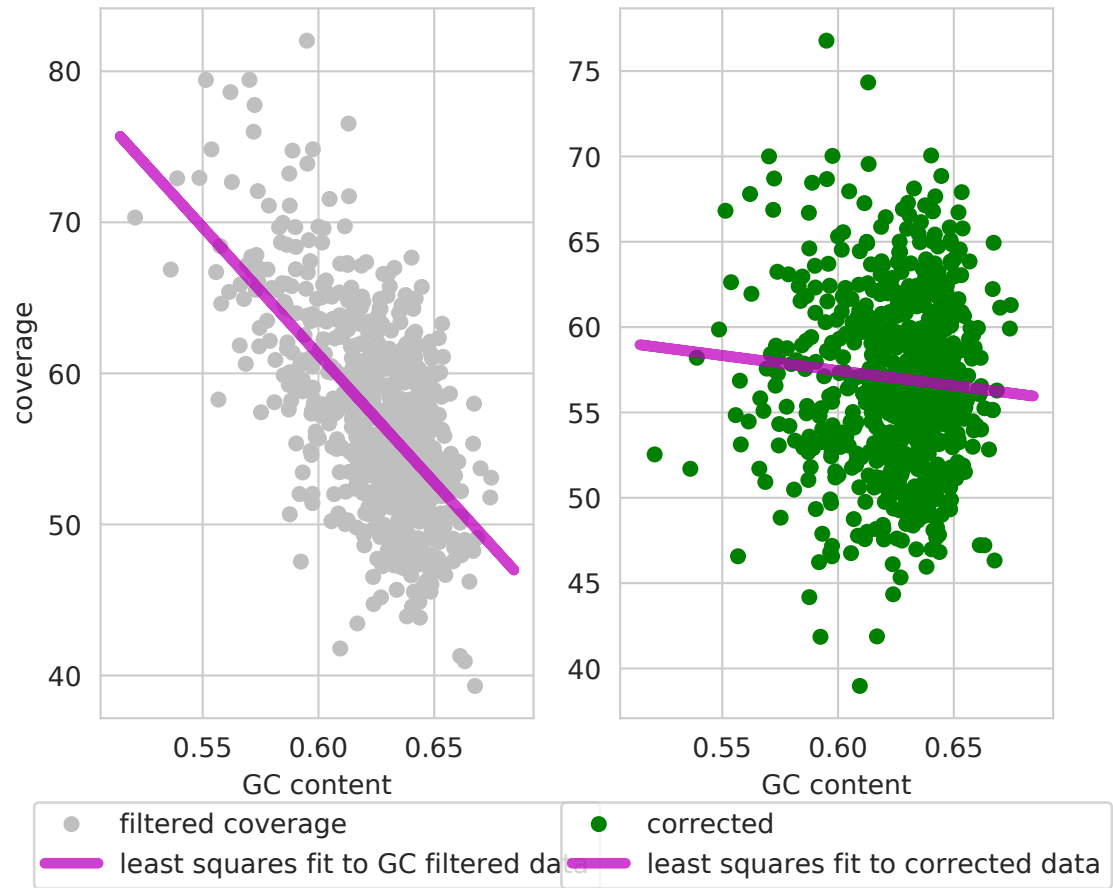


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0727/GCF_005937885.1_ASM593788v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0727-A
iRep: 1.31 raw iRep: 1.39 GC r²: 0.38 GC bias: -64.99 avg. cov: 56.82 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0727/GCF_005937885.1_ASM593788v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0727-A
iRep: 1.31 un-filtered iRep: 1.31 r^2 : 0.99 avg. cov: 56.82 % windows: 100.00

