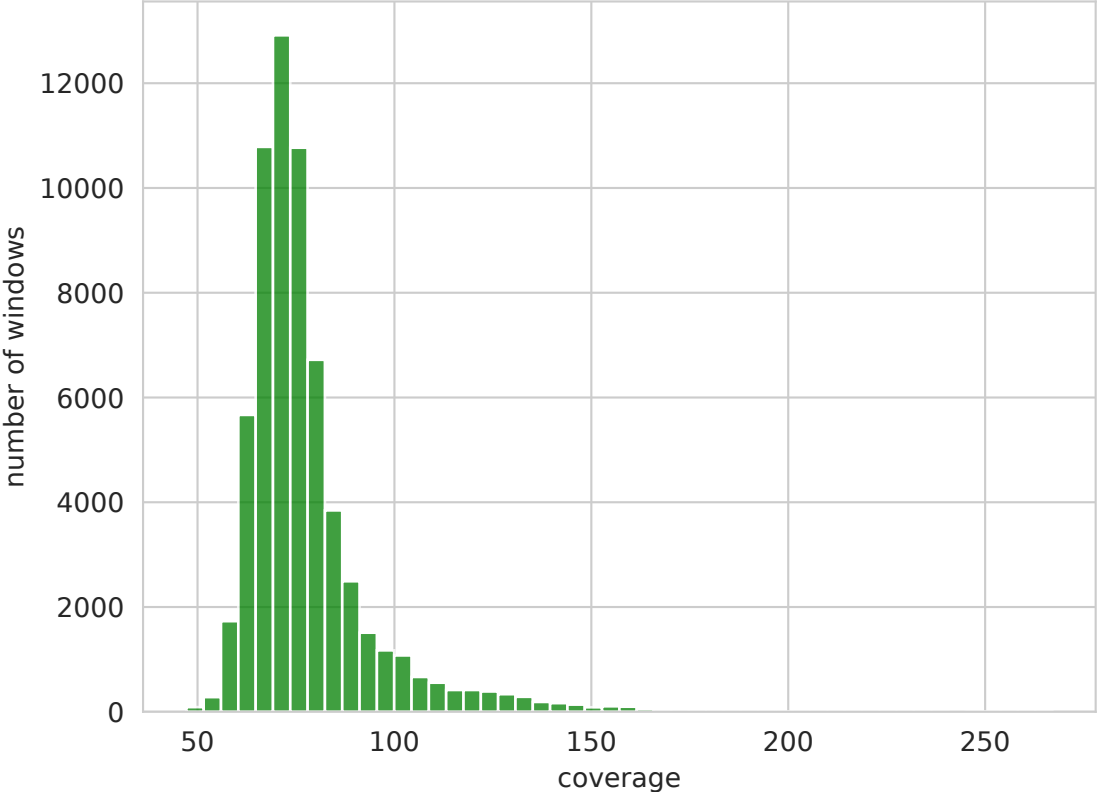
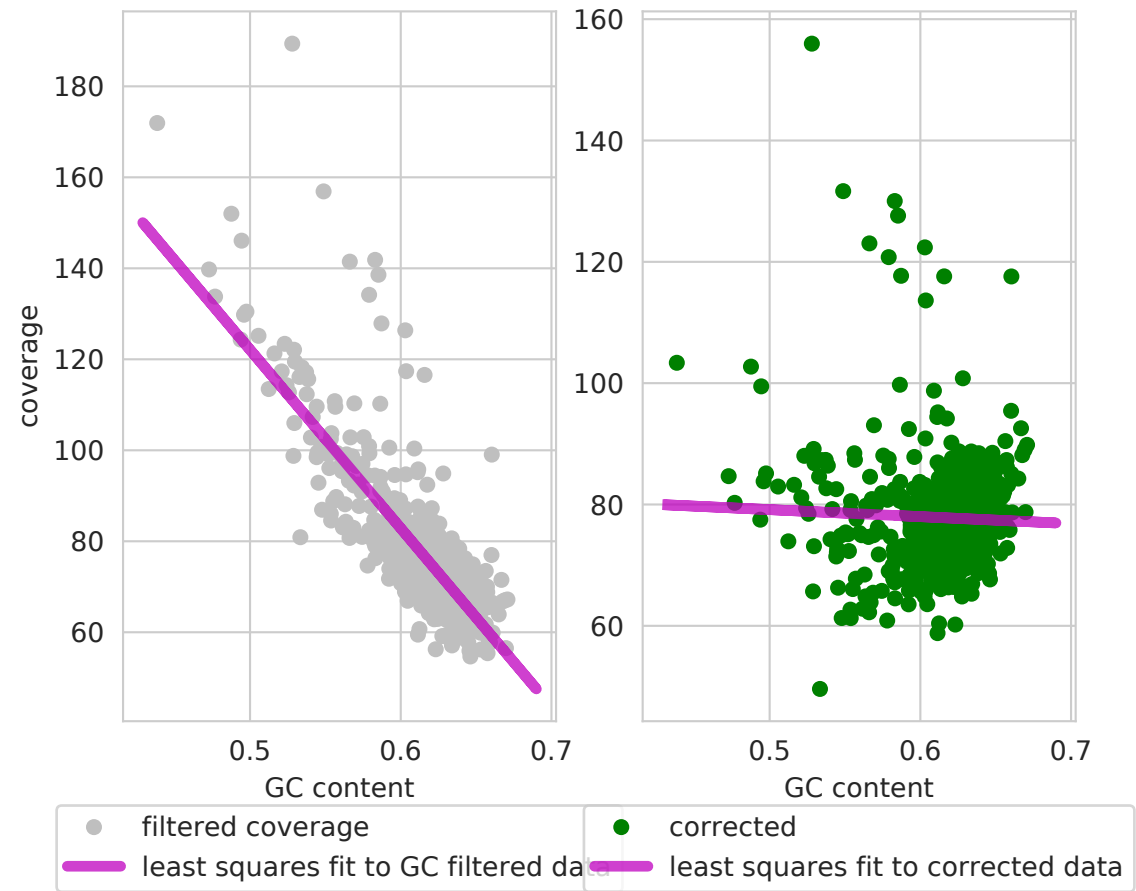


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0802/GCF_005937845_2_ASM593784v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0802-B
iRep: 1.29 r²: 0.98 avg. cov: 77.73 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0802/GCF_005937845_2_ASM593784v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0802-B
iRep: 1.29 raw iRep: 1.56 GC r²: 0.75 GC bias: -295.83 avg. cov: 77.73 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0802/GCF_005937845_2_ASM593784v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0802-B
iRep: 1.29 un-filtered iRep: 1.29 r^2 : 0.98 avg. cov: 77.73 % windows: 100.00

