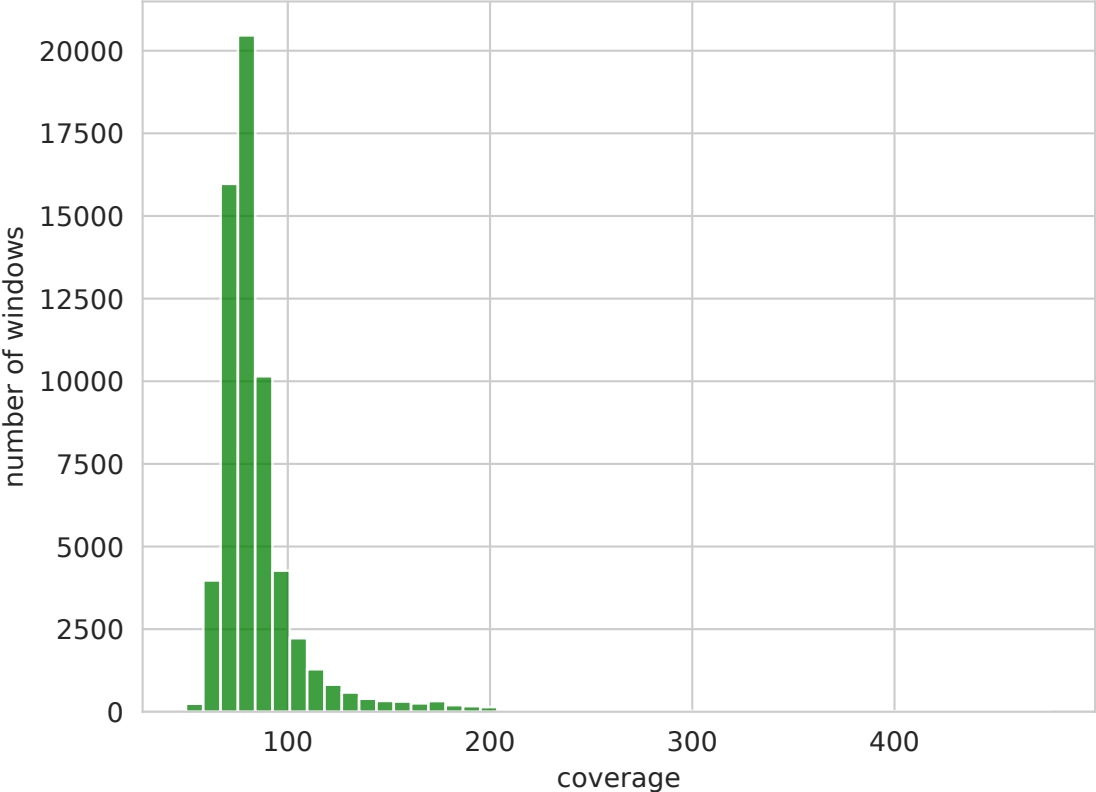
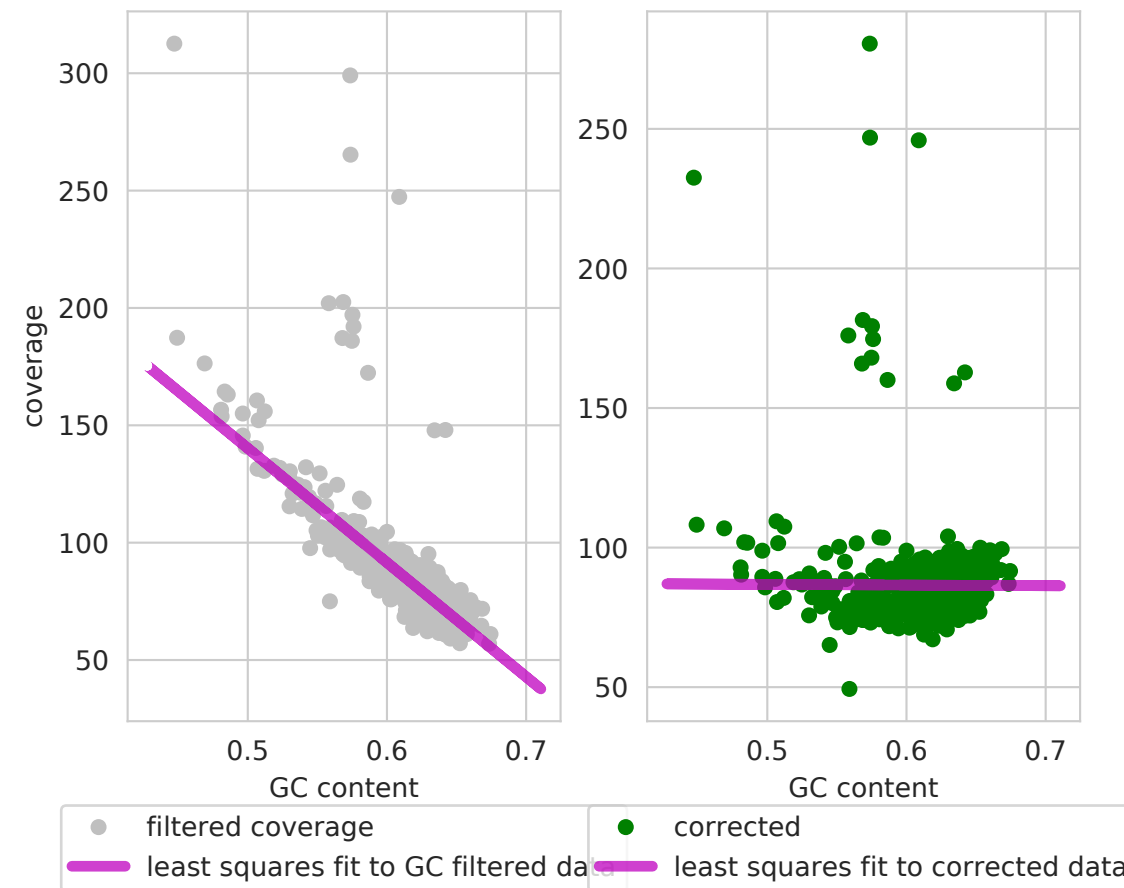


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0802/GCF_005937845.1_ASM593784v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0802-D
iRep: 1.28 r^2 : 0.98 avg. cov: 85.87 % windows: 100.00



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-D
iRep: 1.28 raw iRep: 1.61 GC r^2 : 0.70 GC bias: -341.41 avg. cov: 85.87 % 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-D
iRep: 1.28 un-filtered iRep: 1.28 r^2 : 0.98 avg. cov: 85.87 % windows: 100.00

