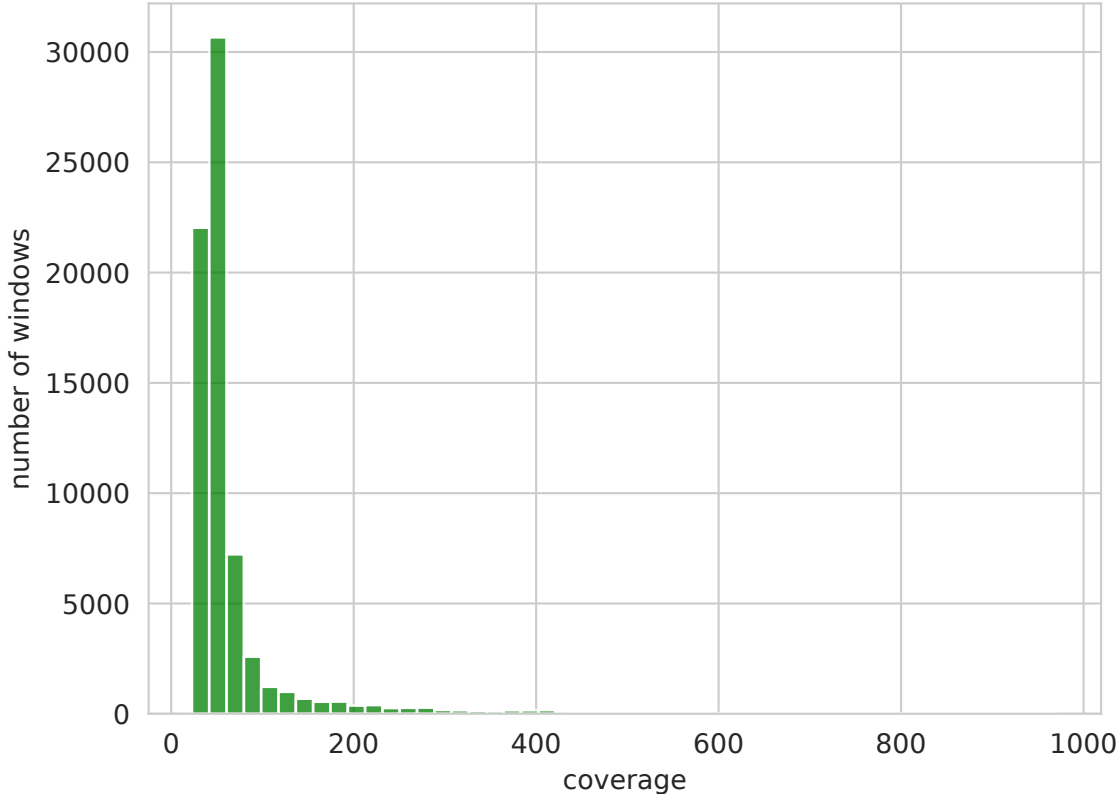
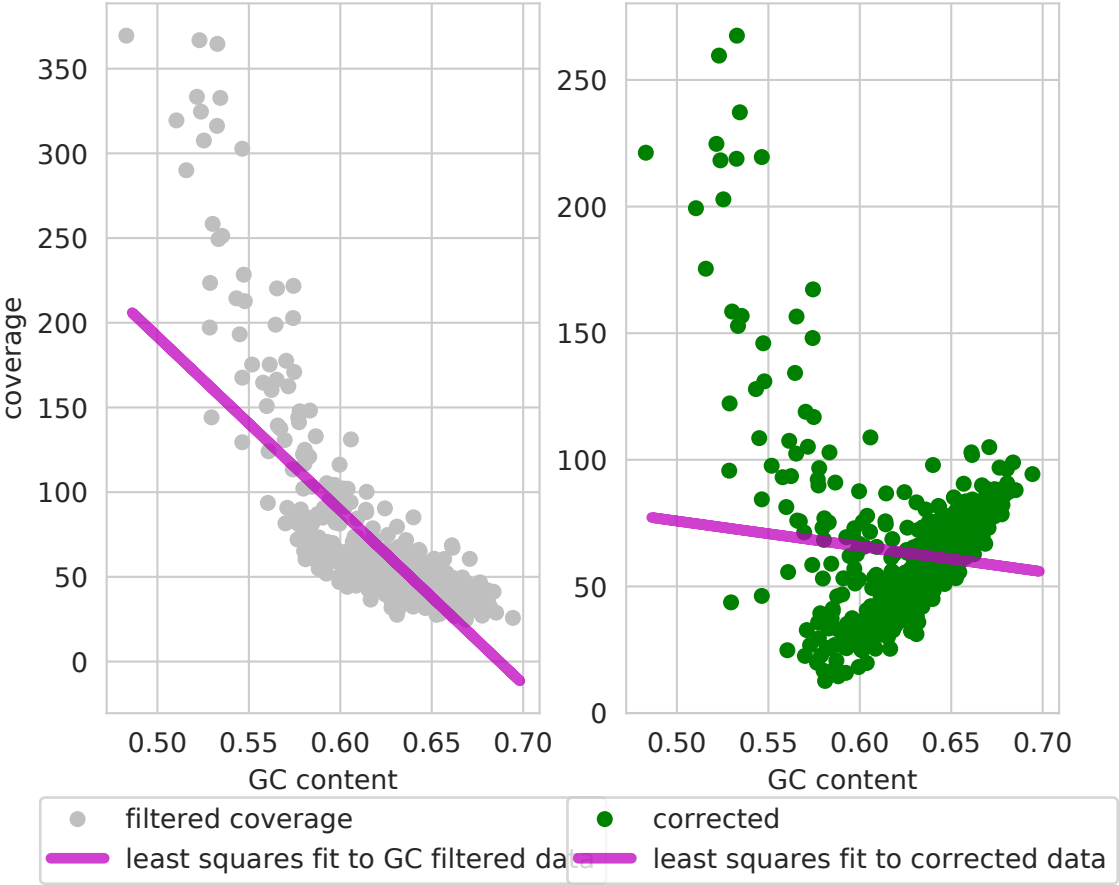


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/ATCC13985/GCF\_006322025.1\_ASM632202v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/ATCC13985-D  
iRep: 2.69    $r^2$ : 0.98   avg. cov: 68.85   % windows: 98.31



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/ATCC13985/GCF\_006322025.1\_ASM632202v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/ATCC13985-D  
iRep: 2.69 raw iRep: 2.98 GC  $r^2$ : 0.64 GC bias: -651.99 avg. cov: 68.85 % windows: 98.31



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/ATCC13985/GCF\_006322025.1\_ASM632202v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/ATCC13985-D  
iRep: 2.69 un-filtered iRep: 2.69  $r^2$ : 0.98 avg. cov: 68.85 % windows: 98.31

