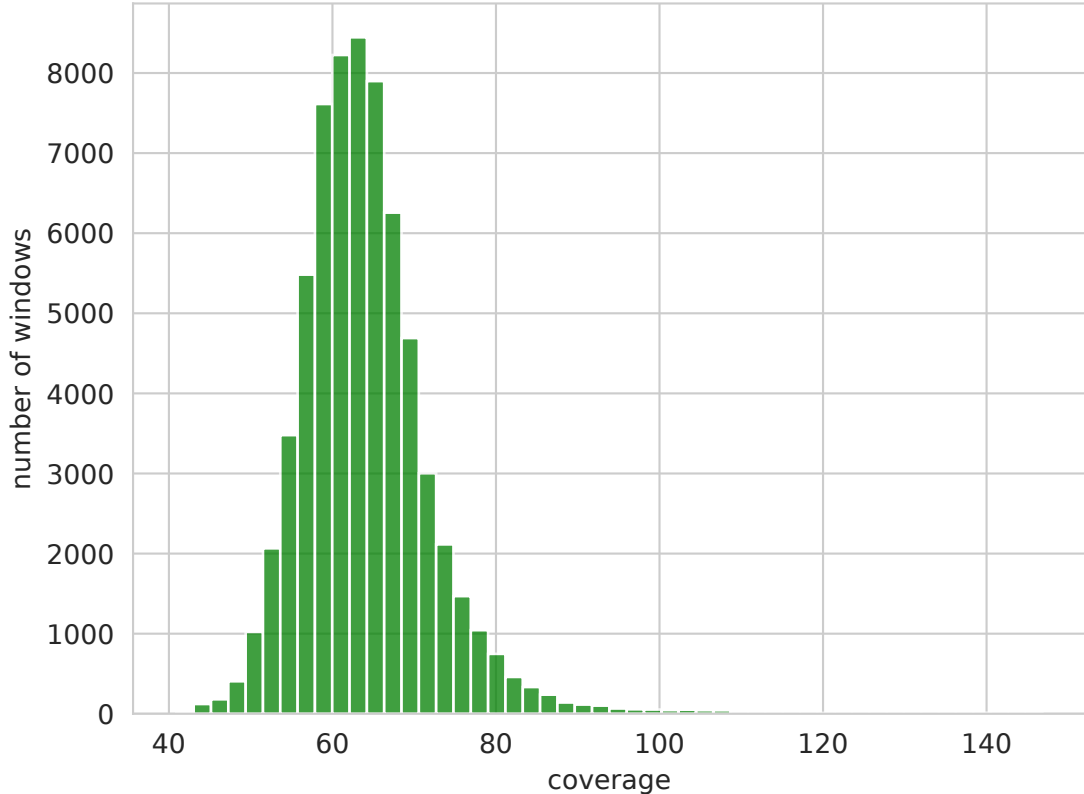
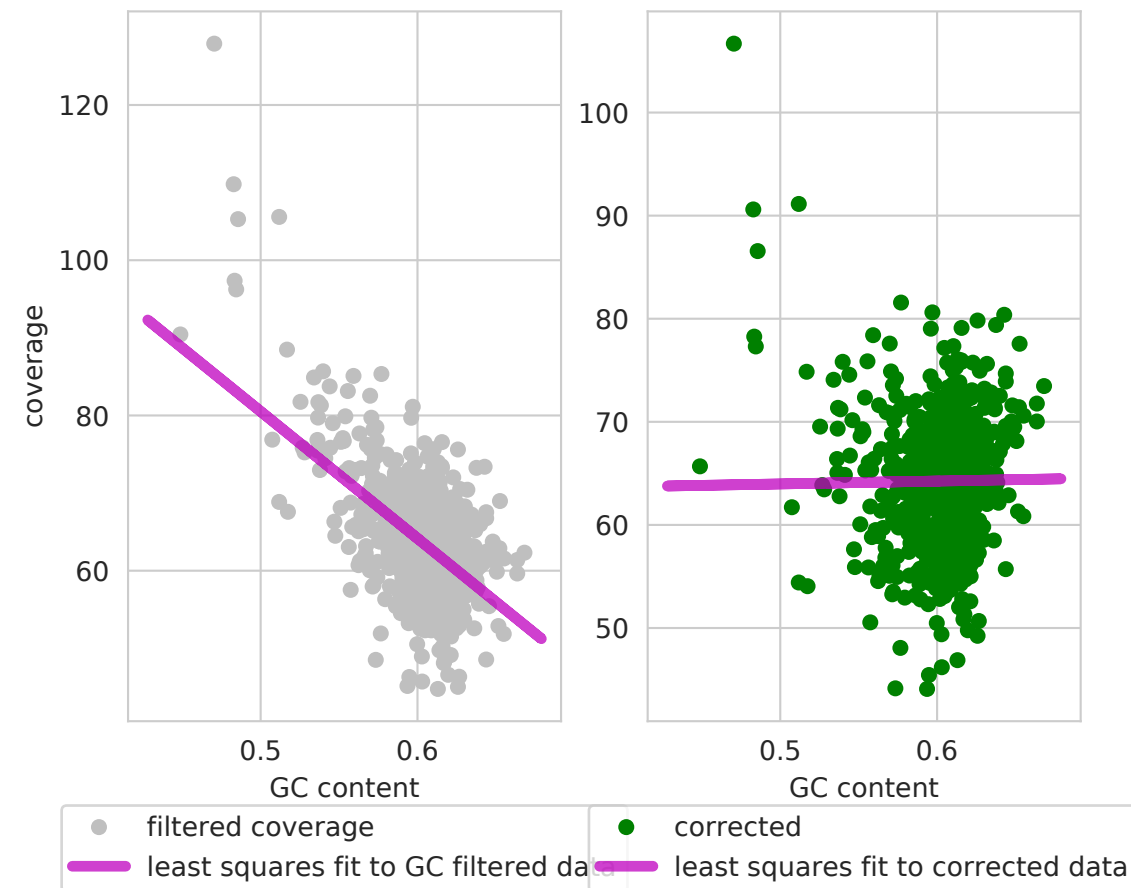


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0710/GCF\_005938045\_2\_ASM593804v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0710-B  
iRep: 1.35   r<sup>2</sup>: 0.99   avg. cov: 64.14   % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0710/GCF\_005938045\_2\_ASM593804v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0710-B  
iRep: 1.35 raw iRep: 1.39 GC  $r^2$ : 0.30 GC bias: -49.67 avg. cov: 64.14 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0710/GCF\_005938045\_2\_ASM593804v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0710-B  
iRep: 1.35 un-filtered iRep: 1.35  $r^2$ : 0.99 avg. cov: 64.14 % windows: 100.00

