Sample sim_k27_4.histo k= 27

Legend:
- Observed (obs.): 152.095 Mb (error−excluded: 139.575 Mb)
  - * k−mer_cov obs.: 29
  - * signal_error_border 11
- Fitted count with fitted k−mer_cov: 127.745 Mb
  - * k−mer_cov fit.: 28.75
  - * 1st−sd:5.44
  - * 1st−skewness: 1.08
- Fitted+obs. with corrected k−mer_cov: 135.589 Mb
  - * k−mer_cov cor.: 29.86
  - * repetitive_ratio 0.14: 18.385 Mb

Y-axis: Number of k-mers

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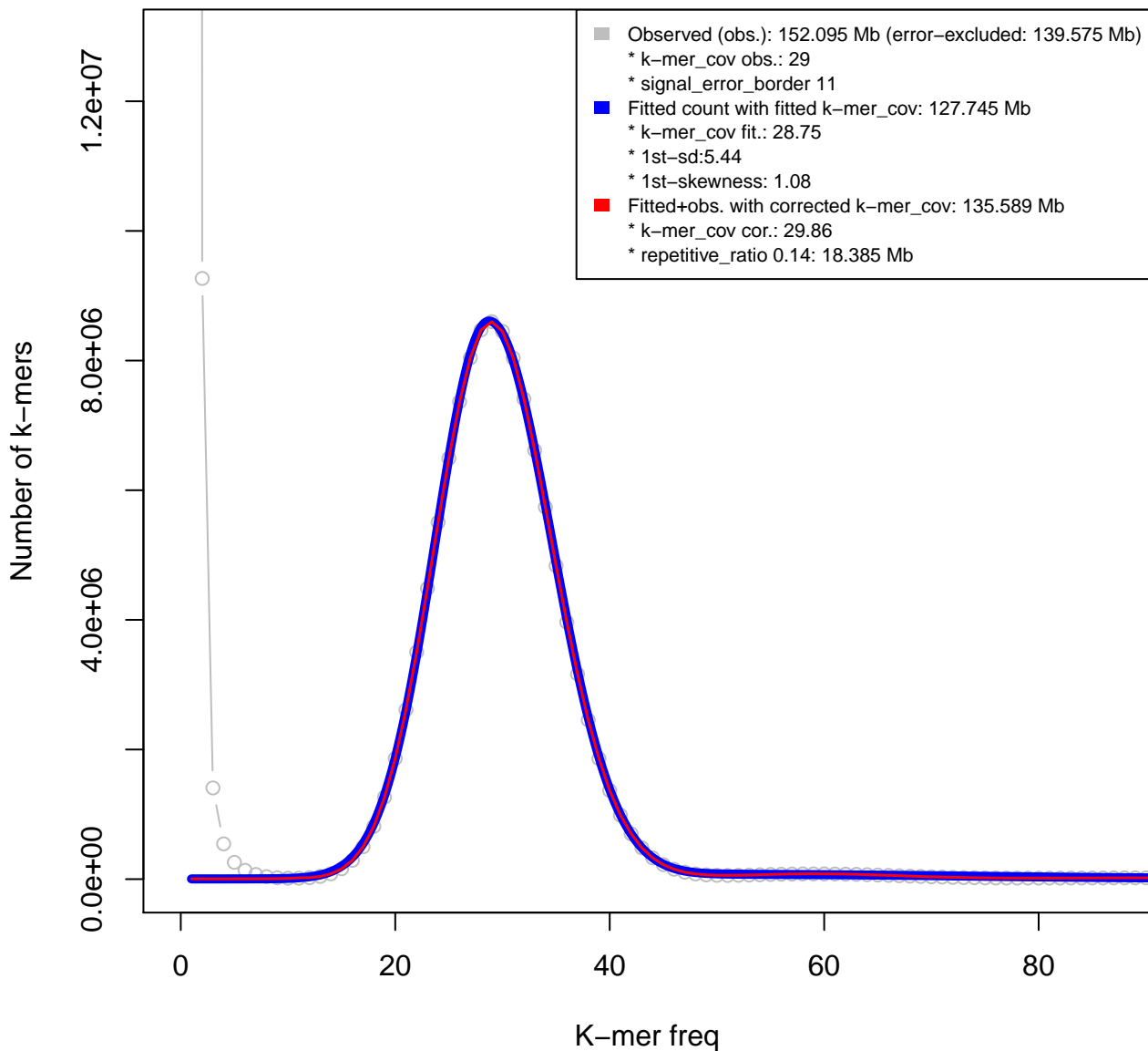
Genome size estimation by error−excluding, fitting, and correcting