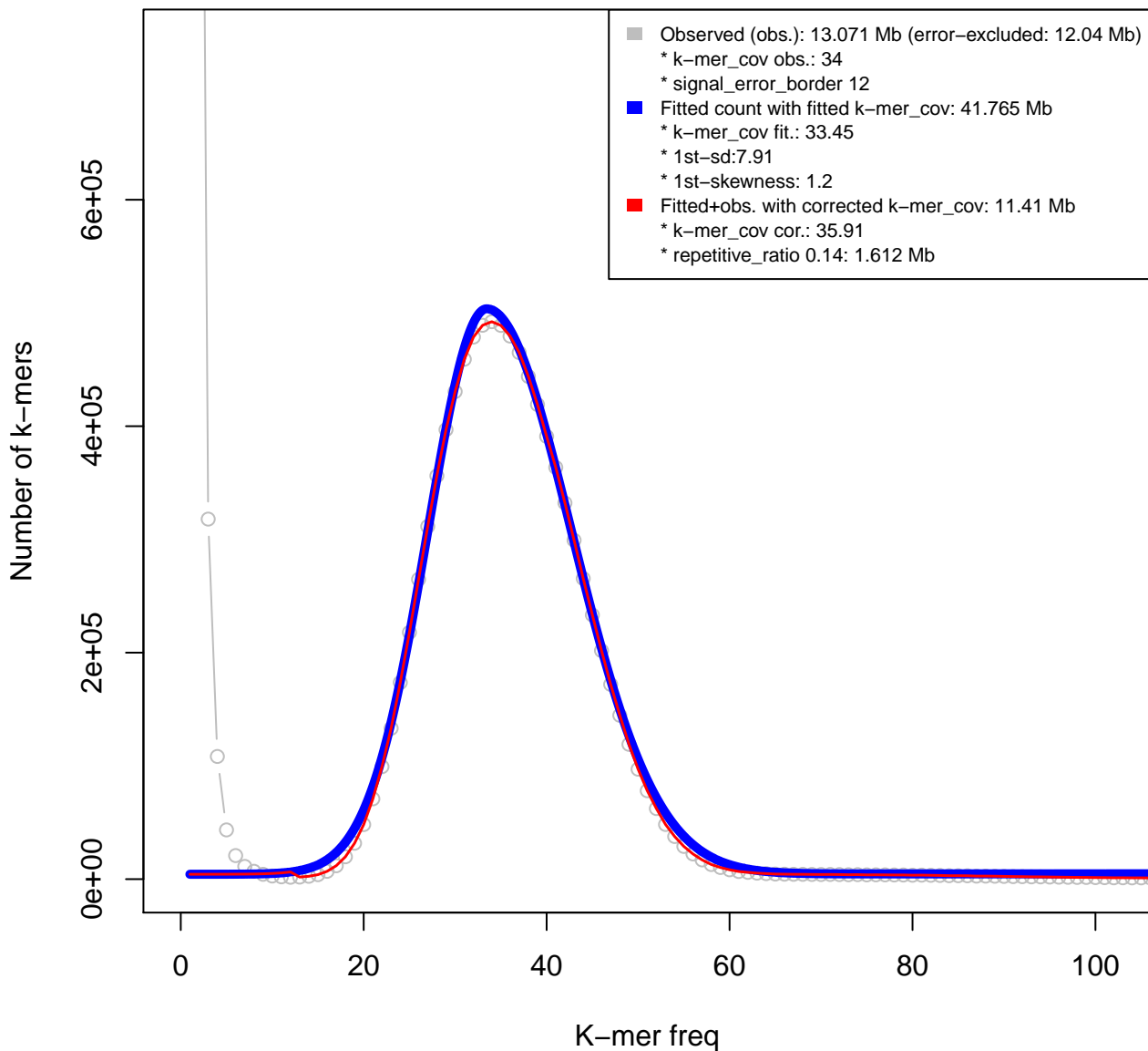


Sample sim_m23_5.histo k= 23



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

