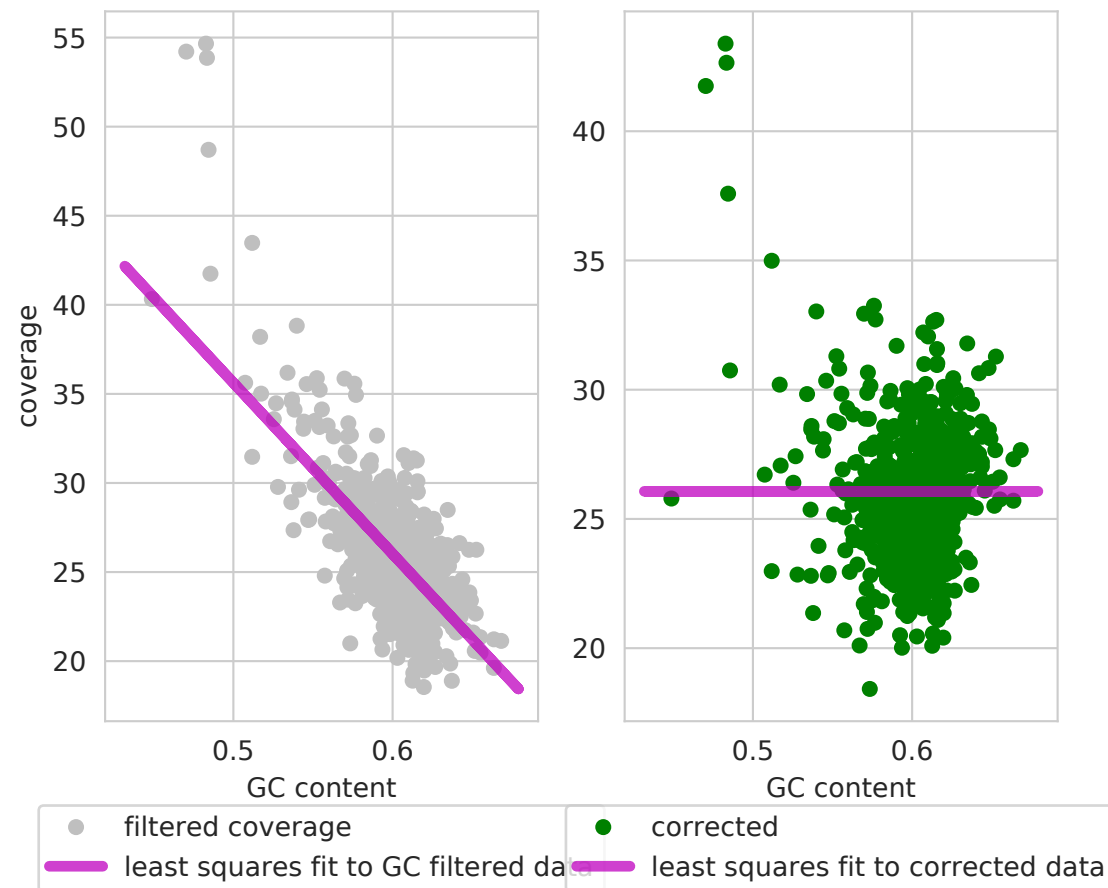


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0710/GCF\_005938045\_2\_ASM593804v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0710-C  
iRep: 1.33 raw iRep: 1.43 GC  $r^2$ : 0.50 GC bias: -47.69 avg. cov: 26.00 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0710/GCF\_005938045\_2\_ASM593804v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0710-C  
iRep: 1.33 un-filtered iRep: 1.33  $r^2$ : 0.99 avg. cov: 26.00 % windows: 100.00

