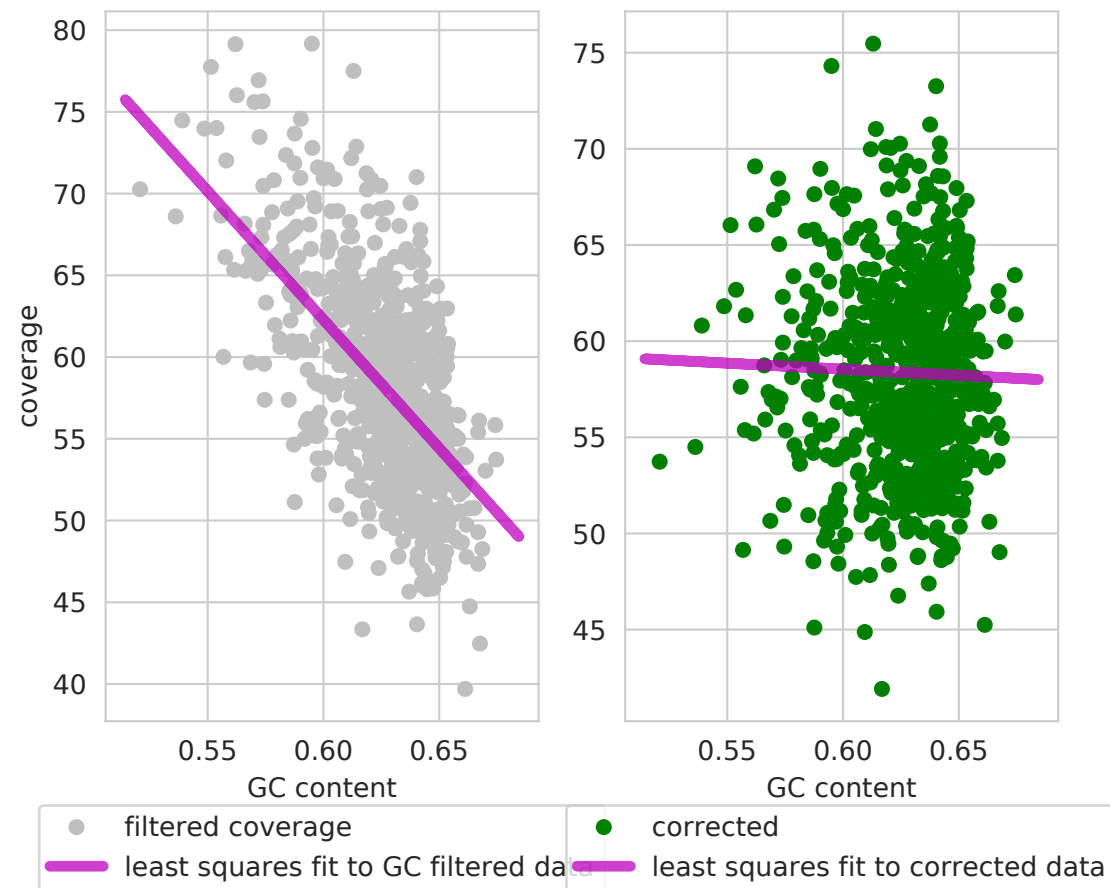


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0727/GCF_005937885.1_ASM593788v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0727-B
iRep: 1.30 raw iRep: 1.37 GC r^2 : 0.35 GC bias: -55.29 avg. cov: 58.20 % windows: 100.00



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-B
iRep: 1.30 un-filtered iRep: 1.30 r^2 : 0.99 avg. cov: 58.20 % windows: 100.00

