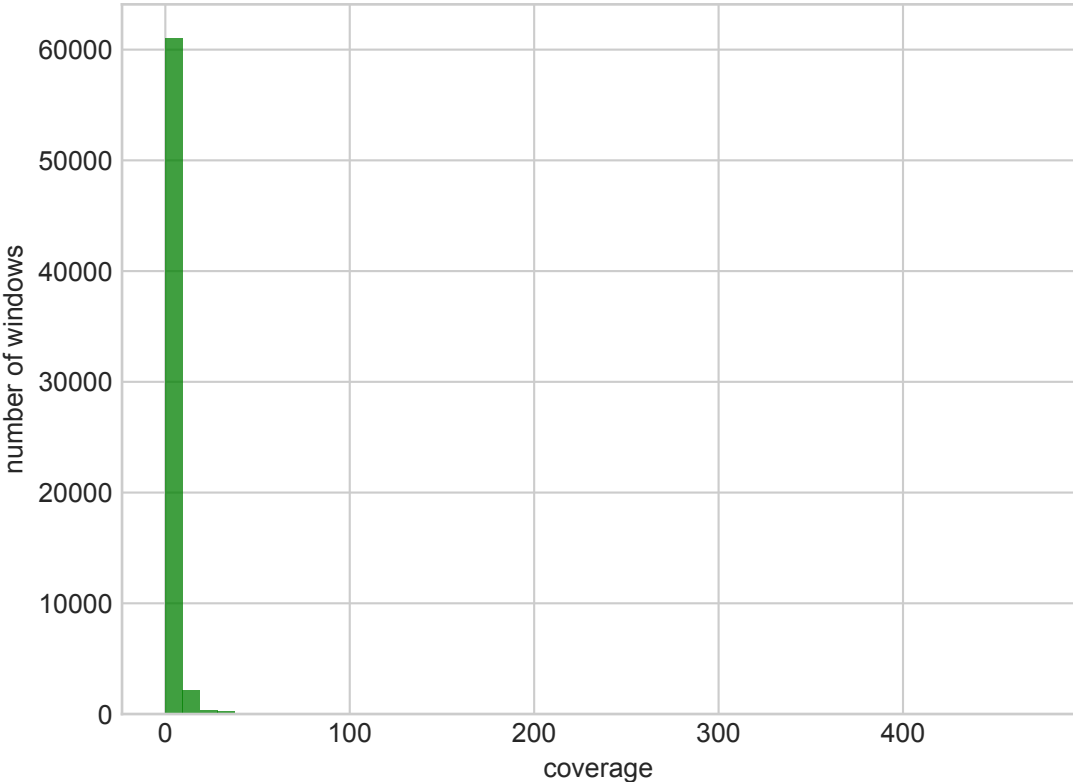


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/ATCC43928/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/ATCC43928-D
iRep: n/a r^2: 0.97 avg. cov: 3.72 % windows: 80.11



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/ATCC43928/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/ATCC43928-D
iRep: n/a raw iRep: 14.70 GC r²: 0.01 GC bias: -0.21 avg. cov: 3.72 % windows: 80.11

