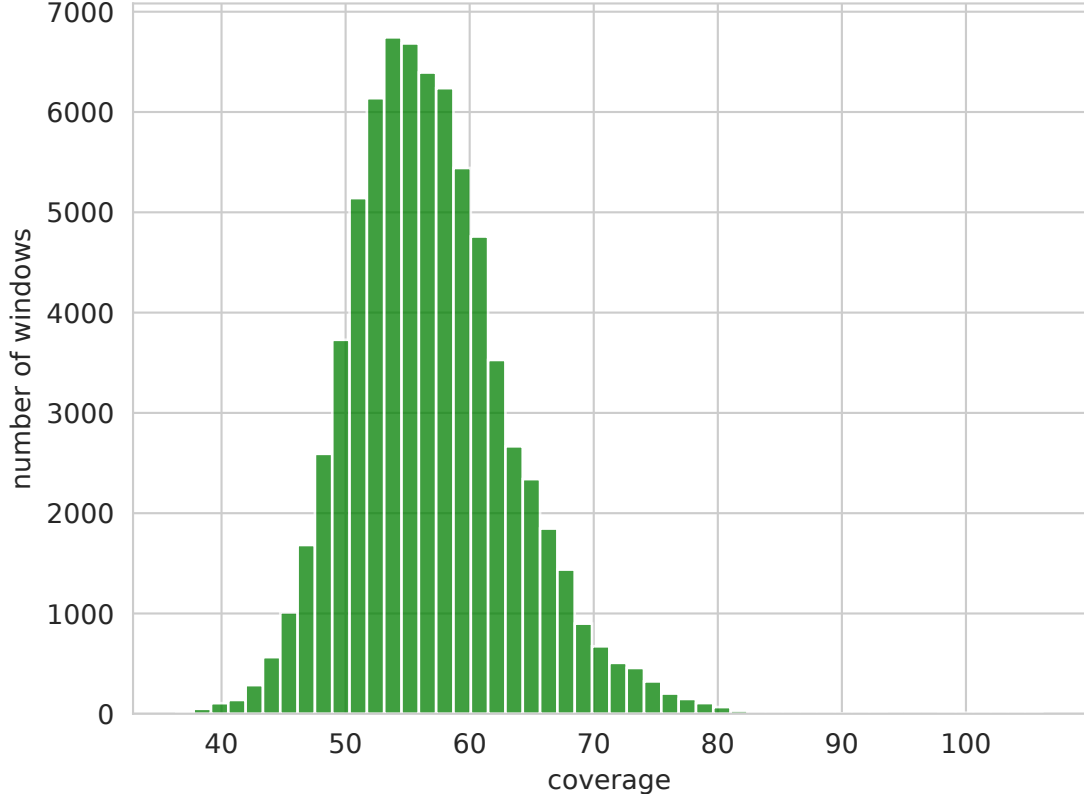
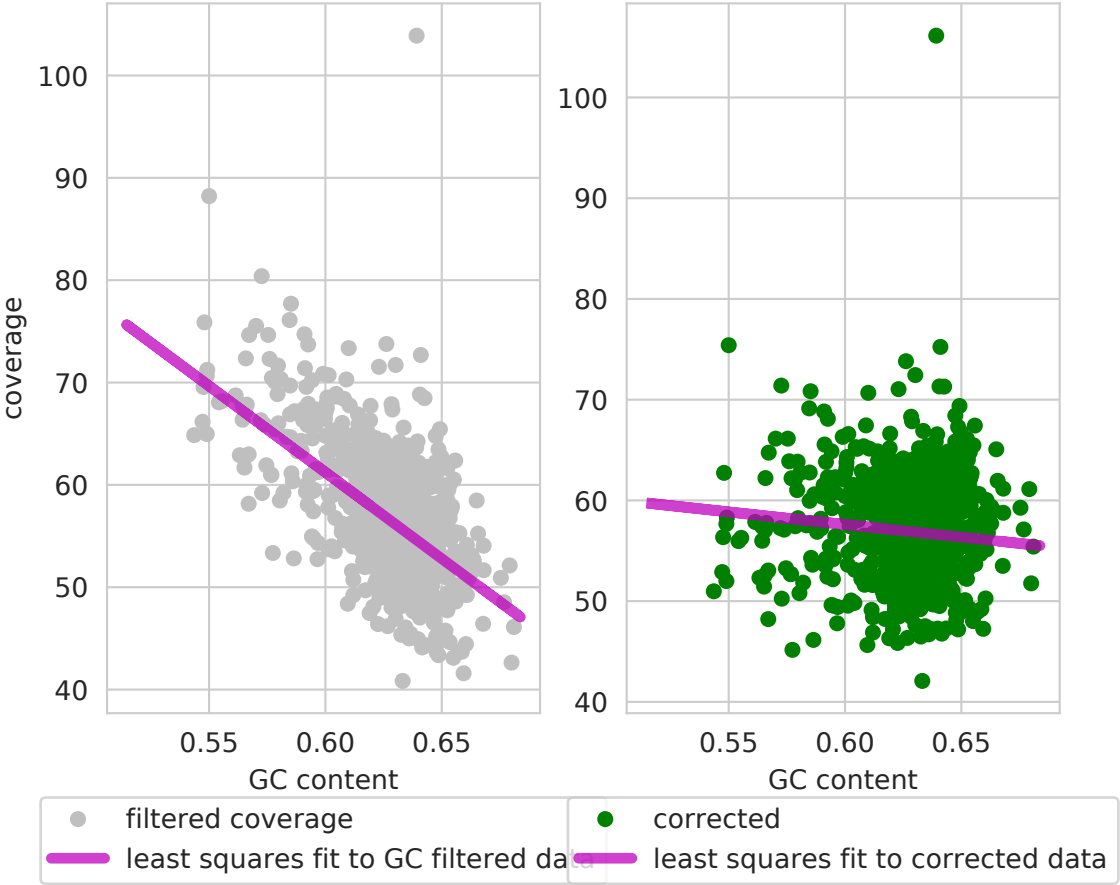


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0727/GCF_005937885_2_ASM593788v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0727-A
iRep: 1.31 r²: 0.99 avg. cov: 56.86 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0727/GCF_005937885_2_ASM593788v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0727-A
iRep: 1.31 raw iRep: 1.39 GC r^2 : 0.38 GC bias: -64.83 avg. cov: 56.86 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0727/GCF_005937885_2_ASM593788v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0727-A
iRep: 1.31 un-filtered iRep: 1.31 r^2 : 0.99 avg. cov: 56.86 % windows: 100.00

