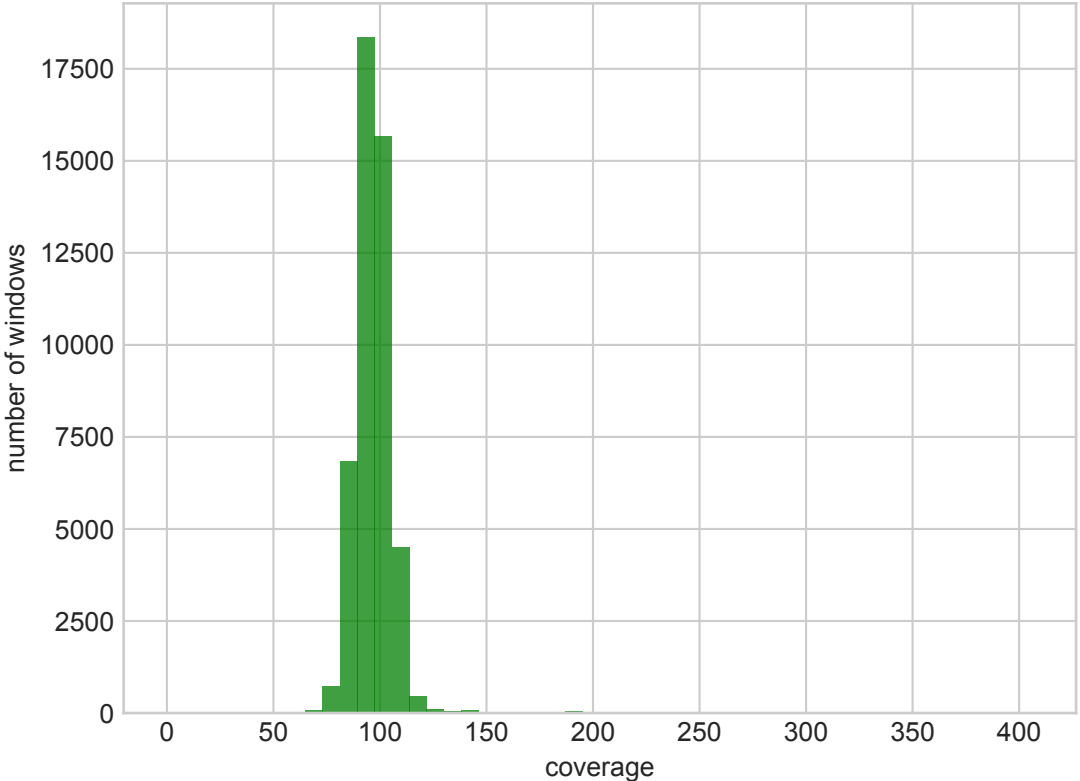
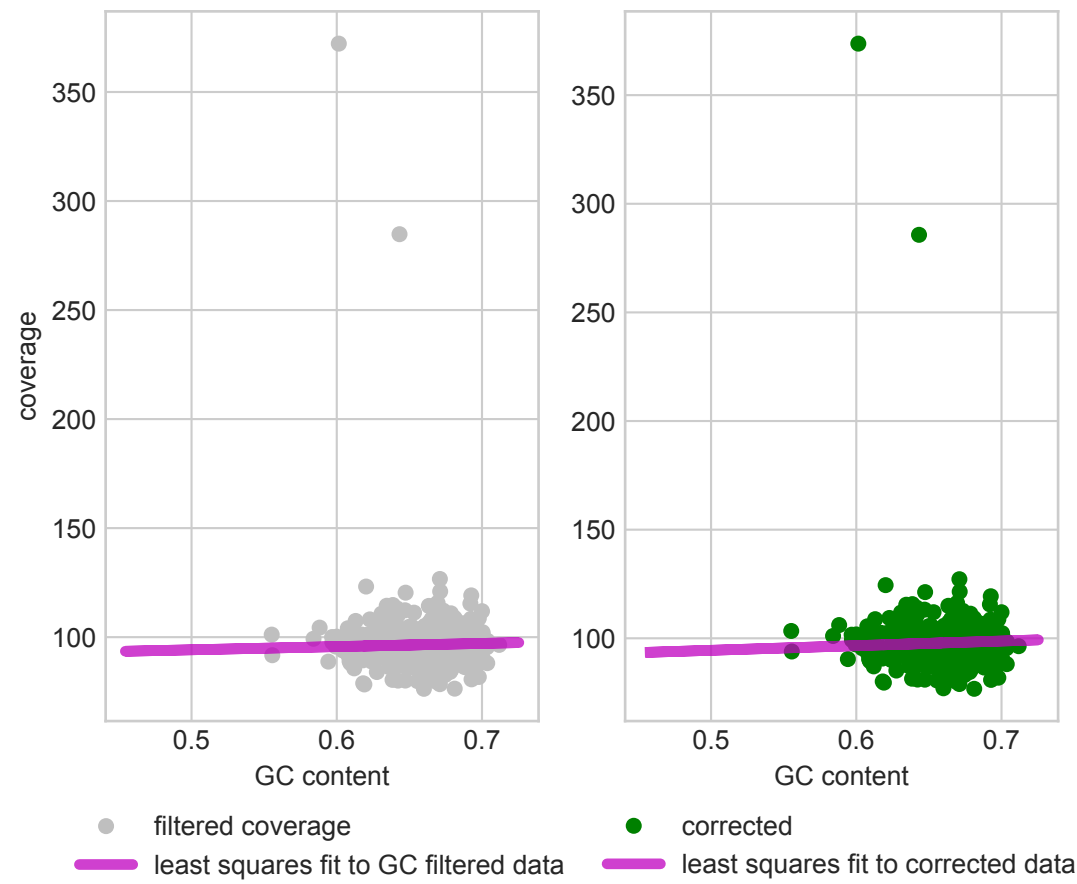


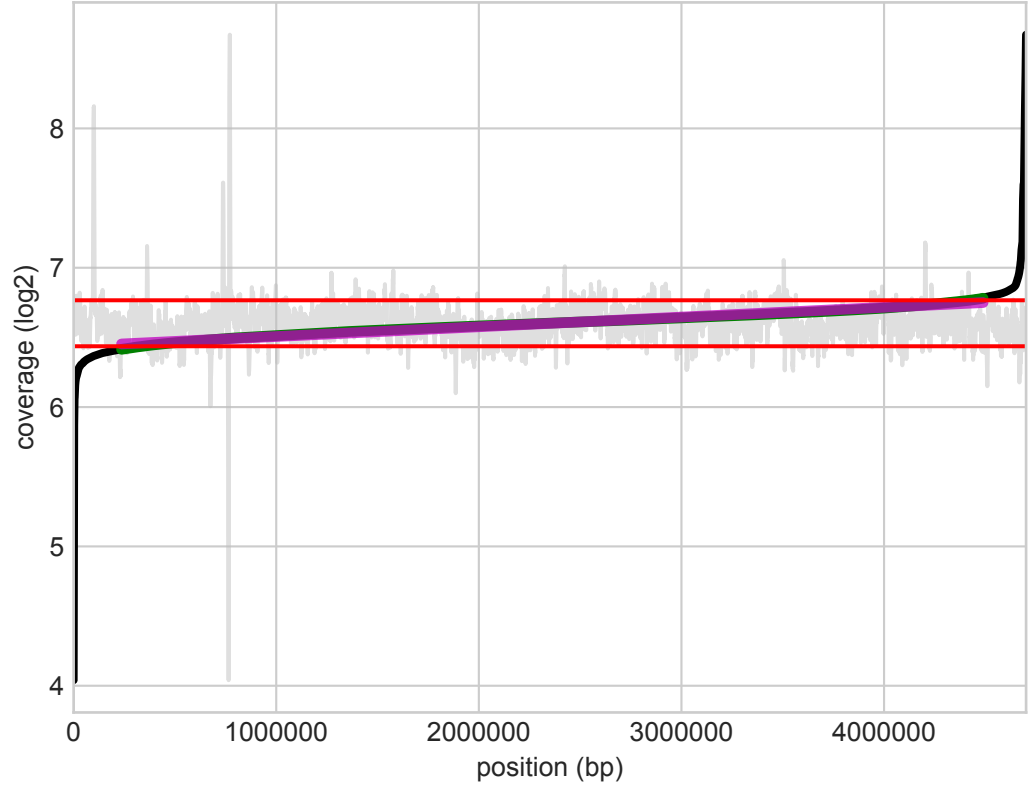
genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0703/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0703-B
iRep: 1.26 r^2: 0.99 avg. cov: 97.19 % windows: 99.98



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0703/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0703-B
iRep: 1.26 raw iRep: 1.26 GC r^2: 0.00 GC bias: 0.03 avg. cov: 97.19 % 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-B
iRep: 1.26 un-filtered iRep: 1.26 r²: 0.99 avg. cov: 97.19 % windows: 99.98



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