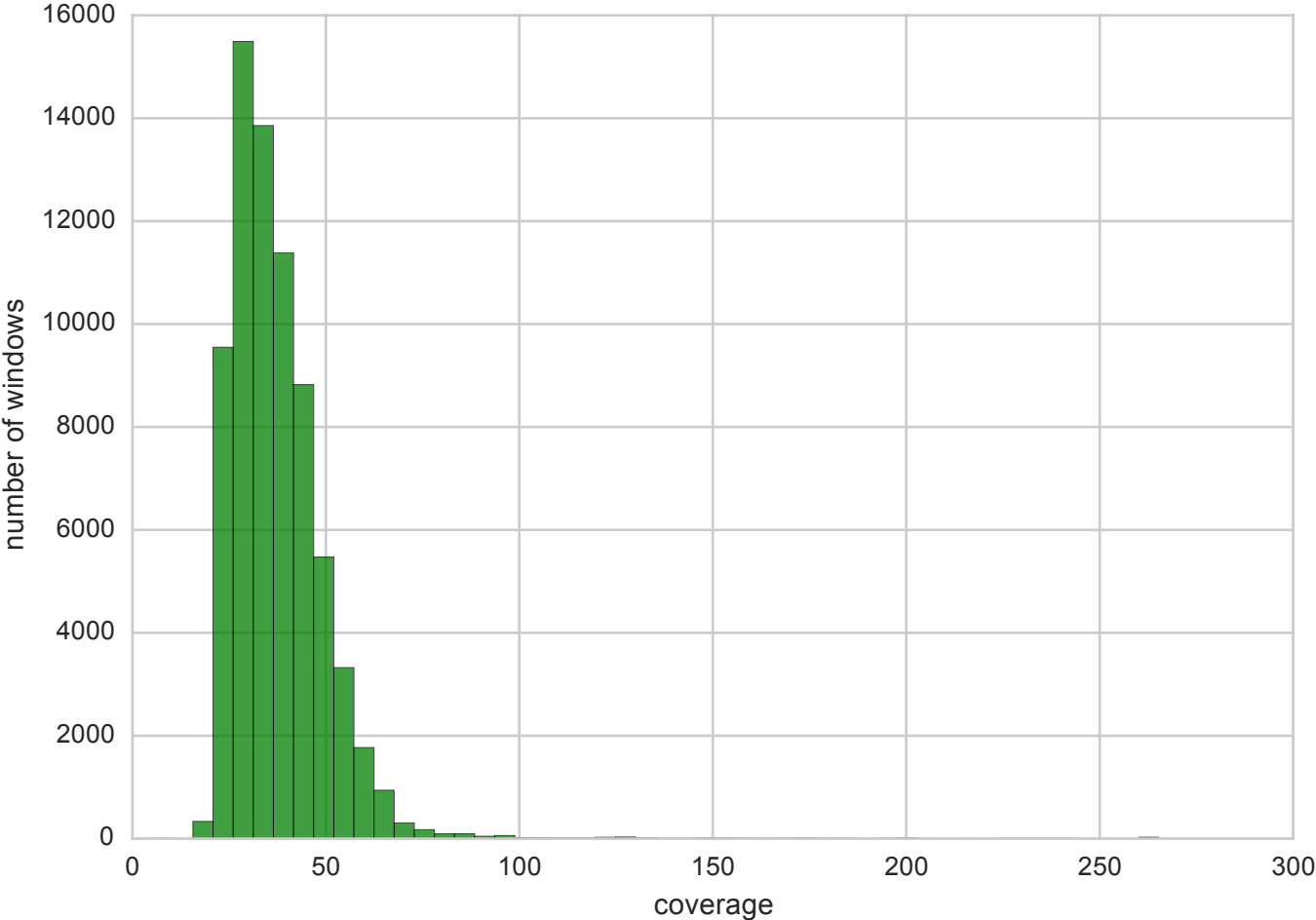
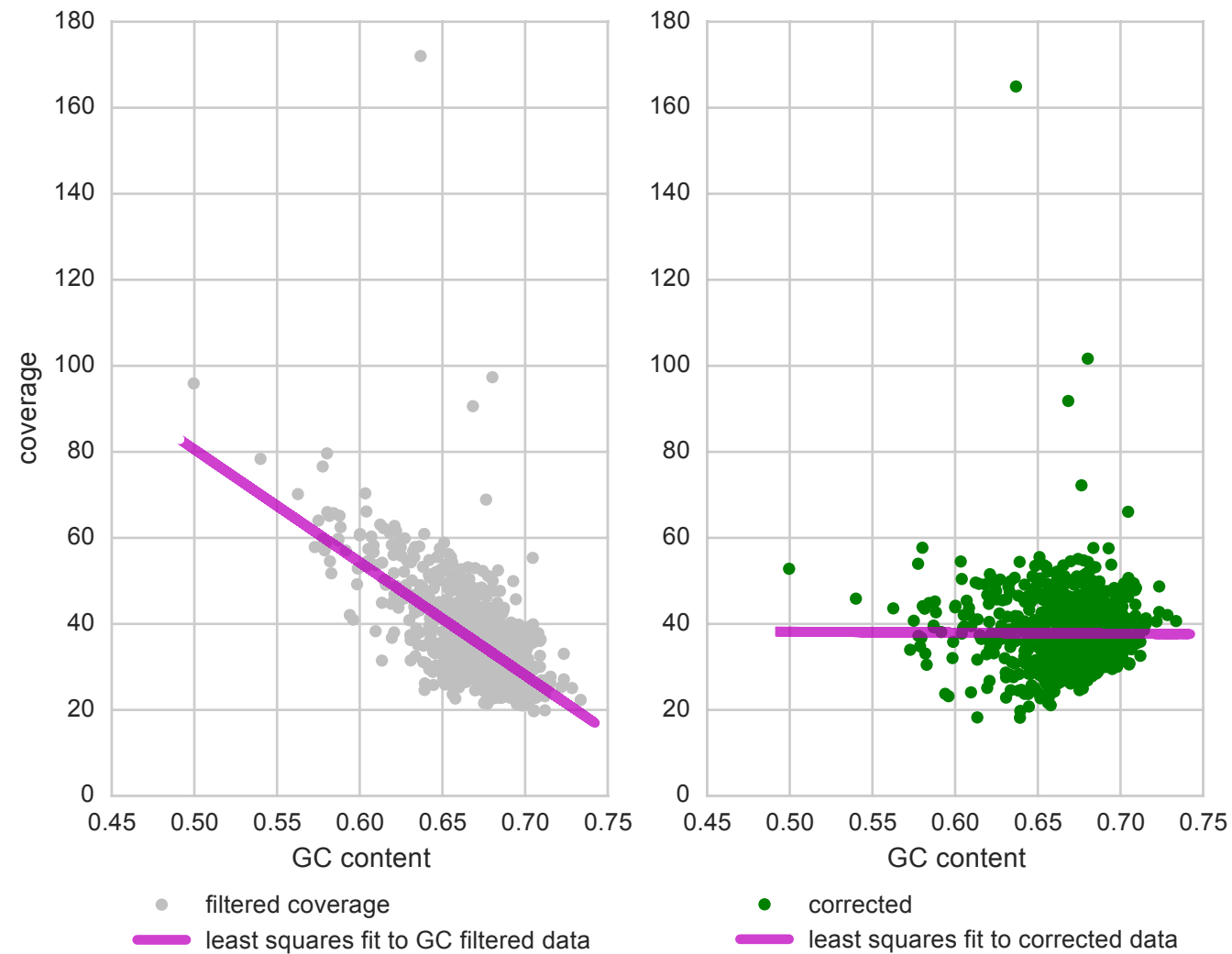
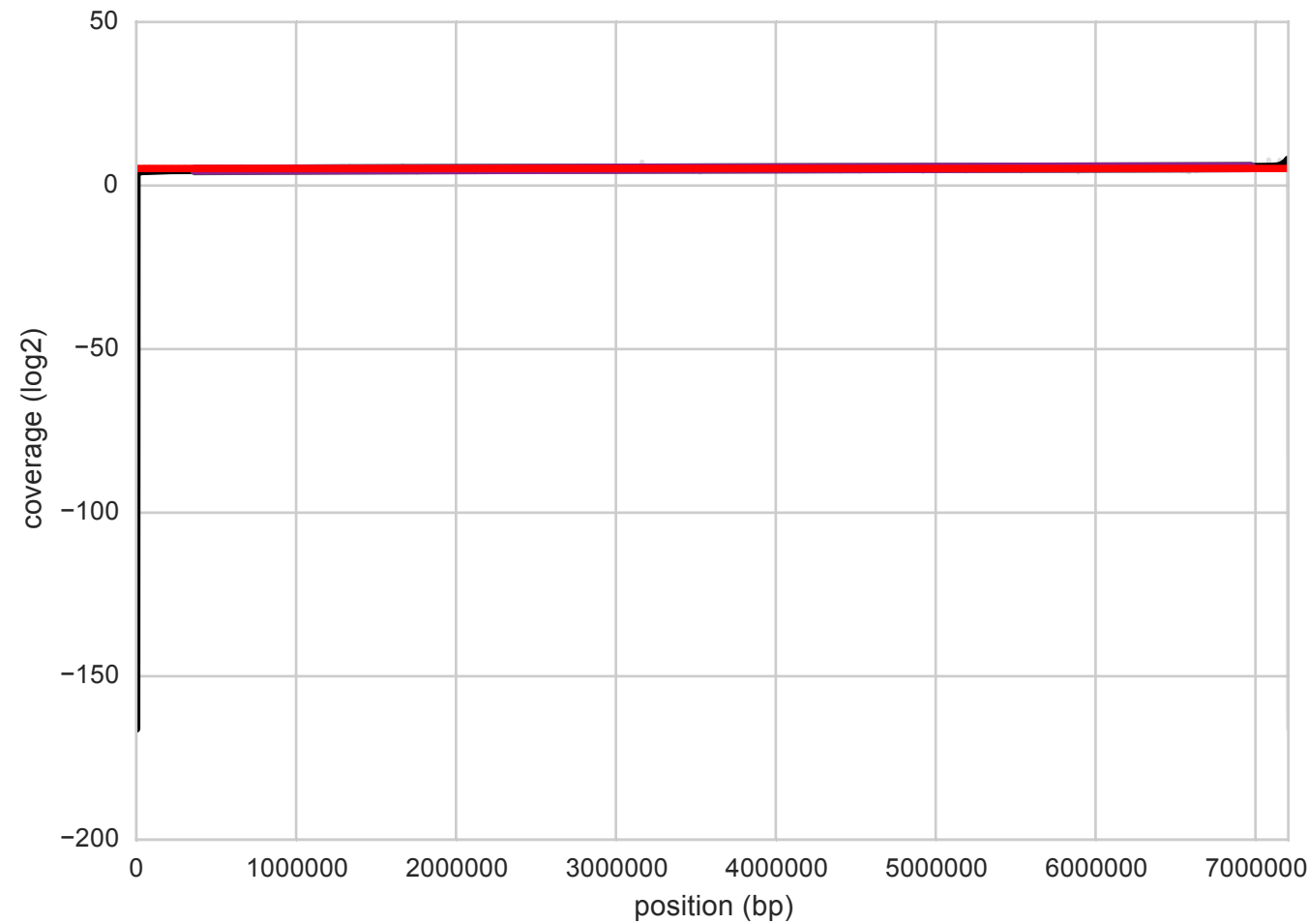


genome: /Users/WRShoemaker/github/Task2/LTDE/data/2016_KBSGenomes_Annotate/KBS0801/G-Chr1 sample: /Users/WRShoemaker/github/Task2/LTDE/data/iRep/KBS0801/GSF1046-KBS0801-B_S82_mapped_sort_NOdup_
iRep: 1.86 r^2: 0.99 avg. cov: 37.57 % windows: 100.00







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