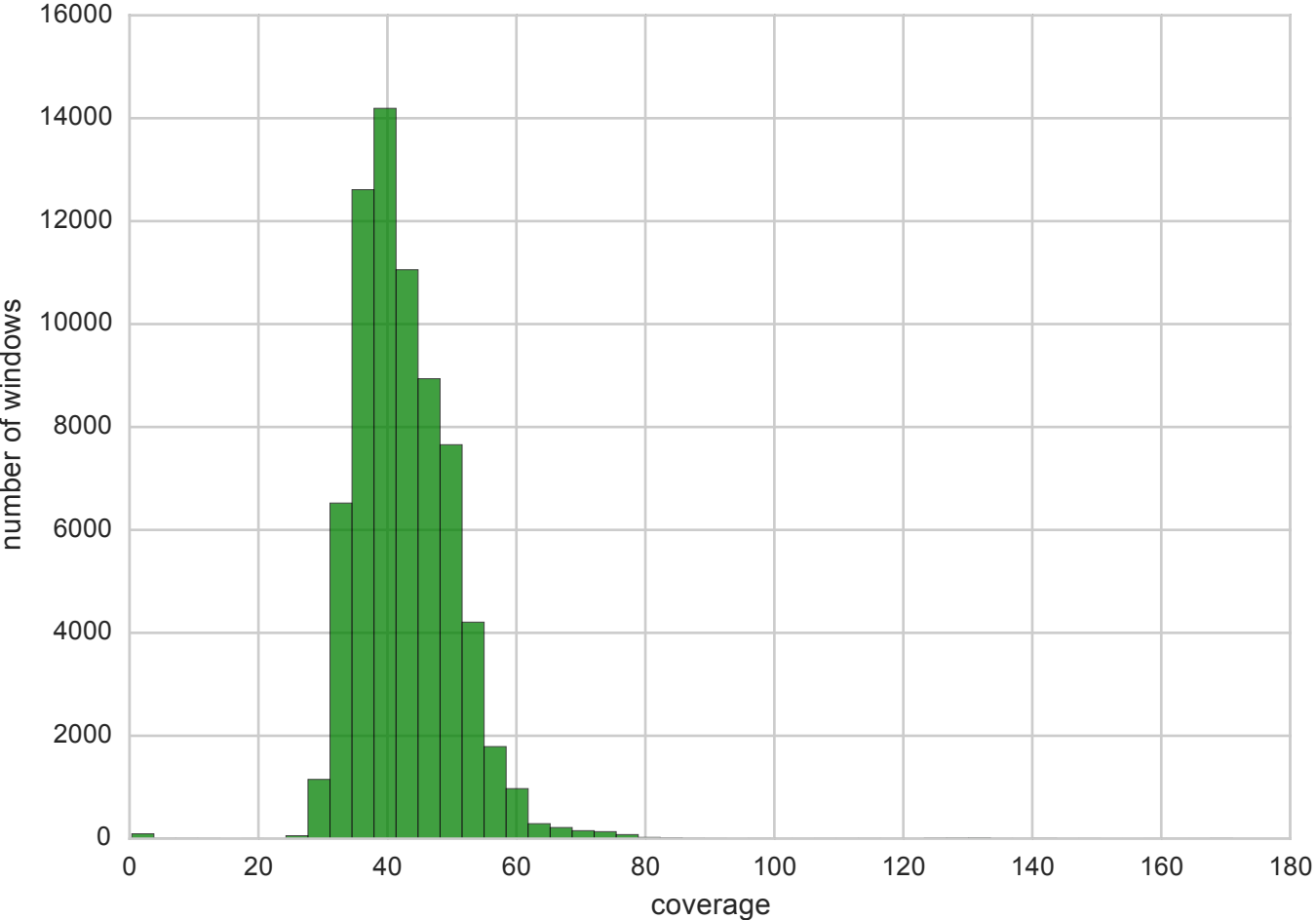
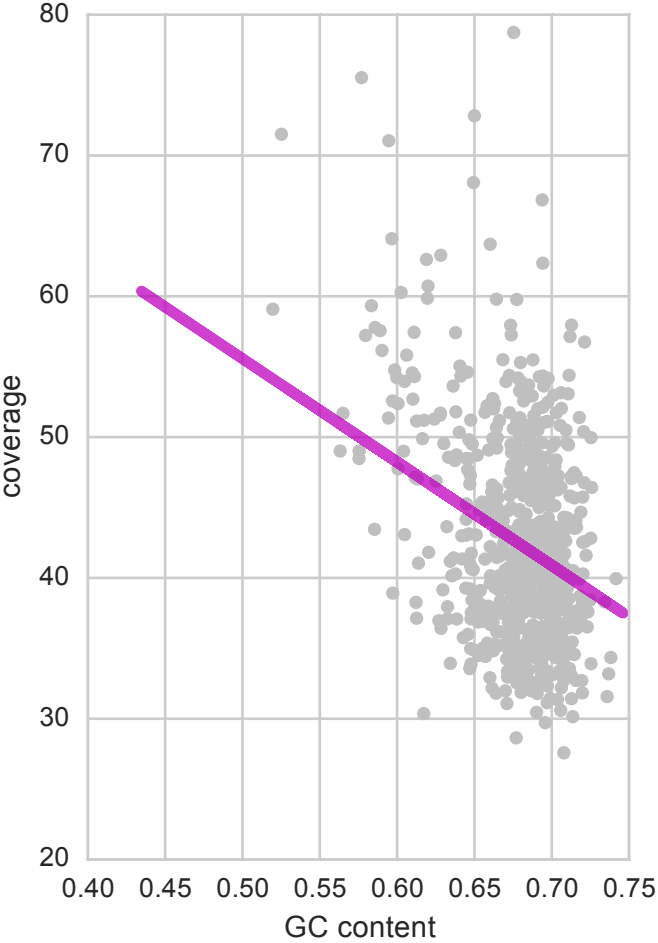
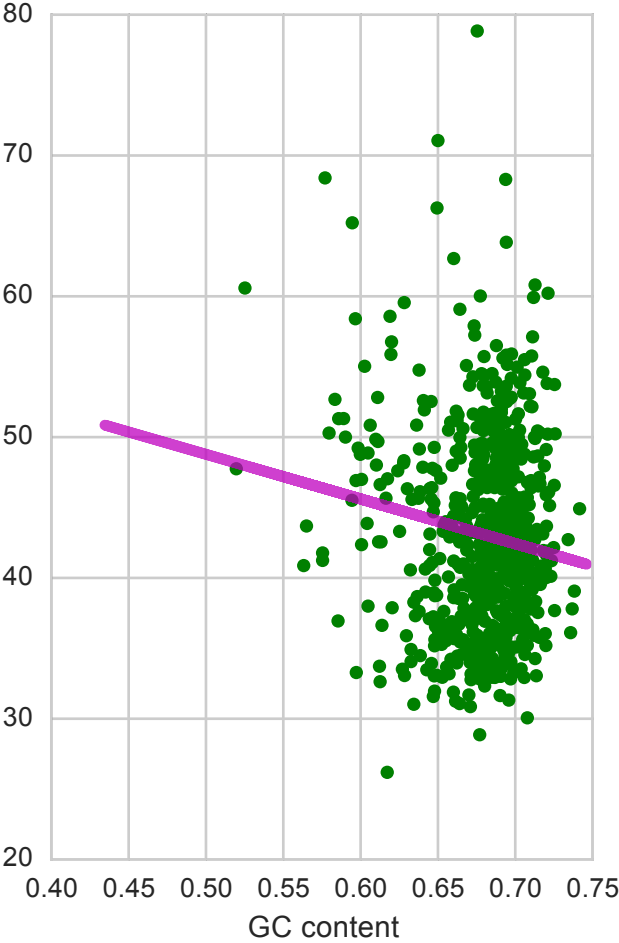


genome: /Users/WRShoemaker/github/Task2/LTDE/data/2016_KBSGenomes_Annotate/KBS0712/G-Chr1 sample: /Users/WRShoemaker/github/Task2/LTDE/data/iRep/KBS0712/GSF1046-KBS0712-B_S60_mapped_sort_NOdup_
iRep: 1.64 r^2: 1.00 avg. cov: 42.73 % windows: 99.86





● filtered coverage
— least squares fit to GC filtered data



● corrected
— least squares fit to corrected data

