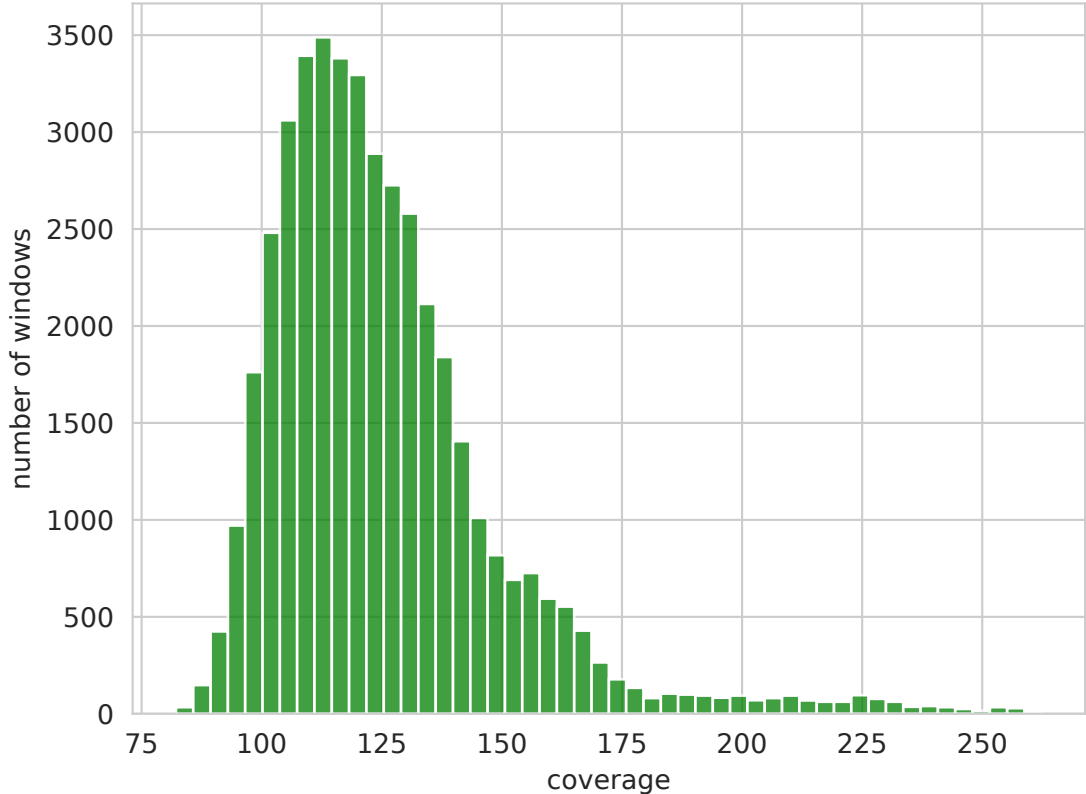
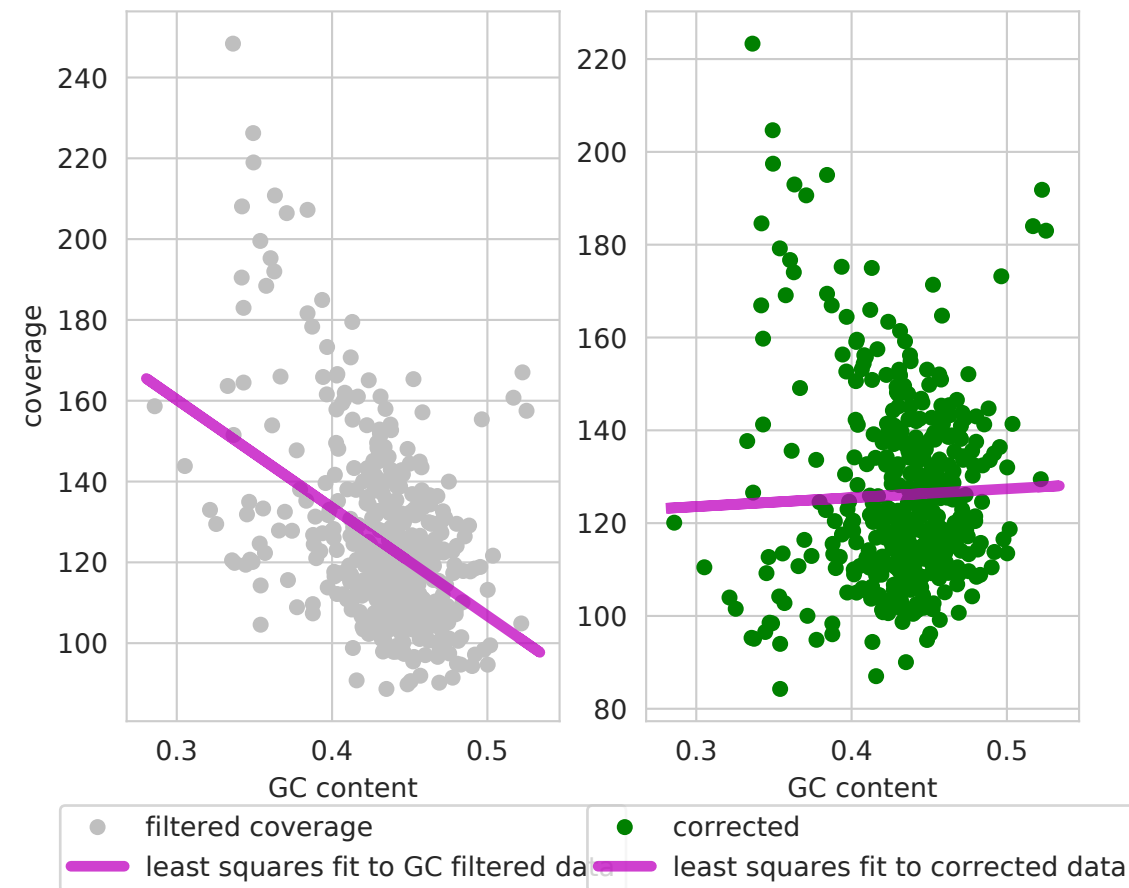


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0812/GCF_005937795.1_ASM593779v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0812-C
iRep: 1.58 r^2 : 0.98 avg. cov: 125.53 % windows: 100.00



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-C
iRep: 1.58 raw iRep: 1.62 GC r²: 0.19 GC bias: -51.47 avg. cov: 125.53 % 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-C
iRep: 1.58 un-filtered iRep: 1.58 r^2 : 0.98 avg. cov: 125.53 % windows: 100.00

