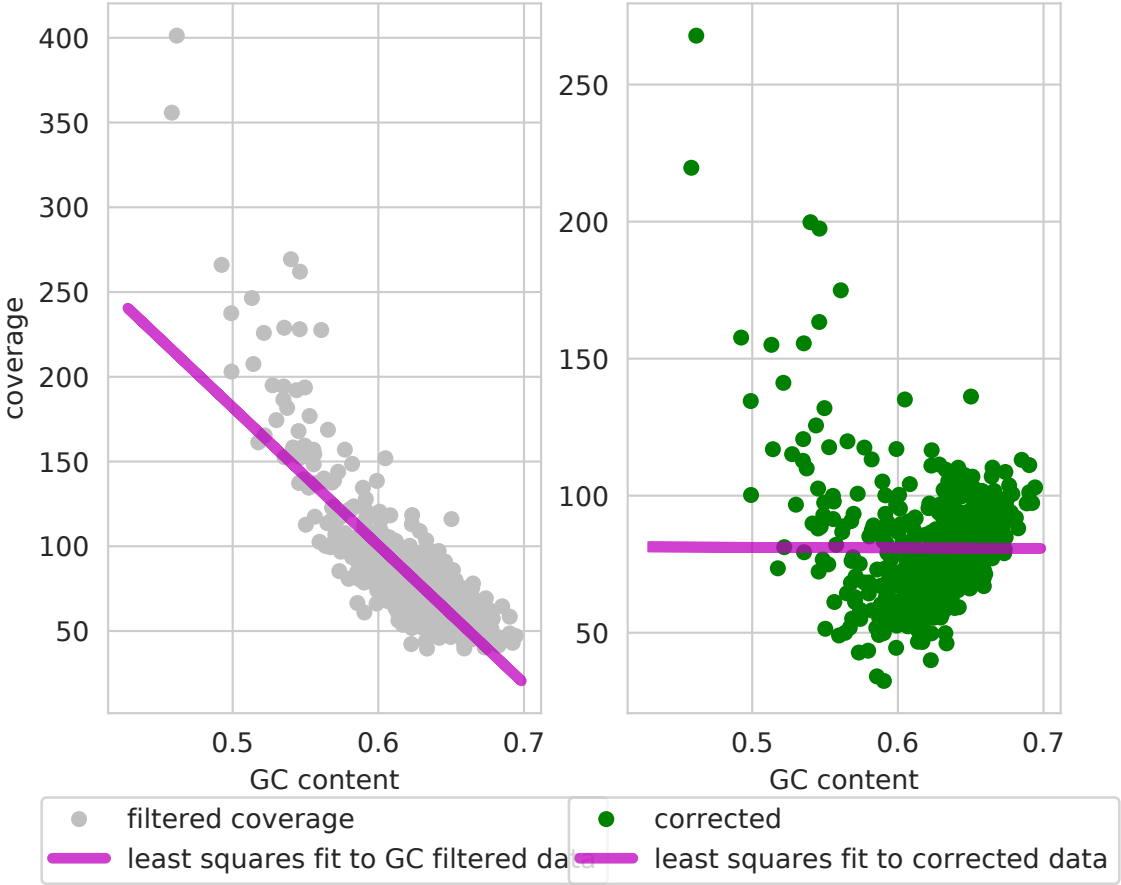


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/ATCC13985/GCF\_006322025\_2\_ASM632202v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/ATCC13985-C  
iRep: 1.79 raw iRep: 2.34 GC r<sup>2</sup>: 0.74 GC bias: -600.92 avg. cov: 79.74 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/ATCC13985/GCF\_006322025\_2\_ASM632202v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/ATCC13985-C  
iRep: 1.79 un-filtered iRep: 1.79  $r^2$ : 0.98 avg. cov: 79.74 % windows: 100.00

