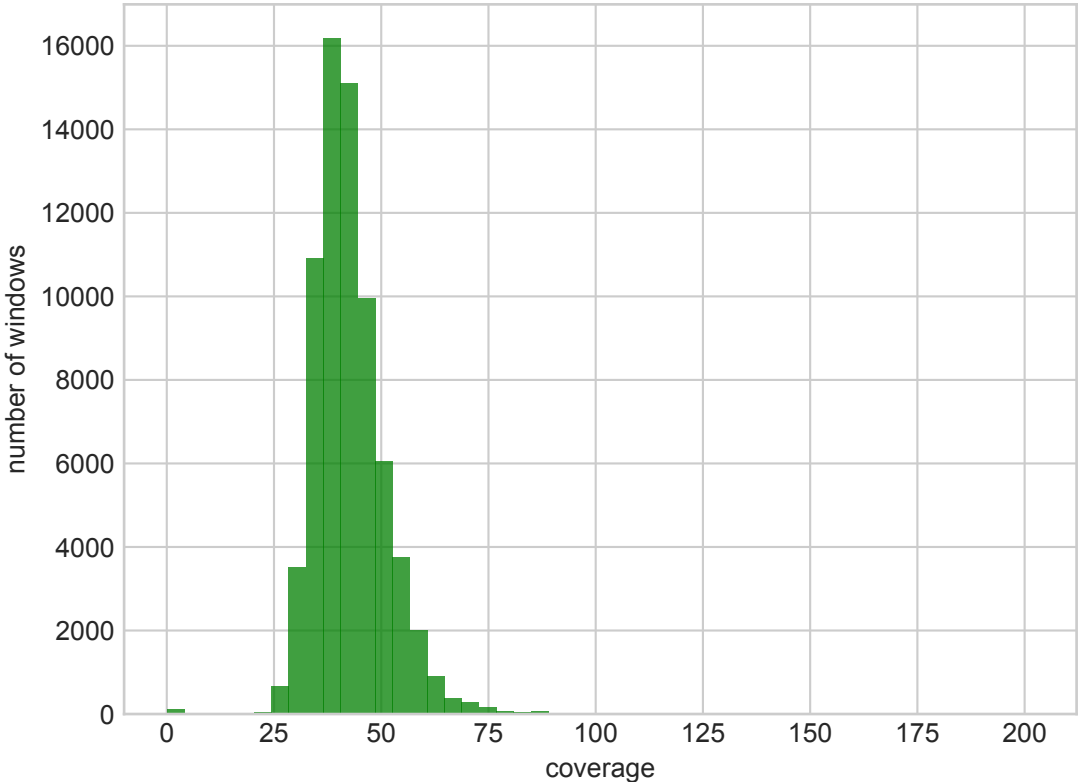


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016\_KBSGenomes\_Annotate/KBS0712/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0712-A  
iRep: 1.60 r^2: 0.99 avg. cov: 42.74 % windows: 99.84



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016\_KBSGenomes\_Annotate/KBS0712/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0712-A  
iRep: 1.60 raw iRep: 1.70 GC r^2: 0.31 GC bias: -44.10 avg. cov: 42.74 % windows: 99.84

