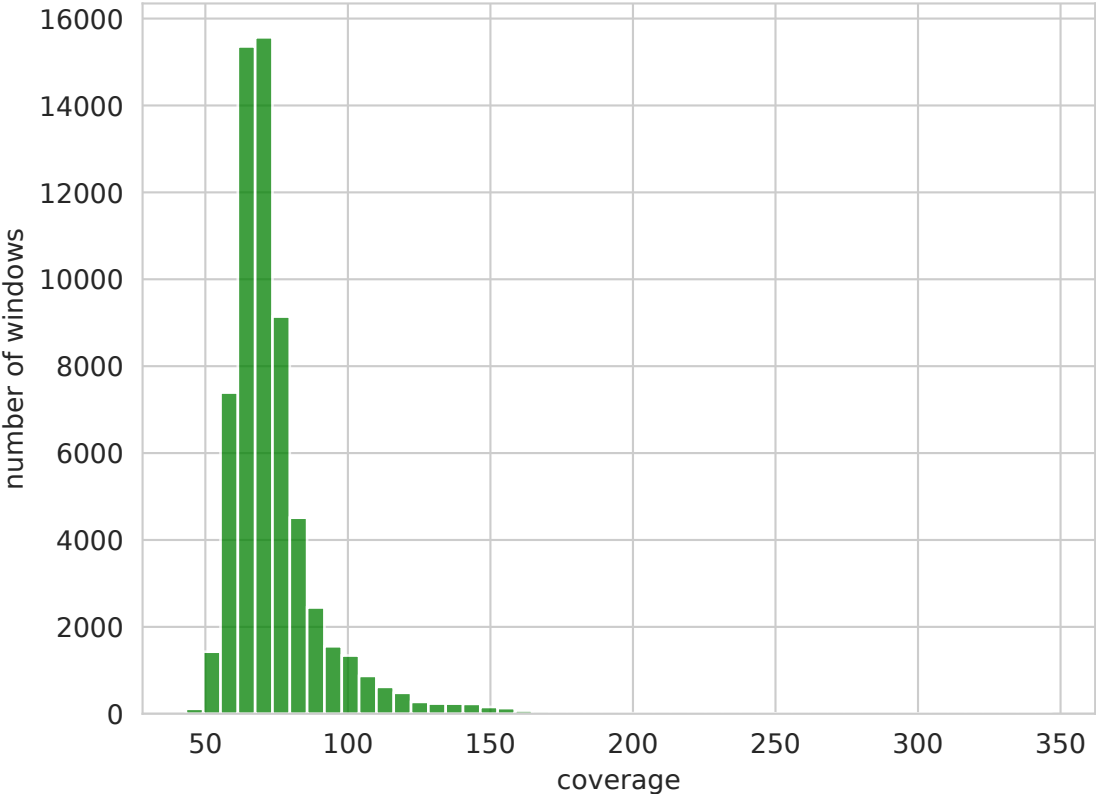
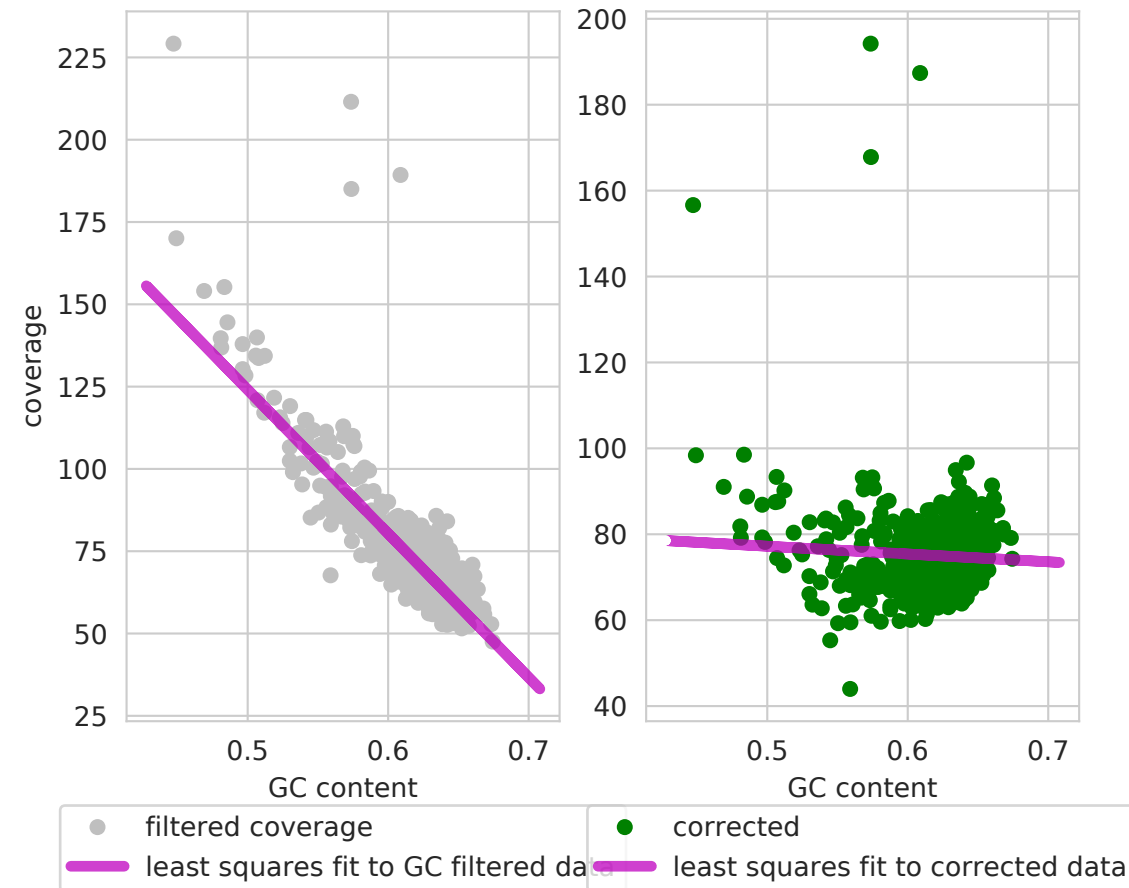


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0802/GCF_005937845.1_ASM593784v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0802-A
iRep: 1.30 r²: 0.98 avg. cov: 74.61 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0802/GCF_005937845.1_ASM593784v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0802-A
iRep: 1.30 raw iRep: 1.63 GC r²: 0.82 GC bias: -357.75 avg. cov: 74.61 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0802/GCF_005937845.1_ASM593784v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0802-A
iRep: 1.30 un-filtered iRep: 1.30 r^2 : 0.98 avg. cov: 74.61 % windows: 100.00

