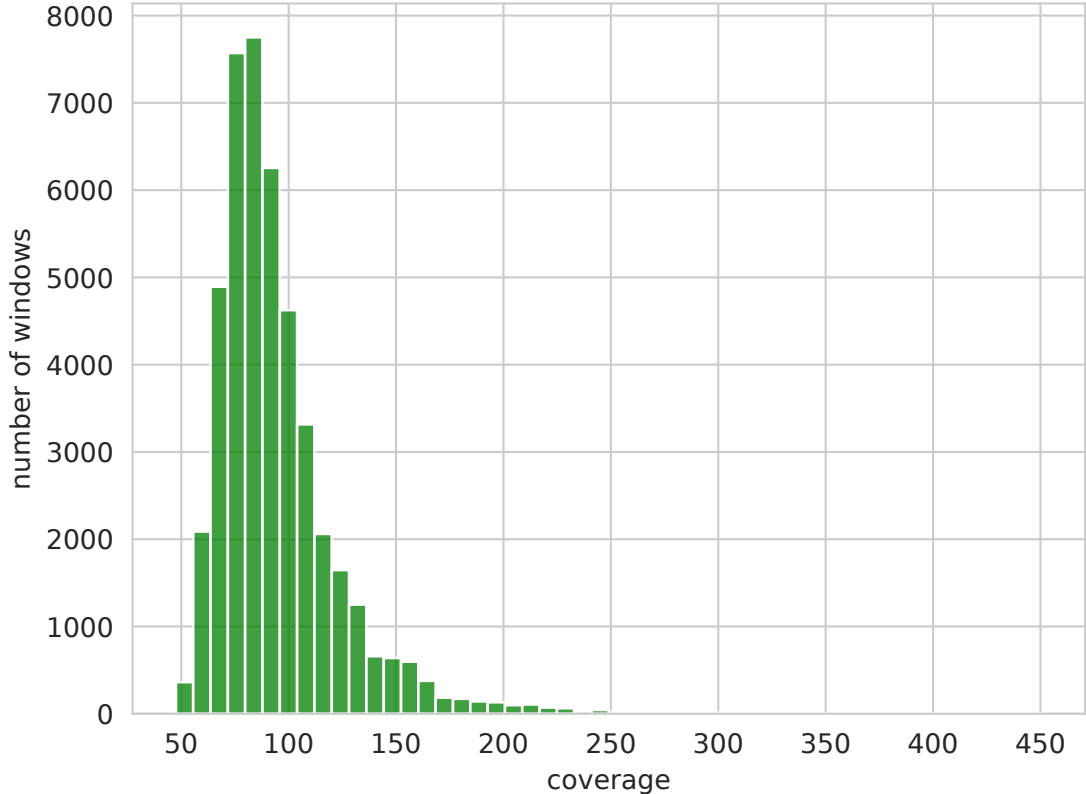
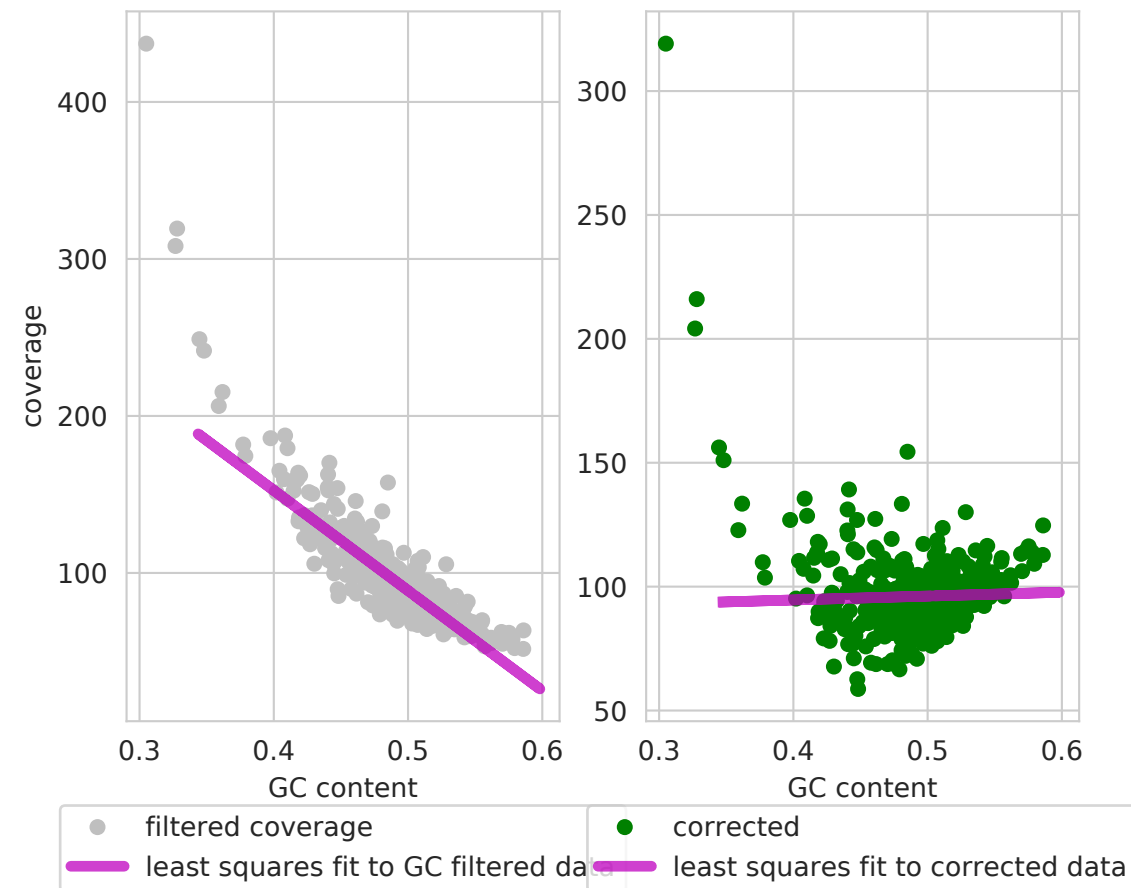


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0713/GCF_005937895.1_ASM593789v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0713-A
iRep: 1.42 r²: 0.98 avg. cov: 95.16 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0713/GCF_005937895.1_ASM593789v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0713-A
iRep: 1.42 raw iRep: 2.12 GC r²: 0.79 GC bias: -505.70 avg. cov: 95.16 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0713/GCF_005937895.1_ASM593789v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0713-A
iRep: 1.42 un-filtered iRep: 1.42 r^2 : 0.98 avg. cov: 95.16 % windows: 100.00

