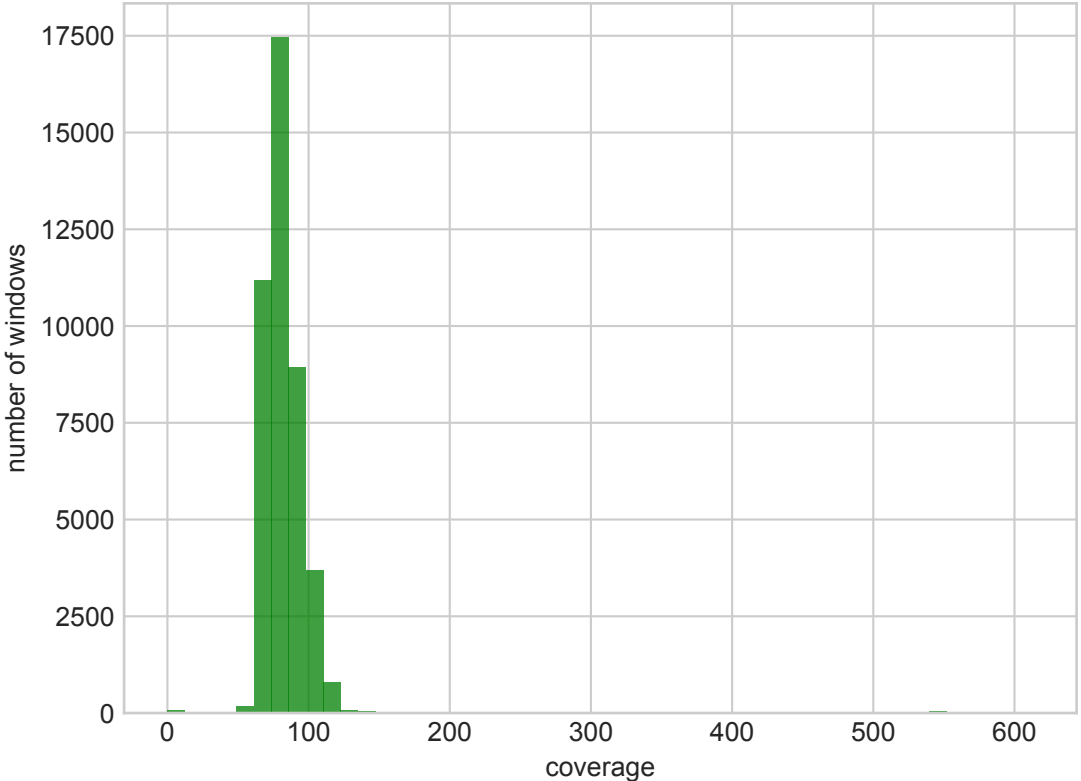
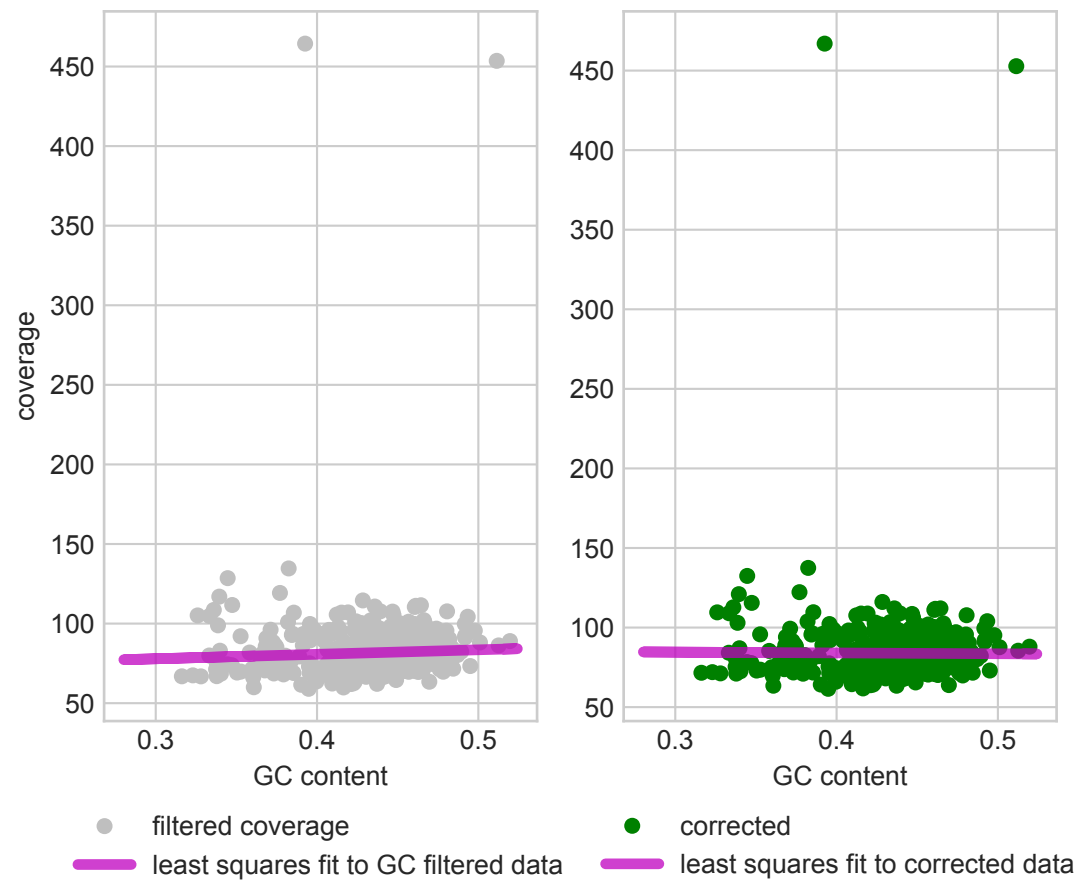
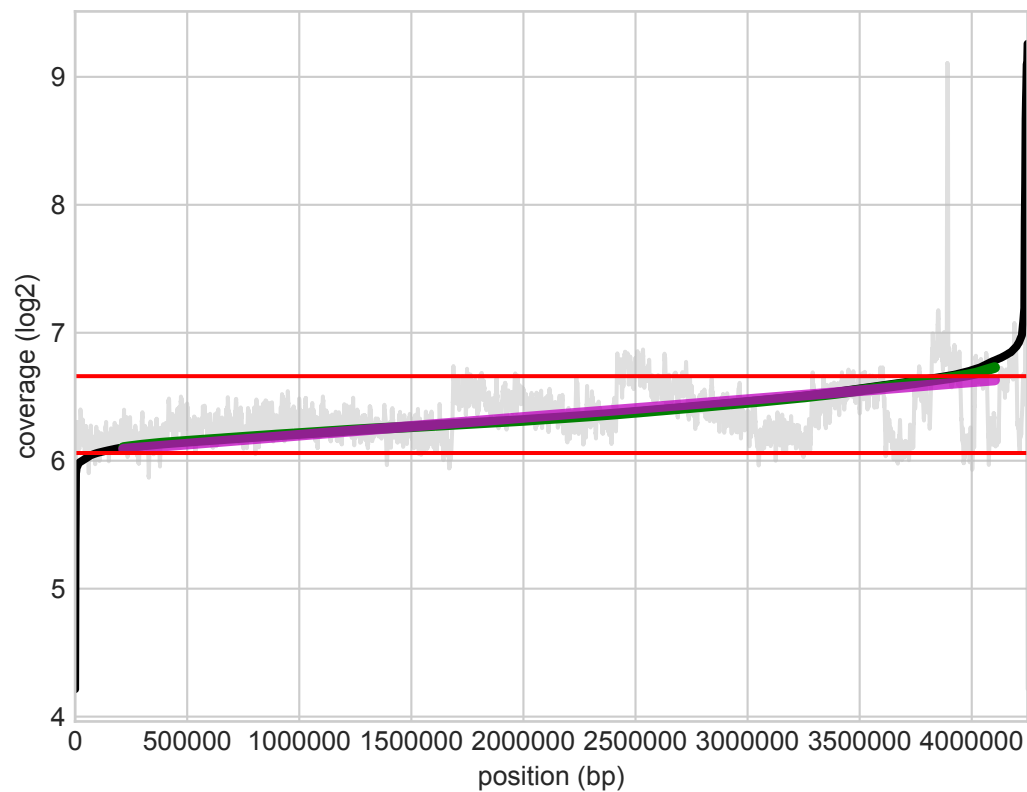


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0812/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0812-B
iRep: 1.52 r^2: 0.97 avg. cov: 82.87 % windows: 99.85



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0812/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0812-B
iRep: 1.52 raw iRep: 1.54 GC r^2: 0.01 GC bias: 0.22 avg. cov: 82.87 % windows: 99.85





- filtered
- sorted
- trimmed
- least squares fit to trimmed
- Ori/Ter