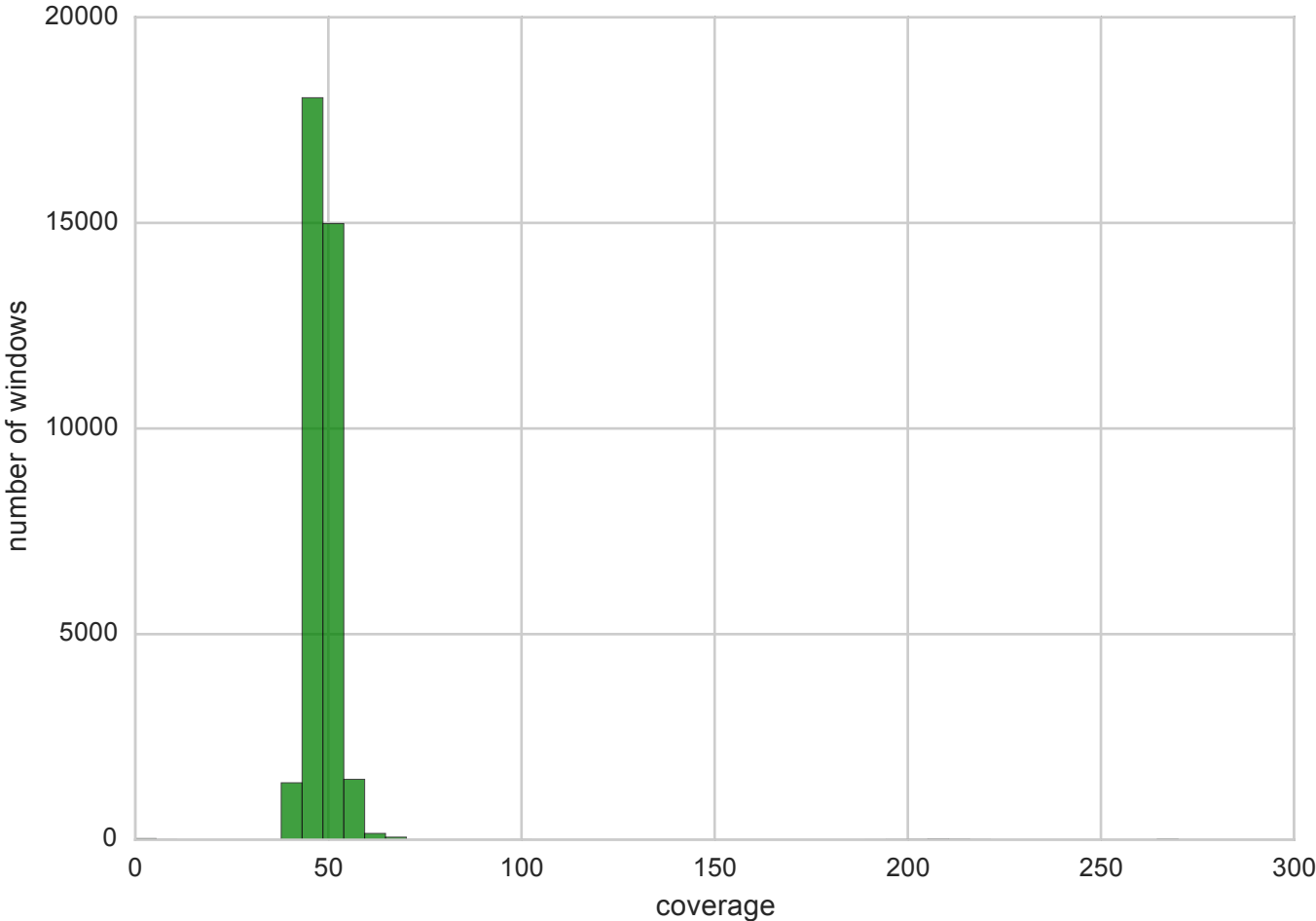
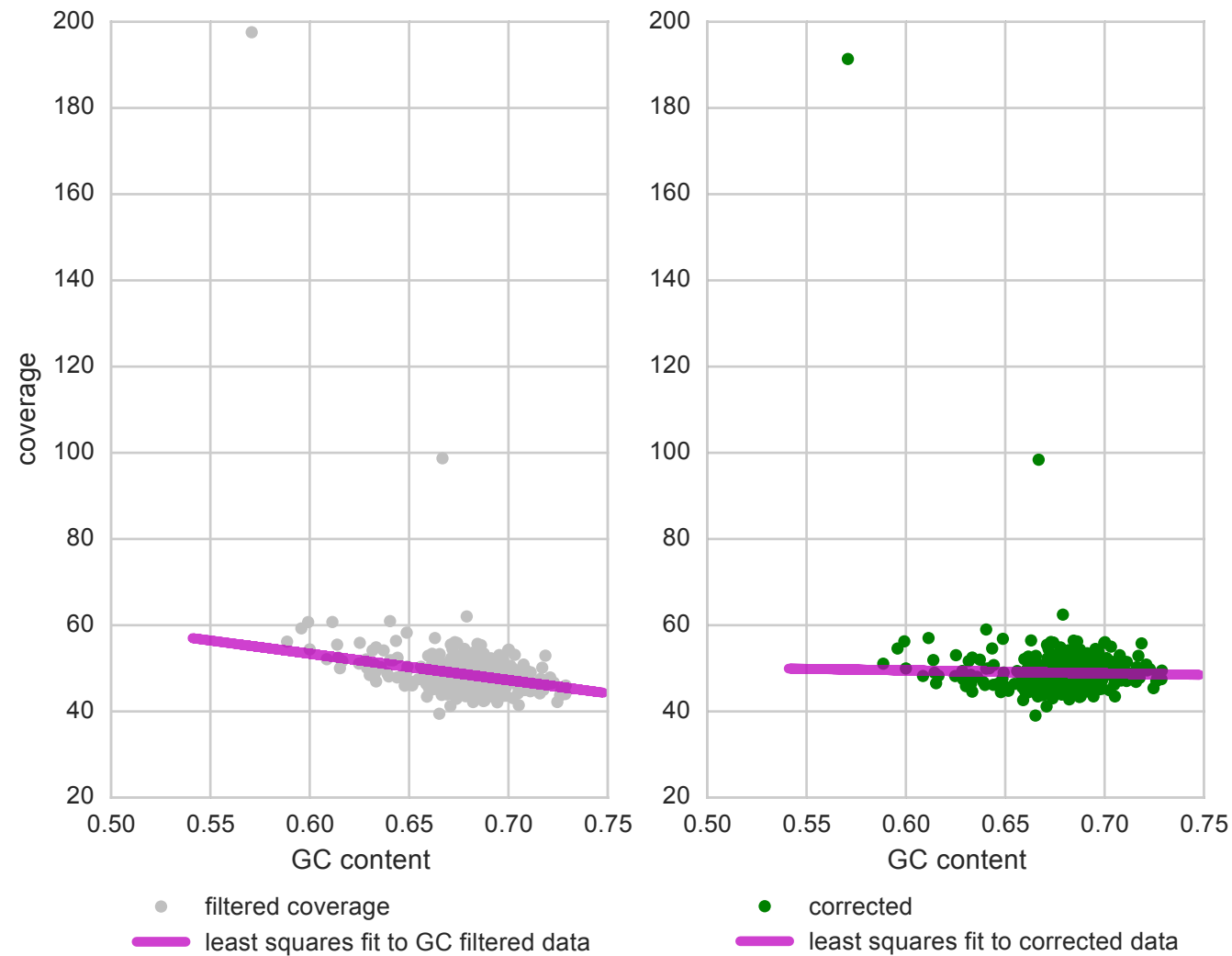
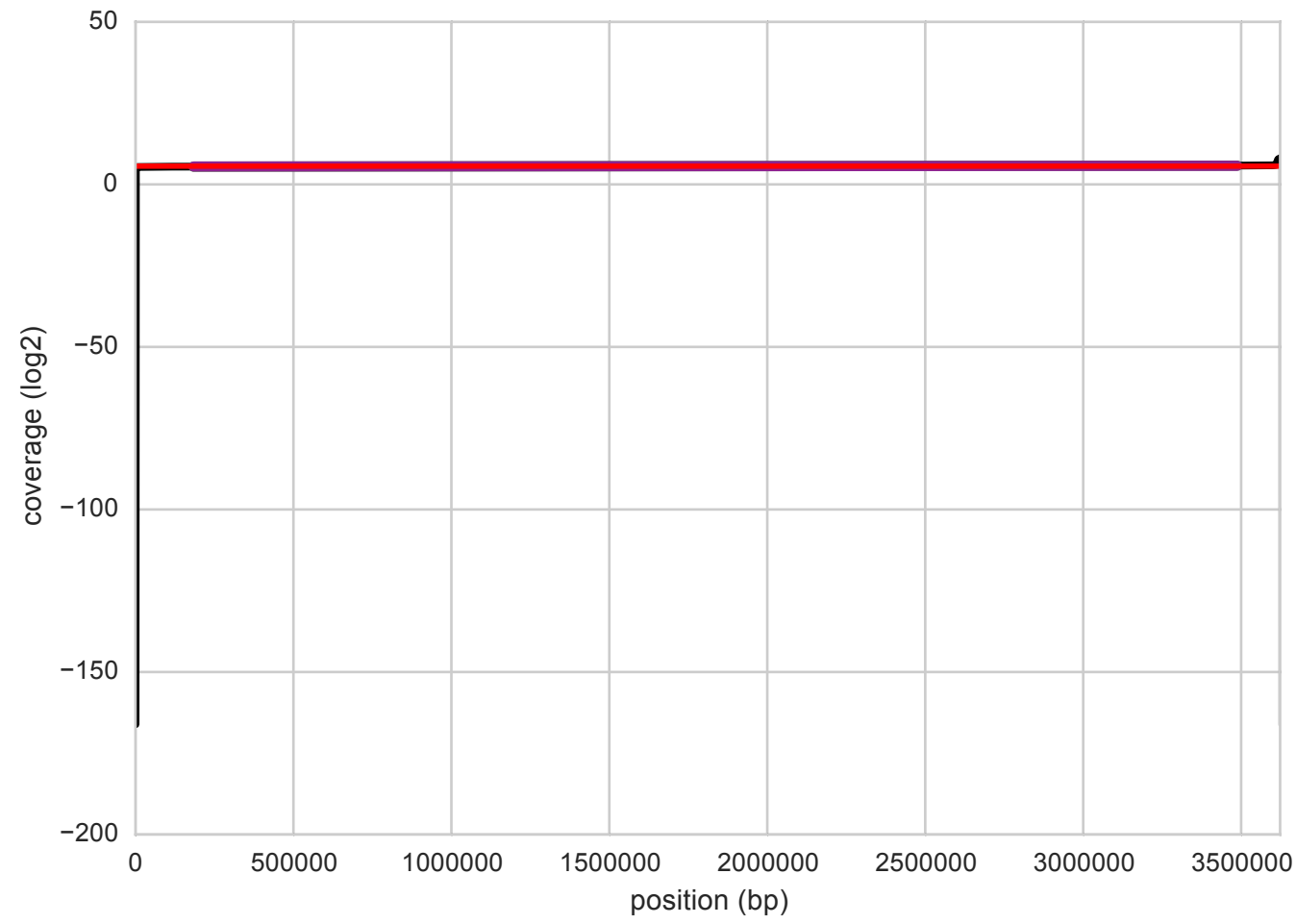


genome: /Users/WRShoemaker/github/Task2/LTDE/data/2016\_KBSGenomes\_Annotate/KBS0702/G-Chr1 sample: /Users/WRShoemaker/github/Task2/LTDE/data/iRep/KBS0702/GSF1046-KBS0702-F\_S16\_mapped\_sort\_NOdup\_  
iRep: 1.20 r^2: 0.99 avg. cov: 48.93 % windows: 99.92





genome: /Users/WRShoemaker/github/Task2/LTDE/data/2016\_KBSGenomes\_Annotate/KBS0702/G-Chr1 sample: /Users/WRShoemaker/github/Task2/LTDE/data/iRep/KBS0702/GSF1046-KBS0702-F\_S16\_mapped\_sort\_NOdup\_  
iRep: 1.20 un-filtered iRep: 1.20 r^2: 0.99 avg. cov: 48.93 % windows: 99.92



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