Sample list1.jelly.27.25x.kmer.histo k= 27

Legend:
- Observed (obs.): 3698.426 Mb (error-excluded: 3296.217 Mb)
  - * k-mer_cov obs.: 24
  - * signal_error_border 7
- Fitted count with fitted k-mer_cov: 3041.414 Mb
  - * k-mer_cov fit.: 23.8
  - * 1st-sd:7.07
  - * 1st-skewness: 1.11
- Fitted+obs. with corrected k-mer_cov: 3129.4 Mb
  - * k-mer_cov cor.: 25.28
  - * repetitive_ratio 0.24: 747.172 Mb

X-axis: K-mer freq

Y-axis: Number of k-mers
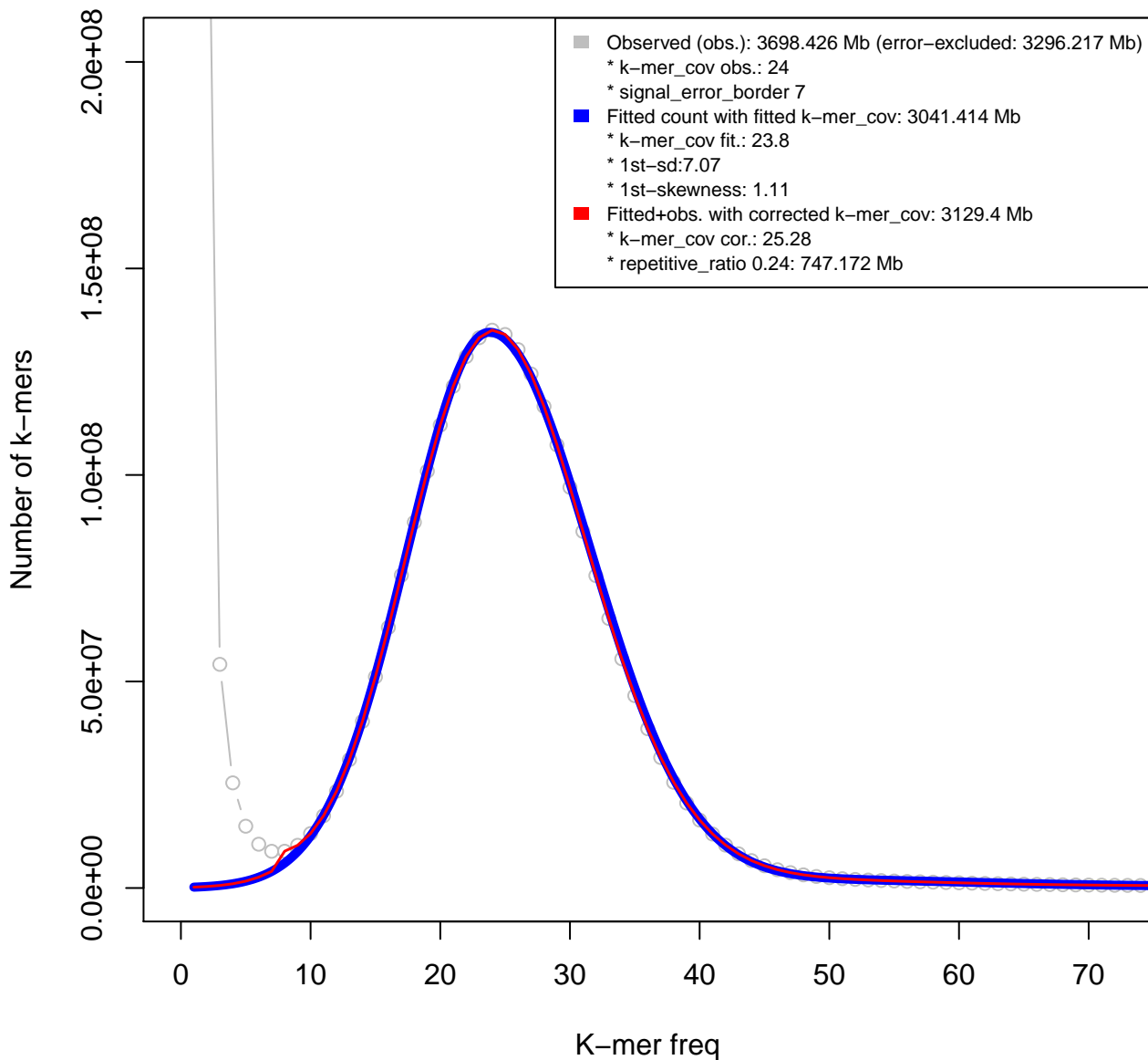
Genome size estimation by error−excluding, fitting, and correcting

Legend:
- EXL: 3296.217 Mb
- FIT: 3041.414 Mb
- COR: 3129.4 Mb

Y-axis: Genome size (bp)

X-axis: Size k