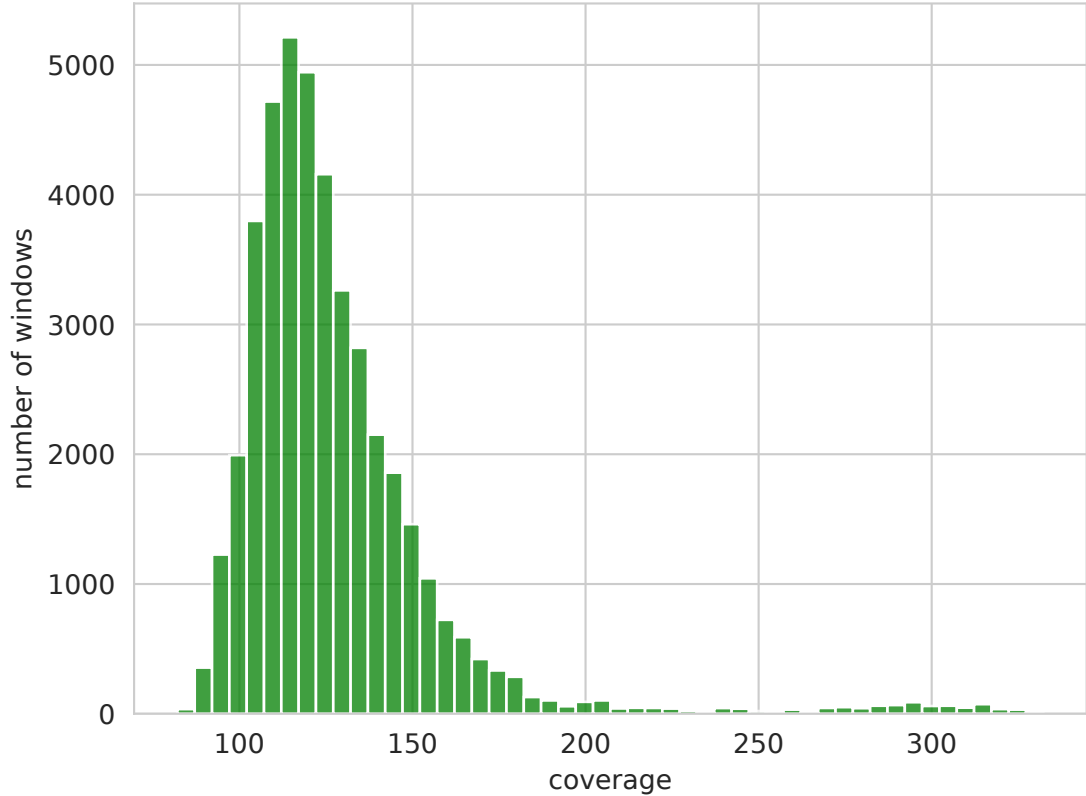
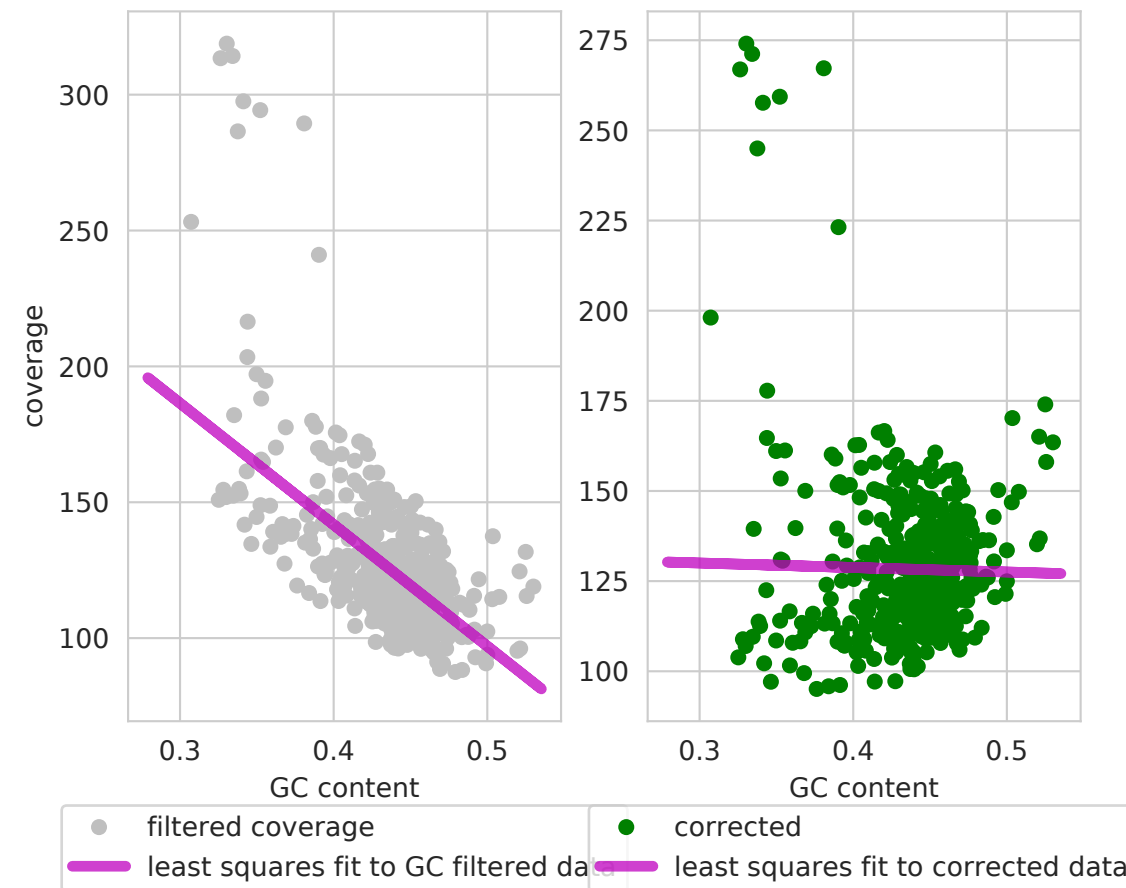


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0812/GCF_005937795_2_ASM593779v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0812-D
iRep: 1.51 r^2 : 0.99 avg. cov: 128.24 % windows: 100.00



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-D
iRep: 1.51 raw iRep: 1.62 GC r^2 : 0.40 GC bias: -178.73 avg. cov: 128.24 % 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-D
iRep: 1.51 un-filtered iRep: 1.51 r^2 : 0.99 avg. cov: 128.24 % windows: 100.00

