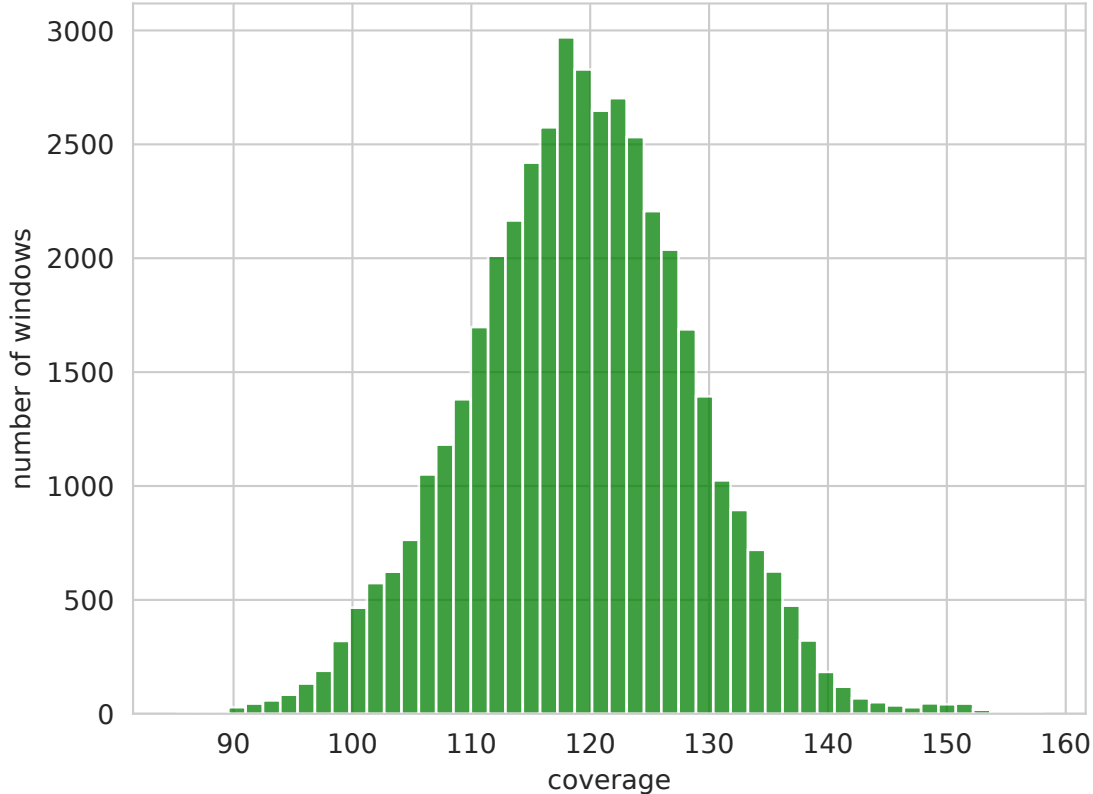


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0722/GCF_005937995_2_ASM593799v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0722-A
iRep: 1.25 r²: 0.98 avg. cov: 119.16 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0722/GCF_005937995_2_ASM593799v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0722-A
iRep: 1.25 raw iRep: 1.26 GC r^2 : 0.12 GC bias: -16.59 avg. cov: 119.16 % windows: 100.00

