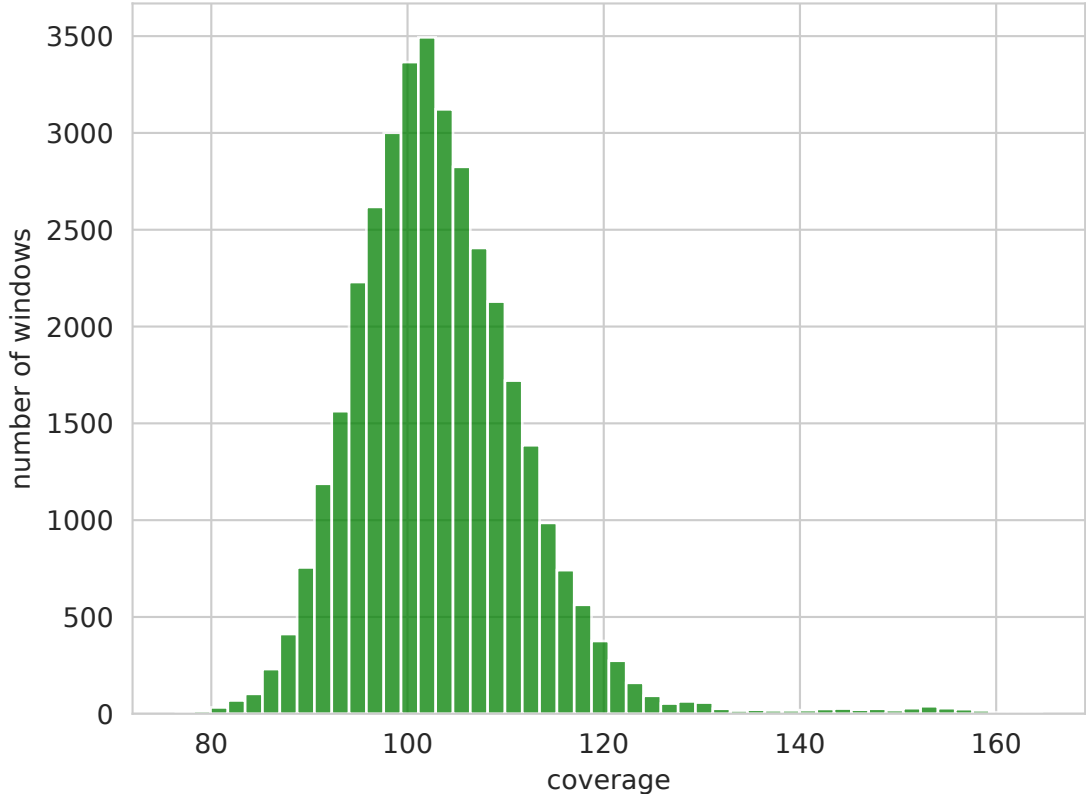
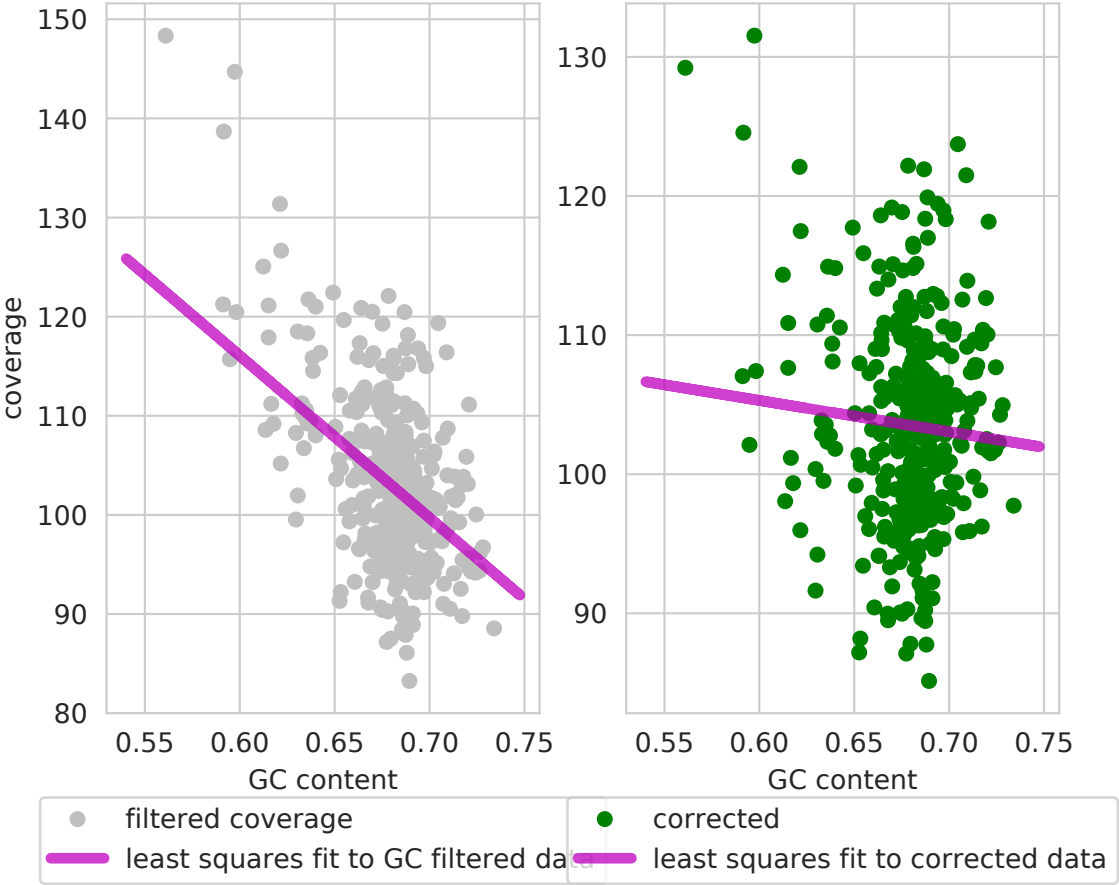


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0702/GCF\_005937985\_2\_ASM593798v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0702-F  
iRep: 1.23   r<sup>2</sup>: 0.99   avg. cov: 103.34   % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0702/GCF\_005937985\_2\_ASM593798v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0702-F  
iRep: 1.23 raw iRep: 1.26 GC r<sup>2</sup>: 0.24 GC bias: -38.83 avg. cov: 103.34 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0702/GCF\_005937985\_2\_ASM593798v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0702-F  
iRep: 1.23 un-filtered iRep: 1.23 r^2: 0.99 avg. cov: 103.34 % windows: 100.00

