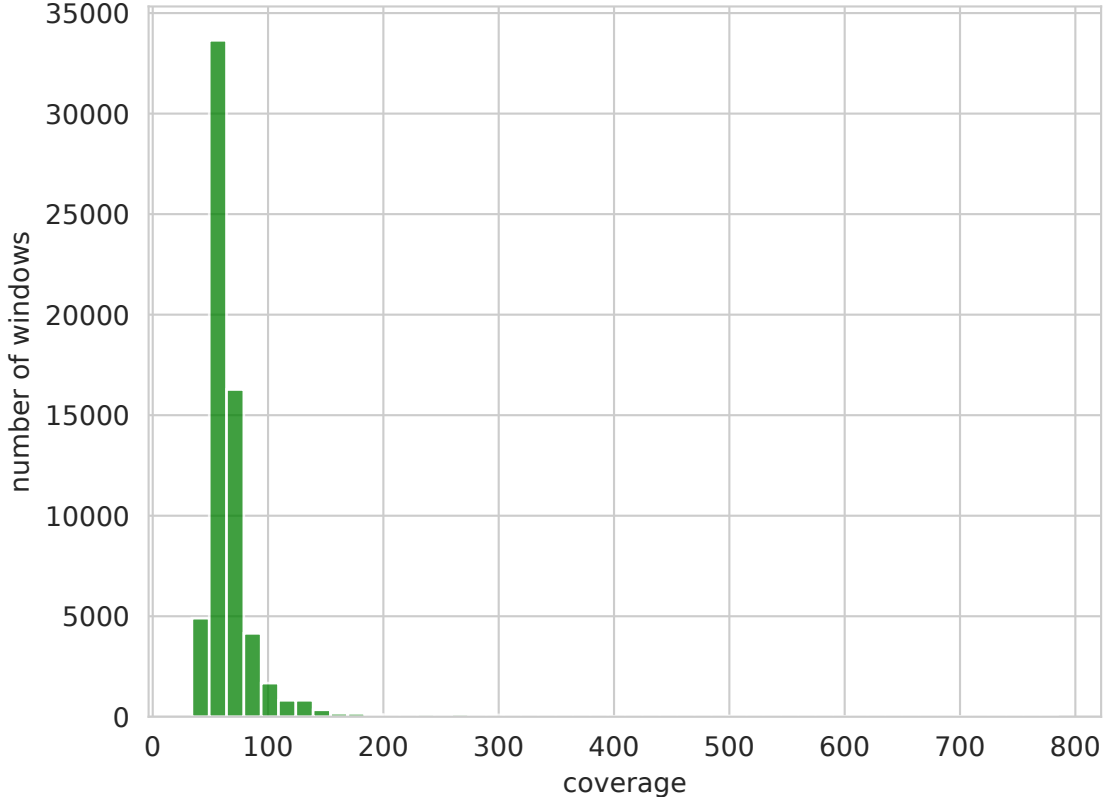
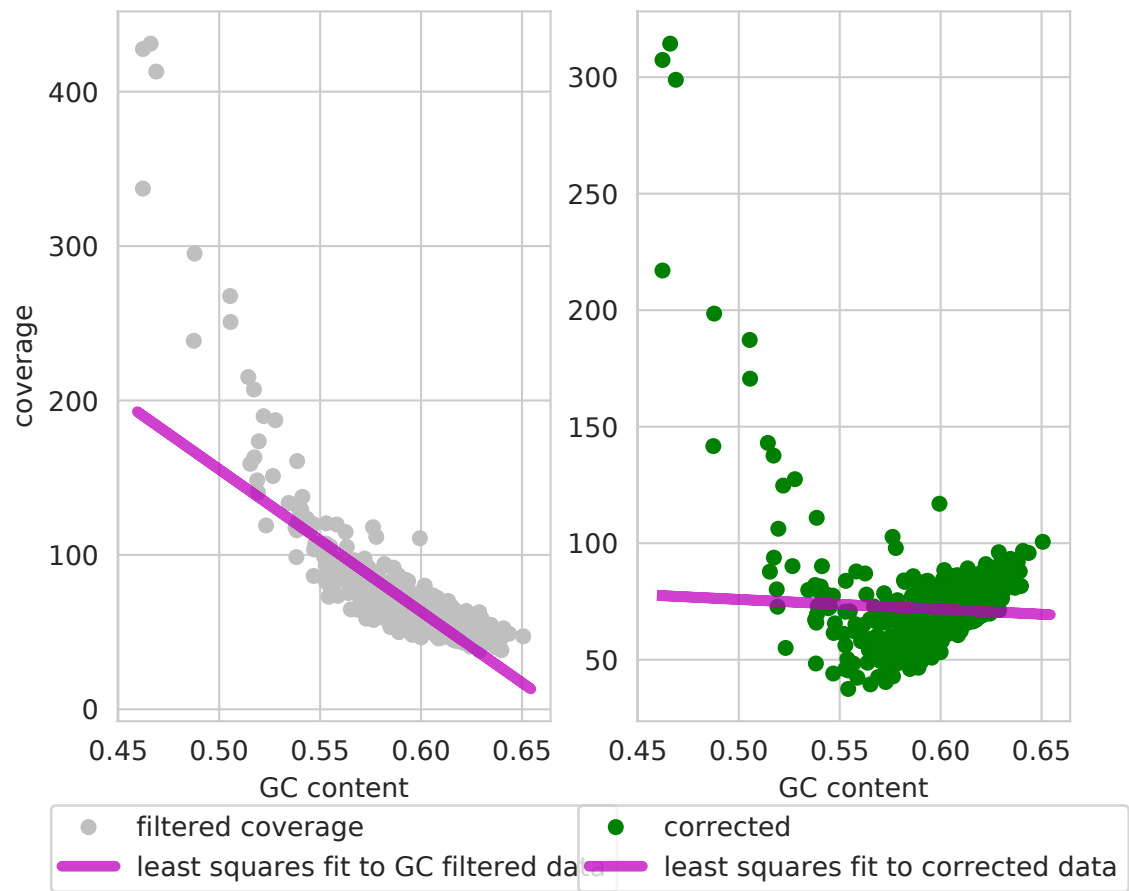


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/ATCC43928/GCF\_005938665.1\_ASM593866v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/ATCC43928-B  
iRep: 1.61     $r^2$ : 0.98    avg. cov: 72.03    % windows: 99.62



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-B  
iRep: 1.61 raw iRep: 1.91 GC  $r^2$ : 0.70 GC bias: -641.31 avg. cov: 72.03 % windows: 99.62



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-B  
iRep: 1.61 un-filtered iRep: 1.61  $r^2$ : 0.98 avg. cov: 72.03 % windows: 99.62

