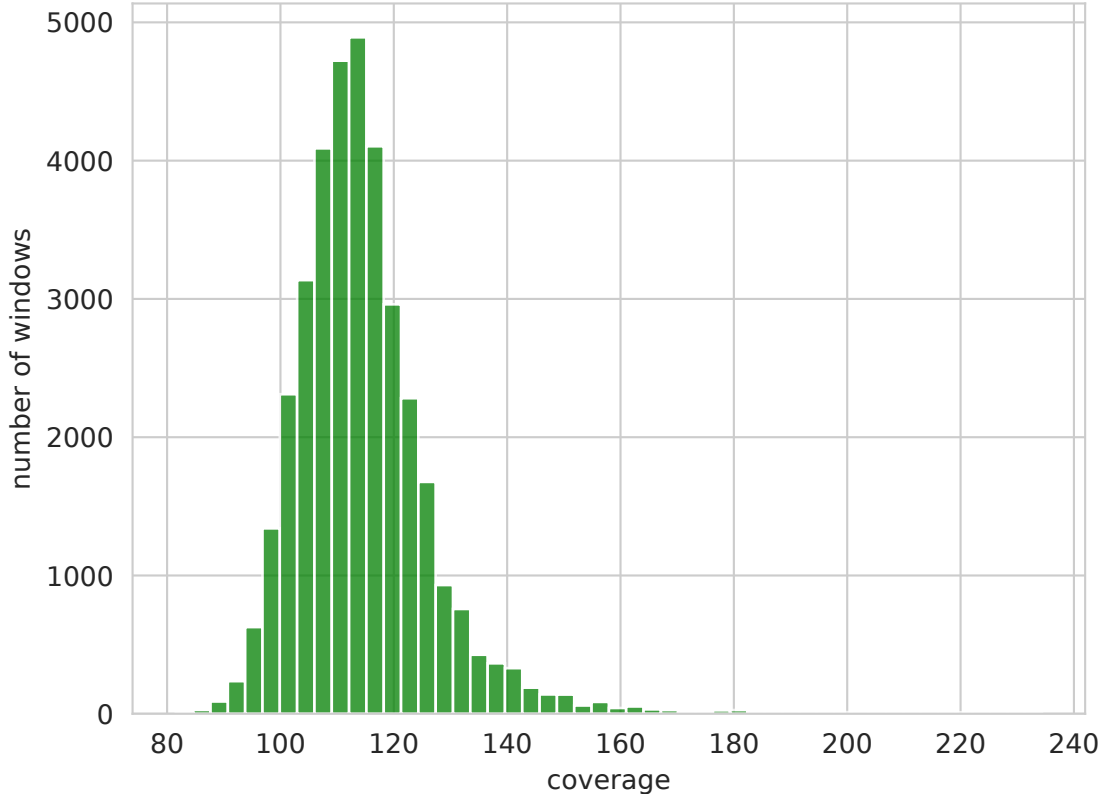genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0702/GCF_005937985.1_ASM593798v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0702-A
iRep: 1.24   r^2: 0.98   avg. cov: 114.92   % windows: 100.00
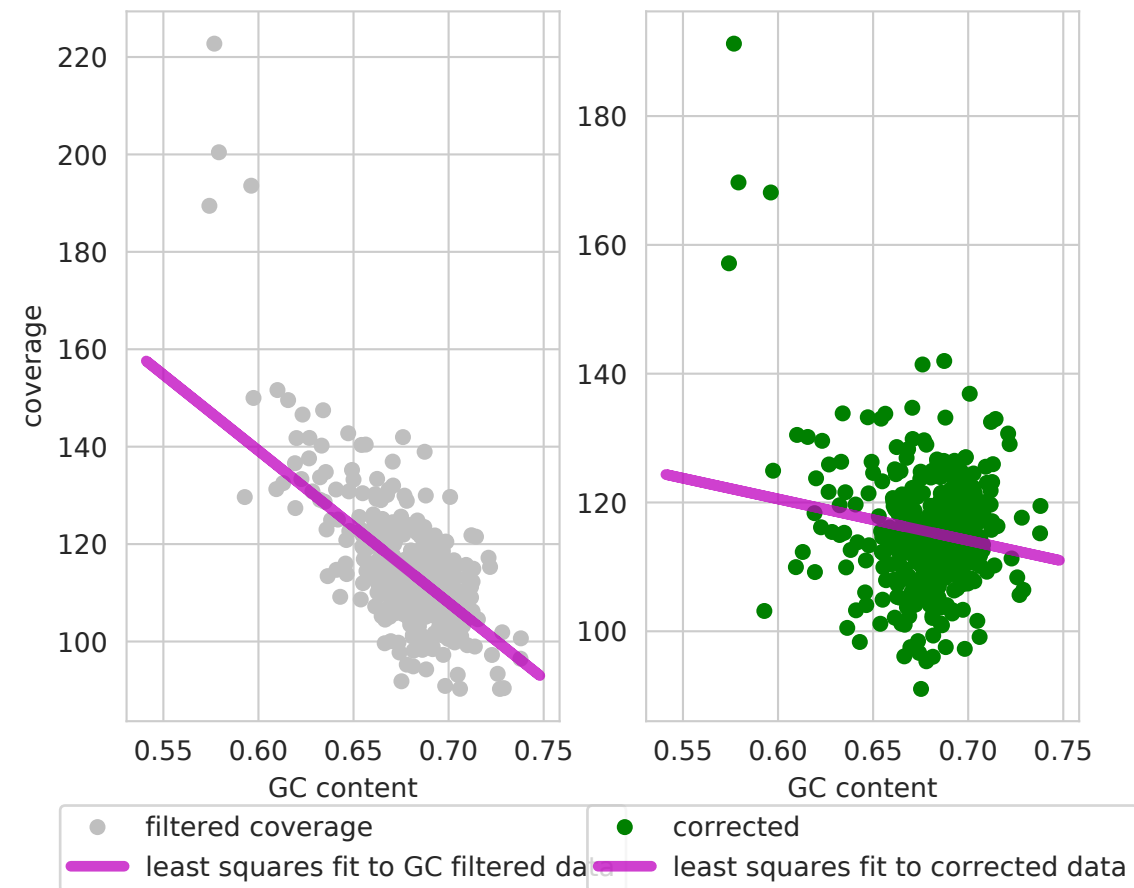
genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0702/GCF_005937985.1_ASM593798v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0702-A
iRep: 1.24   raw iRep: 1.31   GC r^2: 0.46   GC bias: -142.42   avg. cov: 114.92   % windows: 100.00

genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0702/GCF_005937985.1_ASM593798v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0702-A
iRep: 1.24   un-filtered iRep: 1.24   r^2: 0.98   avg. cov: 114.92   % windows: 100.00