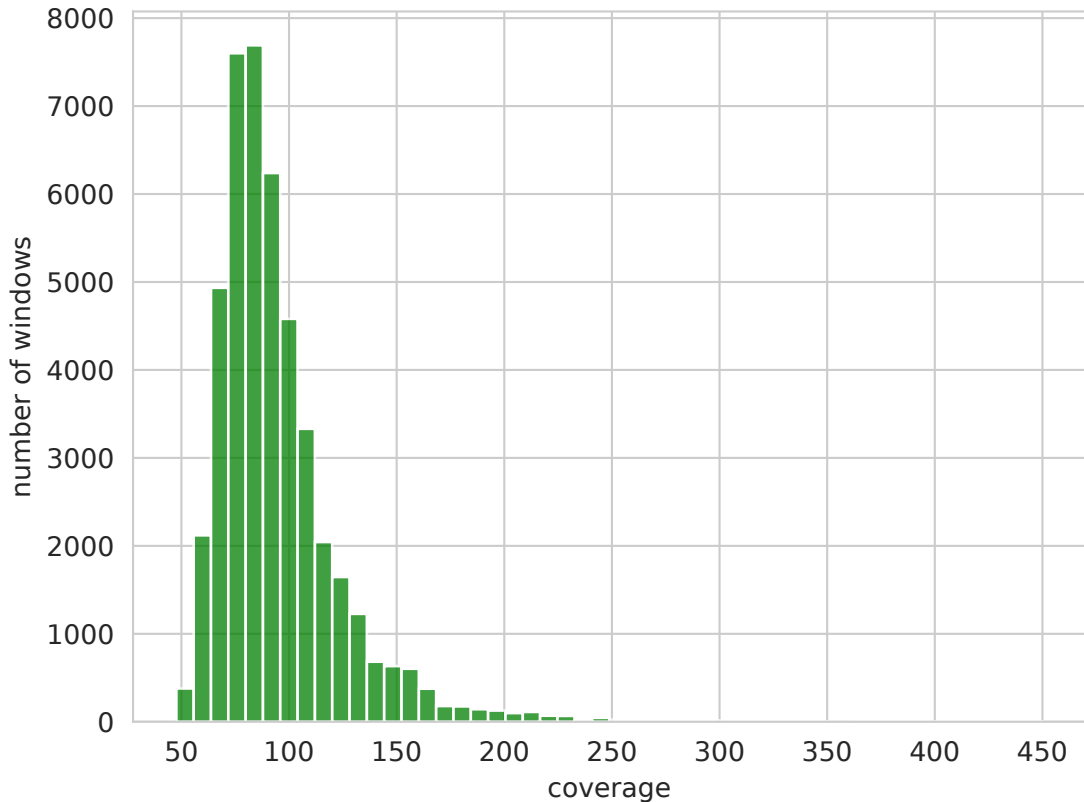
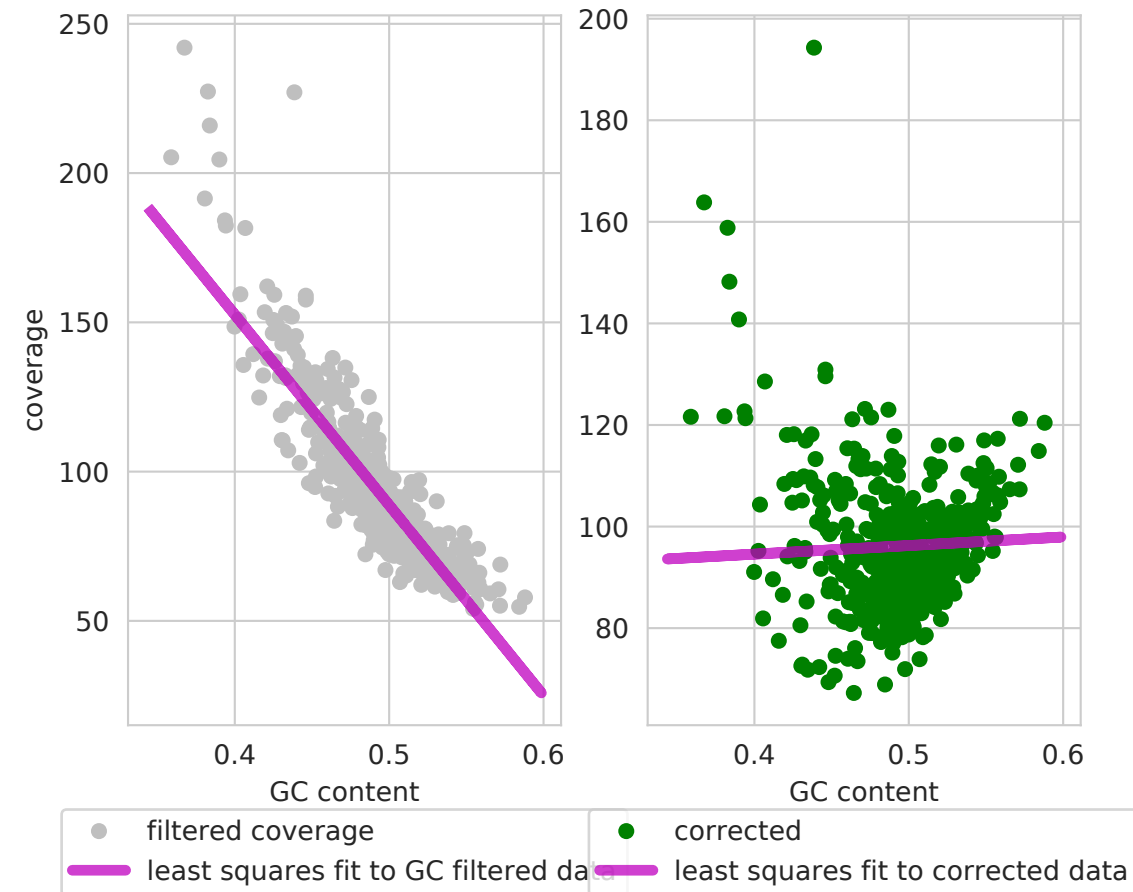


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0713/GCF_005937895_2_ASM593789v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0713-A
iRep: 1.42 r^2 : 0.98 avg. cov: 95.17 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0713/GCF_005937895_2_ASM593789v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0713-A
iRep: 1.42 raw iRep: 2.12 GC r^2 : 0.79 GC bias: -506.02 avg. cov: 95.17 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0713/GCF_005937895_2_ASM593789v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0713-A
iRep: 1.42 un-filtered iRep: 1.42 r^2 : 0.98 avg. cov: 95.17 % windows: 100.00

