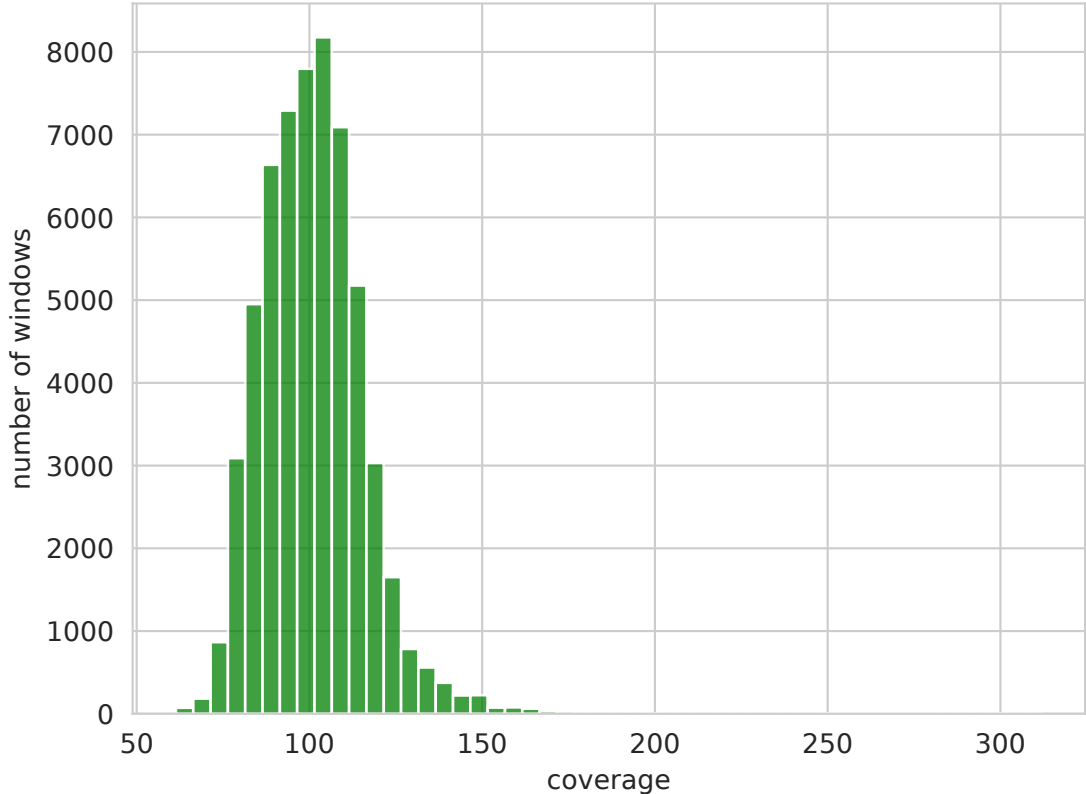
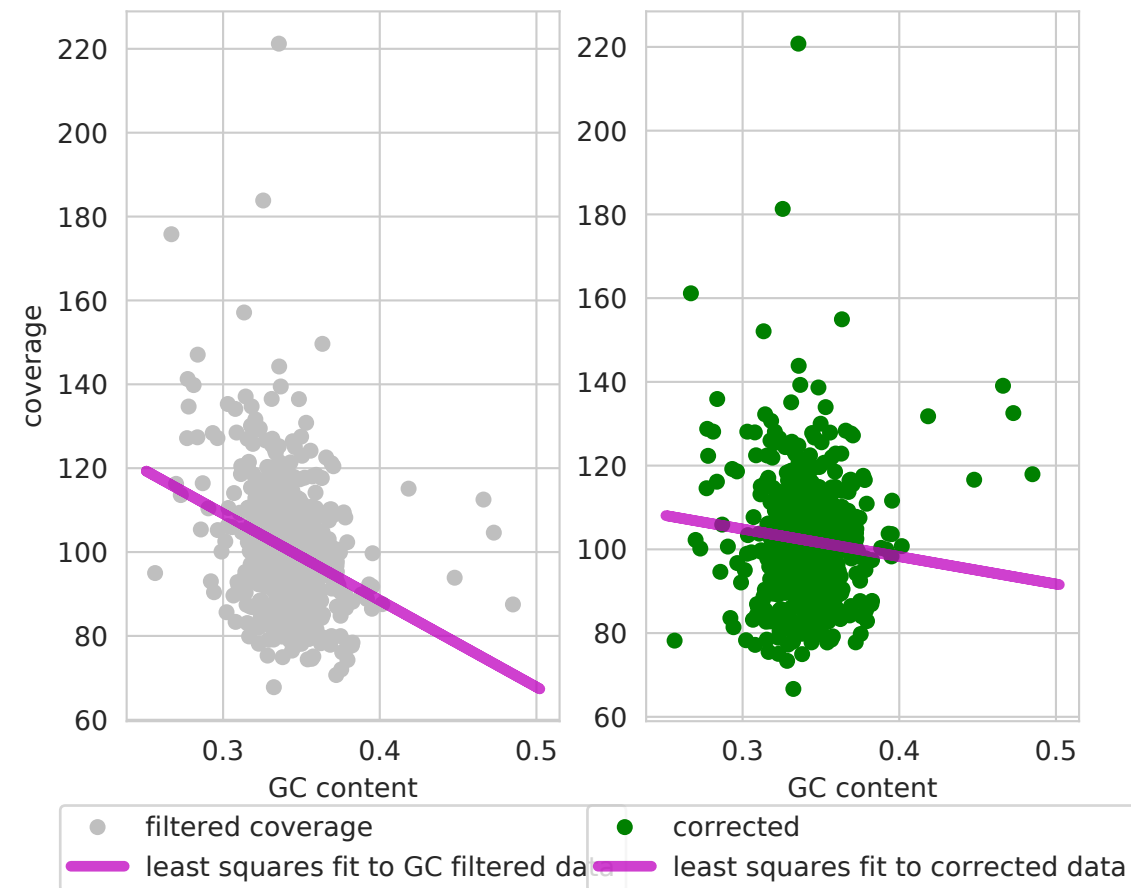


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0721/GCF_002007065.2_ASM200706v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0721-C
iRep: 1.48 r²: 0.99 avg. cov: 101.35 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0721/GCF_002007065.2_ASM200706v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0721-C
iRep: 1.48 raw iRep: 1.52 GC r^2 : 0.11 GC bias: -23.48 avg. cov: 101.35 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0721/GCF_002007065.2_ASM200706v2_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: 101.35 % windows: 100.00

