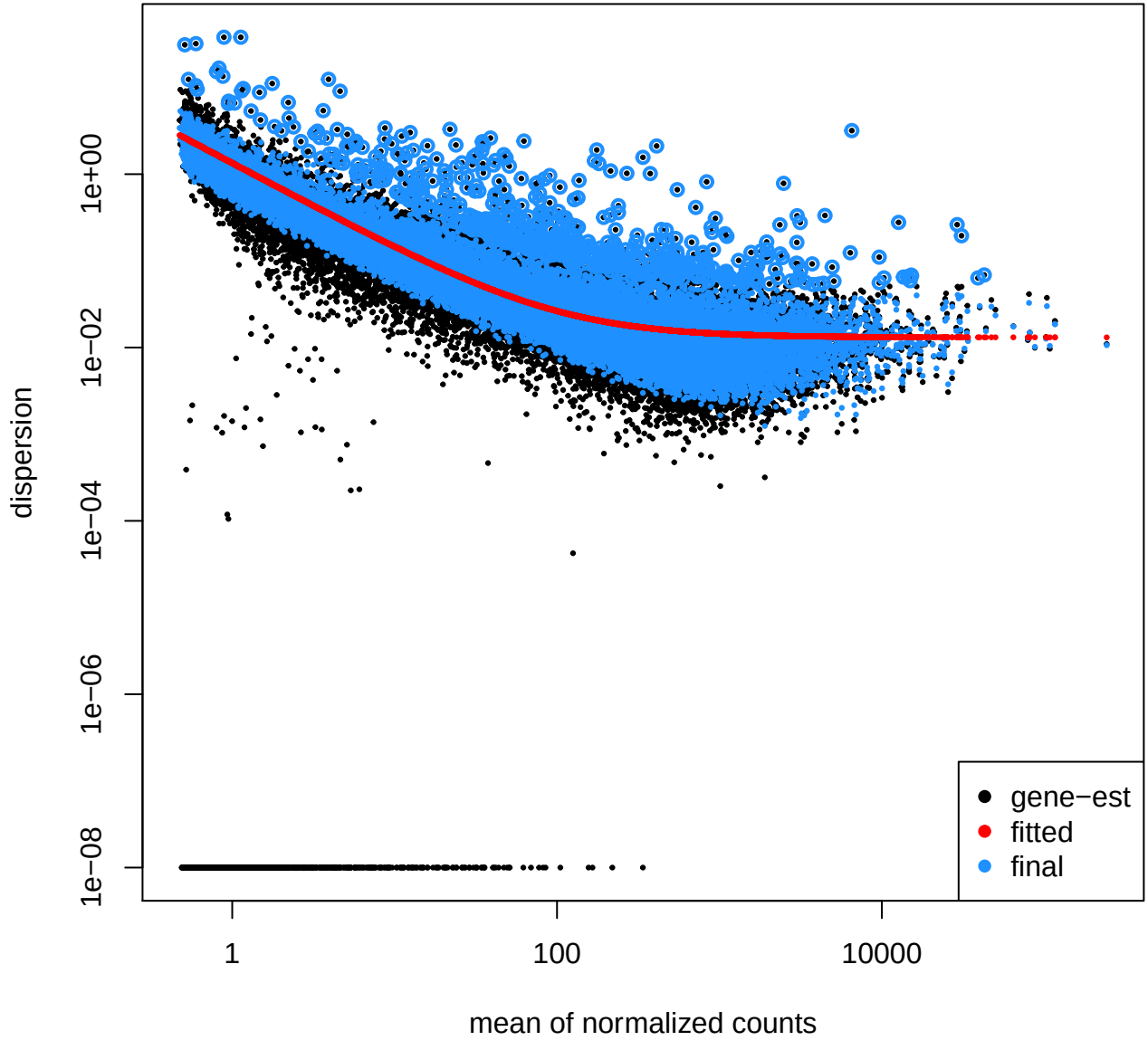
dispersion
1e+00
1e−02
1e−04
1e−06
1e−08
1
100
10000
mean of normalized counts
gene−est
fitted
final