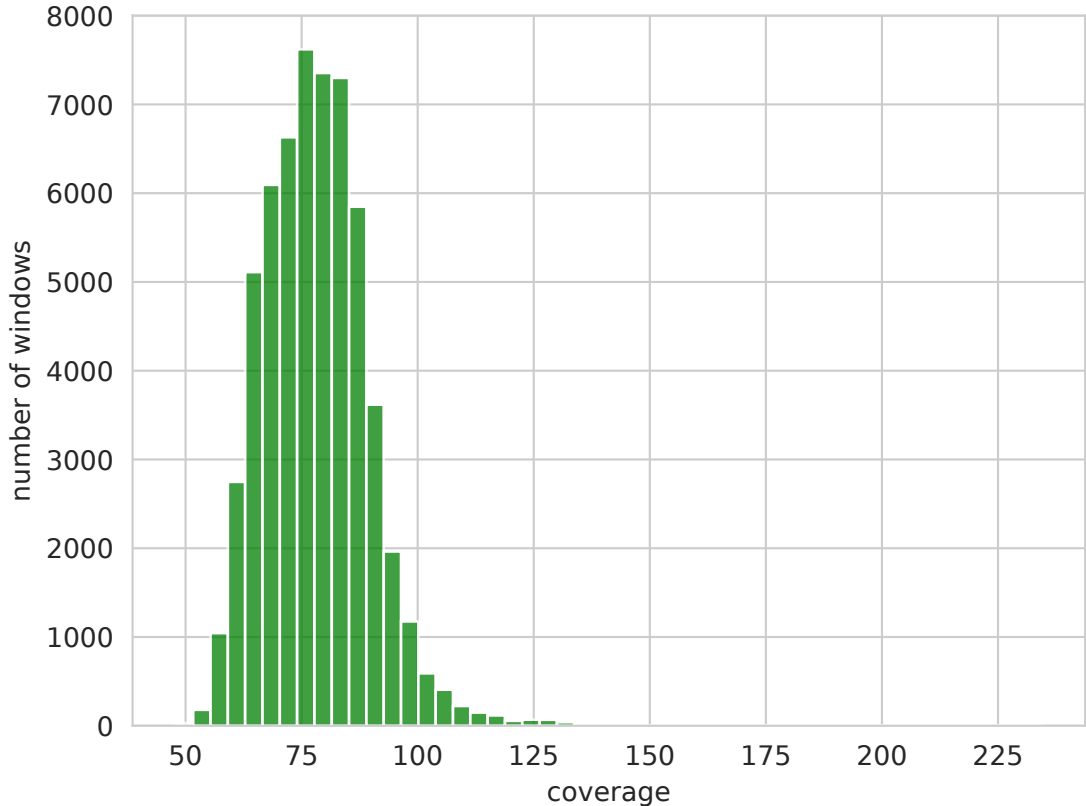
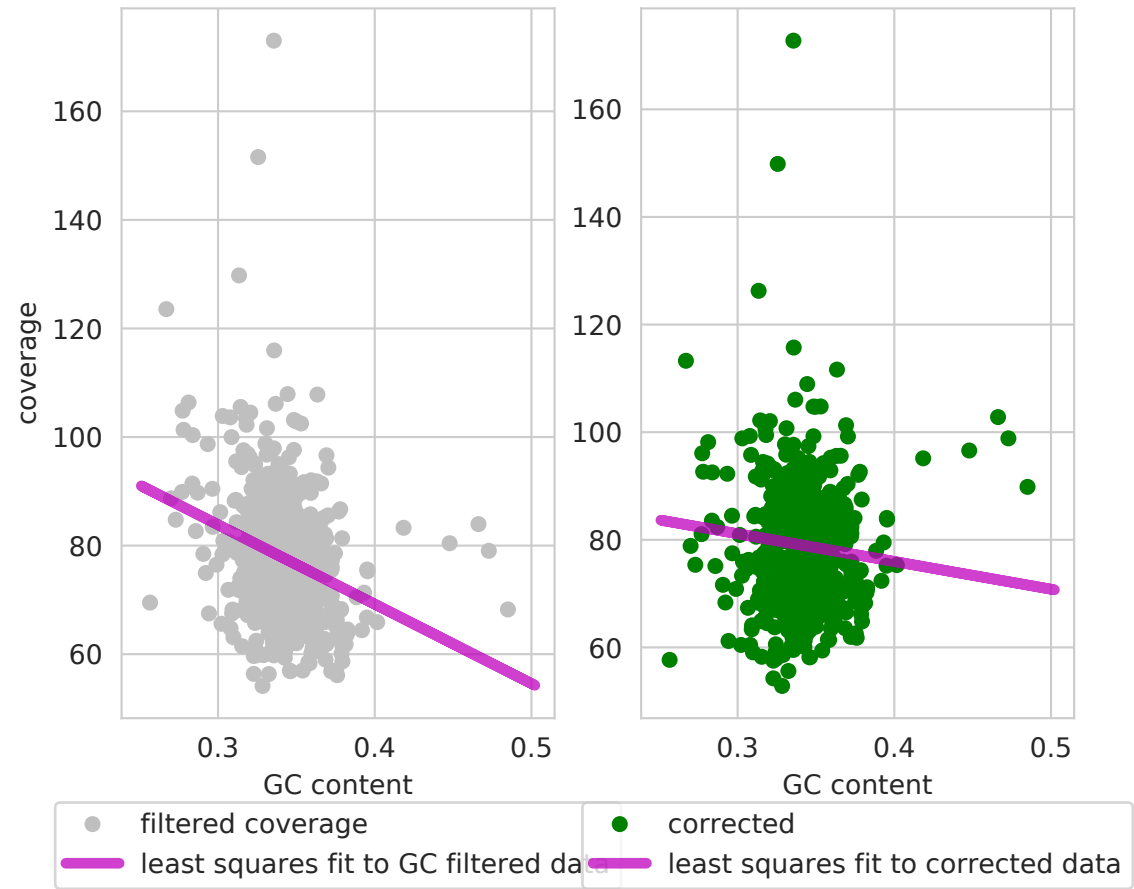


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0721/GCF_002007065.2_ASM200706v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0721-B
iRep: 1.49 r²: 0.99 avg. cov: 78.38 % windows: 100.00



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-B
iRep: 1.49 raw iRep: 1.53 GC r^2: 0.10 GC bias: -14.18 avg. cov: 78.38 % 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-B
iRep: 1.49 un-filtered iRep: 1.49 r^2 : 0.99 avg. cov: 78.38 % windows: 100.00

