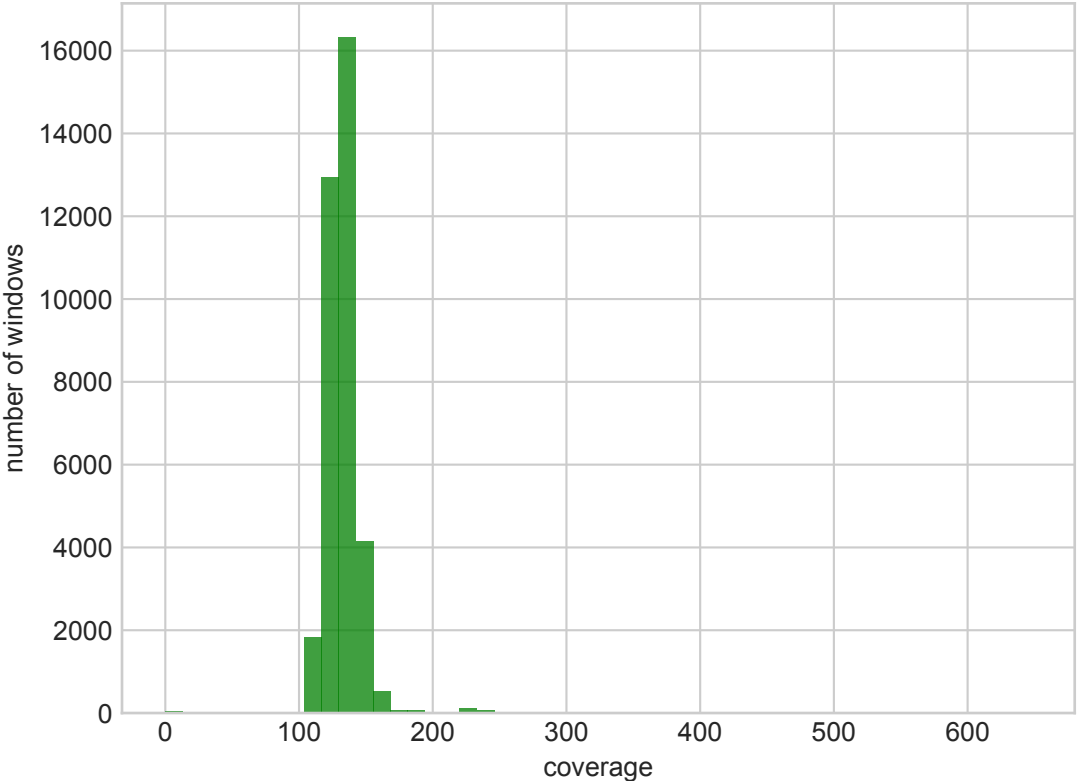


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0702/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0702-D
iRep: 1.23 r^2: 0.99 avg. cov: 133.52 % windows: 99.90



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0702/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0702-D
iRep: 1.23 raw iRep: 1.25 GC r²: 0.20 GC bias: -39.52 avg. cov: 133.52 % windows: 99.90

