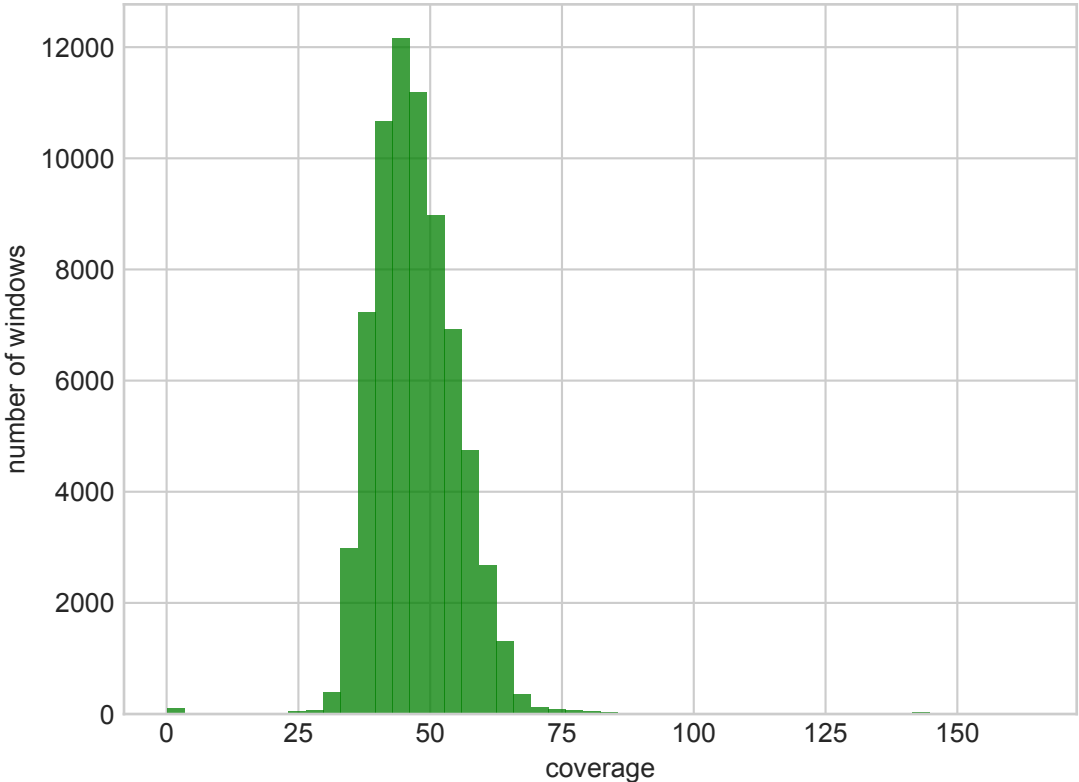


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0712/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0712-D
iRep: 1.60 r^2: 0.99 avg. cov: 47.36 % windows: 99.85



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0712/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0712-D
iRep: 1.60 raw iRep: 1.64 GC r^2: 0.11 GC bias: -8.77 avg. cov: 47.36 % windows: 99.85

