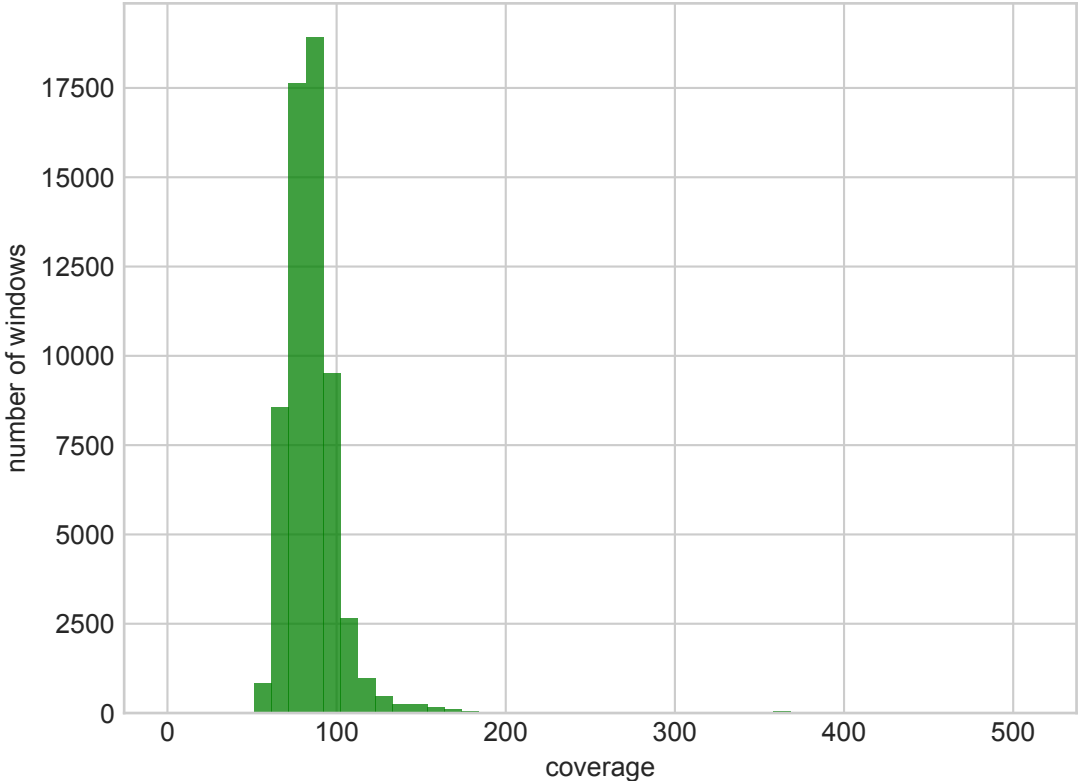
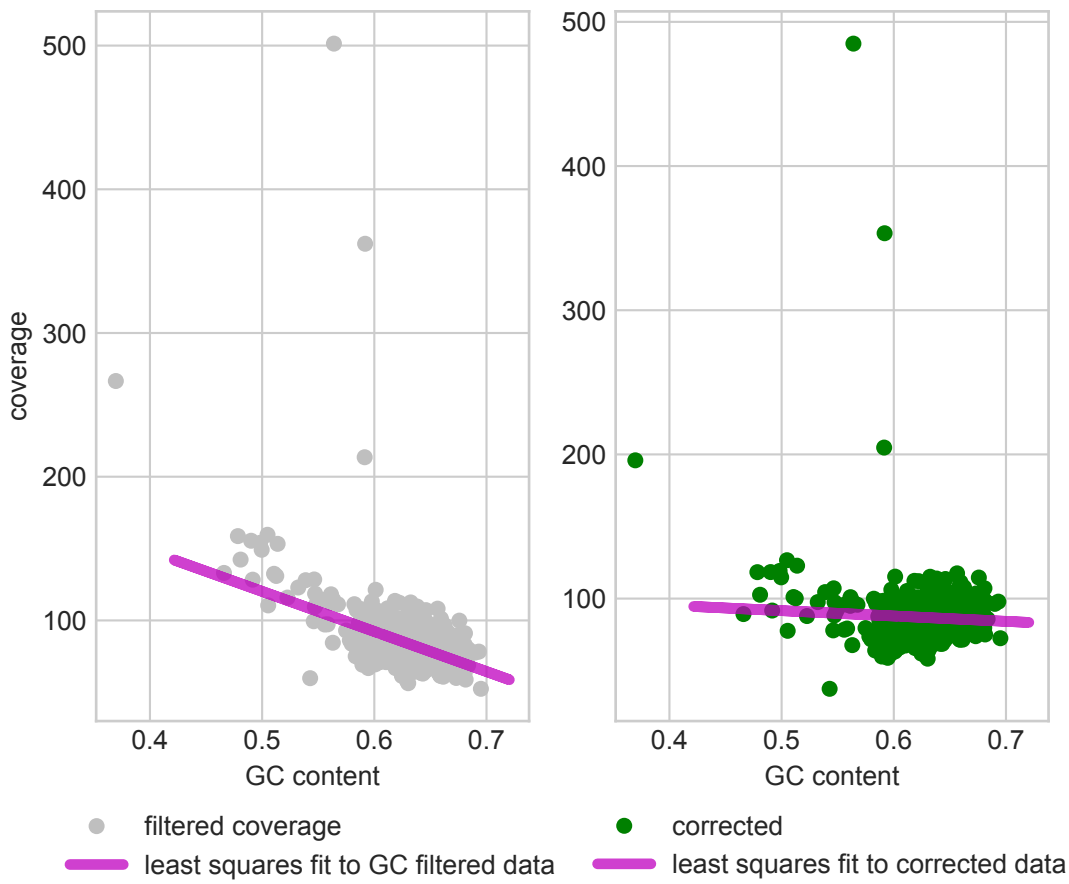


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0711/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0711-WB
iRep: 1.49 r^2 : 0.99 avg. cov: 86.08 % windows: 99.98



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0711/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0711-WB
iRep: 1.49 raw iRep: 1.57 GC r^2: 0.37 GC bias: -103.67 avg. cov: 86.08 % windows: 99.98



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0711/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0711-WB
iRep: 1.49 un-filtered iRep: 1.49 r^2: 0.99 avg. cov: 86.08 % windows: 99.98

