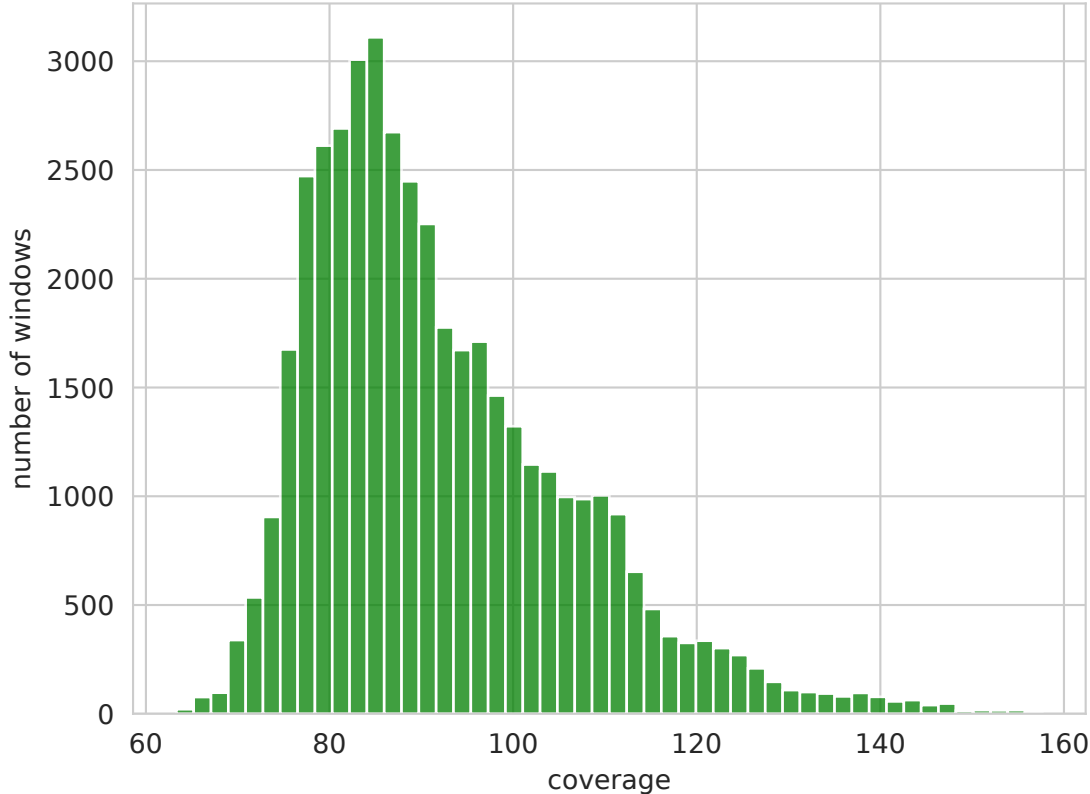
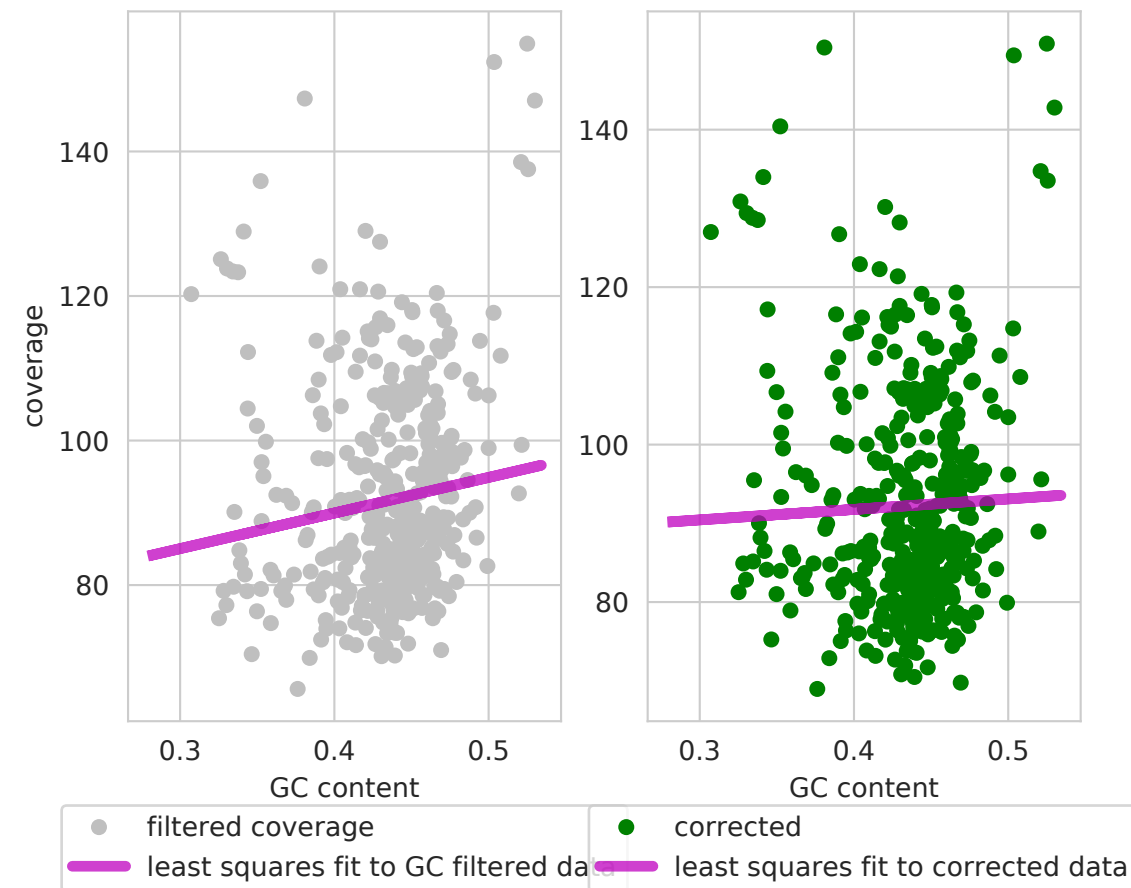


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0812/GCF_005937795_2_ASM593779v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0812-B
iRep: 1.54 r^2 : 0.97 avg. cov: 92.16 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0812/GCF_005937795_2_ASM593779v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0812-B
iRep: 1.54 raw iRep: 1.56 GC r^2 : 0.02 GC bias: 0.86 avg. cov: 92.16 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0812/GCF_005937795_2_ASM593779v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0812-B
iRep: 1.54 un-filtered iRep: 1.54 r²: 0.97 avg. cov: 92.16 % windows: 100.00

