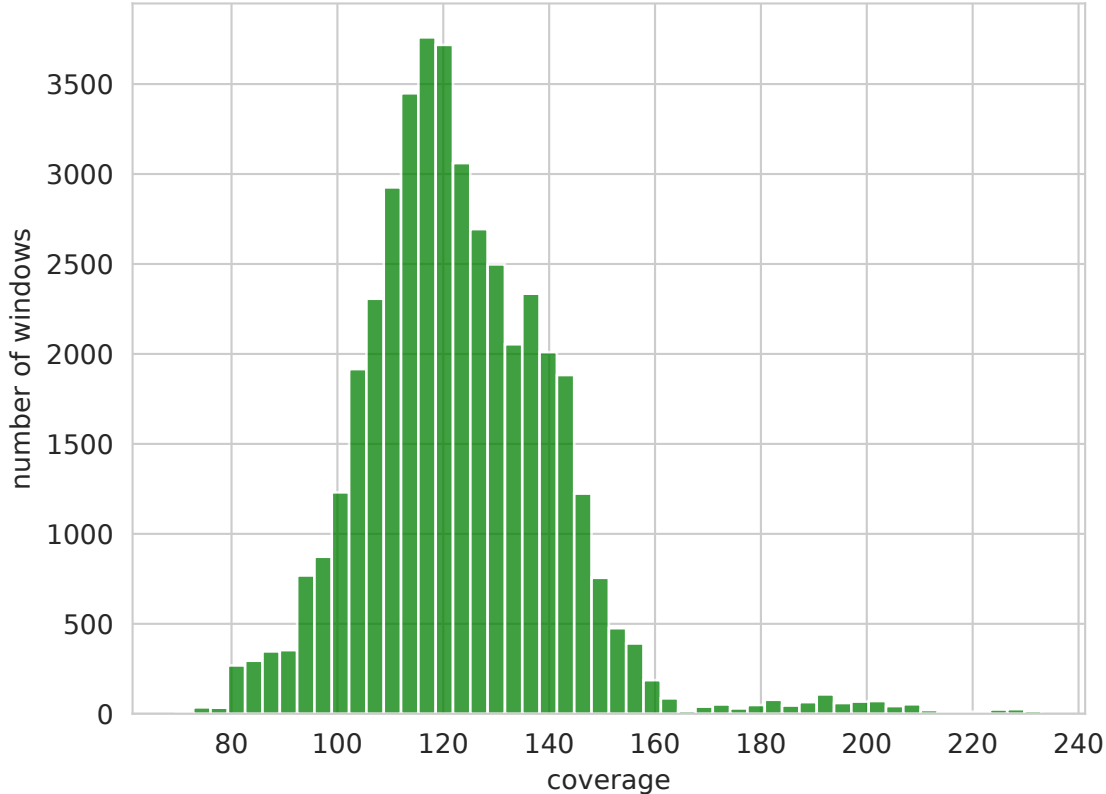
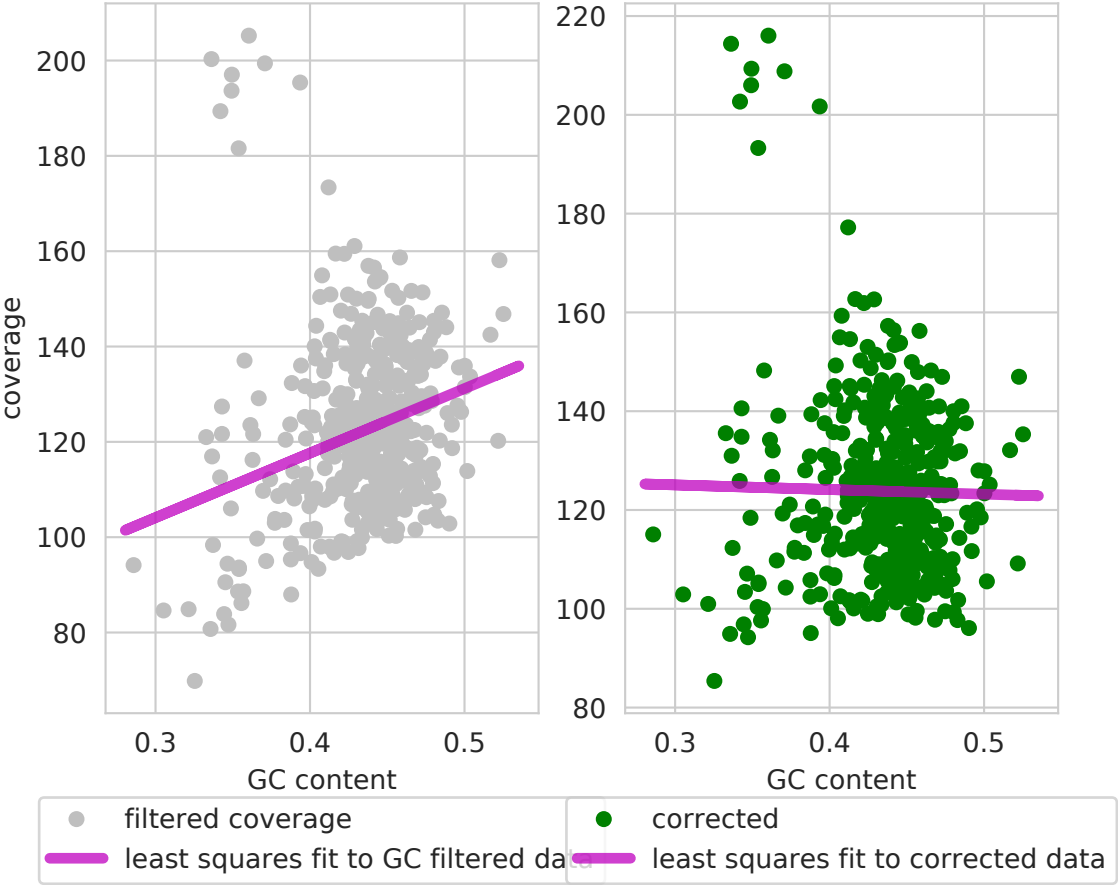


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0812/GCF_005937795.1_ASM593779v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0812-A
iRep: 1.47 r^2 : 0.99 avg. cov: 123.10 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0812/GCF_005937795.1_ASM593779v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0812-A
iRep: 1.47 raw iRep: 1.51 GC r²: 0.08 GC bias: 11.05 avg. cov: 123.10 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0812/GCF_005937795.1_ASM593779v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0812-A
iRep: 1.47 un-filtered iRep: 1.47 r^2 : 0.99 avg. cov: 123.10 % windows: 100.00

