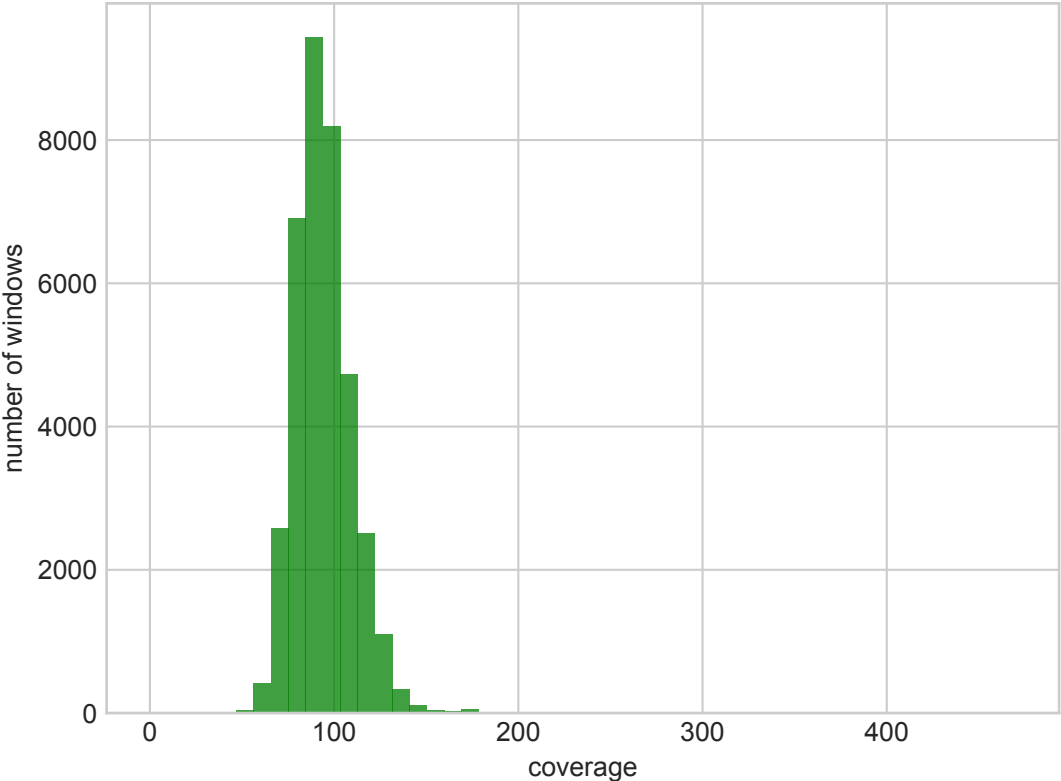
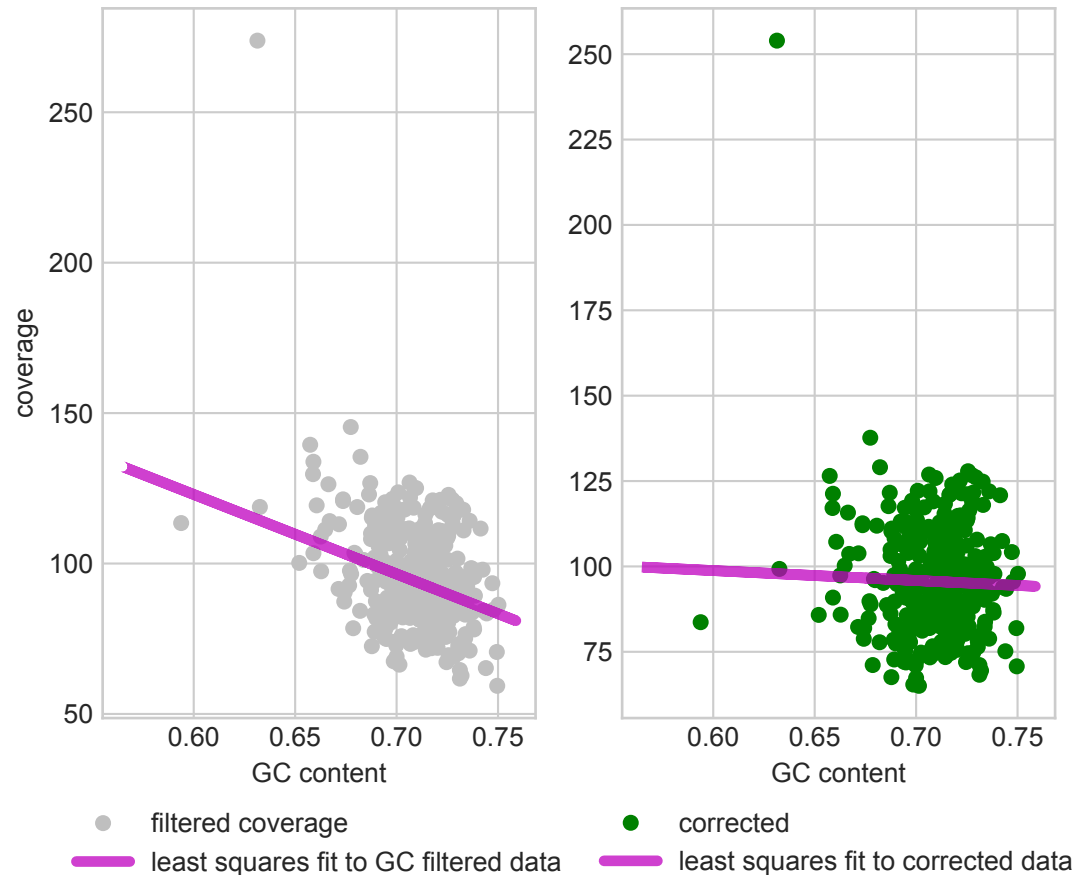


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0715/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0715-A
iRep: 1.57 r^2: 0.99 avg. cov: 94.71 % windows: 99.97



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0715/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0715-A
iRep: 1.57 raw iRep: 1.61 GC r^2: 0.13 GC bias: -34.61 avg. cov: 94.71 % windows: 99.97



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0715/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0715-A
iRep: 1.57 un-filtered iRep: 1.57 r^2: 0.99 avg. cov: 94.71 % windows: 99.97

