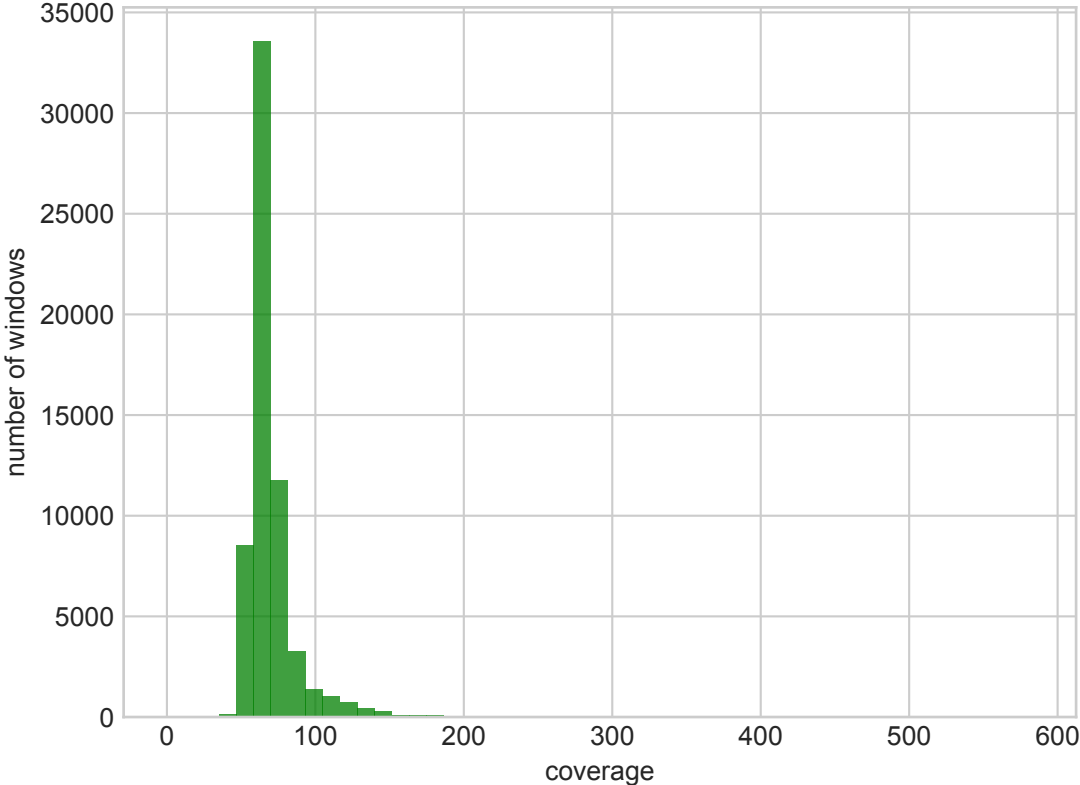
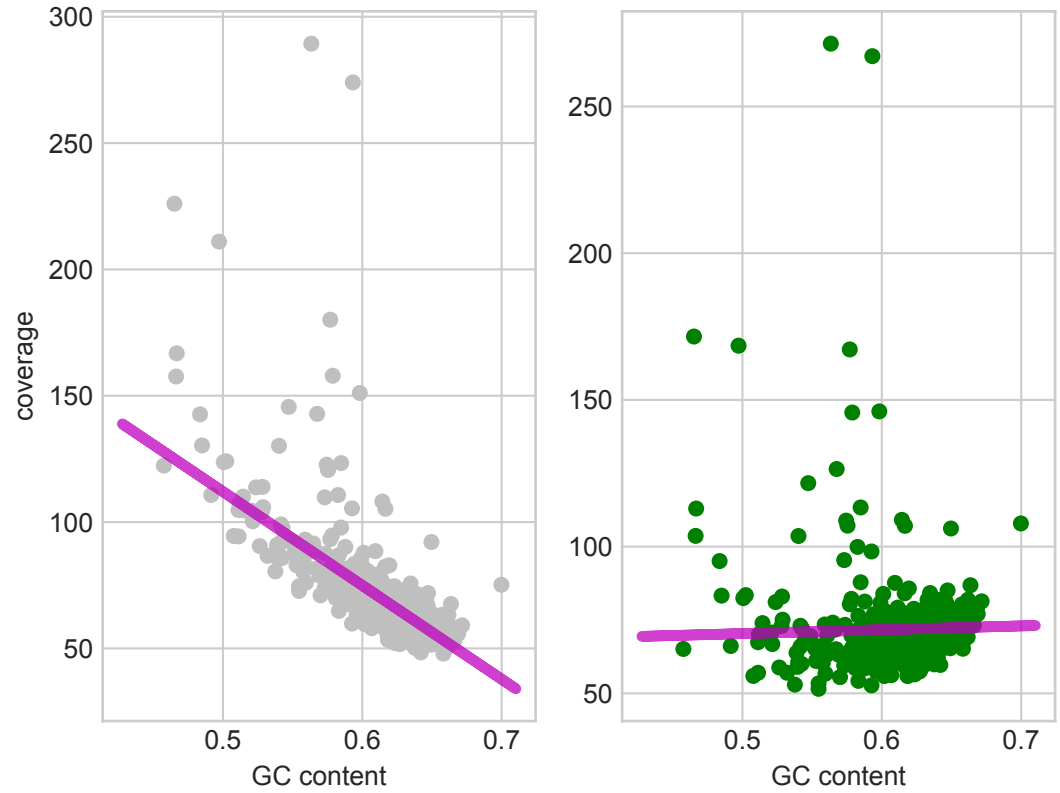


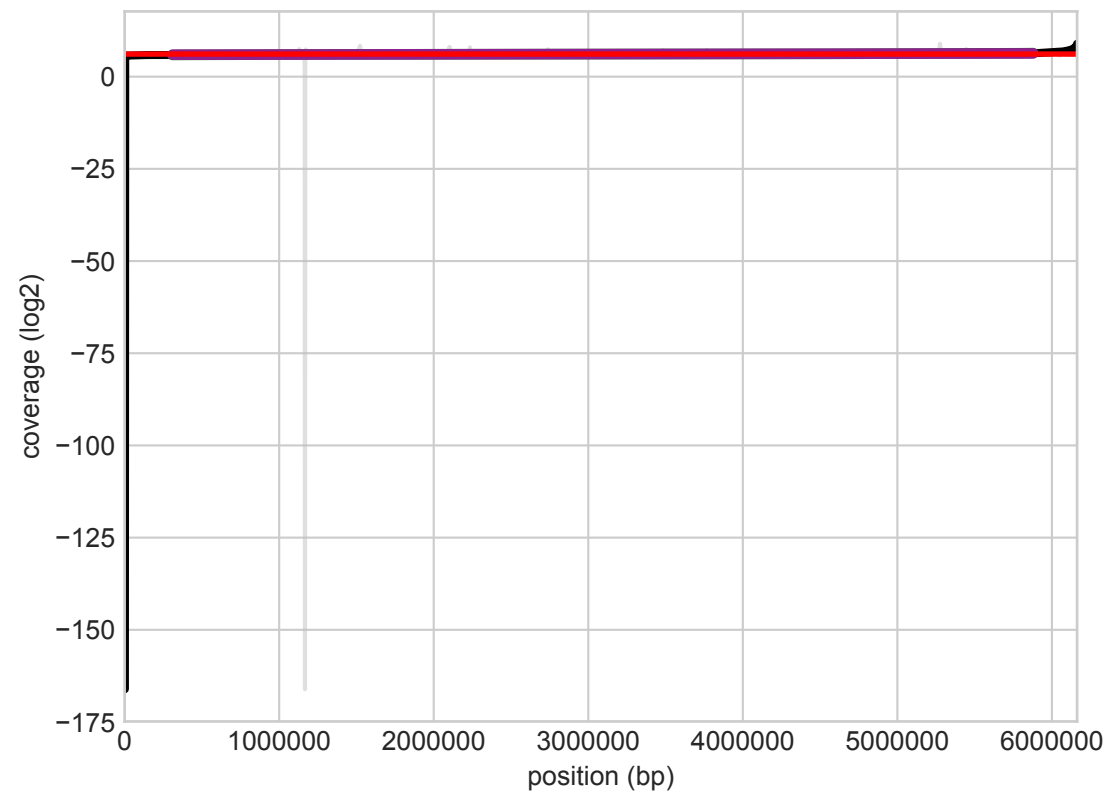
genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0802/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0802-B  
iRep: 1.30 r^2: 0.98 avg. cov: 70.70 % windows: 99.96



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0802/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0802-B  
iRep: 1.30 raw iRep: 1.57 GC r^2: 0.67 GC bias: -249.68 avg. cov: 70.70 % windows: 99.96



● filtered coverage      ● corrected  
— least squares fit to GC filtered data      — least squares fit to corrected data



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