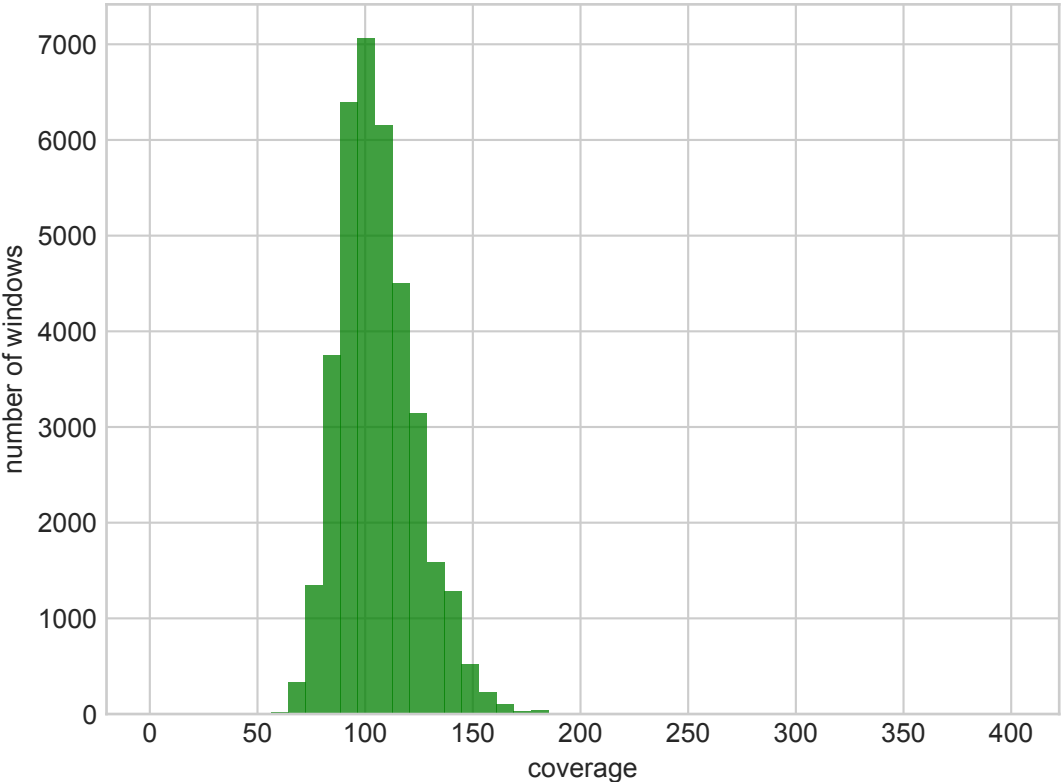
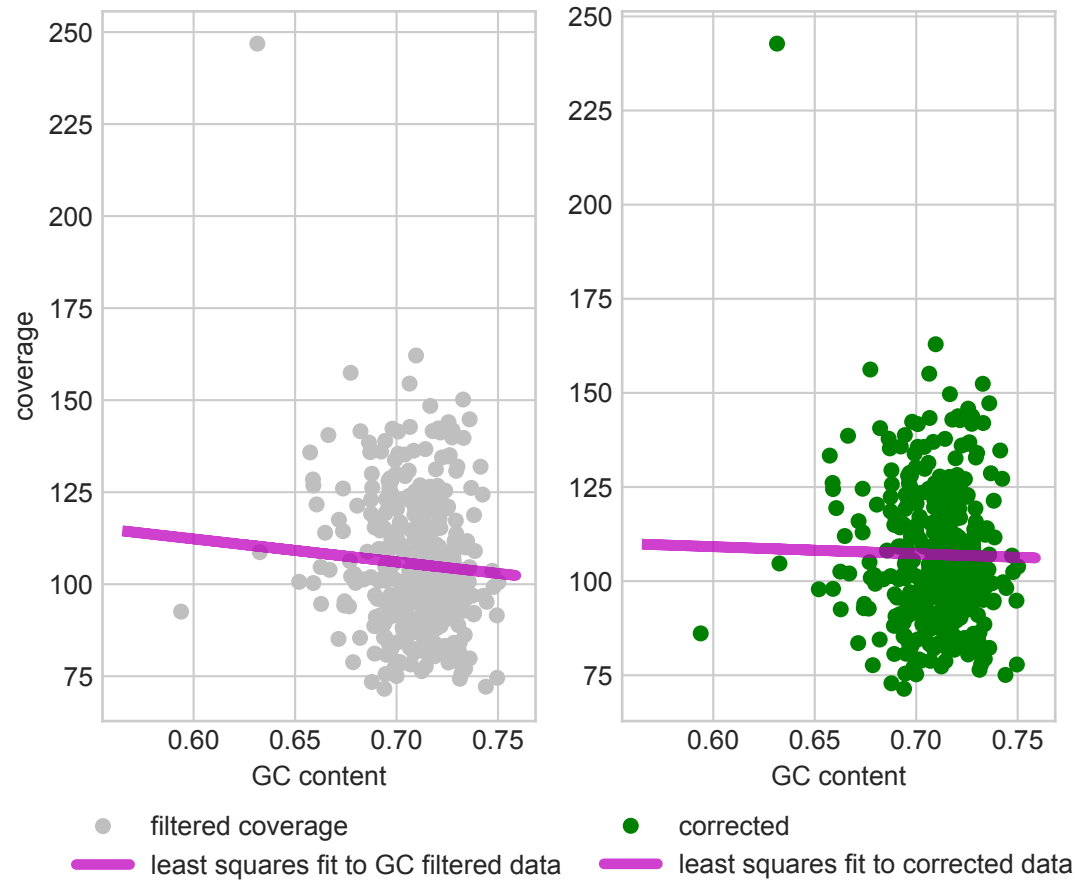
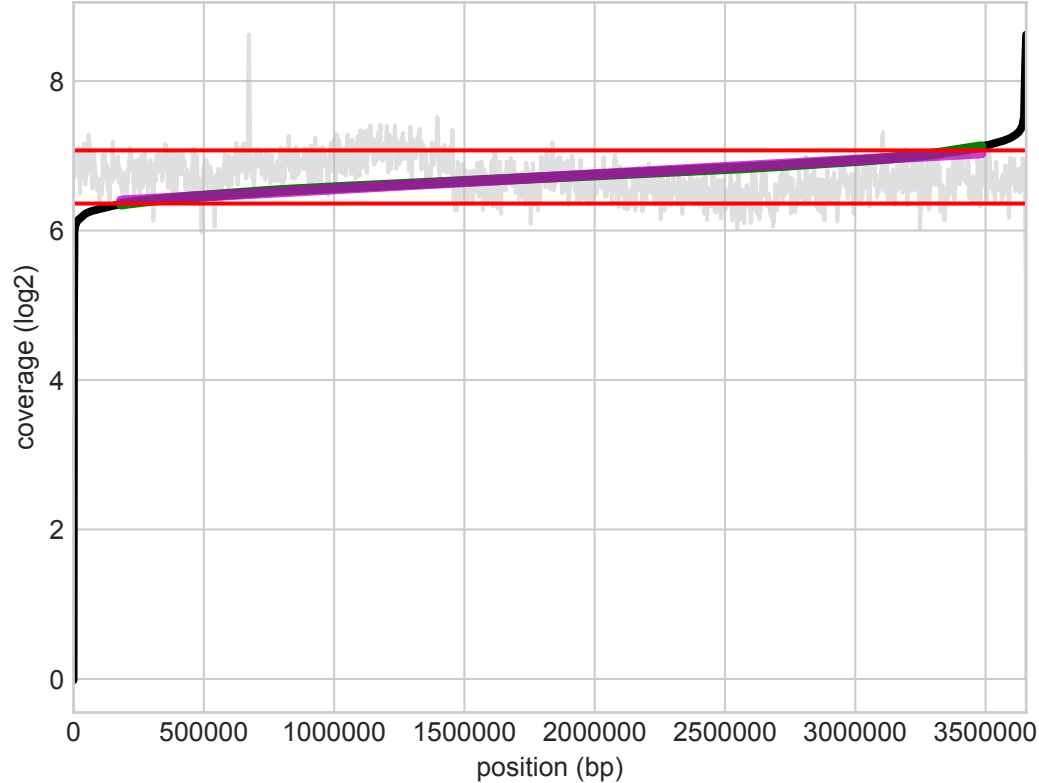


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0715/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0715-C
iRep: 1.64 r^2: 0.99 avg. cov: 106.13 % windows: 99.97



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0715/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0715-C
iRep: 1.64 raw iRep: 1.64 GC r^2: 0.01 GC bias: -0.34 avg. cov: 106.13 % windows: 99.97





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