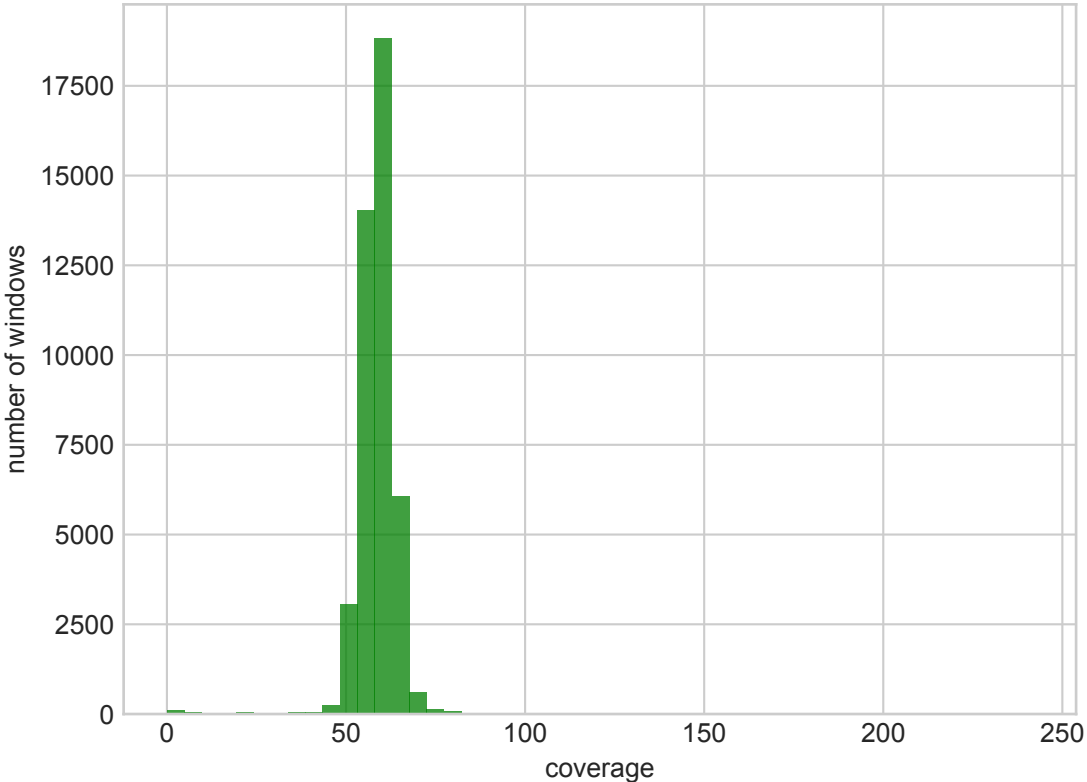
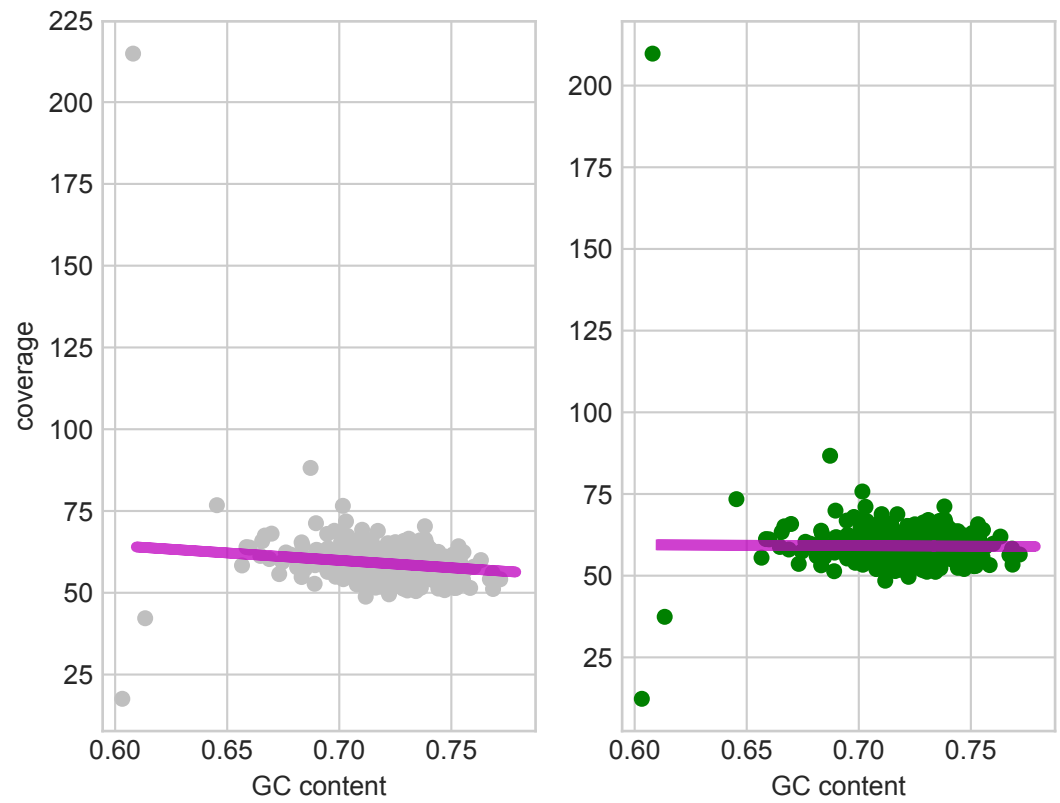


genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0722/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0722-B
iRep: 1.23 r²: 0.98 avg. cov: 58.85 % windows: 99.72



genome: /Users/WRShoemaker/GitHub/LTDE/data/genomes/2015_SoilGenomes_Annotate/KBS0722/G-Chr1 sample: /Users/WRShoemaker/GitHub/LTDE/data/bwa_sam/KBS0722-B
iRep: 1.23 raw iRep: 1.23 GC r^2: 0.05 GC bias: -2.33 avg. cov: 58.85 % windows: 99.72



● filtered coverage ● corrected
— least squares fit to GC filtered data — least squares fit to corrected data

