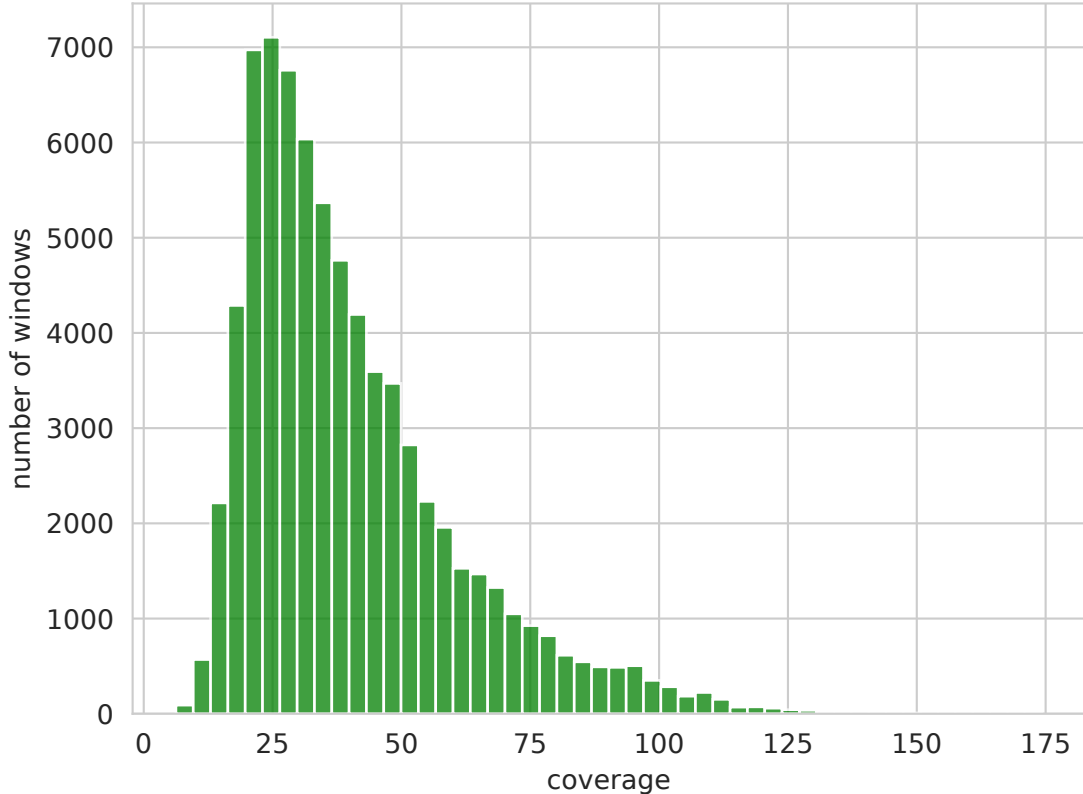
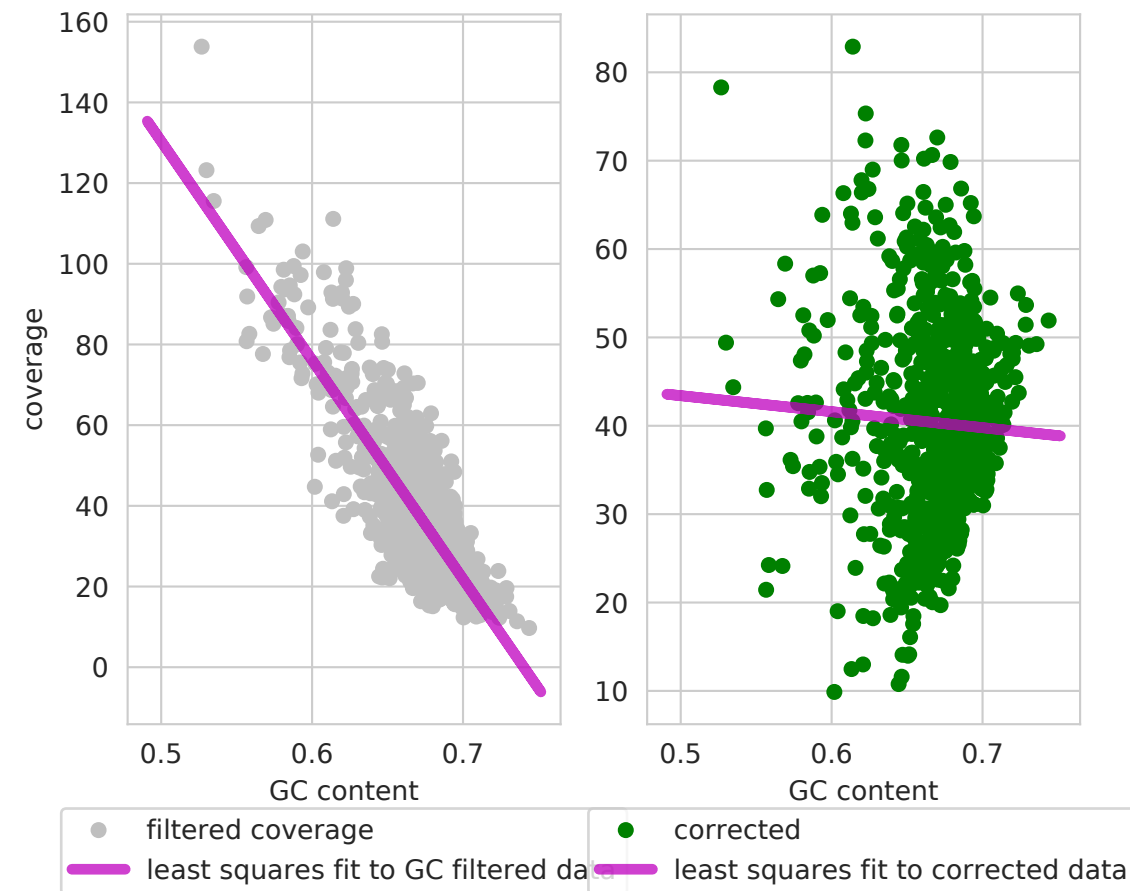


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0801/GCF_005937855.1_ASM593785v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0801-D
iRep: 2.42 r²: 0.98 avg. cov: 40.30 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0801/GCF_005937855.1_ASM593785v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0801-D
iRep: 2.42 raw iRep: 4.48 GC r^2 : 0.68 GC bias: -371.01 avg. cov: 40.30 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0801/GCF_005937855.1_ASM593785v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0801-D
iRep: 2.42 un-filtered iRep: 2.42 r^2 : 0.98 avg. cov: 40.30 % windows: 100.00

