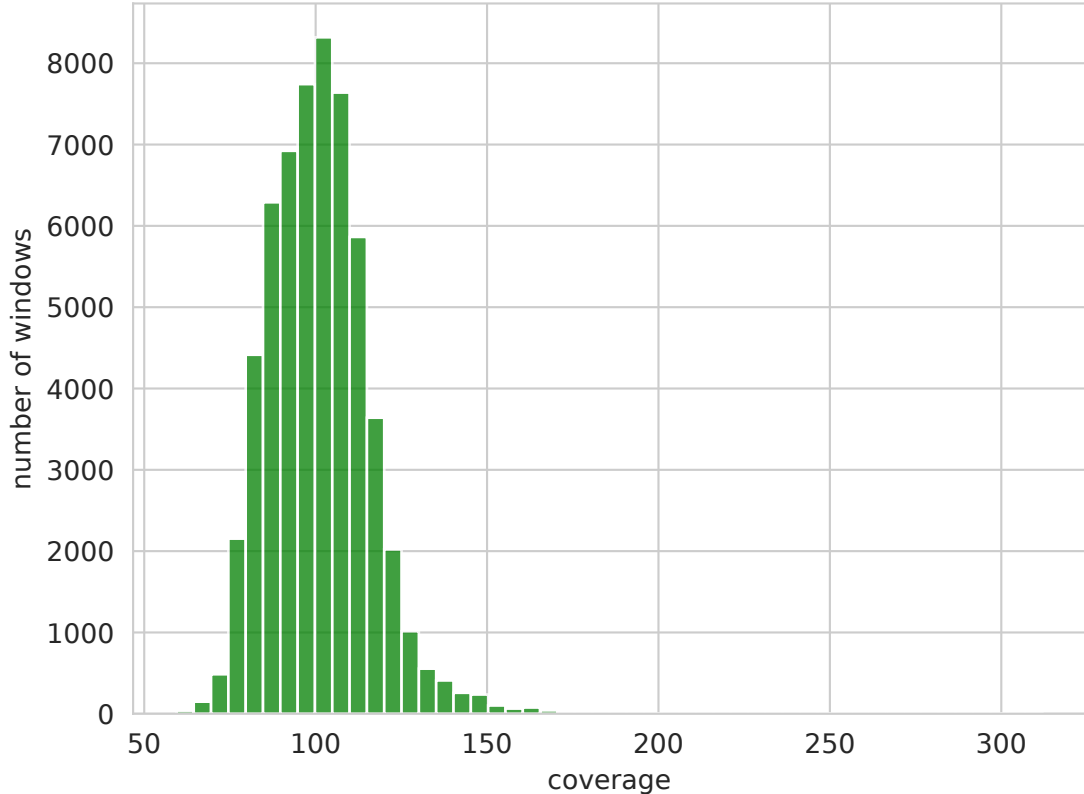
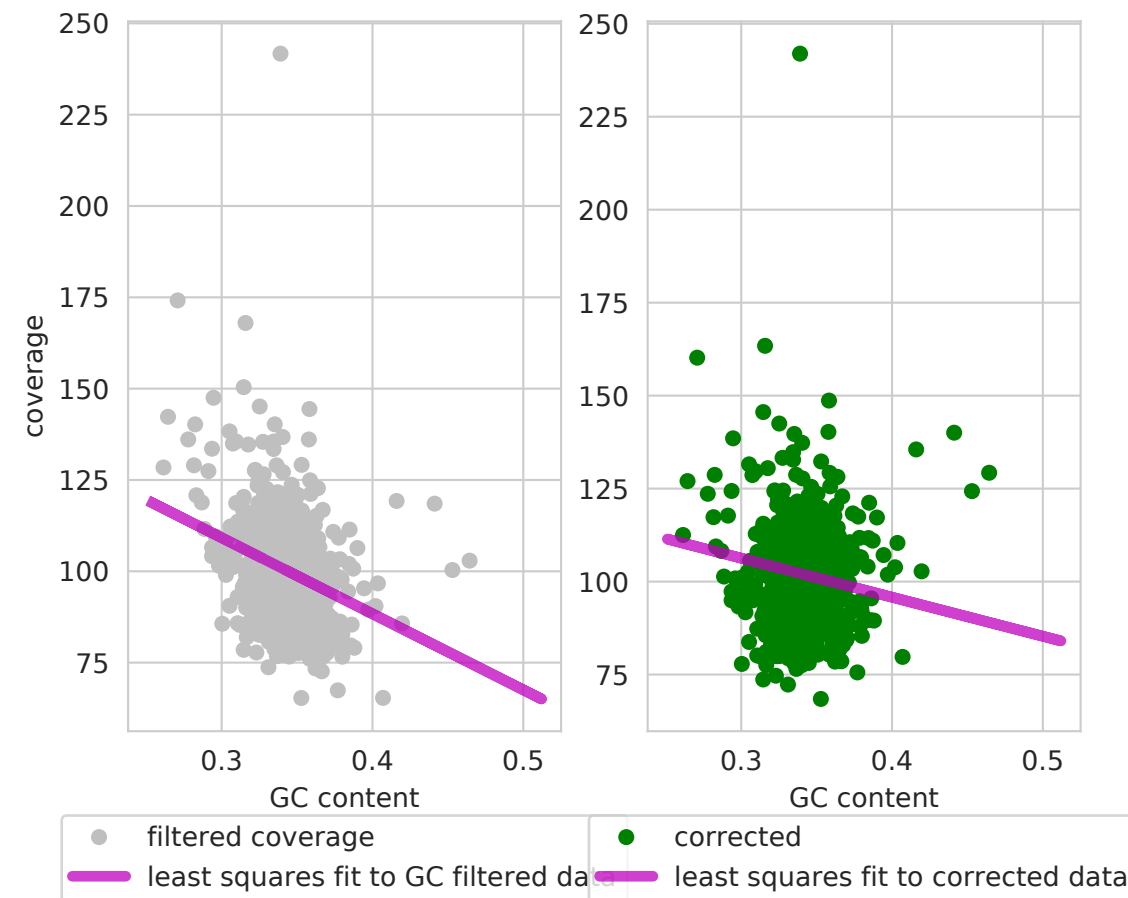


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0721/GCF_002007065_3_ASM200706v3_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0721-C
iRep: 1.48 r^2 : 0.99 avg. cov: 101.32 % windows: 100.00



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-C
iRep: 1.48 raw iRep: 1.52 GC r²: 0.12 GC bias: -24.18 avg. cov: 101.32 % 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-C
iRep: 1.48 un-filtered iRep: 1.48 r^2 : 0.99 avg. cov: 101.32 % windows: 100.00

