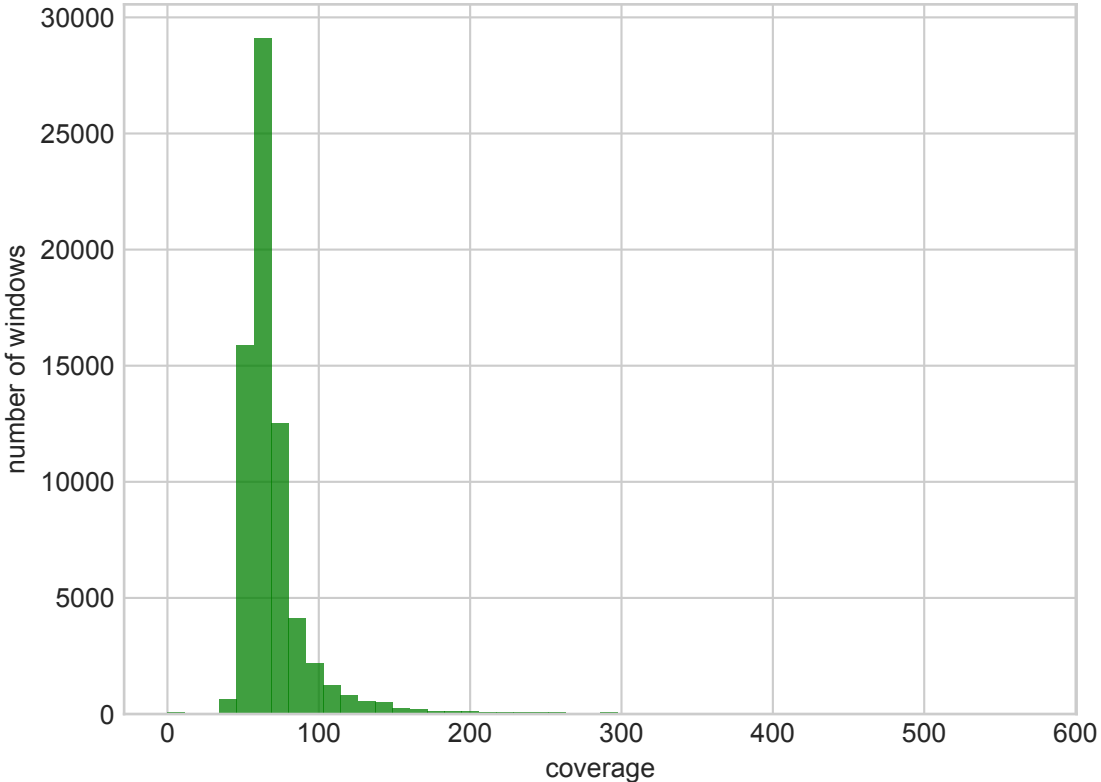
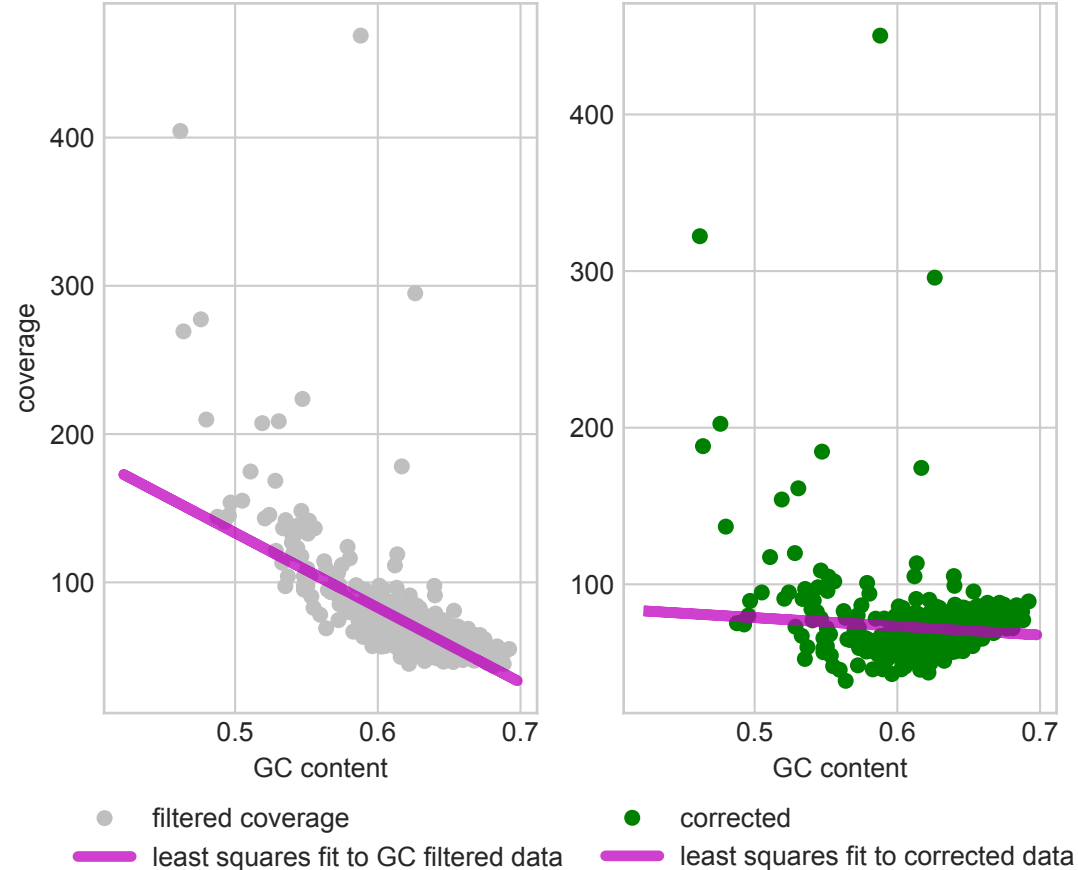


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/ATCC13985/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/ATCC13985-A
iRep: 1.52 r^2: 0.98 avg. cov: 70.67 % windows: 99.88



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/ATCC13985/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/ATCC13985-A
iRep: 1.52 raw iRep: 1.83 GC r^2: 0.68 GC bias: -340.60 avg. cov: 70.67 % windows: 99.88



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2016_KBSGenomes_Annotate/ATCC13985/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/ATCC13985-A
iRep: 1.52 un-filtered iRep: 1.52 r^2: 0.98 avg. cov: 70.67 % windows: 99.88

