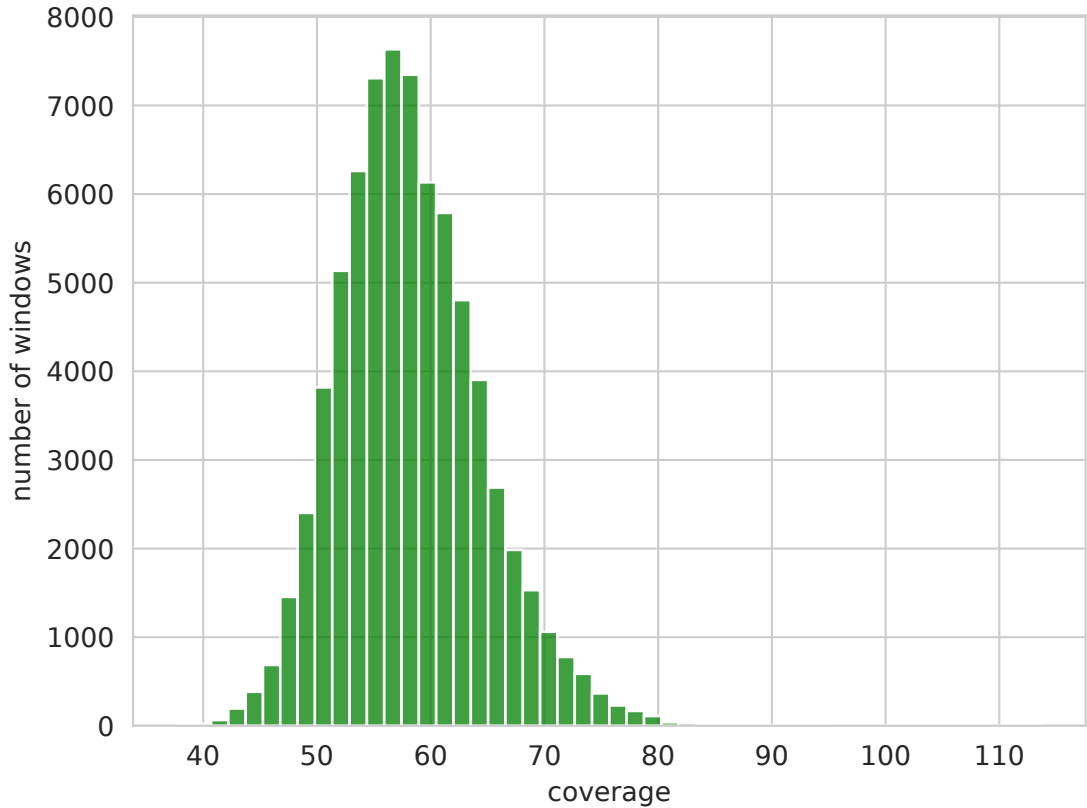


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0727/GCF\_005937885\_2\_ASM593788v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0727-B  
iRep: 1.30    r<sup>2</sup>: 0.99    avg. cov: 58.24    % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0727/GCF\_005937885\_2\_ASM593788v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0727-B  
iRep: 1.30 raw iRep: 1.37 GC  $r^2$ : 0.35 GC bias: -55.23 avg. cov: 58.24 % windows: 100.00

