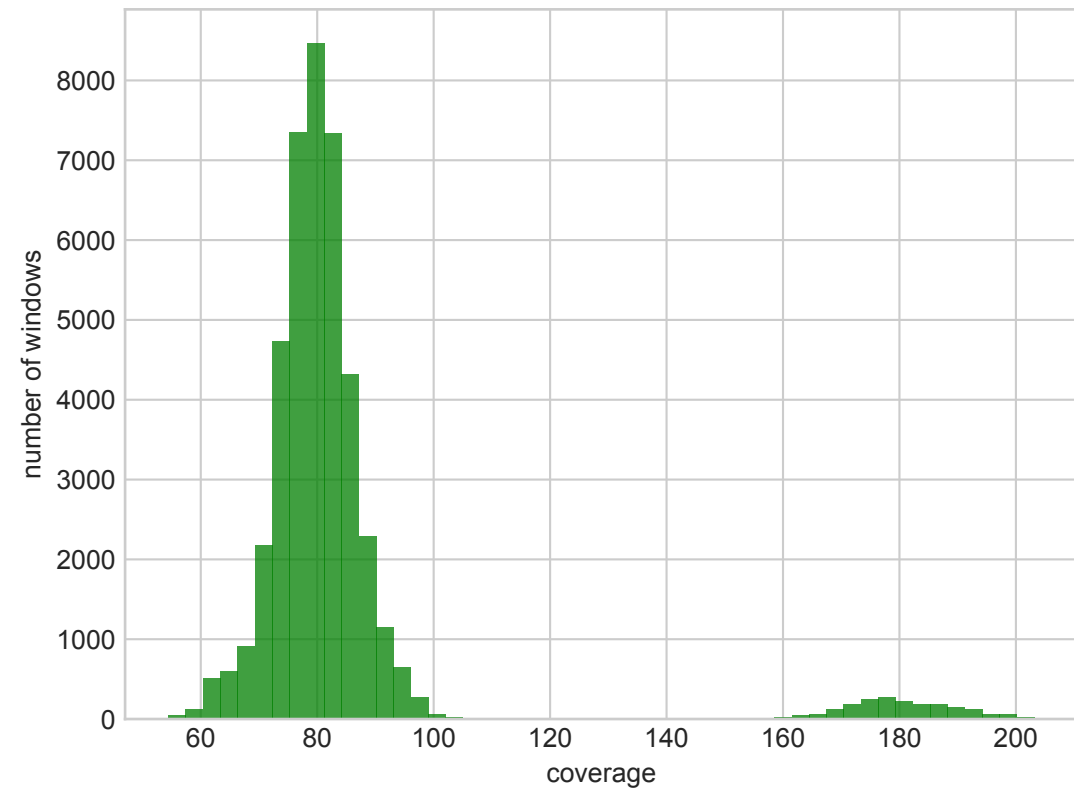
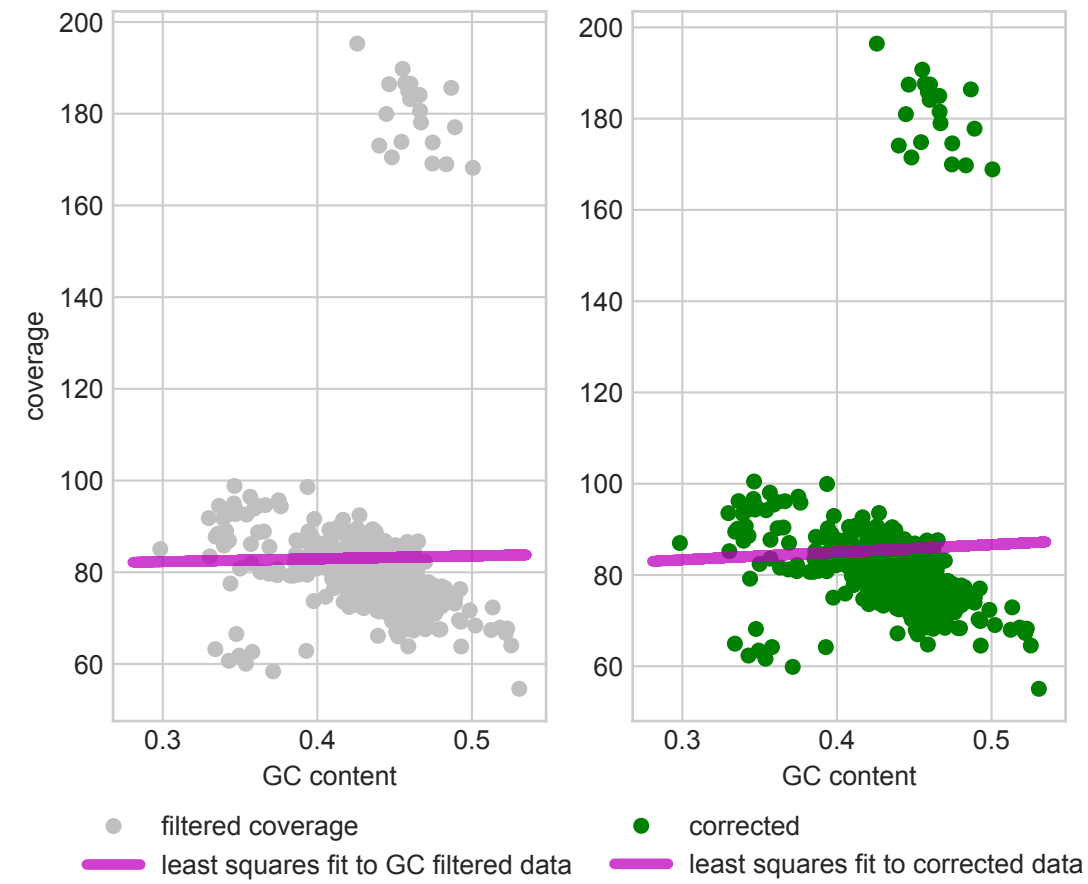


genome: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/reference_assemblies_task2/Bacillus_subtilis_NCIB_3610/GCA_002055965.1_ASM205596v1_genomic sample: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/bwa_fna/D300/Sample_L2B1-300_mapped_sort_NOdup_sort_merged_sort
iRep: 1.29 r^2: 0.96 avg. cov: 84.23 % 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_L2B1-300_mapped_sort_NOdup_sort_merged_sort
iRep: 1.29 un-filtered iRep: 1.29 r^2: 0.96 avg. cov: 84.23 % windows: 100.00

