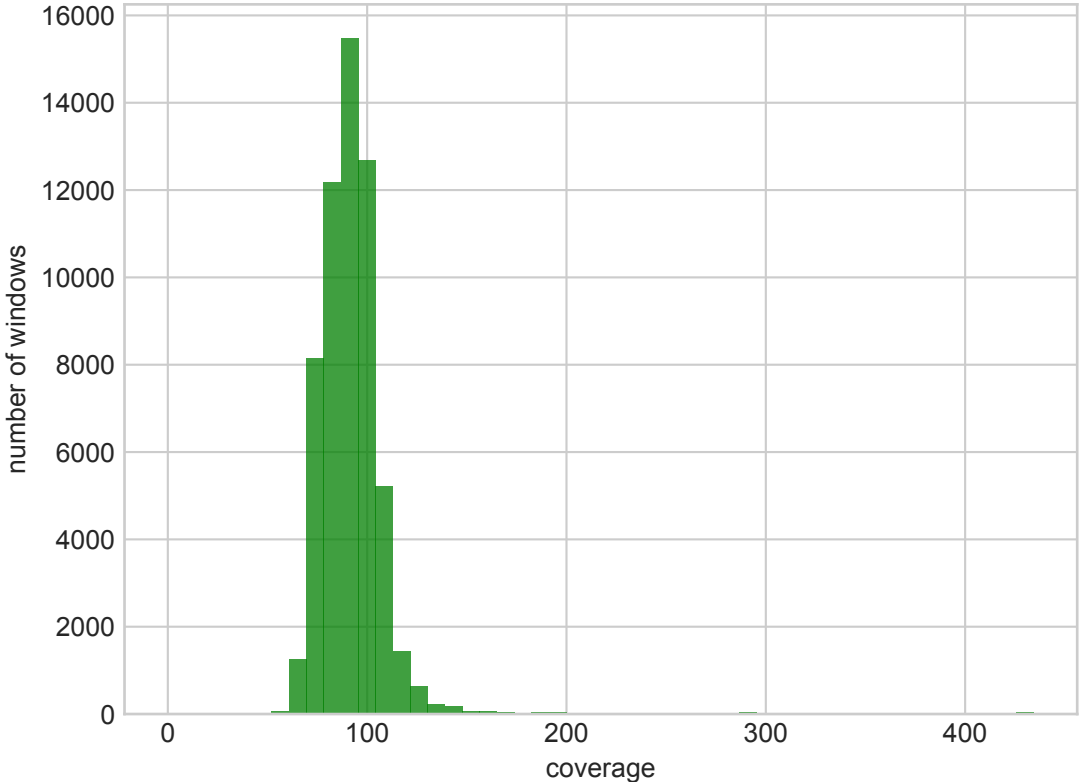
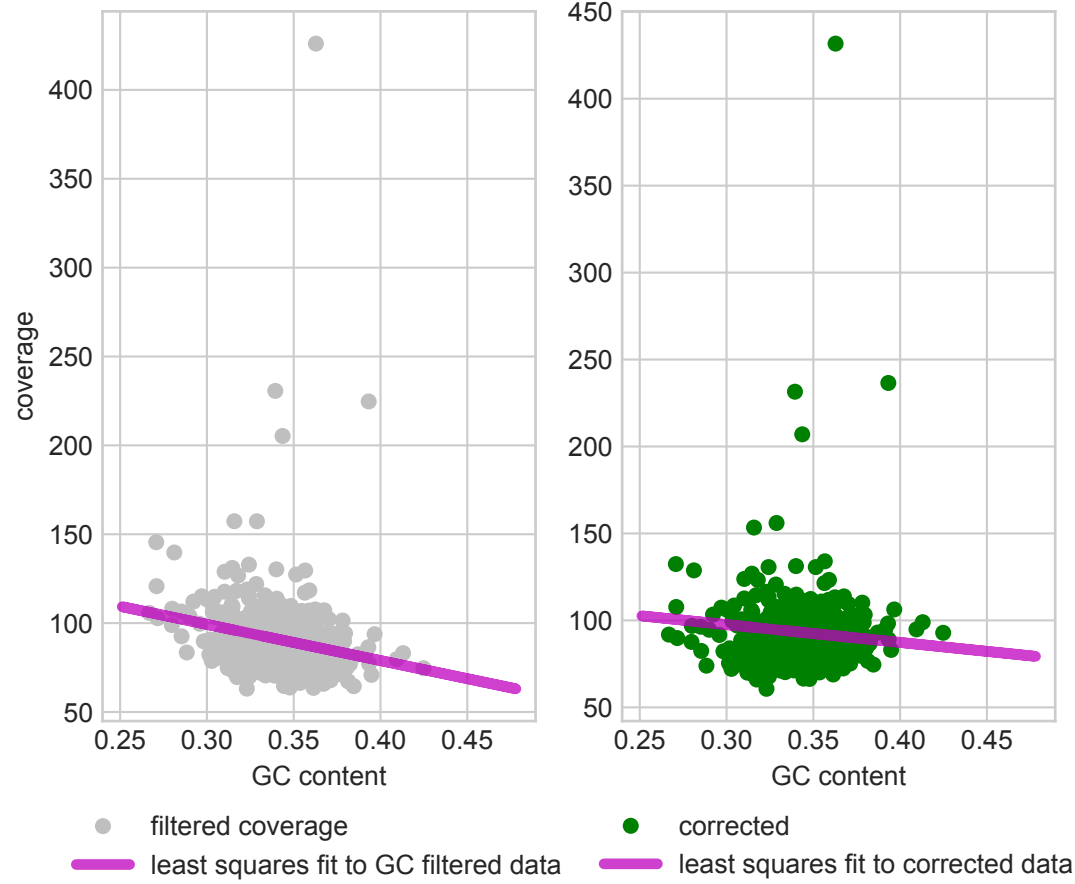
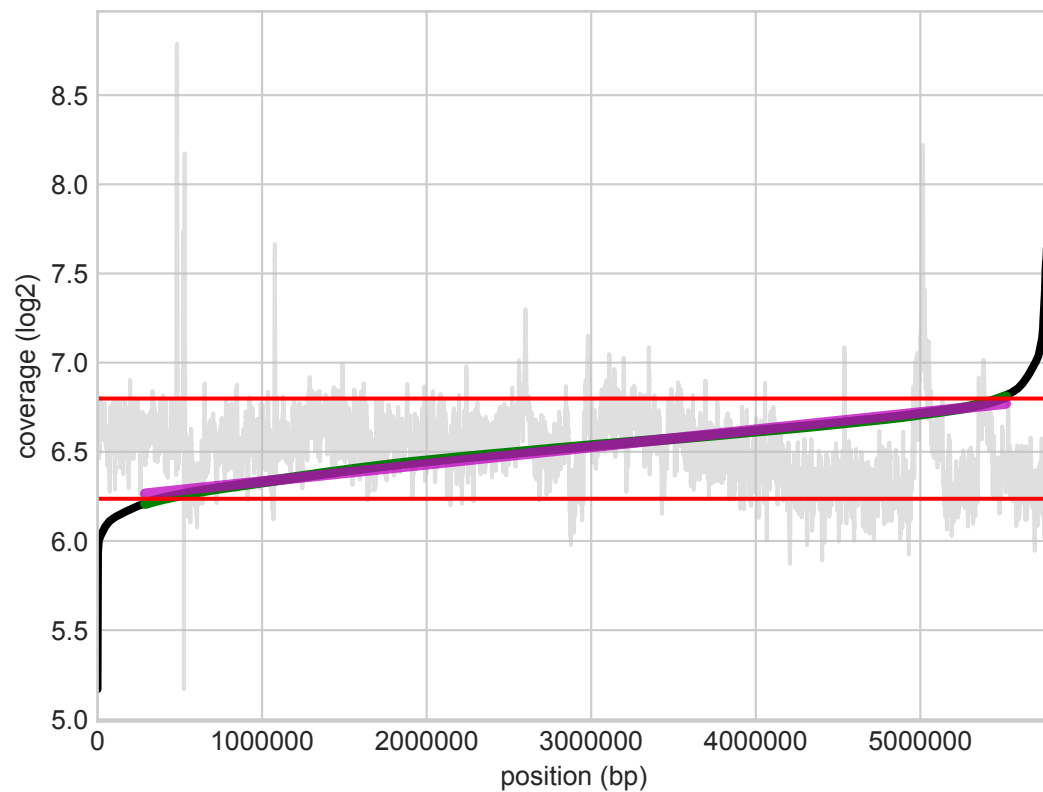


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0721/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0721-D
iRep: 1.48 r^2: 0.99 avg. cov: 92.12 % windows: 99.99





genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0721/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0721-D
iRep: 1.48 un-filtered iRep: 1.48 r^2 : 0.99 avg. cov: 92.12 % windows: 99.99



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