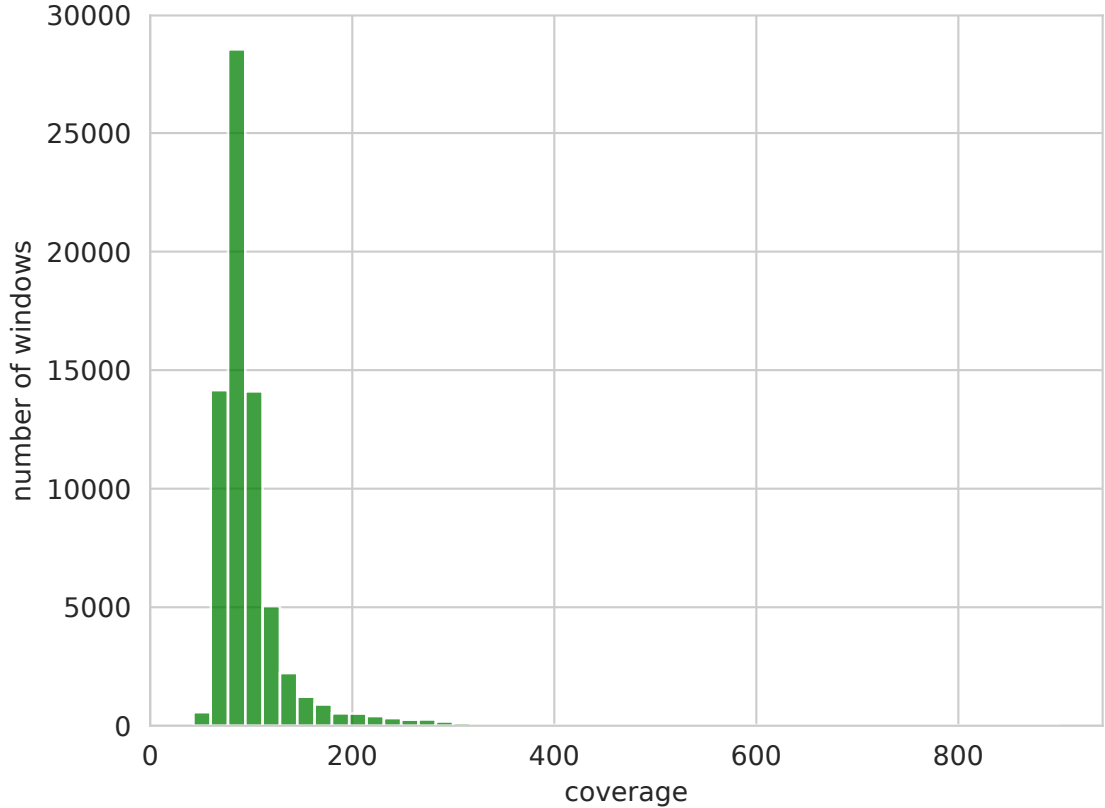
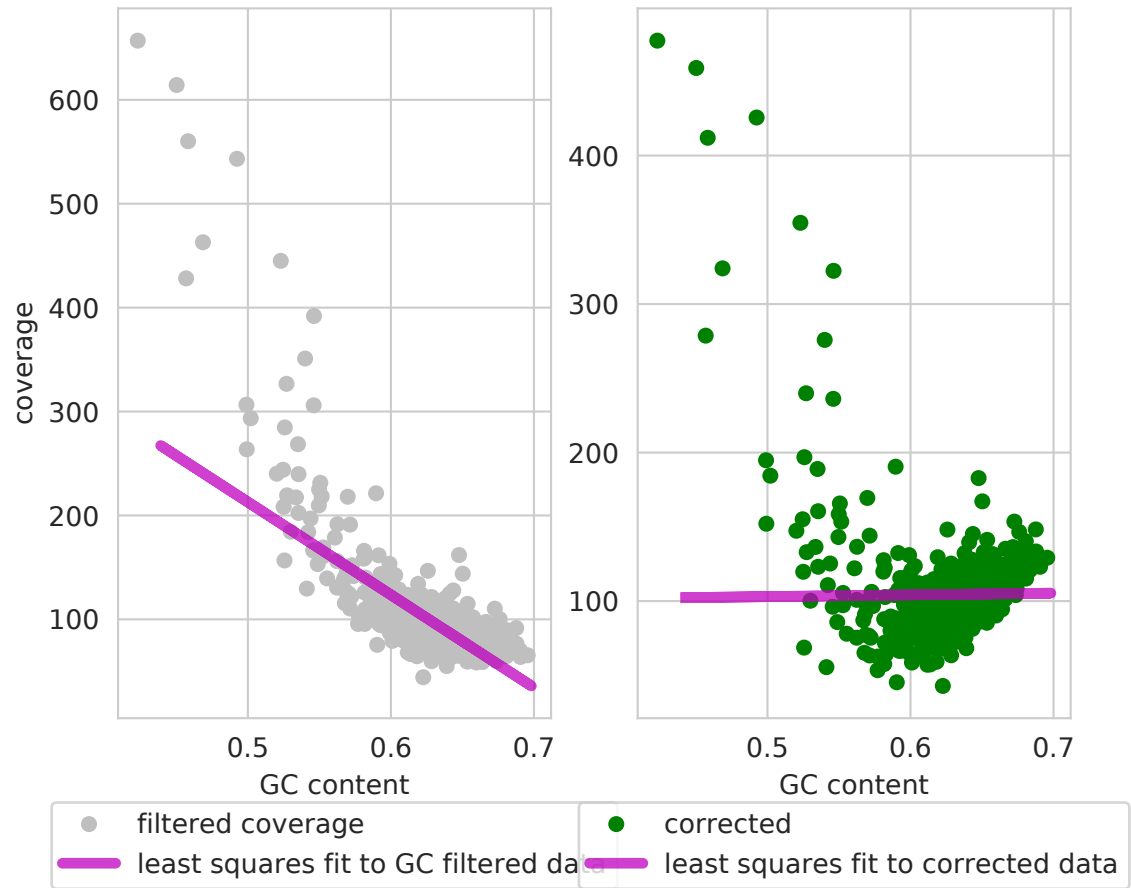


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/ATCC13985/GCF\_006322025\_2\_ASM632202v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/ATCC13985-B  
iRep: 1.73   r<sup>2</sup>: 0.98   avg. cov: 102.67   % windows: 99.90



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/ATCC13985/GCF\_006322025\_2\_ASM632202v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/ATCC13985-B  
iRep: 1.73 raw iRep: 2.02 GC r^2: 0.64 GC bias: -569.23 avg. cov: 102.67 % windows: 99.90



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/ATCC13985/GCF\_006322025\_2\_ASM632202v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/ATCC13985-B  
iRep: 1.73 un-filtered iRep: 1.73  $r^2$ : 0.98 avg. cov: 102.67 % windows: 99.90

