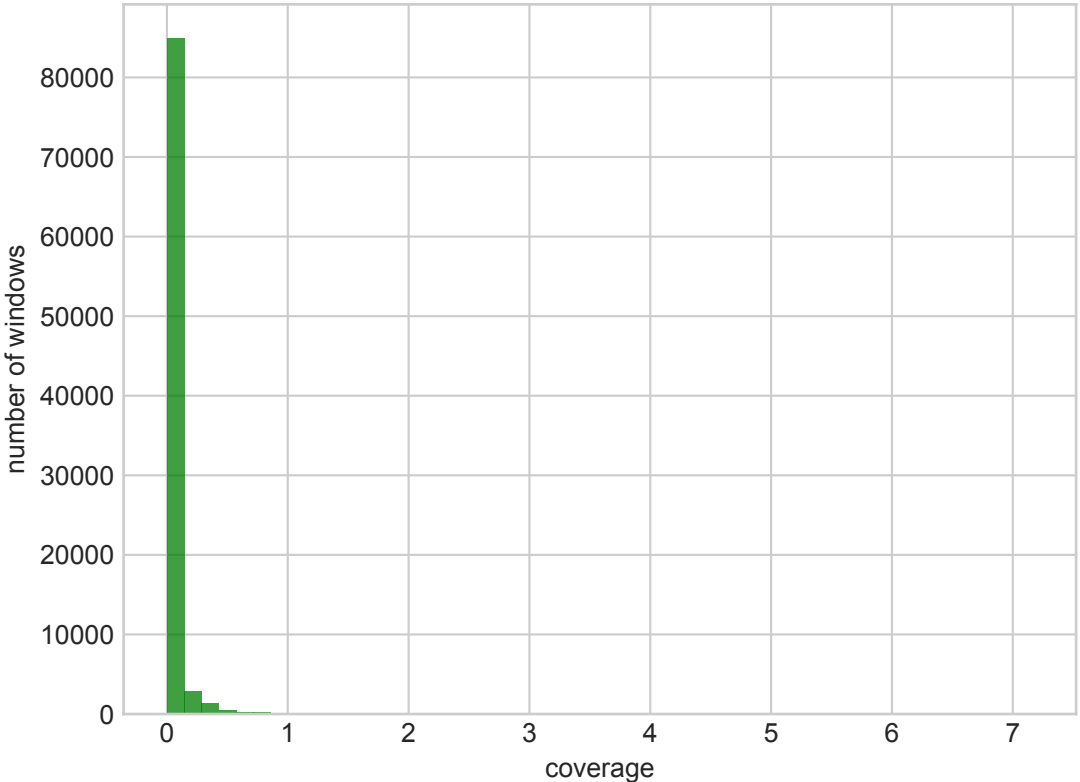
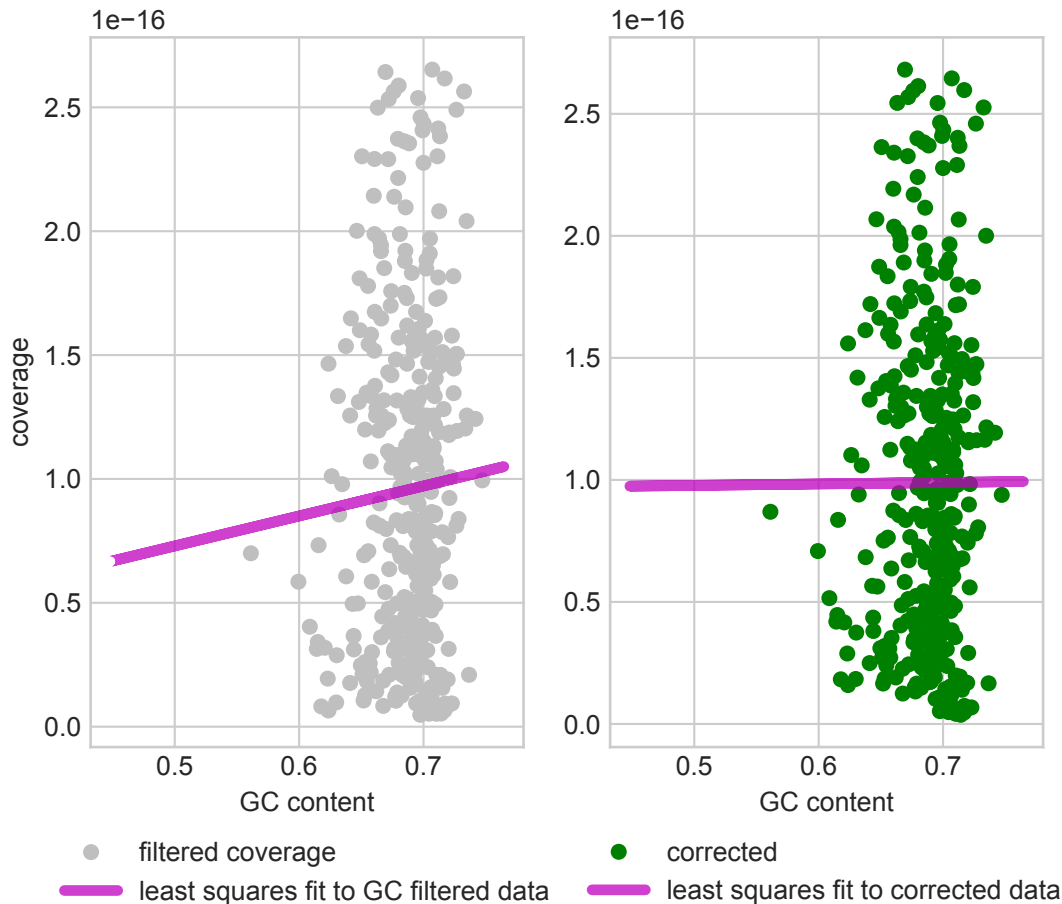


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0706/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0706-A  
iRep: n/a r^2: 0.96 avg. cov: 0.03 % windows: 39.35



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0706/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0706-A  
iRep: n/a raw iRep: 15.88 GC r^2: 0.00 GC bias: 0.00 avg. cov: 0.03 % windows: 39.35



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0706/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0706-A  
iRep: n/a un-filtered iRep: 14.51 r^2: 0.96 avg. cov: 0.03 % windows: 39.35

