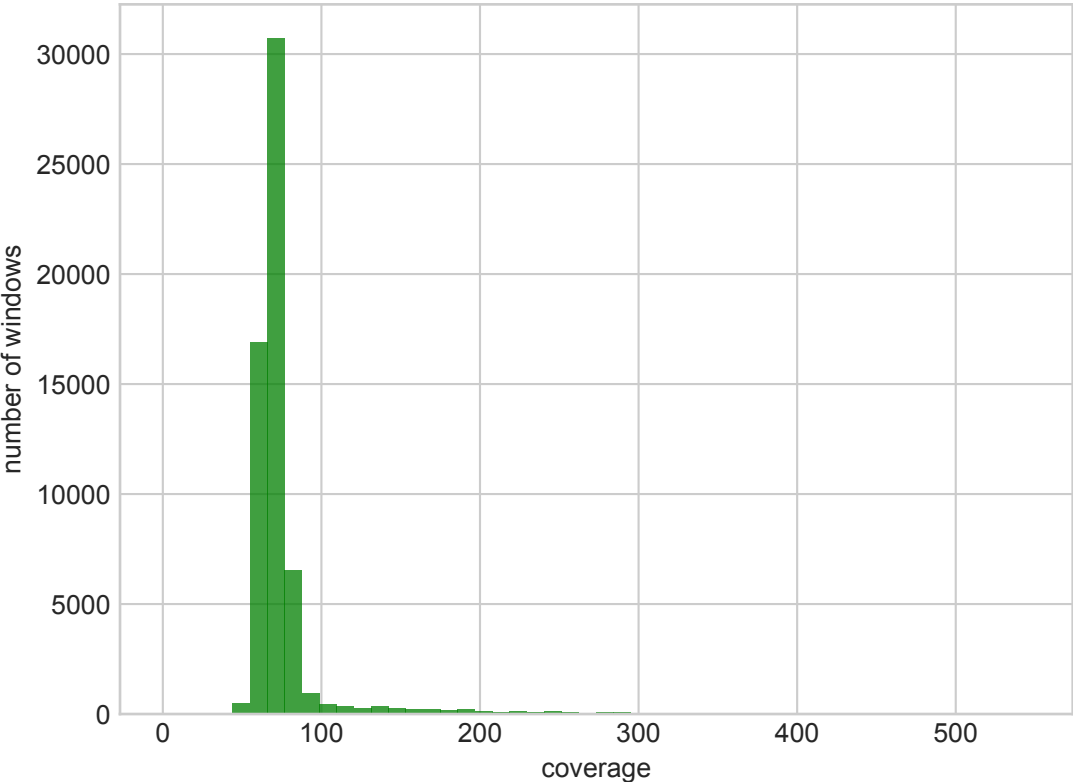genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0707/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0707-D
iRep: 1.57   r^2: 0.97   avg. cov: 75.23   % windows: 99.98

genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0707/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0707-D
iRep: 1.57   raw iRep: 1.42   GC r^2: 0.10   GC bias: -22.99   avg. cov: 75.23   % windows: 99.98

genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/KBS0707/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0707-D
iRep: 1.57   un-filtered iRep: 1.57   r^2: 0.97   avg. cov: 75.23   % windows: 99.98