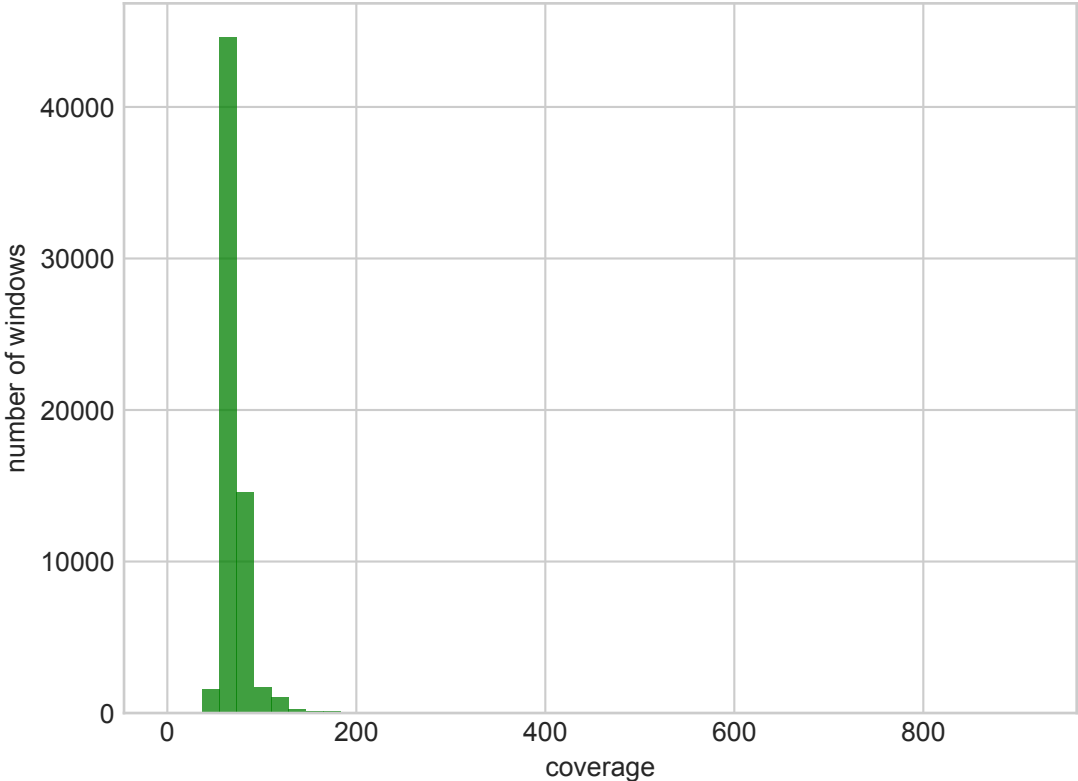
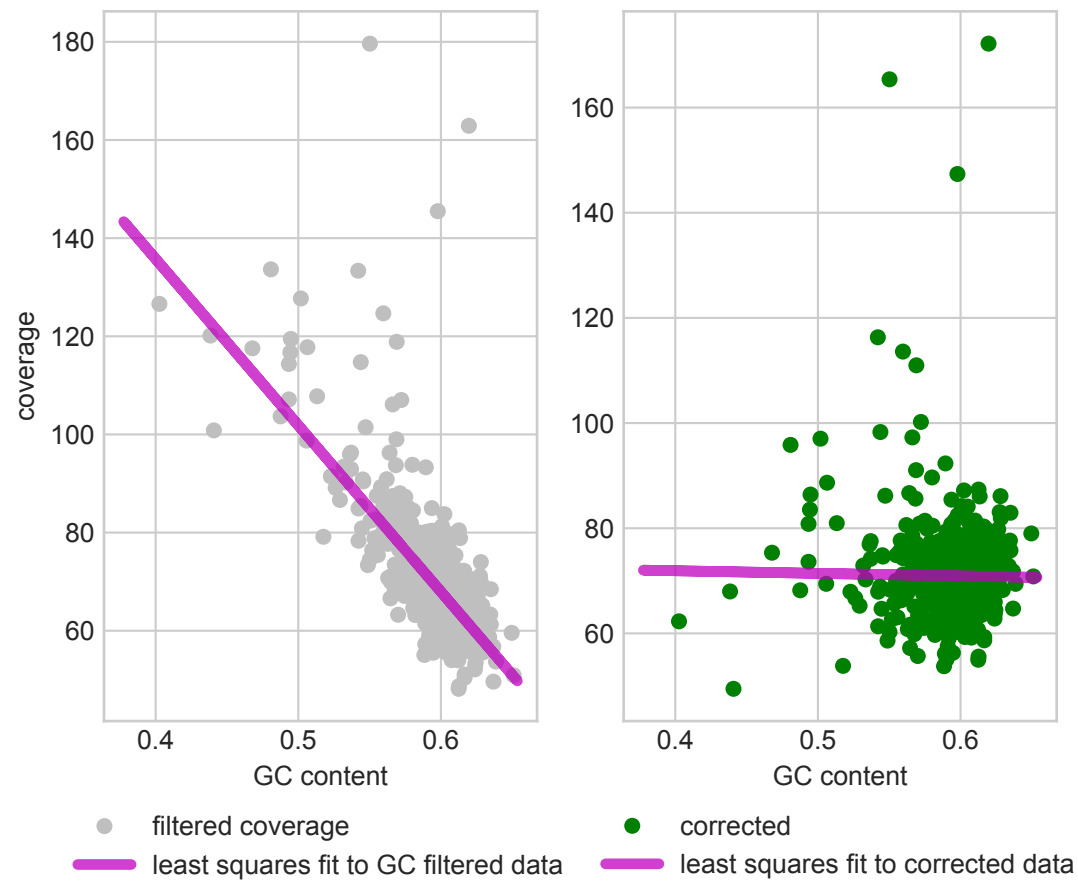


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/ATCC43928/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/ATCC43928-C
iRep: 1.29 r^2: 0.98 avg. cov: 71.31 % windows: 99.78



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/ATCC43928/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/ATCC43928-C
iRep: 1.29 raw iRep: 1.47 GC r^2: 0.68 GC bias: -230.25 avg. cov: 71.31 % windows: 99.78



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/ATCC43928/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/ATCC43928-C
iRep: 1.29 un-filtered iRep: 1.29 r^2: 0.98 avg. cov: 71.31 % windows: 99.78

