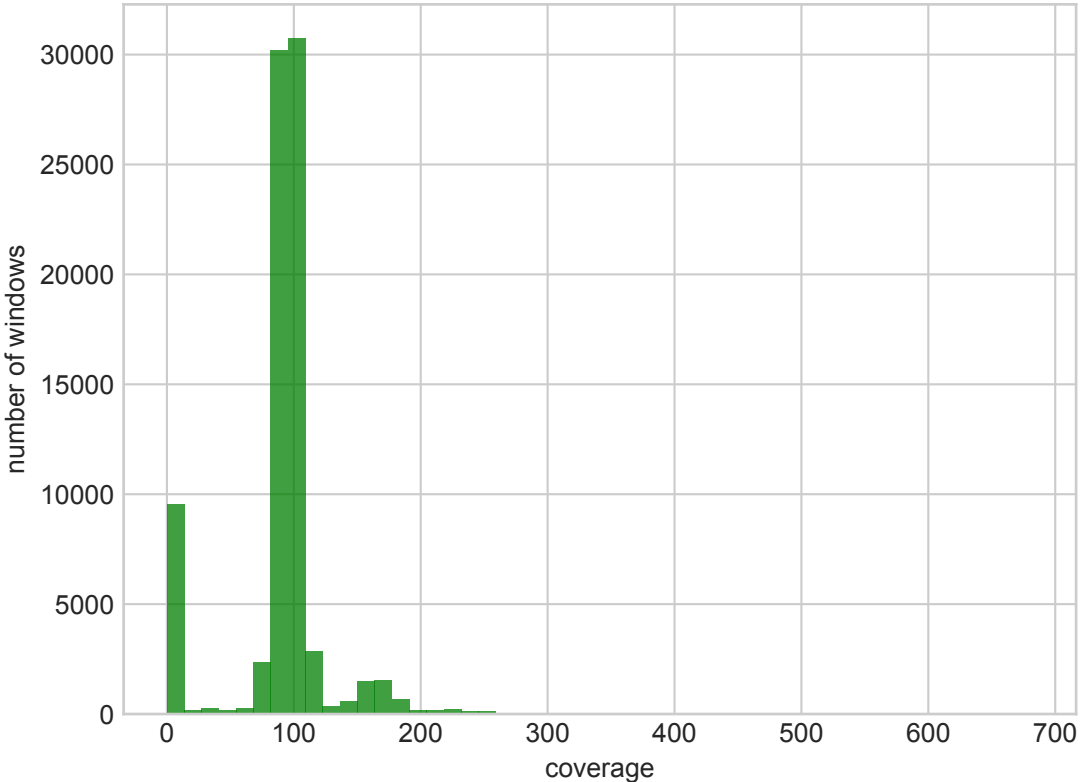
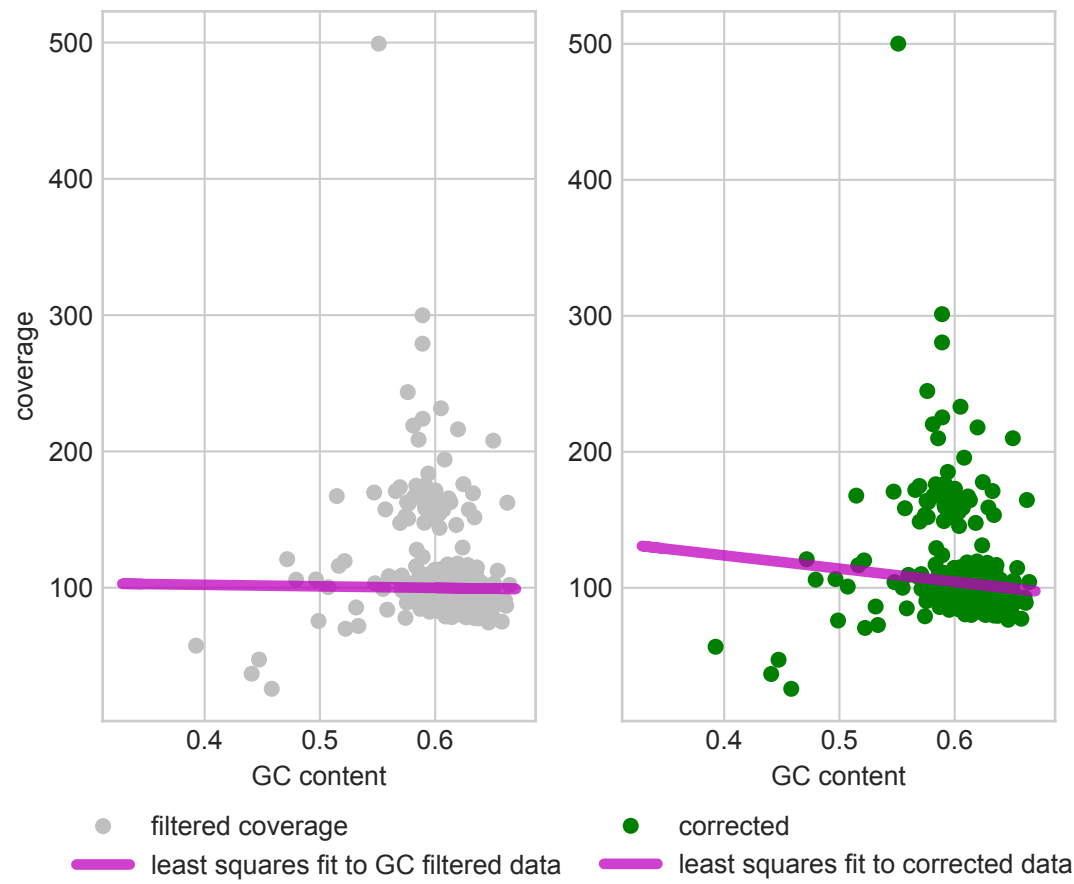


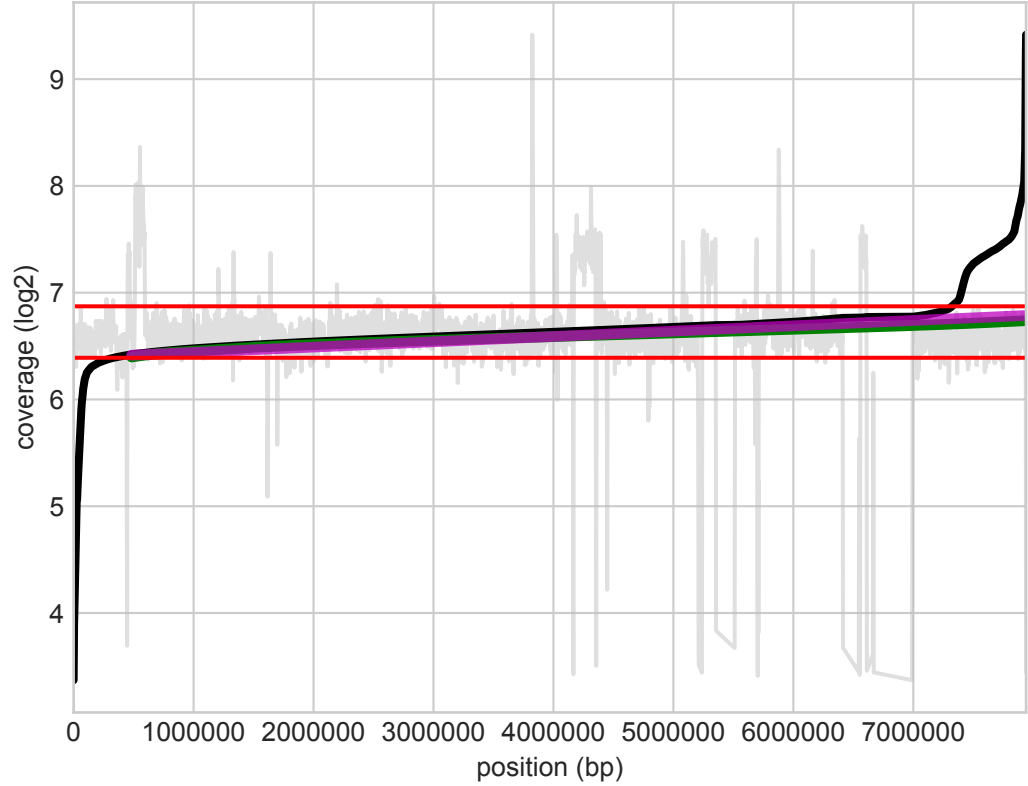
genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0724/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0724-B
iRep: n/a r^2 : 0.75 avg. cov: 89.58 % windows: 88.39



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0724/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0724-B
iRep: n/a raw iRep: 1.41 GC r^2: 0.00 GC bias: -0.00 avg. cov: 89.58 % windows: 88.39



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0724/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0724-B
iRep: n/a un-filtered iRep: 1.40 r^2: 0.75 avg. cov: 89.58 % windows: 88.39



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