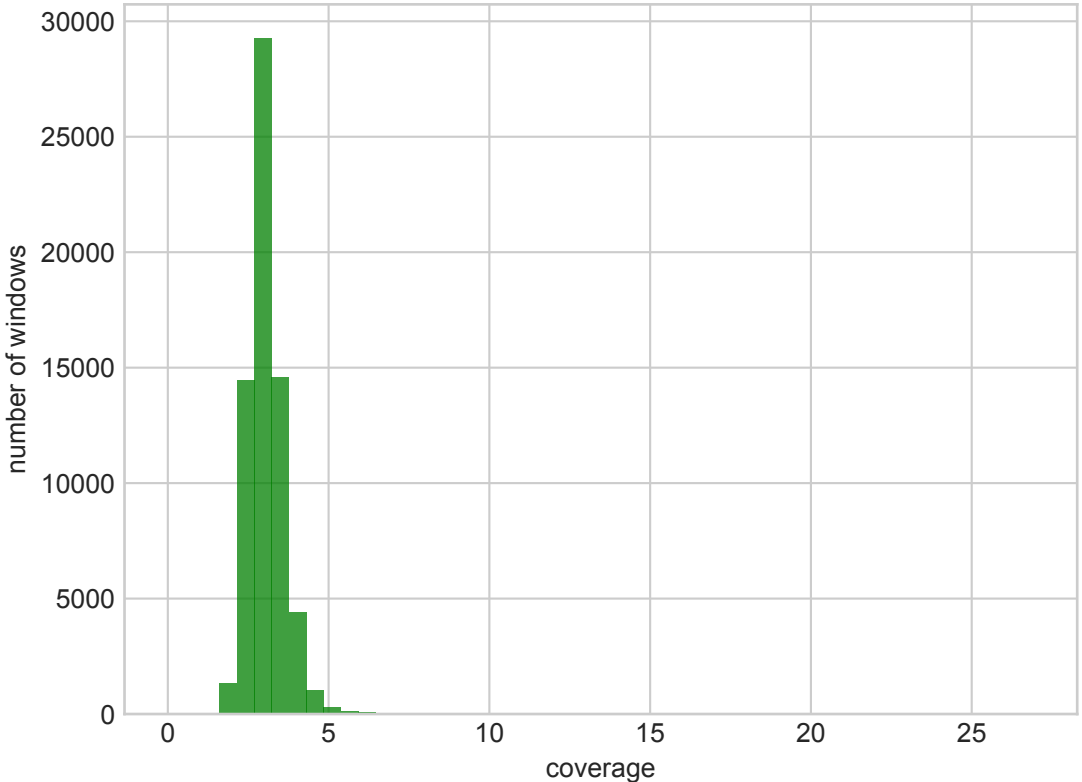
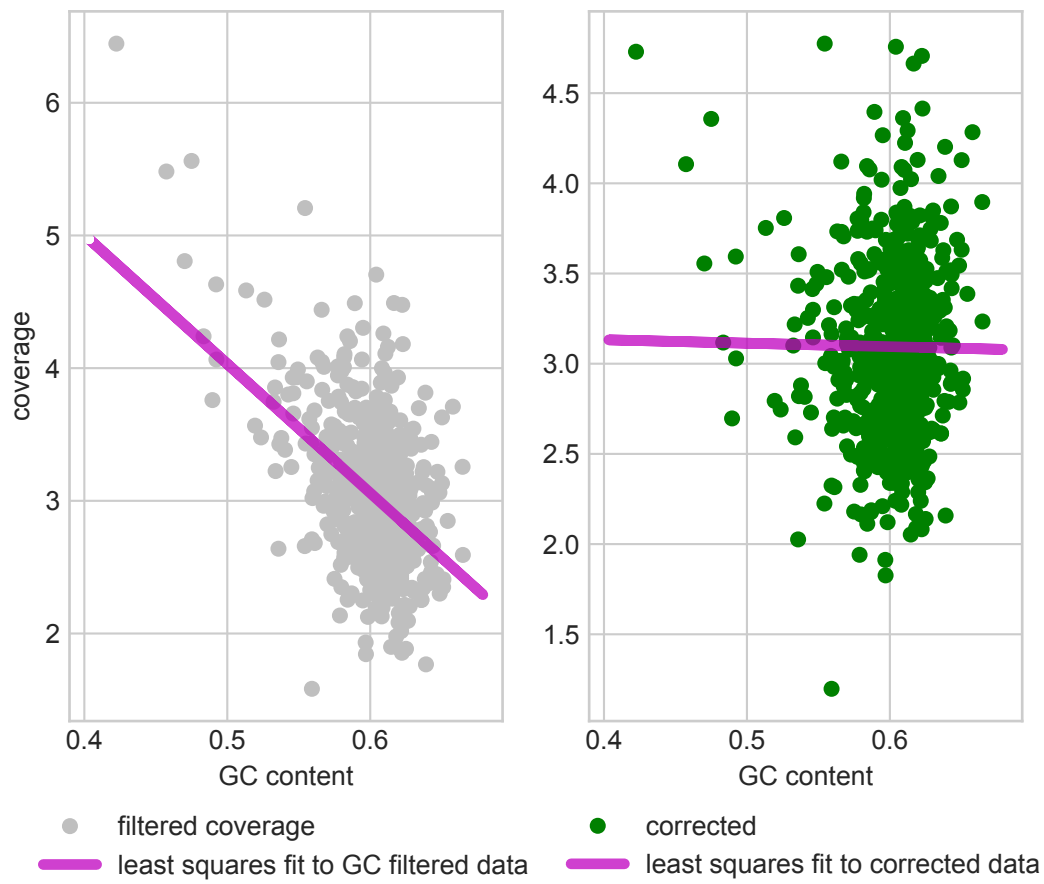


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0710/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0710-D
iRep: n/a r^2: 0.99 avg. cov: 3.07 % windows: 99.96



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0710/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0710-D
iRep: n/a raw iRep: 1.62 GC r^2: 0.24 GC bias: -2.35 avg. cov: 3.07 % windows: 99.96



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0710/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0710-D
iRep: n/a un-filtered iRep: 1.56 r^2: 0.99 avg. cov: 3.07 % windows: 99.96

