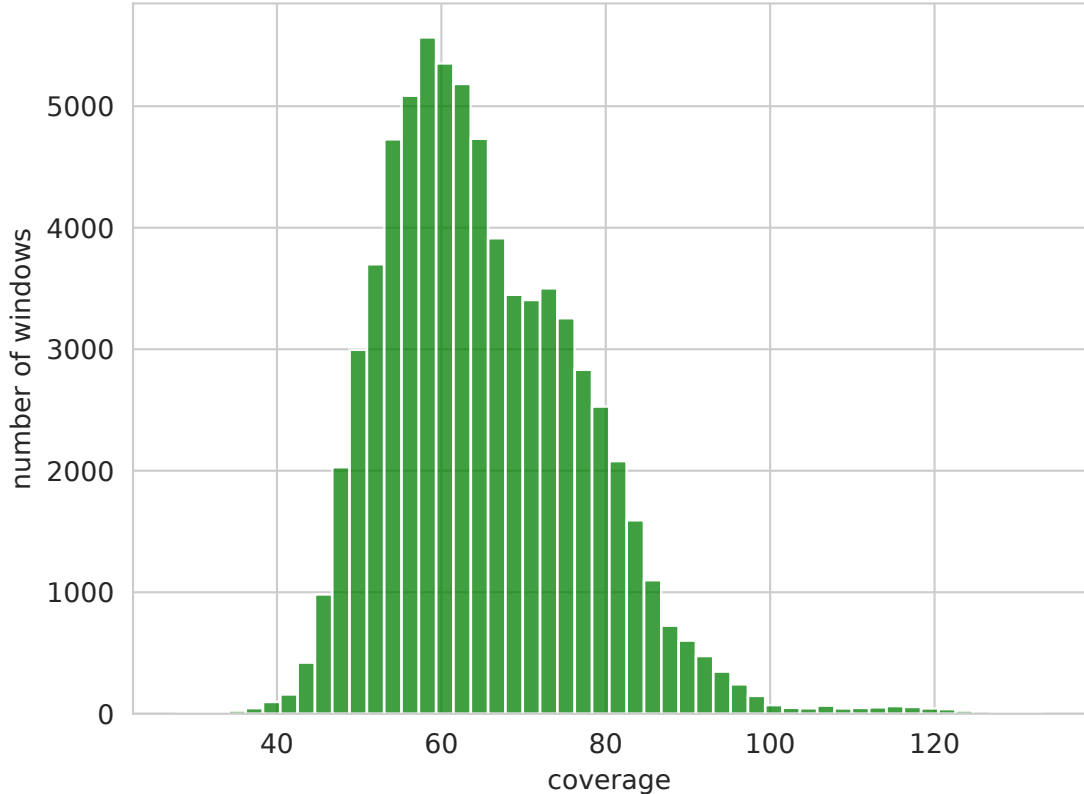
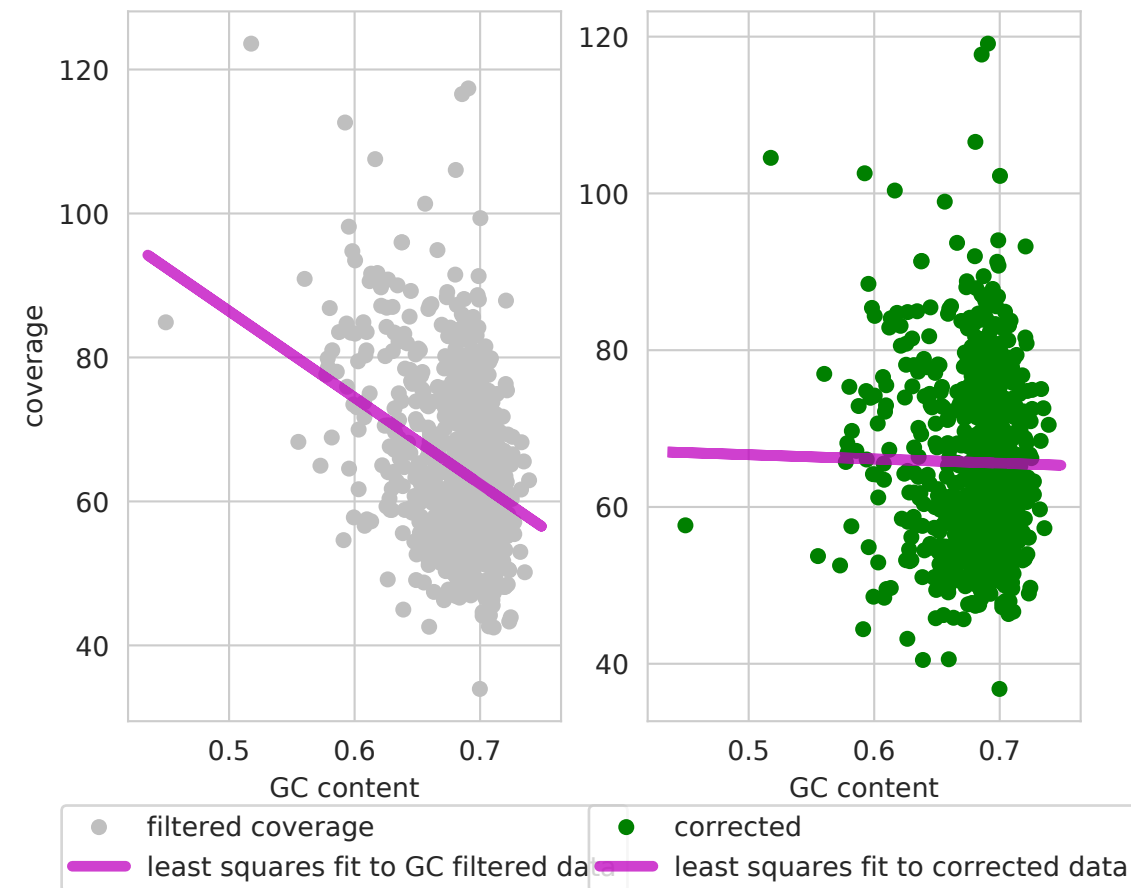


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0712/GCF_006323185_2_ASM632318v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0712-B
iRep: 1.71 r^2 : 1.00 avg. cov: 65.26 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0712/GCF_006323185_2_ASM632318v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0712-B
iRep: 1.71 raw iRep: 1.75 GC r^2 : 0.11 GC bias: -13.17 avg. cov: 65.26 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0712/GCF_006323185_2_ASM632318v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0712-B
iRep: 1.71 un-filtered iRep: 1.71 r^2 : 1.00 avg. cov: 65.26 % windows: 100.00

