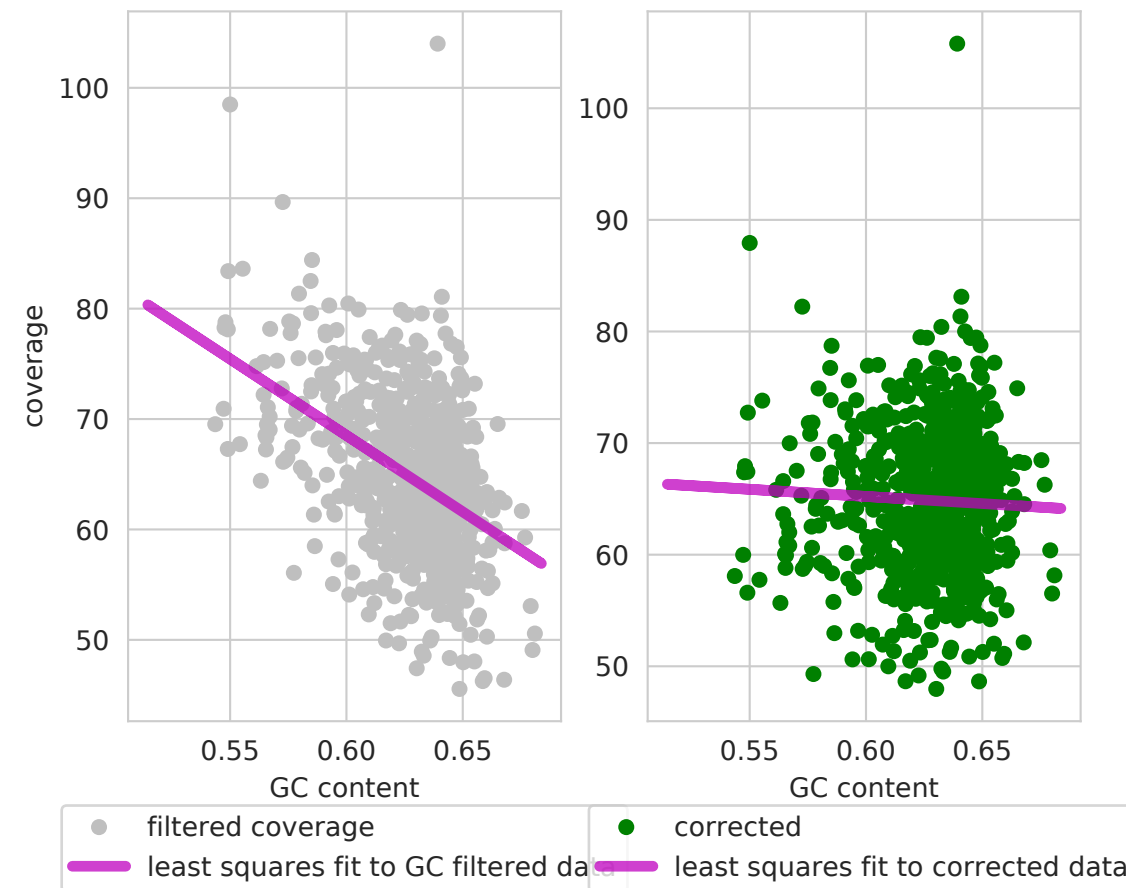


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0727/GCF\_005937885\_2\_ASM593788v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0727-D  
iRep: 1.34 raw iRep: 1.39 GC  $r^2$ : 0.22 GC bias: -30.28 avg. cov: 64.89 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0727/GCF\_005937885\_2\_ASM593788v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0727-D  
iRep: 1.34 un-filtered iRep: 1.34  $r^2$ : 0.99 avg. cov: 64.89 % windows: 100.00

