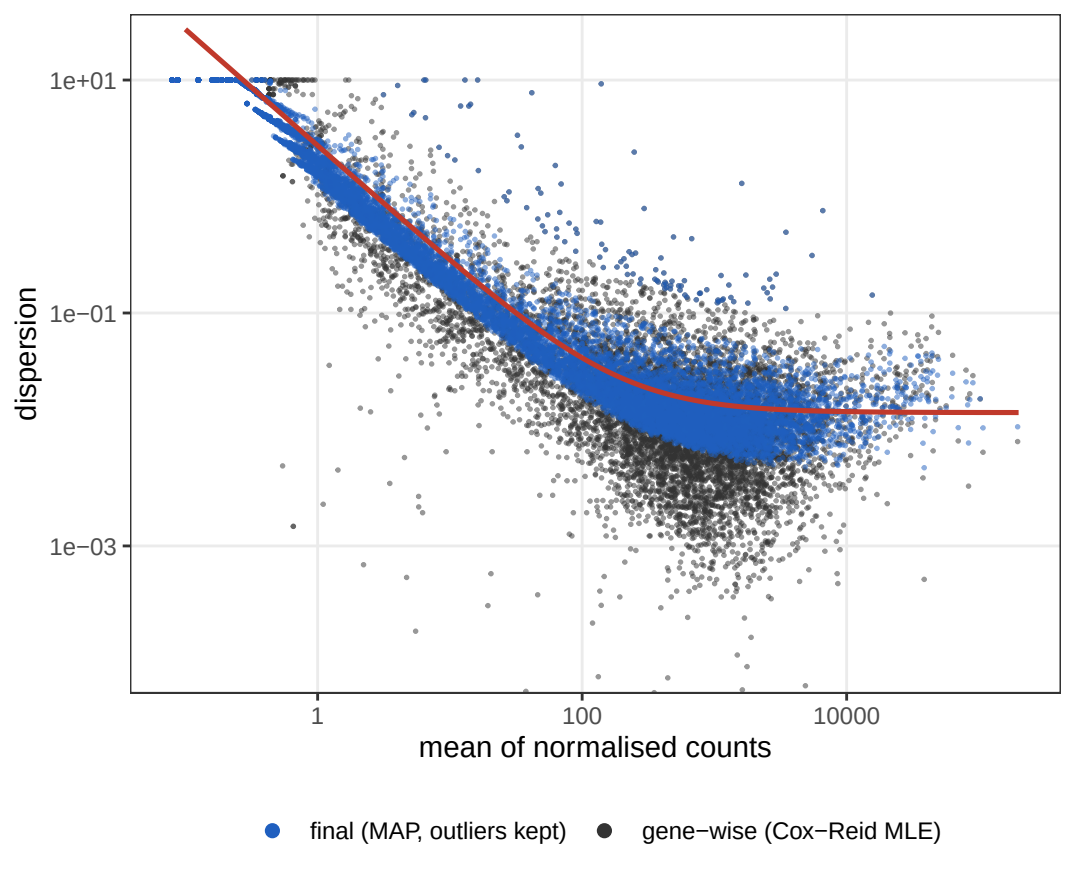

dispersion
1e+01
1e−01
1e−03
1
100
10000
mean of normalised counts
final (MAP, outliers kept)
gene−wise (Cox−Reid MLE)