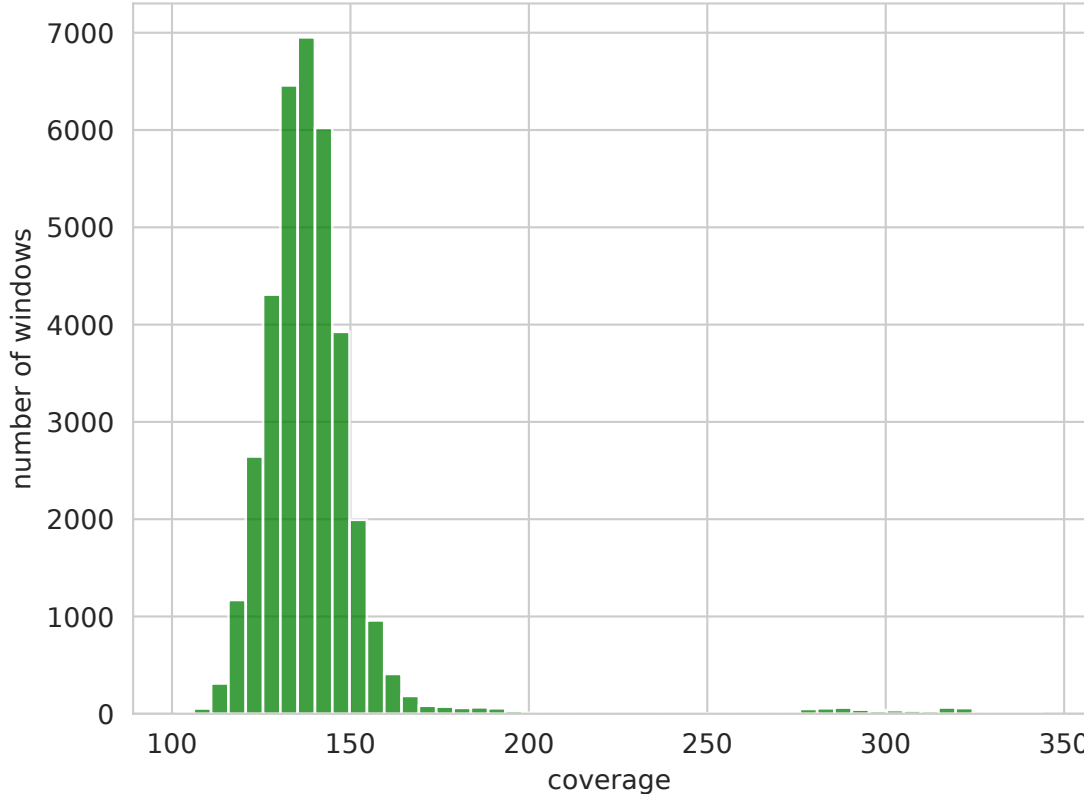
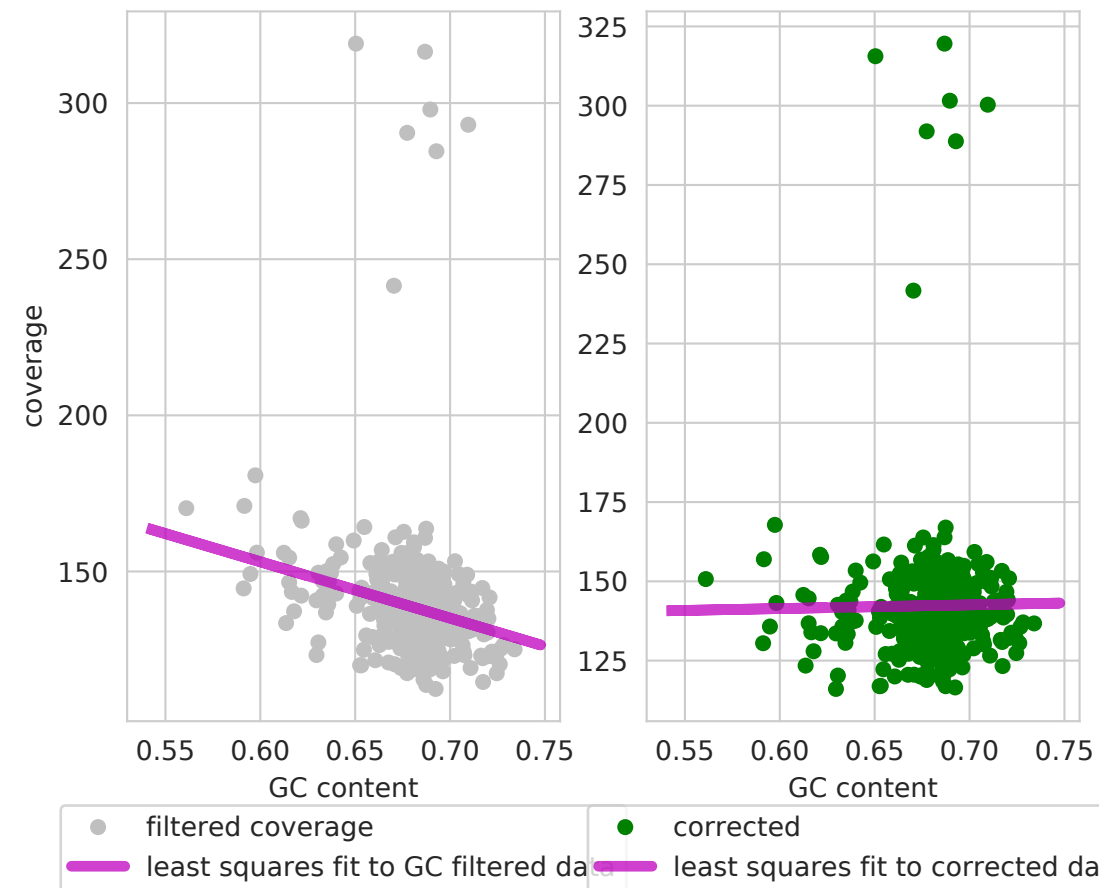


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0702/GCF_005937985_2_ASM593798v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0702-E
iRep: 1.23 r²: 0.98 avg. cov: 140.61 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0702/GCF_005937985_2_ASM593798v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0702-E
iRep: 1.23 raw iRep: 1.26 GC r²: 0.08 GC bias: -14.32 avg. cov: 140.61 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0702/GCF_005937985_2_ASM593798v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0702-E
iRep: 1.23 un-filtered iRep: 1.23 r^2: 0.98 avg. cov: 140.61 % windows: 100.00

