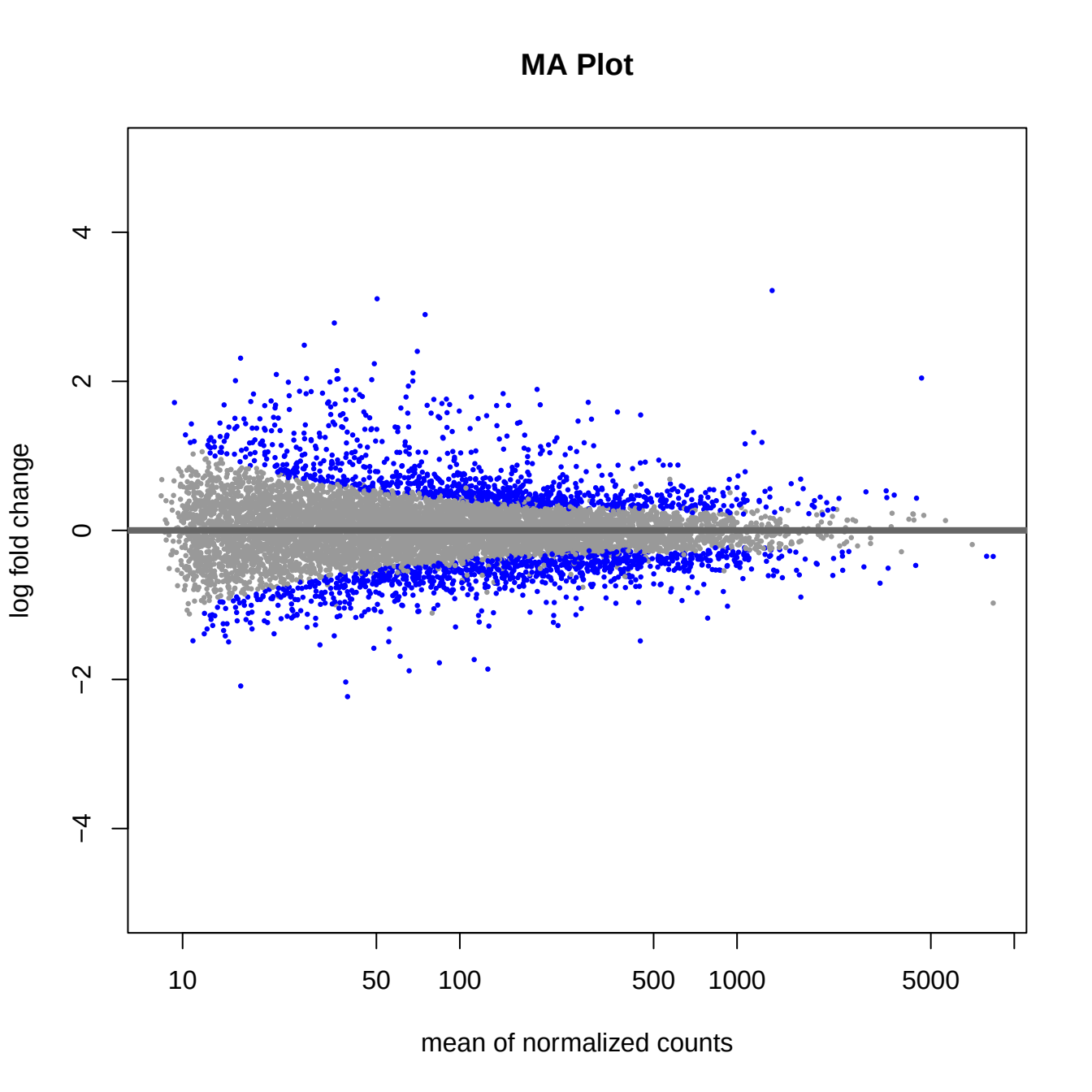
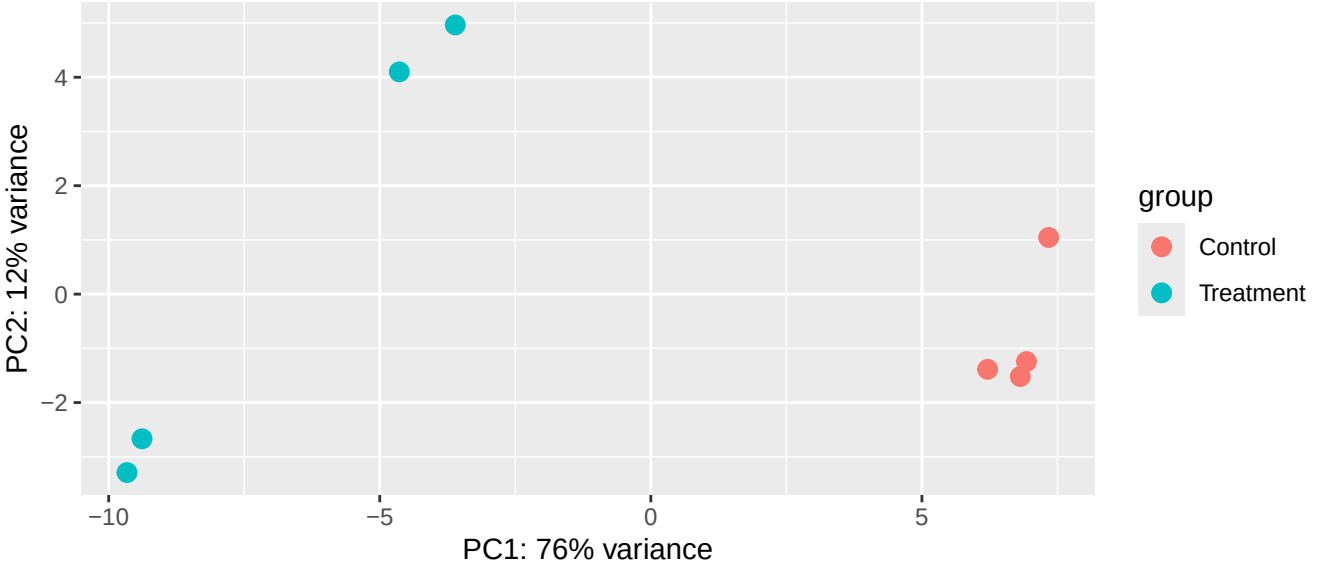
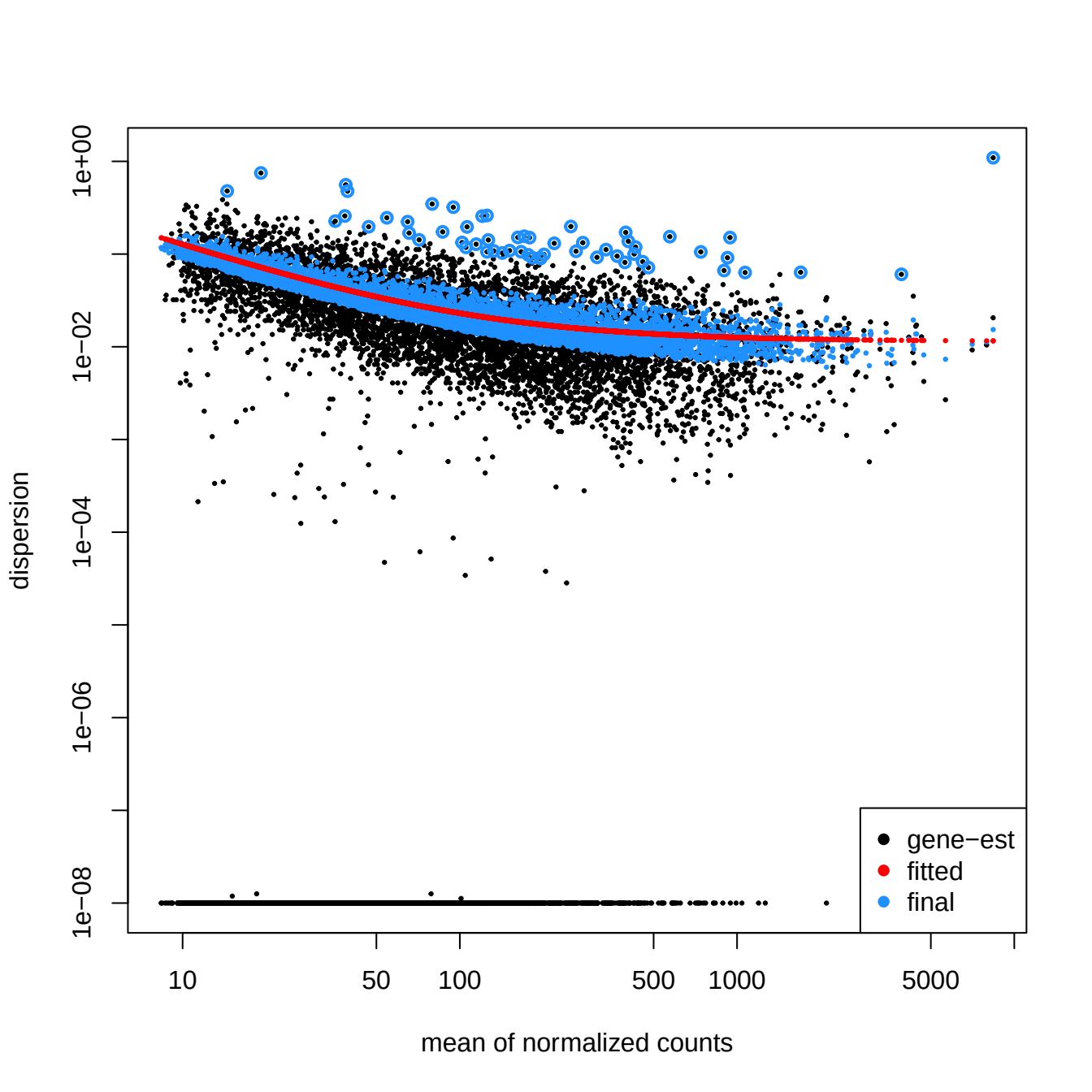
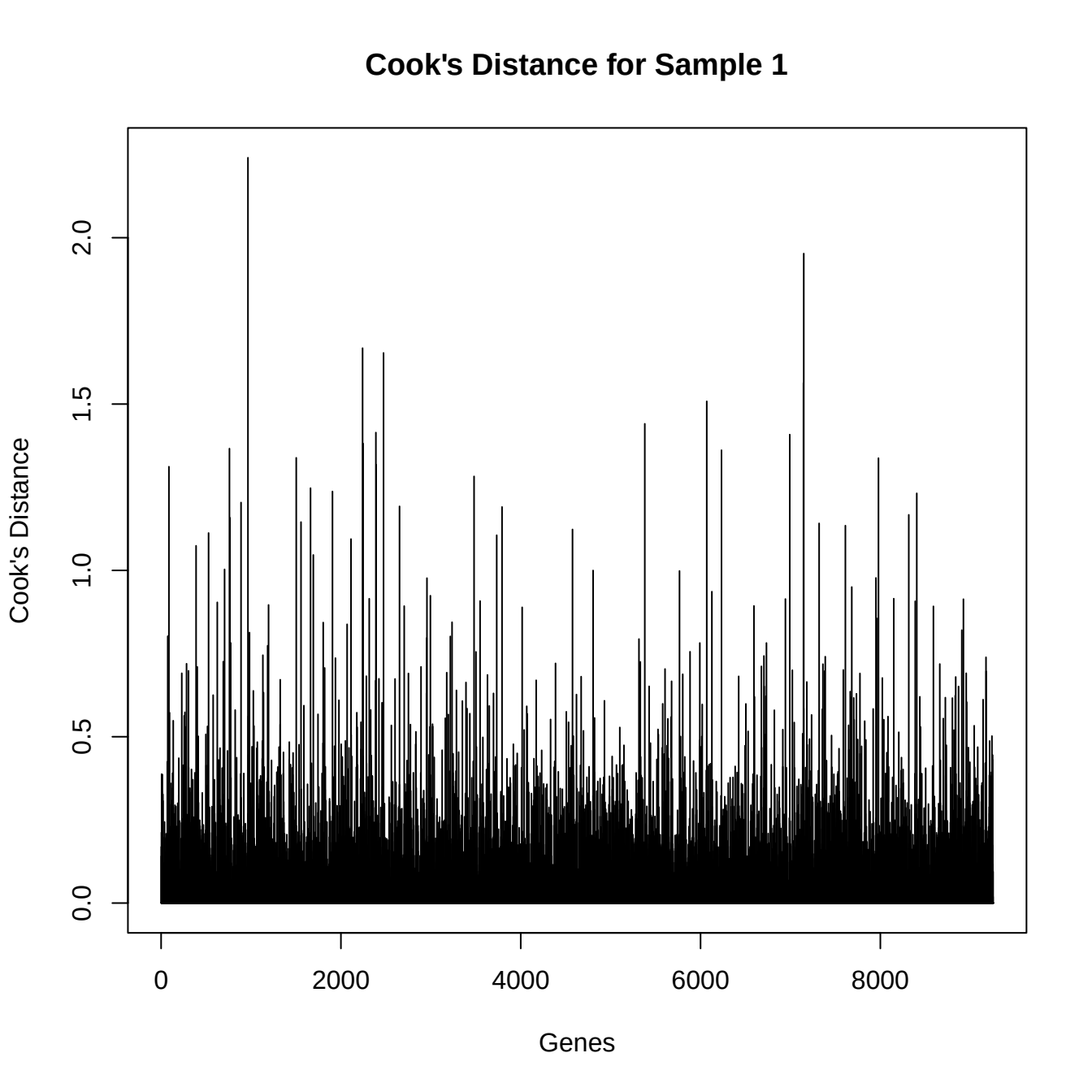
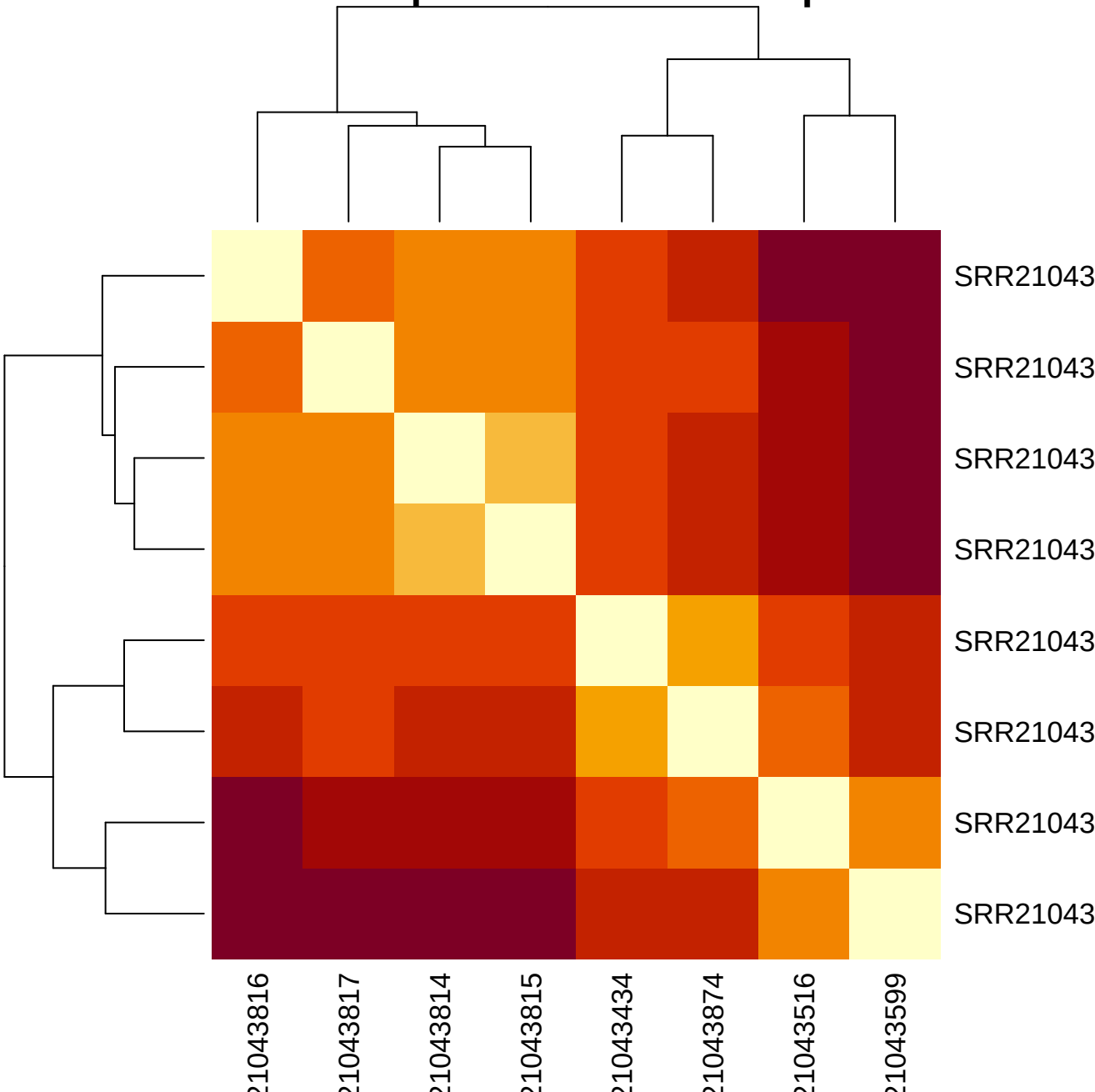

MA Plot
log fold change
4
2
0
−2
−4
10
50
100
500
1000
5000
mean of normalized counts


PC2: 12% variance
4
2
0
−2
−10
−5
0
5
PC1: 76% variance
group
Control
Treatment


dispersion
1e+00
1e−02
1e−04
1e−06
1e−08
10
50
100
500
1000
5000
mean of normalized counts
gene−est
fitted
final


Cook's Distance for Sample 1
Cook's Distance
0.0
0.5
1.0
1.5
2.0
0
2000
4000
6000
8000
Genes


Sample Distance Heatmap
SRR21043
SRR21043
SRR21043
SRR21043
SRR21043
SRR21043
SRR21043
SRR21043
21043816
21043817
21043814
21043815
21043434
21043874
21043516
21043599