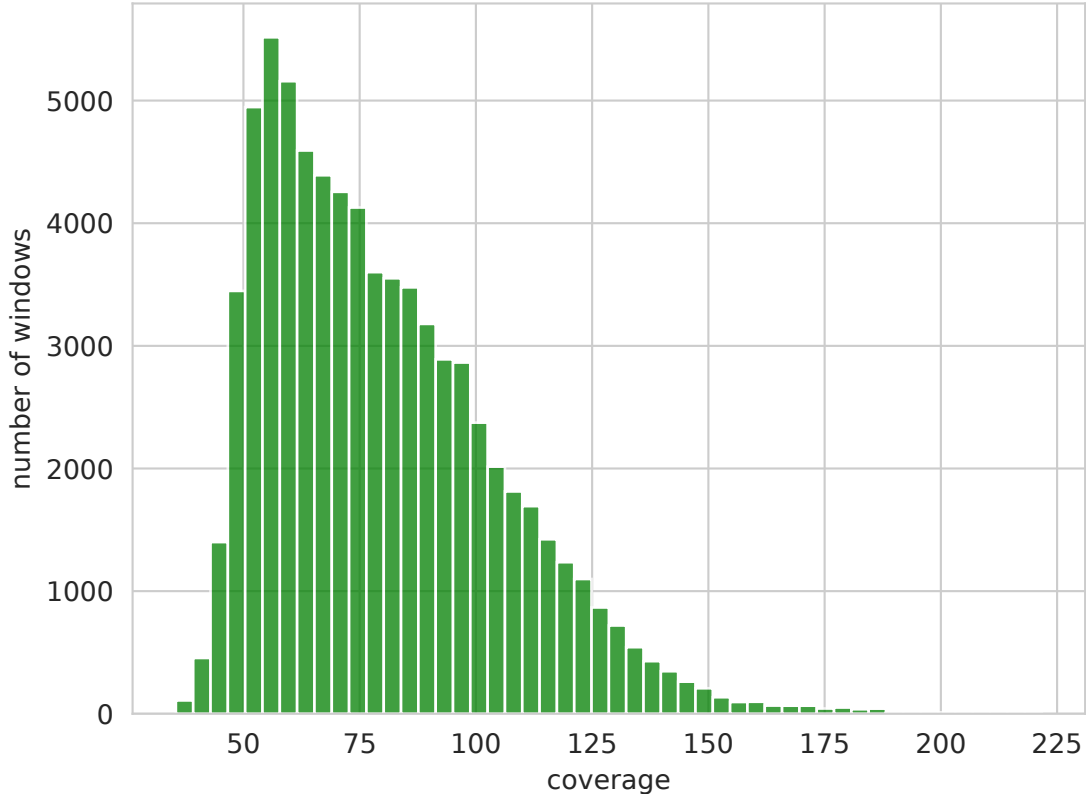
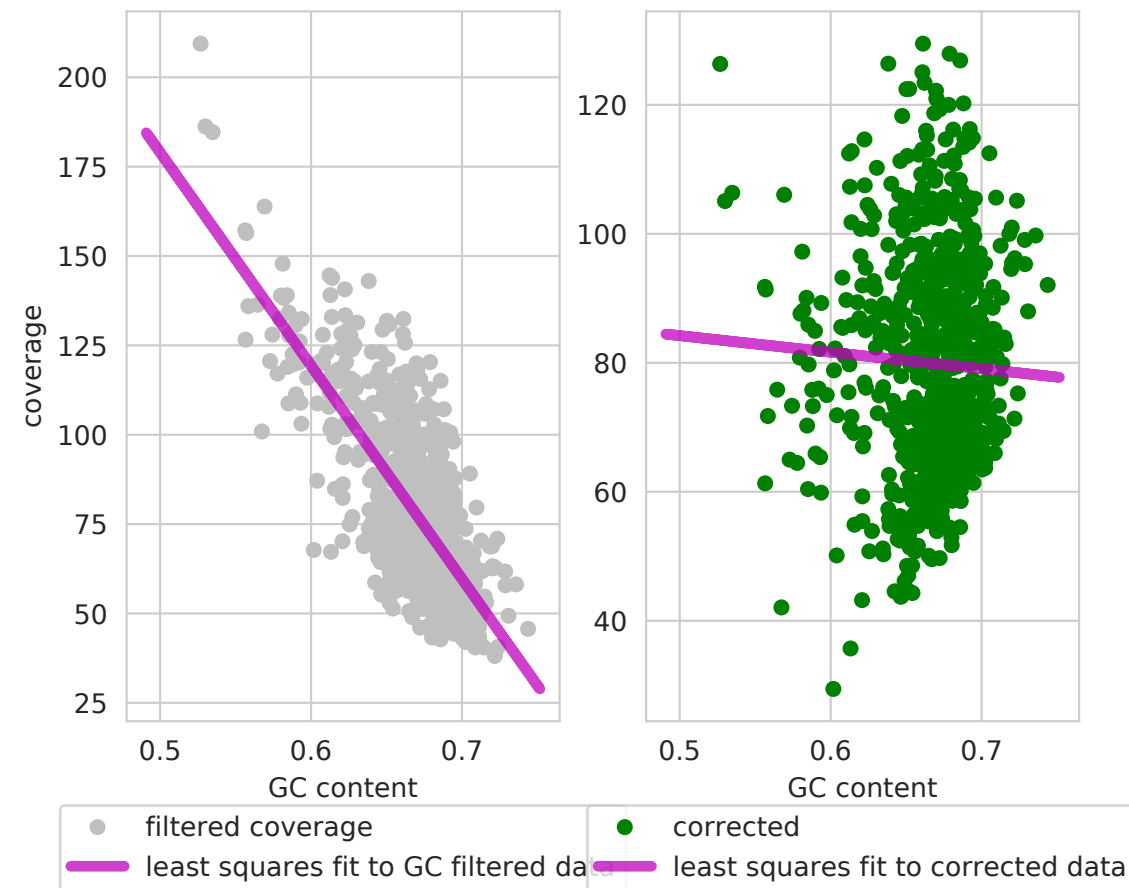


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0801/GCF_005937855.1_ASM593785v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0801-B
iRep: 1.98 r^2 : 0.99 avg. cov: 79.91 % windows: 100.00



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-B
iRep: 1.98 raw iRep: 2.65 GC r^2 : 0.55 GC bias: -326.47 avg. cov: 79.91 % 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-B
iRep: 1.98 un-filtered iRep: 1.98 r^2 : 0.99 avg. cov: 79.91 % windows: 100.00

