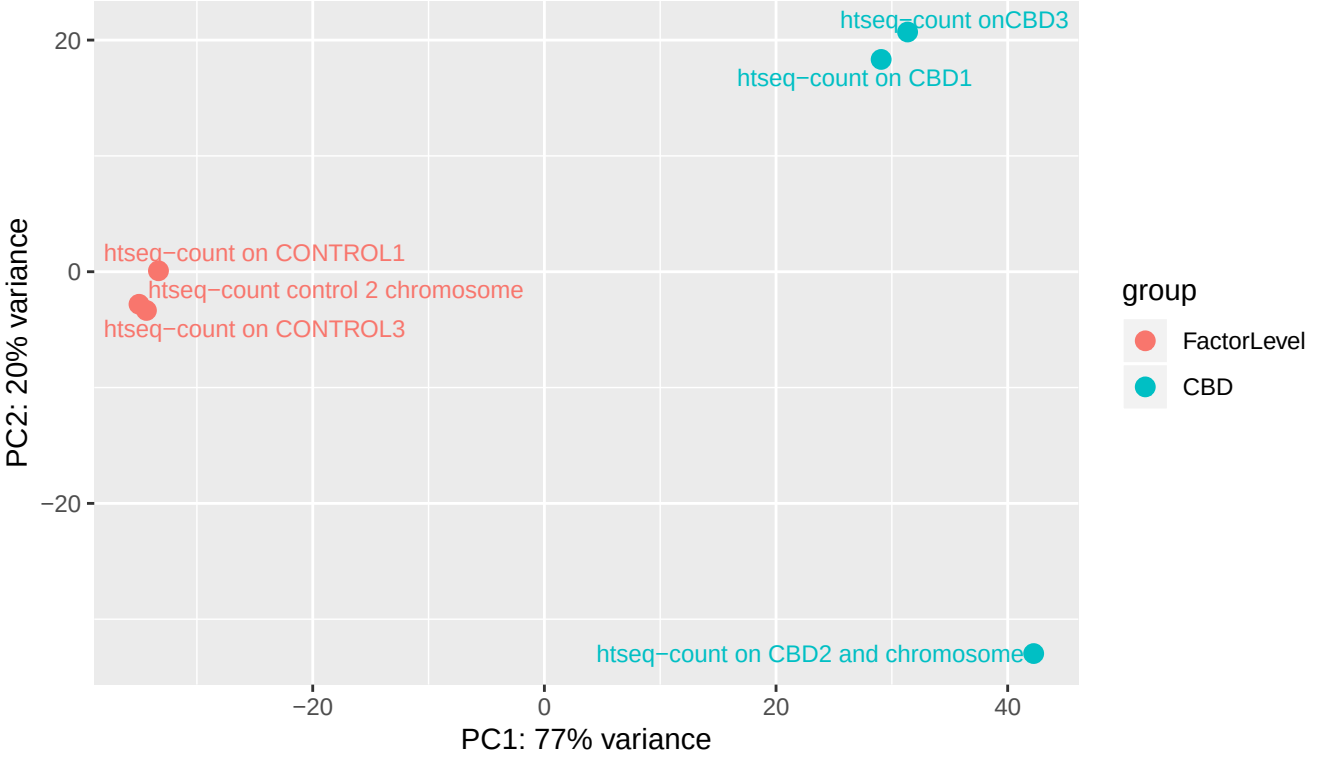
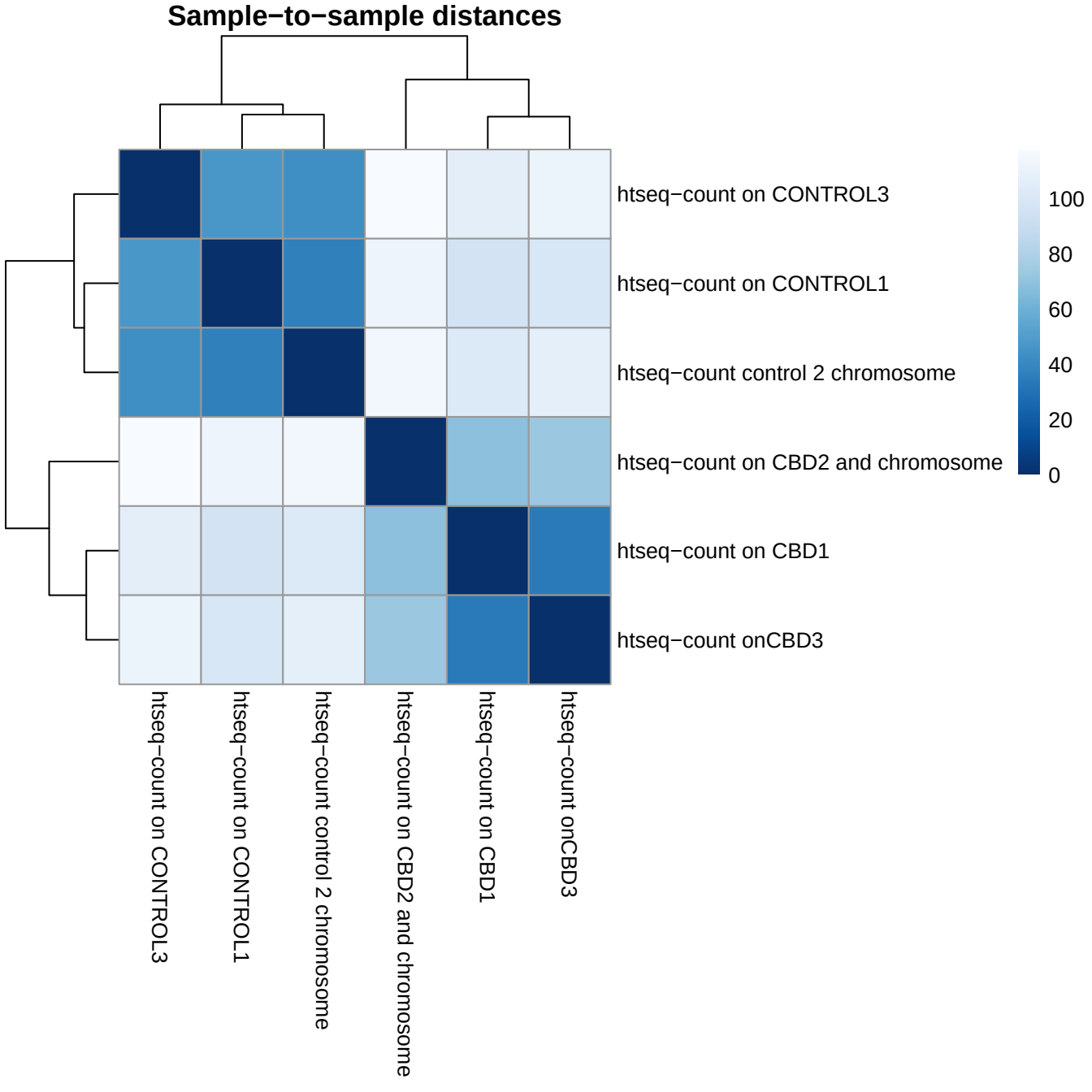
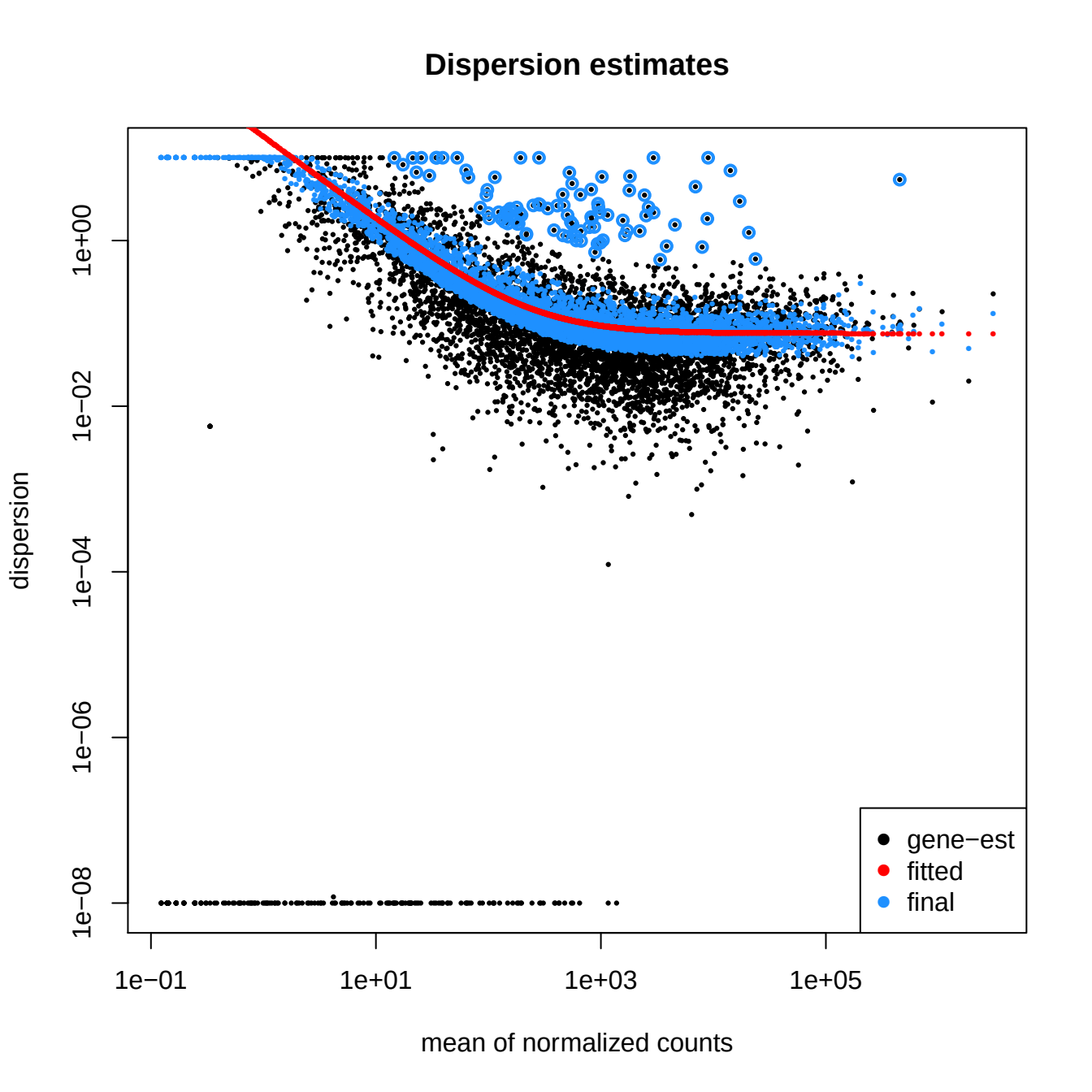
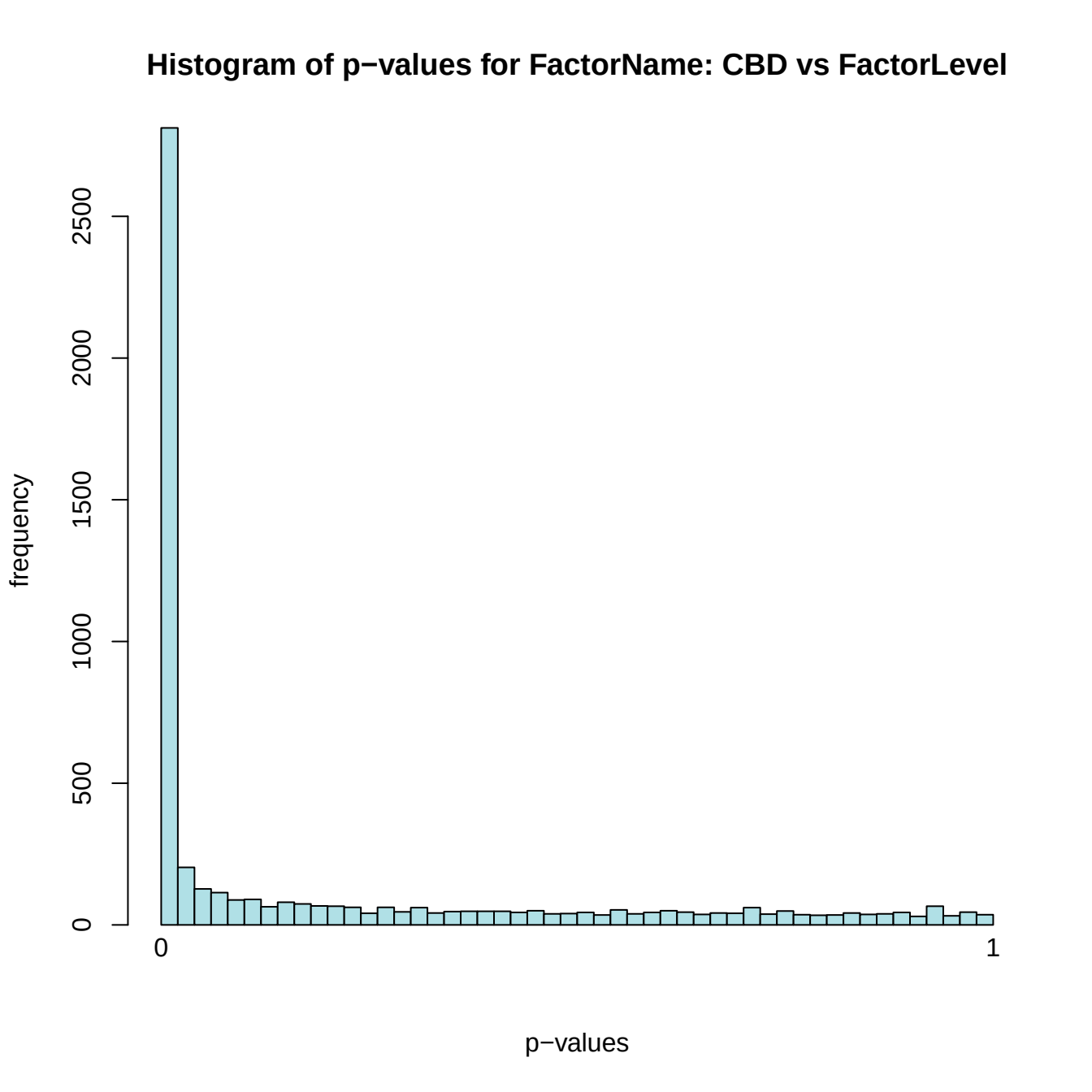
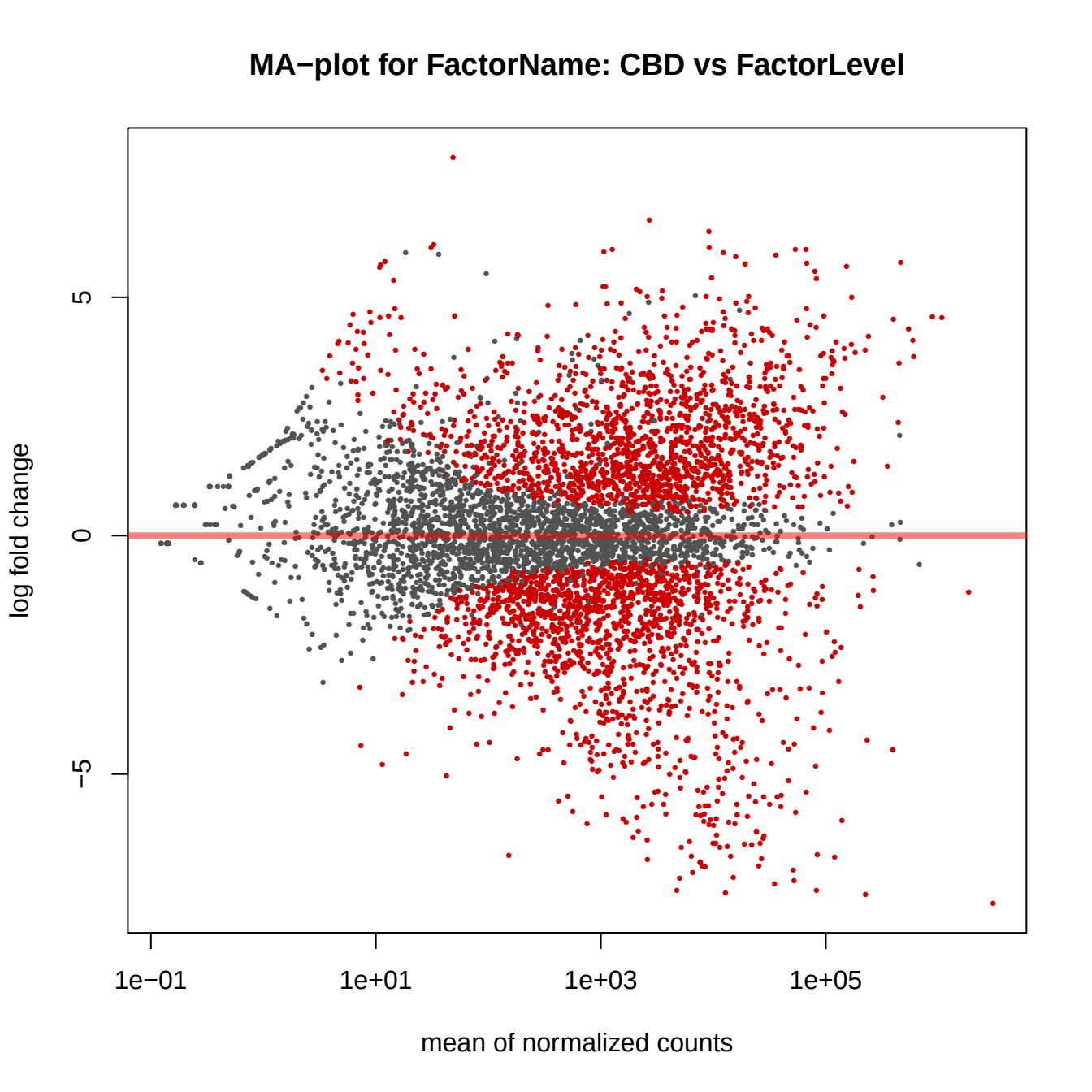

htseq−count onCBD3
htseq−count on CBD1
htseq−count on CONTROL1
htseq−count control 2 chromosome
htseq−count on CONTROL3
htseq−count on CBD2 and chromosome
20
0
−20
−20
0
20
40
PC2: 20% variance
PC1: 77% variance
group
FactorLevel
CBD


Sample−to−sample distances
htseq−count on CONTROL3
htseq−count on CONTROL1
htseq−count control 2 chromosome
htseq−count on CBD2 and chromosome
htseq−count on CBD1
htseq−count onCBD3
100
80
60
40
20
0


Dispersion estimates
dispersion
1e+00
1e−02
1e−04
1e−06
1e−08
1e−01
1e+01
1e+03
1e+05
mean of normalized counts
gene−est
fitted
final


Histogram of p−values for FactorName: CBD vs FactorLevel
frequency
0
500
1000
1500
2000
2500
0
1
p−values


MA−plot for FactorName: CBD vs FactorLevel
log fold change
5
0
−5
1e−01
1e+01
1e+03
1e+05
mean of normalized counts