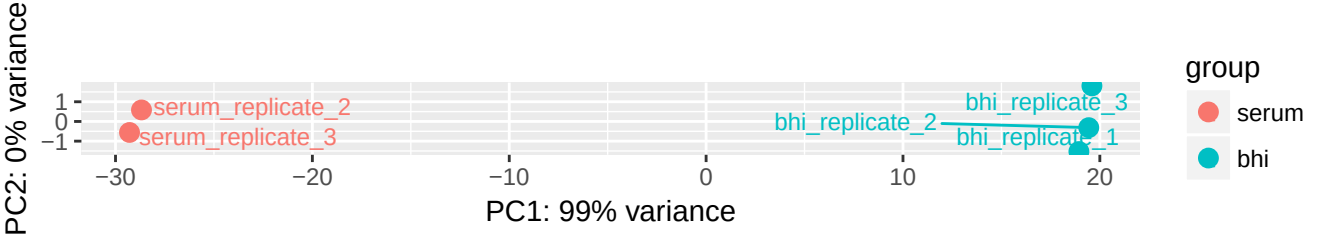
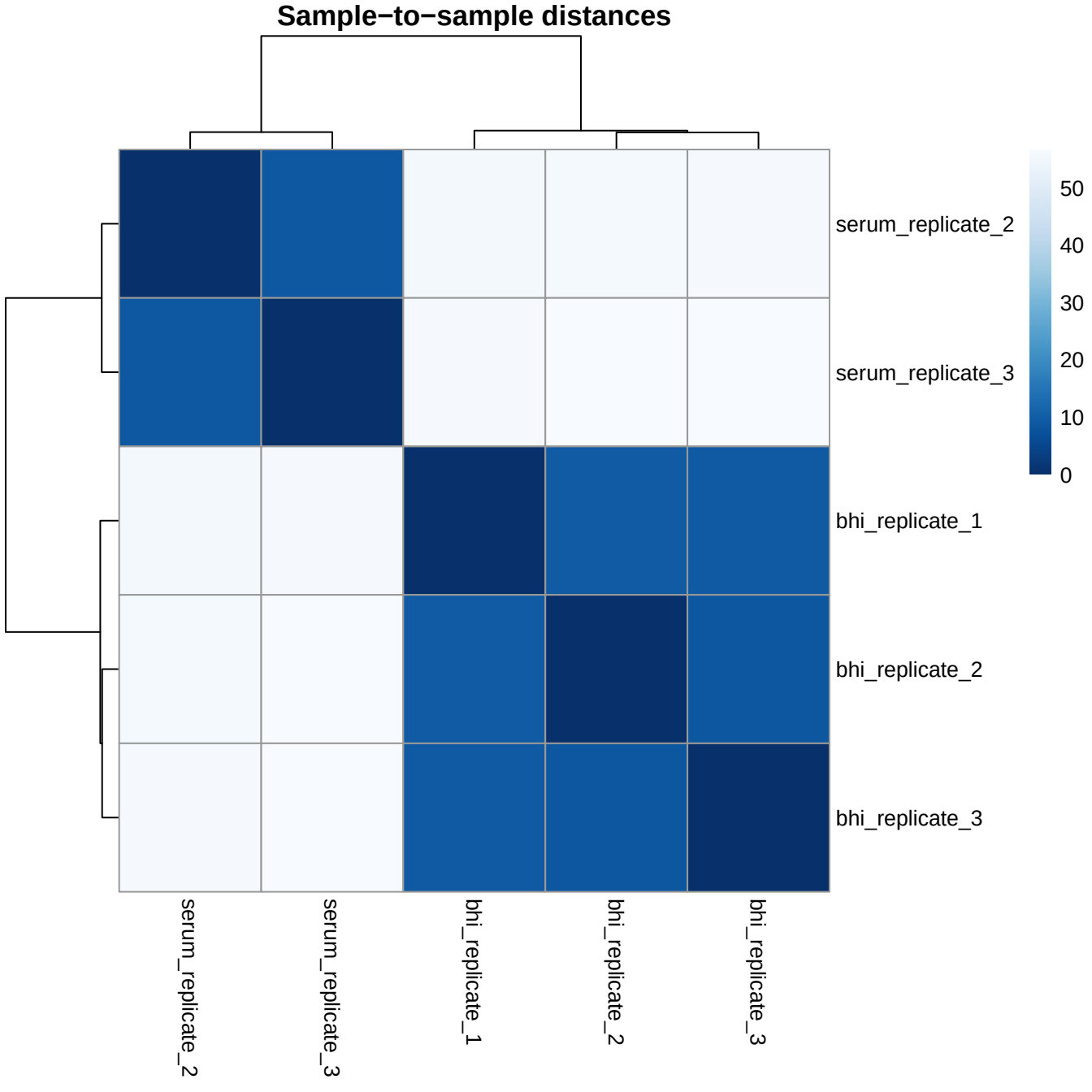
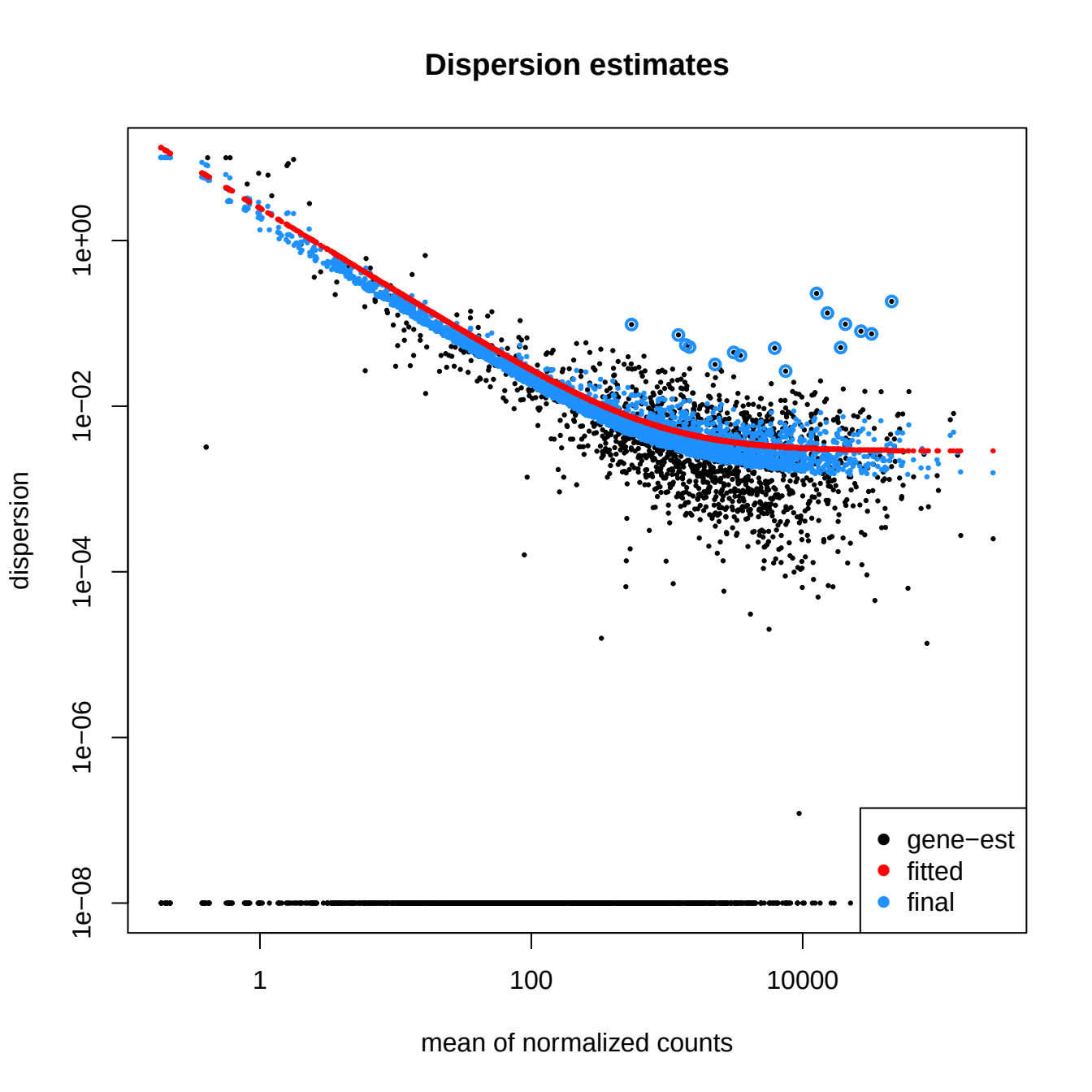
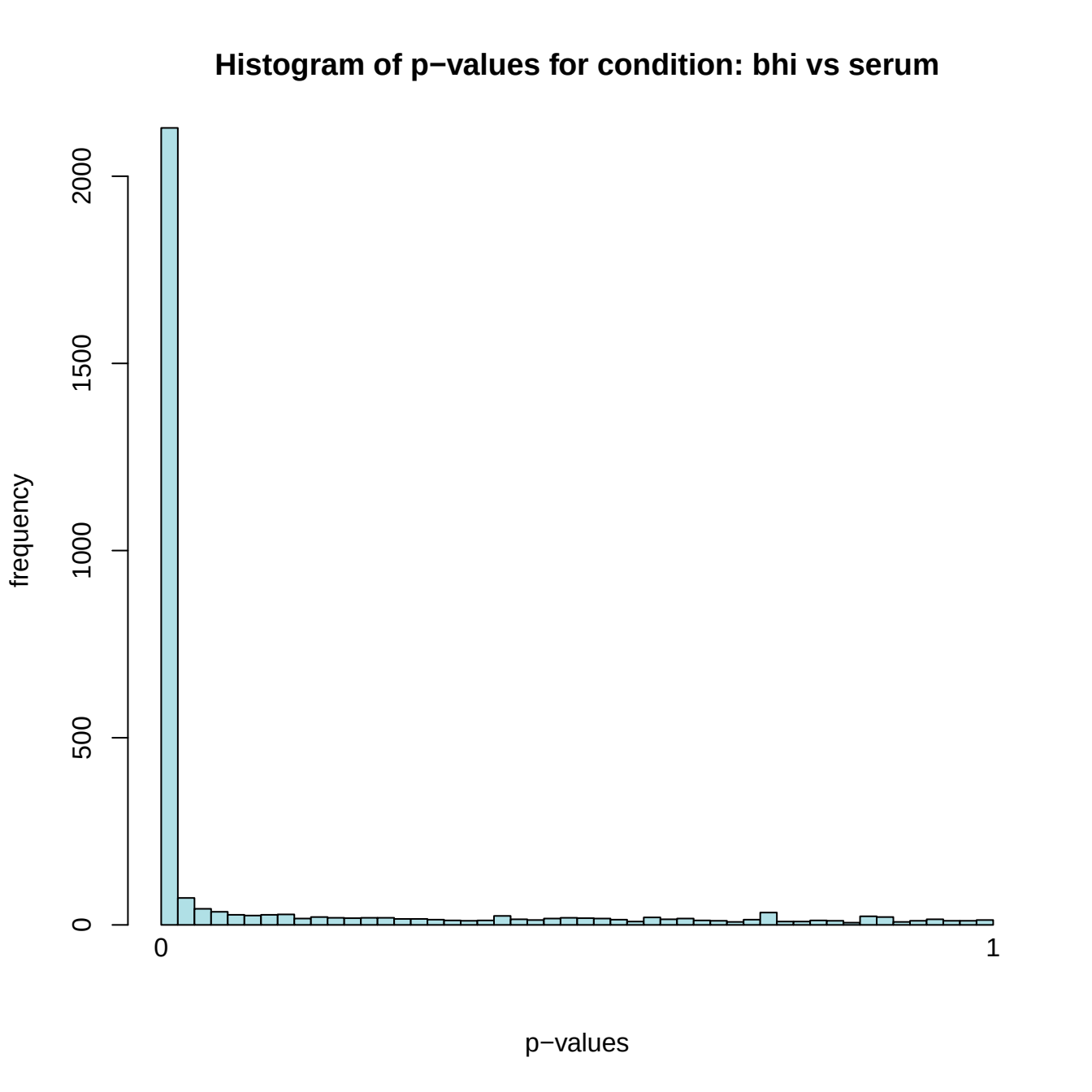
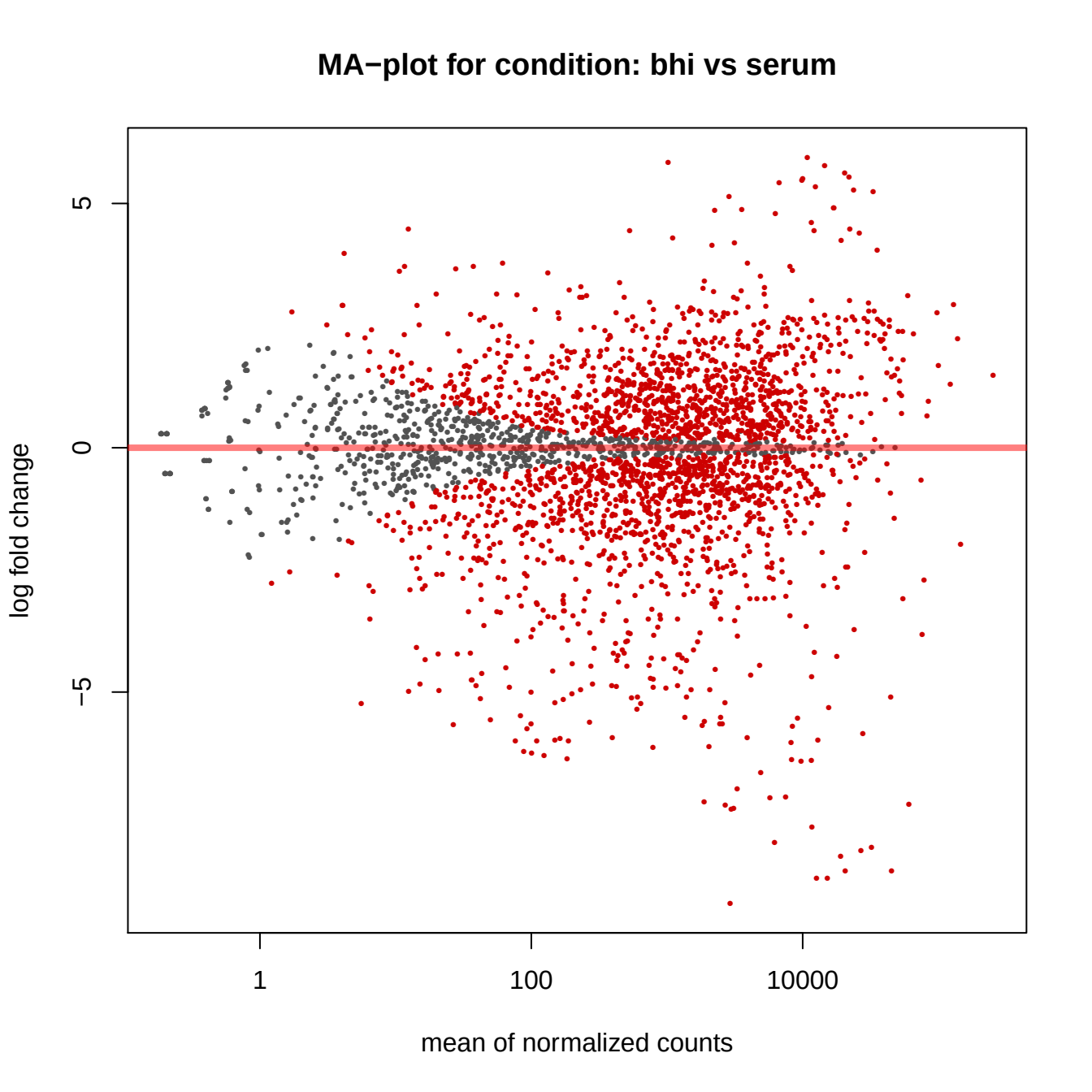

PC2: 0% variance
1
0
−1
serum_replicate_2
serum_replicate_3
bhi_replicate_3
bhi_replicate_2
bhi_replicate_1
−30
−20
−10
0
10
20
PC1: 99% variance
group
serum
bhi


Sample−to−sample distances
serum_replicate_2
serum_replicate_3
bhi_replicate_1
bhi_replicate_2
bhi_replicate_3
50
40
30
20
10
0


Dispersion estimates
dispersion
1e+00
1e−02
1e−04
1e−06
1e−08
1
100
10000
mean of normalized counts
gene−est
fitted
final


Histogram of p−values for condition: bhi vs serum
frequency
0
500
1000
1500
2000
0
1
p−values


MA−plot for condition: bhi vs serum
log fold change
5
0
−5
1
100
10000
mean of normalized counts