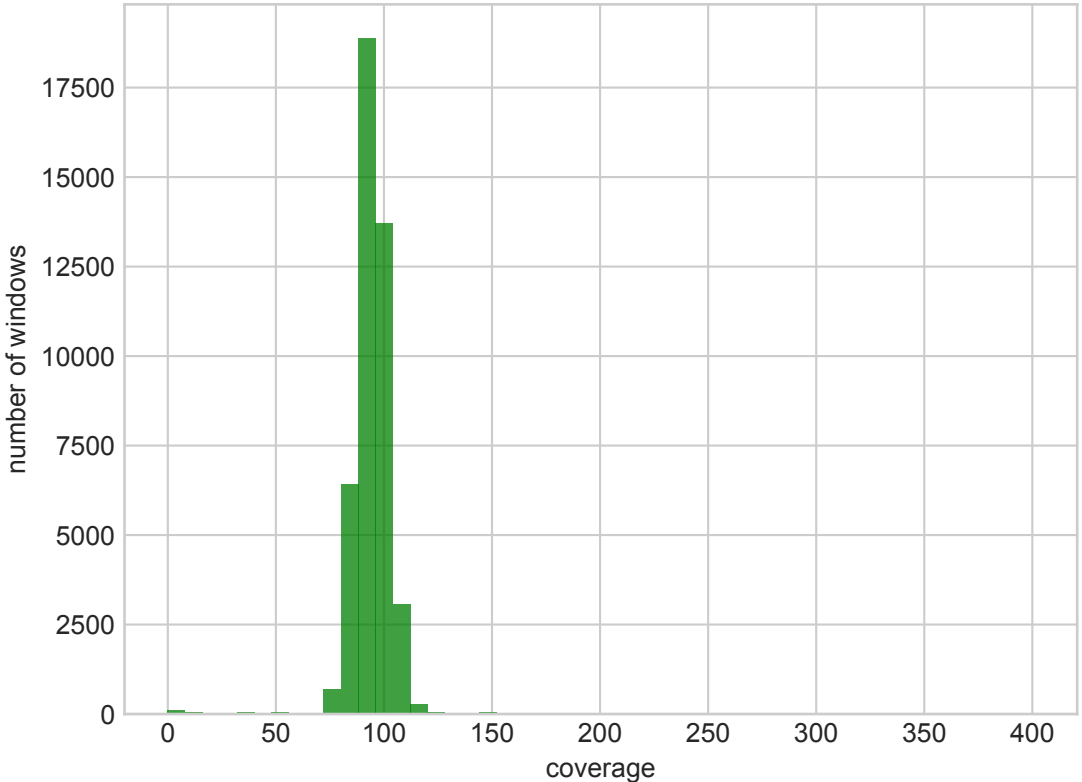
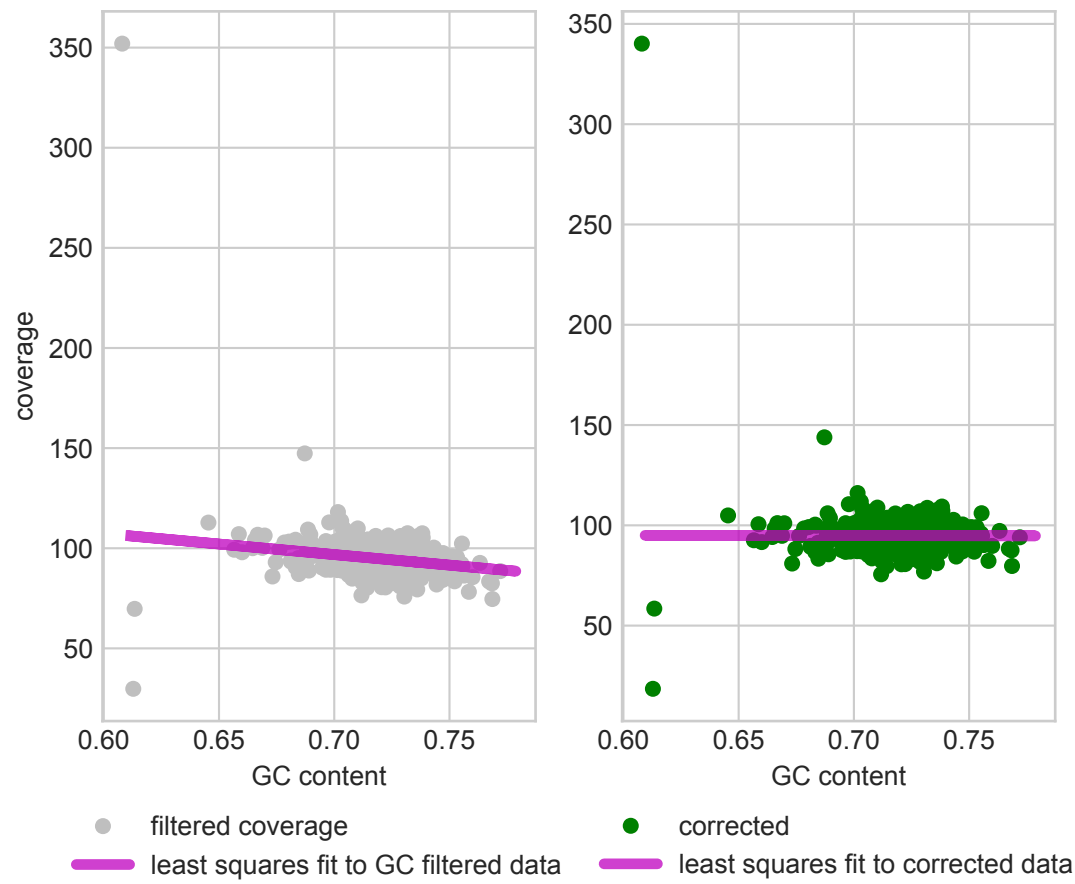
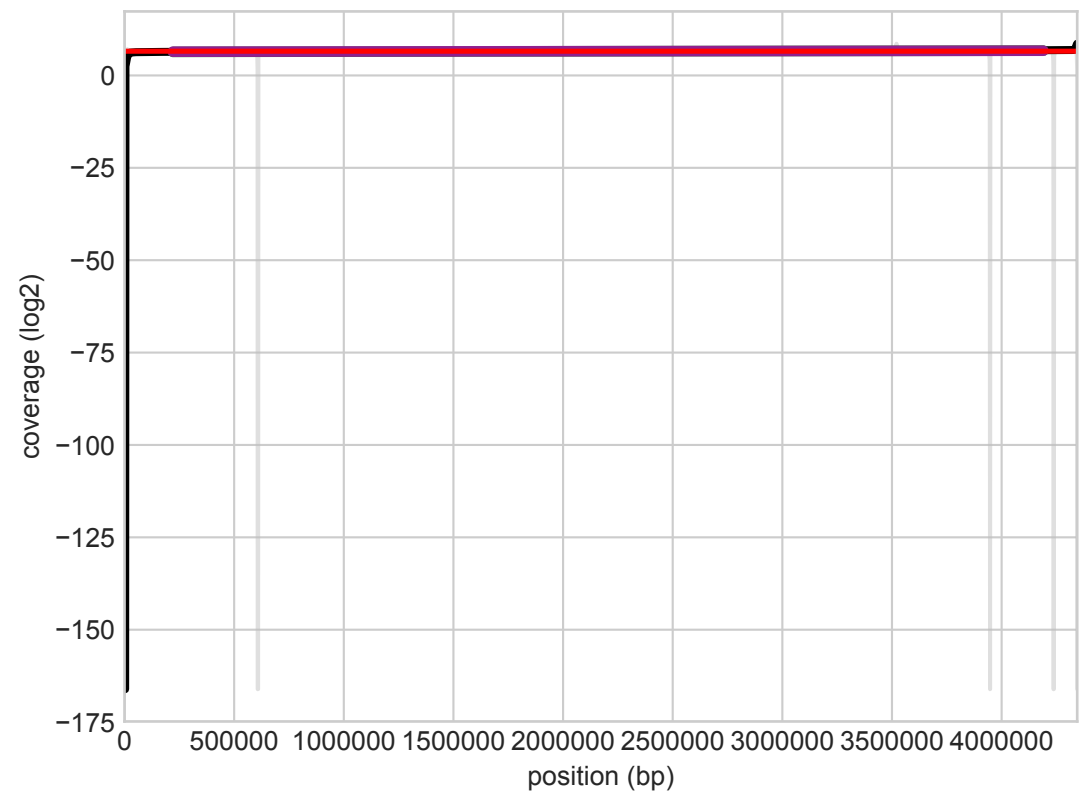


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0722/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0722-D
iRep: 1.23 r^2: 0.99 avg. cov: 94.42 % windows: 99.73



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0722/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0722-D
iRep: 1.23 raw iRep: 1.24 GC r²: 0.11 GC bias: -11.15 avg. cov: 94.42 % windows: 99.73





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