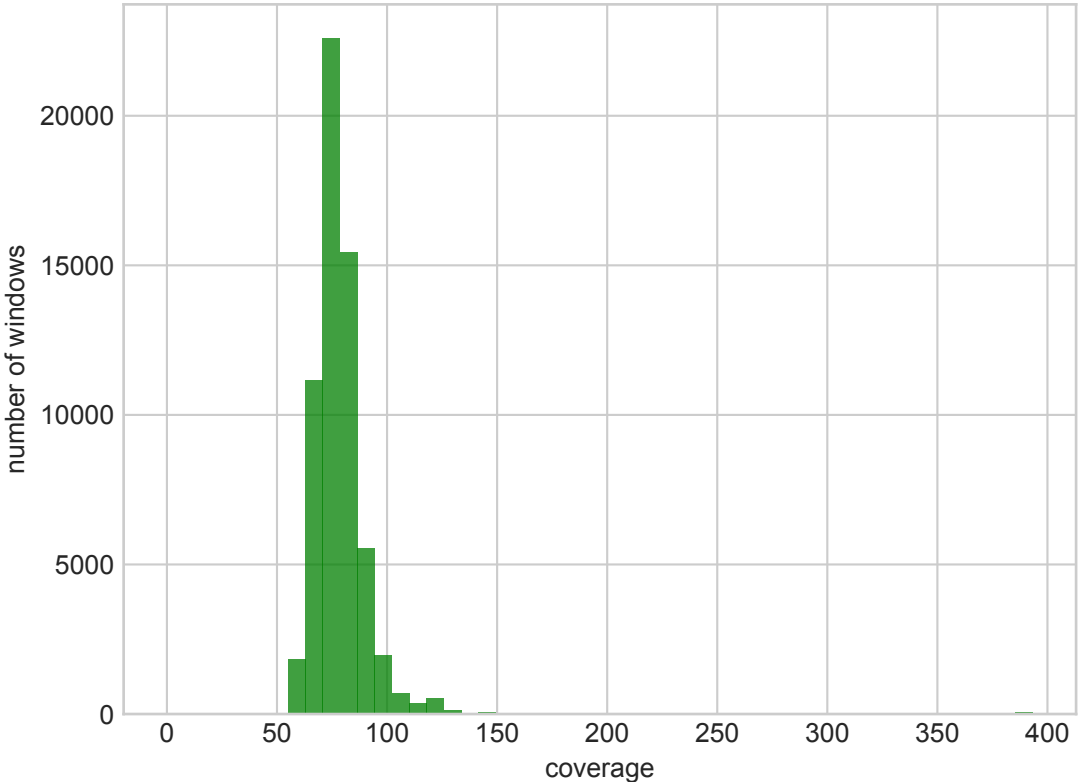
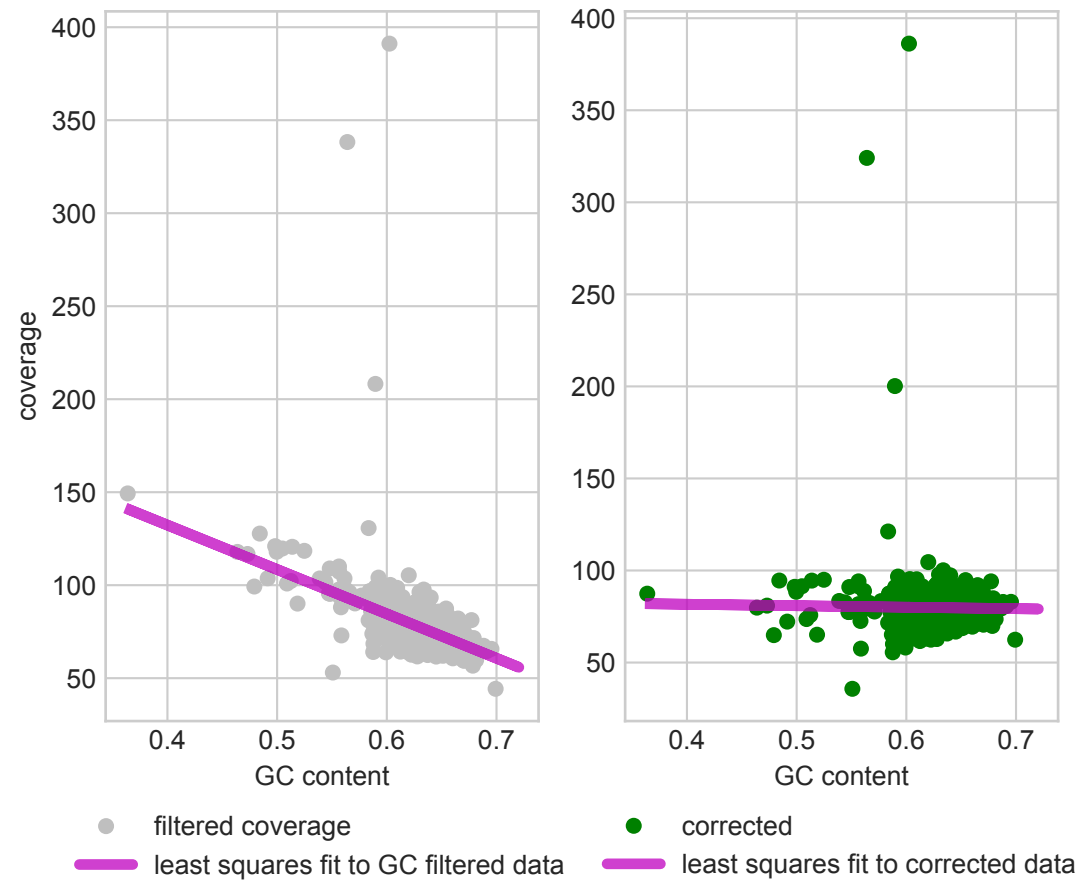


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0711/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0711-K  
iRep: 1.30 r^2: 0.98 avg. cov: 79.15 % windows: 99.97



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0711/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0711-K  
iRep: 1.30 raw iRep: 1.41 GC r^2: 0.54 GC bias: -128.75 avg. cov: 79.15 % windows: 99.97



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0711/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0711-K  
iRep: 1.30 un-filtered iRep: 1.30 r^2: 0.98 avg. cov: 79.15 % windows: 99.97

