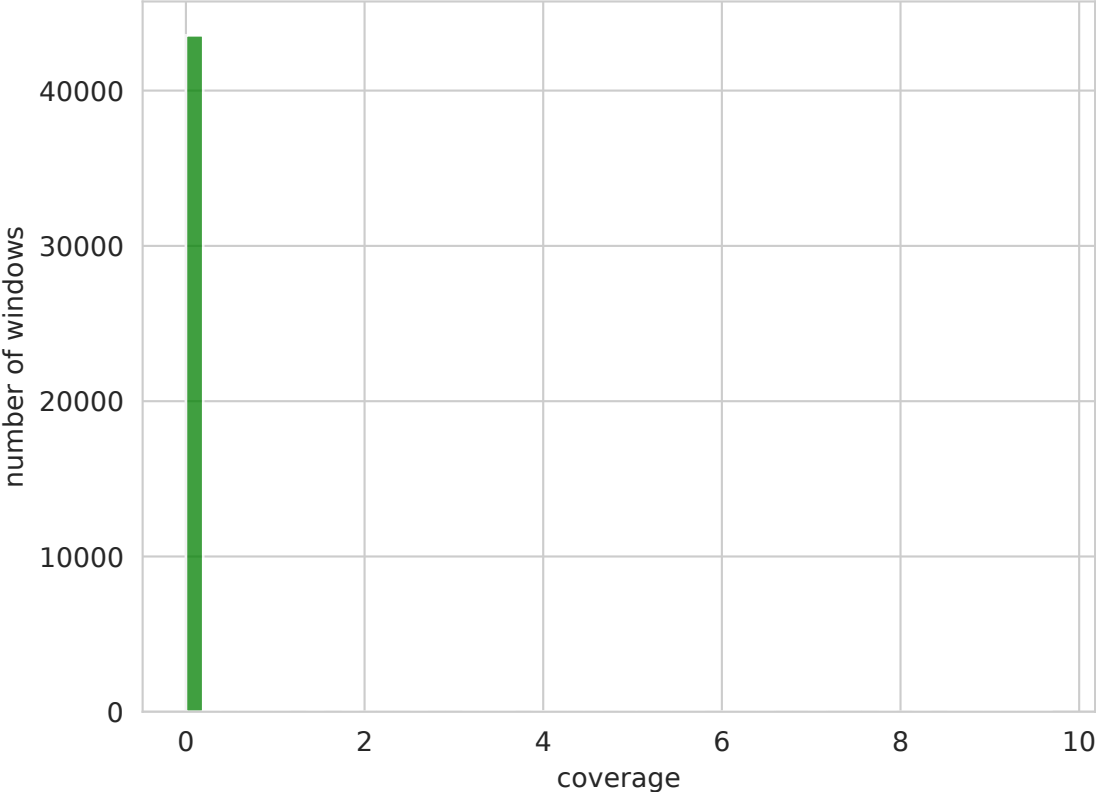
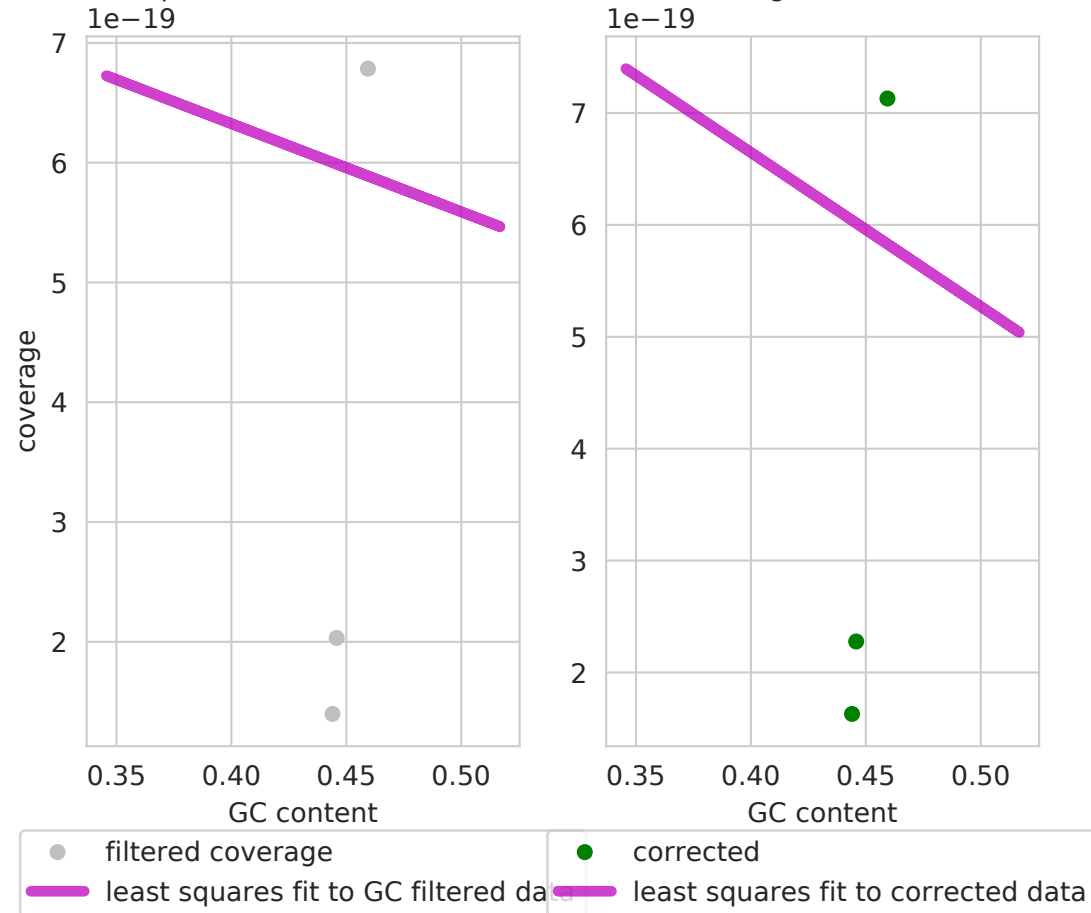


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0705/GCF_005938025.1_ASM593802v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0705-D
iRep: n/a r^2 : 0.90 avg. cov: 0.01 % windows: 0.50



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0705/GCF_005938025.1_ASM593802v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0705-D

iRep: n/a raw iRep: 10.07 GC r²: 0.00 GC bias: -0.00 avg. cov: 0.01 % windows: 0.50



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0705/GCF_005938025.1_ASM593802v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0705-D
iRep: n/a un-filtered iRep: 9.52 r^2: 0.90 avg. cov: 0.01 % windows: 0.50

