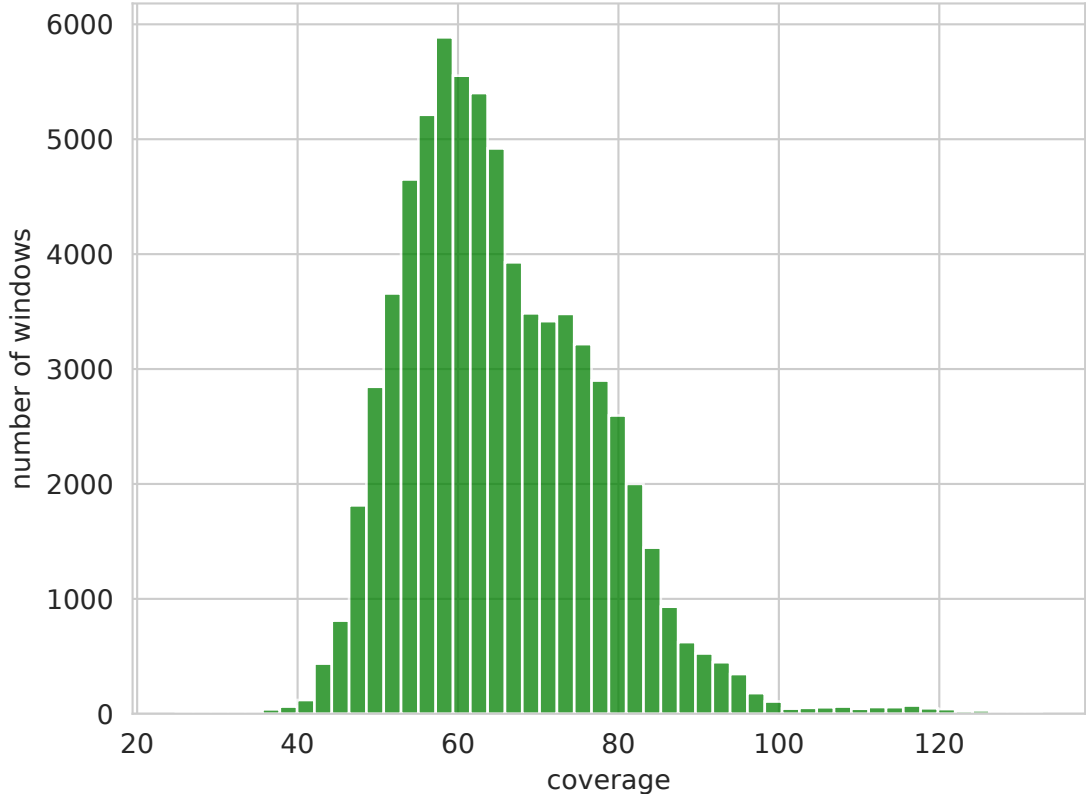
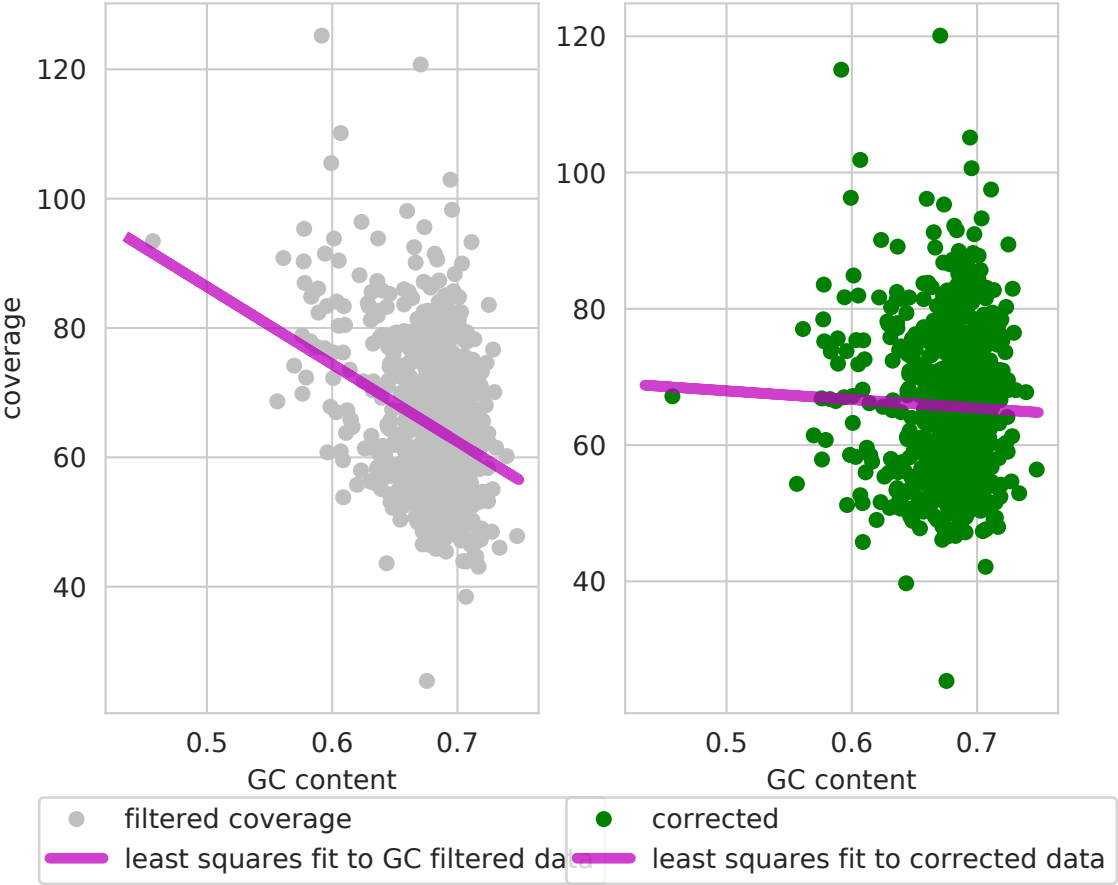


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0712/GCF_006323185.1_ASM632318v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0712-B
iRep: 1.70 r^2 : 1.00 avg. cov: 65.29 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0712/GCF_006323185.1_ASM632318v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0712-B
iRep: 1.70 raw iRep: 1.73 GC r²: 0.11 GC bias: -13.07 avg. cov: 65.29 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0712/GCF_006323185.1_ASM632318v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0712-B
iRep: 1.70 un-filtered iRep: 1.70 r^2 : 1.00 avg. cov: 65.29 % windows: 100.00

