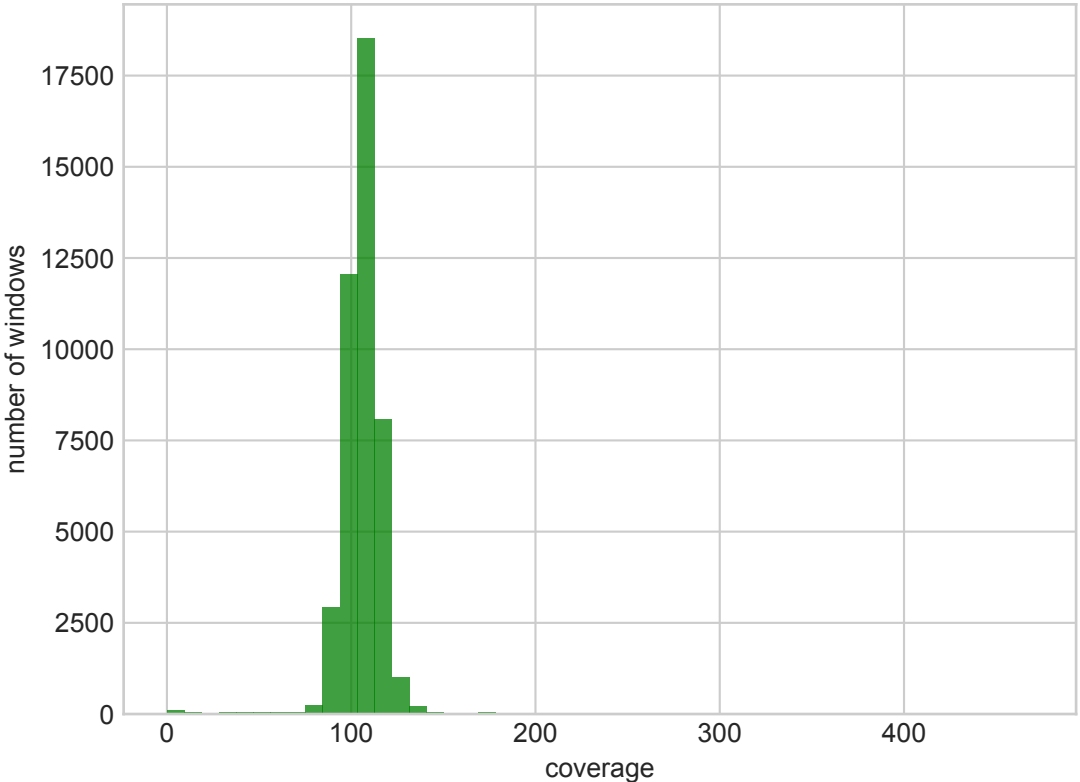
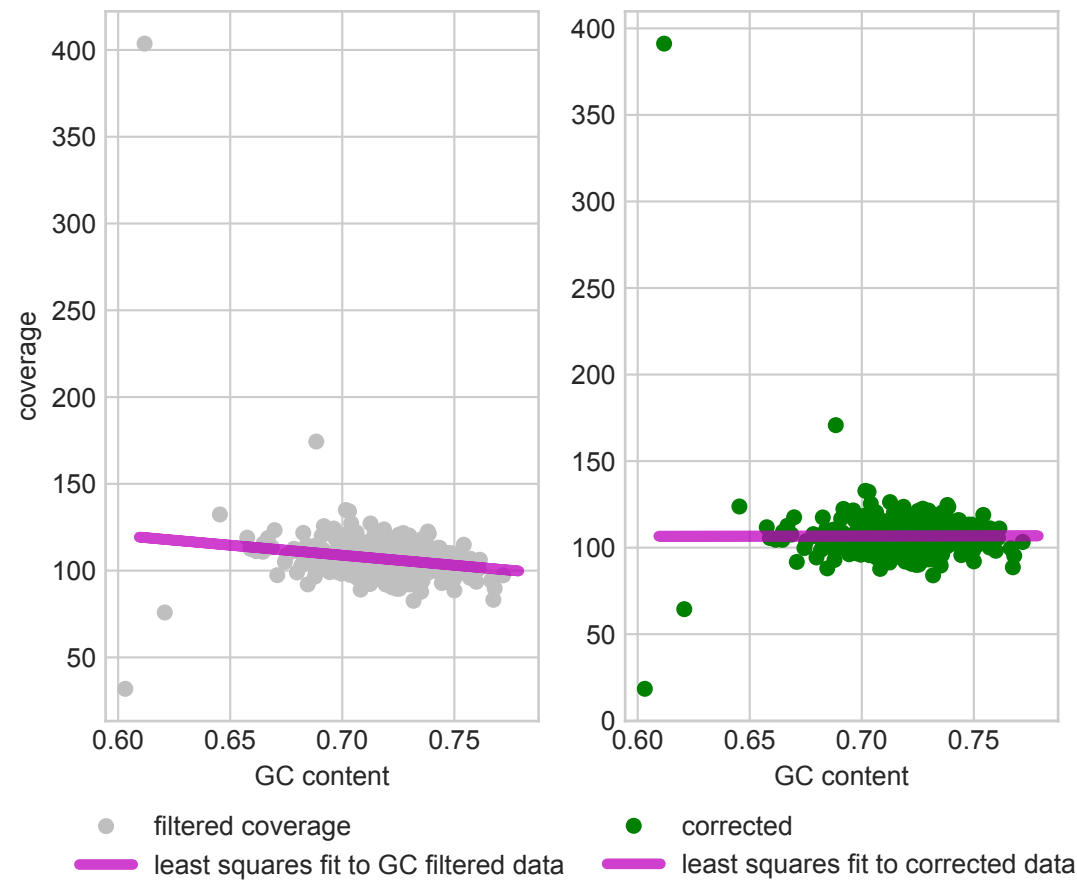


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0722/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0722-A
iRep: 1.26 r^2: 0.98 avg. cov: 106.24 % windows: 99.73



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0722/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0722-A
iRep: 1.26 raw iRep: 1.27 GC r^2: 0.08 GC bias: -9.17 avg. cov: 106.24 % windows: 99.73



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0722/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0722-A
iRep: 1.26 un-filtered iRep: 1.26 r^2: 0.98 avg. cov: 106.24 % windows: 99.73

