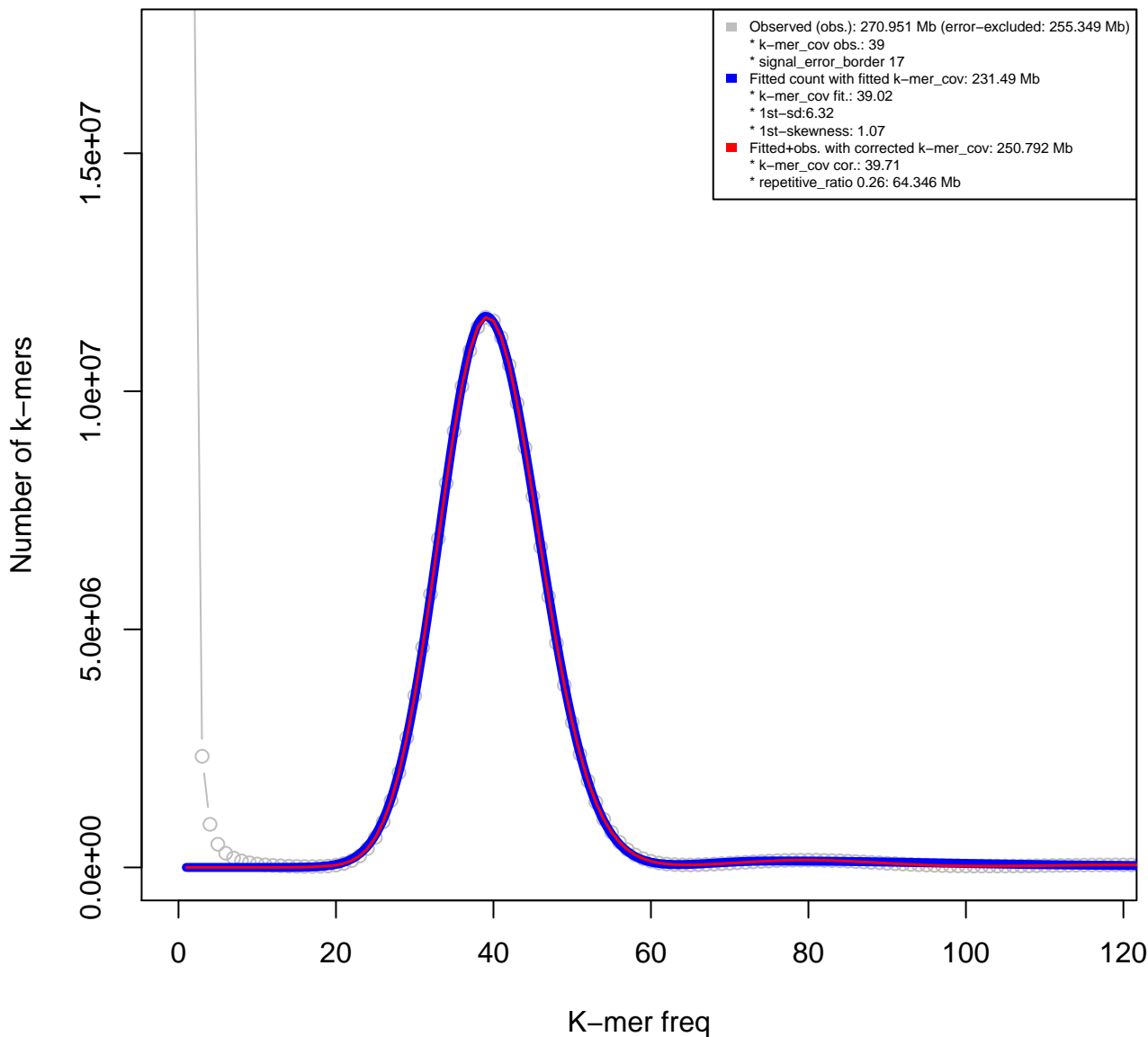


Sample K\_21\_4.histo k=21



# Genome size estimation by error-excluding, fitting, and correcting

