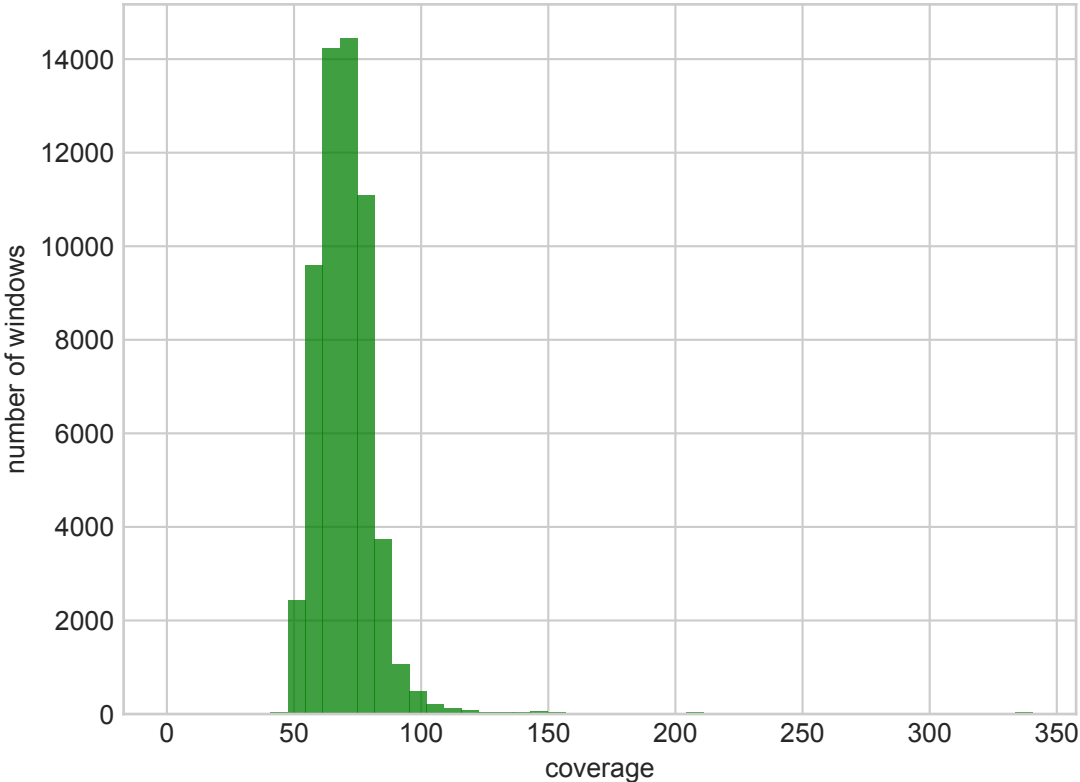
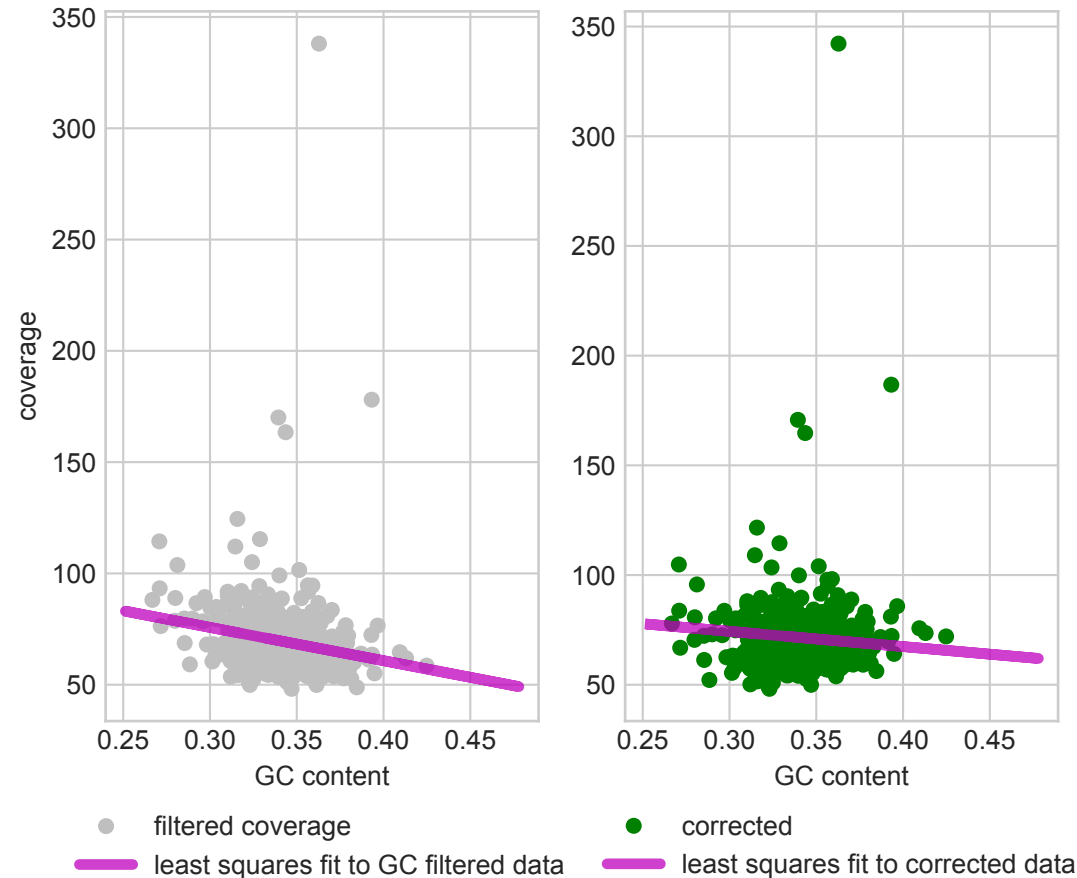
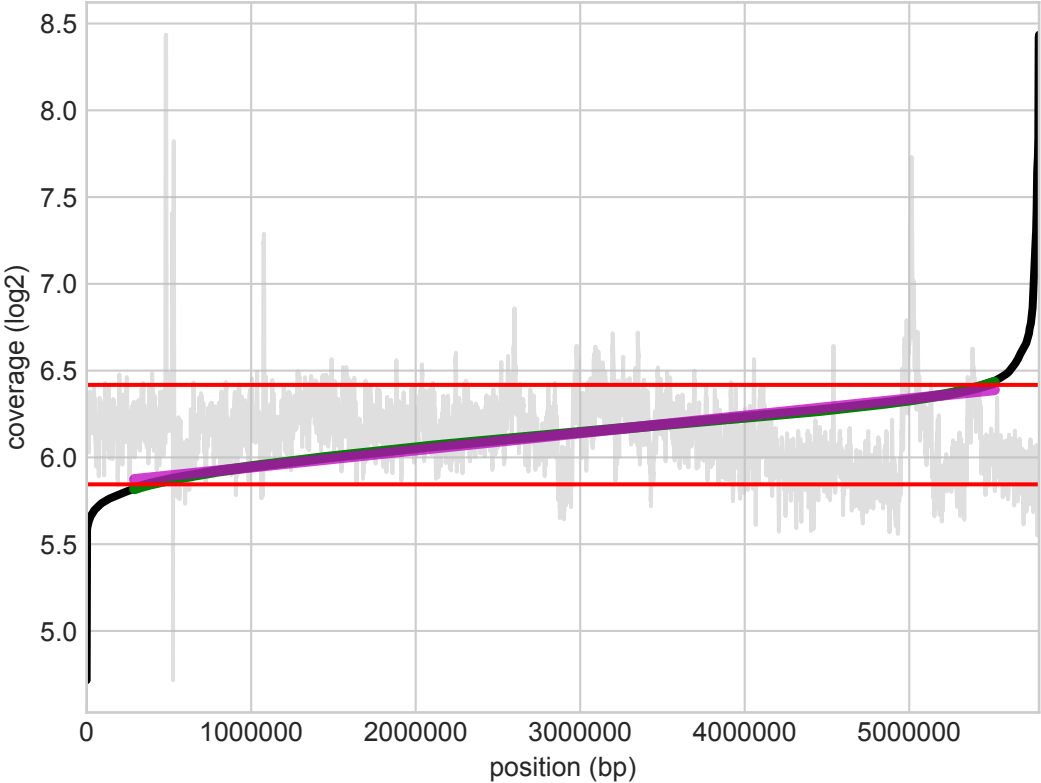


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0721/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0721-B  
iRep: 1.49 r^2: 0.99 avg. cov: 70.53 % windows: 99.99



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0721/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0721-B  
iRep: 1.49 raw iRep: 1.53 GC r^2: 0.10 GC bias: -15.43 avg. cov: 70.53 % windows: 99.99





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