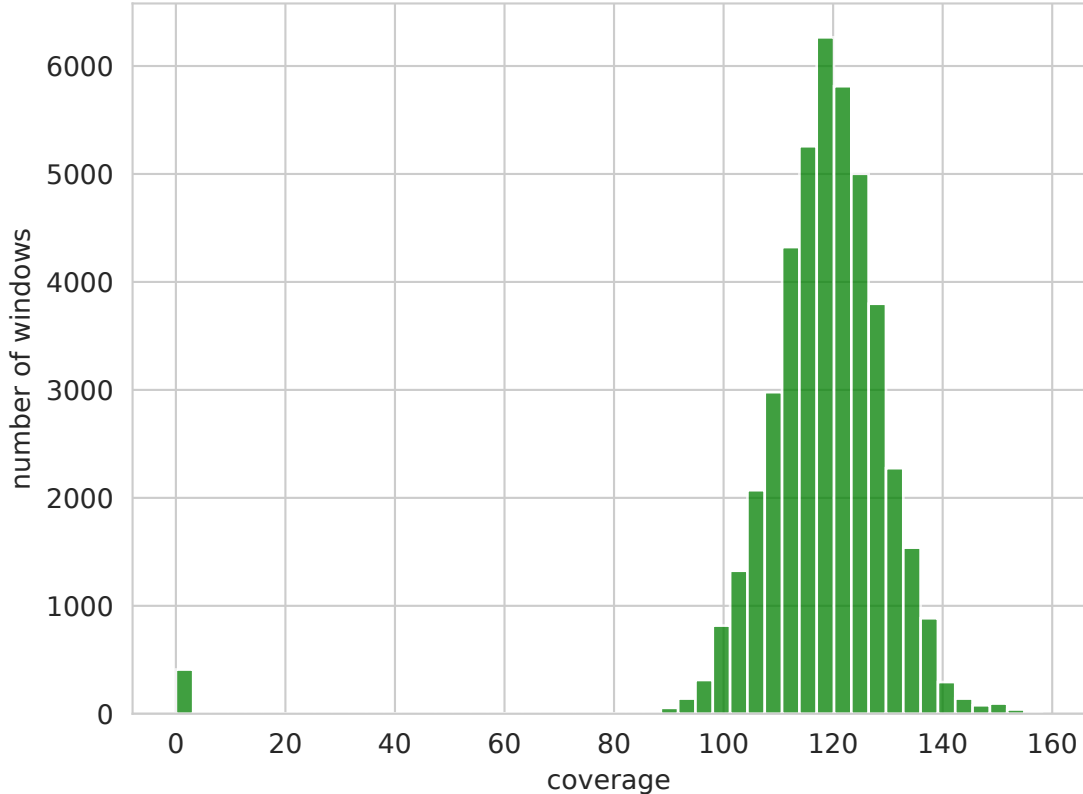
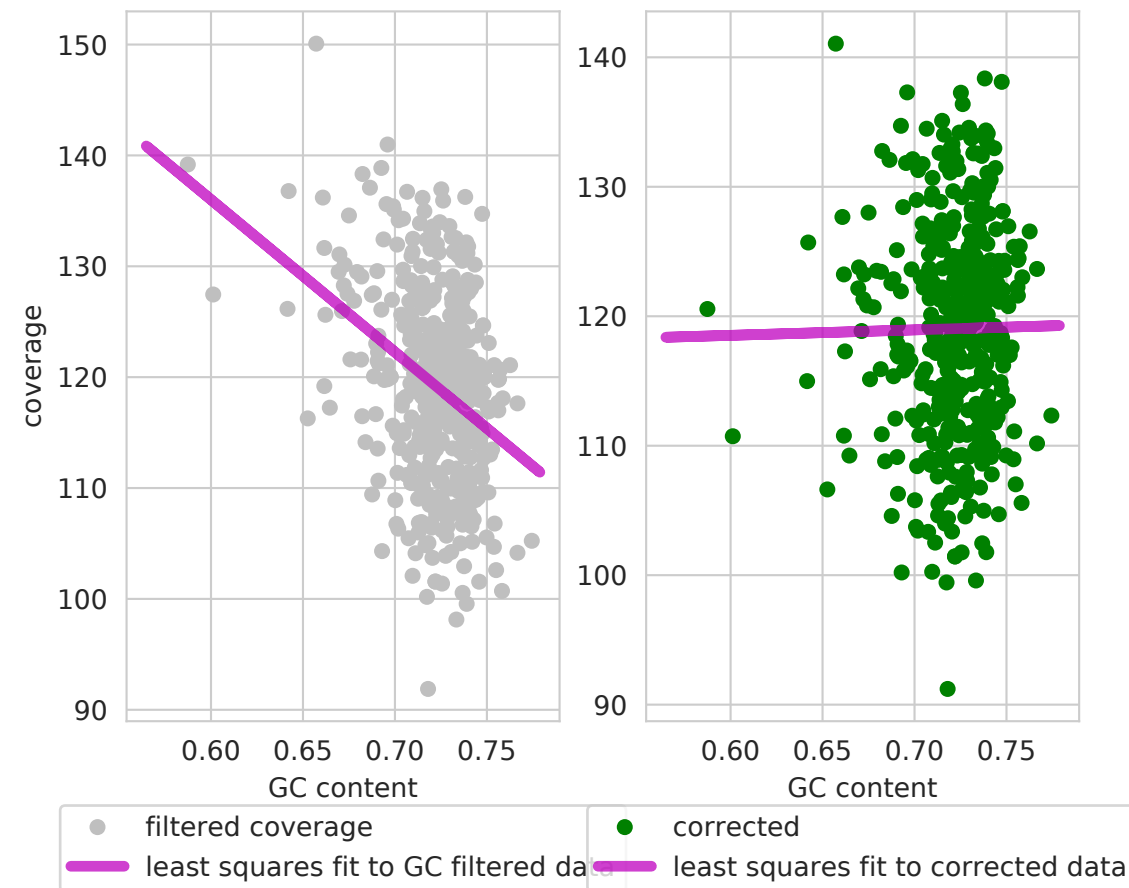


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0722/GCF_005937995.1_ASM593799v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0722-A
iRep: 1.25 r^2 : 0.98 avg. cov: 117.93 % windows: 99.05



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-A
iRep: 1.25 raw iRep: 1.26 GC r^2 : 0.12 GC bias: -16.11 avg. cov: 117.93 % windows: 99.05



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-A
iRep: 1.25 un-filtered iRep: 1.25 r^2 : 0.98 avg. cov: 117.93 % windows: 99.05

