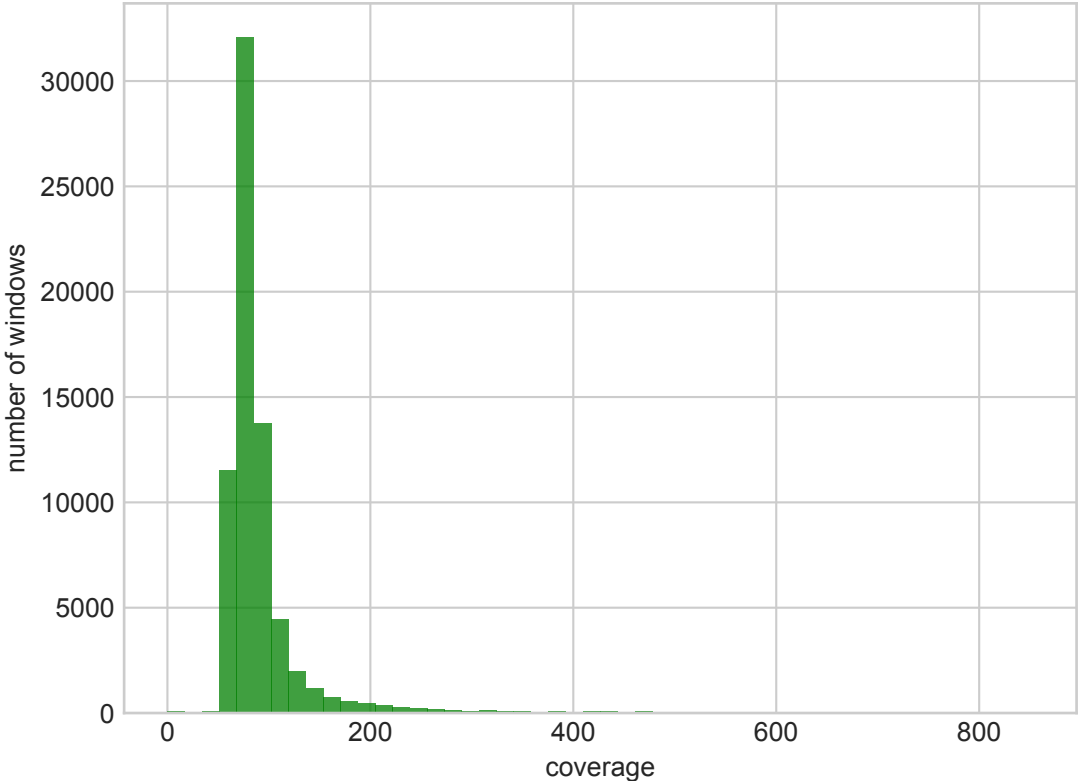


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016\_KBSGenomes\_Annotate/ATCC13985/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/ATCC13985-B  
iRep: 1.72 r^2: 0.98 avg. cov: 92.79 % windows: 99.80



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016\_KBSGenomes\_Annotate/ATCC13985/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/ATCC13985-B  
iRep: 1.72 raw iRep: 1.98 GC r^2: 0.61 GC bias: -505.30 avg. cov: 92.79 % windows: 99.80

