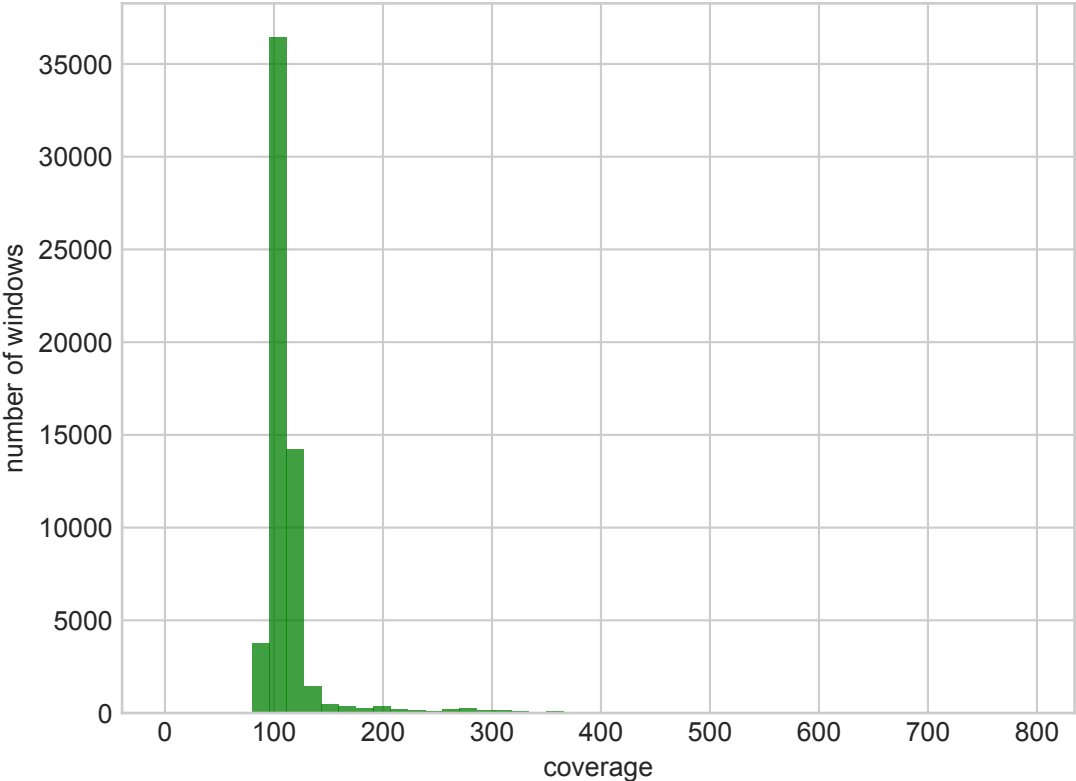
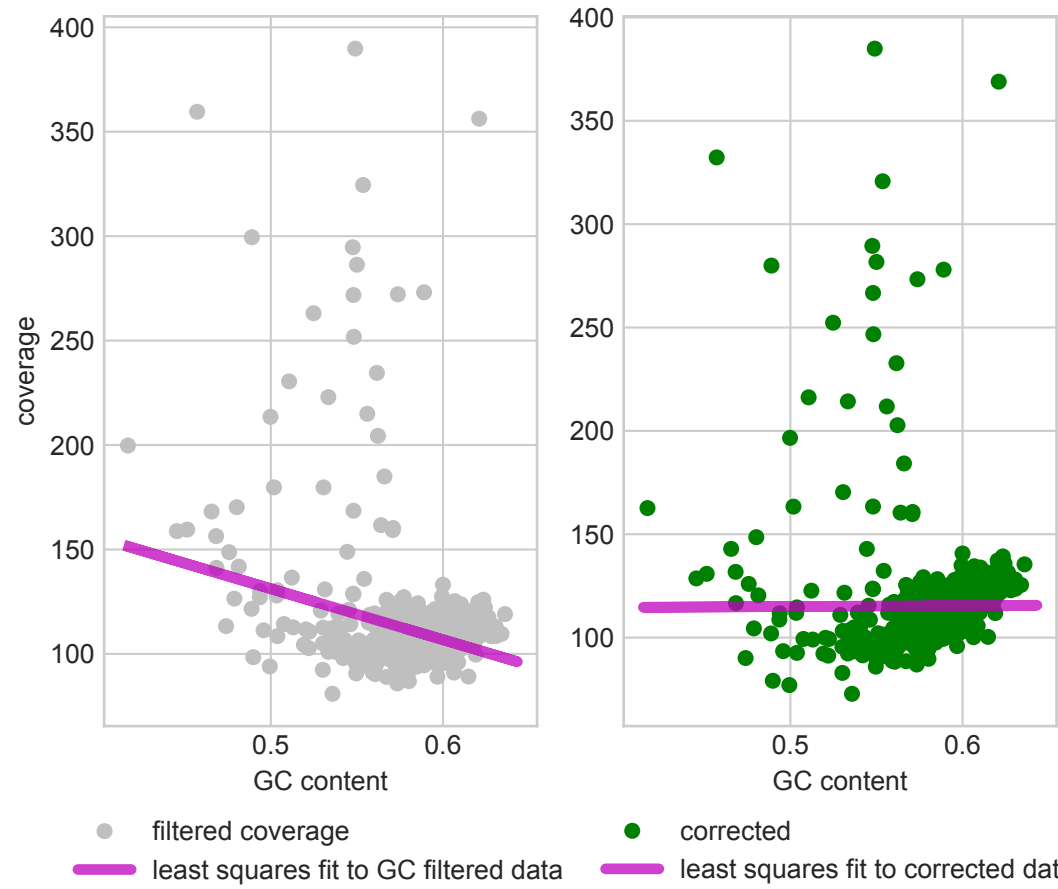


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016\_KBSGenomes\_Annotate/KBS0707/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0707-B  
iRep: 1.38 r^2: 0.97 avg. cov: 114.31 % windows: 99.98



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016\_KBSGenomes\_Annotate/KBS0707/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0707-B  
iRep: 1.38 raw iRep: 1.31 GC r^2: 0.09 GC bias: -21.35 avg. cov: 114.31 % windows: 99.98



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016\_KBSGenomes\_Annotate/KBS0707/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0707-B  
iRep: 1.38 un-filtered iRep: 1.38 r^2: 0.97 avg. cov: 114.31 % windows: 99.98

