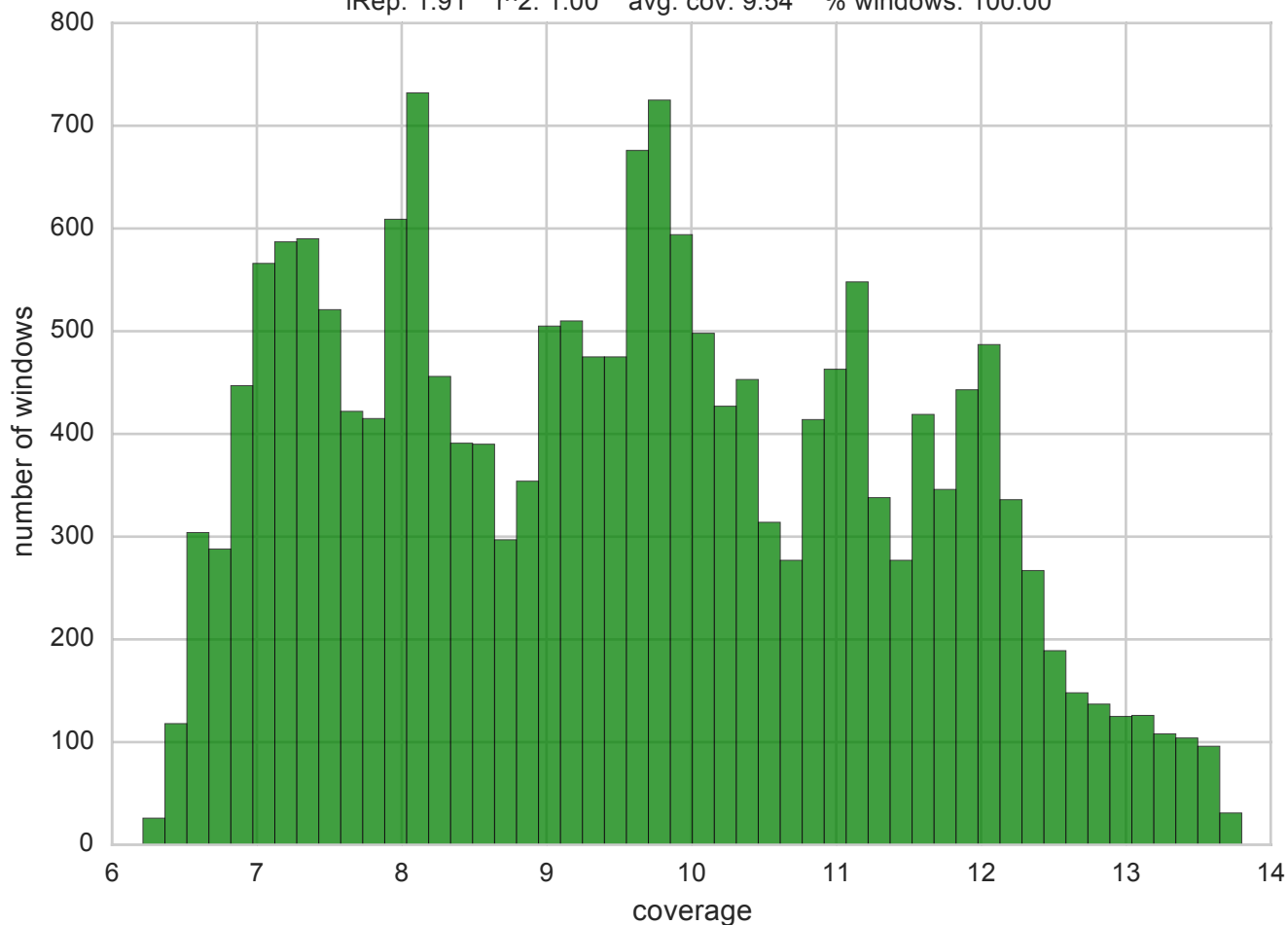


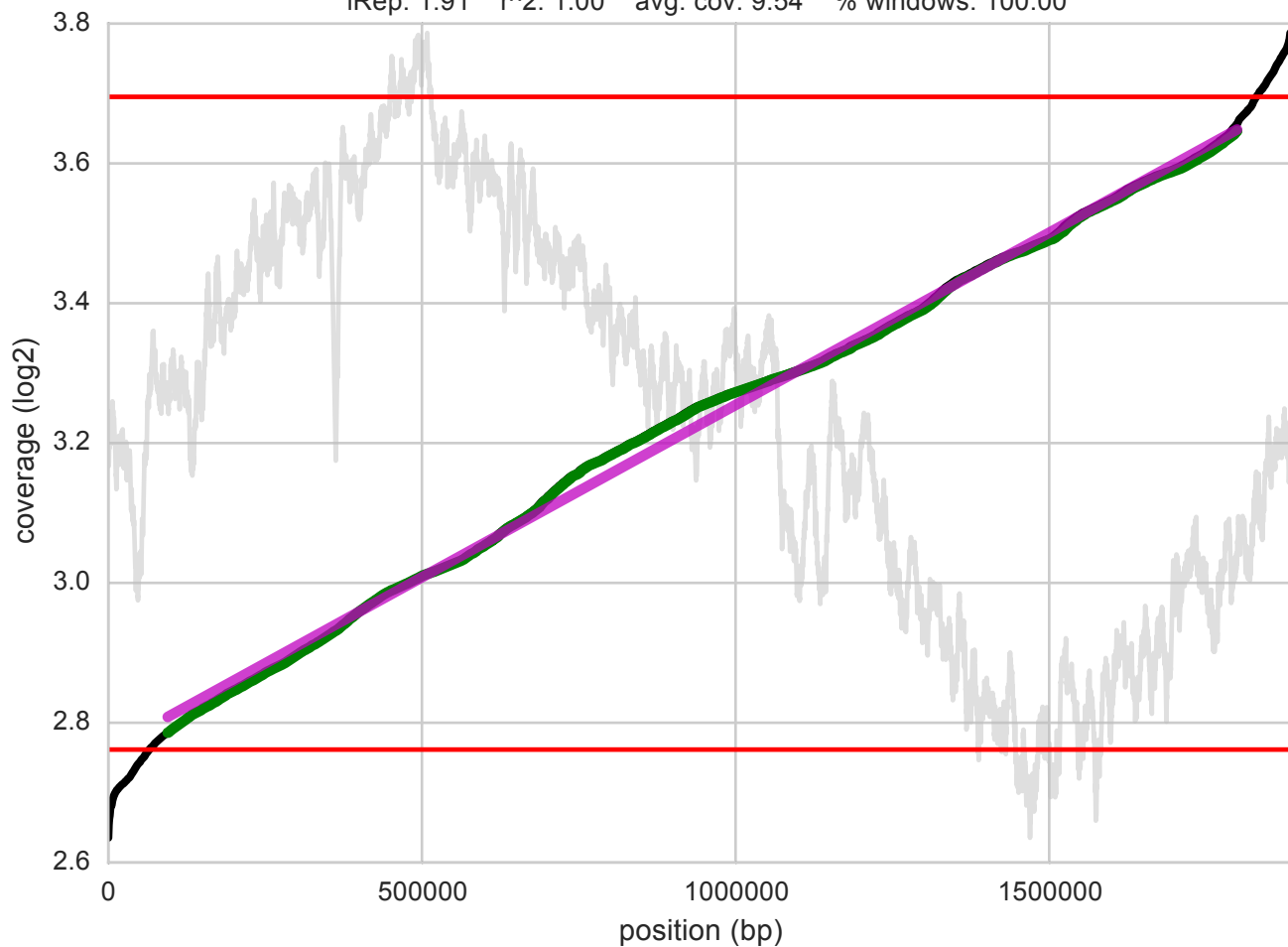
genome: sample_data/l_gasseri sample: sample_data/l_gasseri.fna-vs-l_gasseri_sample1-shrun

iRep: 1.91 r²: 1.00 avg. cov: 9.54 % windows: 100.00



genome: sample_data/l_gasseri sample: sample_data/l_gasseri.fna-vs-l_gasseri_sample1-shrun

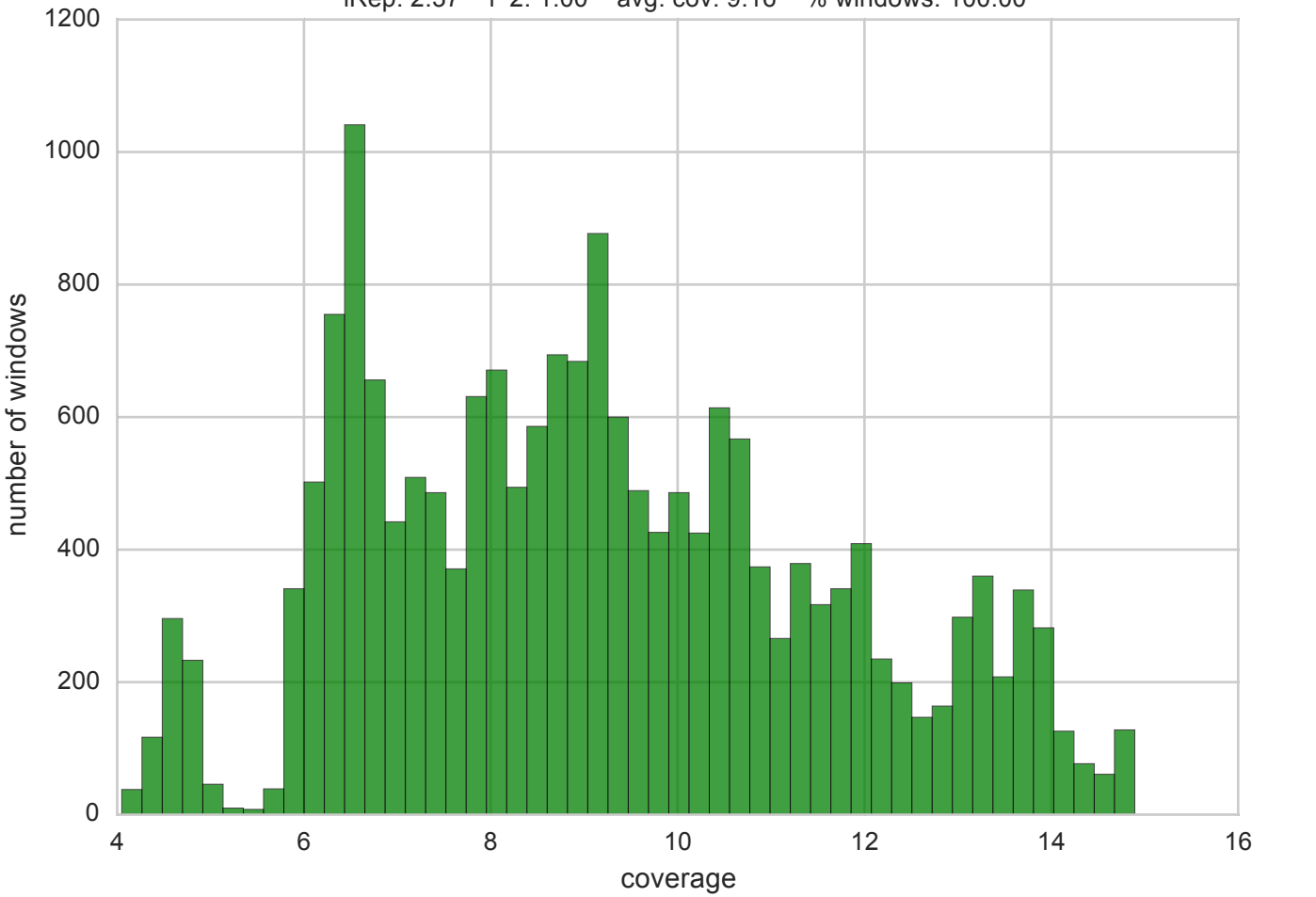
iRep: 1.91 r²: 1.00 avg. cov: 9.54 % windows: 100.00



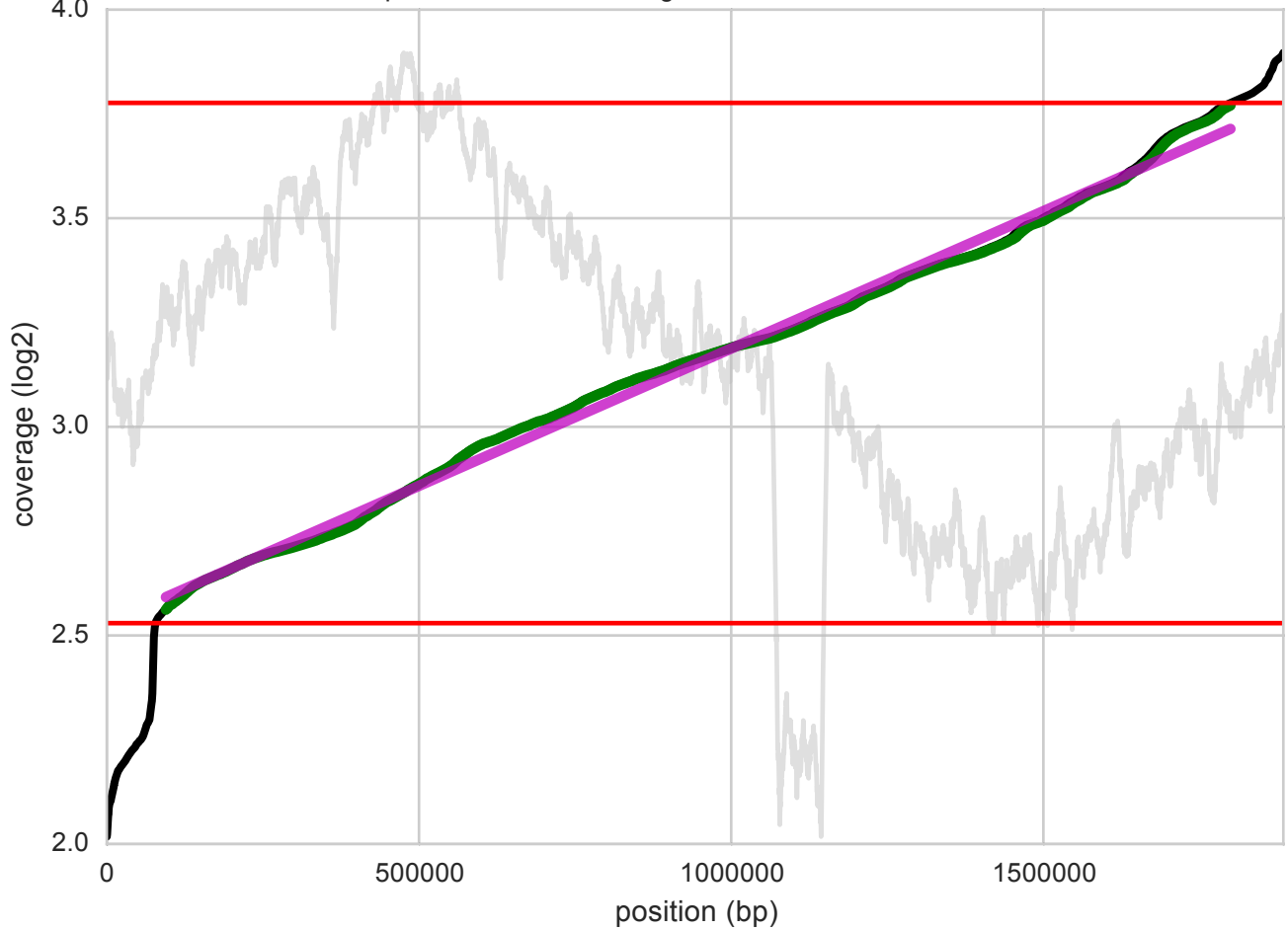
- filtered
- sorted
- trimmed
- least squares fit to trimmed
- Ori/Ter

genome: sample_data/l_gasseri sample: sample_data/l_gasseri.fna-vs-l_gasseri_sample2-shrun

iRep: 2.37 r^2: 1.00 avg. cov: 9.16 % windows: 100.00



iRep: 2.37 r^2: 1.00 avg. cov: 9.16 % windows: 100.00



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