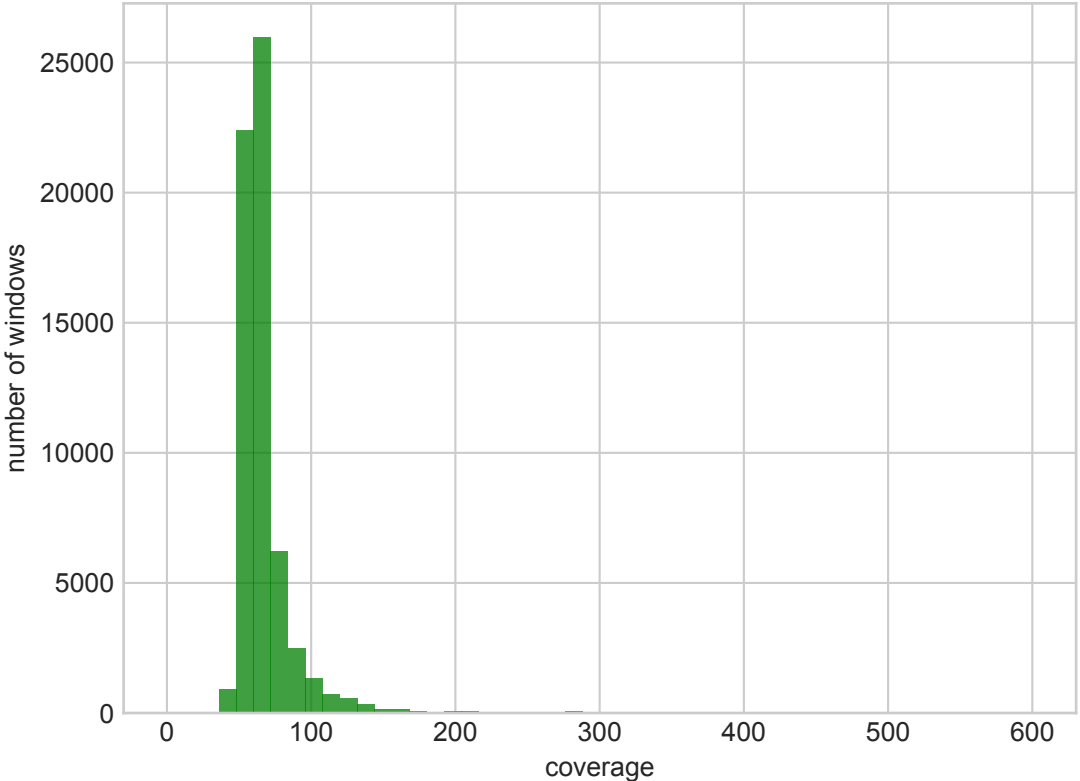


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0802/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0802-A  
iRep: 1.31   r^2: 0.98   avg. cov: 67.47   % windows: 99.94



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0802/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0802-A  
iRep: 1.31 raw iRep: 1.64 GC r^2: 0.78 GC bias: -318.77 avg. cov: 67.47 % windows: 99.94

