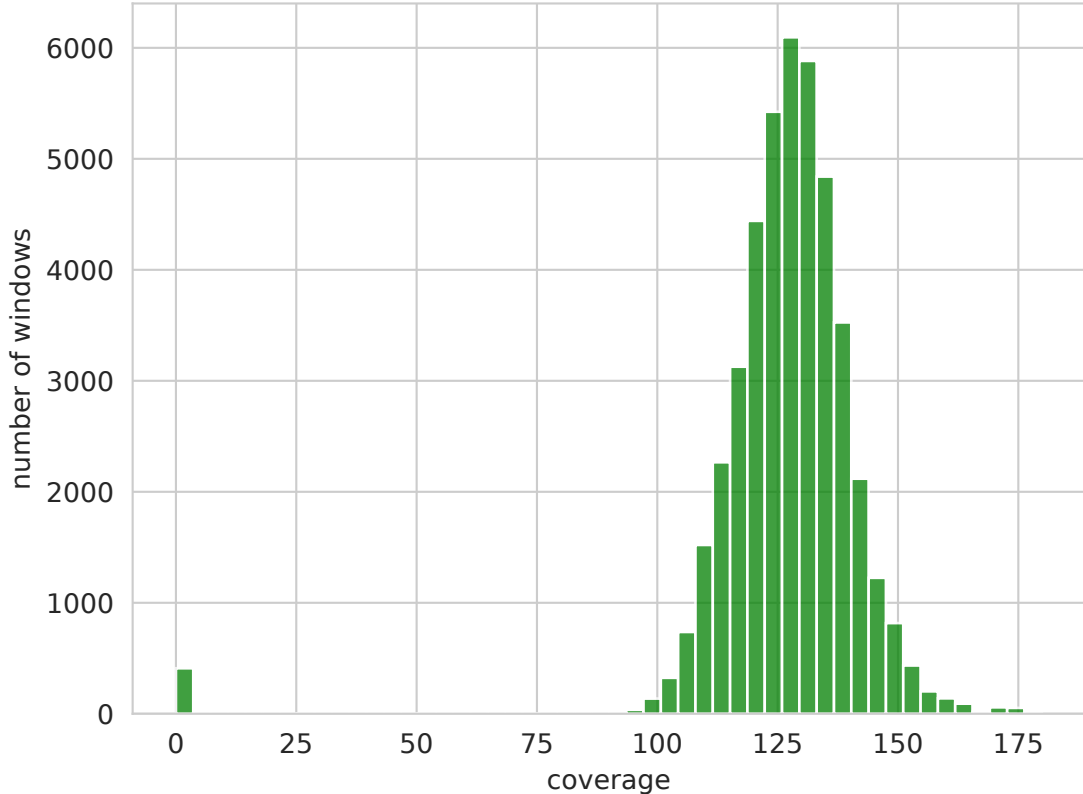
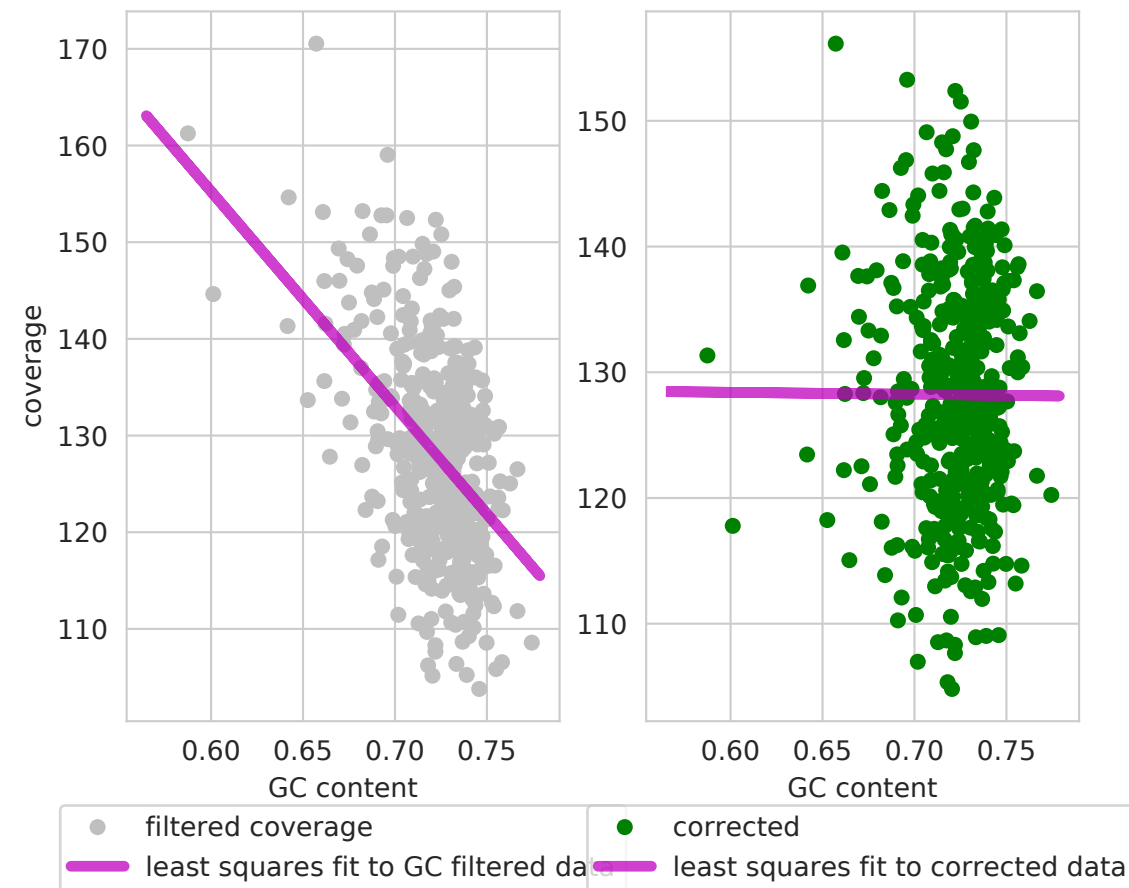


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0722/GCF_005937995.1_ASM593799v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0722-C
iRep: 1.25 r^2 : 0.99 avg. cov: 126.86 % windows: 99.06



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-C
iRep: 1.25 raw iRep: 1.28 GC r^2 : 0.23 GC bias: -50.10 avg. cov: 126.86 % 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-C
iRep: 1.25 un-filtered iRep: 1.25 r^2 : 0.99 avg. cov: 126.86 % windows: 99.06

