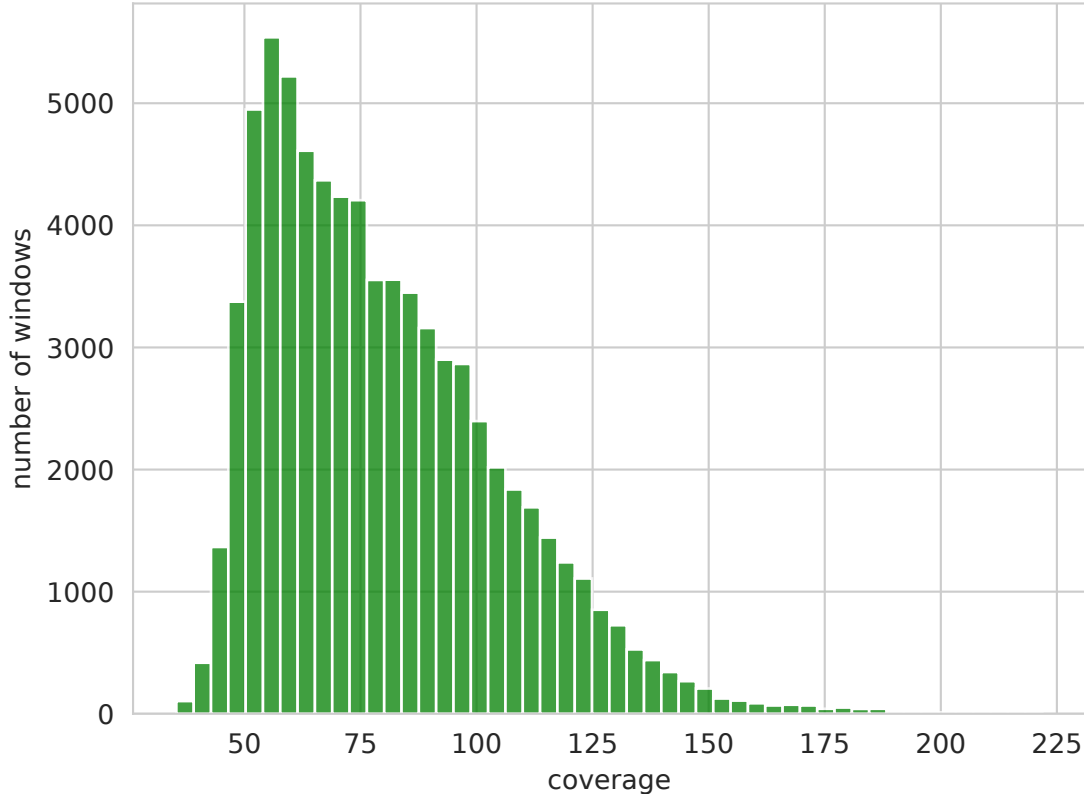
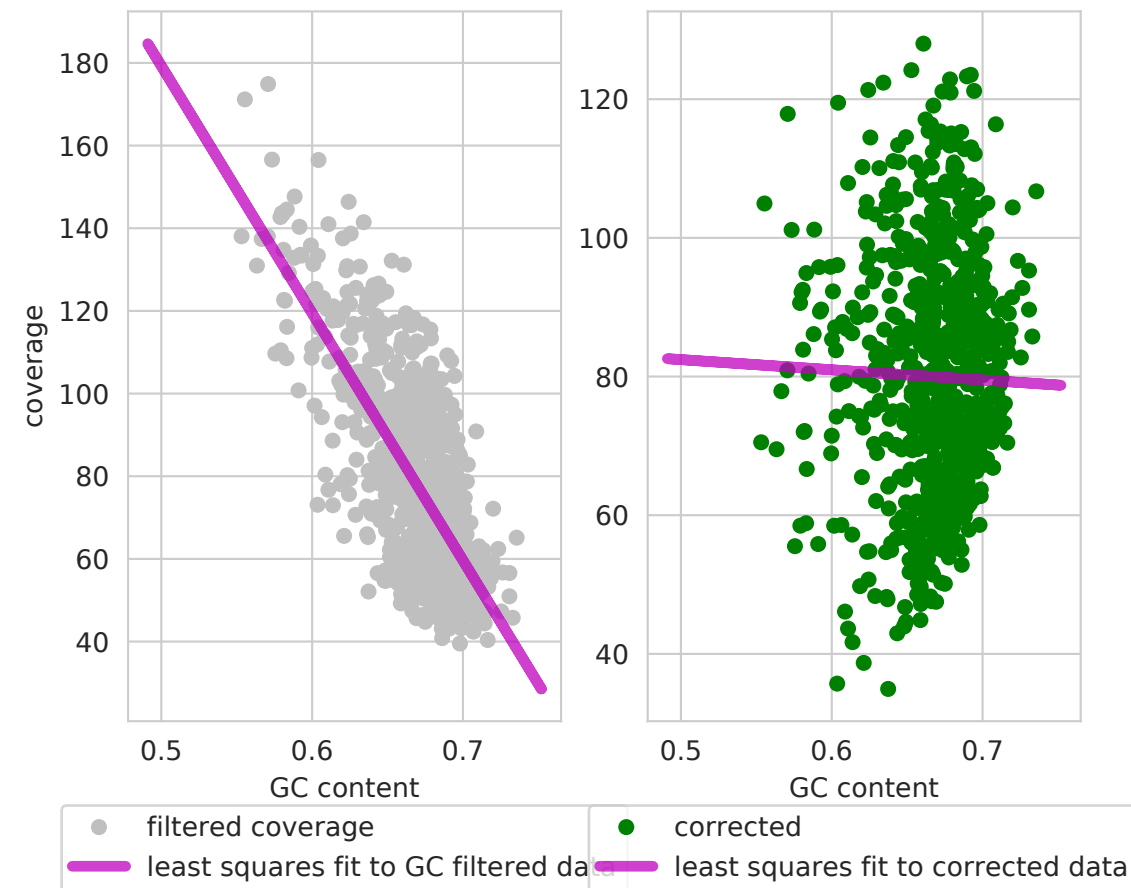


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0801/GCF\_005937855\_2\_ASM593785v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0801-B  
iRep: 1.98     $r^2$ : 0.99    avg. cov: 79.92    % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0801/GCF\_005937855\_2\_ASM593785v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0801-B  
iRep: 1.98 raw iRep: 2.65 GC r<sup>2</sup>: 0.55 GC bias: -327.90 avg. cov: 79.92 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0801/GCF\_005937855\_2\_ASM593785v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0801-B  
iRep: 1.98 un-filtered iRep: 1.98  $r^2$ : 0.99 avg. cov: 79.92 % windows: 100.00

