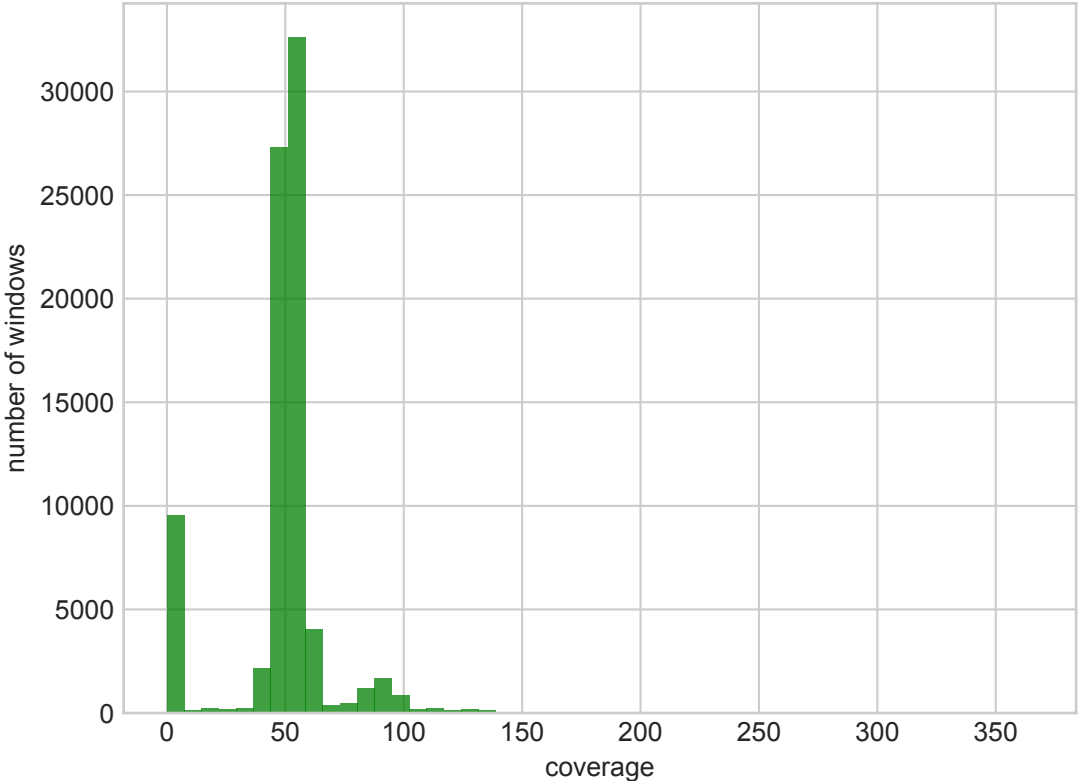
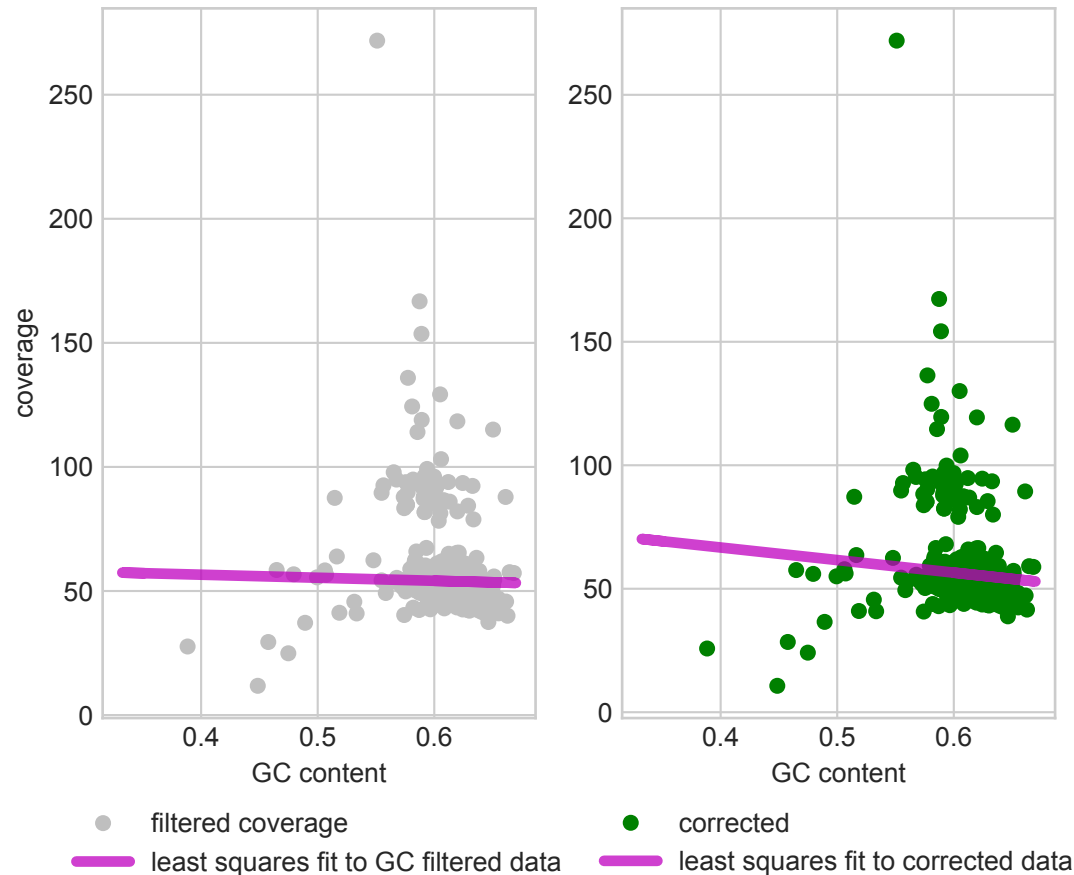
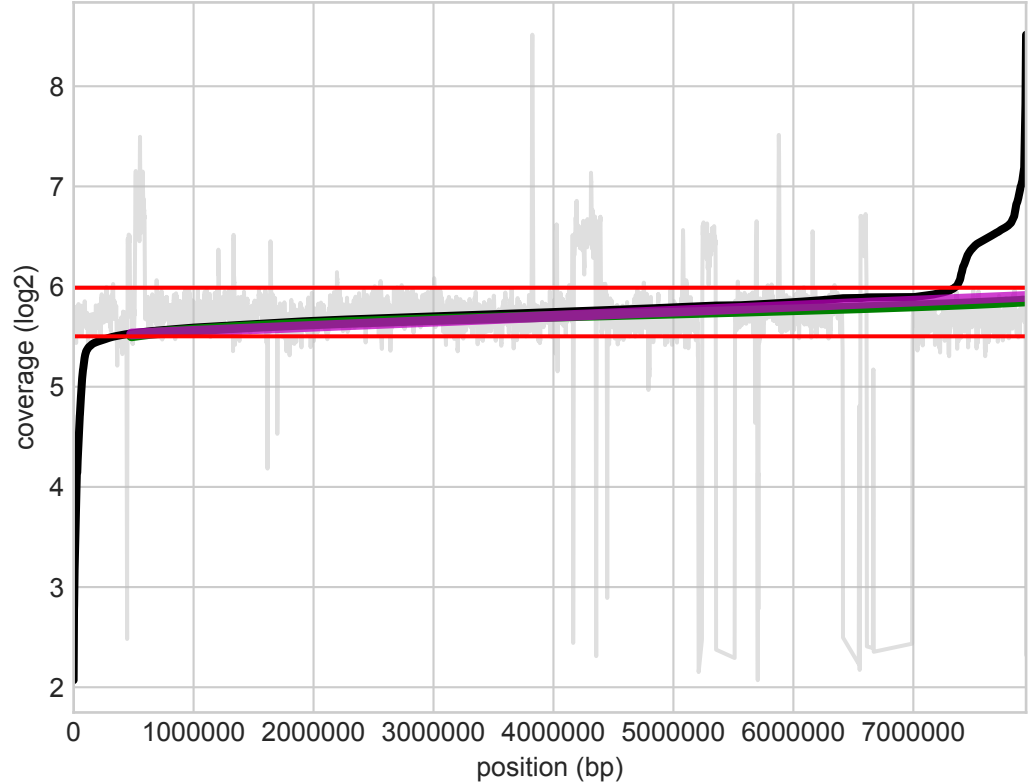


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0724/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0724-A
iRep: n/a r^2: 0.75 avg. cov: 48.49 % windows: 88.38



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0724/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0724-A
iRep: n/a raw iRep: 1.41 GC r^2: 0.00 GC bias: -0.01 avg. cov: 48.49 % windows: 88.38





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