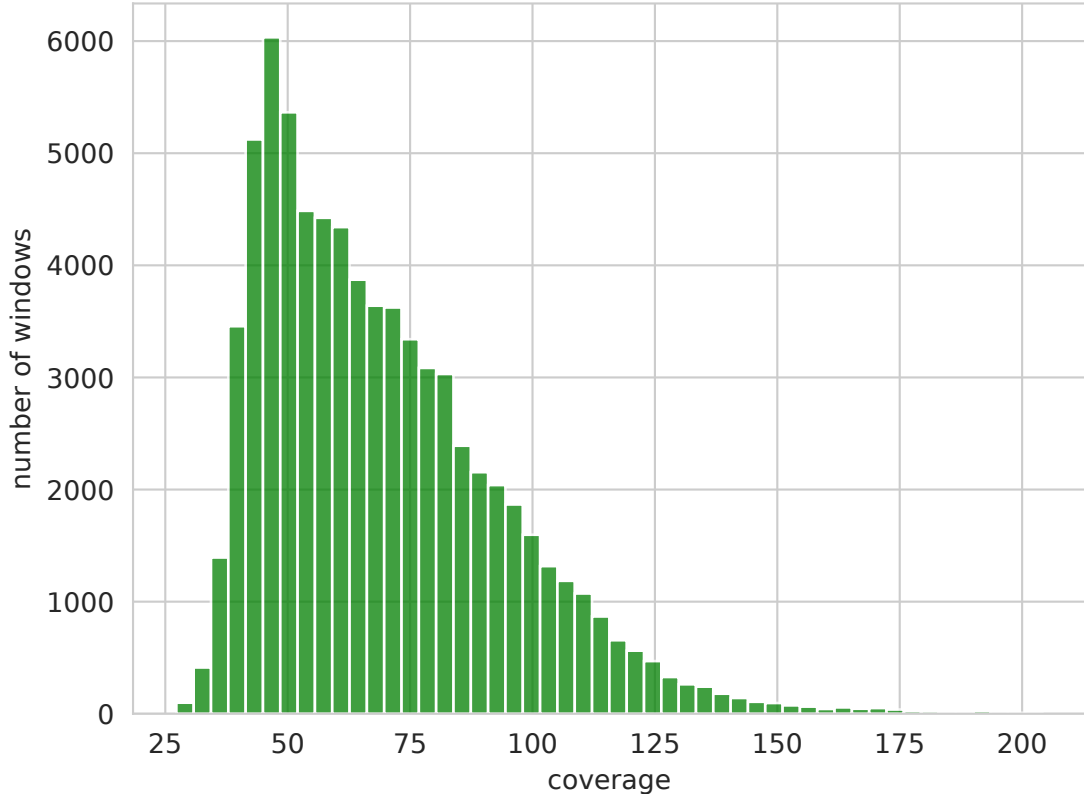
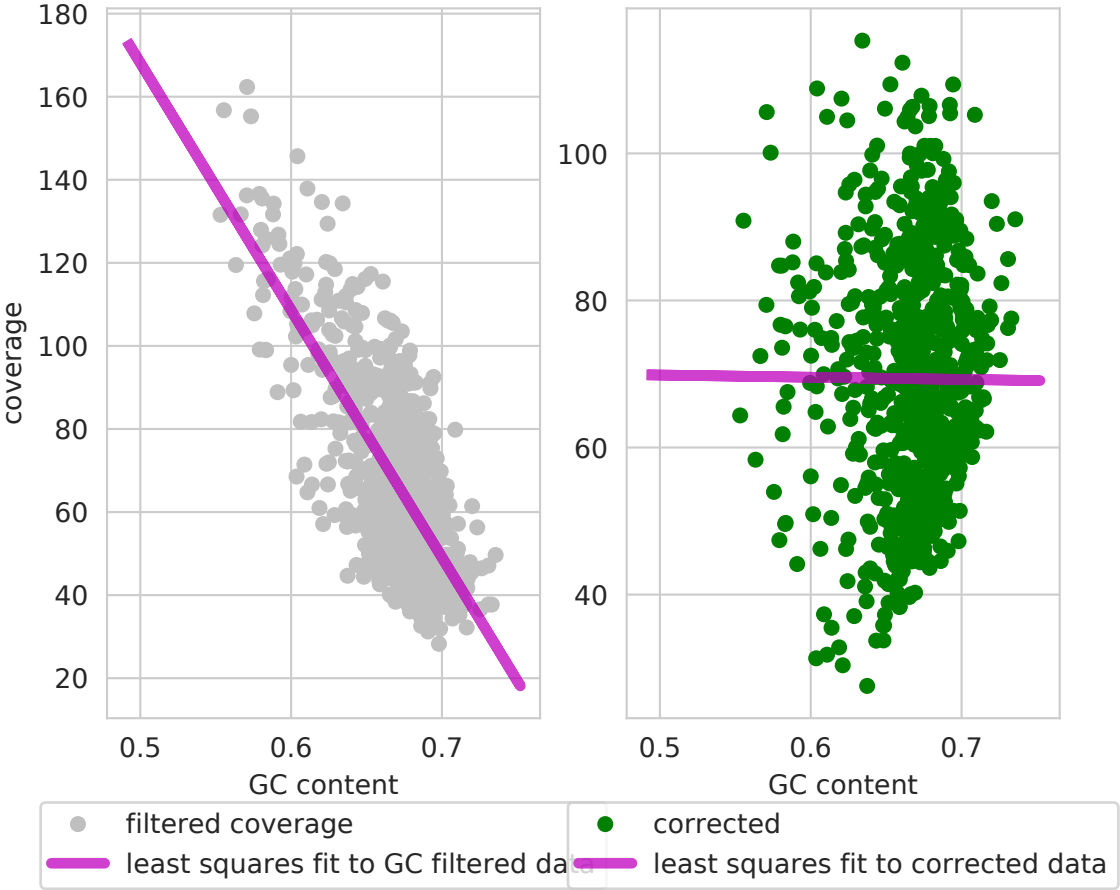


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0801/GCF_005937855_2_ASM593785v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0801-A
iRep: 2.10 r^2 : 0.99 avg. cov: 69.27 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0801/GCF_005937855_2_ASM593785v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0801-A
iRep: 2.10 raw iRep: 2.93 GC r²: 0.58 GC bias: -344.63 avg. cov: 69.27 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0801/GCF_005937855_2_ASM593785v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0801-A
iRep: 2.10 un-filtered iRep: 2.10 r^2 : 0.99 avg. cov: 69.27 % windows: 100.00

