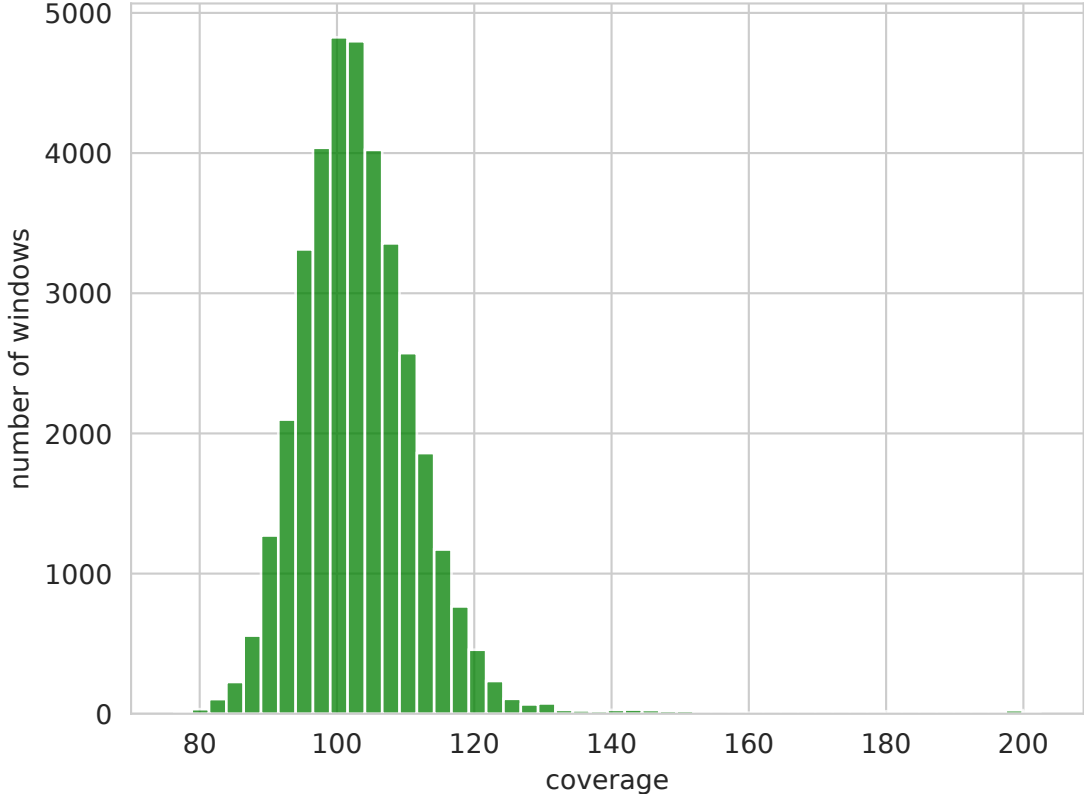
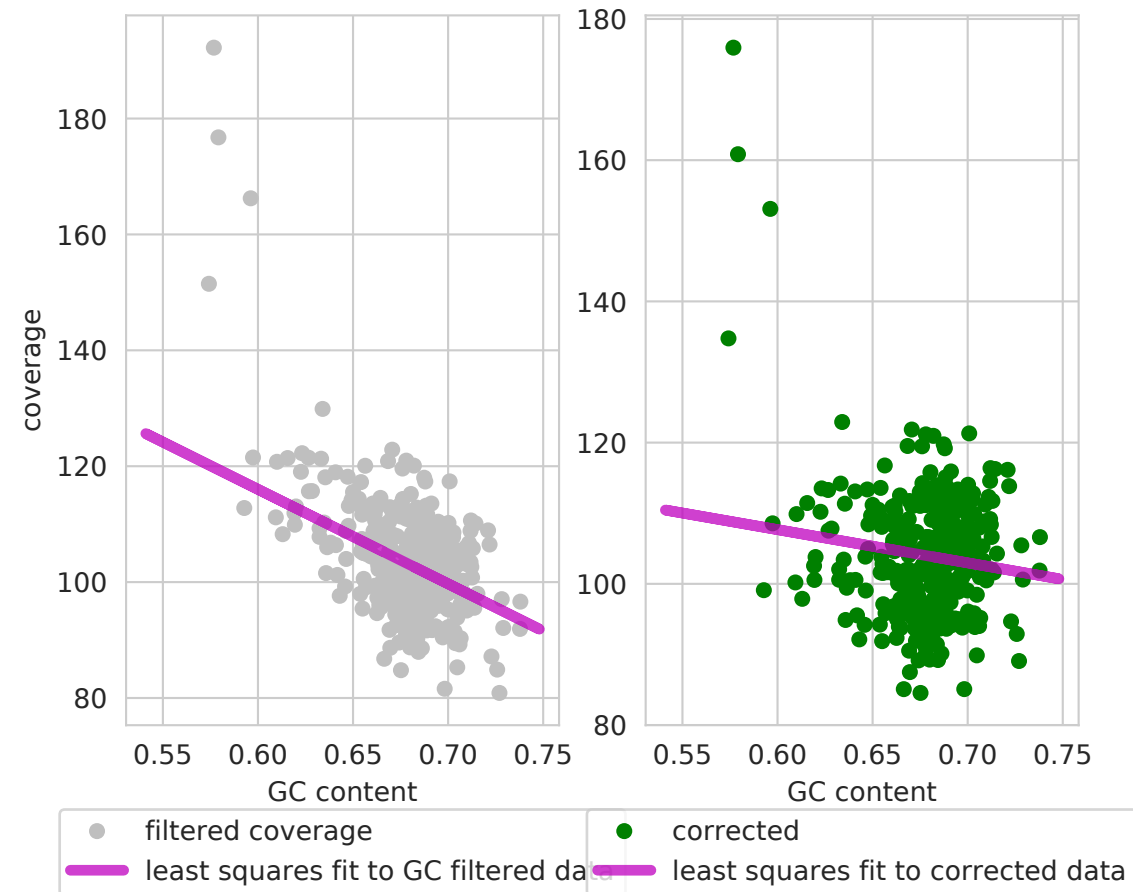


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0702/GCF_005937985.1_ASM593798v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0702-F
iRep: 1.23 r²: 0.99 avg. cov: 103.48 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0702/GCF_005937985.1_ASM593798v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0702-F
iRep: 1.23 raw iRep: 1.26 GC r^2 : 0.23 GC bias: -37.68 avg. cov: 103.48 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0702/GCF_005937985.1_ASM593798v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0702-F
iRep: 1.23 un-filtered iRep: 1.23 r^2 : 0.99 avg. cov: 103.48 % windows: 100.00

