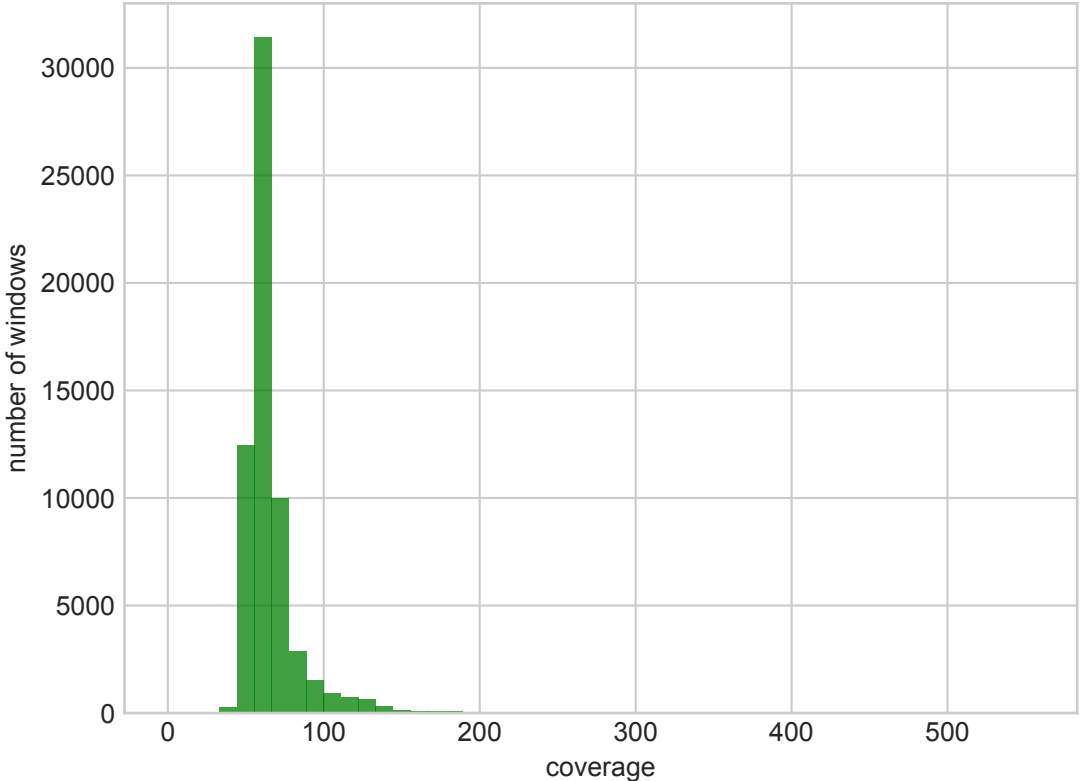


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0802/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0802-C
iRep: 1.30 r^2: 0.98 avg. cov: 66.34 % windows: 99.95



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0802/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0802-C
iRep: 1.30 raw iRep: 1.63 GC r^2: 0.72 GC bias: -284.34 avg. cov: 66.34 % windows: 99.95

