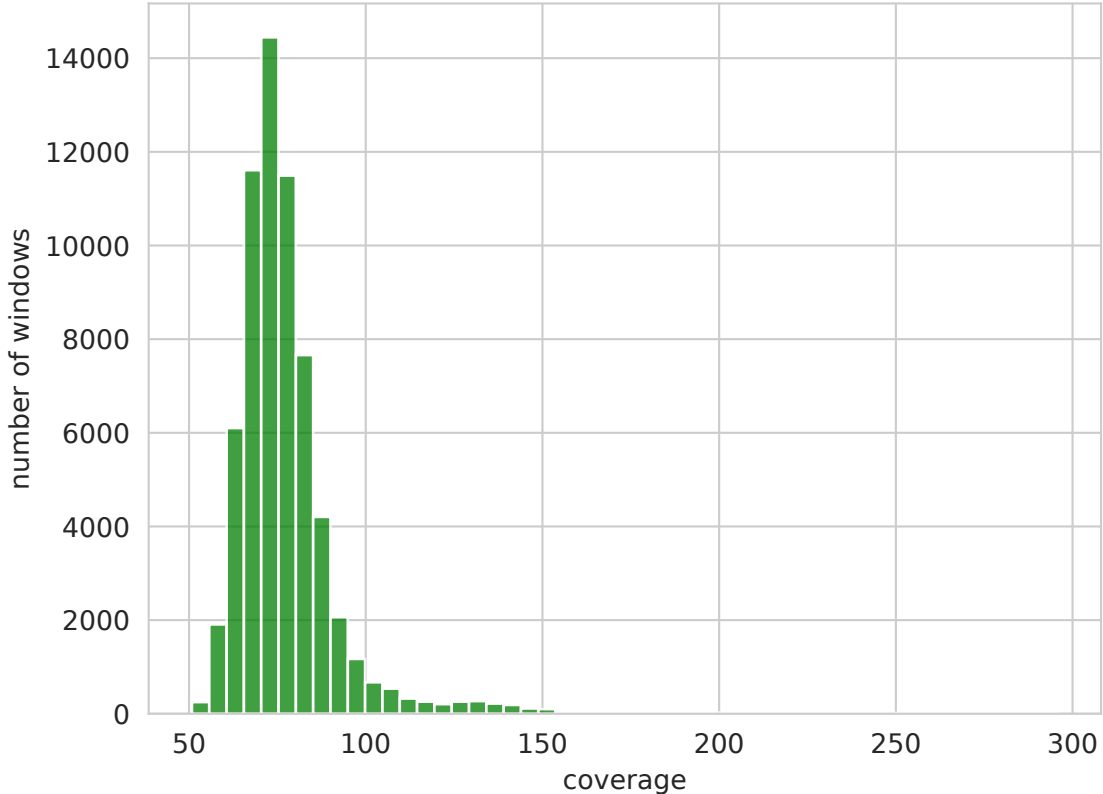
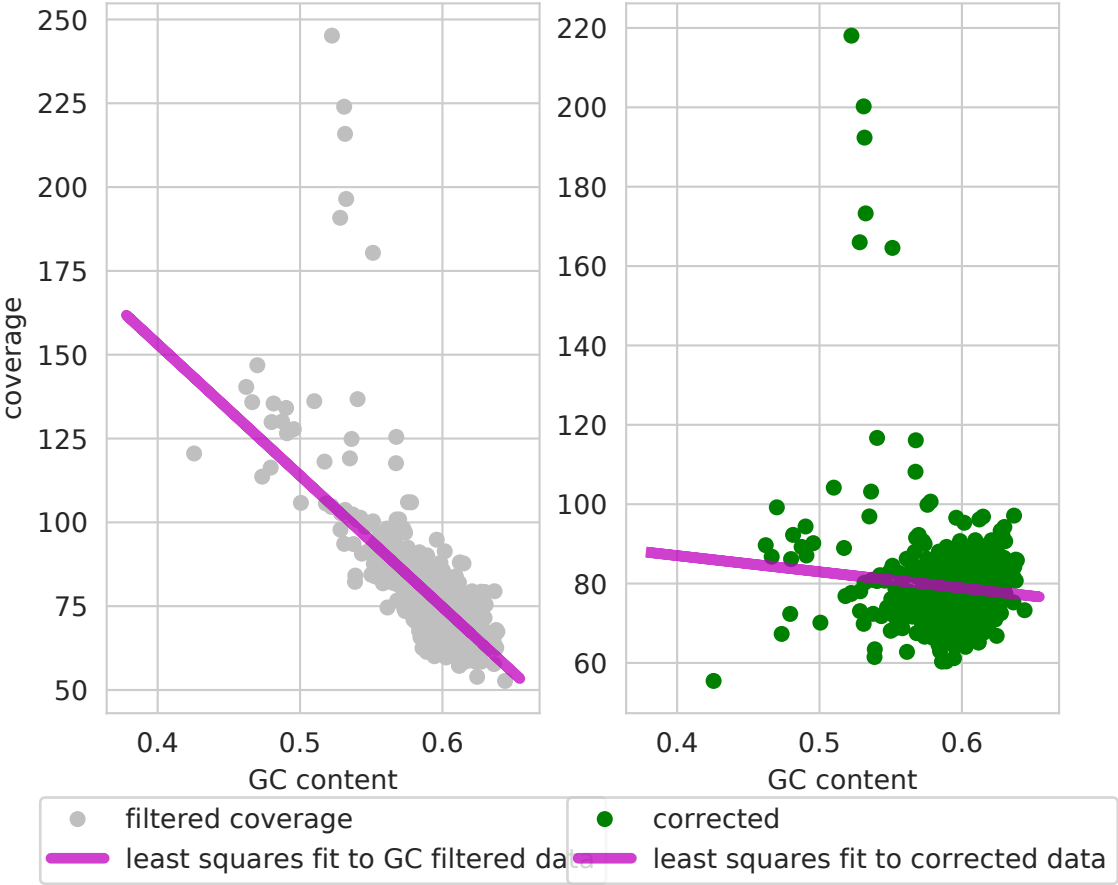


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/ATCC43928/GCF\_005938665.1\_ASM593866v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/ATCC43928-C  
iRep: 1.31   r<sup>2</sup>: 0.98   avg. cov: 78.05   % windows: 100.00





genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/ATCC43928/GCF\_005938665.1\_ASM593866v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/ATCC43928-C  
iRep: 1.31 un-filtered iRep: 1.31  $r^2$ : 0.98 avg. cov: 78.05 % windows: 100.00

