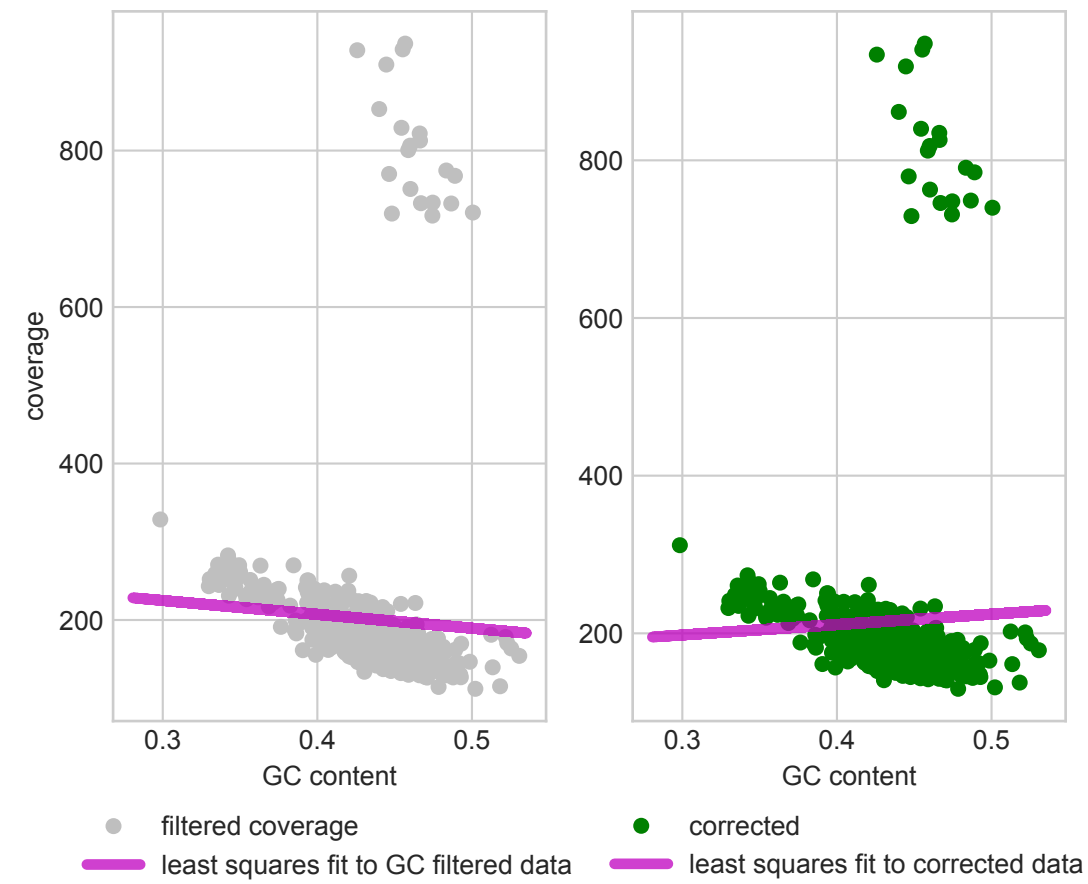


genome: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/reference_assemblies_task2/Bacillus_subtilis_NCIB_3610/GCA_002055965.1_ASM205596v1_genomic sample: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/bwa_fna/D300/Sample_L1B3-300_mapped_sort_NOdup_sort_merged_sort
iRep: 1.68 raw iRep: 1.85 GC r^2: 0.00 GC bias: -0.55 avg. cov: 208.55 % windows: 100.00



genome: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/reference_assemblies_task2/Bacillus_subtilis_NCIB_3610/GCA_002055965.1_ASM205596v1_genomic sample: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/bwa_fna/D300/Sample_L1B3-300_mapped_sort_NOdup_sort_merged_sort
iRep: 1.68 un-filtered iRep: 1.68 r²: 0.94 avg. cov: 208.55 % windows: 100.00

