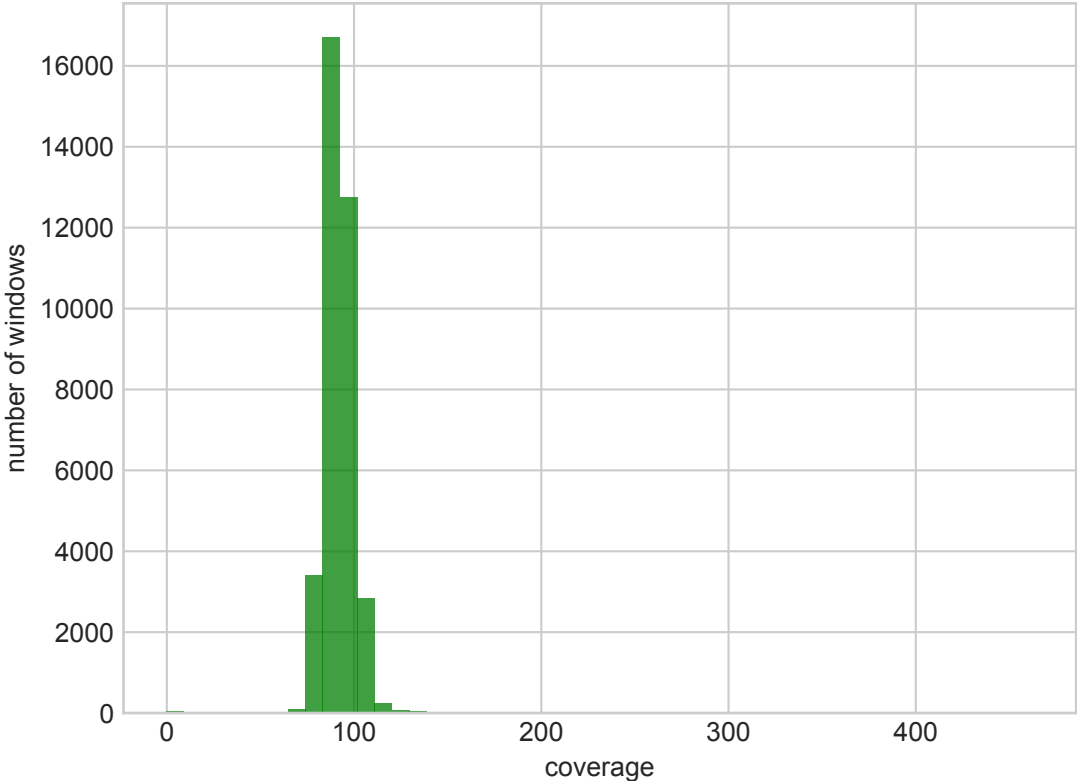


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0702/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0702-F
iRep: 1.23 r^2: 0.99 avg. cov: 92.67 % windows: 99.90



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0702/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0702-F
iRep: 1.23 raw iRep: 1.26 GC r^2: 0.23 GC bias: -33.72 avg. cov: 92.67 % windows: 99.90

