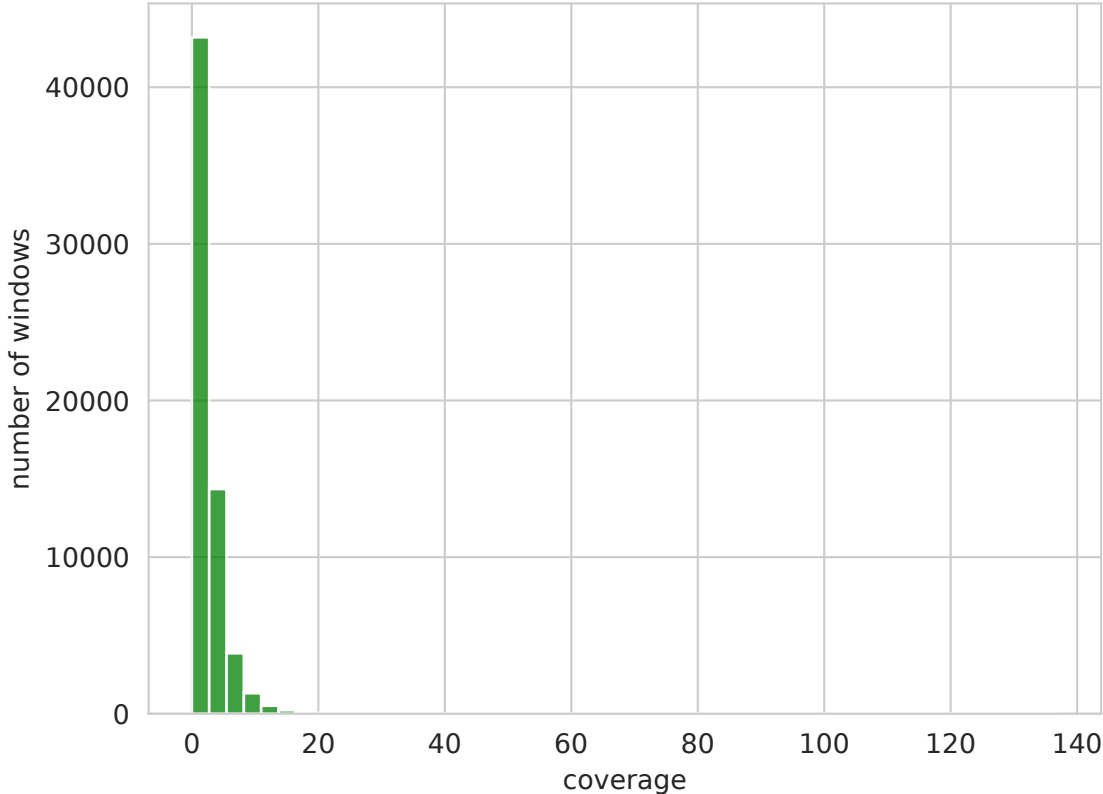
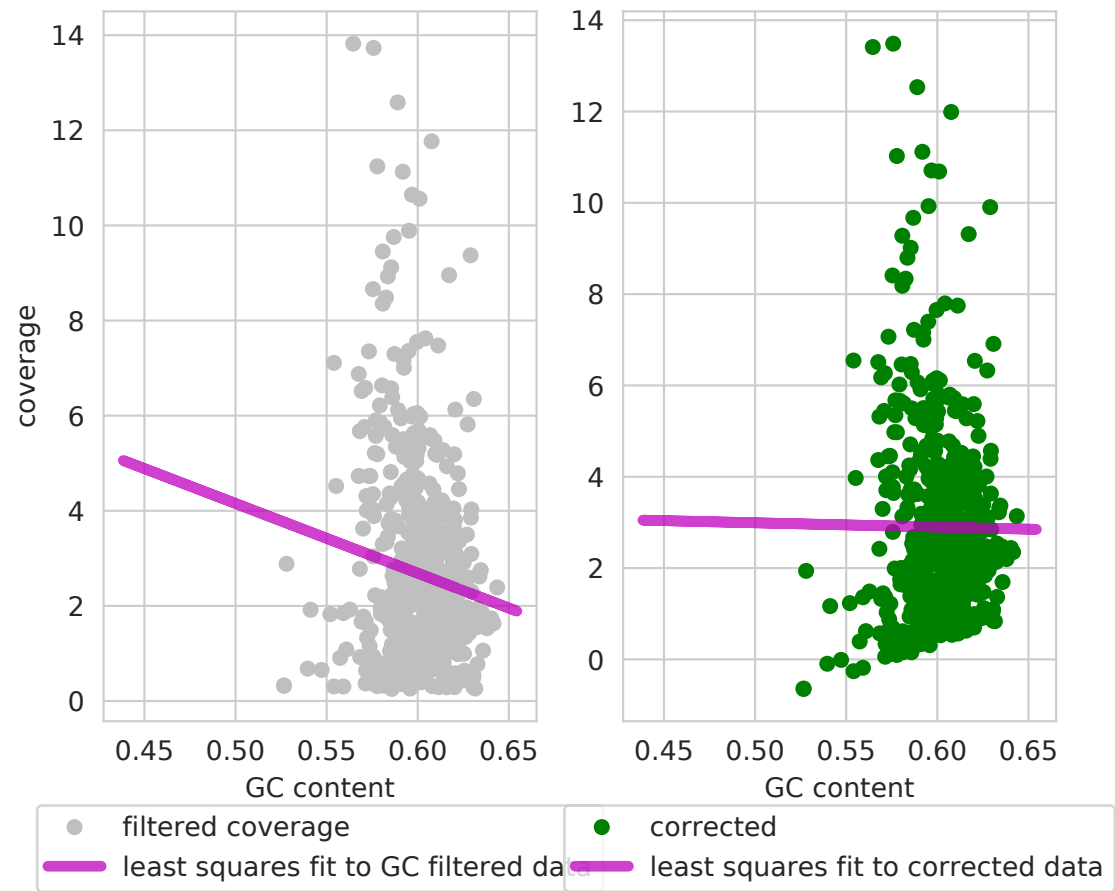


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/ATCC43928/GCF_005938665.1_ASM593866v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/ATCC43928-A
iRep: n/a r^2 : 0.97 avg. cov: 3.23 % windows: 80.15



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/ATCC43928/GCF_005938665.1_ASM593866v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/ATCC43928-A
iRep: n/a raw iRep: 12.88 GC r^2: 0.02 GC bias: -0.28 avg. cov: 3.23 % windows: 80.15



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/ATCC43928/GCF_005938665.1_ASM593866v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/ATCC43928-A
iRep: n/a un-filtered iRep: 11.16 r^2 : 0.97 avg. cov: 3.23 % windows: 80.15

