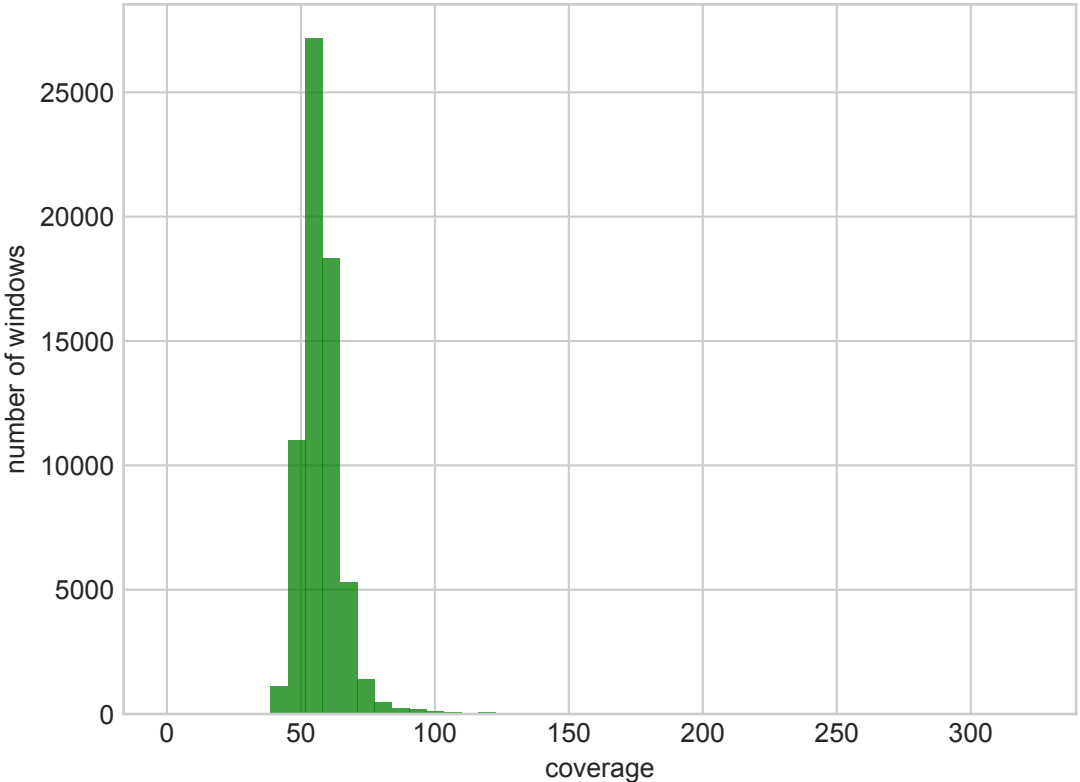


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0710/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0710-B  
iRep: 1.35   r^2: 0.99   avg. cov: 57.67   % windows: 99.98



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015\_SoilGenomes\_Annotate/KBS0710/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa\_sam/KBS0710-B  
iRep: 1.35 raw iRep: 1.39 GC r^2: 0.30 GC bias: -46.03 avg. cov: 57.67 % windows: 99.98

