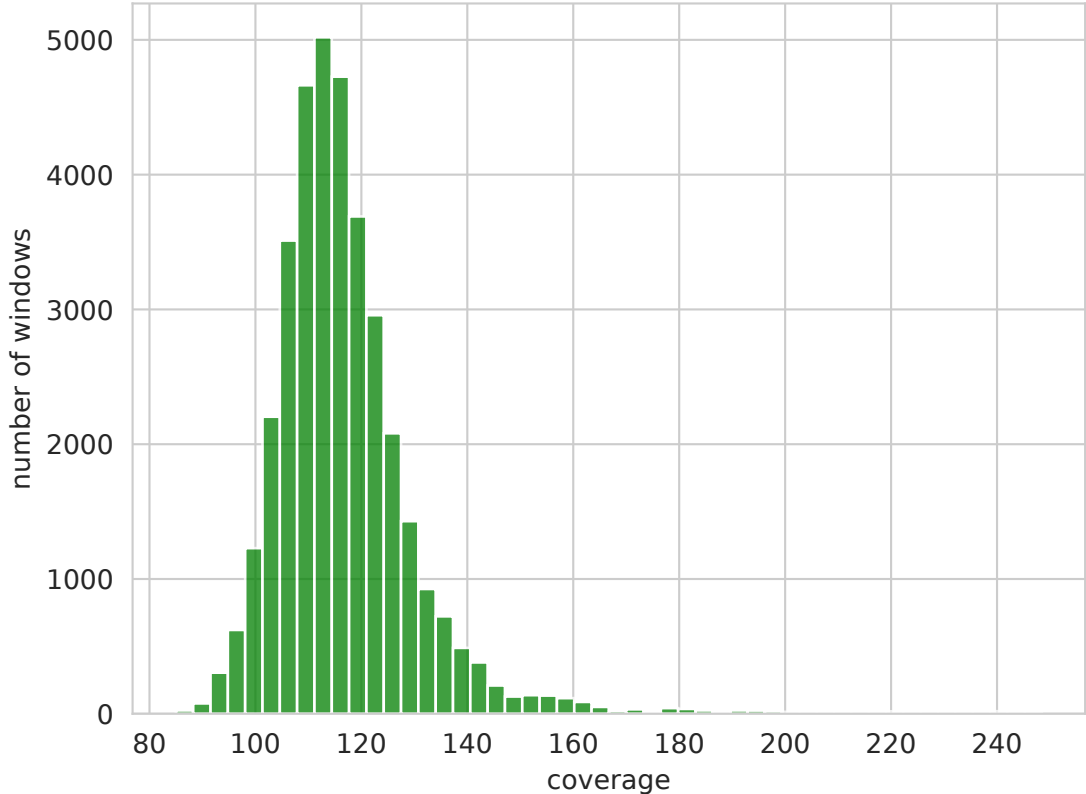
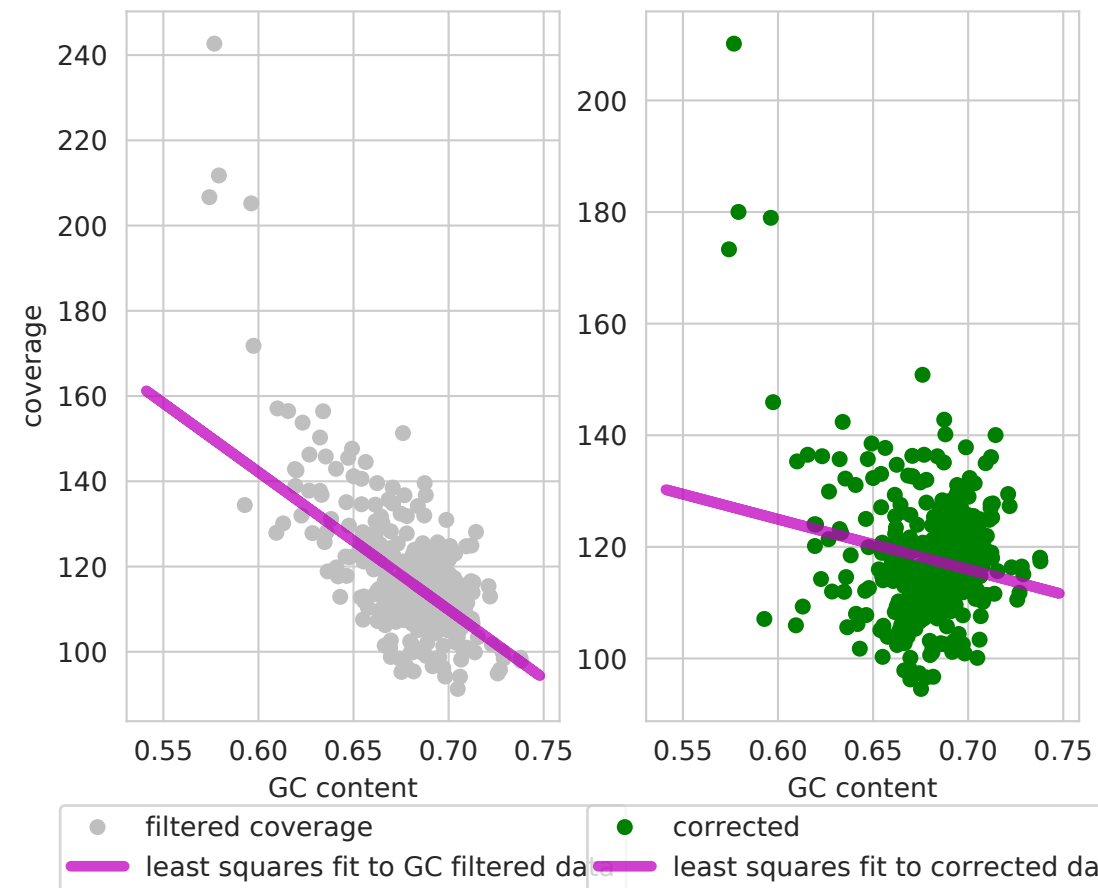


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0702/GCF\_005937985.1\_ASM593798v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0702-B  
iRep: 1.26   r<sup>2</sup>: 0.99   avg. cov: 117.17   % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0702/GCF\_005937985.1\_ASM593798v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0702-B  
iRep: 1.26 raw iRep: 1.32 GC  $r^2$ : 0.41 GC bias: -133.60 avg. cov: 117.17 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0702/GCF\_005937985.1\_ASM593798v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0702-B  
iRep: 1.26 un-filtered iRep: 1.26  $r^2$ : 0.99 avg. cov: 117.17 % windows: 100.00

