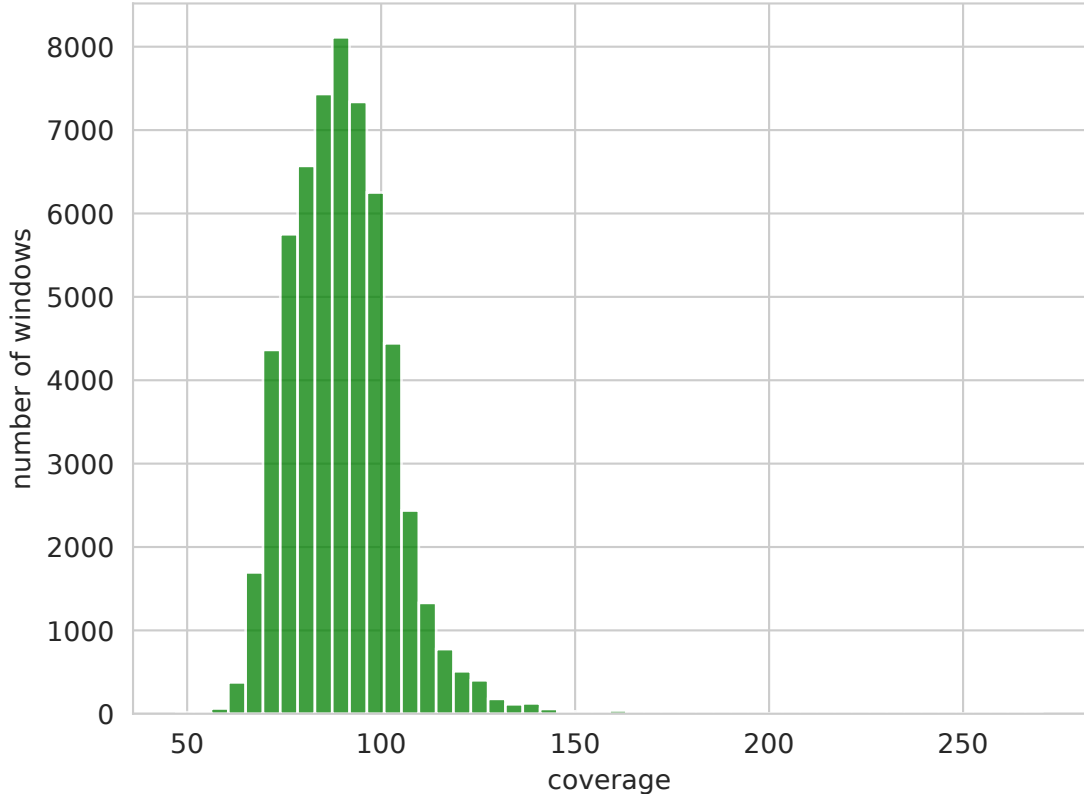
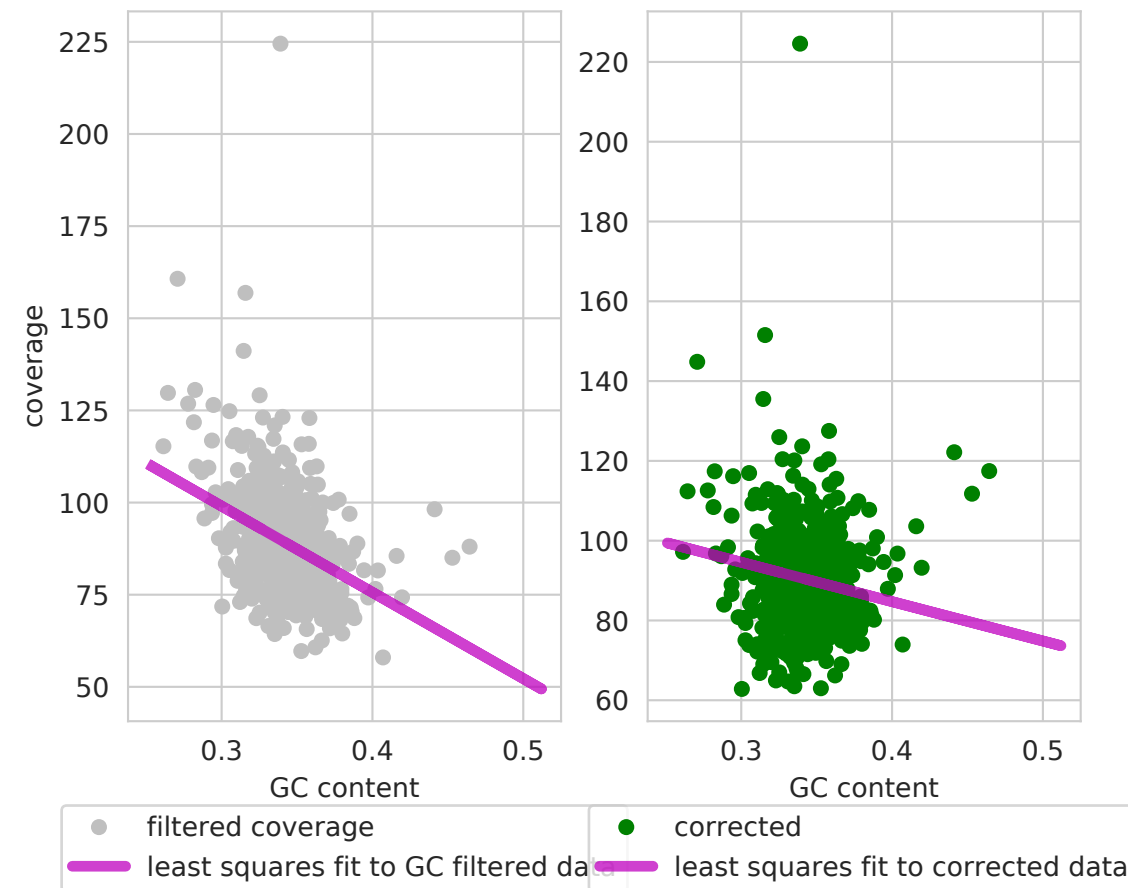


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0721/GCF_002007065_3_ASM200706v3_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0721-A
iRep: 1.49 r²: 0.99 avg. cov: 89.99 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0721/GCF_002007065_3_ASM200706v3_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0721-A
iRep: 1.49 raw iRep: 1.55 GC r^2 : 0.17 GC bias: -40.69 avg. cov: 89.99 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0721/GCF_002007065_3_ASM200706v3_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0721-A
iRep: 1.49 un-filtered iRep: 1.49 r^2 : 0.99 avg. cov: 89.99 % windows: 100.00

