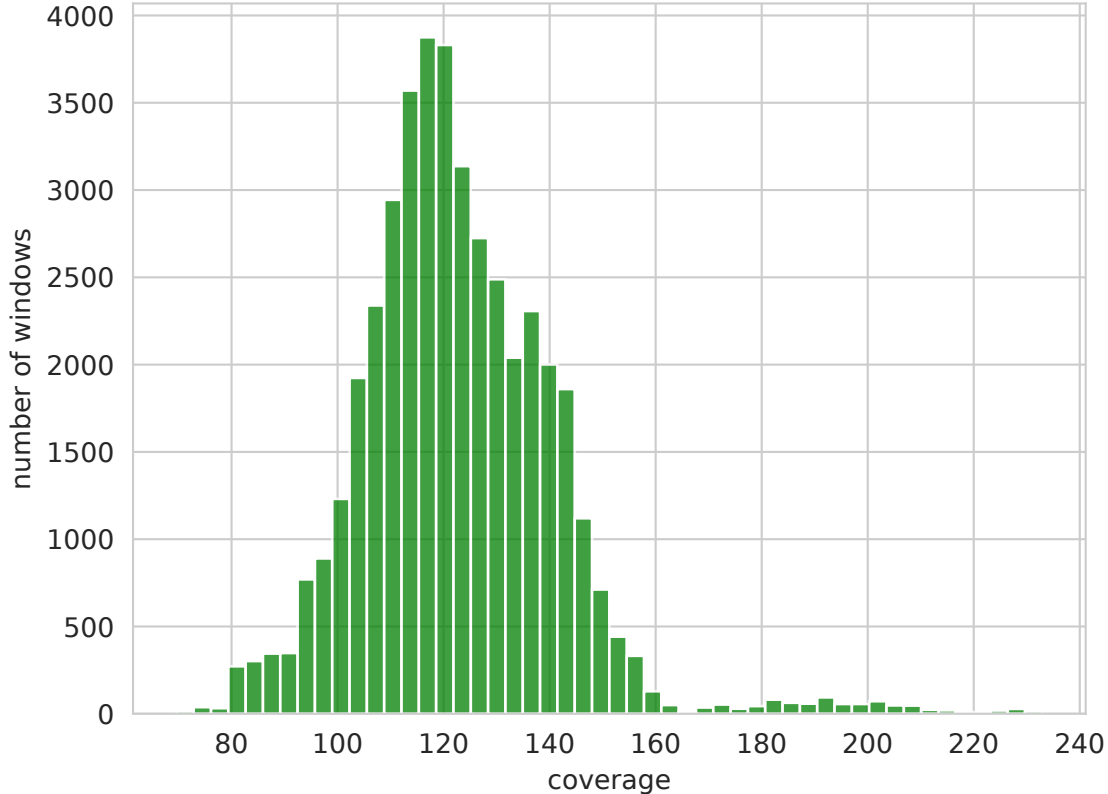
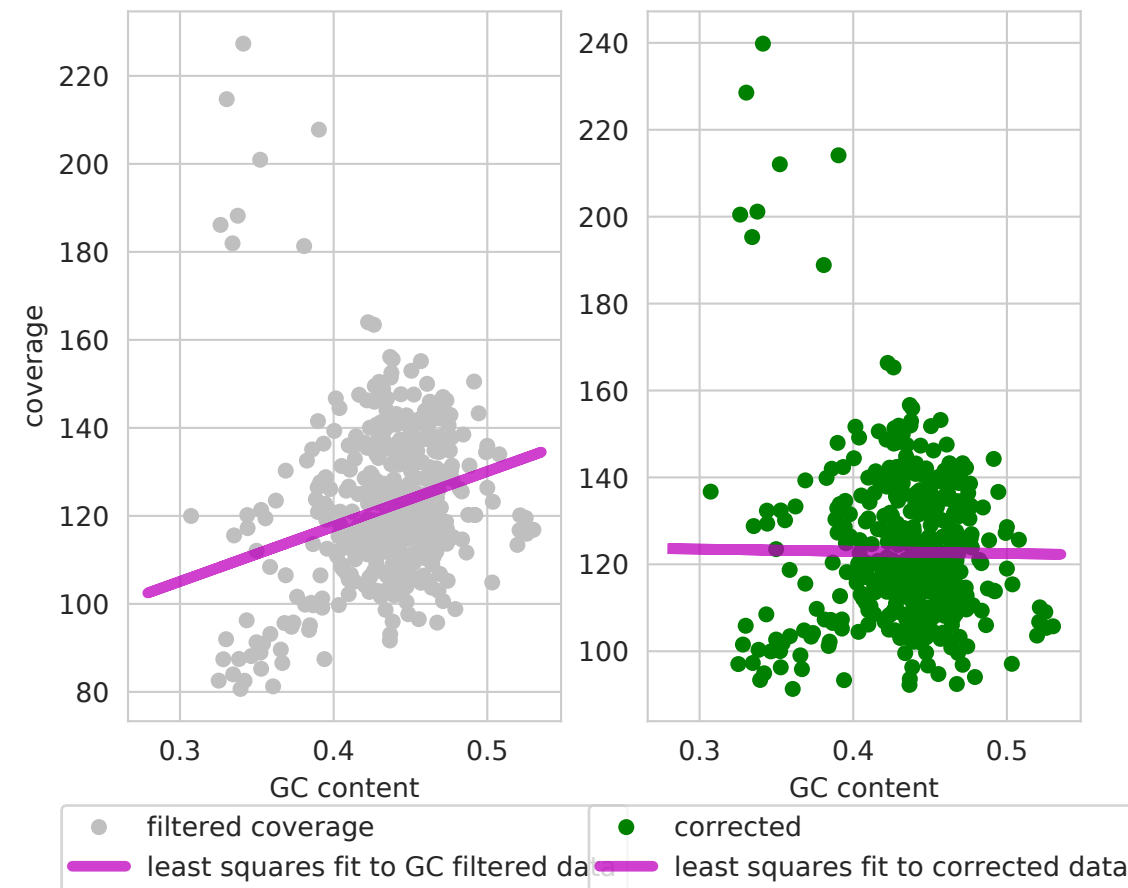


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0812/GCF_005937795_2_ASM593779v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0812-A
iRep: 1.47 r^2 : 0.99 avg. cov: 122.81 % windows: 100.00



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-A
iRep: 1.47 raw iRep: 1.50 GC r^2 : 0.07 GC bias: 9.09 avg. cov: 122.81 % 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-A
iRep: 1.47 un-filtered iRep: 1.47 r^2 : 0.99 avg. cov: 122.81 % windows: 100.00

