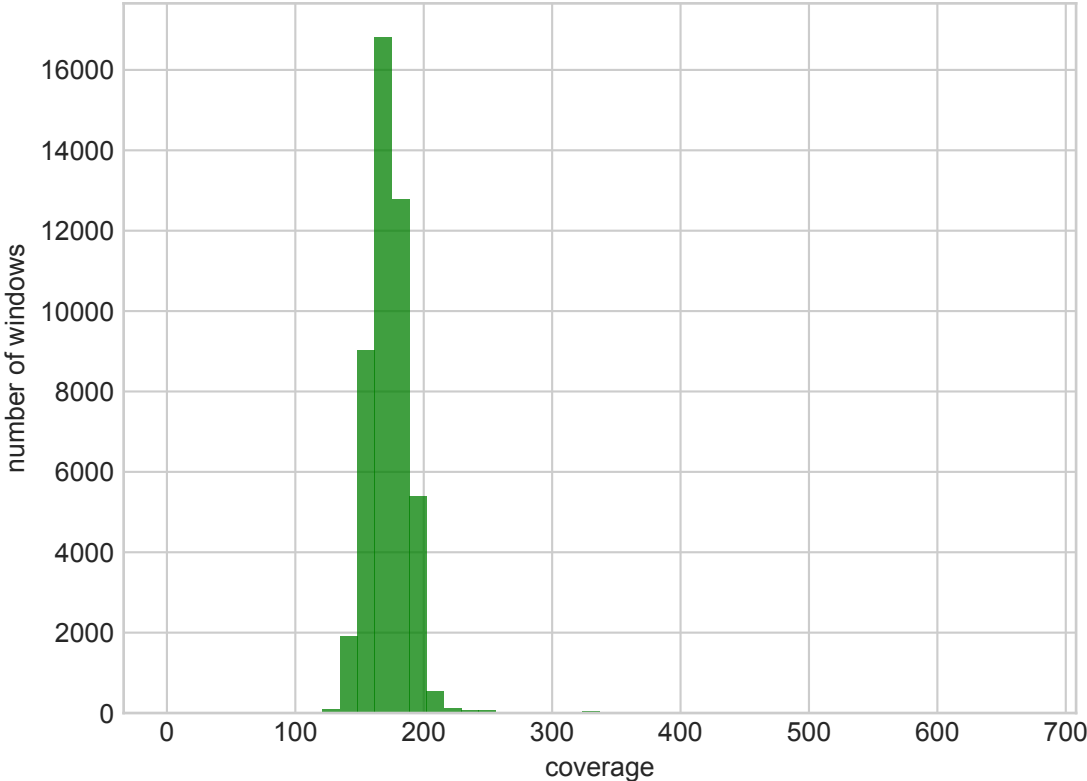
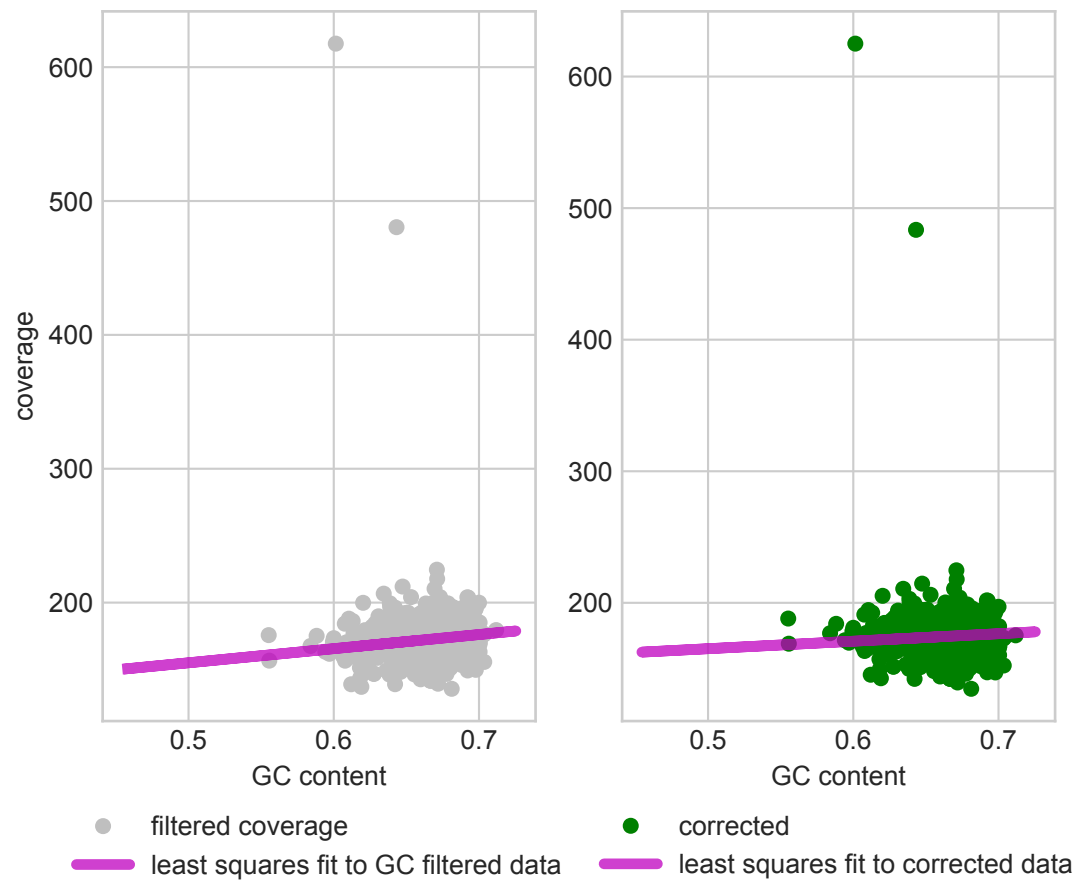
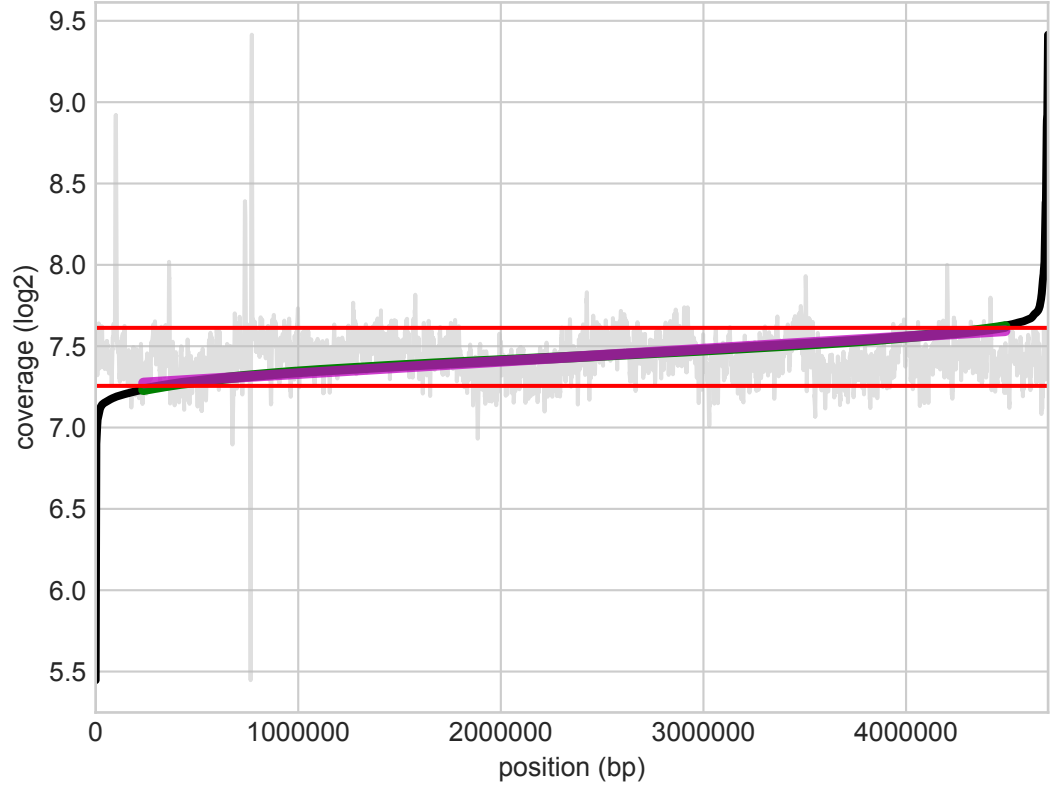


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0703/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0703-A
iRep: 1.28 r^2: 0.99 avg. cov: 173.15 % windows: 99.98





genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0703/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0703-A
iRep: 1.28 un-filtered iRep: 1.28 r^2: 0.99 avg. cov: 173.15 % windows: 99.98



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