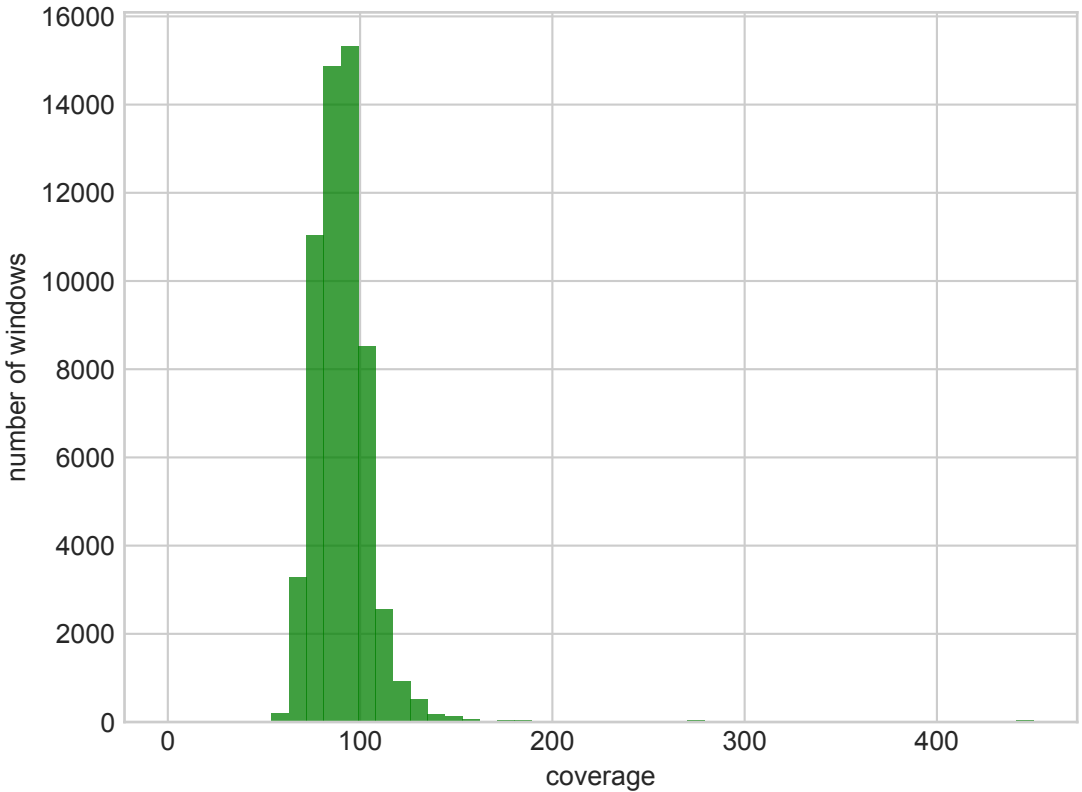
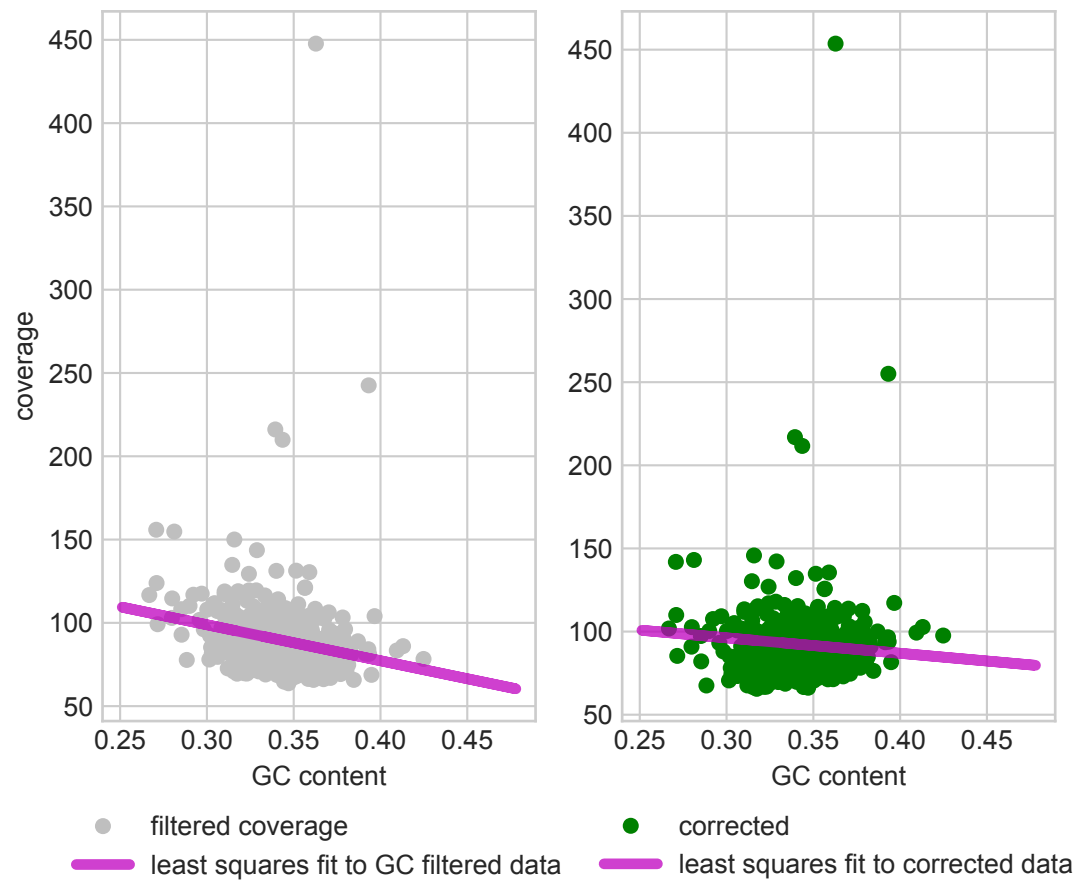
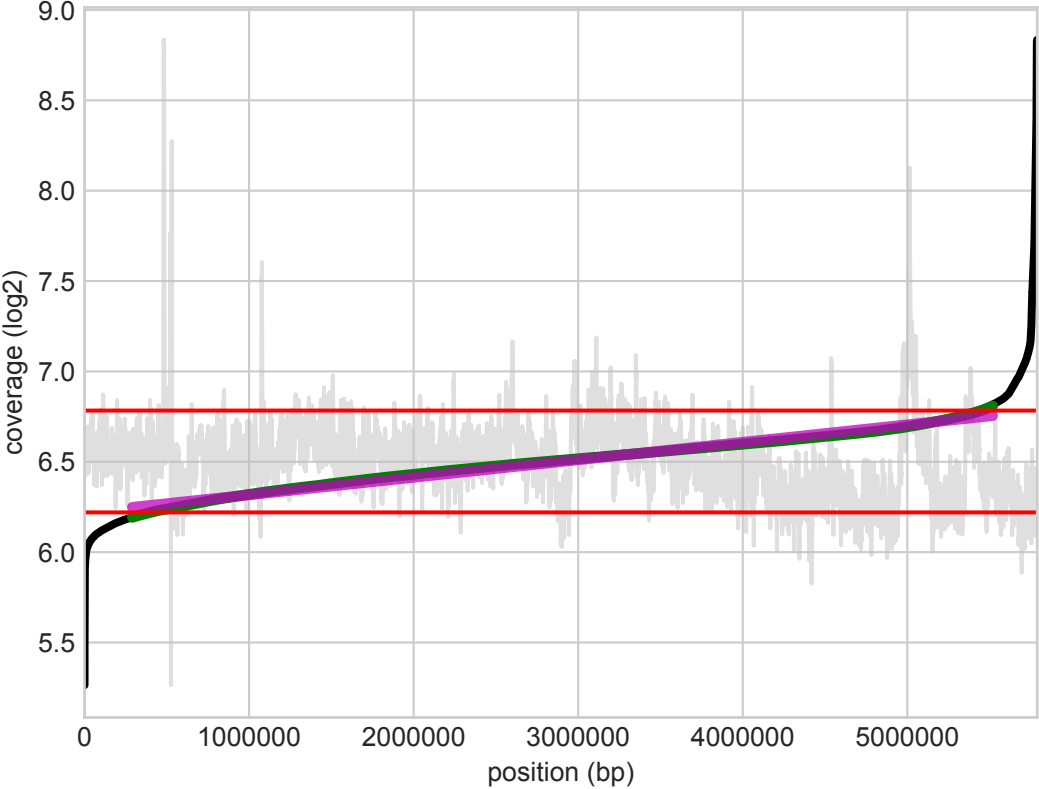


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0721/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0721-C
iRep: 1.48 r^2: 0.99 avg. cov: 91.19 % windows: 99.99



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0721/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0721-C
iRep: 1.48 raw iRep: 1.53 GC r^2: 0.13 GC bias: -27.37 avg. cov: 91.19 % windows: 99.99





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