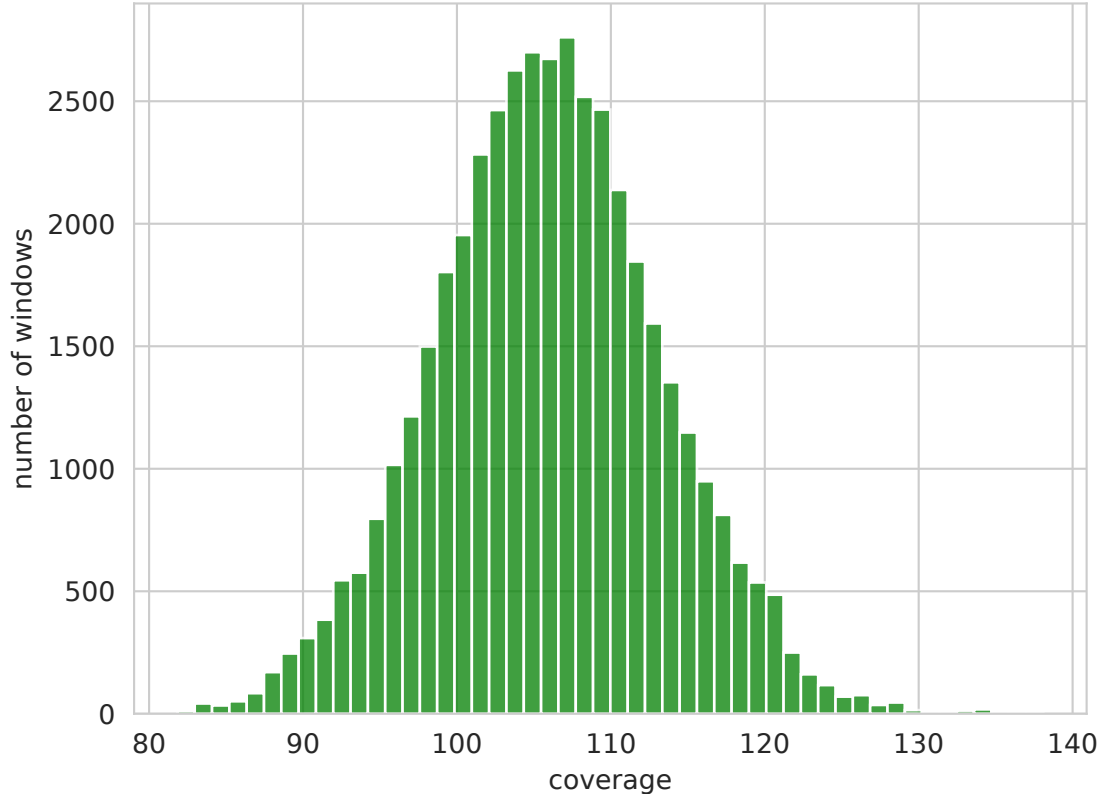
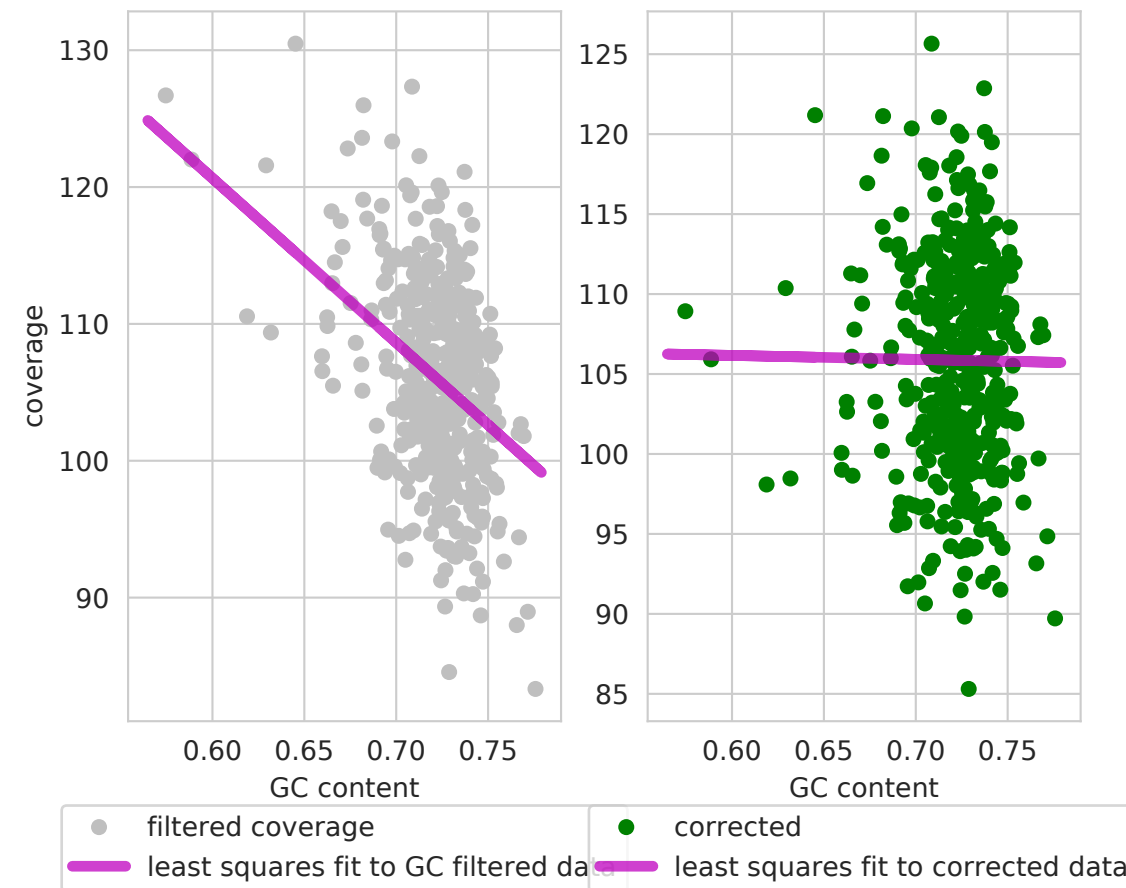


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0722/GCF_005937995_2_ASM593799v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0722-D
iRep: 1.22 r^2 : 0.99 avg. cov: 105.93 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0722/GCF_005937995_2_ASM593799v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0722-D
iRep: 1.22 raw iRep: 1.23 GC r^2 : 0.14 GC bias: -17.25 avg. cov: 105.93 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0722/GCF_005937995.2_ASM593799v2_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: 105.93 % windows: 100.00

