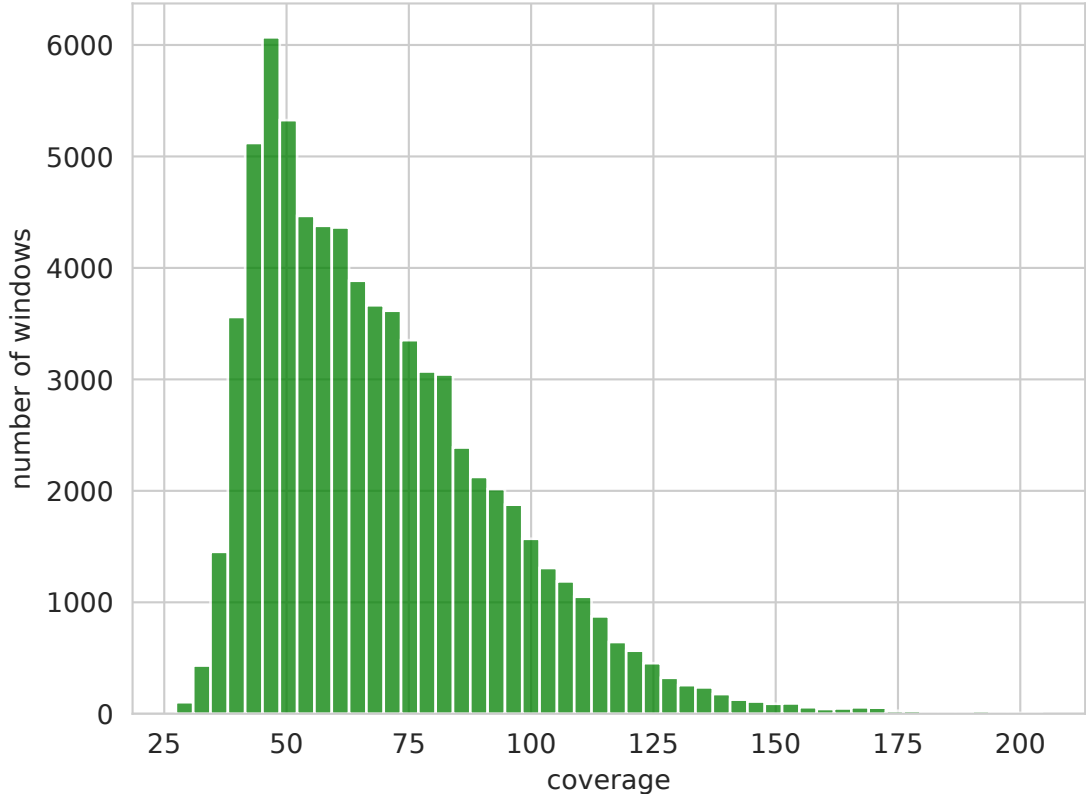
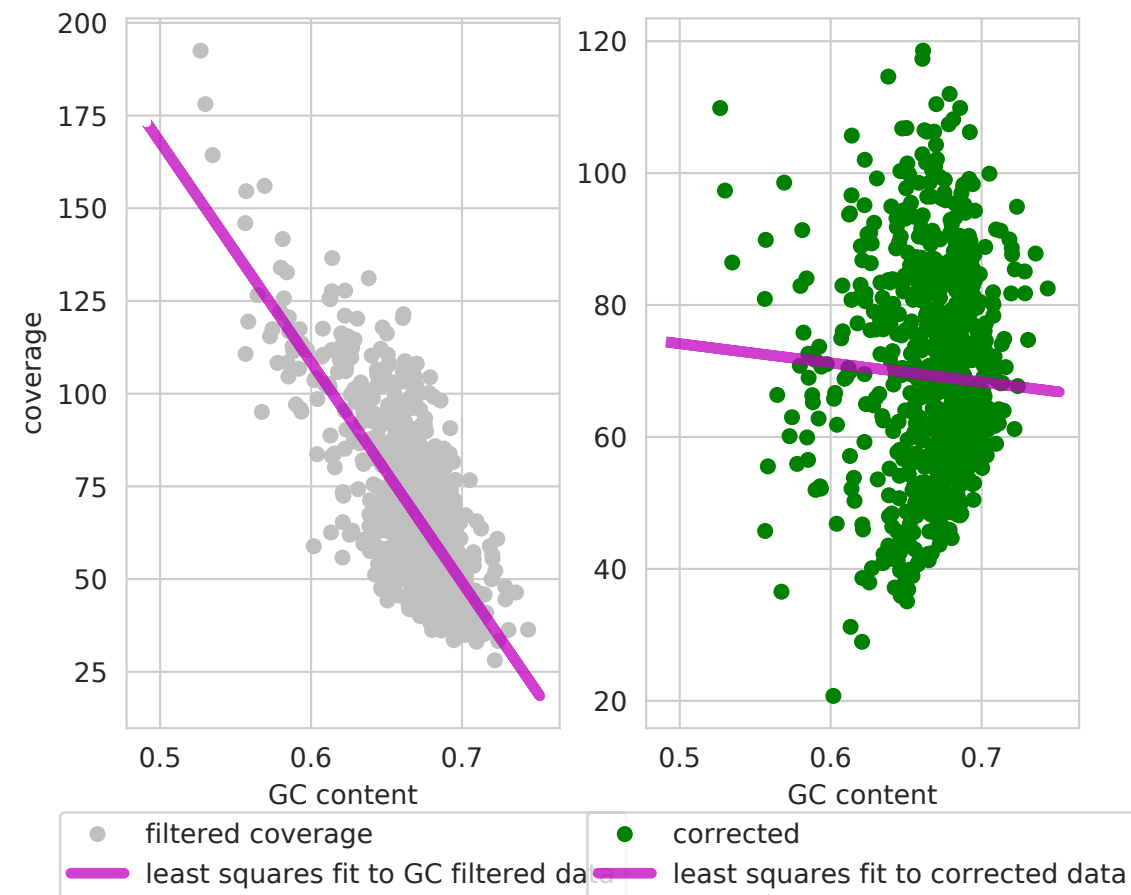


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0801/GCF_005937855.1_ASM593785v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0801-A
iRep: 2.10 r^2 : 0.99 avg. cov: 69.26 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0801/GCF_005937855.1_ASM593785v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0801-A
iRep: 2.10 raw iRep: 2.93 GC r^2 : 0.58 GC bias: -342.92 avg. cov: 69.26 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0801/GCF_005937855.1_ASM593785v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0801-A
iRep: 2.10 un-filtered iRep: 2.10 r^2 : 0.99 avg. cov: 69.26 % windows: 100.00

