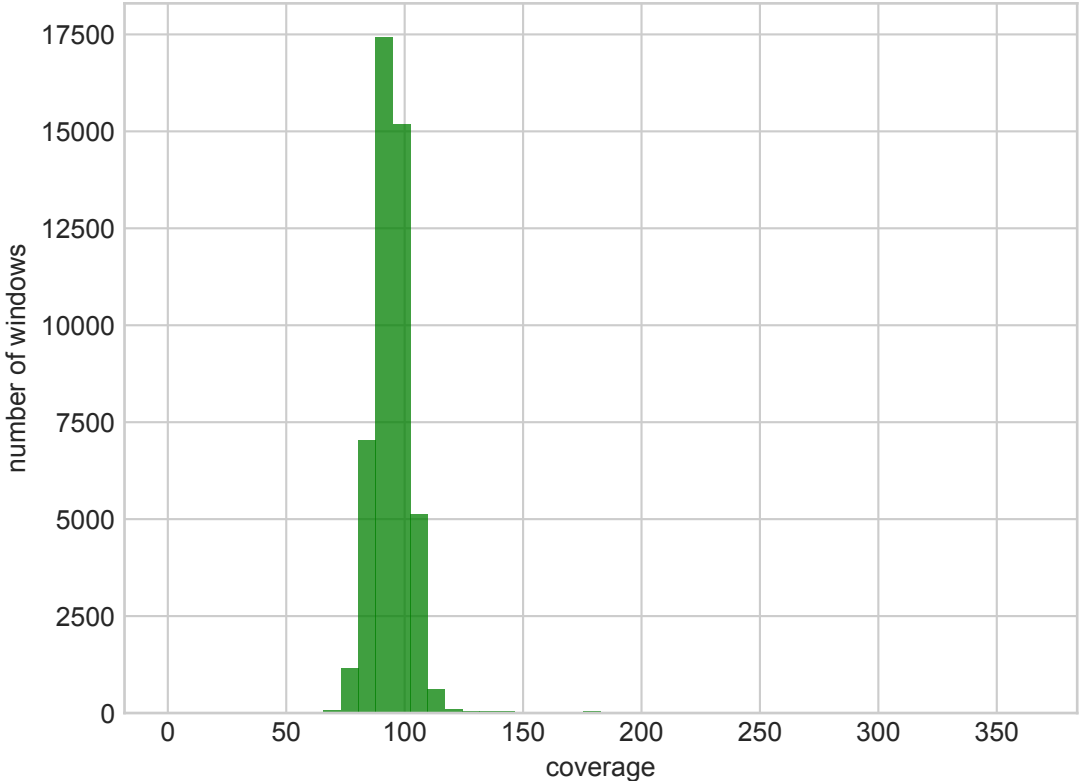
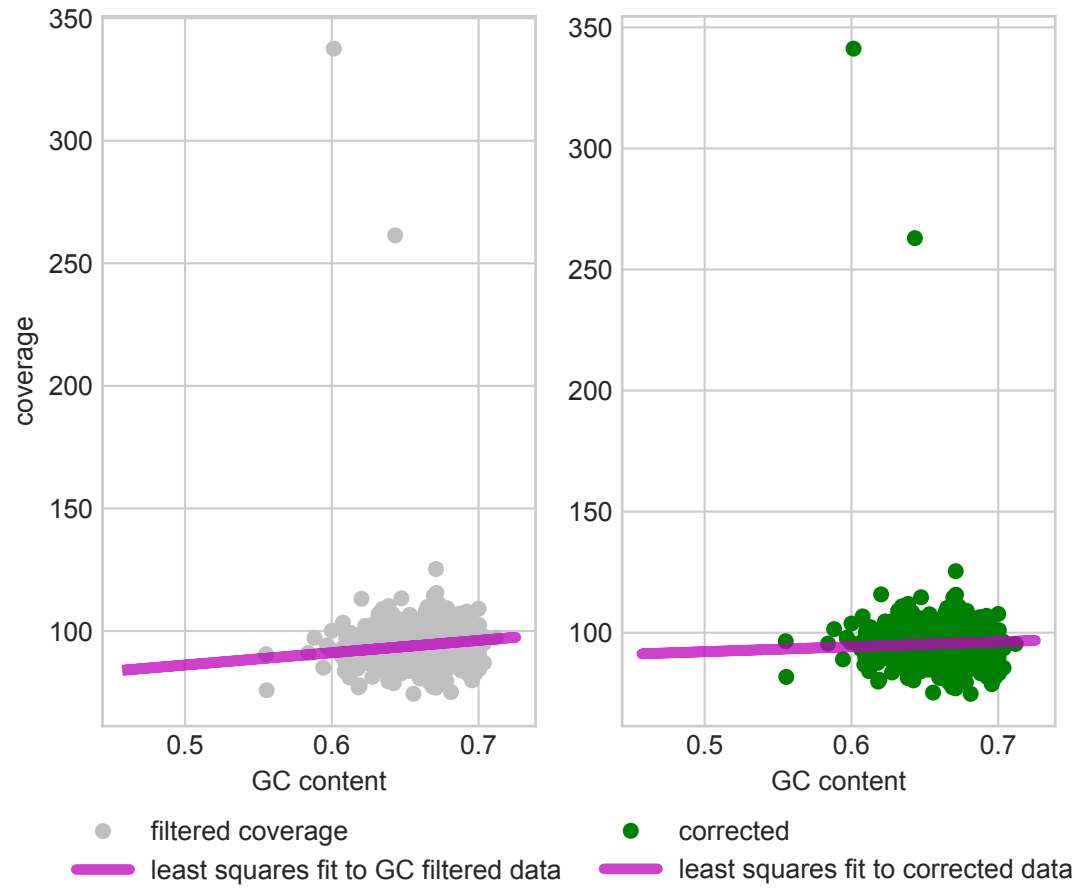
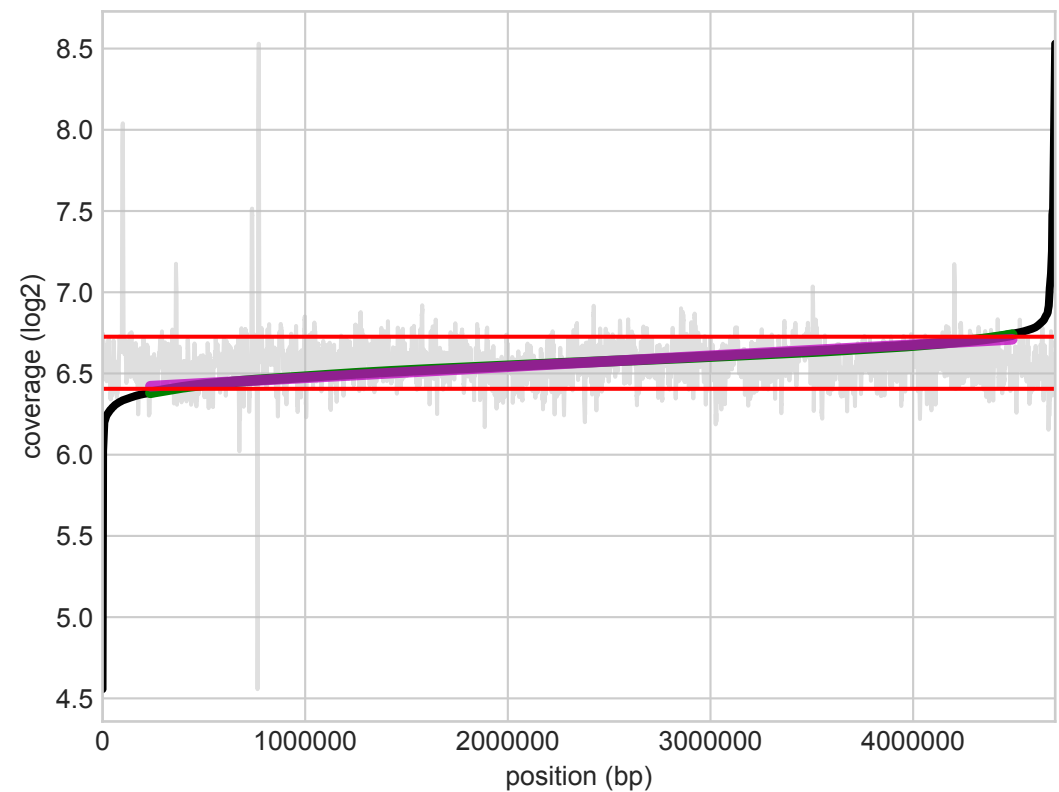


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0703/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0703-D  
iRep: 1.25   r^2: 0.99   avg. cov: 94.81   % windows: 99.98



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0703/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0703-D  
iRep: 1.25 raw iRep: 1.26 GC r^2: 0.03 GC bias: 1.44 avg. cov: 94.81 % windows: 99.98





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