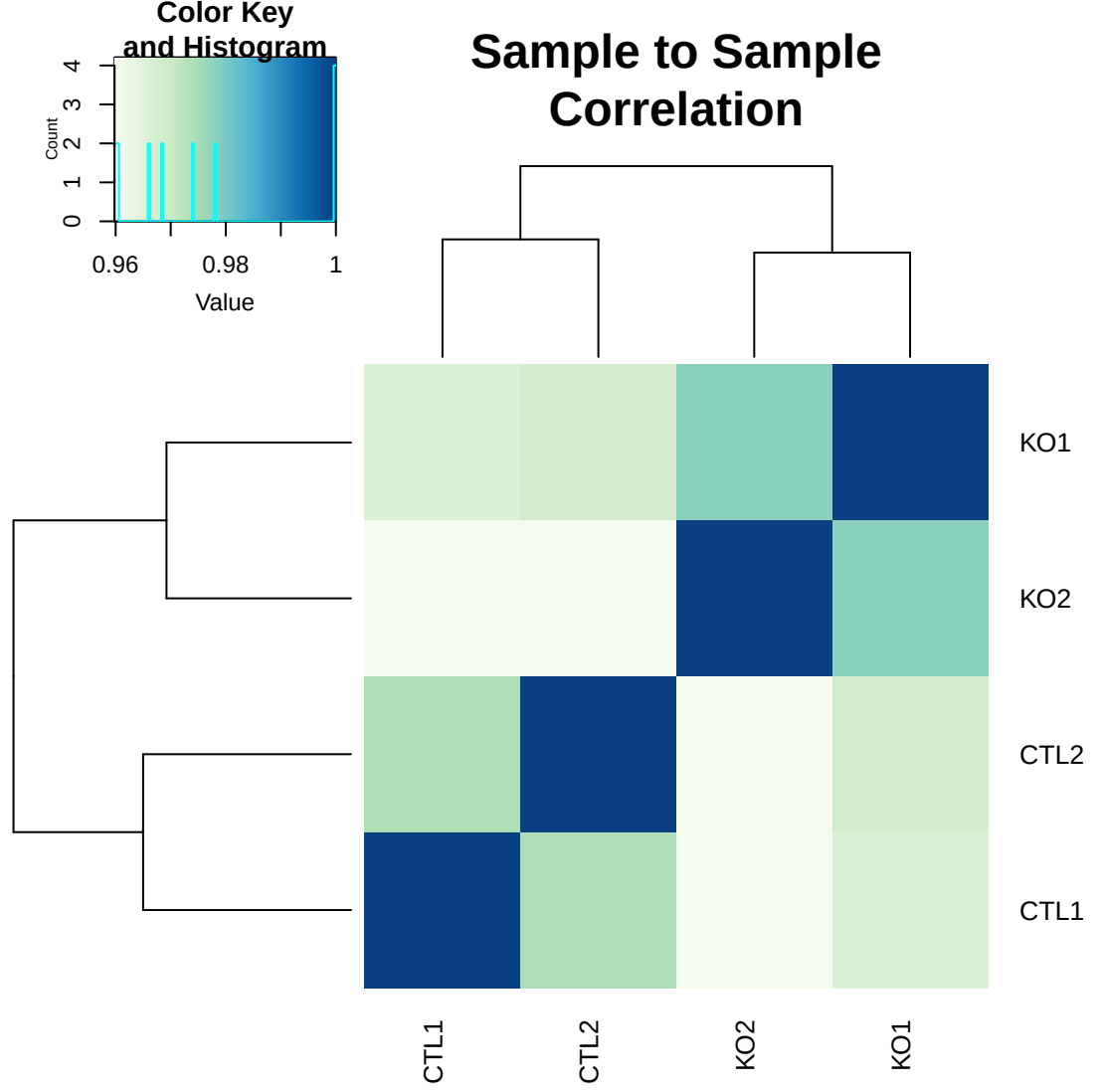
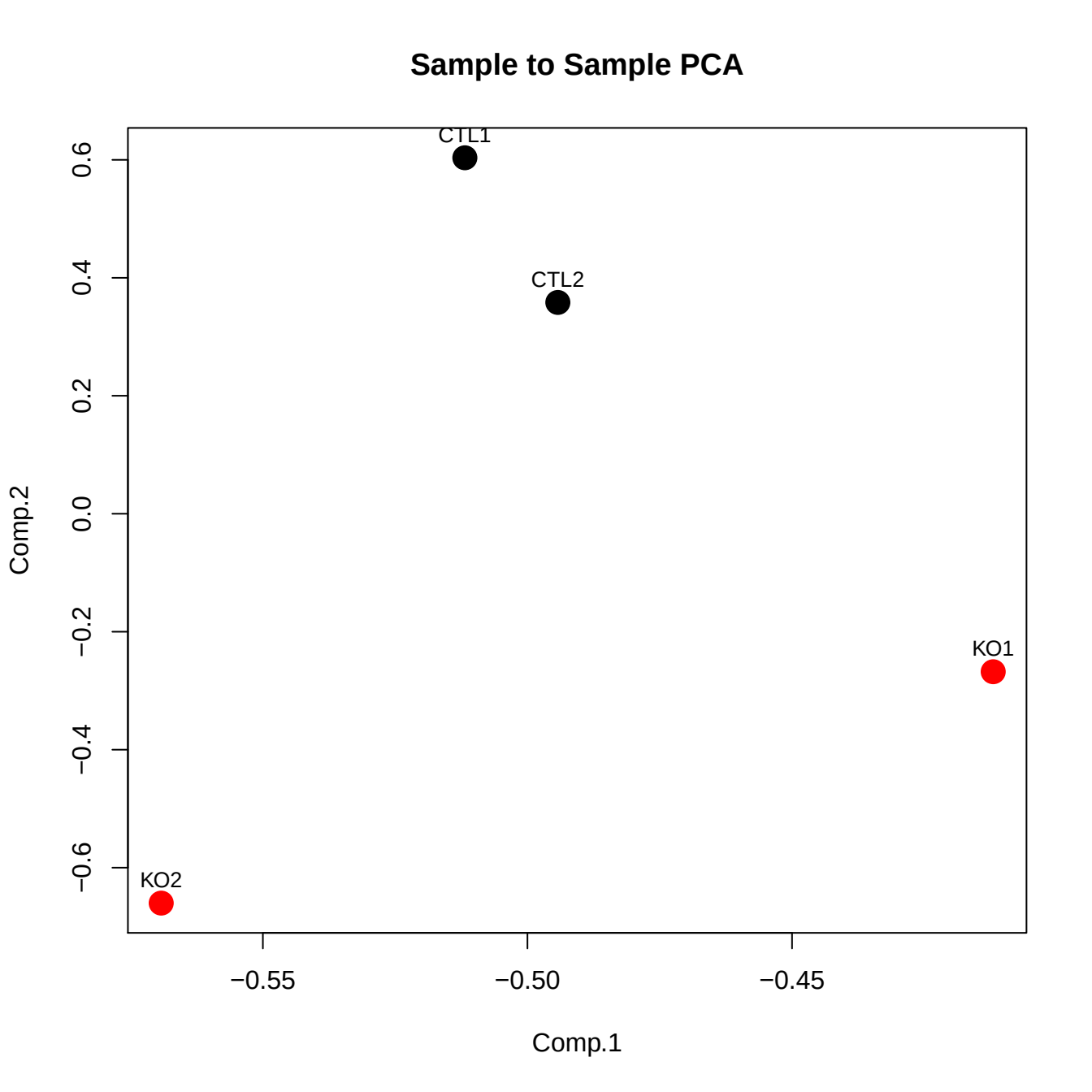
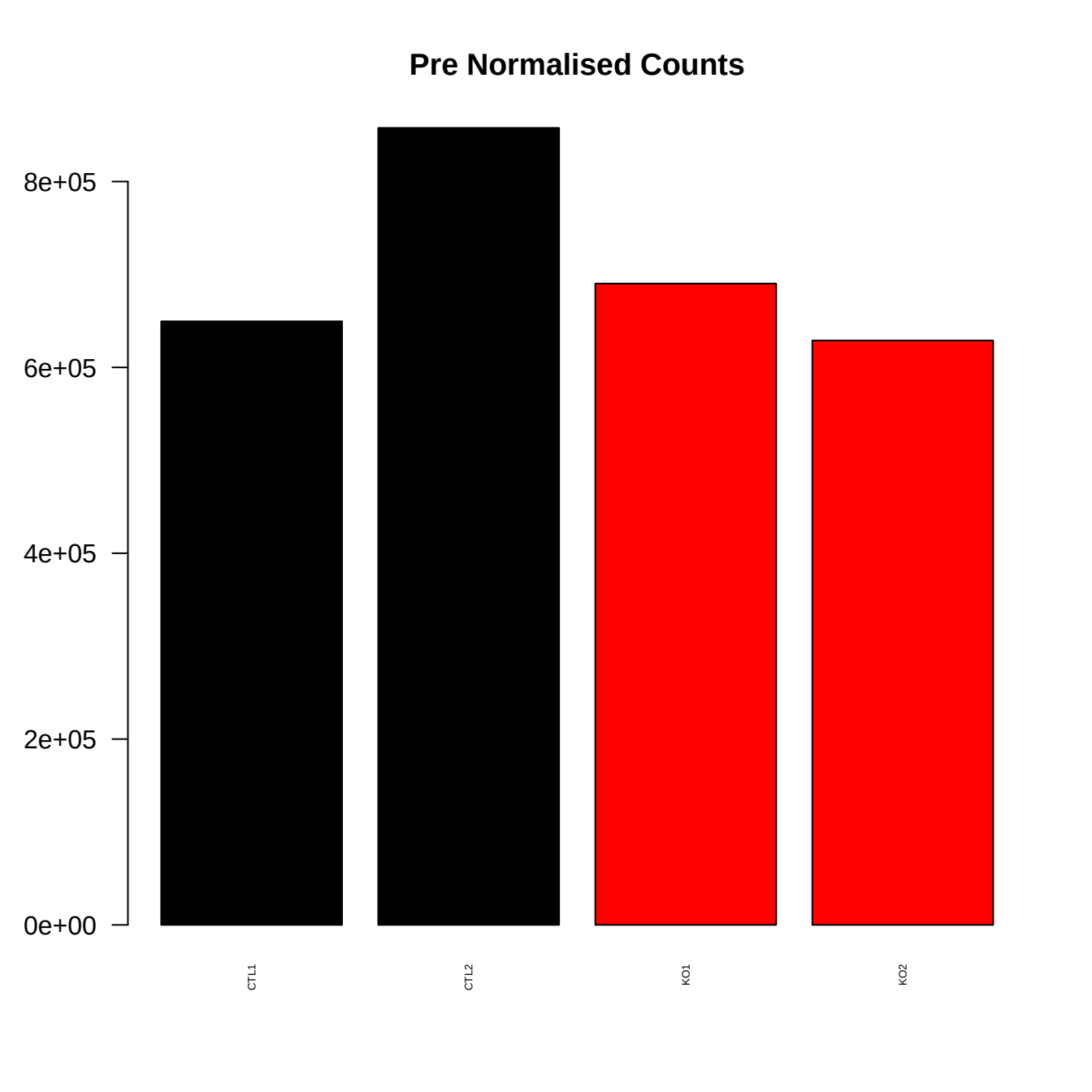
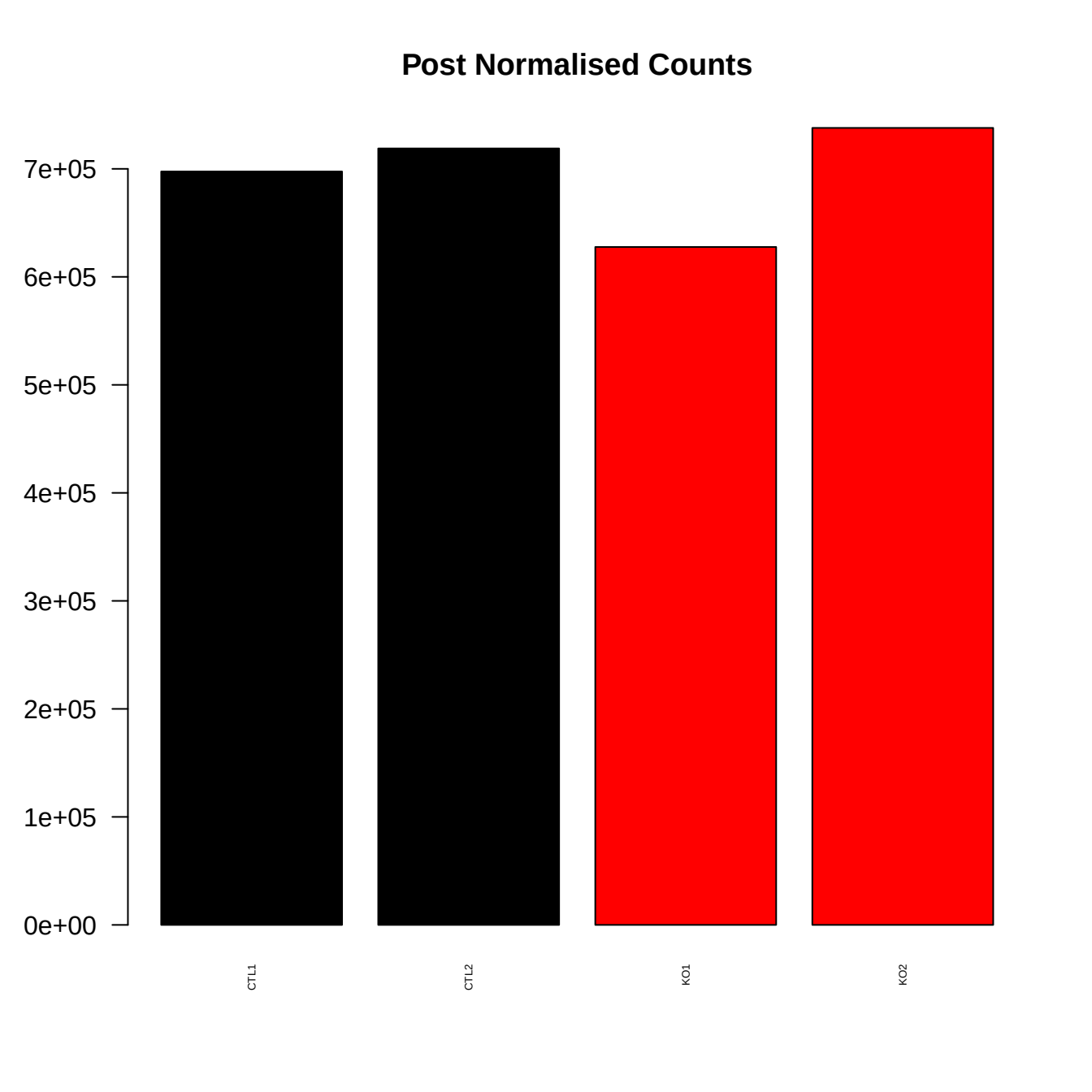
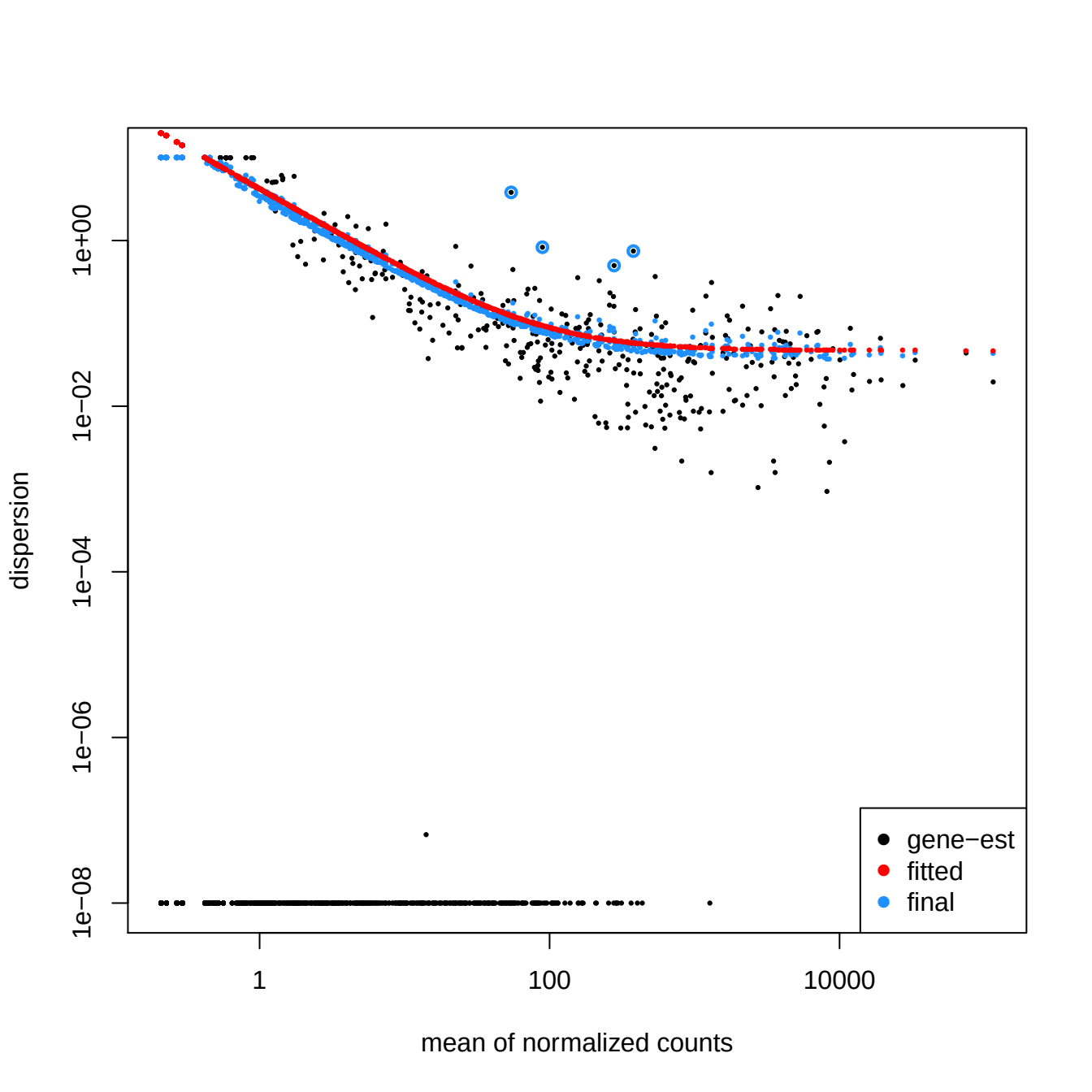

Color Key
and Histogram
Count
0
1
2
3
4
0.96
0.98
1
Value
Sample to Sample
Correlation
KO1
KO2
CTL2
CTL1
CTL1
CTL2
KO2
KO1


Sample to Sample PCA
CTL1
CTL2
KO1
KO2
0.6
0.4
0.2
0.0
−0.2
−0.4
−0.6
Comp.2
−0.55
−0.50
−0.45
Comp.1

CTL
CTL
KO
KO


Pre Normalised Counts
8e+05
6e+05
4e+05
2e+05
0e+00
CTL1
CTL2
KO1
KO2


Post Normalised Counts
7e+05
6e+05
5e+05
4e+05
3e+05
2e+05
1e+05
0e+00
CTL1
CTL2
KO1
KO2


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