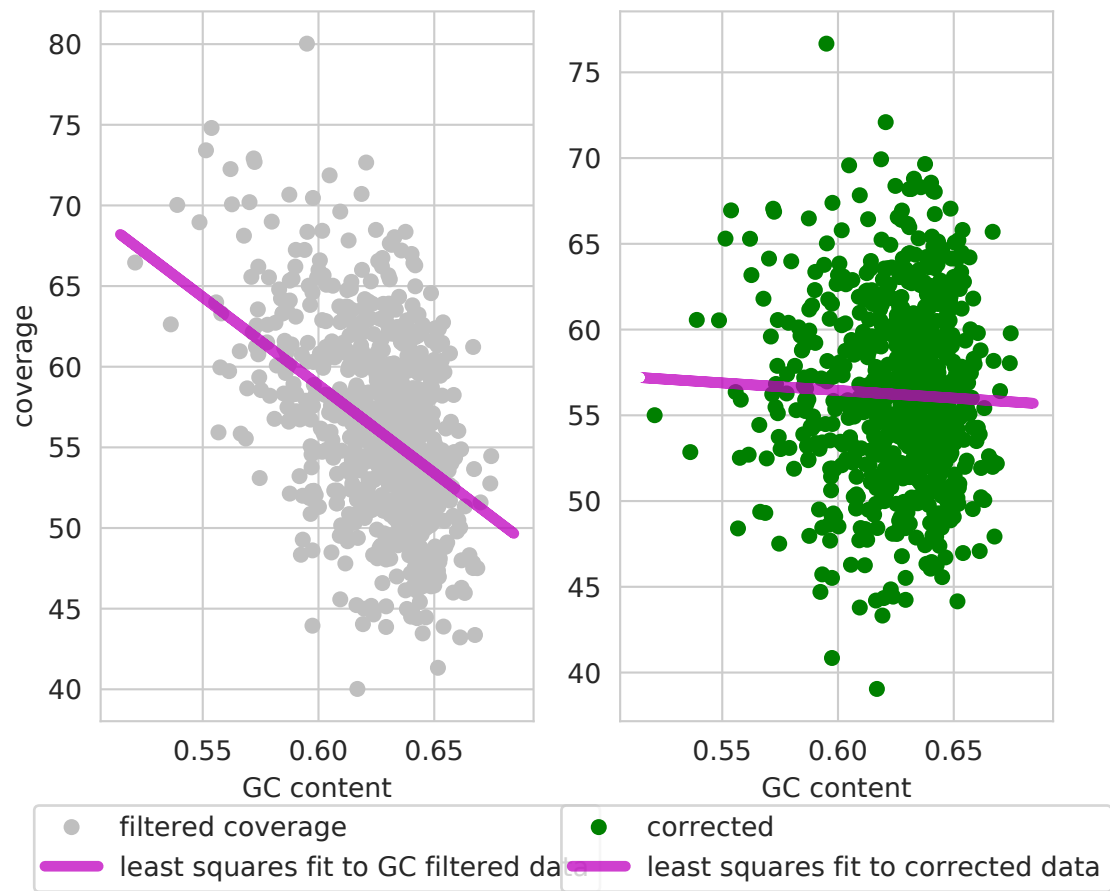


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0727/GCF_005937885.1_ASM593788v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0727-C
iRep: 1.34 raw iRep: 1.38 GC r²: 0.19 GC bias: -20.68 avg. cov: 56.04 % windows: 100.00



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-C
iRep: 1.34 un-filtered iRep: 1.34 r^2 : 0.99 avg. cov: 56.04 % windows: 100.00

