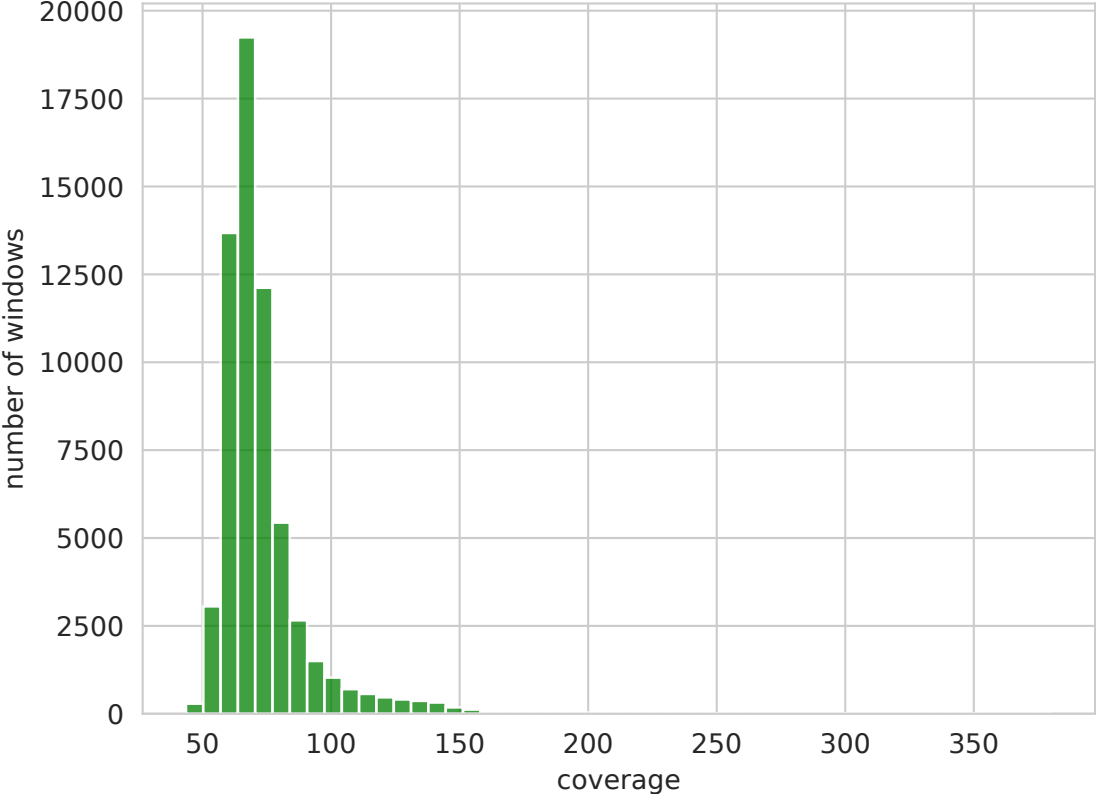
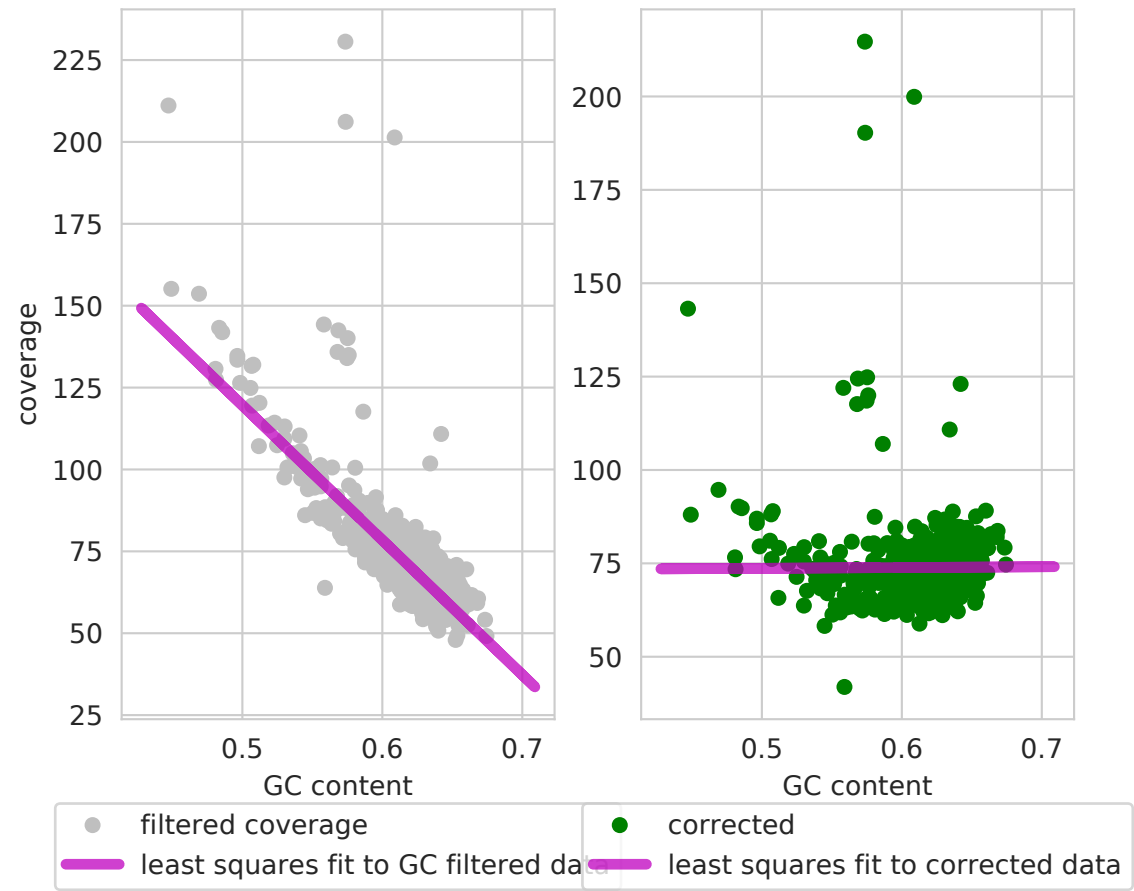


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0802/GCF\_005937845.1\_ASM593784v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0802-C  
iRep: 1.28     $r^2$ : 0.98    avg. cov: 73.37    % windows: 100.00



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-C  
iRep: 1.28 raw iRep: 1.61 GC r<sup>2</sup>: 0.78 GC bias: -319.09 avg. cov: 73.37 % 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-C  
iRep: 1.28 un-filtered iRep: 1.28  $r^2$ : 0.98 avg. cov: 73.37 % windows: 100.00

