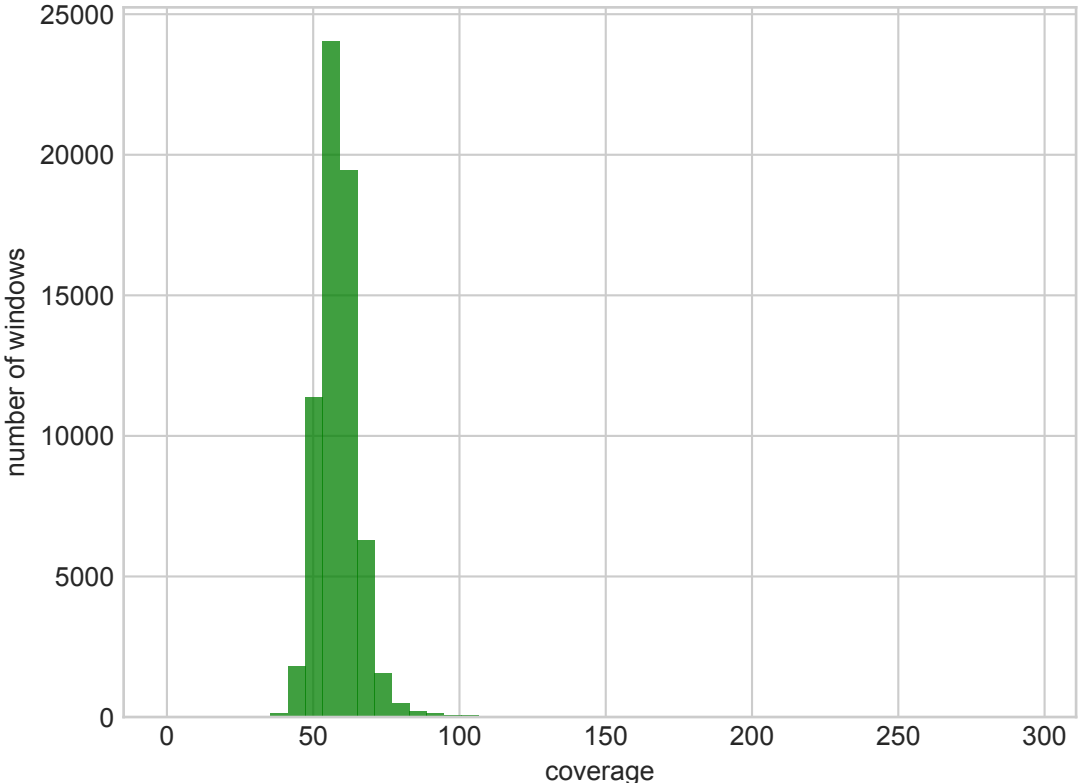


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0710/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0710-A  
iRep: 1.37 r^2: 0.99 avg. cov: 58.71 % windows: 99.97



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0710/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0710-A  
iRep: 1.37 raw iRep: 1.37 GC r^2: 0.10 GC bias: -7.85 avg. cov: 58.71 % windows: 99.97

