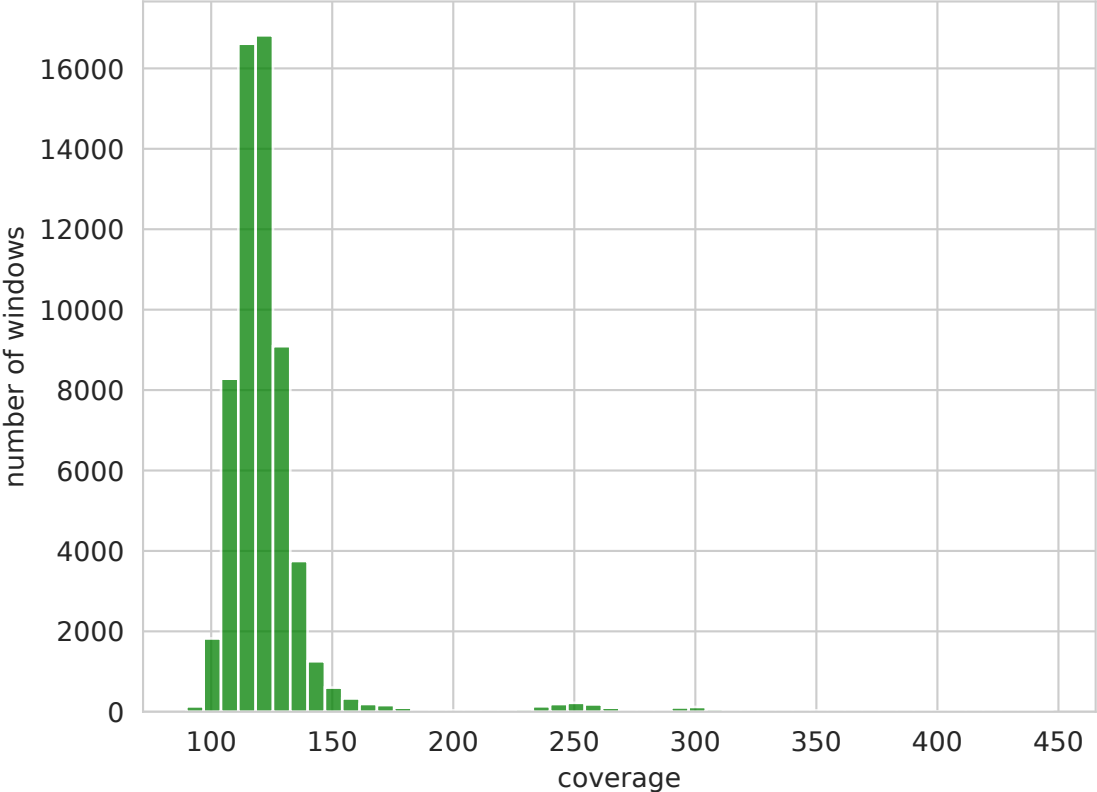
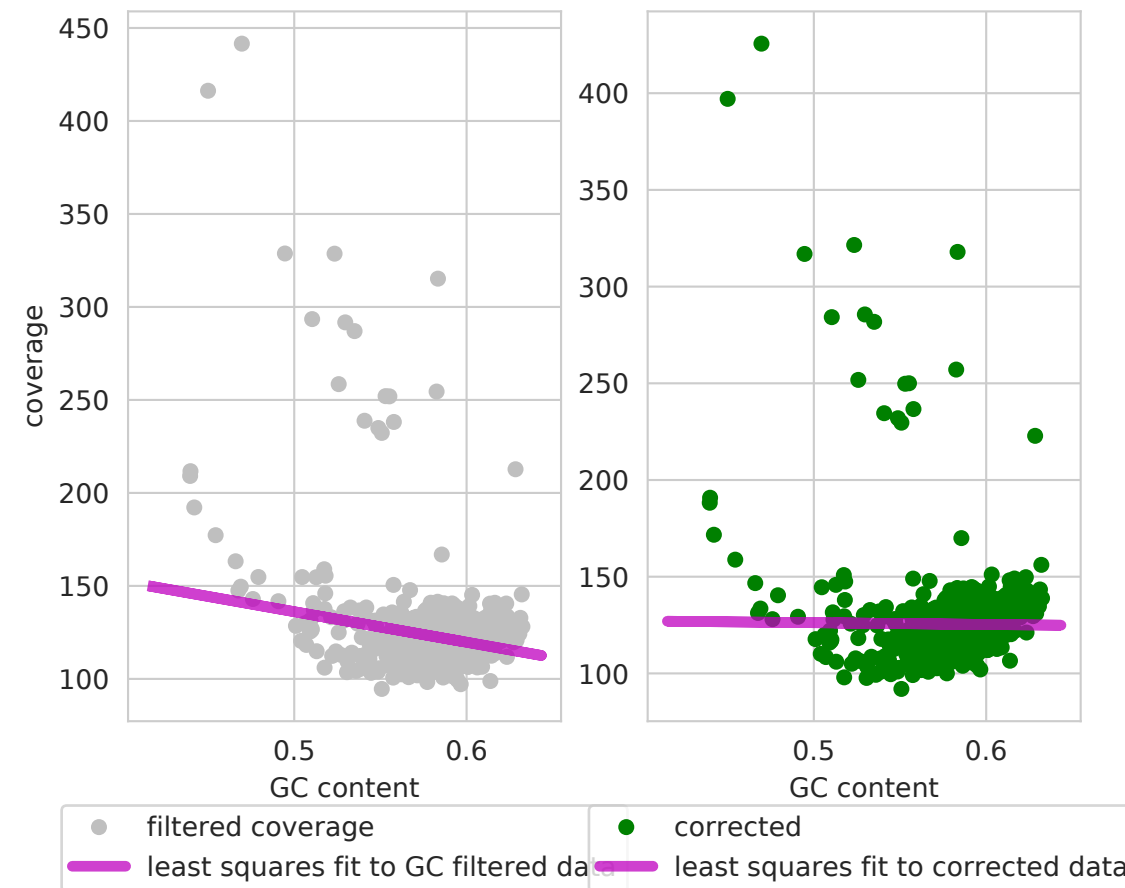


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0707/GCF_005937945_2_ASM593794v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0707-B
iRep: 1.33 r²: 0.98 avg. cov: 125.25 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0707/GCF_005937945_2_ASM593794v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0707-B
iRep: 1.33 raw iRep: 1.30 GC r^2 : 0.05 GC bias: -8.74 avg. cov: 125.25 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0707/GCF_005937945.2_ASM593794v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0707-B
iRep: 1.33 un-filtered iRep: 1.33 r^2 : 0.98 avg. cov: 125.25 % windows: 100.00

