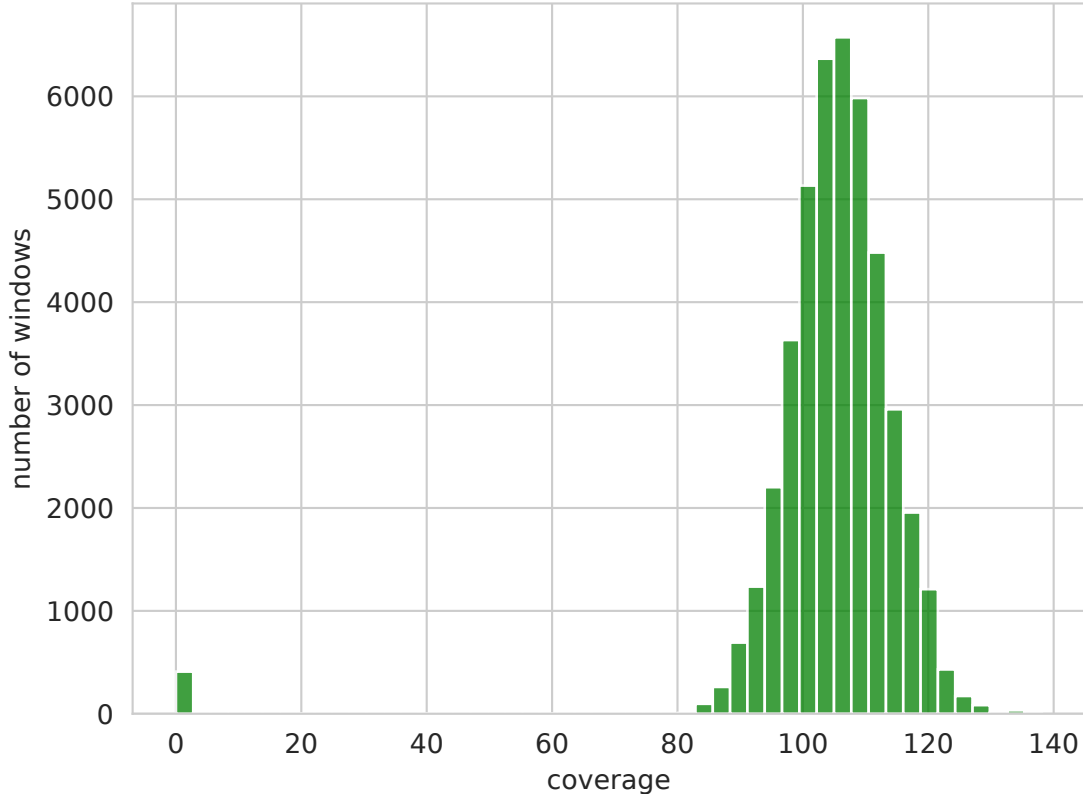
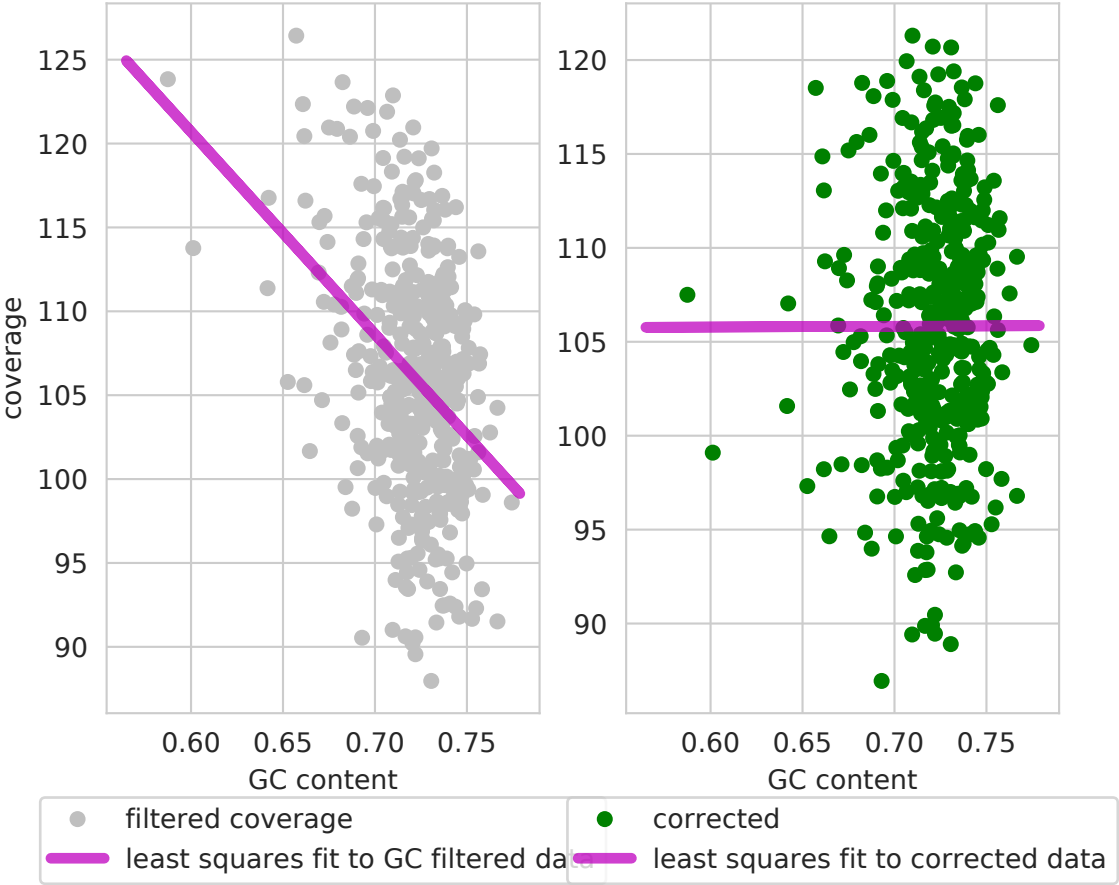


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0722/GCF_005937995.1_ASM593799v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0722-D
iRep: 1.22 r^2 : 0.99 avg. cov: 104.83 % windows: 99.06



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0722/GCF_005937995.1_ASM593799v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0722-D
iRep: 1.22 raw iRep: 1.23 GC r²: 0.14 GC bias: -17.45 avg. cov: 104.83 % windows: 99.06



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0722/GCF_005937995.1_ASM593799v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0722-D
iRep: 1.22 un-filtered iRep: 1.22 r^2 : 0.99 avg. cov: 104.83 % windows: 99.06

