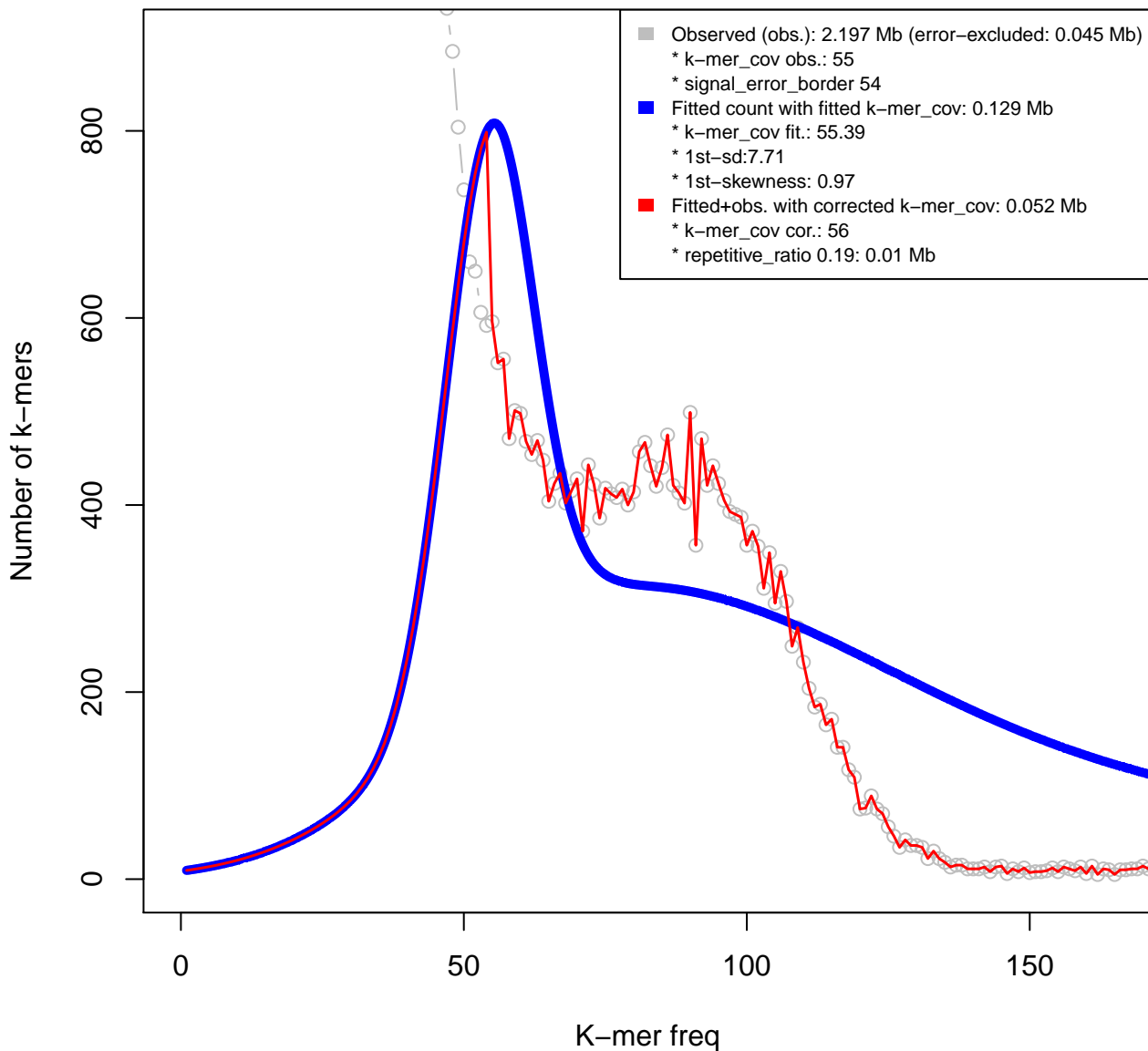


Sample sim_k25_2.histo k= 25



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

