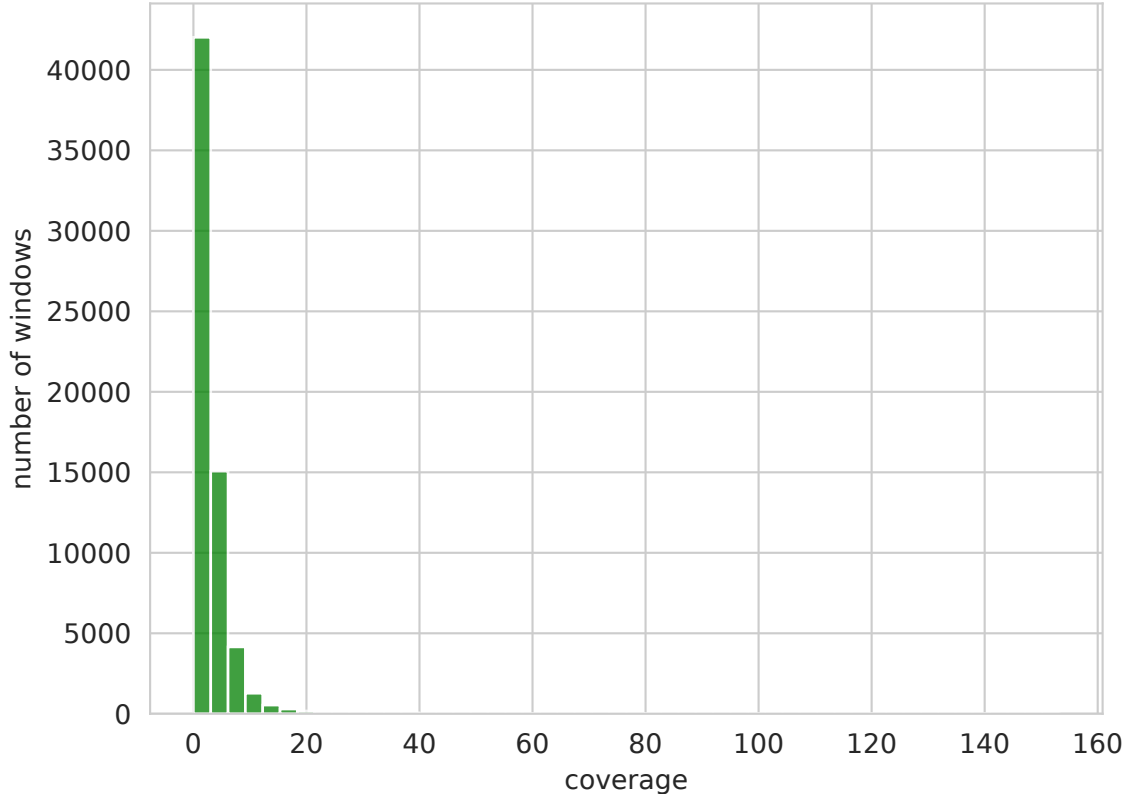
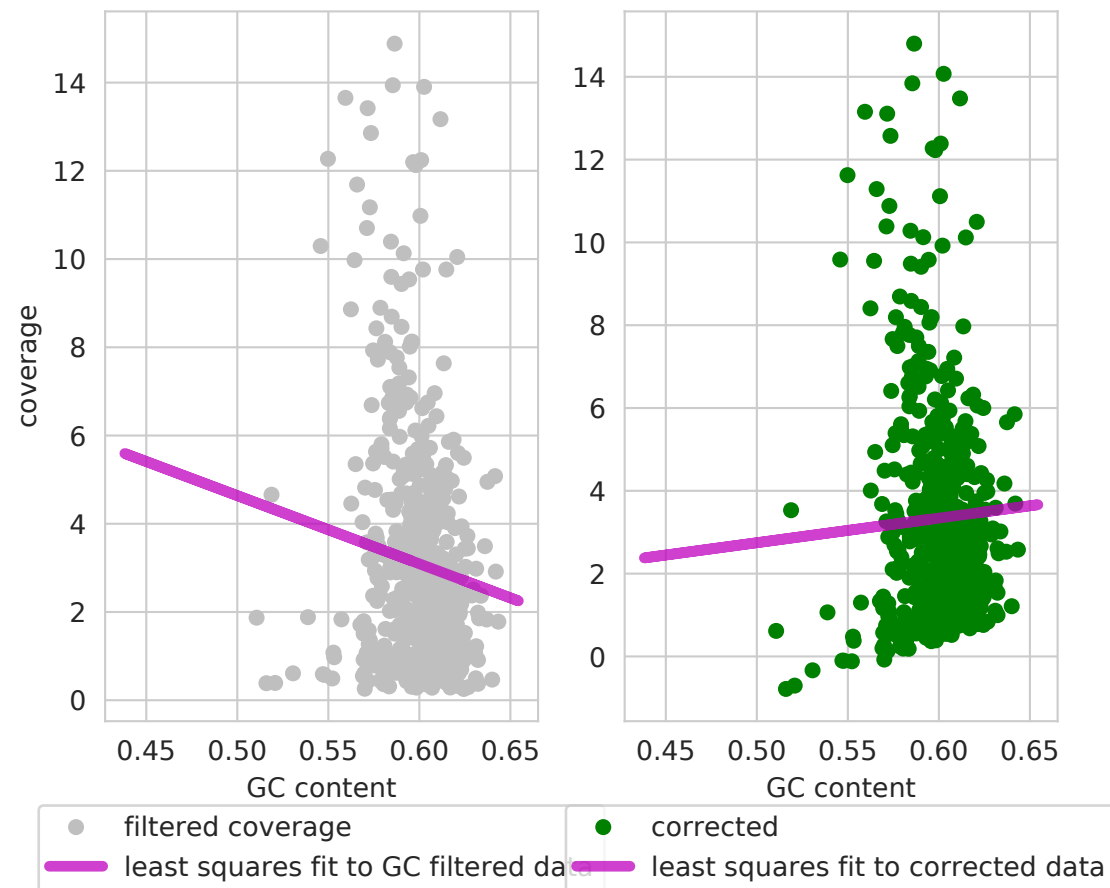


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/ATCC43928/GCF_005938665_2_ASM593866v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/ATCC43928-D
iRep: n/a r^2 : 0.98 avg. cov: 3.75 % windows: 79.91



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/ATCC43928/GCF_005938665_2_ASM593866v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/ATCC43928-D
iRep: n/a raw iRep: 13.36 GC r²: 0.02 GC bias: -0.25 avg. cov: 3.75 % windows: 79.91



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/ATCC43928/GCF_005938665_2_ASM593866v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/ATCC43928-D
iRep: n/a un-filtered iRep: 11.35 r^2 : 0.98 avg. cov: 3.75 % windows: 79.91

