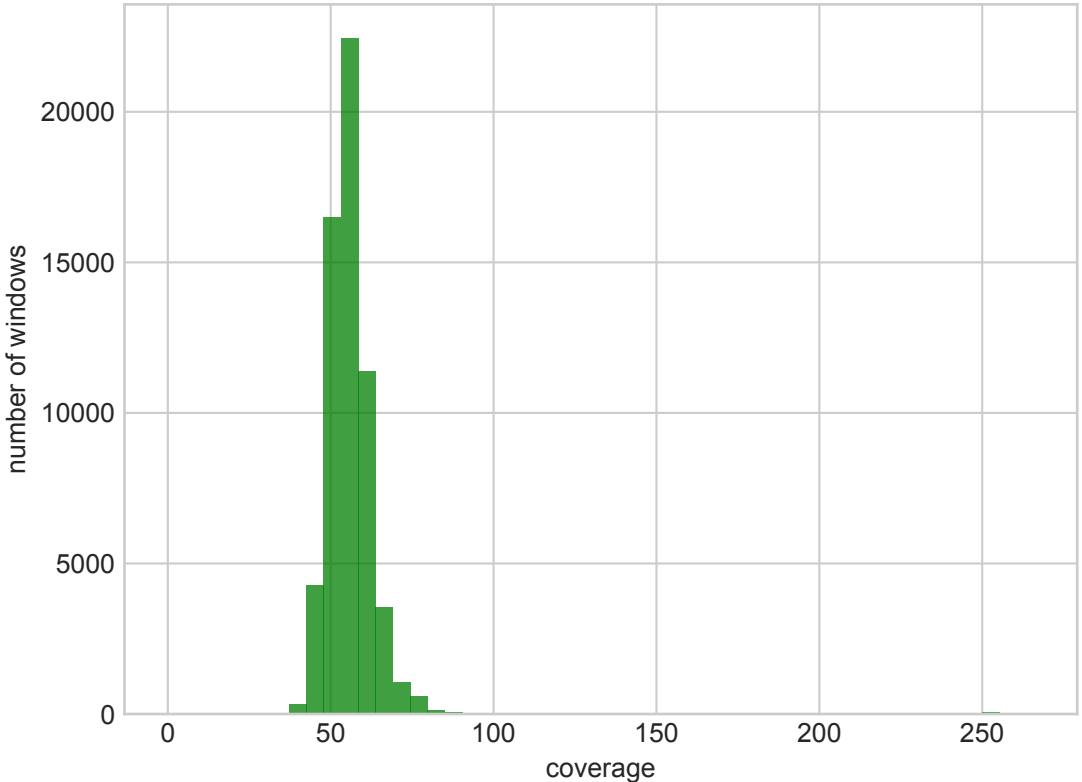
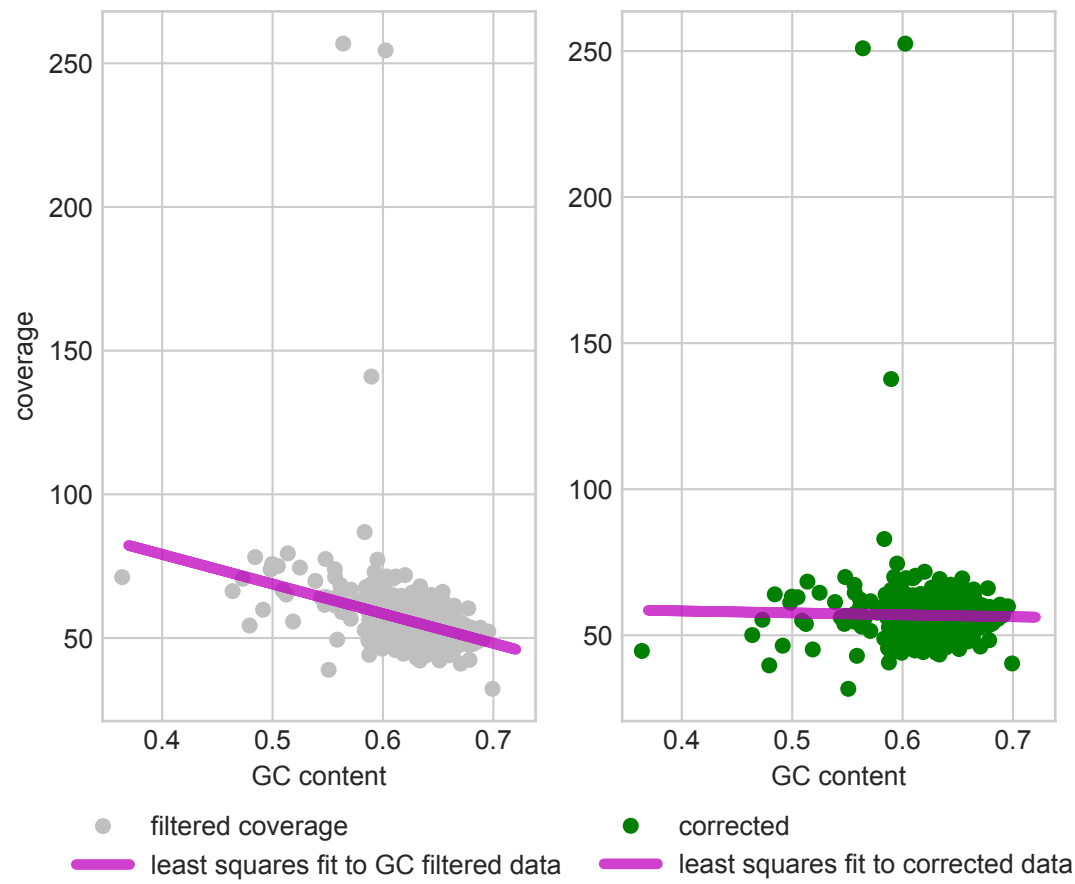


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0711/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0711-D
iRep: 1.32 r^2: 0.99 avg. cov: 56.37 % windows: 99.97



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0711/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0711-D
iRep: 1.32 raw iRep: 1.36 GC r^2: 0.29 GC bias: -29.54 avg. cov: 56.37 % windows: 99.97



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0711/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0711-D
iRep: 1.32 un-filtered iRep: 1.32 r^2: 0.99 avg. cov: 56.37 % windows: 99.97

