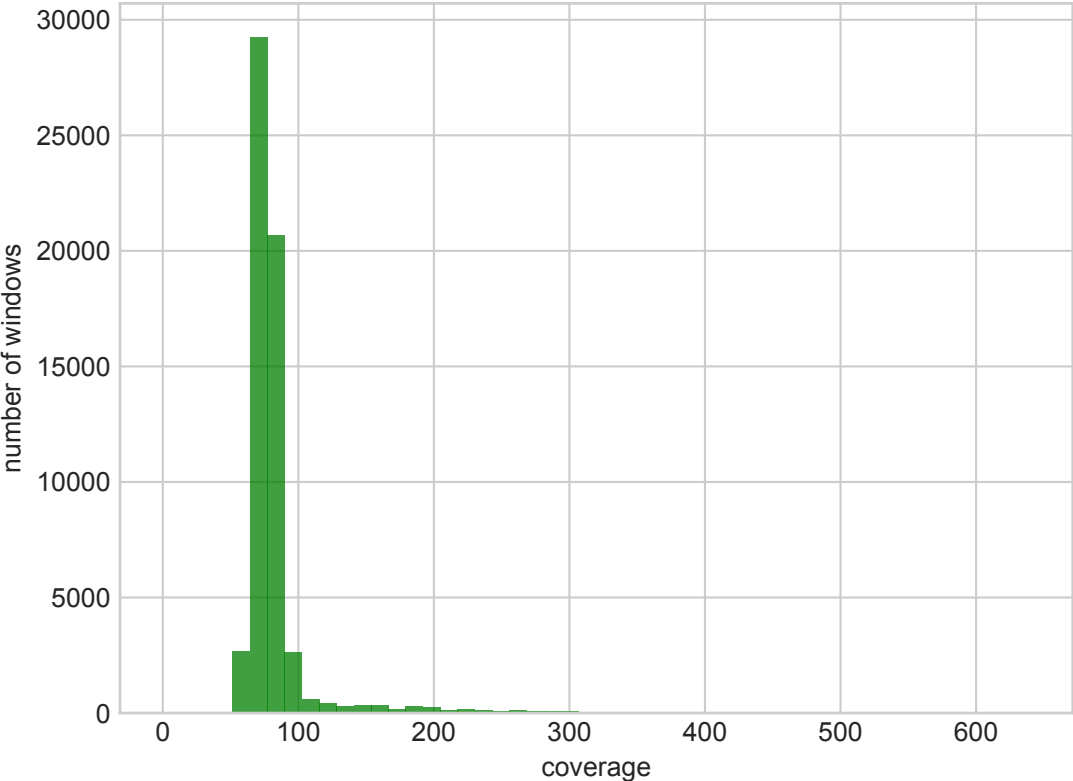


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016\_KBSGenomes\_Annotate/KBS0707/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0707-C  
iRep: 1.59 r^2: 0.97 avg. cov: 83.16 % windows: 99.92



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016\_KBSGenomes\_Annotate/KBS0707/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0707-C  
iRep: 1.59 raw iRep: 1.43 GC r^2: 0.11 GC bias: -30.80 avg. cov: 83.16 % windows: 99.92

