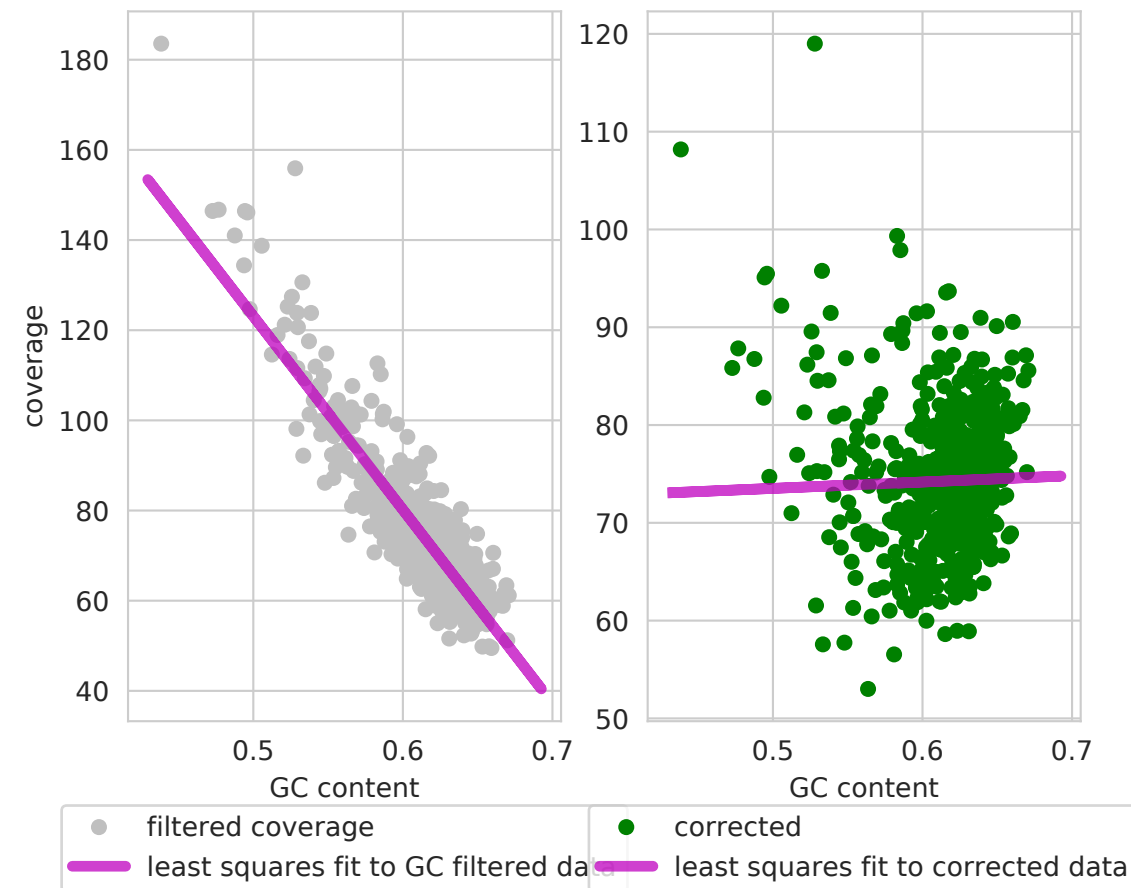


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0802/GCF\_005937845\_2\_ASM593784v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0802-A  
iRep: 1.30 raw iRep: 1.63 GC  $r^2$ : 0.82 GC bias: -349.94 avg. cov: 74.20 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0802/GCF\_005937845\_2\_ASM593784v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0802-A  
iRep: 1.30 un-filtered iRep: 1.30  $r^2$ : 0.98 avg. cov: 74.20 % windows: 100.00

