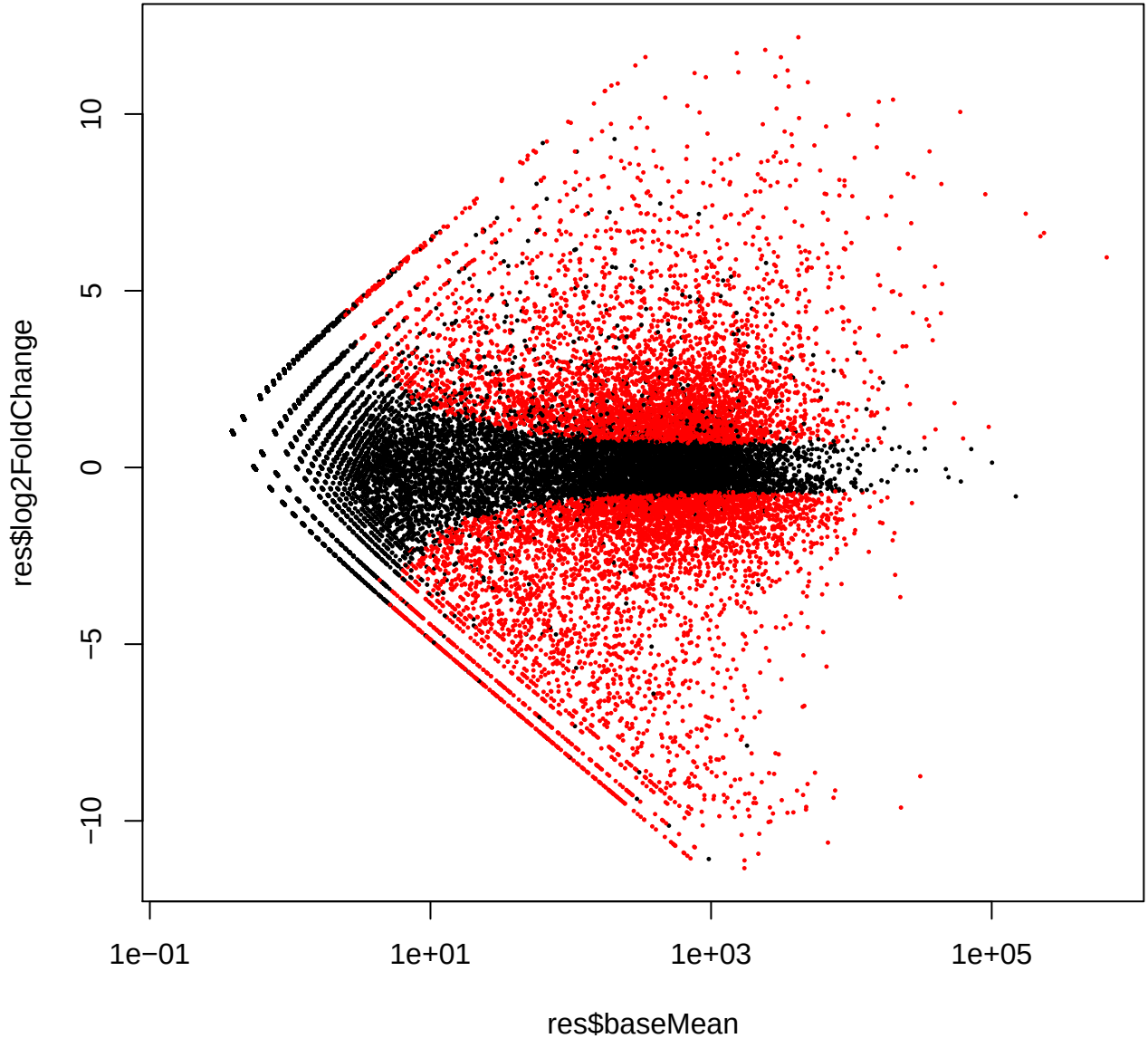
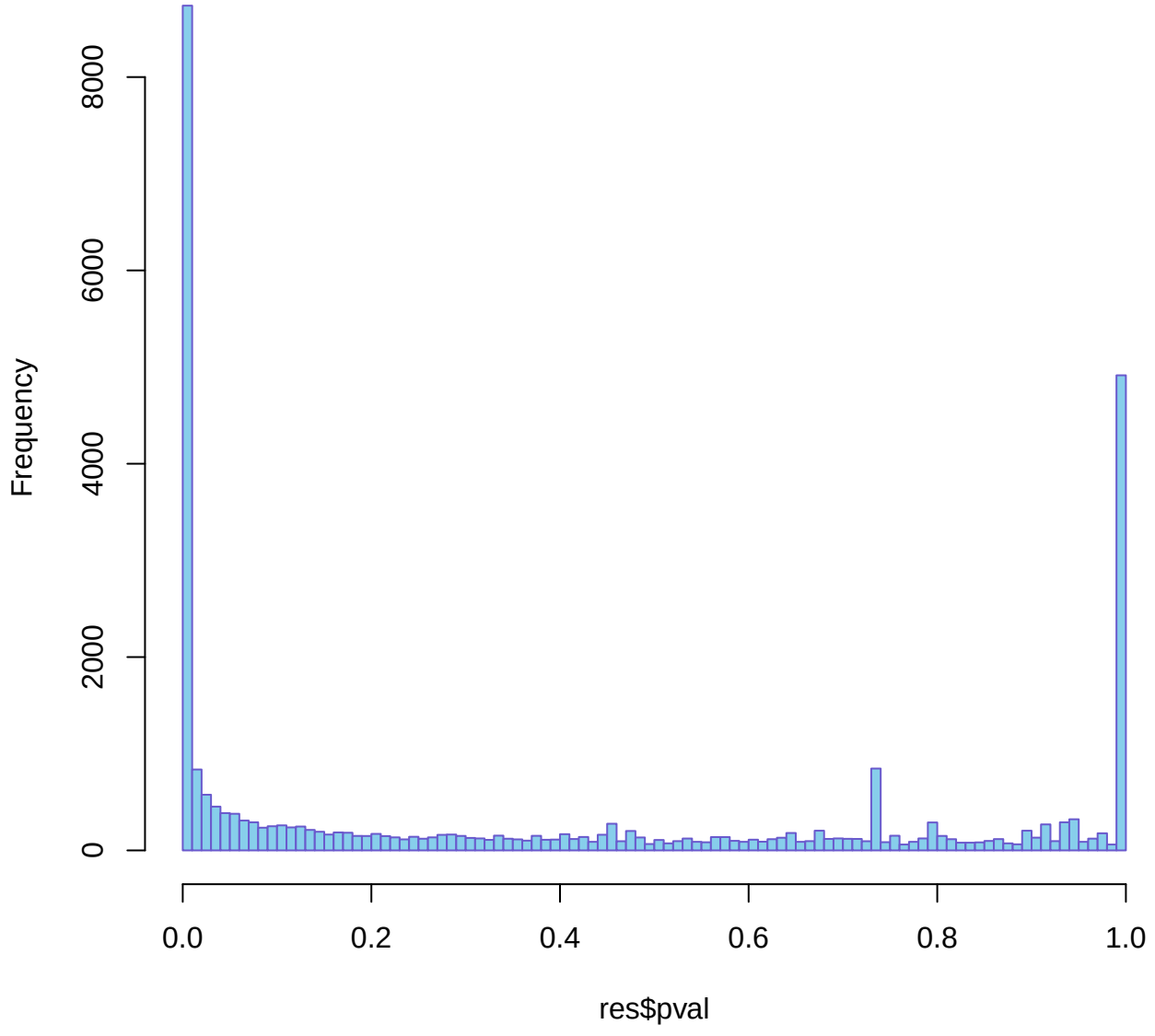
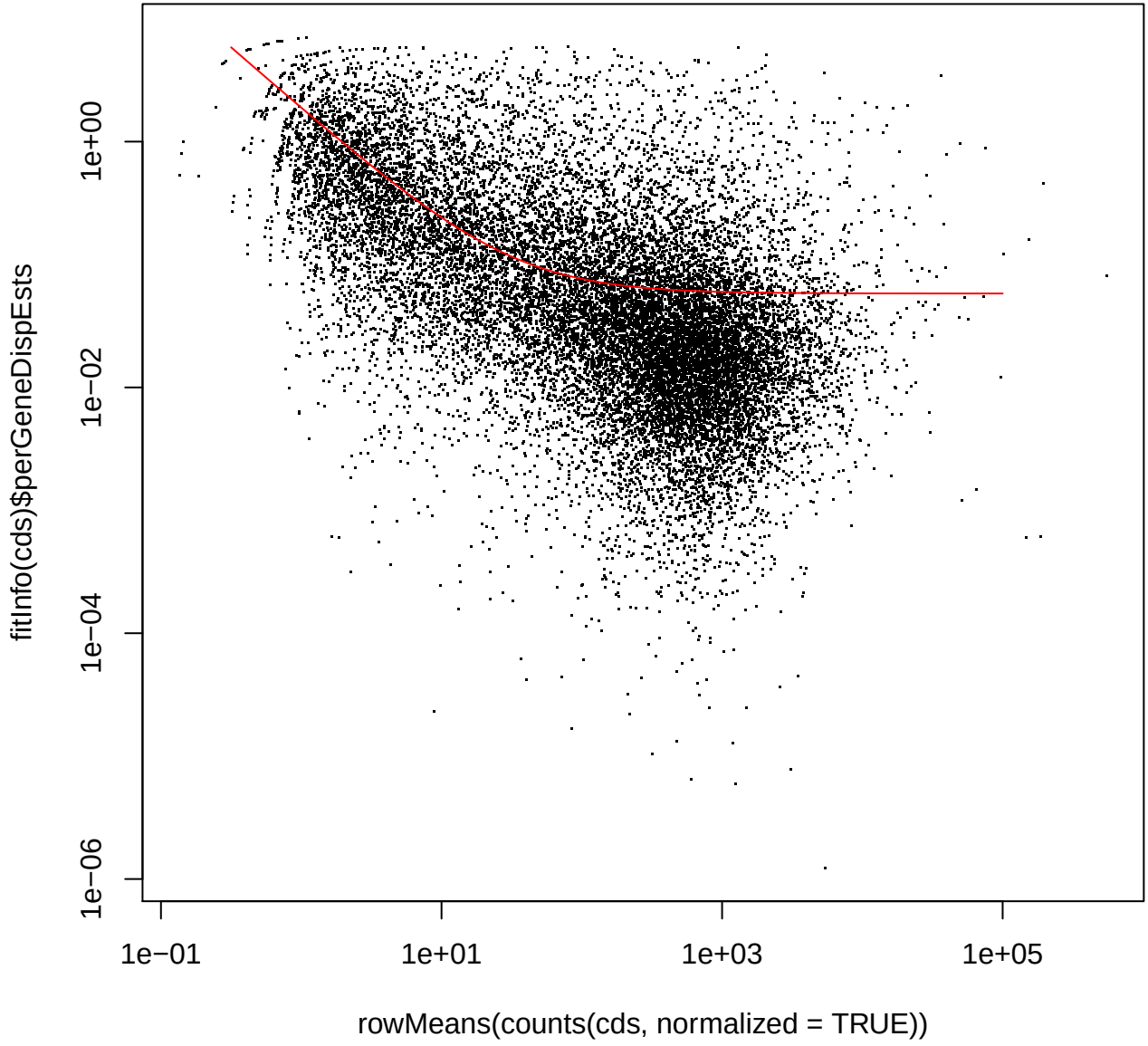

10
5
0
−5
−10
res$log2FoldChange
1e−01
1e+01
1e+03
1e+05
res$baseMean


Frequency
0
2000
4000
6000
8000
0.0
0.2
0.4
0.6
0.8
1.0
res$pval


fitInfo(cds)$perGeneDispEsts
1e+00
1e−02
1e−04
1e−06
1e−01
1e+01
1e+03
1e+05
rowMeans(counts(cds, normalized = TRUE))