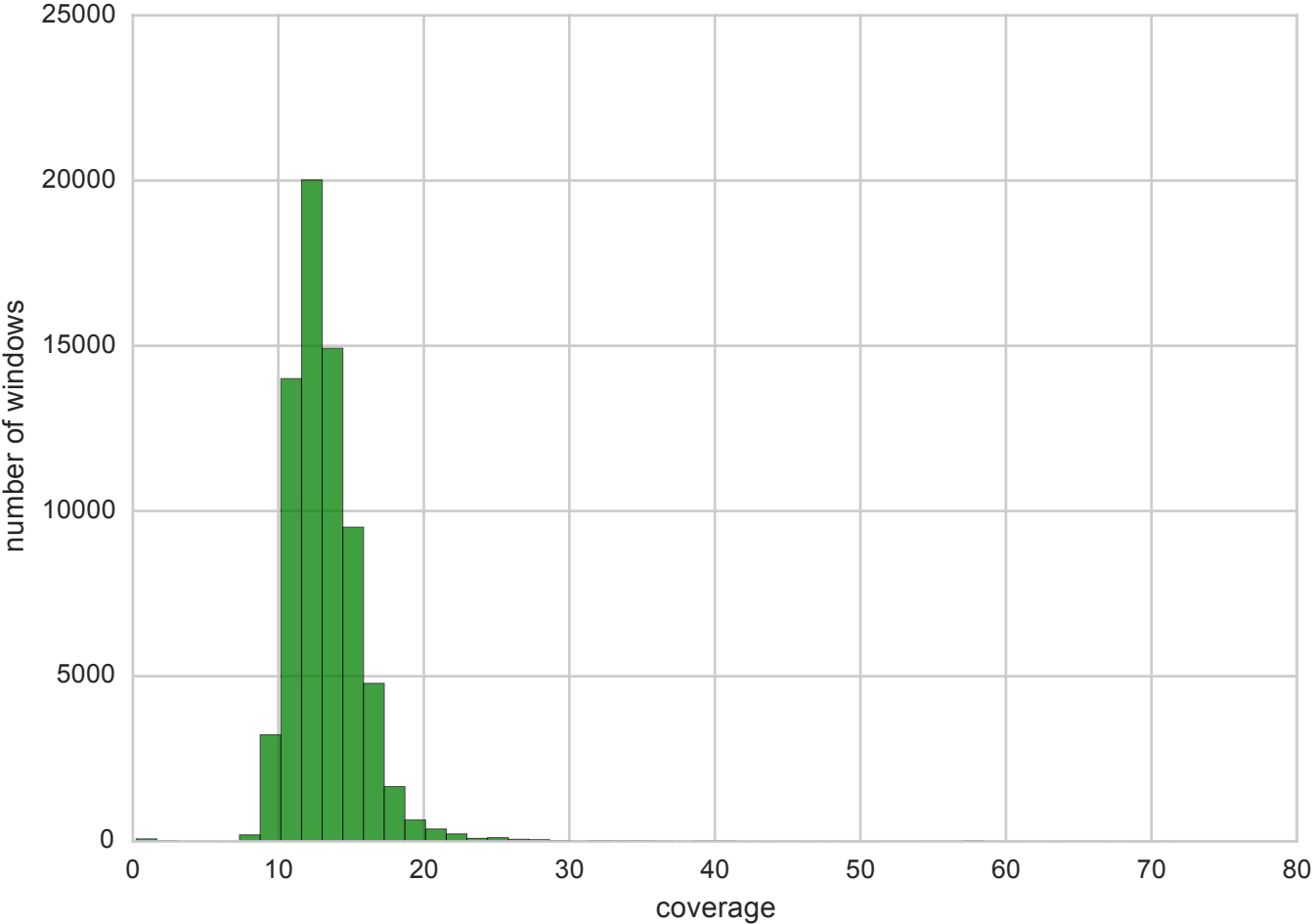
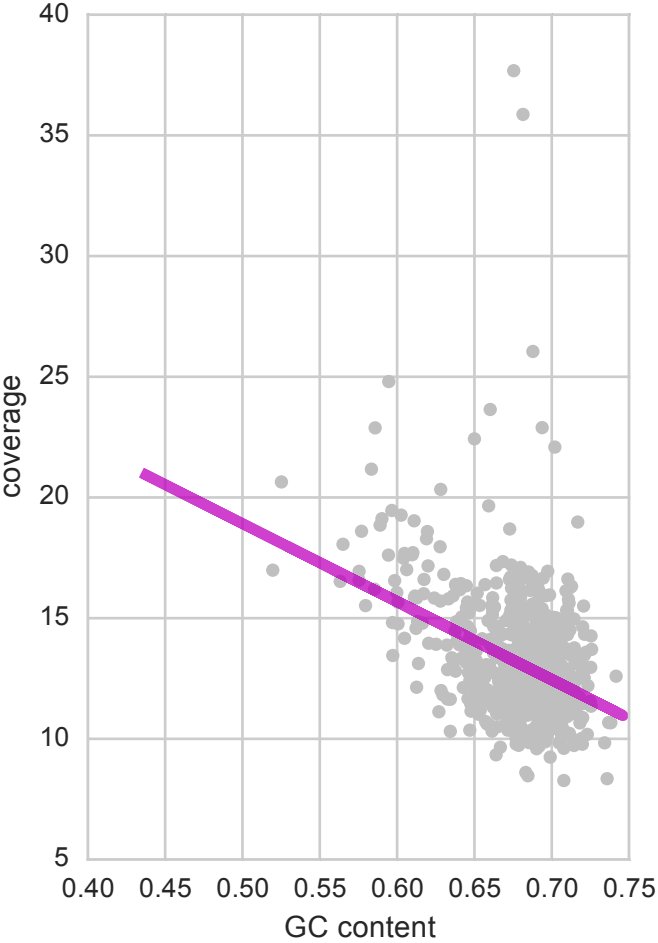
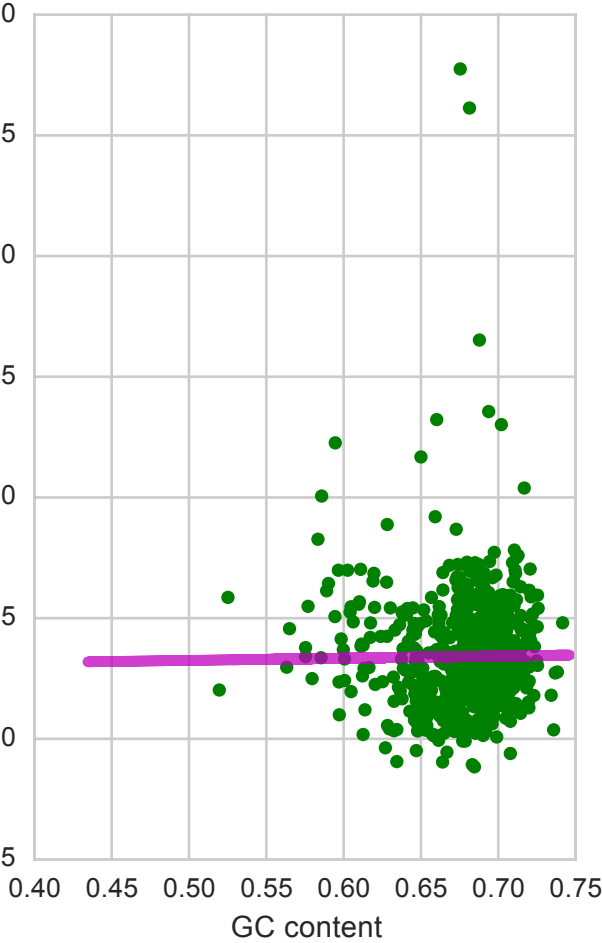


genome: /Users/WRShoemaker/github/Task2/LTDE/data/2016_KBSGenomes_Annotate/KBS0712/G-Chr1 sample: /Users/WRShoemaker/github/Task2/LTDE/data/iRep/KBS0712/GSF1046-KBS0712-A_S53_mapped_sort_NOdup_
iRep: 1.57 r^2: 0.99 avg. cov: 13.30 % windows: 99.89

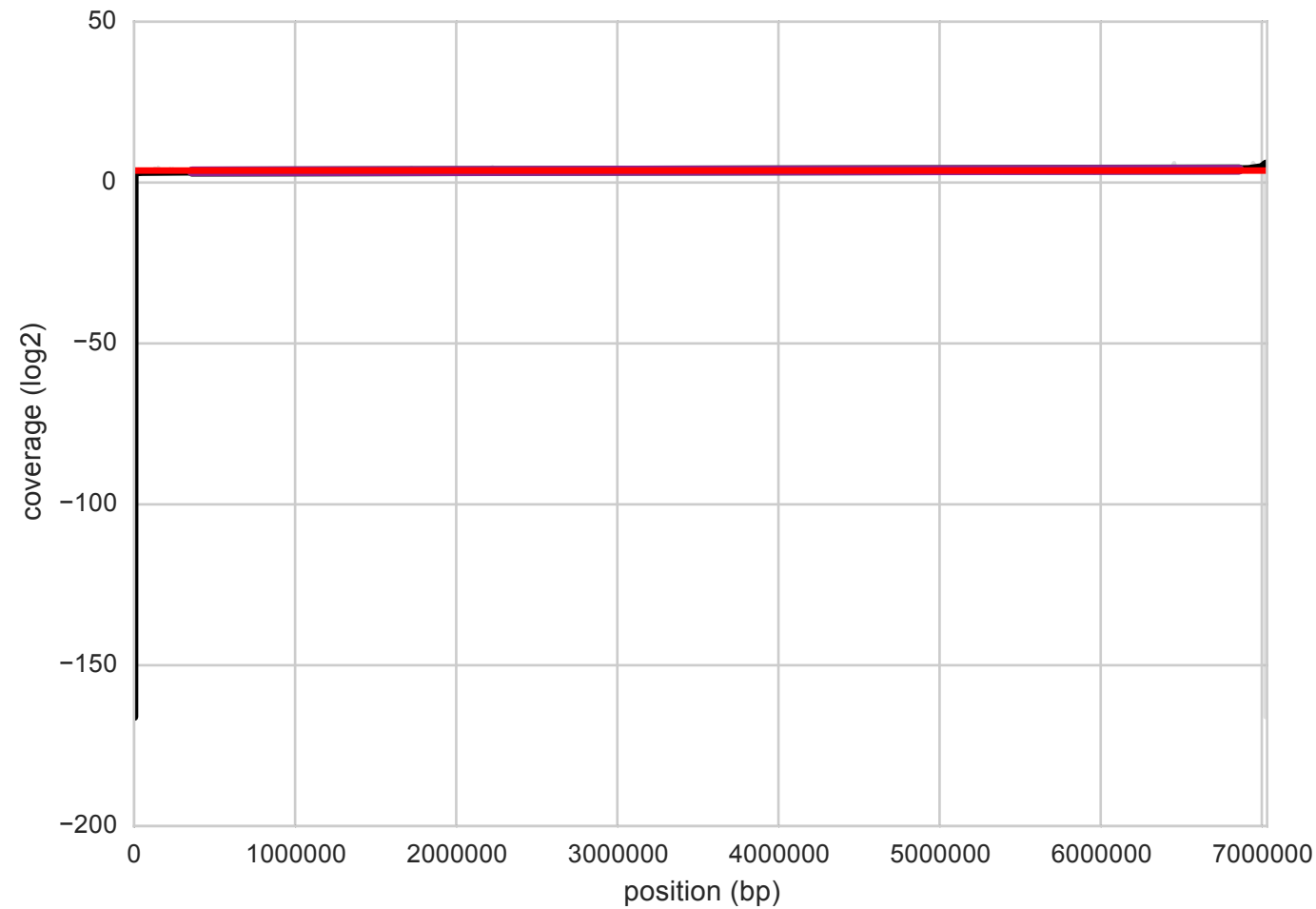




● filtered coverage
— least squares fit to GC filtered data



● corrected
— least squares fit to corrected data



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