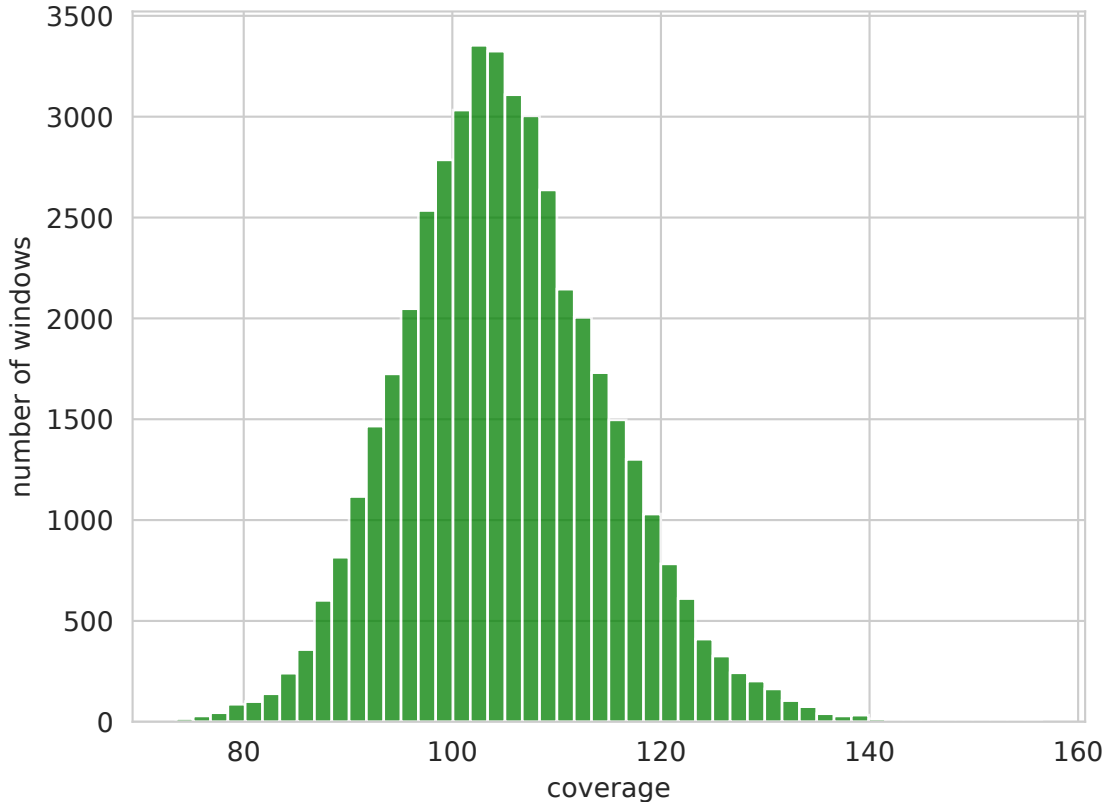
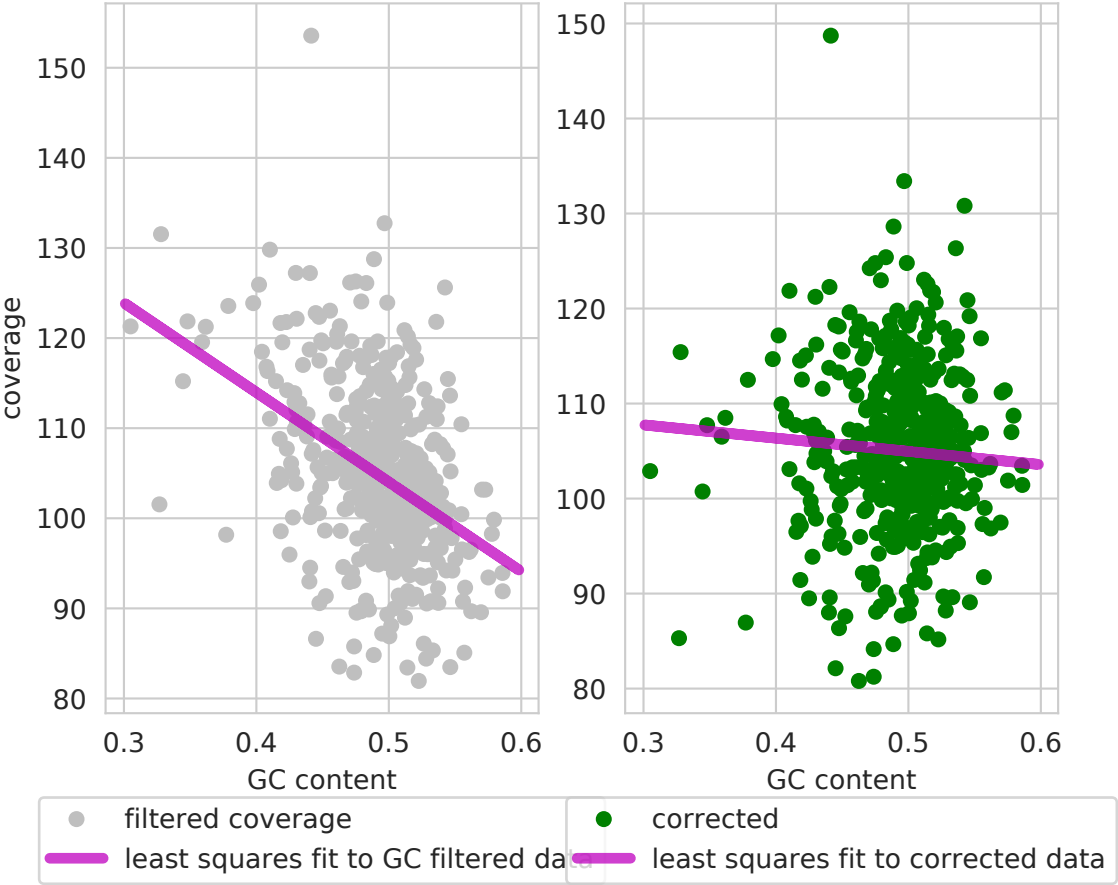


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0713/GCF_005937895.1_ASM593789v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0713-B
iRep: 1.29 r^2 : 0.99 avg. cov: 104.98 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0713/GCF_005937895.1_ASM593789v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0713-B
iRep: 1.29 raw iRep: 1.32 GC r²: 0.17 GC bias: -16.55 avg. cov: 104.98 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0713/GCF_005937895.1_ASM593789v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0713-B
iRep: 1.29 un-filtered iRep: 1.29 r^2: 0.99 avg. cov: 104.98 % windows: 100.00

