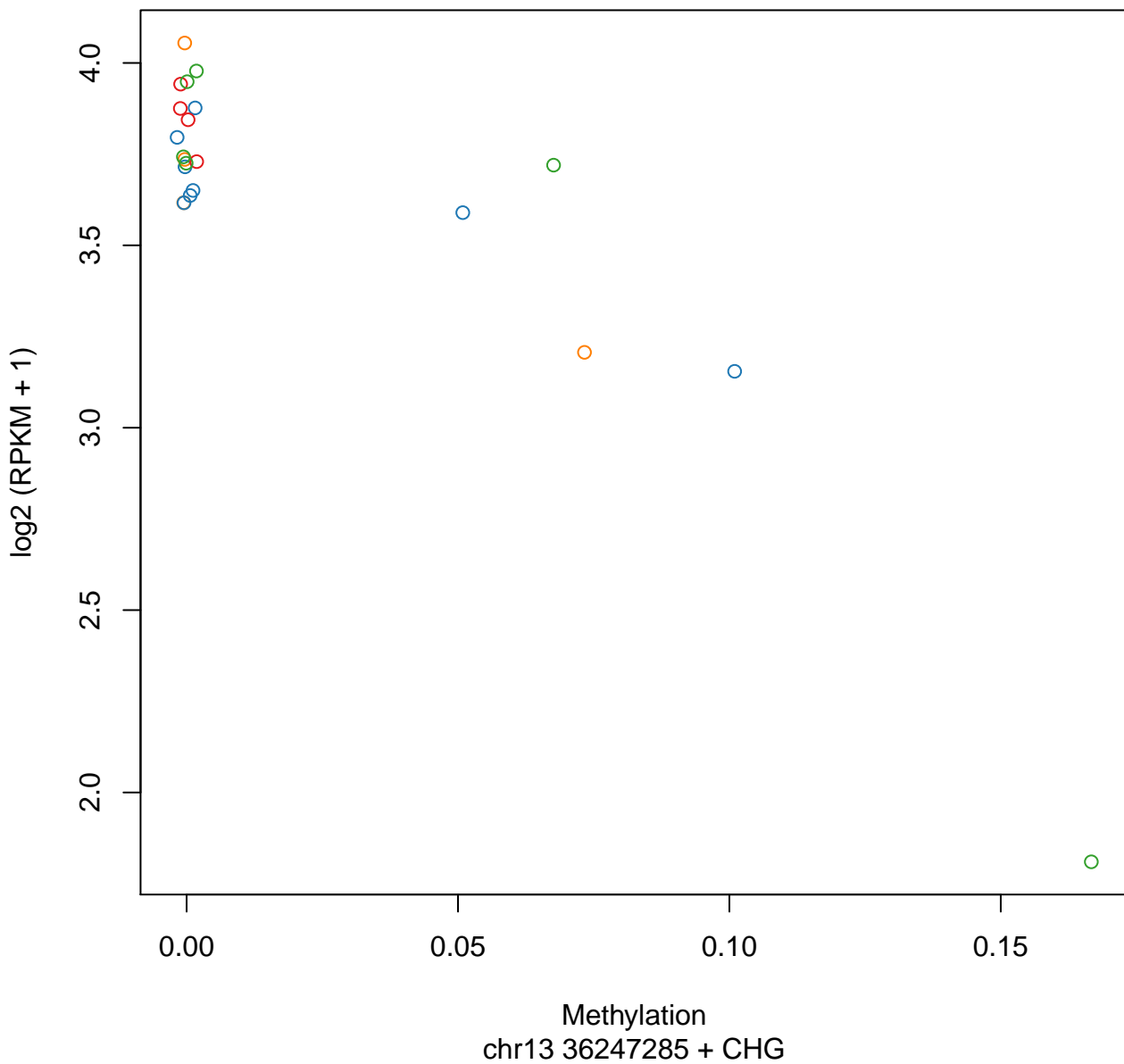
exon ENSE00001483114.1_1 FDR 0.000154
NBEA

log2 (RPKM + 1)

Methylation
chr13 36247285 + CHG
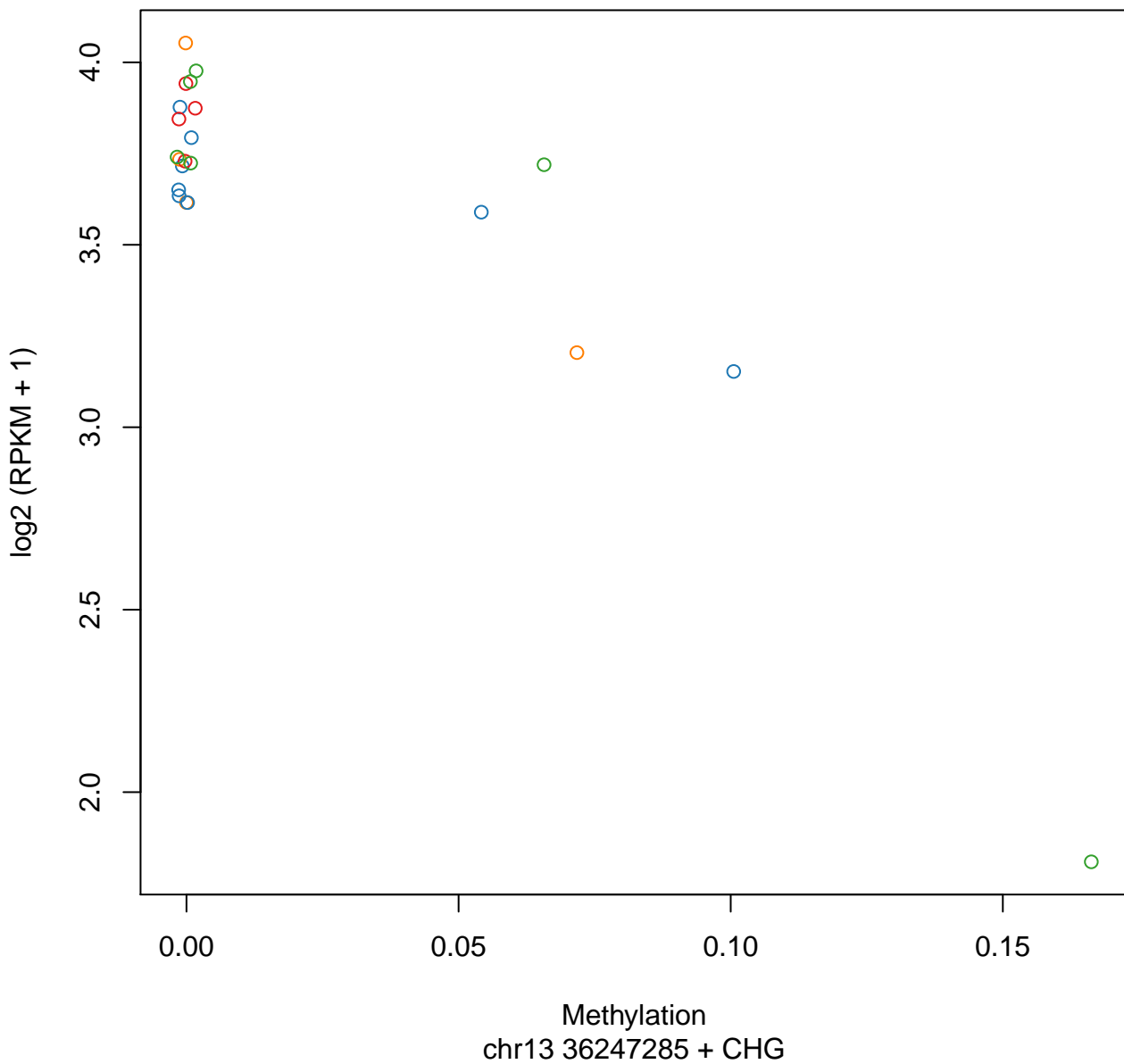
exon ENSE00002047930.1_1 FDR 0.00111
LDB2

log2 (RPKM + 1) vs Methylation
chr4 16505148 − CHG

exon ENSE00001743131.2_1 FDR 0.00201
AC007740.1

log2 (RPKM + 1)

Methylation
chr2 163625451 + CHG
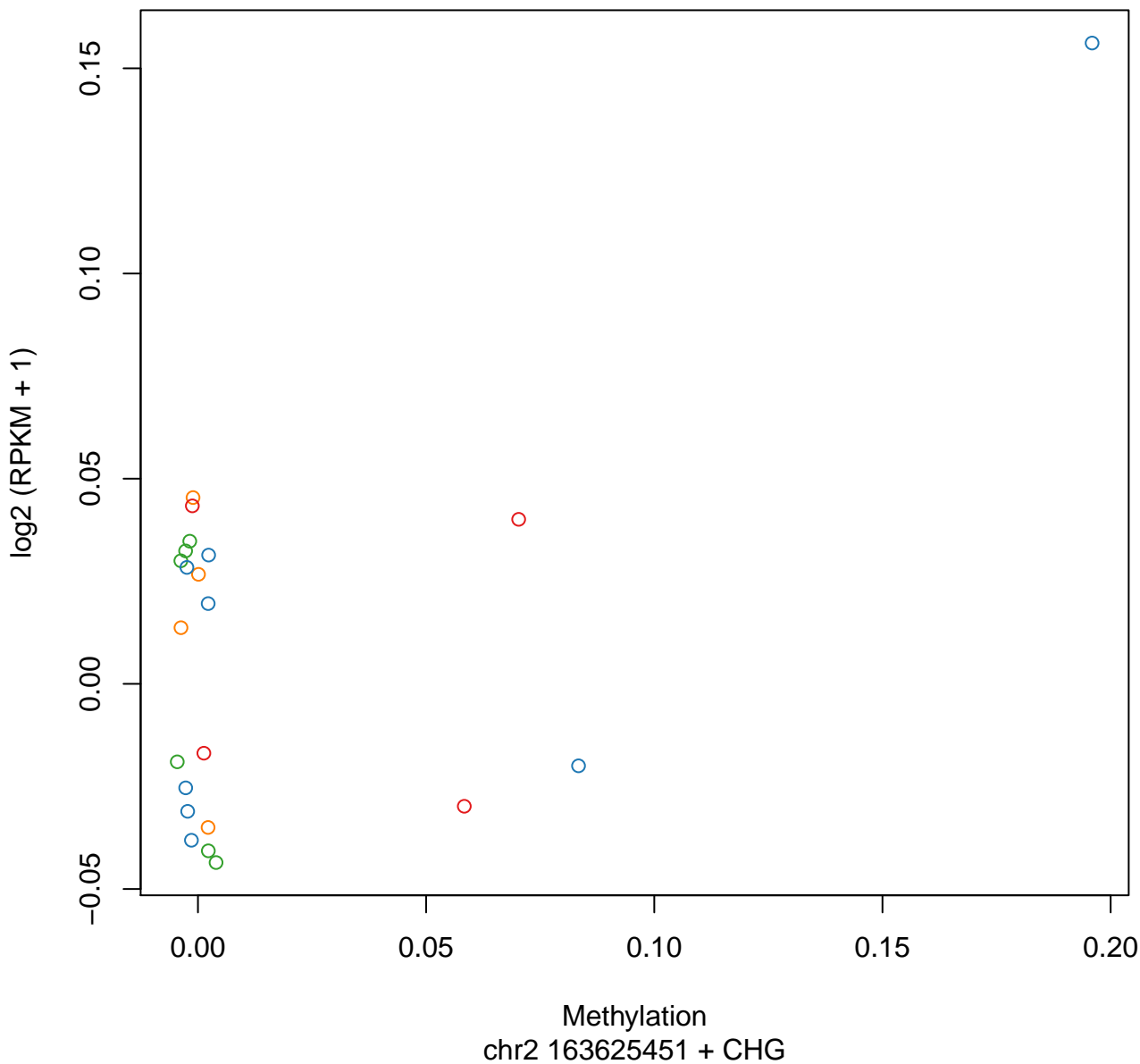
exon ENSE00002081945.1_1 FDR 0.00626
LDB2

Methylation
chr4 16505148 – CHG
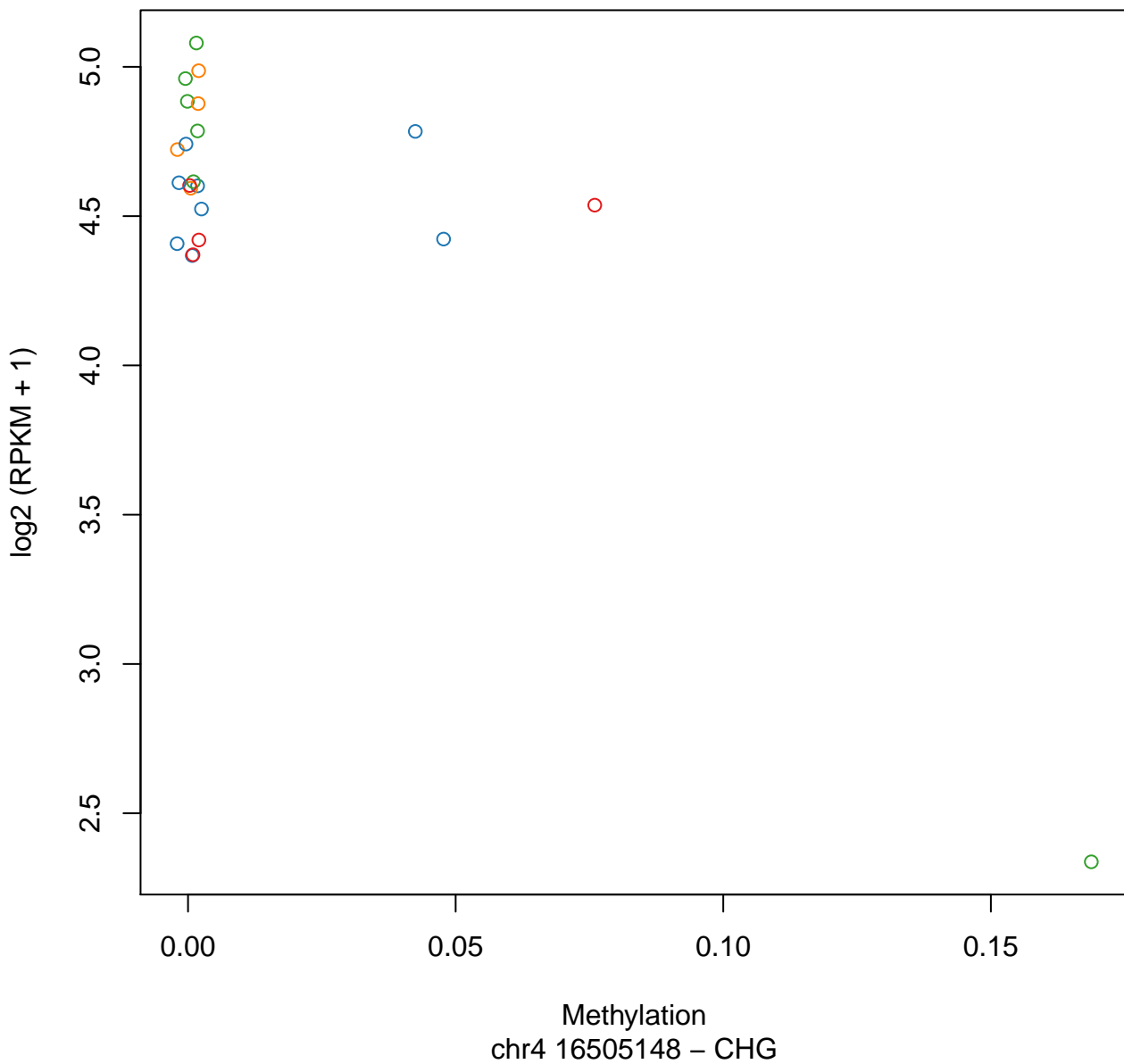
exon ENSE00003668908.1_1 FDR 0.00849
LDB2

log2 (RPKM + 1)

Methylation
chr4 16505148 – CHG

exon ENSE00001779542.1_1 FDR 0.00849
LDB2

Methylation
chr4 16505148 − CHG
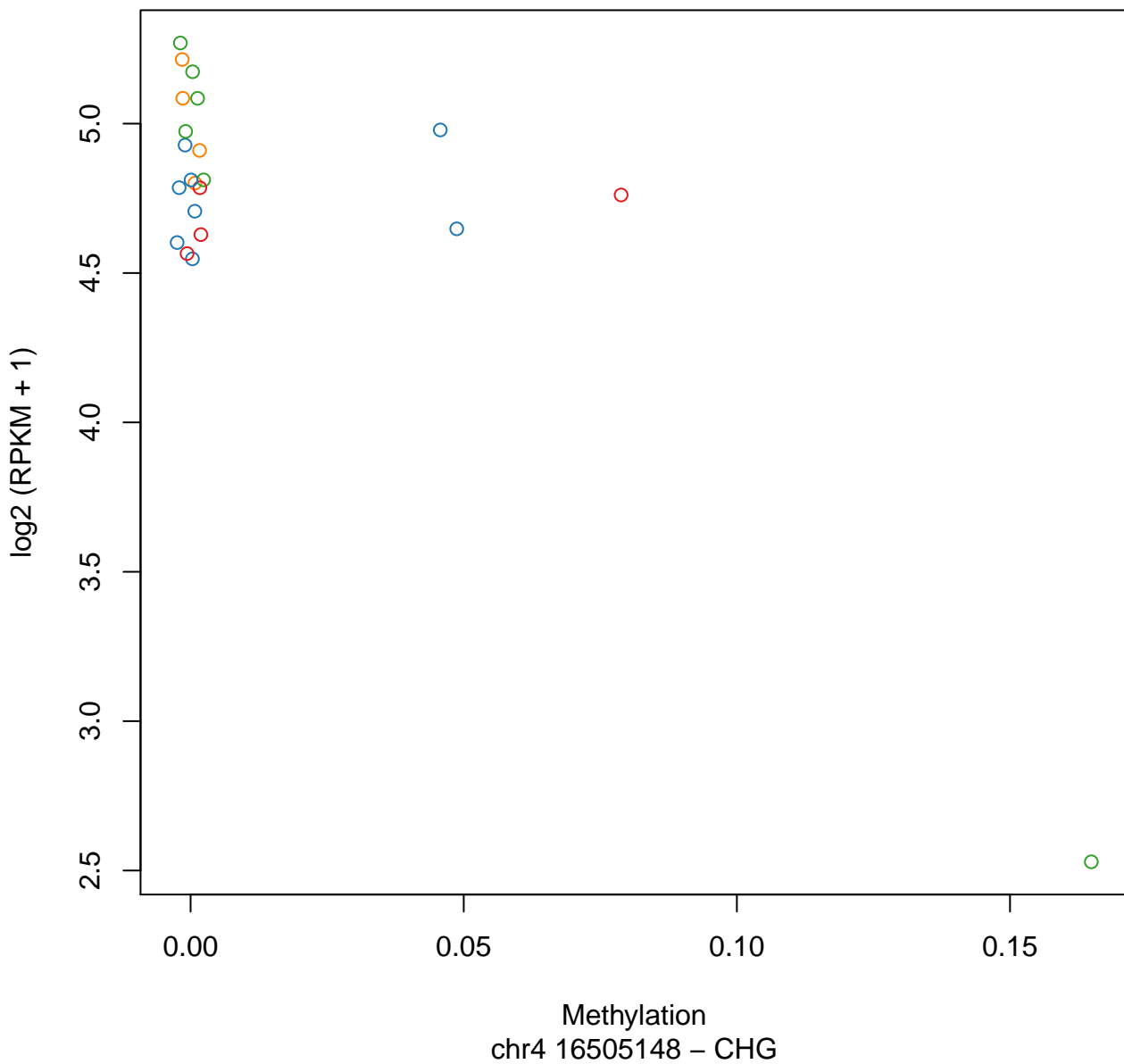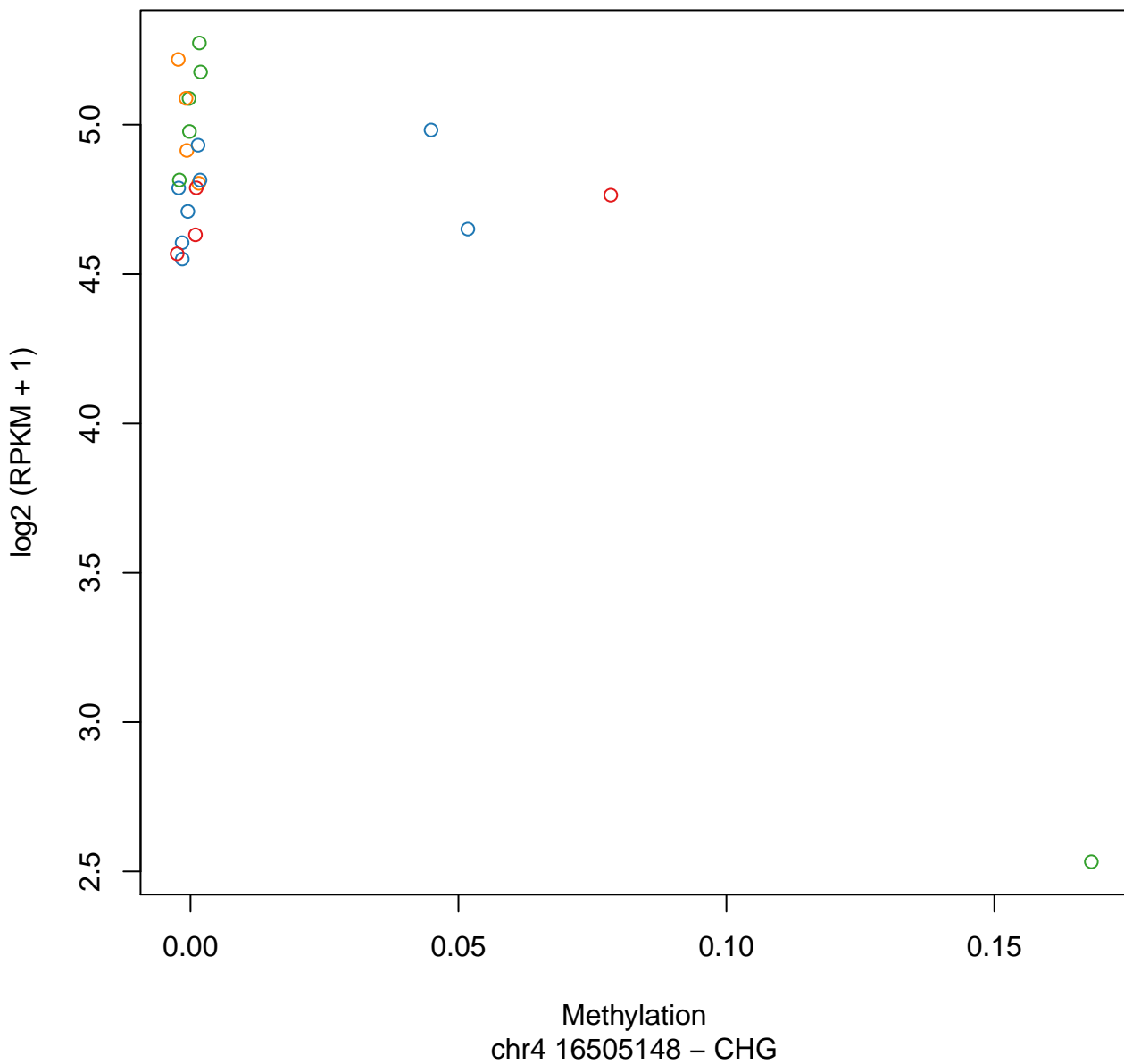
exon ENSE00002308670.1_1 FDR 0.0182
CACNA1C−AS4

log2 (RPKM + 1)

Methylation
chr12 2330000 + CHG
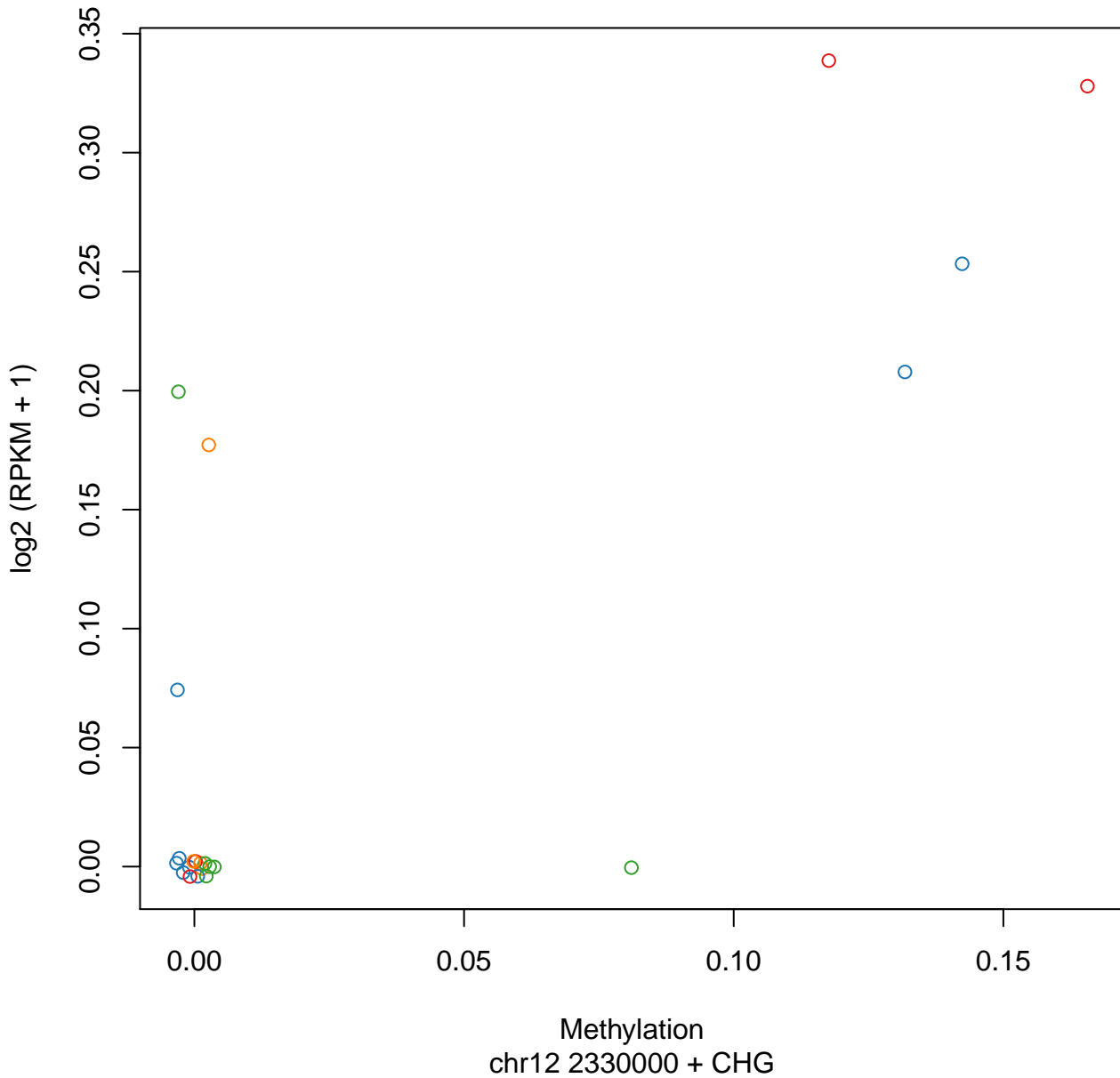
exon ENSE00002188123.1_1 FDR 0.0197
OPCML

log2 (RPKM + 1)

Methylation
chr11 132290458 + CHH
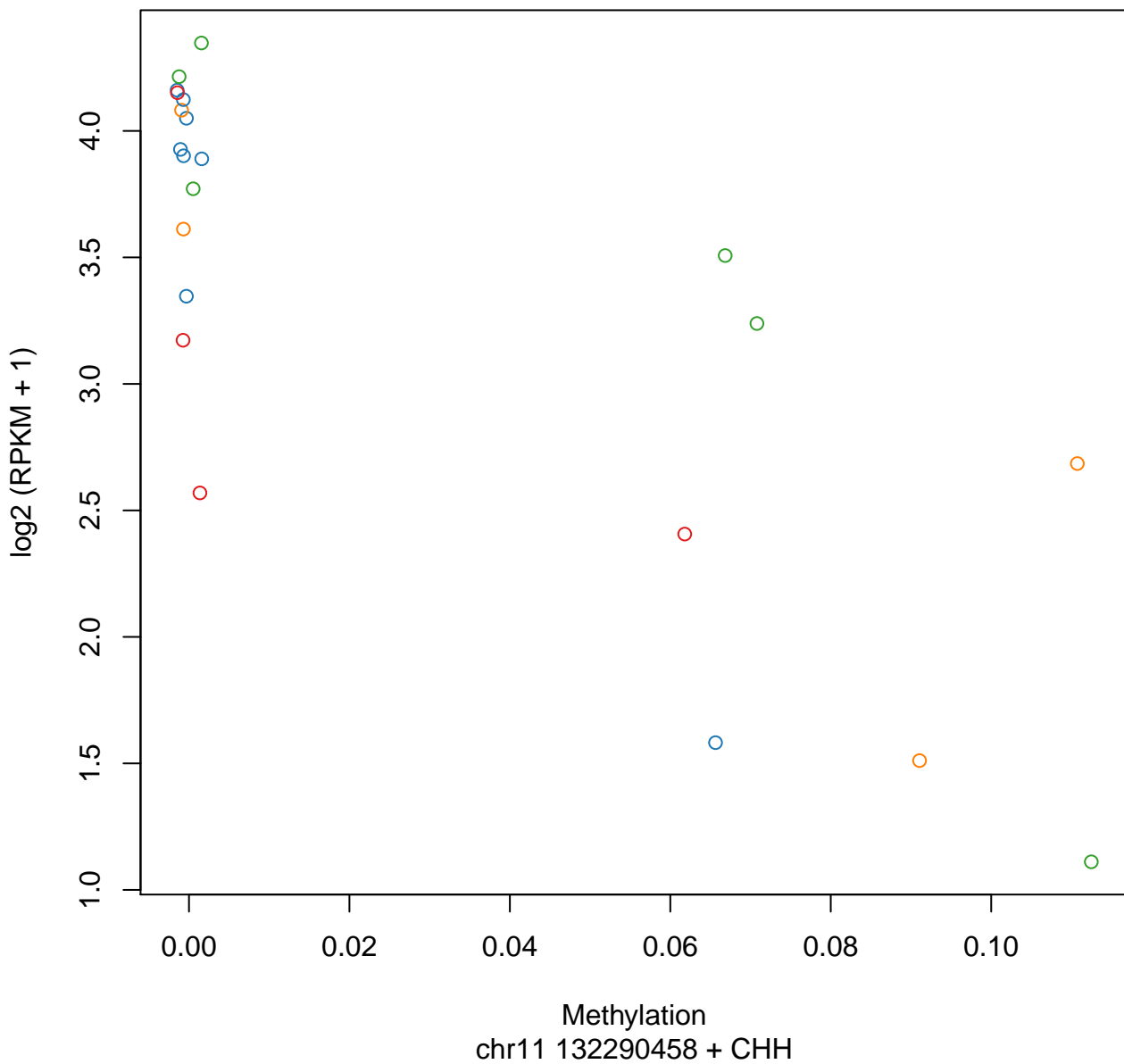
exon ENSE00002621434.1_1 FDR 0.0226
PWRN1

log2 (RPKM + 1)

Methylation
chr15 24779265 + CHG
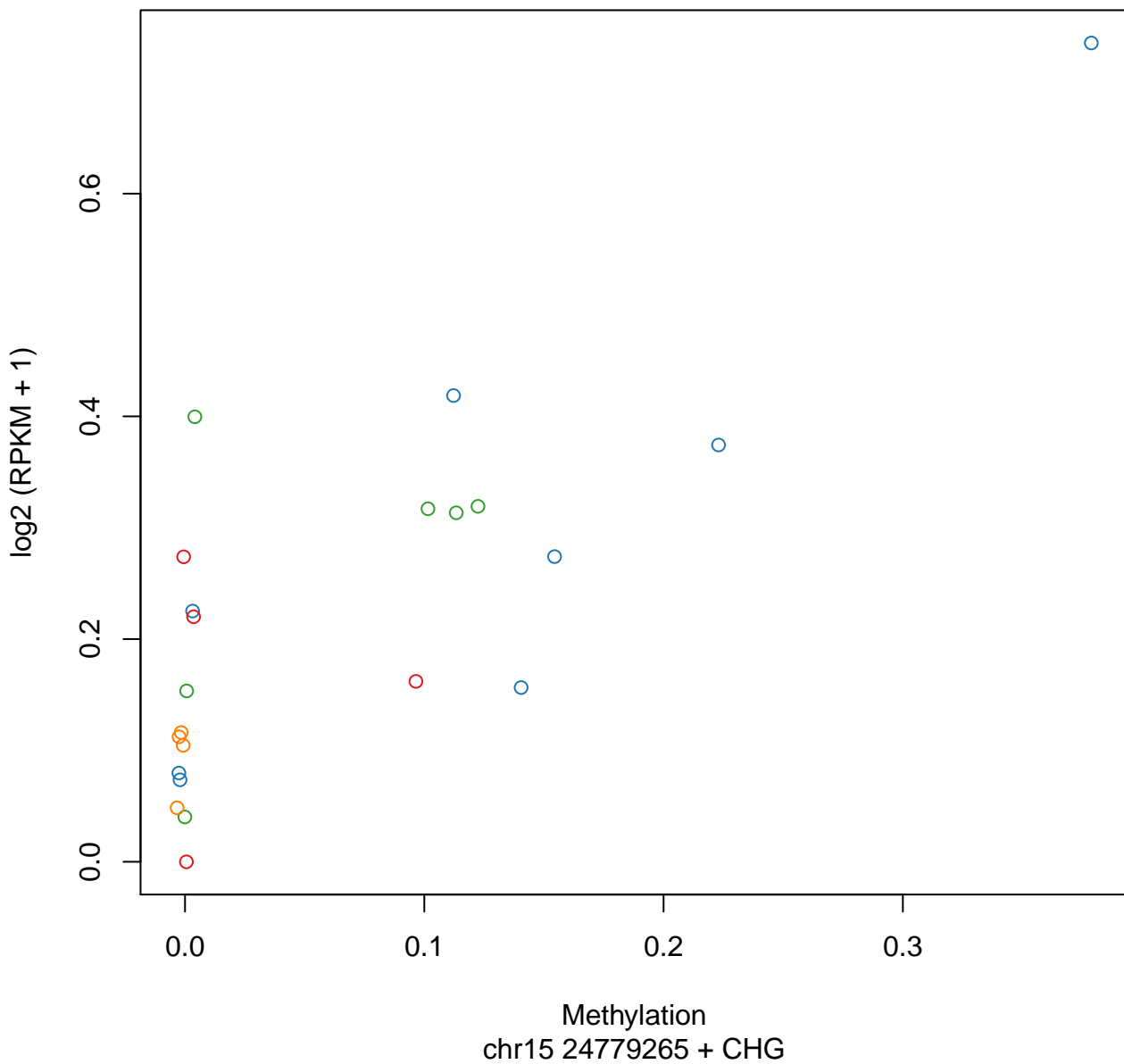
exon ENSE00002905144.1_1 FDR 0.0239
NEDD4L

Methylation
chr18 56067848 – CHH
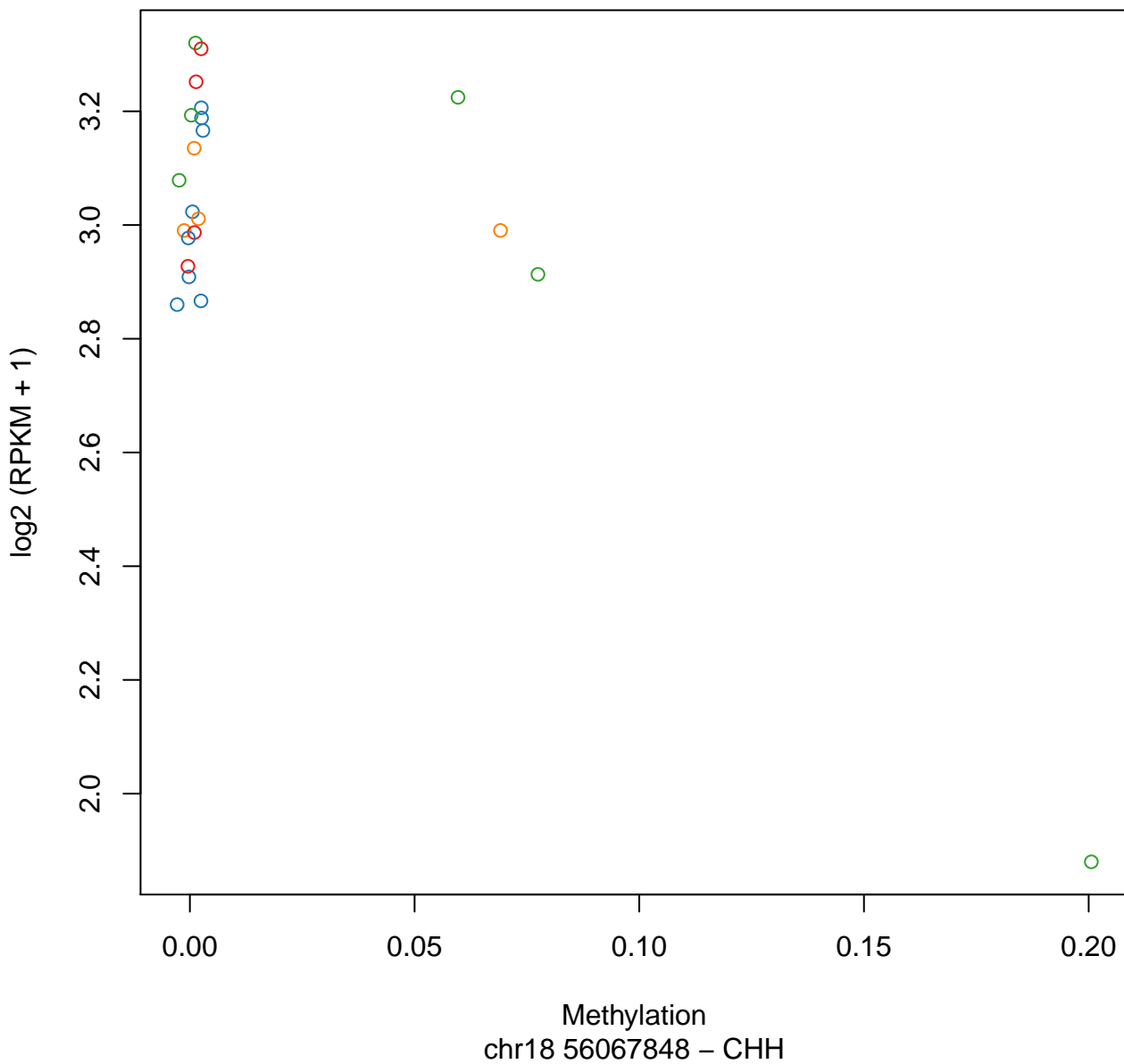
exon ENSE00001341470.2_1 FDR 0.0239
NEDD4L
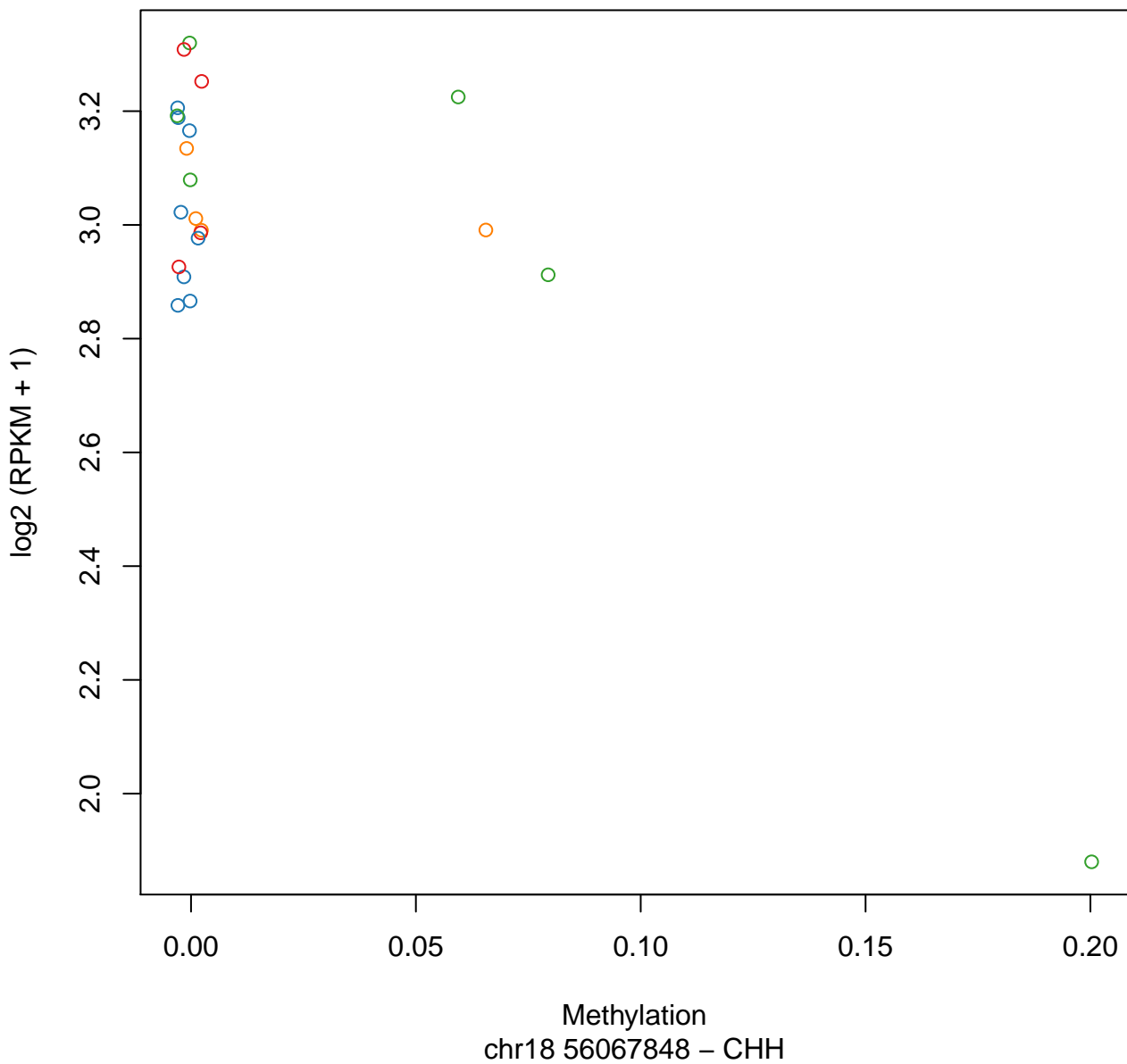
exon ENSE00003638601.2_1 FDR 0.0249
VIPR1

log2 (RPKM + 1)

Methylation
chr3 42578213 − CHG
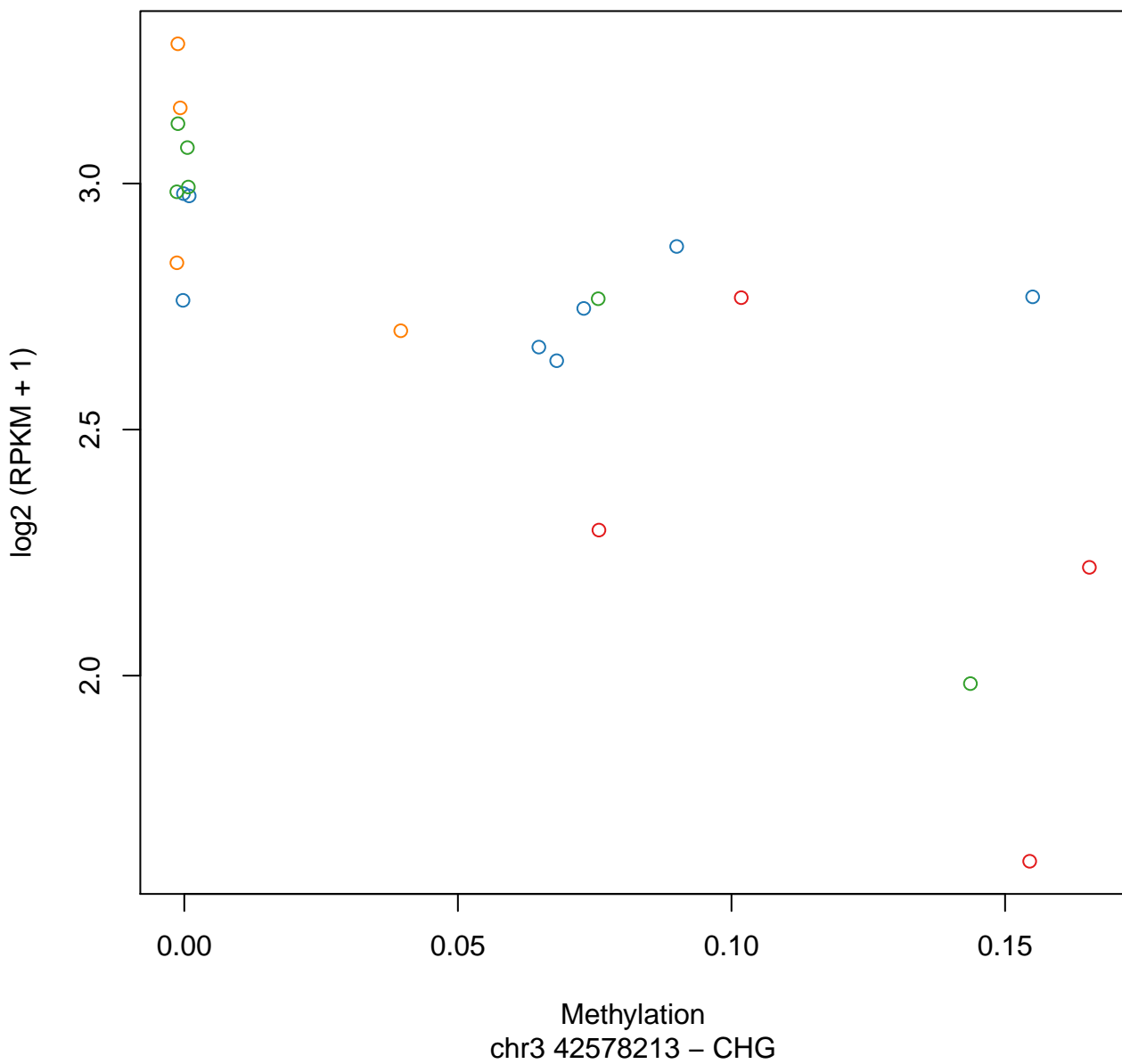
exon ENSE00003546753.1_1 FDR 0.025
VIPR1

log2 (RPKM + 1)

Methylation
chr3 42578213 − CHG
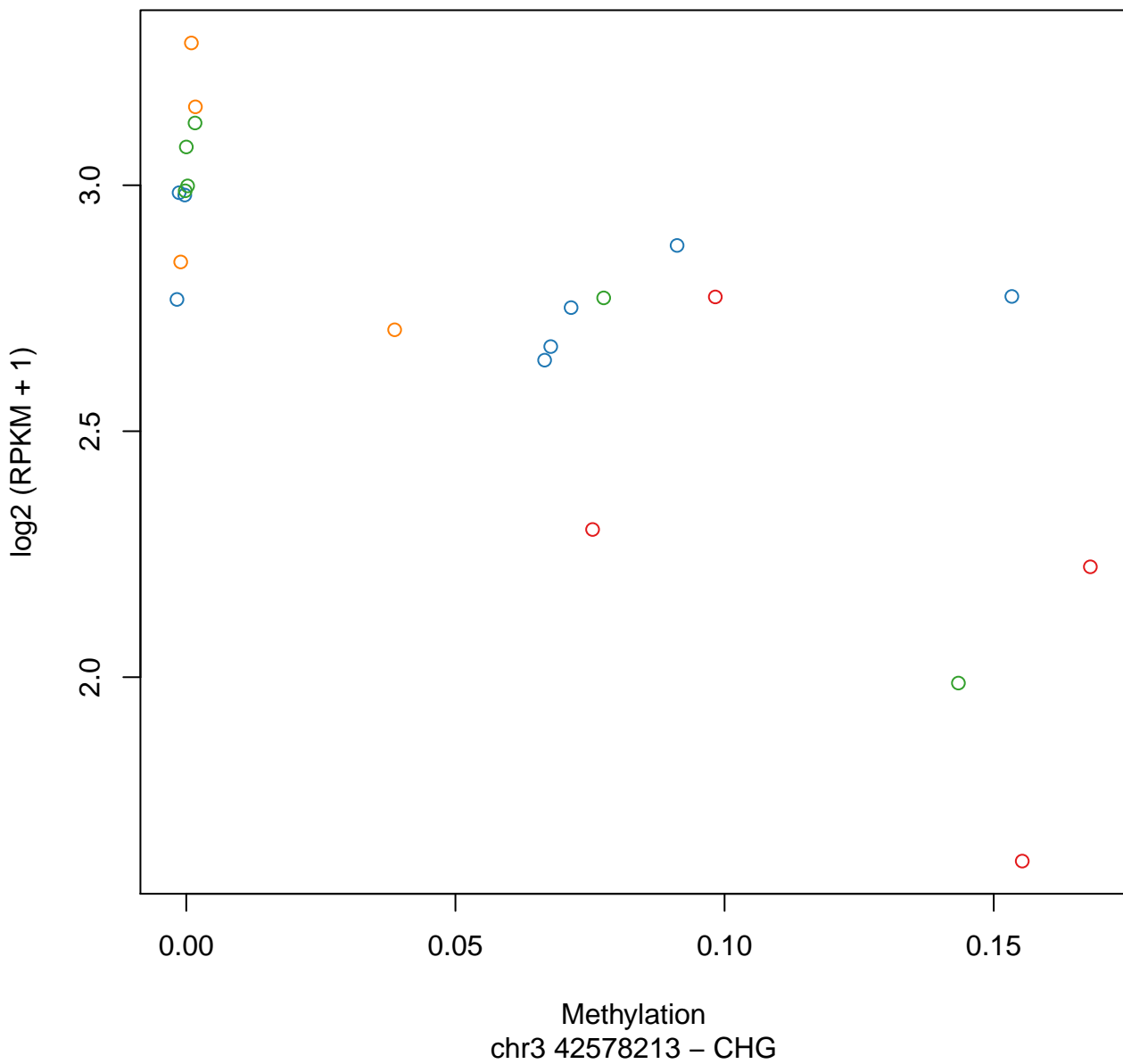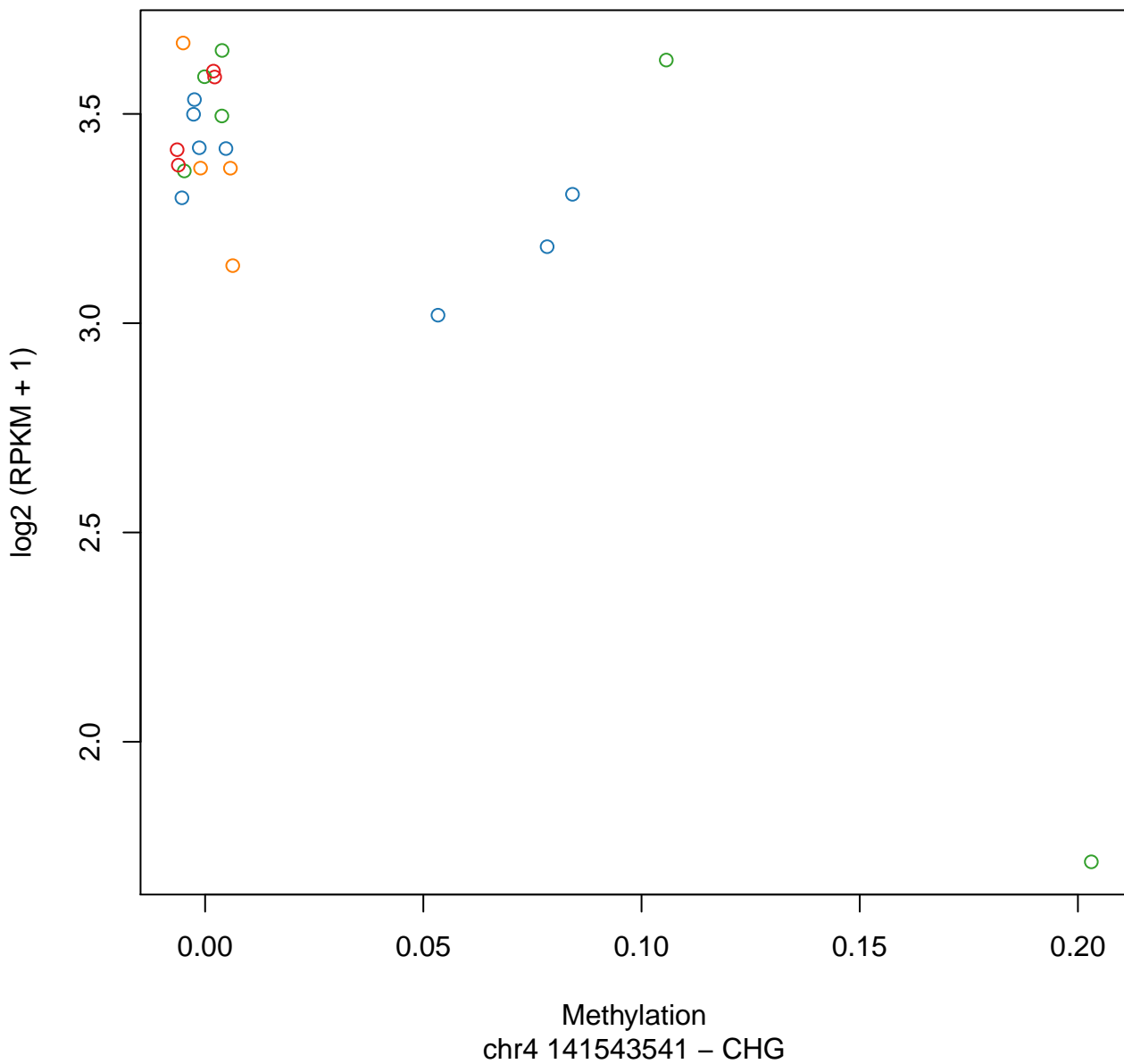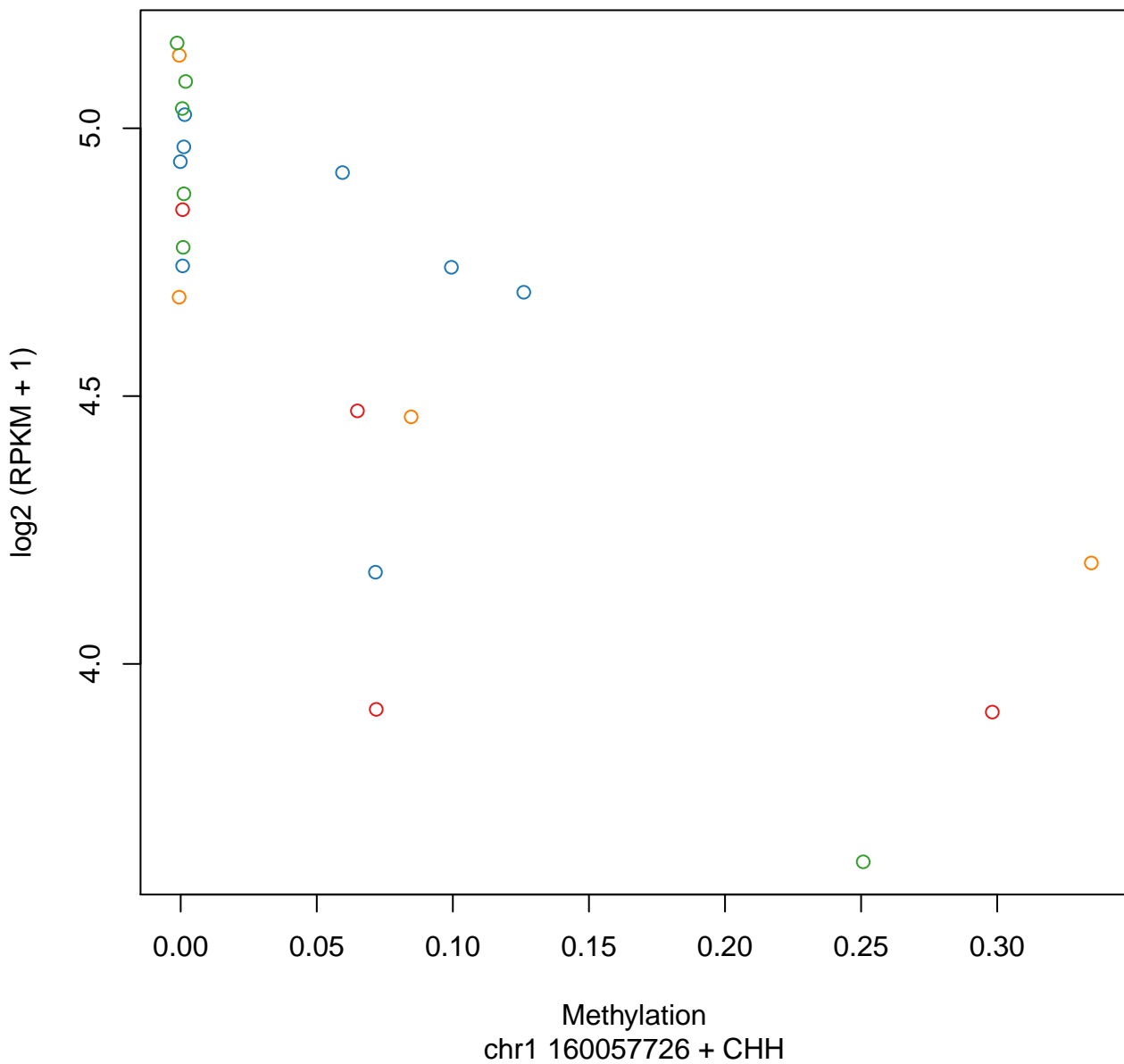
exon ENSE00001444298.3_1 FDR 0.0261
ULBP2
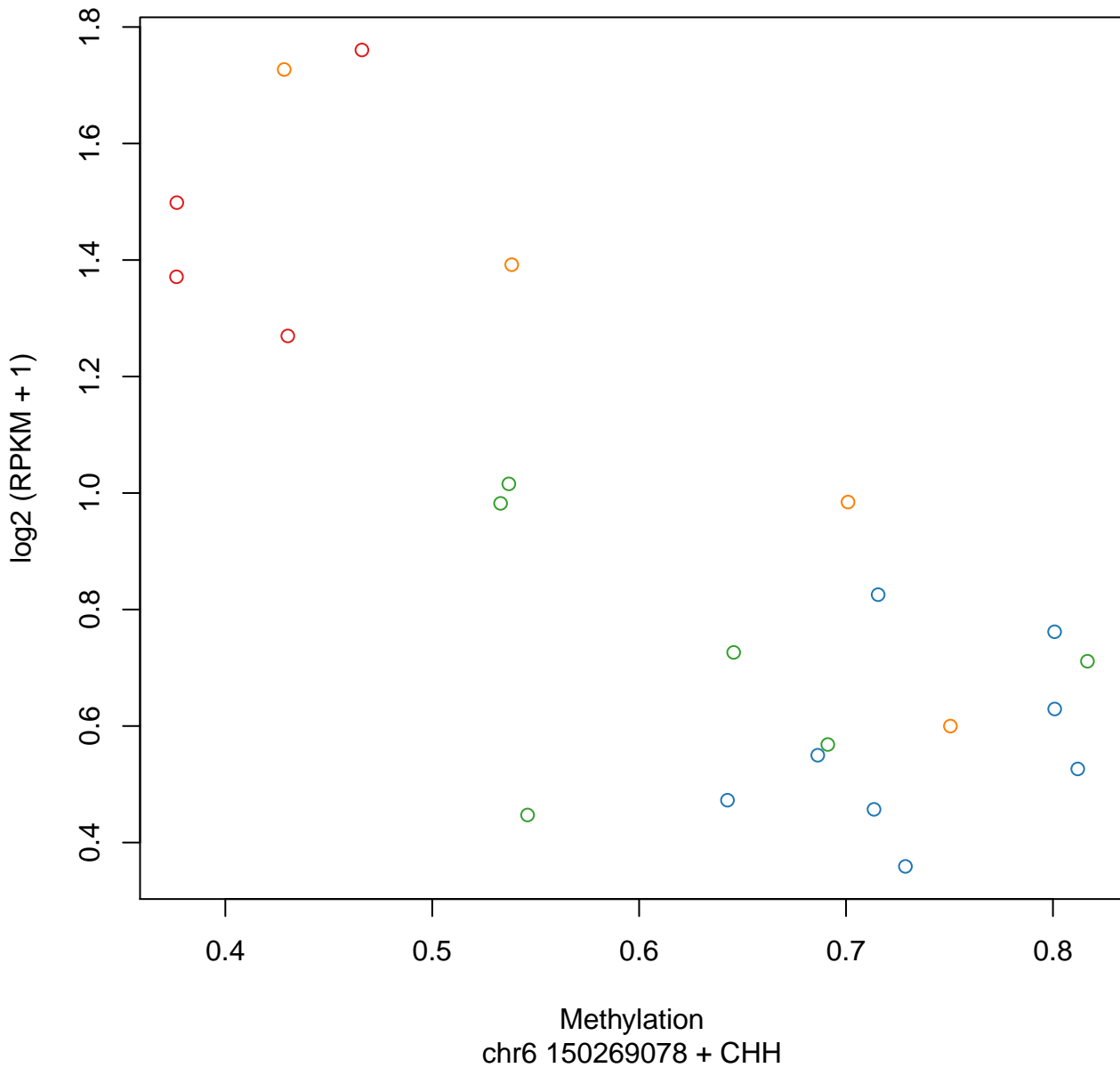
exon ENSE00001747040.2_1 FDR 0.0291
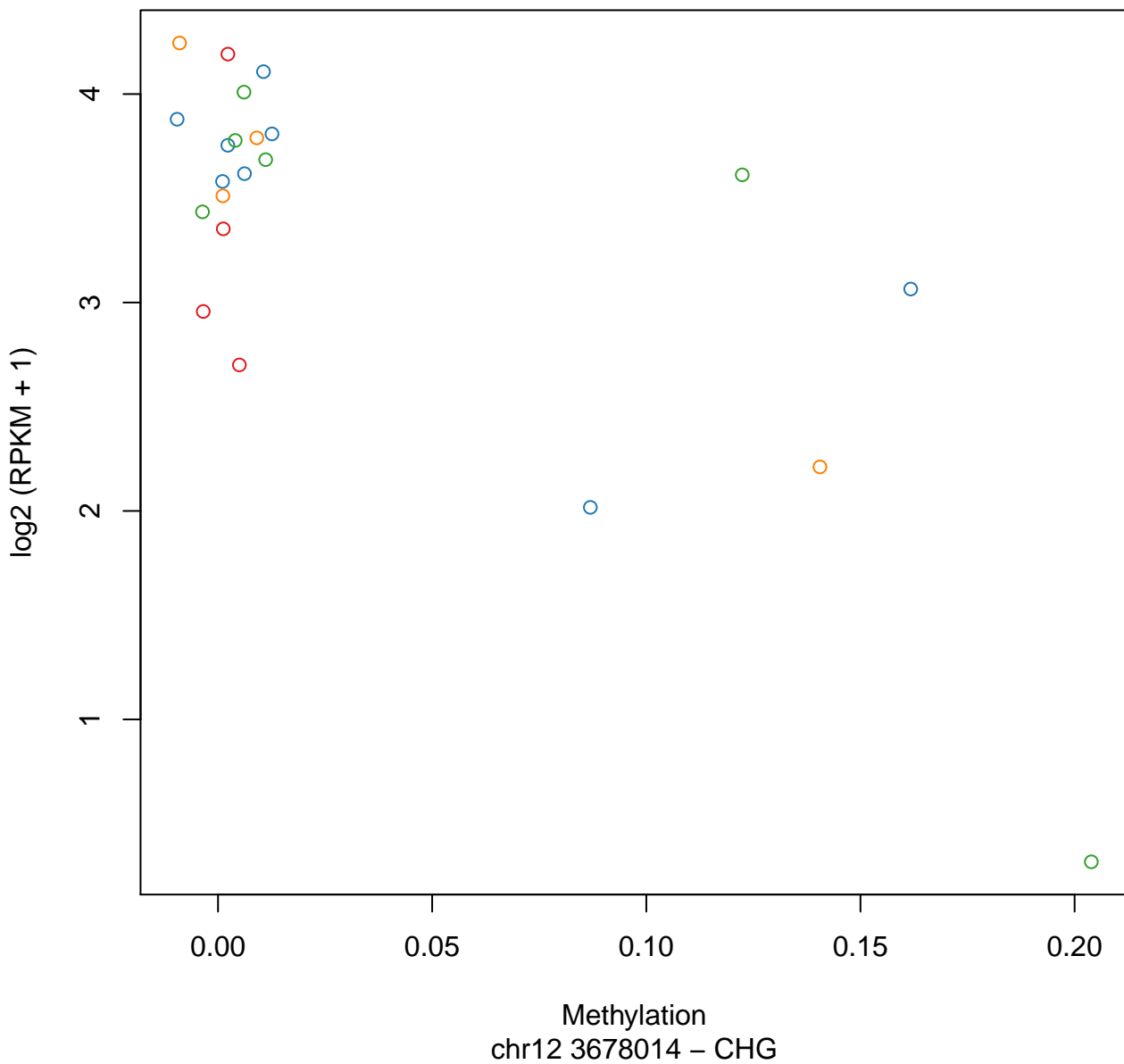TBC1D9

exon ENSE00001446294.3_1 FDR 0.0317
KCNJ9

exon ENSE00001448906.3_1 FDR 0.032
SV2A

log2 (RPKM + 1)
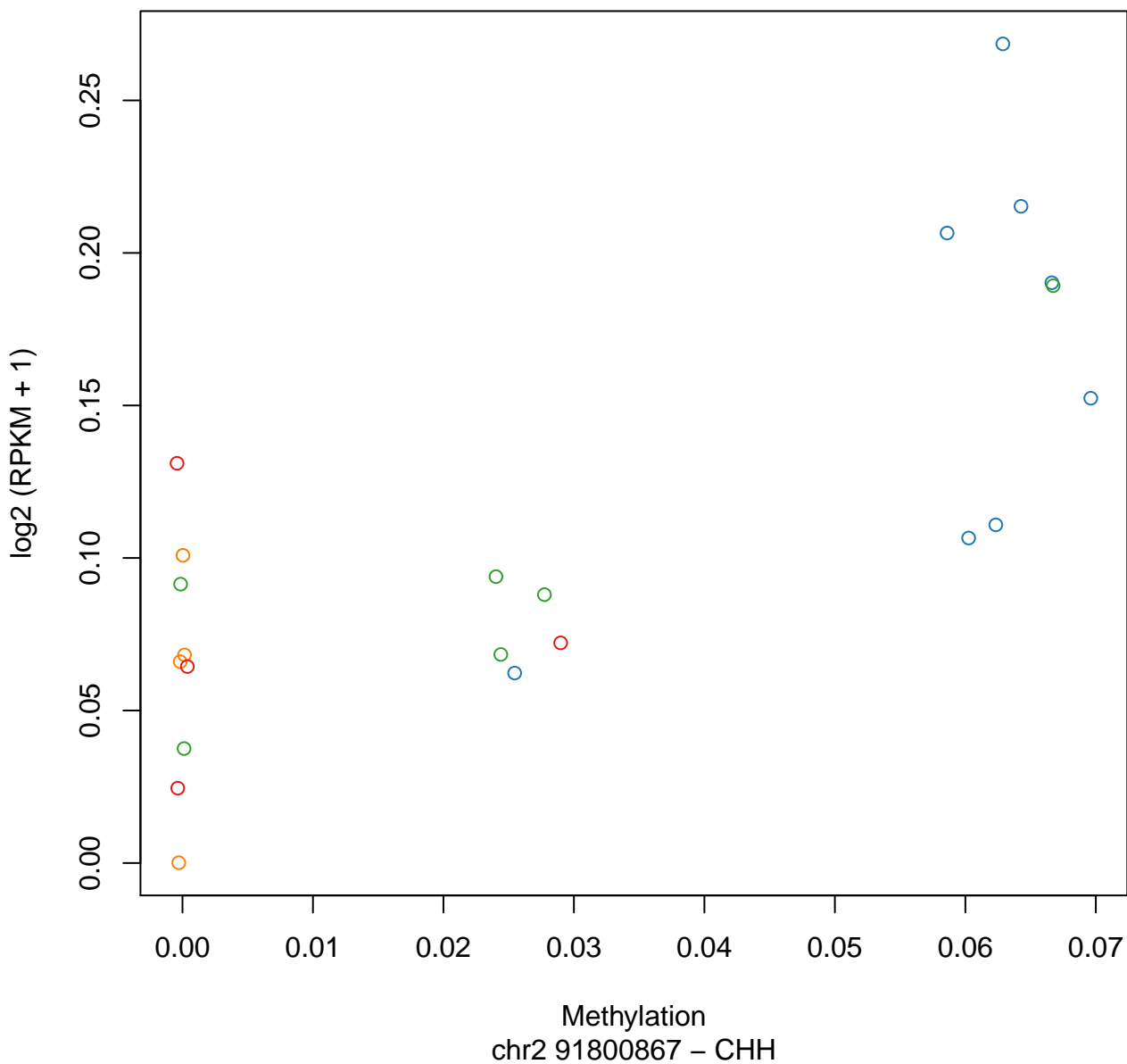
Methylation
chr1 149875242 + CHH
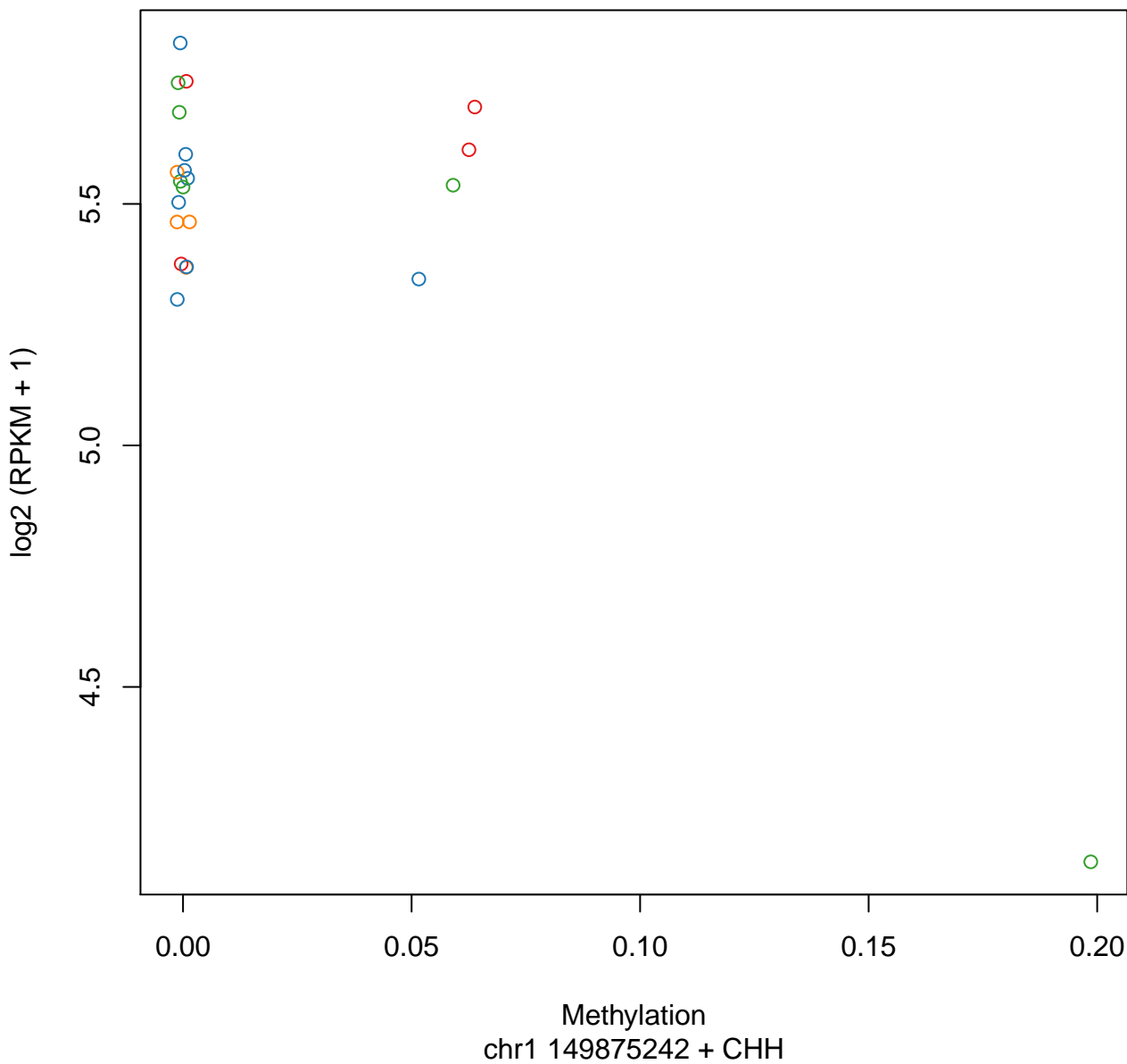
exon ENSE00003651107.1_1 FDR 0.0339
PRMT8

Methylation
chr12 3678014 − CHG

exon ENSE00003716811.1_1 FDR 0.0343
CH17−333M13.1

Methylation
chr2 91800867 − CHH

exon ENSE00001222019.7_1 FDR 0.035
PIANP

log2 (RPKM + 1)

Methylation
chr12 6804357 + CHH
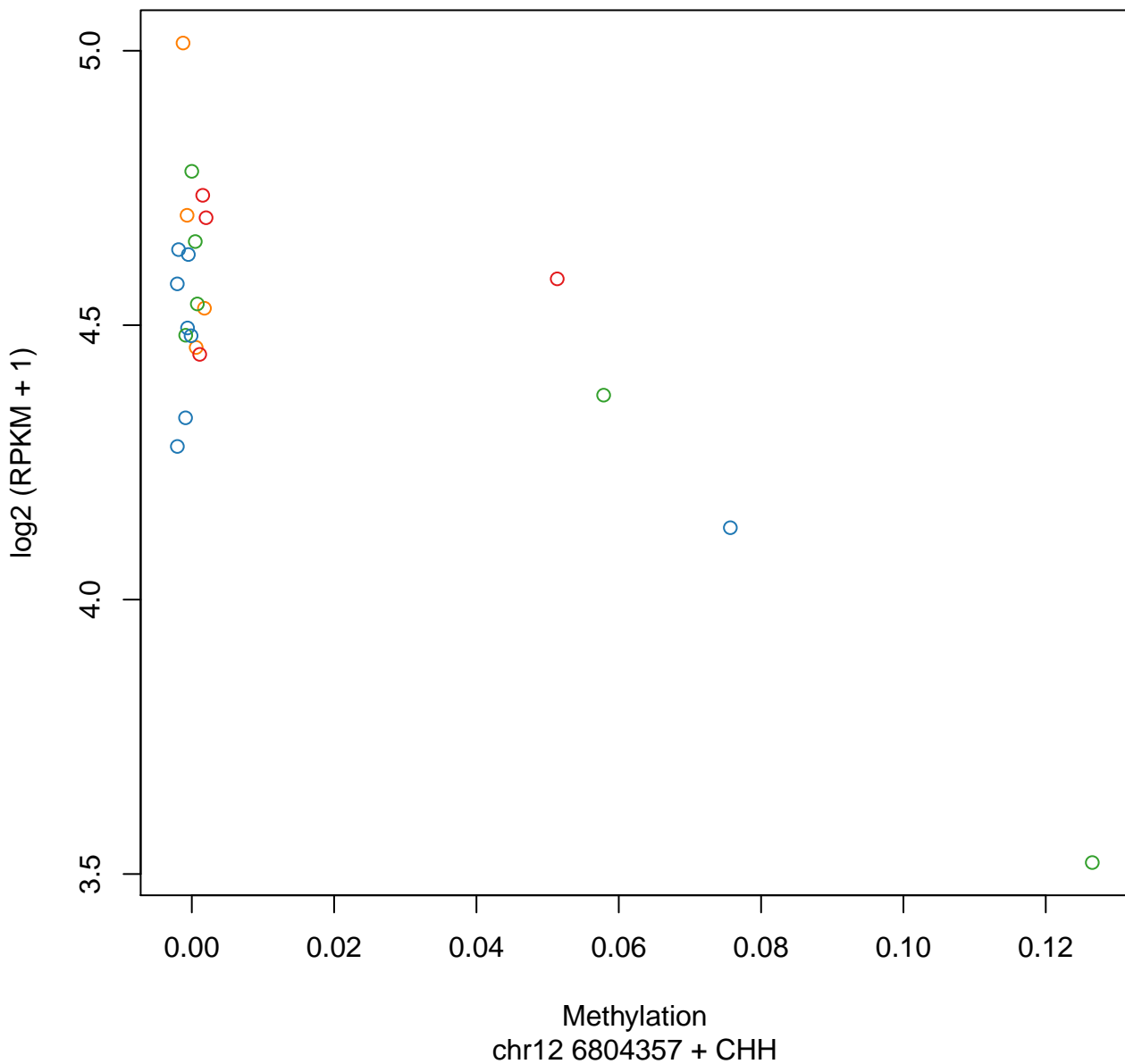
exon ENSE00001838757.1_1 FDR 0.0427
HS3ST2

Methylation
chr16 22927772 + CHH
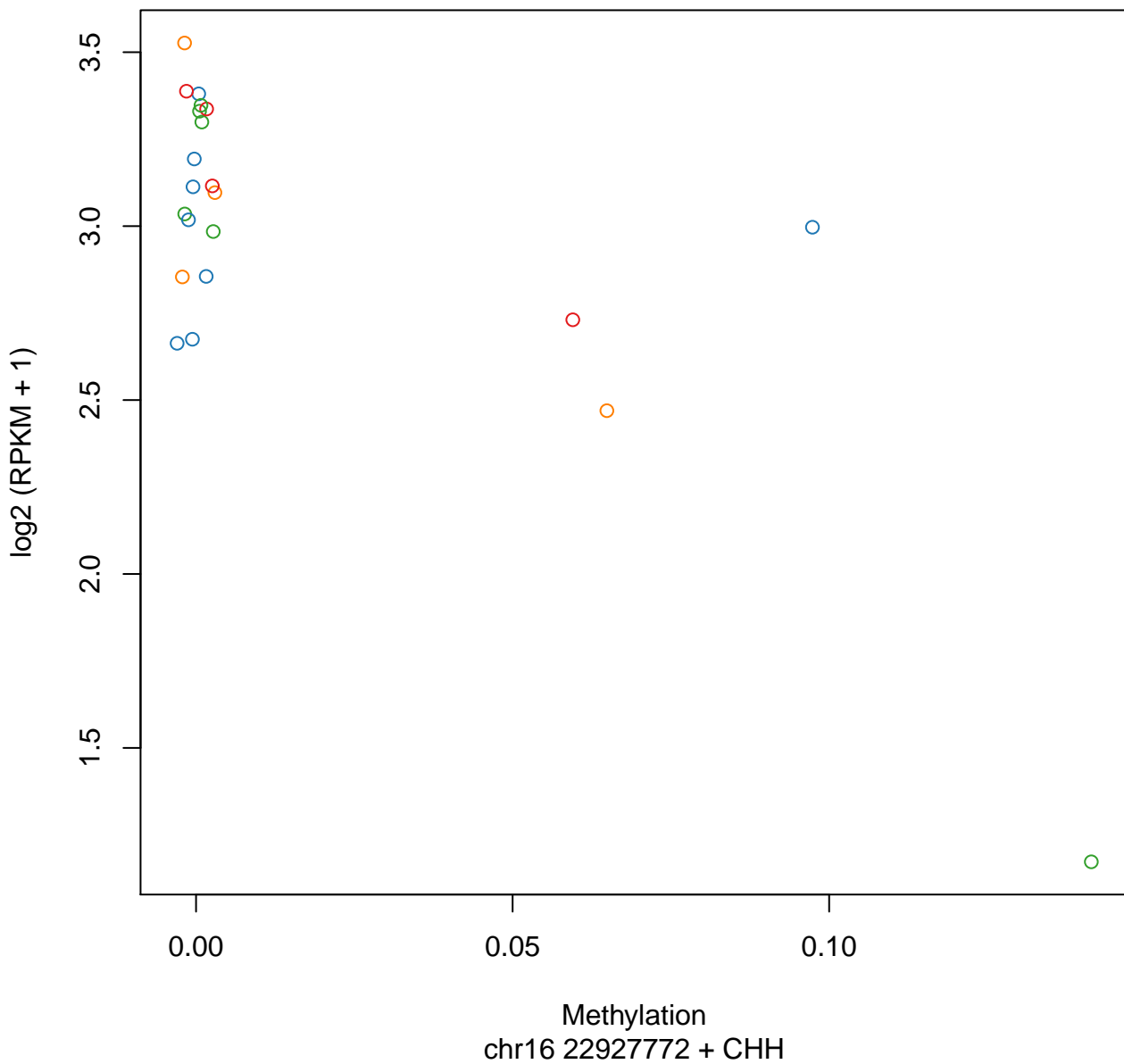
exon ENSE00001188294.4_1 FDR 0.043
HS3ST2

log2 (RPKM + 1)

Methylation
chr16 22927772 + CHH
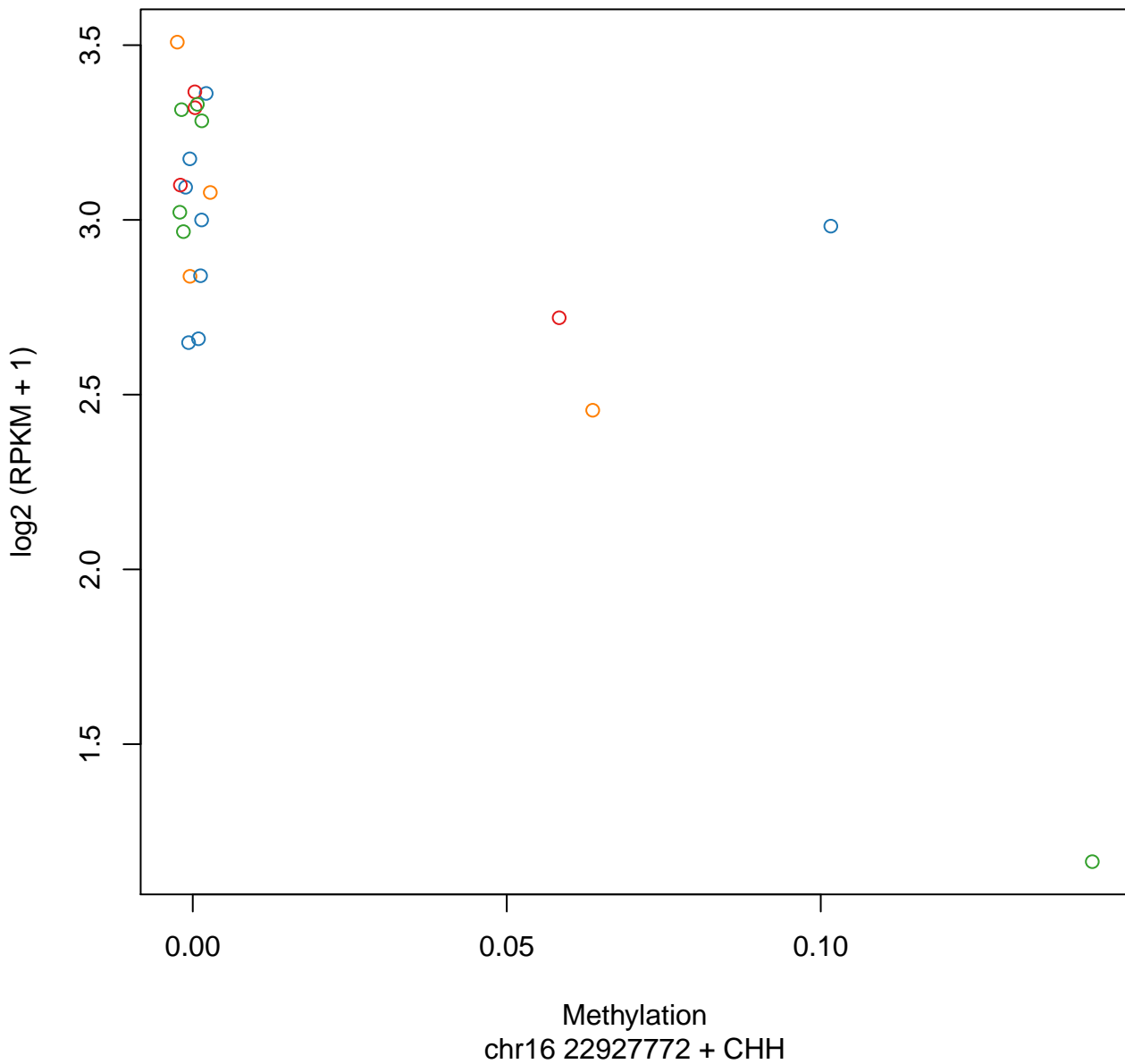
exon ENSE00003799046.1_1 FDR 0.0437
CTC−304I17.6

log2 (RPKM + 1)

Methylation
chr17 32263794 − CHH
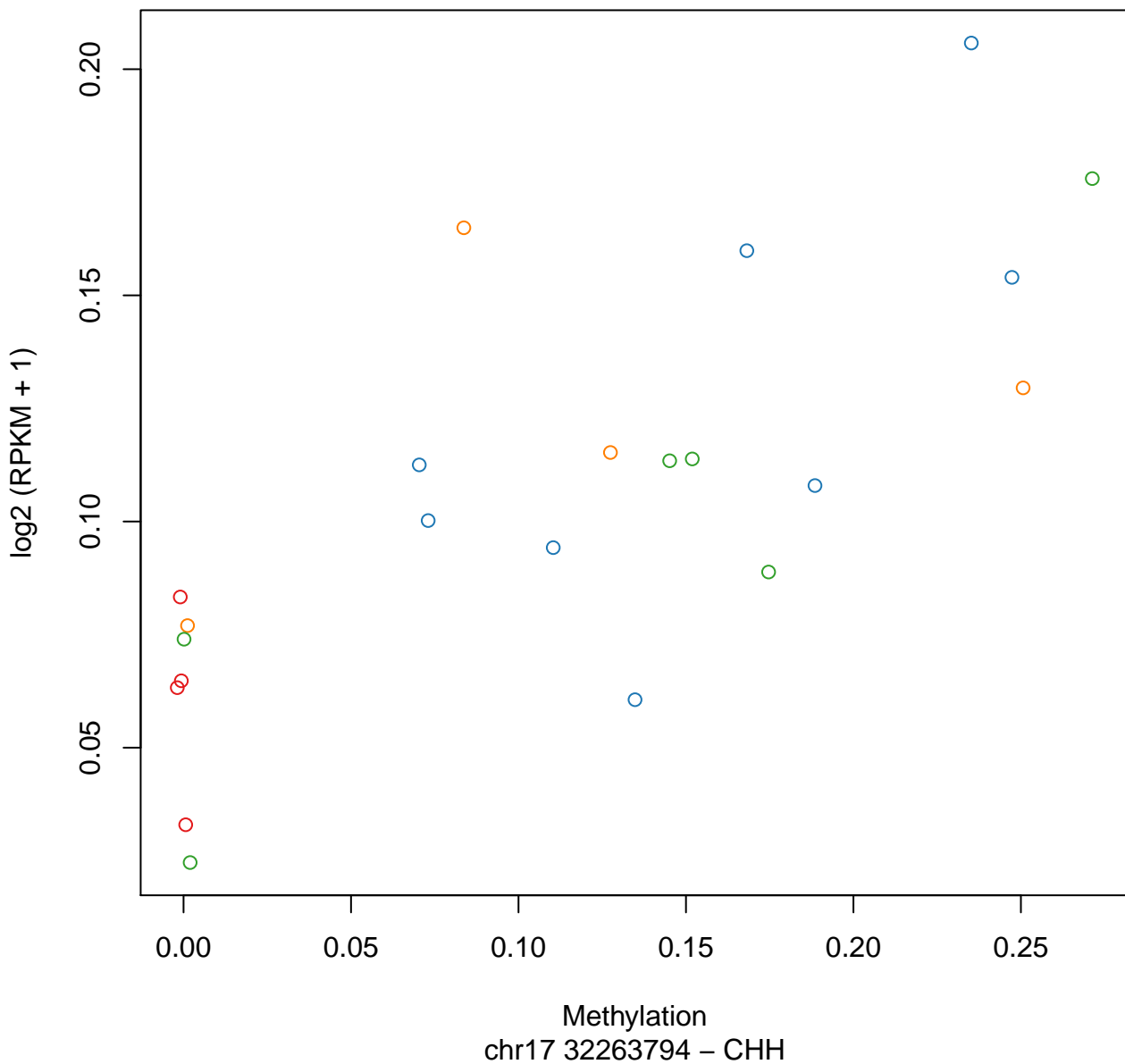
exon ENSE00003793009.1_1 FDR 0.0437
RP11−17M24.3

Methylation
chr17 32263794 − CHH
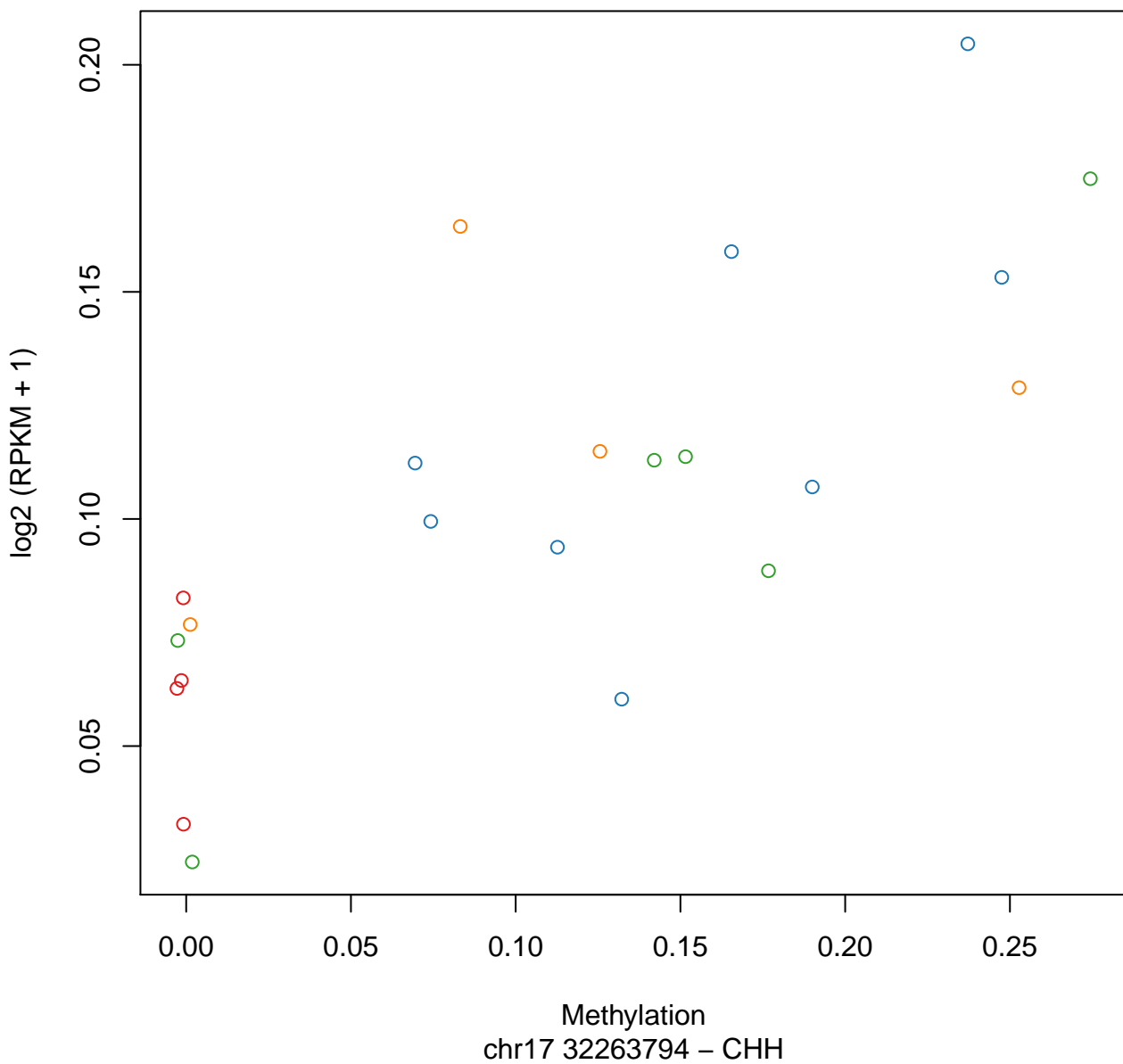
exon ENSE00002088729.1 FDR 0.0456
RNA5SP262

log2 (RPKM + 1)

Methylation
chr8 32049293 + CHG
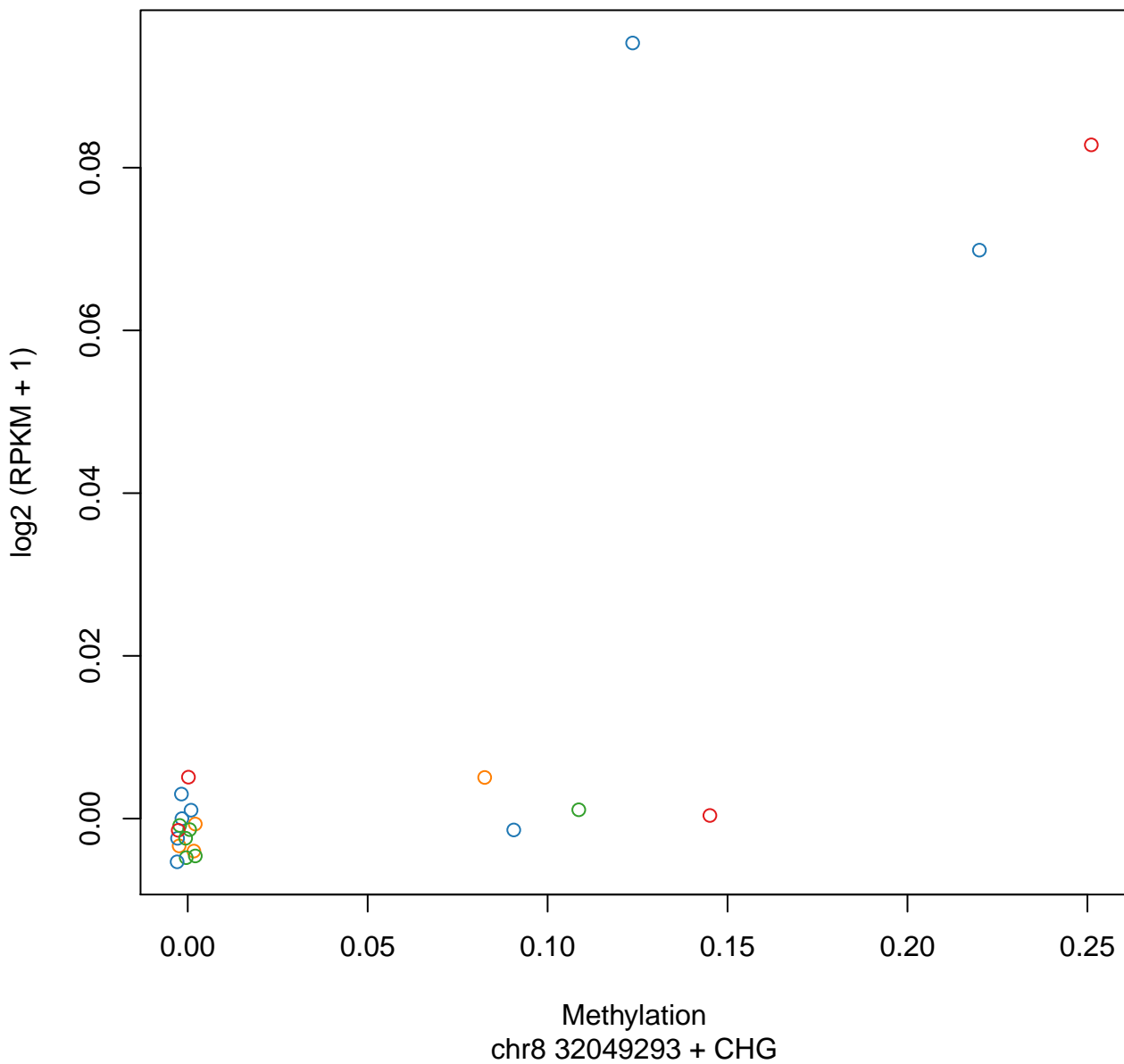
exon ENSE00001188440.1_1 FDR 0.046
COL24A1

Methylation
chr1 86435987 – CHH
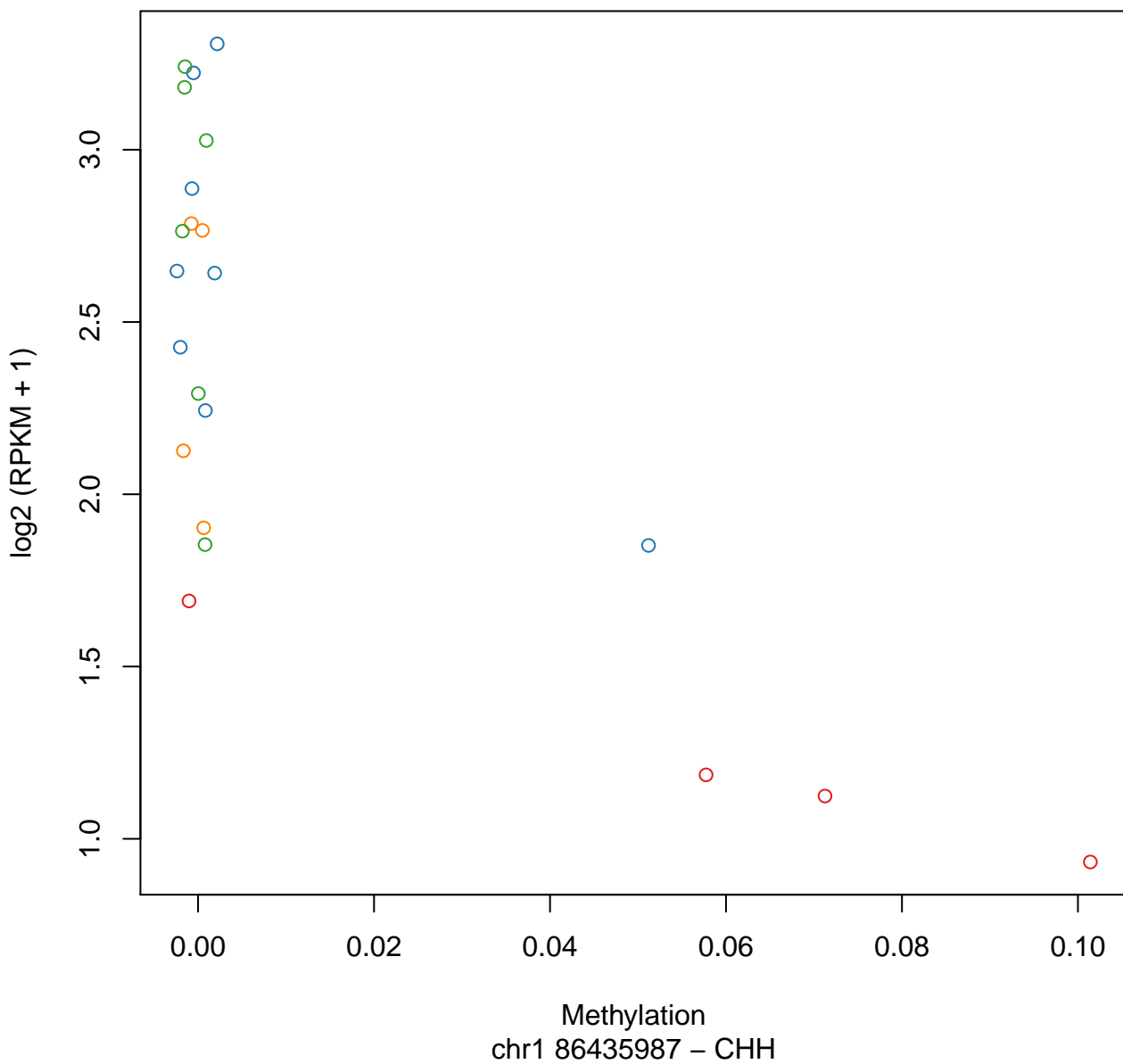
exon ENSE00001613704.1_1 FDR 0.0476
RAB15

log2 (RPKM + 1)

Methylation
chr14 65412146 – CHH
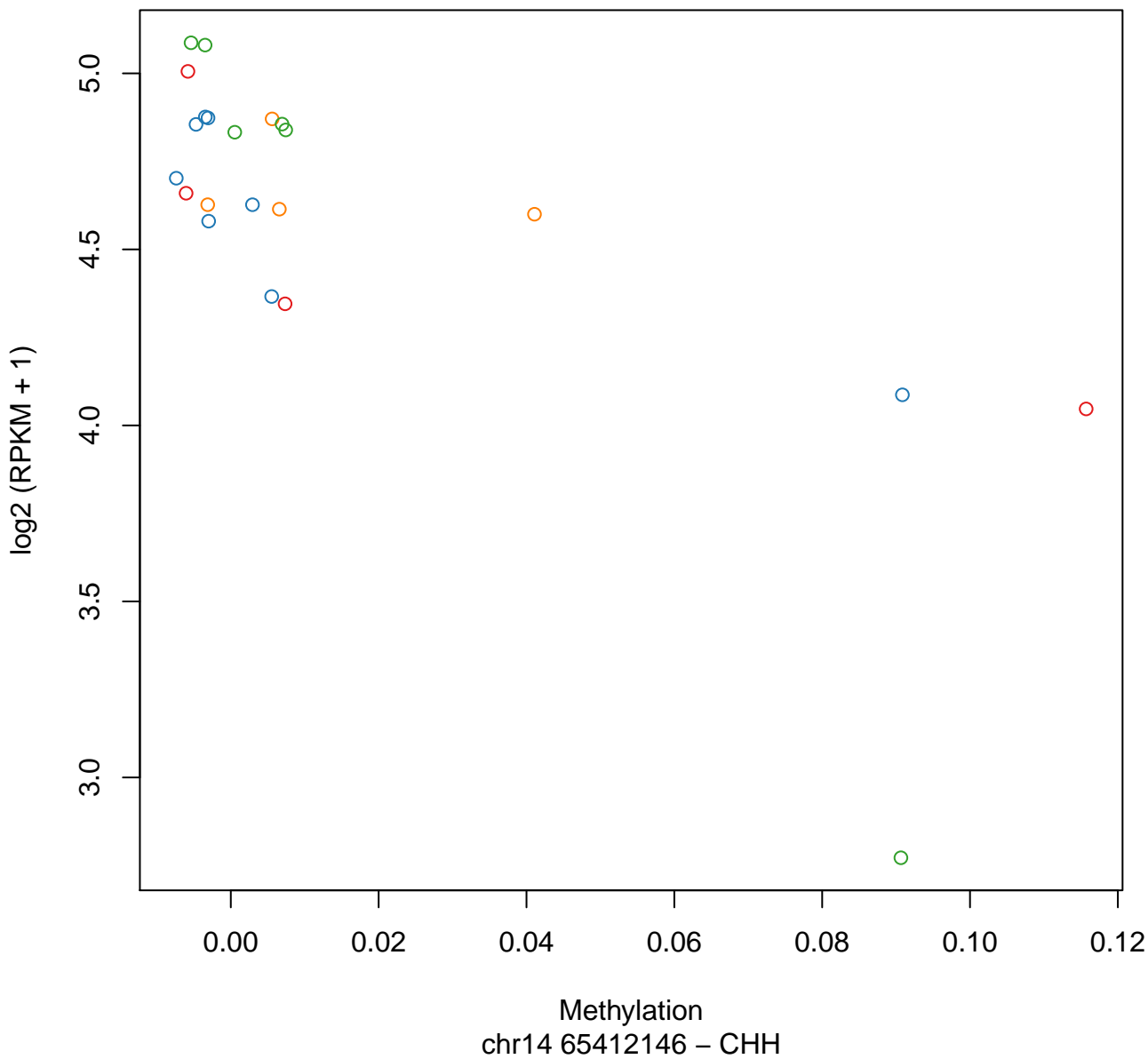
exon ENSE00002200375.2_1 FDR 0.0476
RAB15

Methylation
chr14 65412146 − CHH
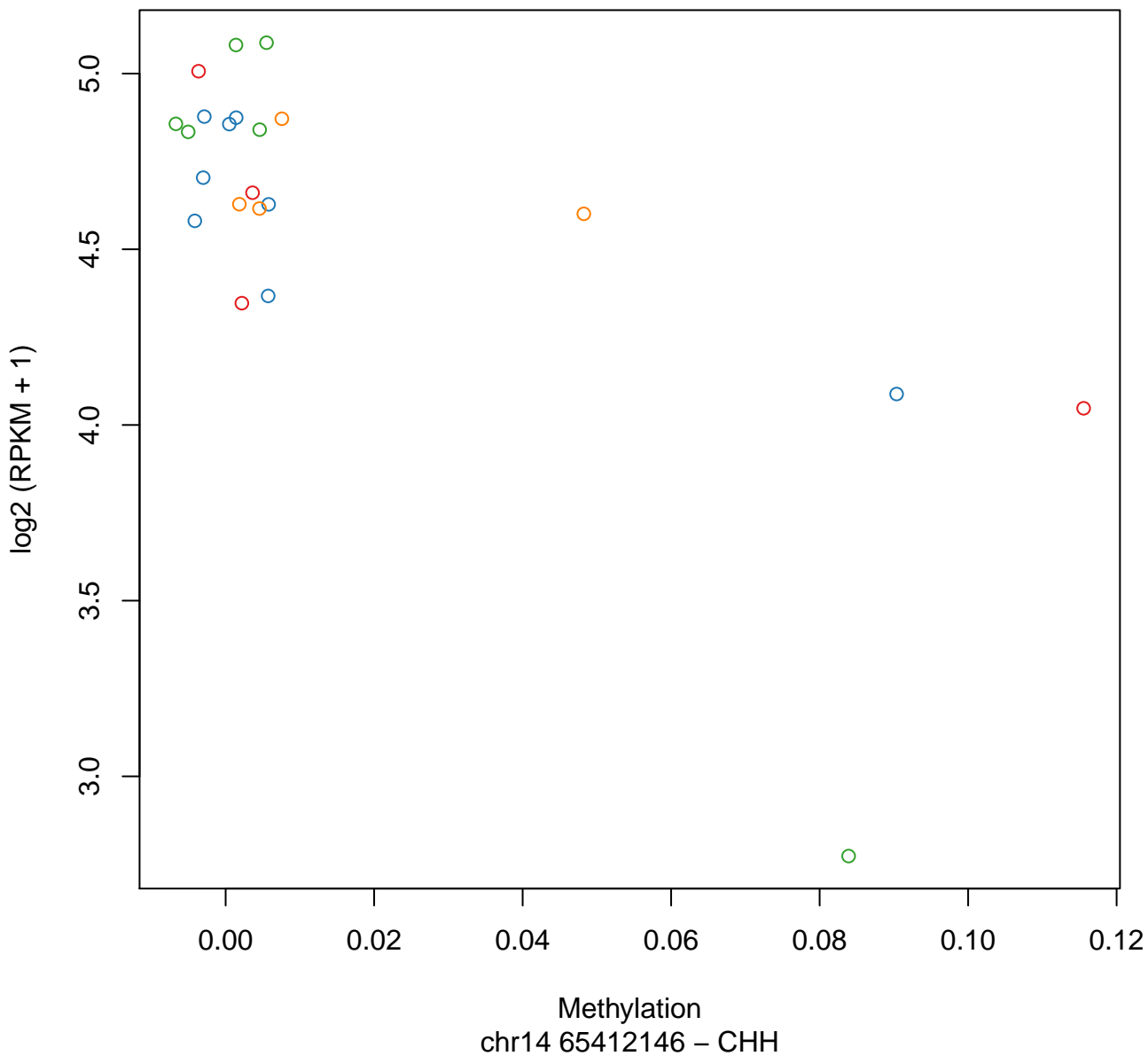
exon ENSE00001489164.1_1 FDR 0.0482
LZTS1

Methylation
chr8 20104130 − CHH
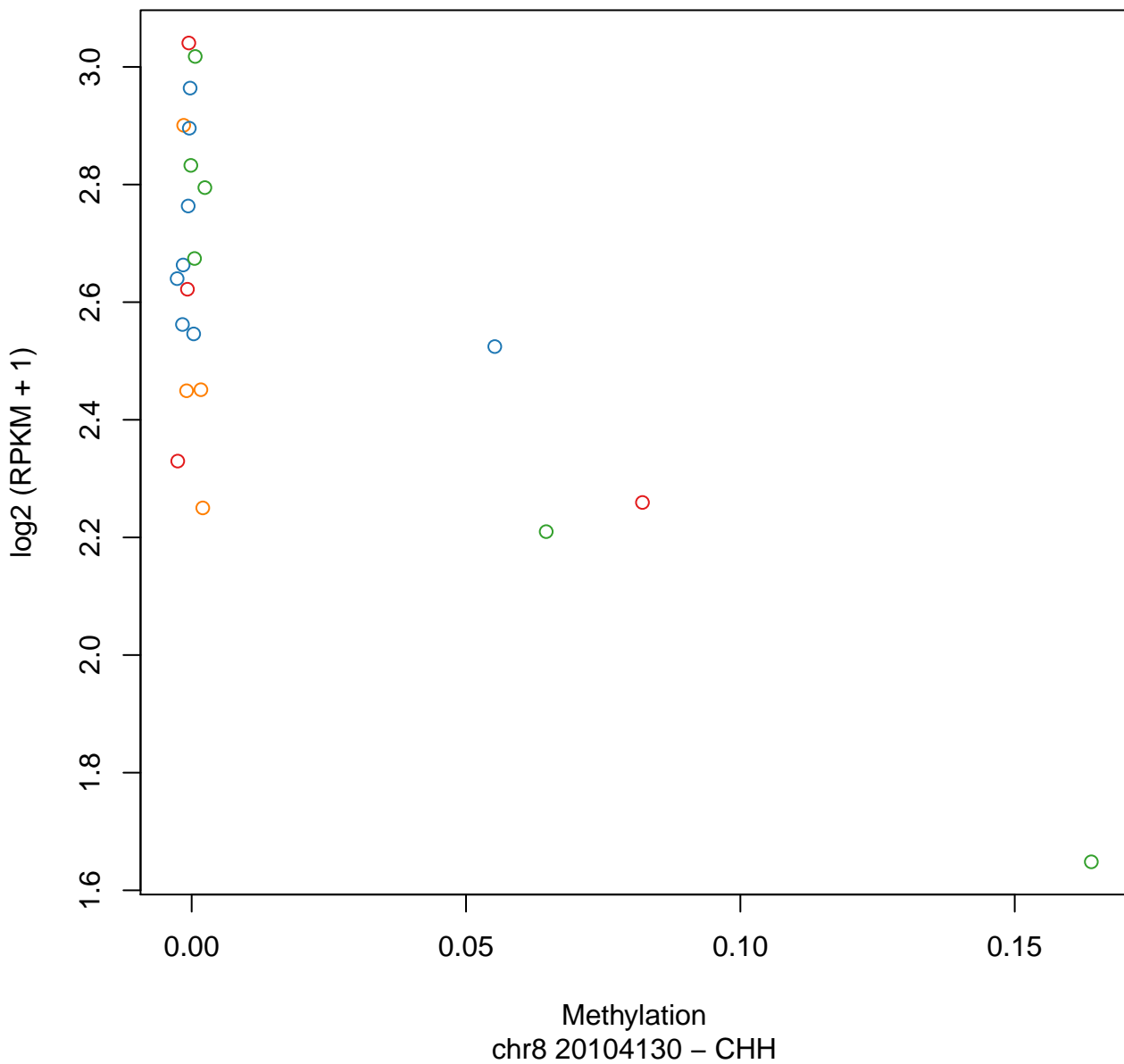
exon ENSE00001871384.1_1 FDR 0.0483
NOS1AP

Methylation
chr1 162336074 + CHH
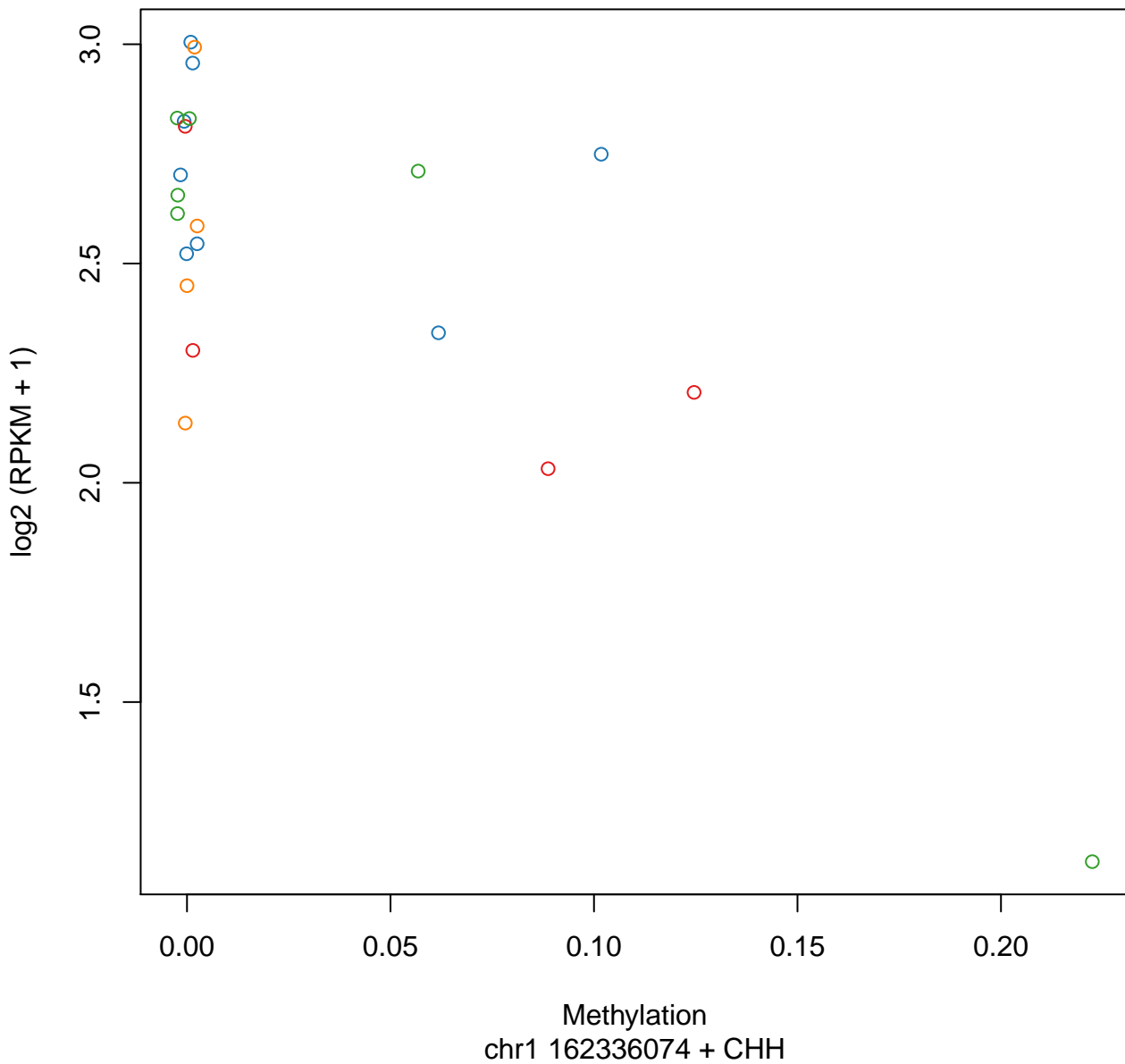
exon ENSE00001967414.2_1 FDR 0.0484
OIP5−AS1

log2 (RPKM + 1)

Methylation
chr15 41598786 + CHG
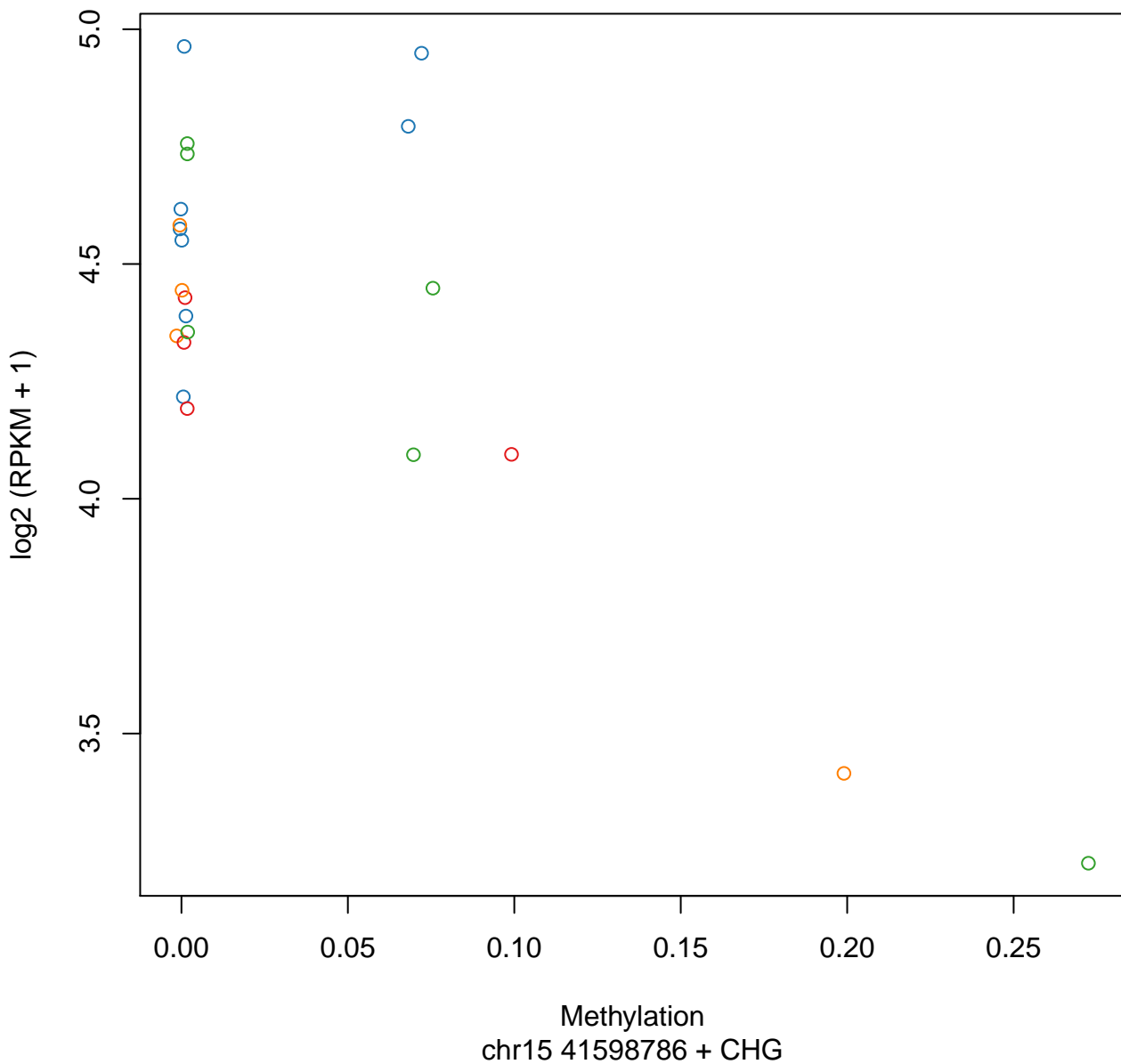
exon ENSE00002286299.2_1 FDR 0.0491
MAPRE2

log2 (RPKM + 1)

Methylation
chr18 32722350 − CHG

exon ENSE00001108348.5_1 FDR 0.0491
MAPRE2

Methylation
chr18 32722350 – CHG