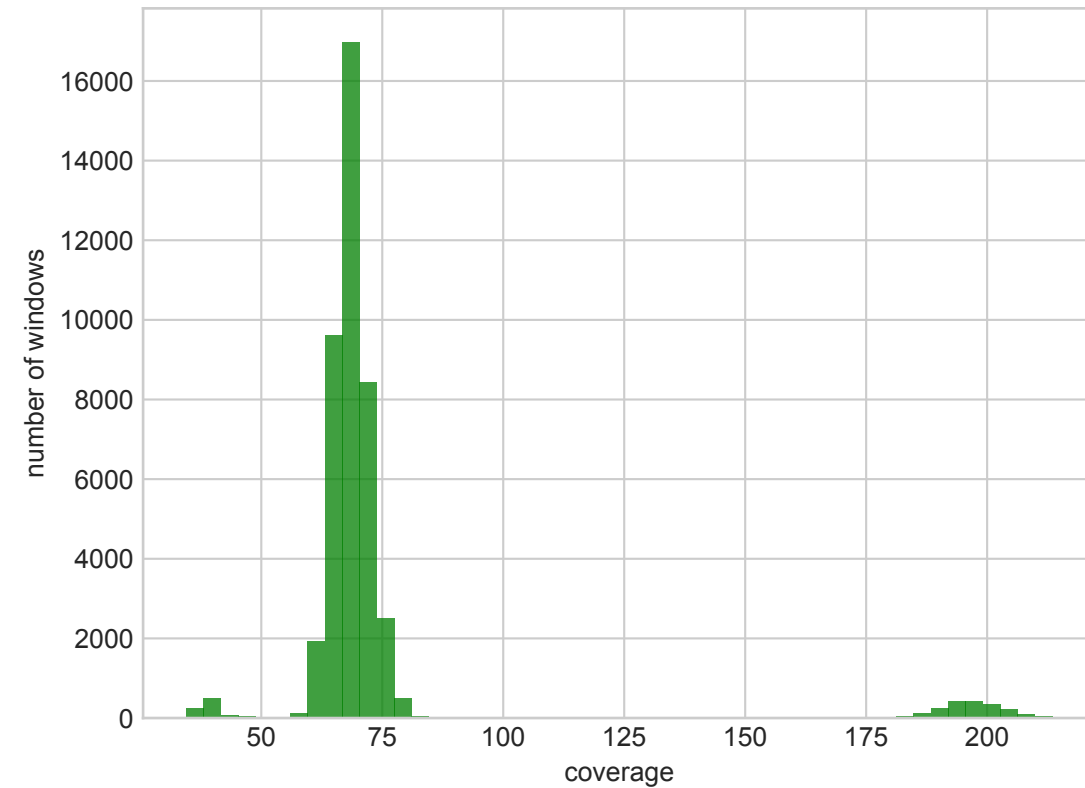


genome: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/reference_assemblies_task2/Bacillus_subtilis_NCIB_3610/GCA_002055965.1_ASM205596v1_genomic sample: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/bwa_fna/D300/Sample_L2B4-300_mapped_sort_NOdup_sort_merged_sort
iRep: 1.34 r^2: 0.95 avg. cov: 73.88 % windows: 100.00



genome: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/reference_assemblies_task2/Bacillus_subtilis_NCIB_3610/GCA_002055965.1_ASM205596v1_genomic sample: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/bwa_fna/D300/Sample_L2B4-300_mapped_sort_NOdup_sort_merged_sort
iRep: 1.34 raw iRep: 1.19 GC r^2: 0.05 GC bias: 6.68 avg. cov: 73.88 % windows: 100.00

