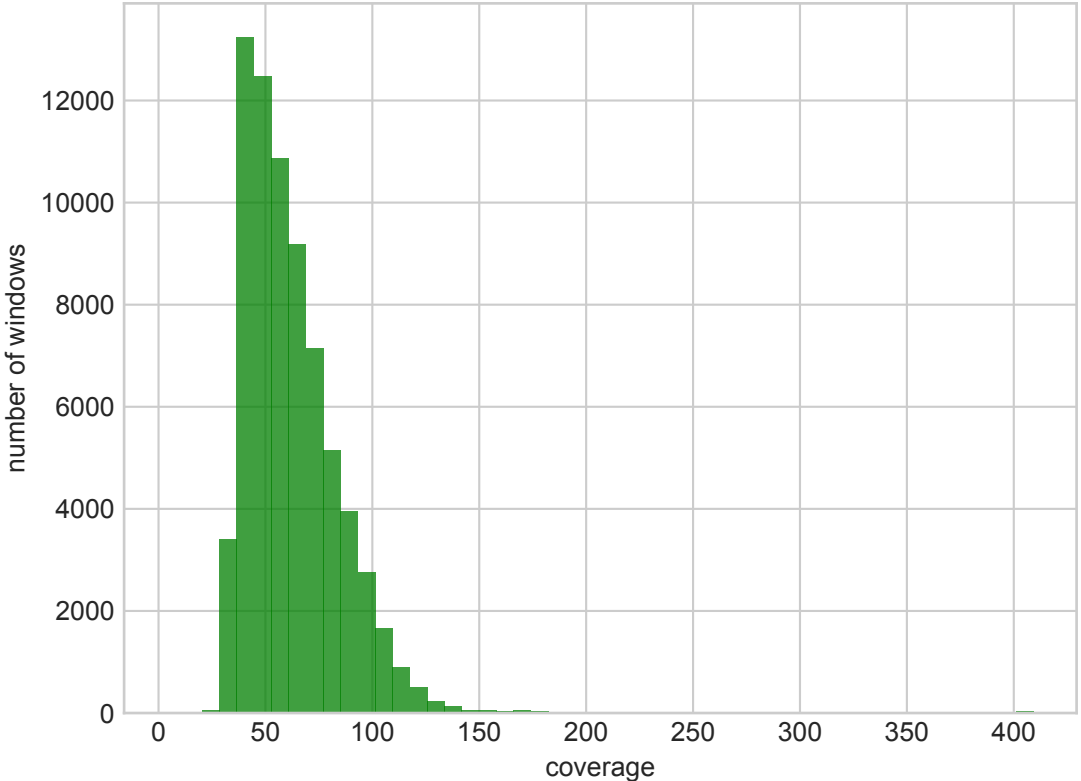


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0801/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0801-A
iRep: 2.05 r^2: 0.99 avg. cov: 62.50 % windows: 99.99



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0801/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0801-A
iRep: 2.05 raw iRep: 2.80 GC r^2: 0.54 GC bias: -282.20 avg. cov: 62.50 % windows: 99.99

