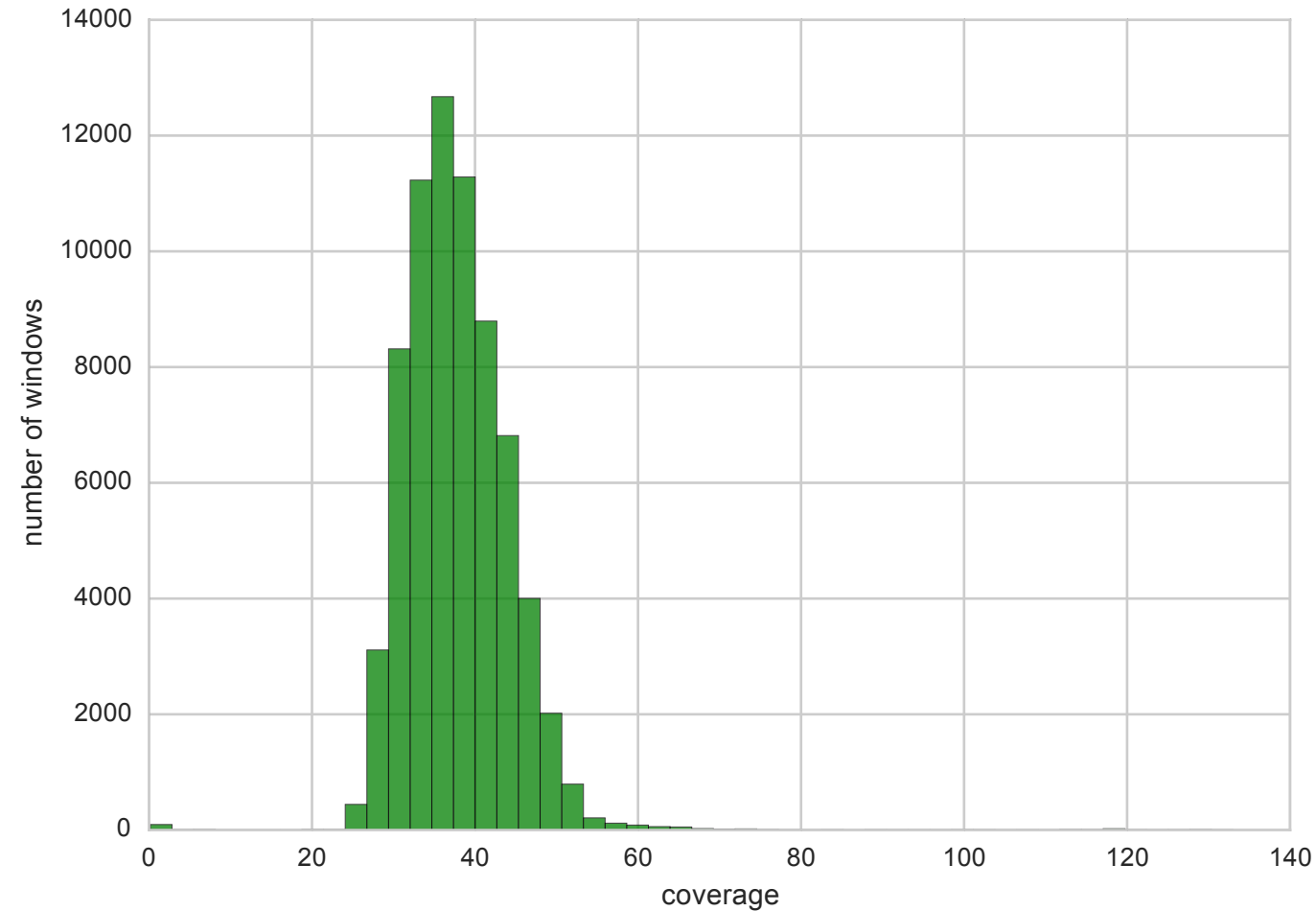
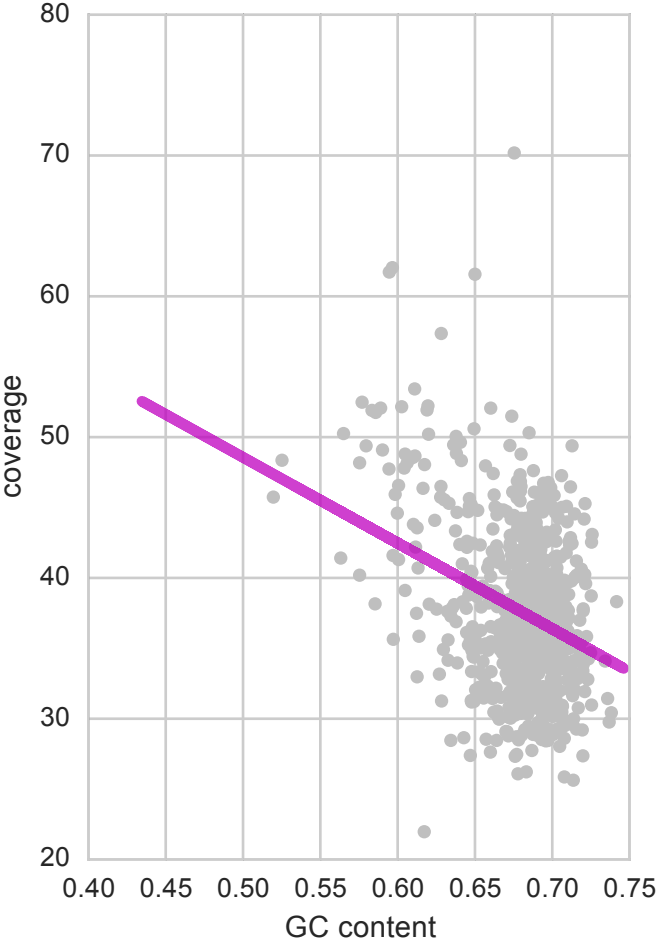
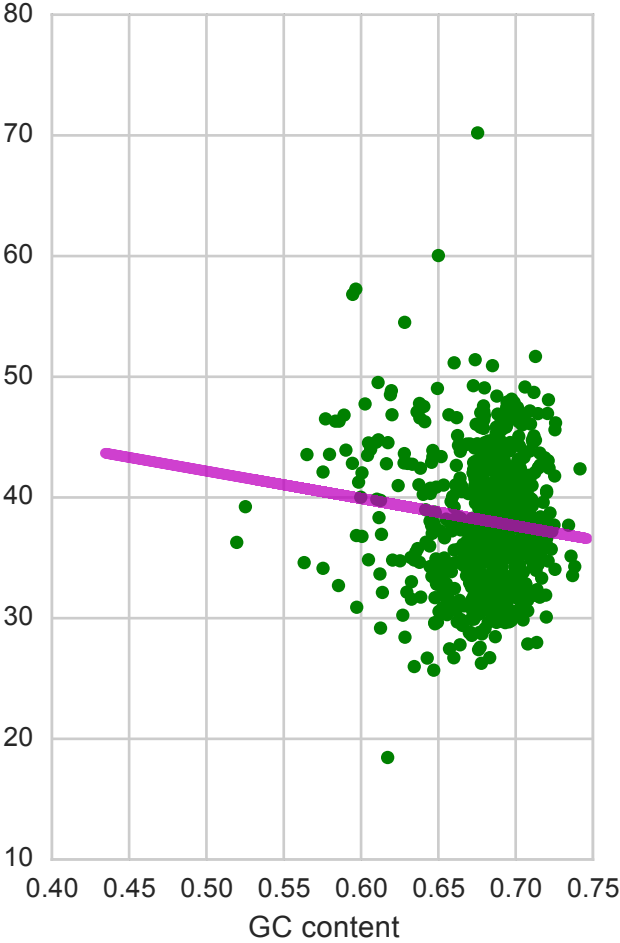


genome: /Users/WRShoemaker/github/Task2/LTDE/data/2016\_KBSGenomes\_Annotate/KBS0712/G-Chr1 sample: /Users/WRShoemaker/github/Task2/LTDE/data/iRep/KBS0712/GSF1046-KBS0712-D\_S74\_mapped\_sort\_NOdup\_  
iRep: 1.57 r^2: 0.99 avg. cov: 37.84 % windows: 99.86

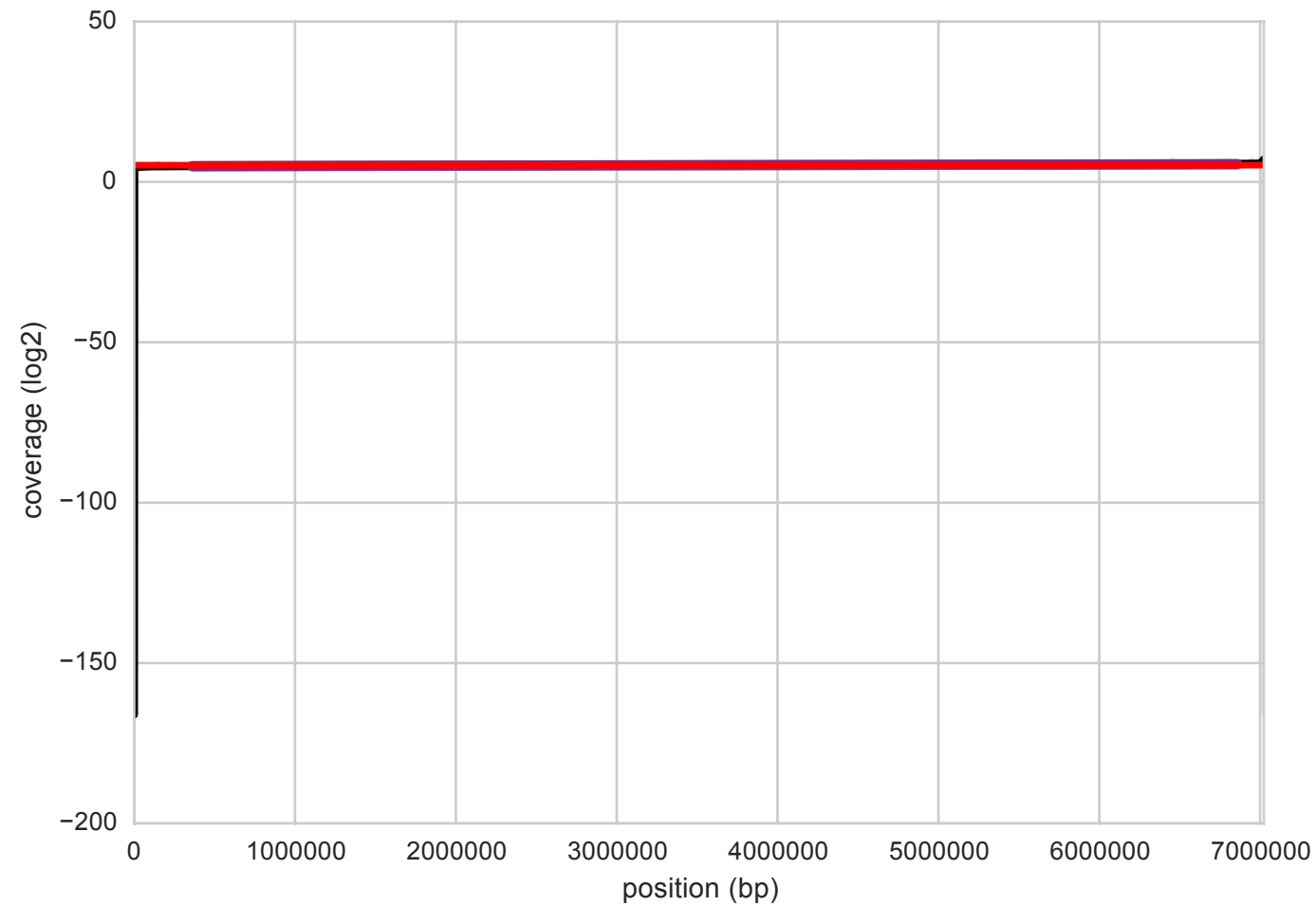




● filtered coverage  
— least squares fit to GC filtered data



● corrected  
— least squares fit to corrected data



- filtered
- sorted
- trimmed
- least squares fit to trimmed
- Ori/Ter