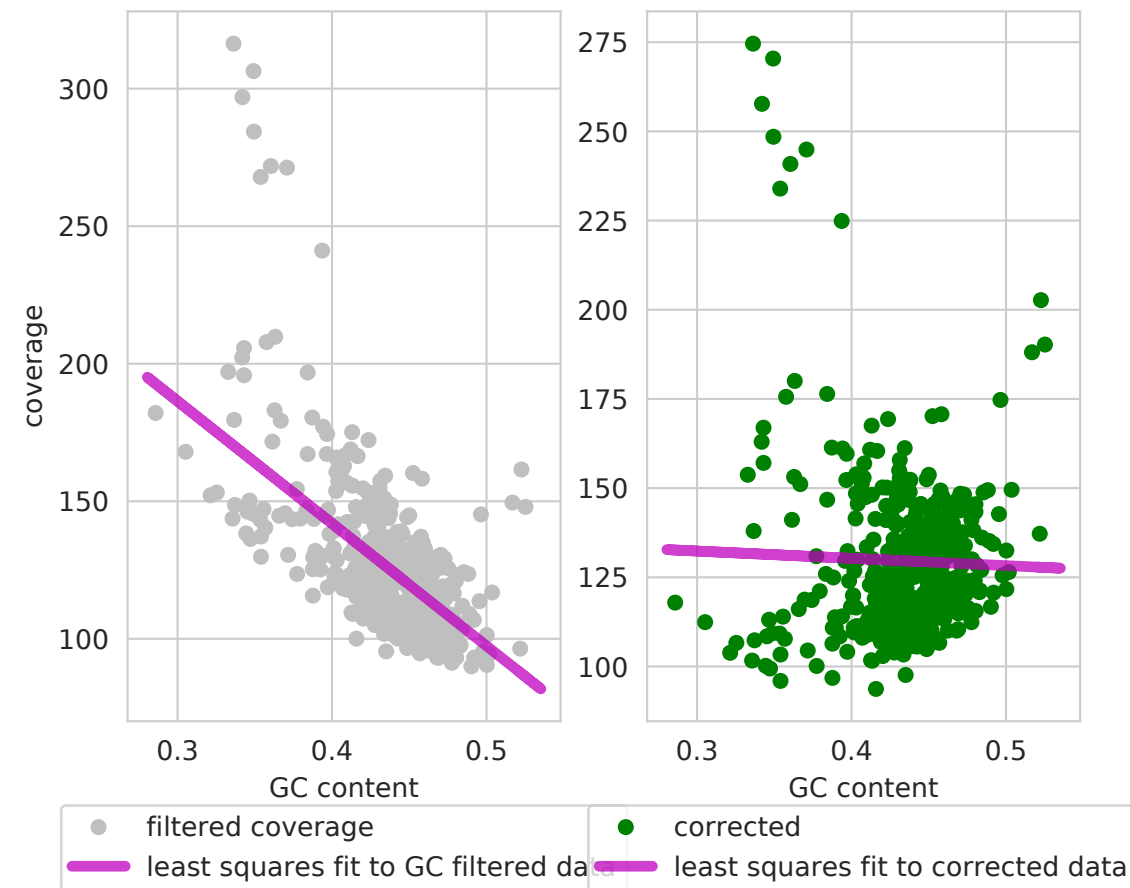


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0812/GCF\_005937795.1\_ASM593779v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0812-D  
iRep: 1.51 raw iRep: 1.63 GC  $r^2$ : 0.38 GC bias: -168.41 avg. cov: 128.52 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0812/GCF\_005937795.1\_ASM593779v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0812-D  
iRep: 1.51 un-filtered iRep: 1.51  $r^2$ : 0.99 avg. cov: 128.52 % windows: 100.00

