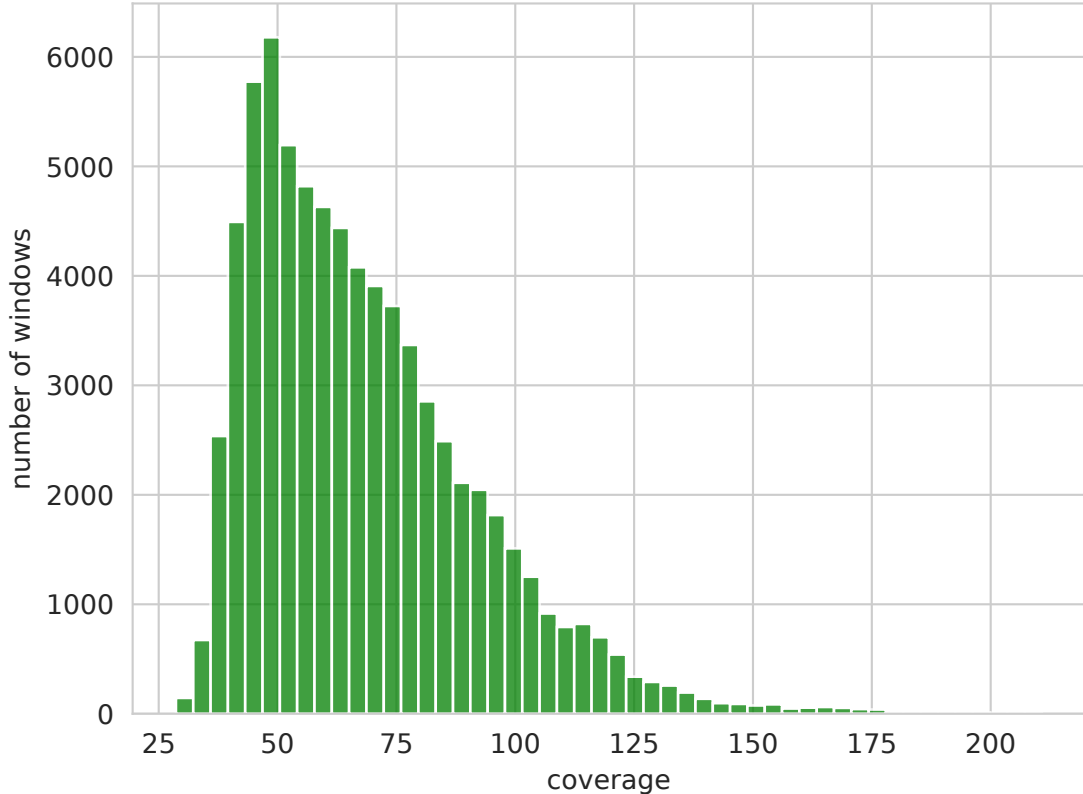
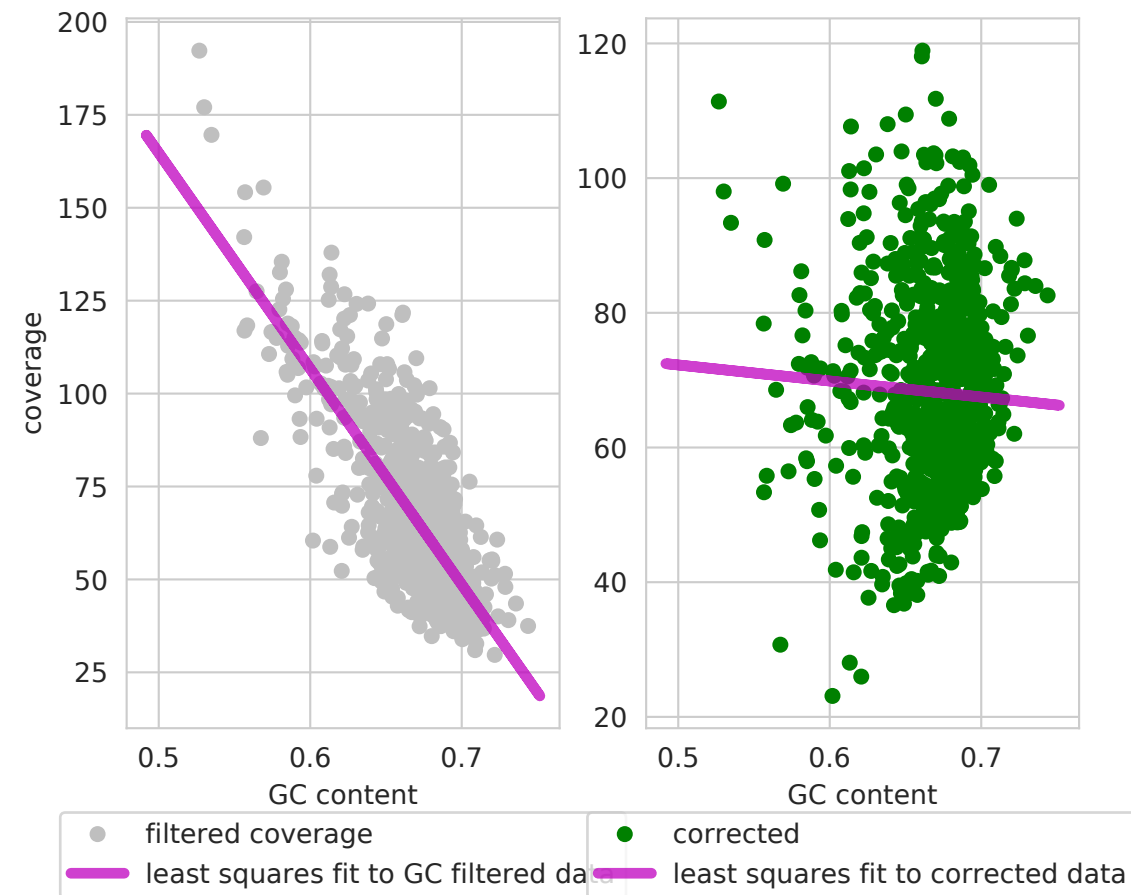


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0801/GCF_005937855.1_ASM593785v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0801-C
iRep: 2.05 r^2 : 0.99 avg. cov: 68.28 % windows: 100.00



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-C
iRep: 2.05 raw iRep: 2.83 GC r²: 0.58 GC bias: -337.73 avg. cov: 68.28 % 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-C
iRep: 2.05 un-filtered iRep: 2.05 r^2 : 0.99 avg. cov: 68.28 % windows: 100.00

