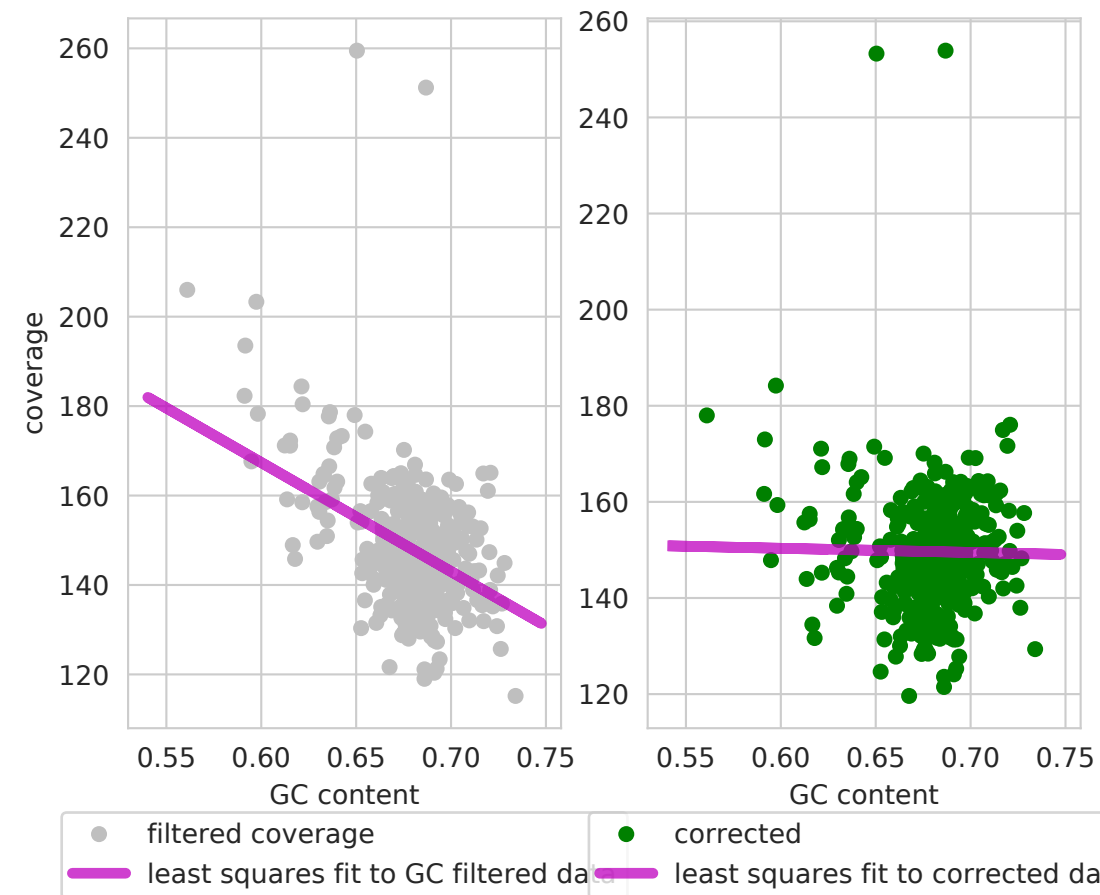


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0702/GCF\_005937985\_2\_ASM593798v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0702-D  
iRep: 1.23 raw iRep: 1.26 GC  $r^2$ : 0.26 GC bias: -62.42 avg. cov: 148.92 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0702/GCF\_005937985\_2\_ASM593798v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0702-D  
iRep: 1.23 un-filtered iRep: 1.23  $r^2$ : 0.99 avg. cov: 148.92 % windows: 100.00

