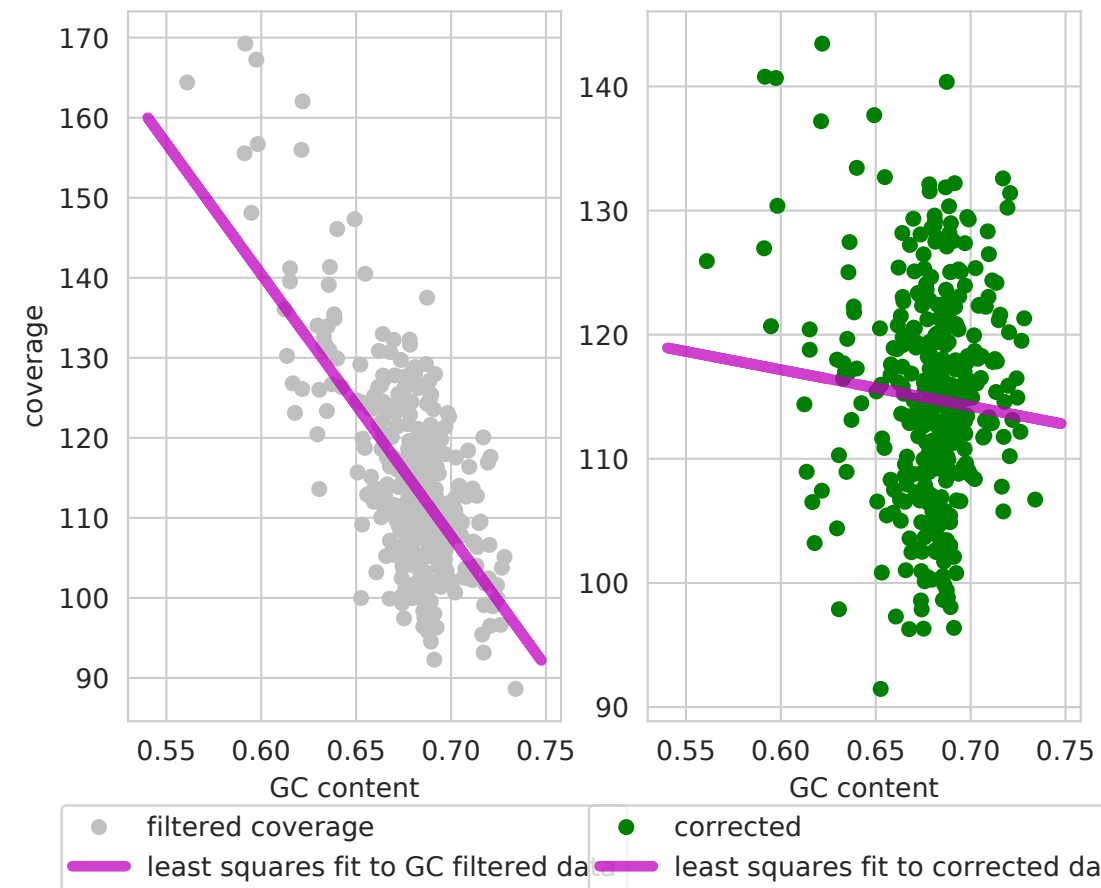


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0702/GCF\_005937985\_2\_ASM593798v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0702-A  
iRep: 1.24 raw iRep: 1.31 GC  $r^2$ : 0.50 GC bias: -164.14 avg. cov: 114.77 % windows: 100.00



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-A  
iRep: 1.24 un-filtered iRep: 1.24 r^2: 0.98 avg. cov: 114.77 % windows: 100.00

