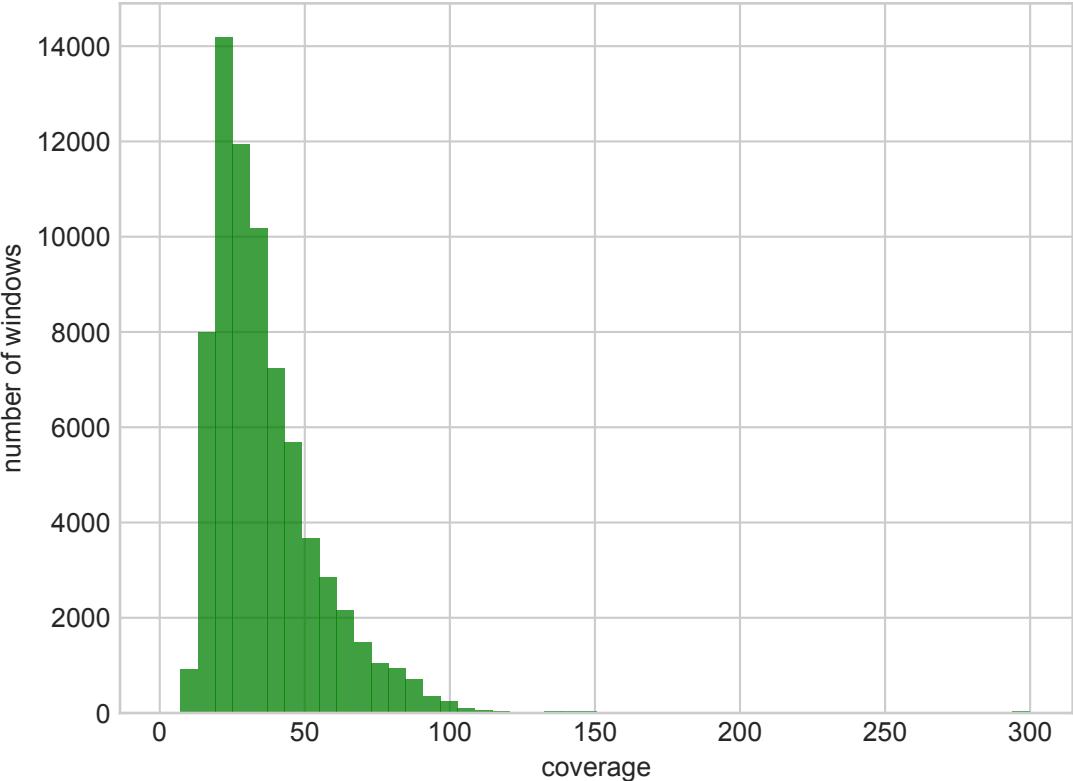


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0801/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0801-D
iRep: 2.40 r^2: 0.98 avg. cov: 36.49 % windows: 99.92



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0801/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0801-D
iRep: 2.40 raw iRep: 4.21 GC r^2: 0.65 GC bias: -319.84 avg. cov: 36.49 % windows: 99.92

