Sample list1.jelly.27.50x.kmer.histo k= 27

Legend:
- Observed (obs.): 3607.265 Mb (error-excluded: 3177.172 Mb)
  - * k-mer_cov obs.: 50
  - * signal_error_border 17
- Fitted count with fitted k-mer_cov: 3216.625 Mb
  - * k-mer_cov fit.: 49.62
  - * 1st-sd:10.91
  - * 1st-skewness: 1.05
- Fitted+obs. with corrected k-mer_cov: 3140.132 Mb
  - * k-mer_cov cor.: 50.6
  - * repetitive_ratio 0.24: 747.055 Mb

Y-axis: Number of k-mers

X-axis: K-mer freq
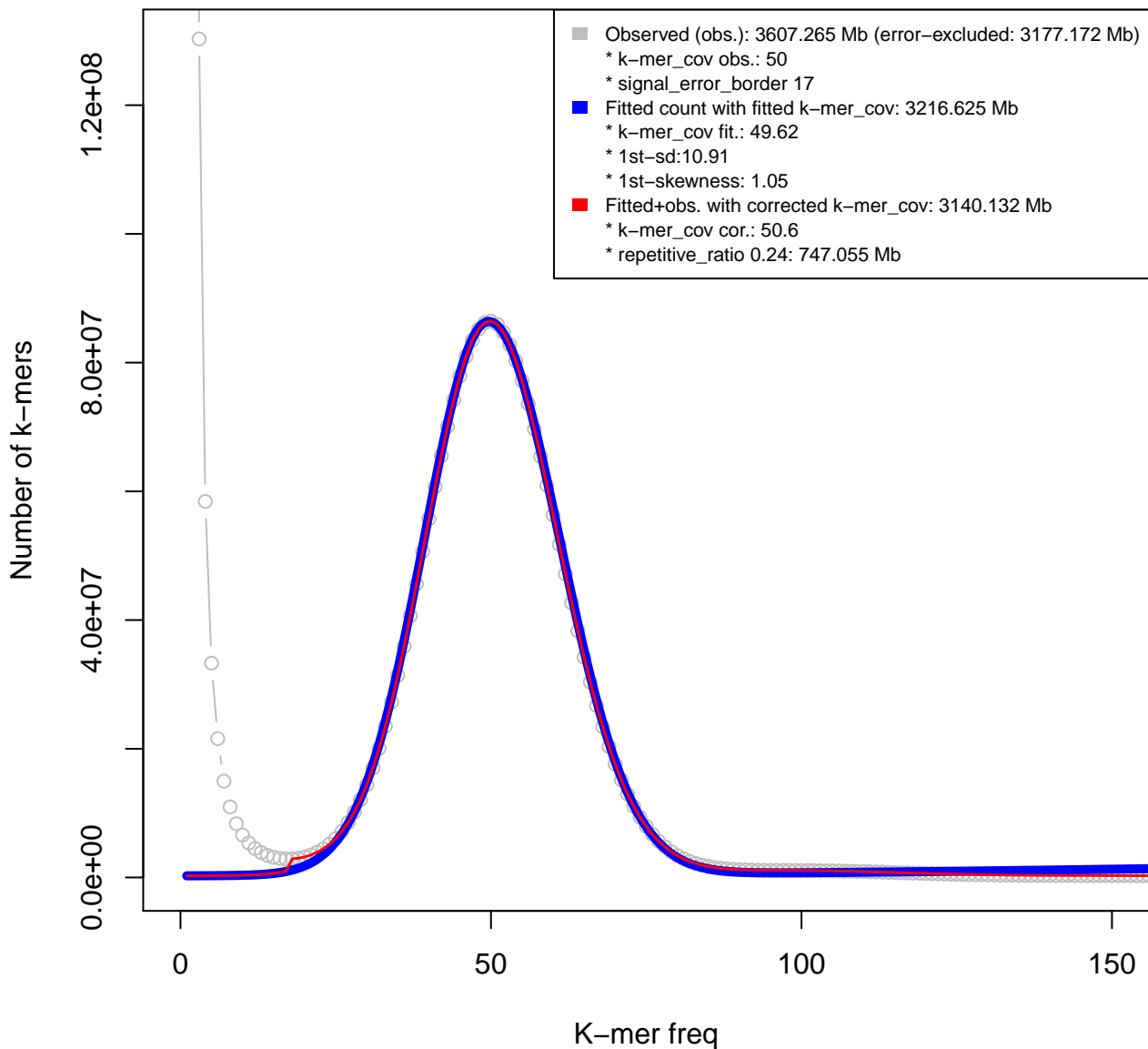
Genome size estimation by error-excluding, fitting, and correcting

Legend:
- EXL: 3177.172 Mb
- FIT: 3216.625 Mb
- COR: 3140.132 Mb

Y-axis: Genome size (bp)

X-axis: Size k

27