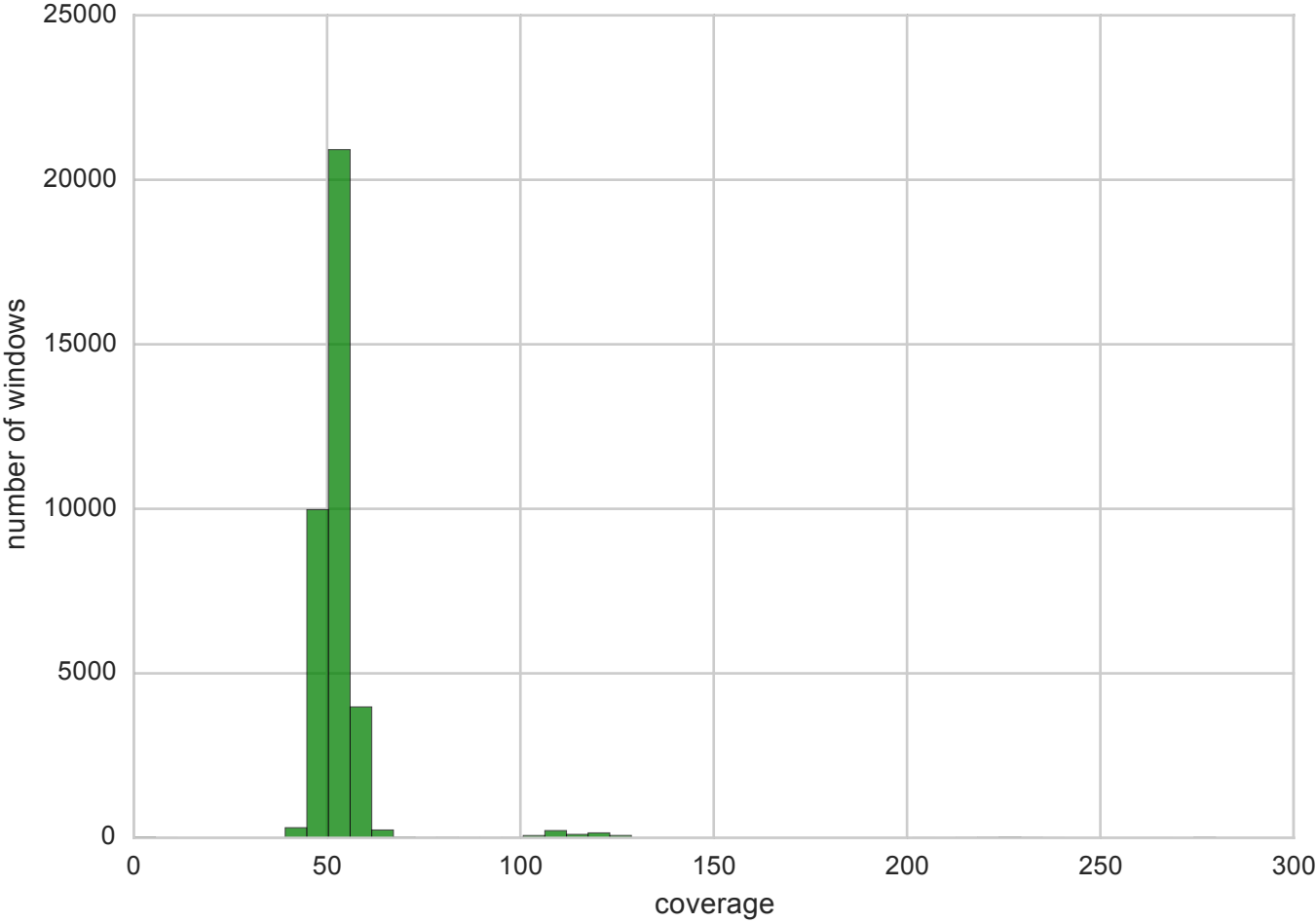
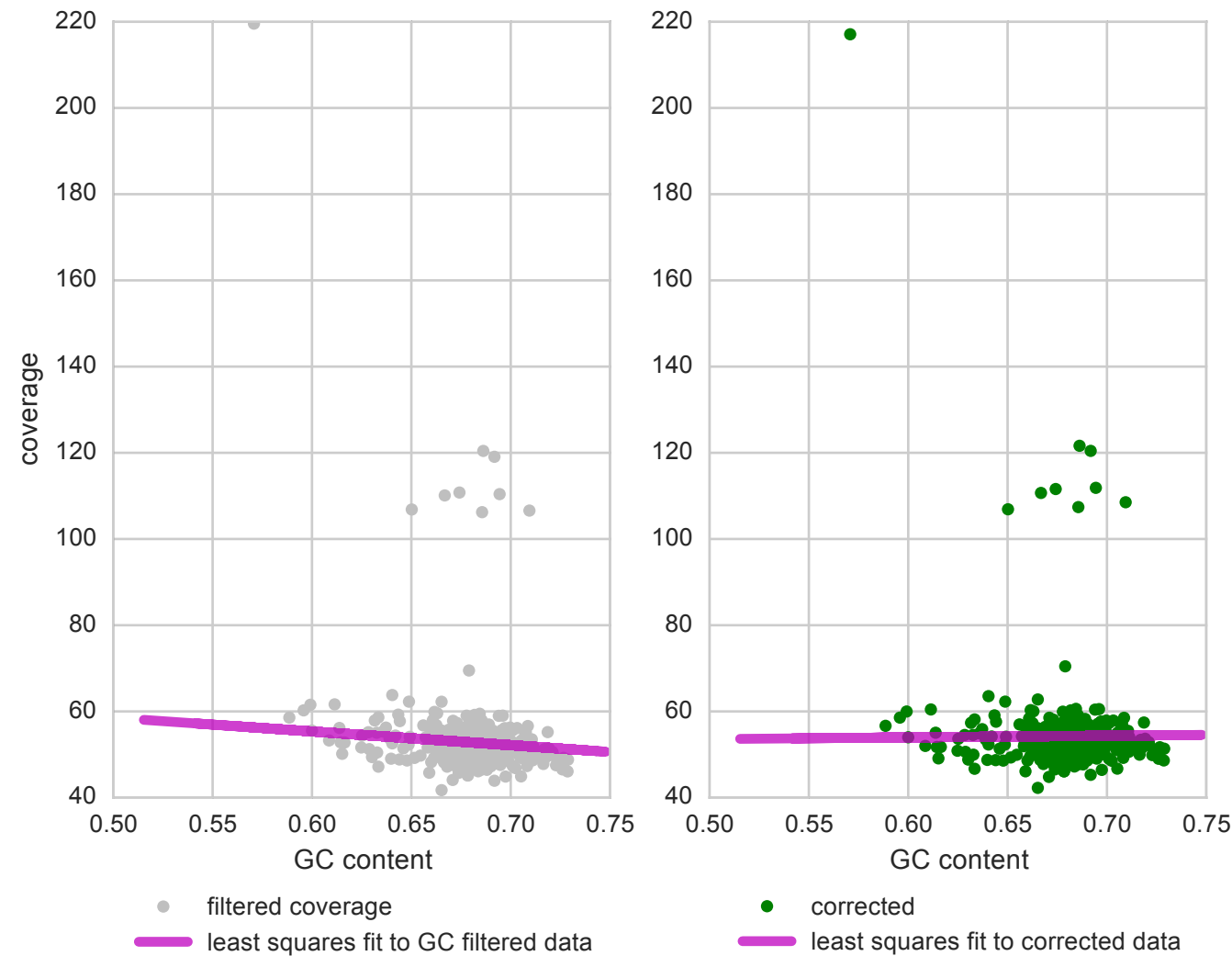
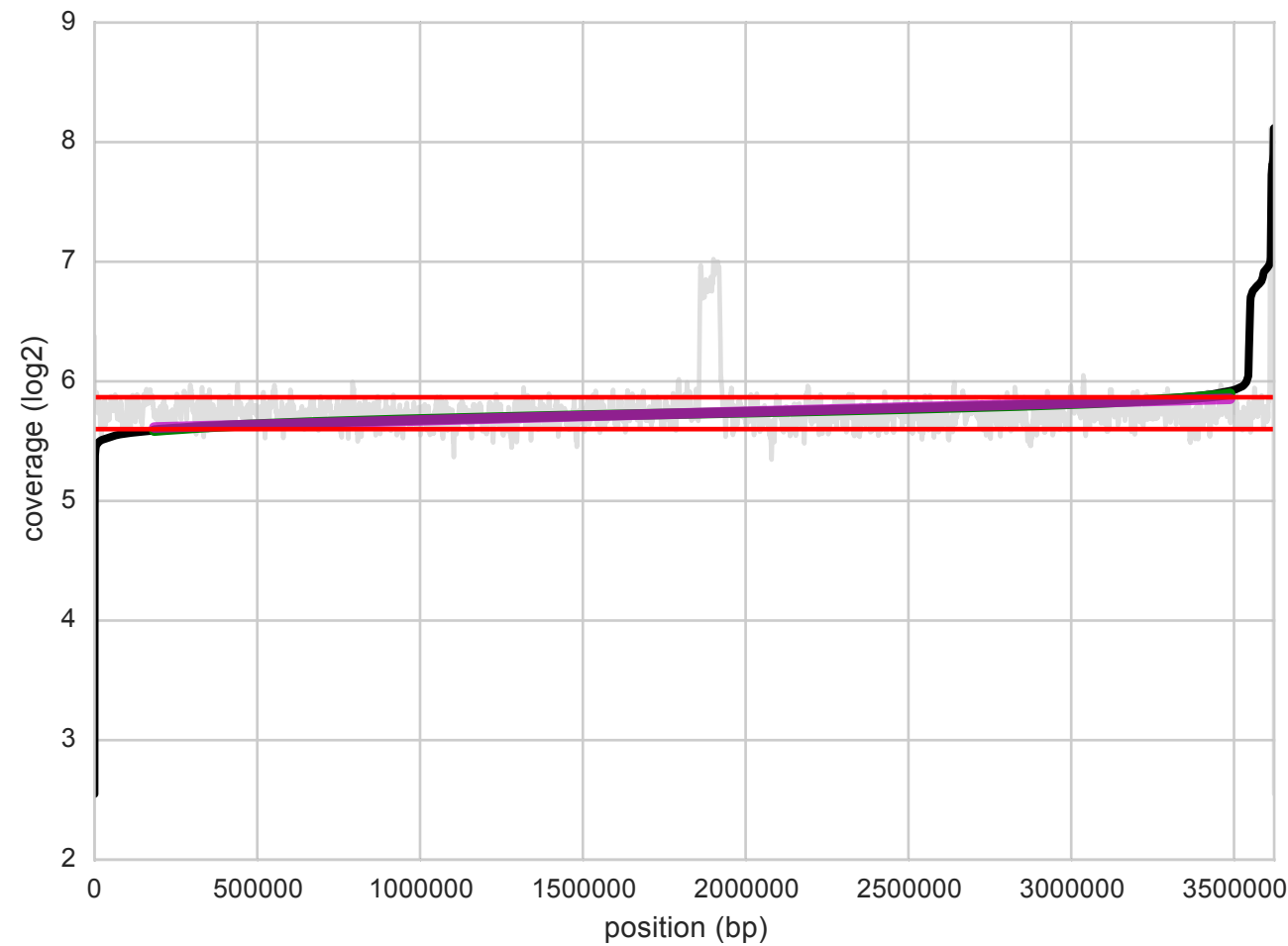


genome: /Users/WRShoemaker/github/Task2/LTDE/data/2016_KBSGenomes_Annotate/KBS0702/G-Chr1 sample: /Users/WRShoemaker/github/Task2/LTDE/data/iRep/KBS0702/GSF1046-KBS0702-E_S9_mapped_sort_NOdup_
iRep: 1.20 r^2: 0.98 avg. cov: 53.76 % windows: 99.92







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