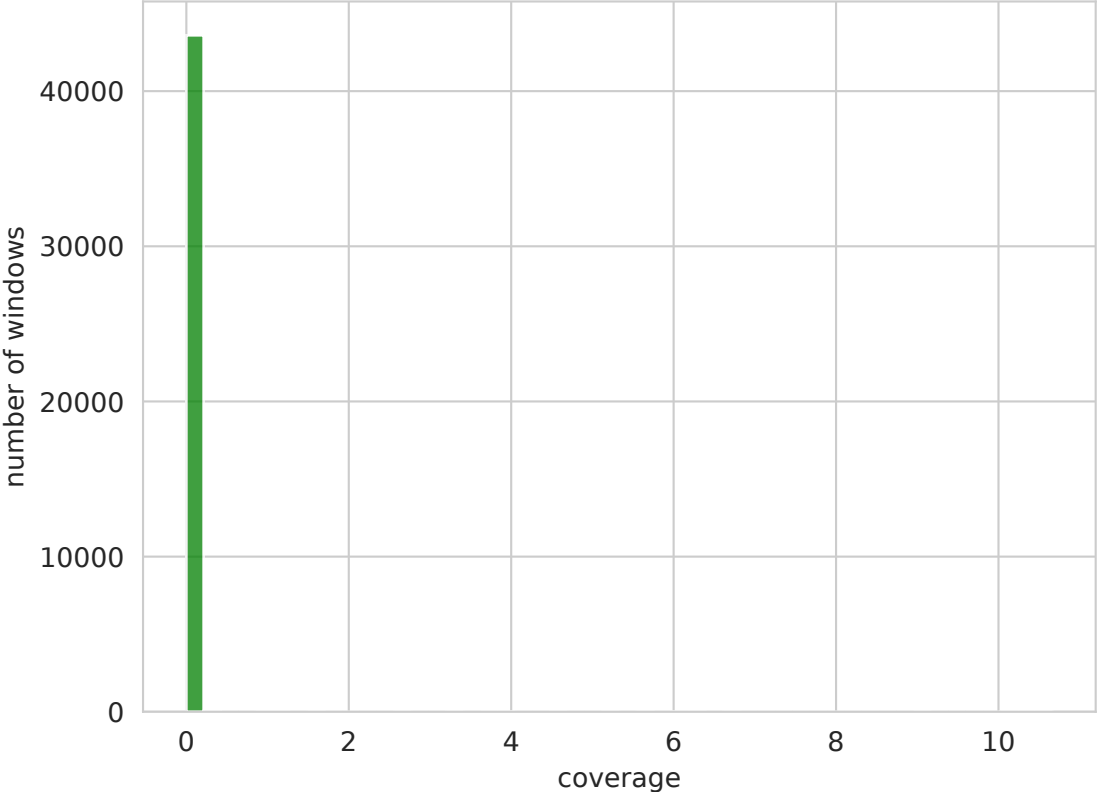


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0705/GCF\_005938025\_2\_ASM593802v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0705-A  
iRep: n/a    $r^2$ : 0.91   avg. cov: 0.01   % windows: 1.02



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0705/GCF\_005938025\_2\_ASM593802v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0705-A  
iRep: n/a raw iRep: 9.56 GC r<sup>2</sup>: 0.01 GC bias: -0.00 avg. cov: 0.01 % windows: 1.02

