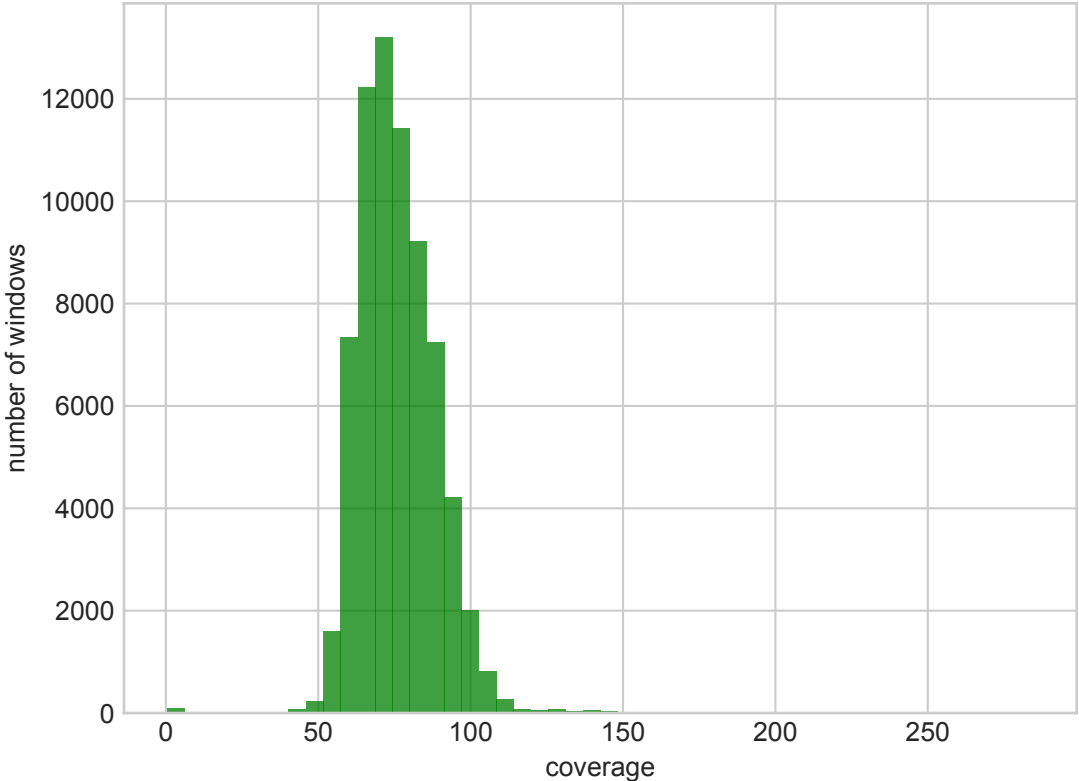


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016\_KBSGenomes\_Annotate/KBS0712/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0712-C  
iRep: 1.59 r^2: 1.00 avg. cov: 76.06 % windows: 99.85



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016\_KBSGenomes\_Annotate/KBS0712/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0712-C  
iRep: 1.59 raw iRep: 1.62 GC r<sup>2</sup>: 0.10 GC bias: -11.22 avg. cov: 76.06 % windows: 99.85

