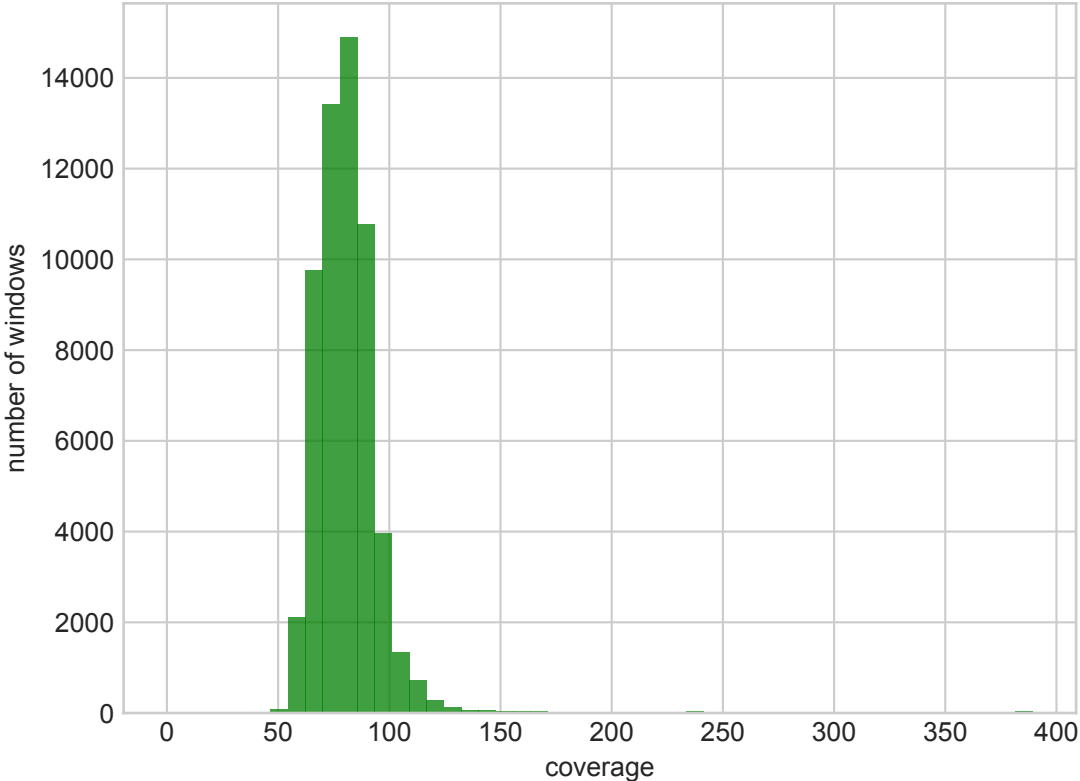
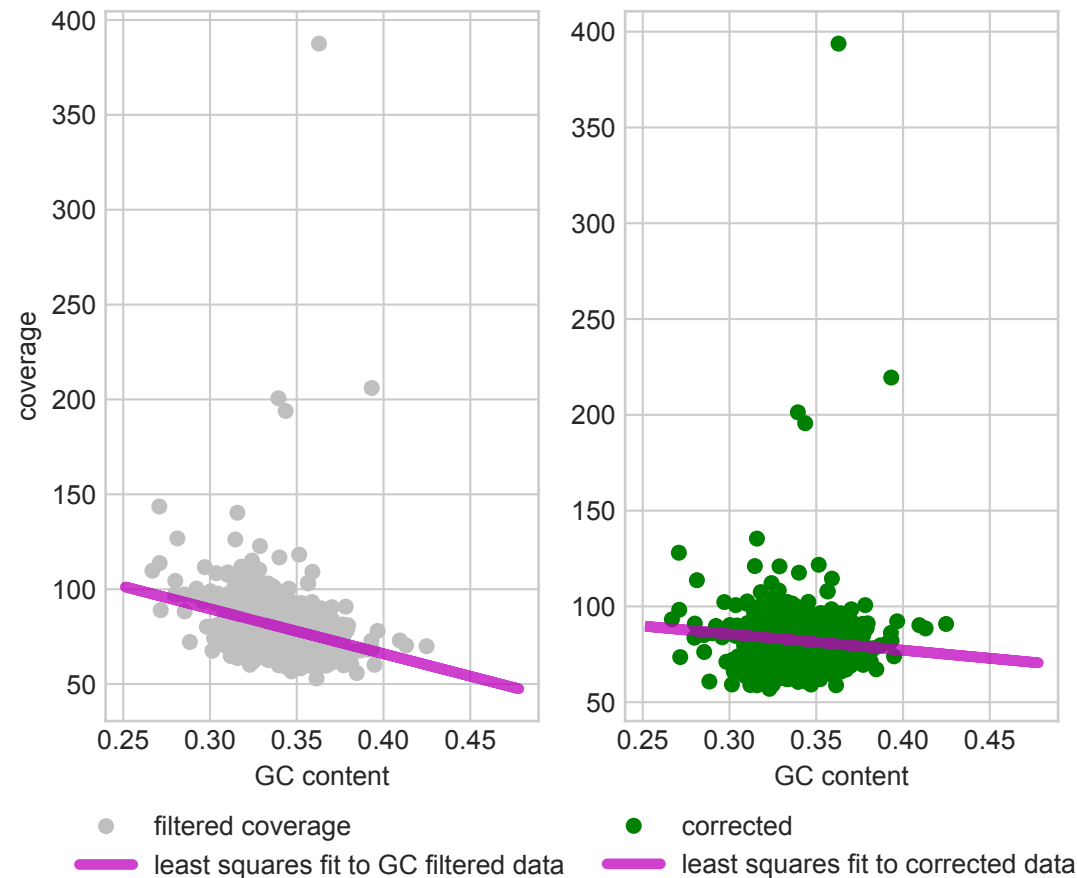
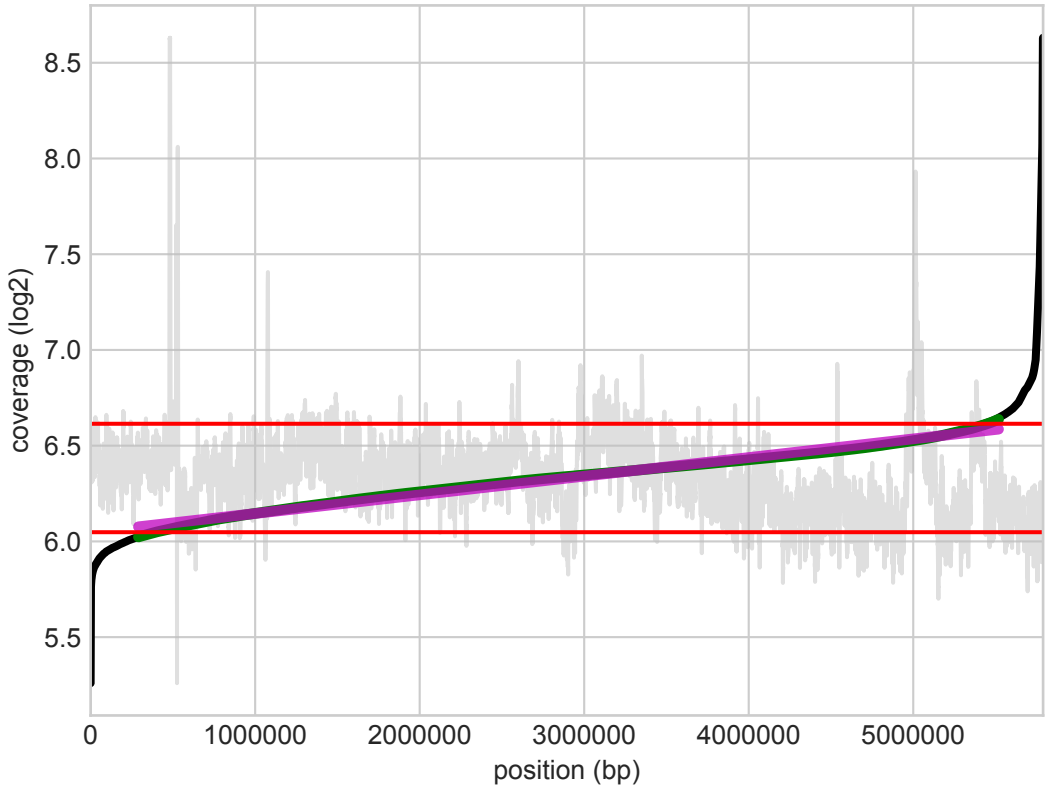


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0721/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0721-A  
iRep: 1.48   r^2: 0.99   avg. cov: 81.00   % windows: 99.99



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0721/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0721-A  
iRep: 1.48 raw iRep: 1.55 GC r^2: 0.18 GC bias: -43.37 avg. cov: 81.00 % windows: 99.99





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