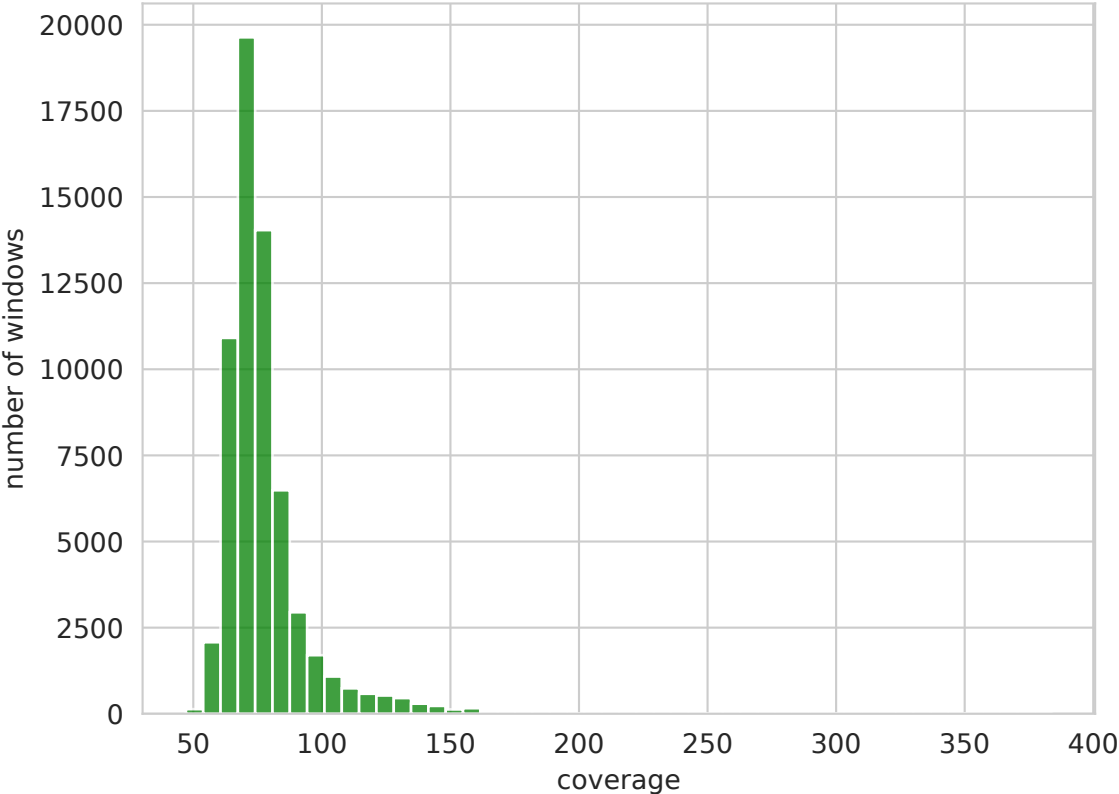
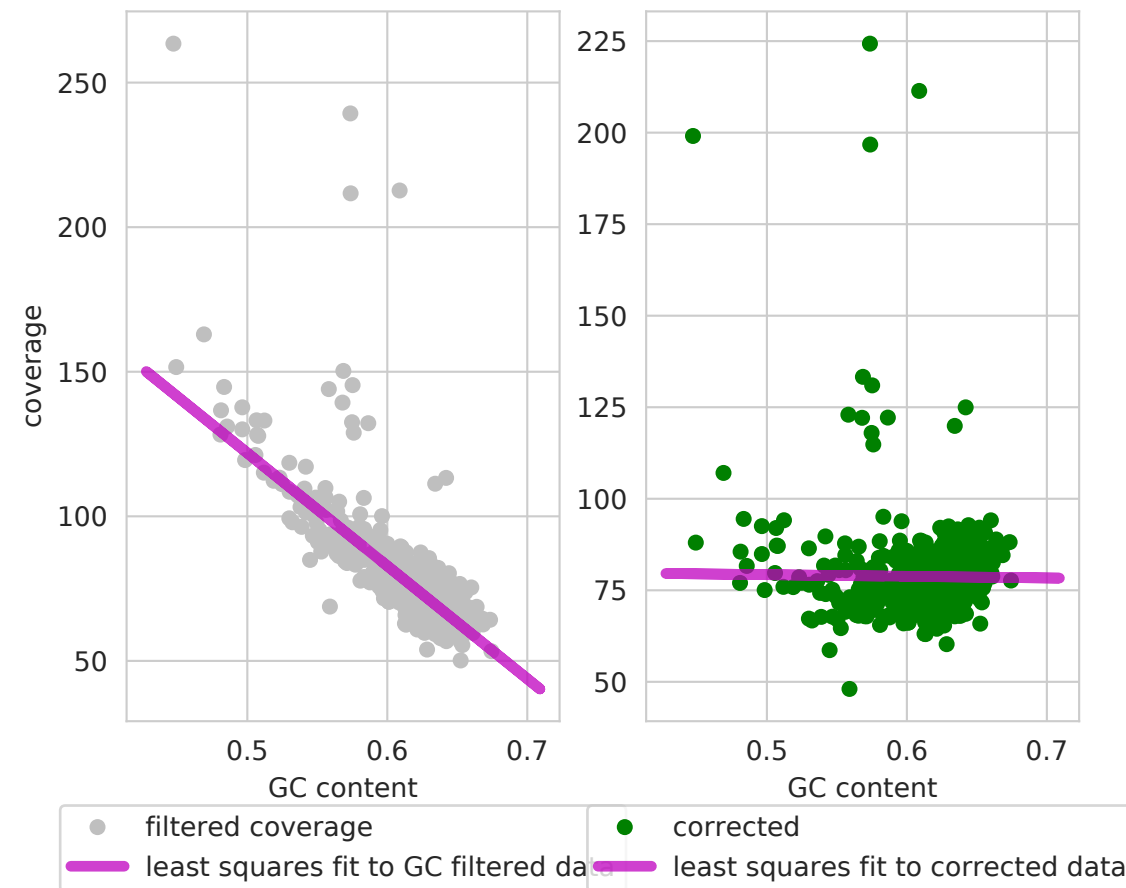


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0802/GCF\_005937845.1\_ASM593784v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0802-B  
iRep: 1.28     $r^2$ : 0.98    avg. cov: 78.15    % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0802/GCF\_005937845.1\_ASM593784v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0802-B  
iRep: 1.28 raw iRep: 1.55 GC r<sup>2</sup>: 0.73 GC bias: -286.58 avg. cov: 78.15 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0802/GCF\_005937845.1\_ASM593784v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0802-B  
iRep: 1.28 un-filtered iRep: 1.28  $r^2$ : 0.98 avg. cov: 78.15 % windows: 100.00

