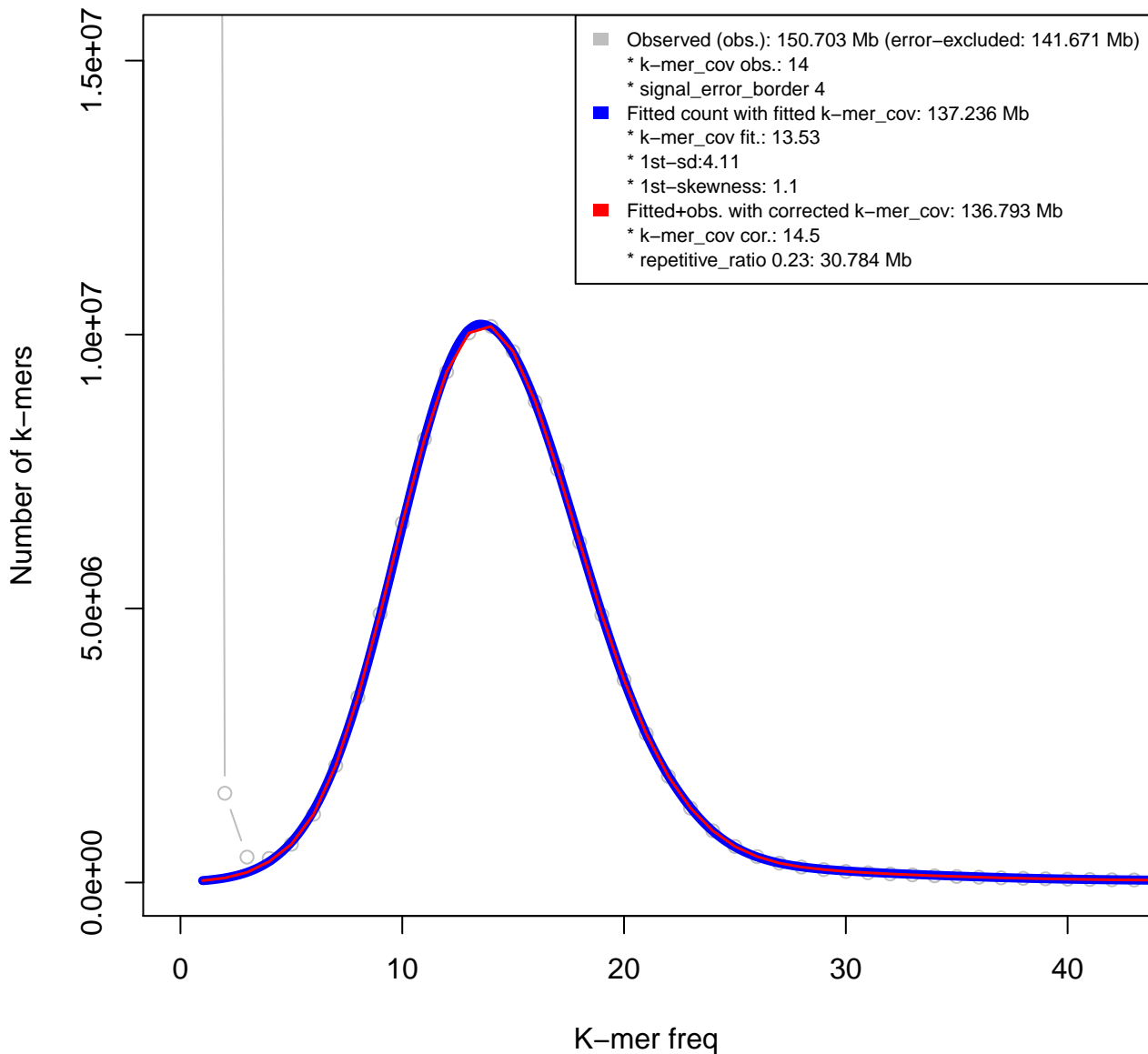


# Sample 6909.21mer.histo k= 21



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

