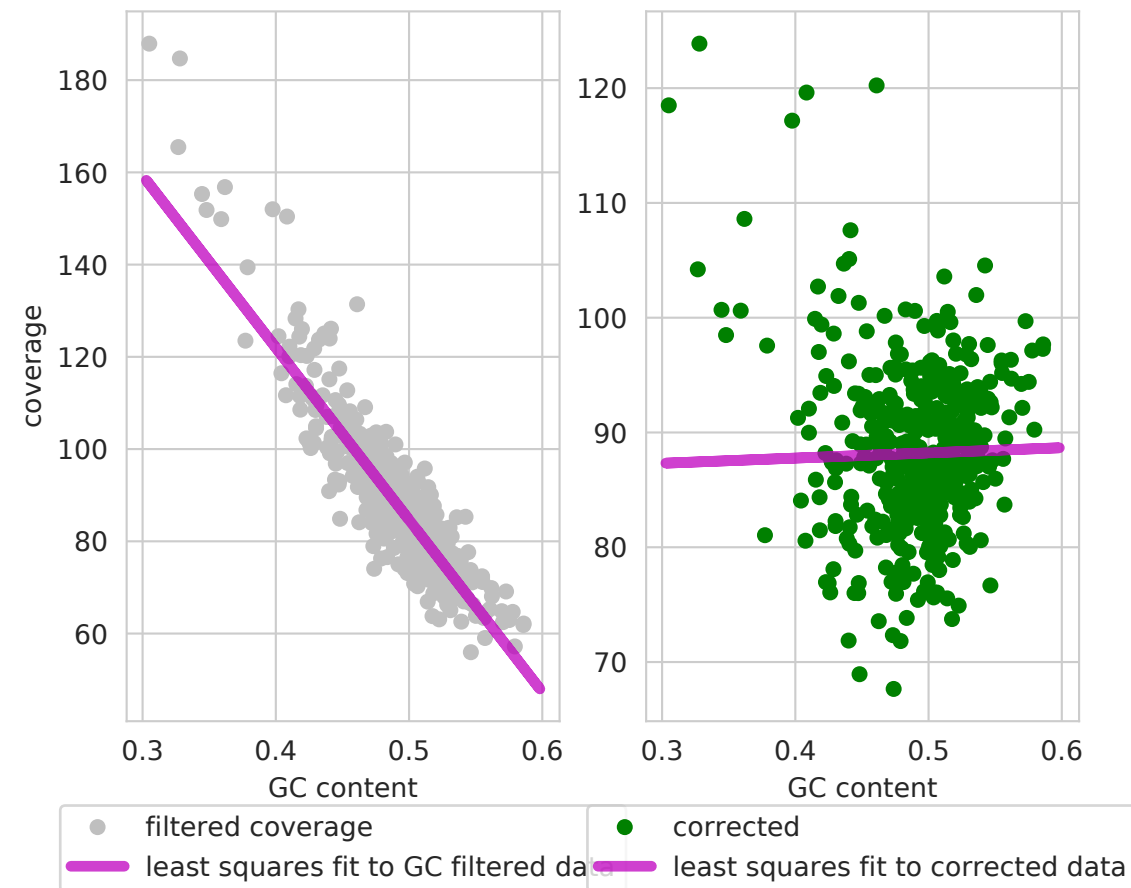


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0713/GCF_005937895.1_ASM593789v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0713-C
iRep: 1.25 raw iRep: 1.62 GC r^2 : 0.82 GC bias: -306.29 avg. cov: 88.02 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0713/GCF_005937895.1_ASM593789v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0713-C
iRep: 1.25 un-filtered iRep: 1.25 r^2 : 0.99 avg. cov: 88.02 % windows: 100.00

