Sample universeK21.50x.histo k= 21

Legend:
- Observed (obs.): 0.434 Mb (error−excluded: 0.434 Mb)
  - * k−mer_cov obs.: 43
  - * signal_error_border 10
- Fitted count with fitted k−mer_cov: 0.458 Mb
  - * k−mer_cov fit.: 43.53
  - * 1st−sd:6.61
  - * 1st−skewness: 1.07
- Fitted+obs. with corrected k−mer_cov: 0.42 Mb
  - * k−mer_cov cor.: 44.4
  - * repetitive_ratio 0.02: 0.01 Mb

X-axis: K−mer freq

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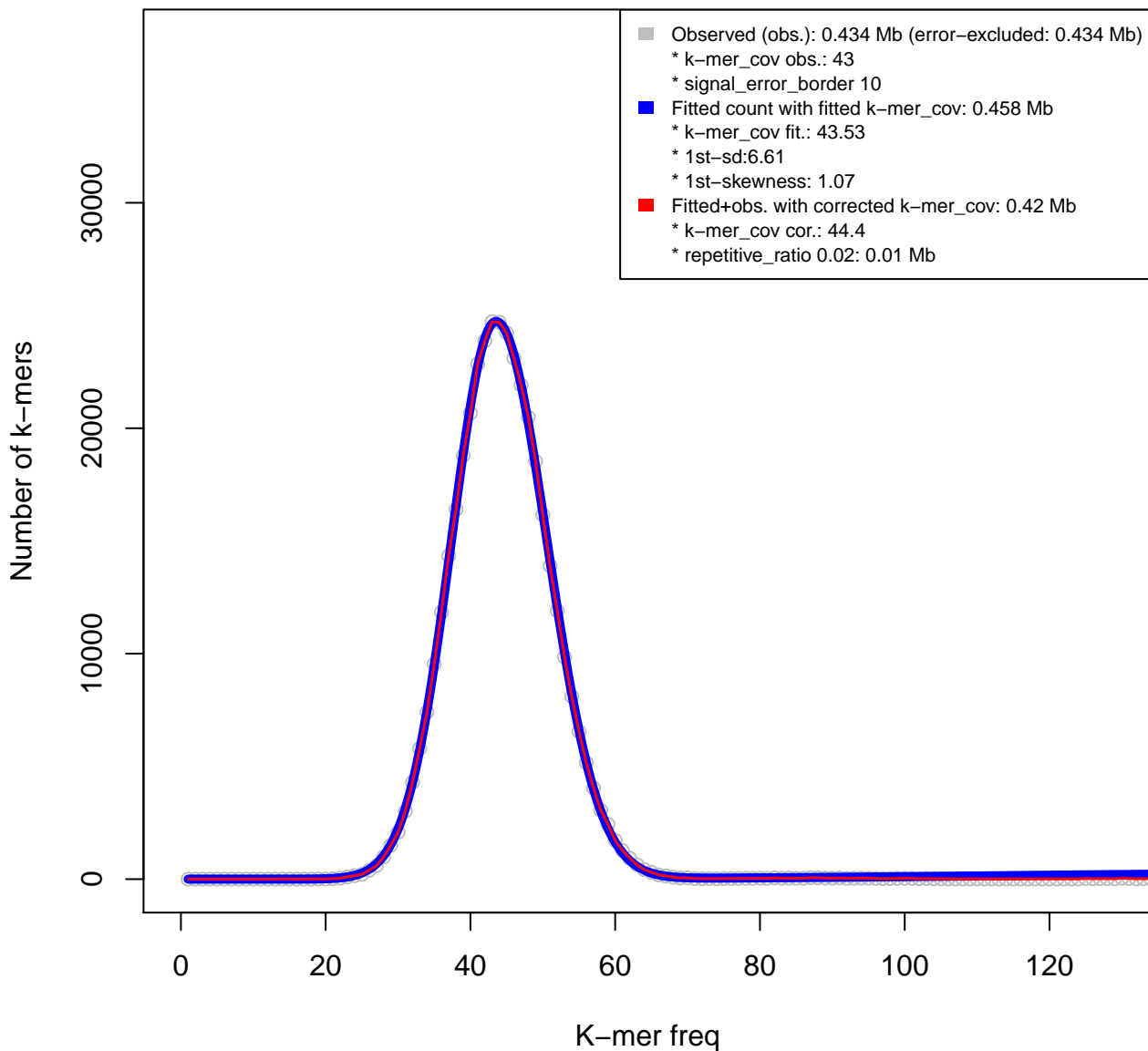
Genome size estimation by error−excluding, fitting, and correcting

Legend:
- EXL: 0.434 Mb
- FIT: 0.458 Mb
- COR: 0.42 Mb

Genome size (bp)

2¹

Size k