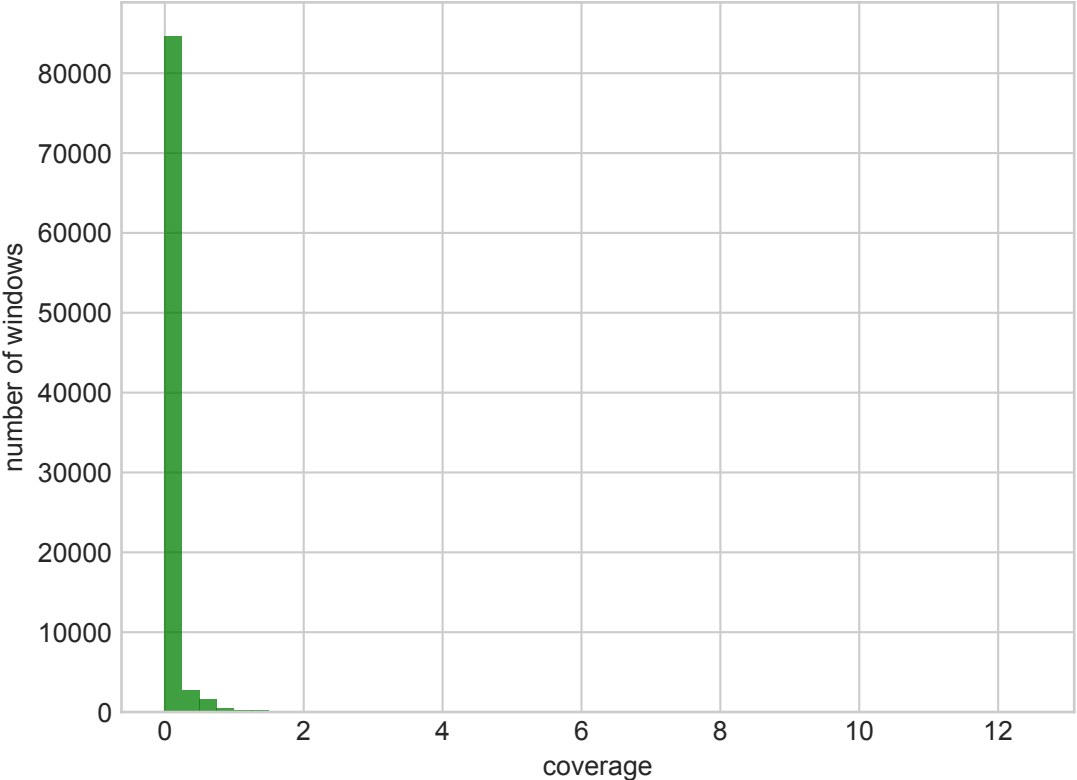
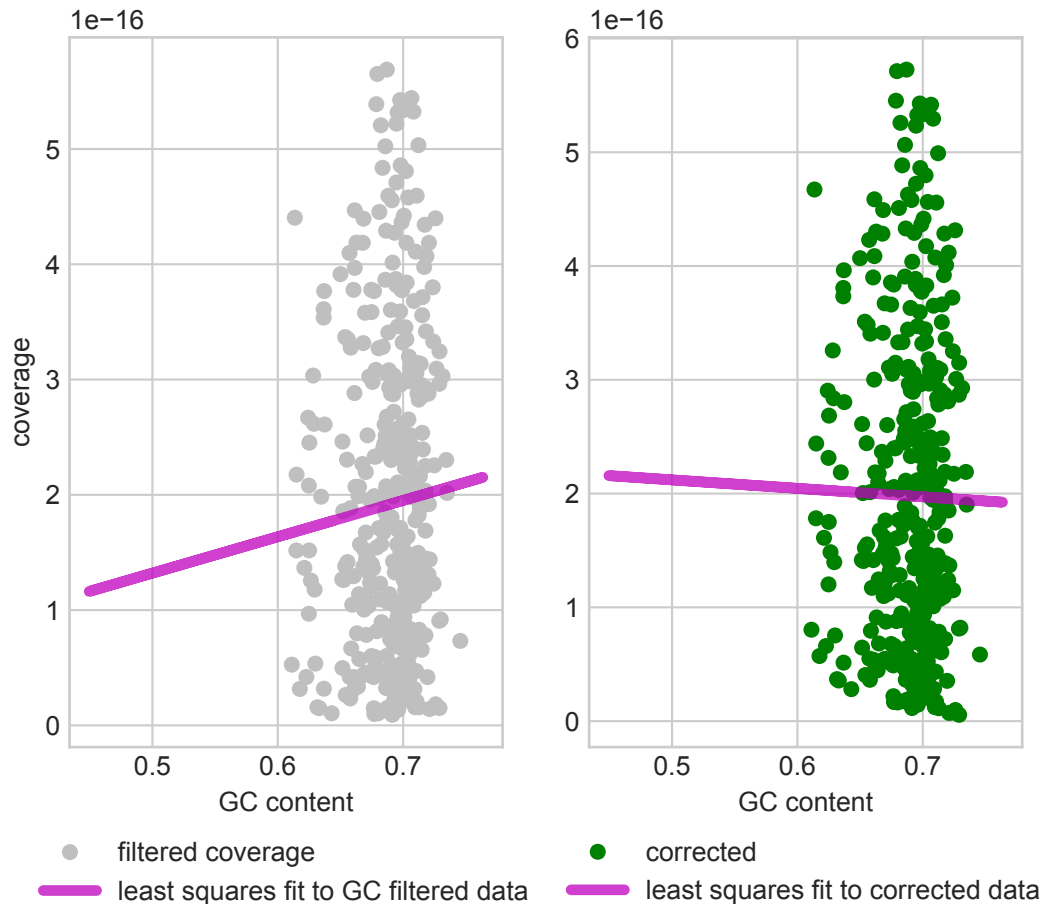


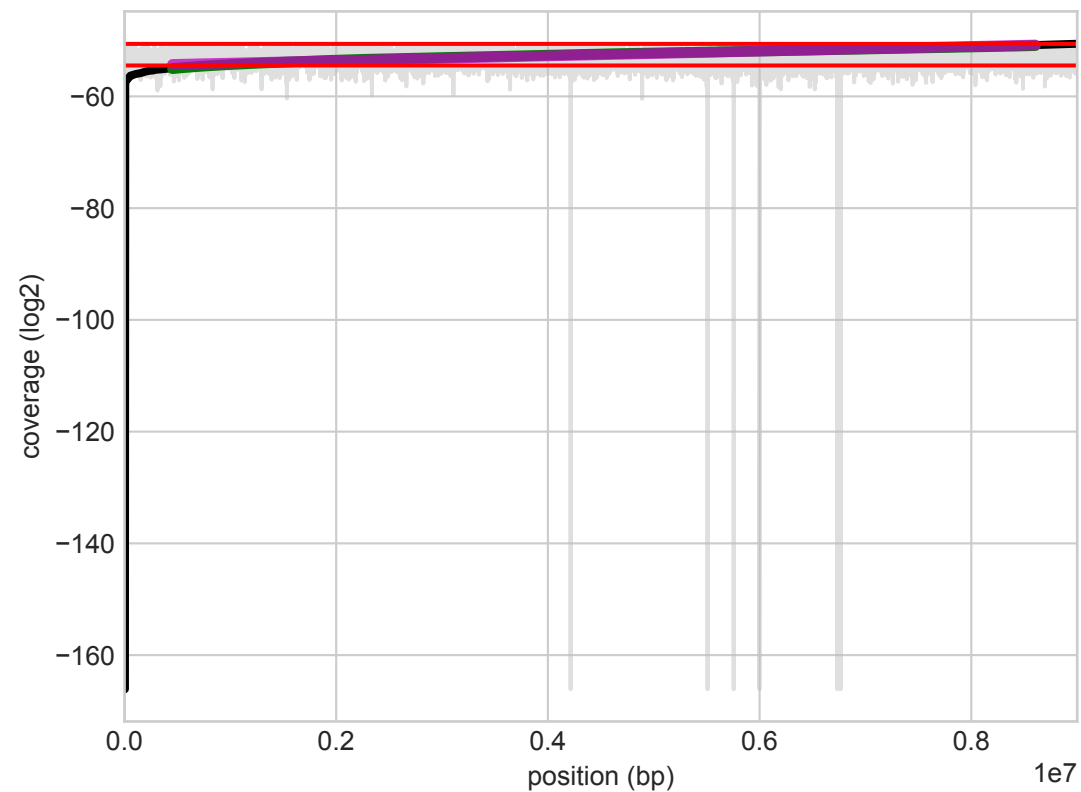
genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0706/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0706-D
iRep: n/a r^2: 0.97 avg. cov: 0.05 % windows: 40.43



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0706/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0706-D
iRep: n/a raw iRep: 16.14 GC r^2: 0.00 GC bias: 0.00 avg. cov: 0.05 % windows: 40.43



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0706/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0706-D
iRep: n/a un-filtered iRep: 14.57 r^2: 0.97 avg. cov: 0.05 % windows: 40.43



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