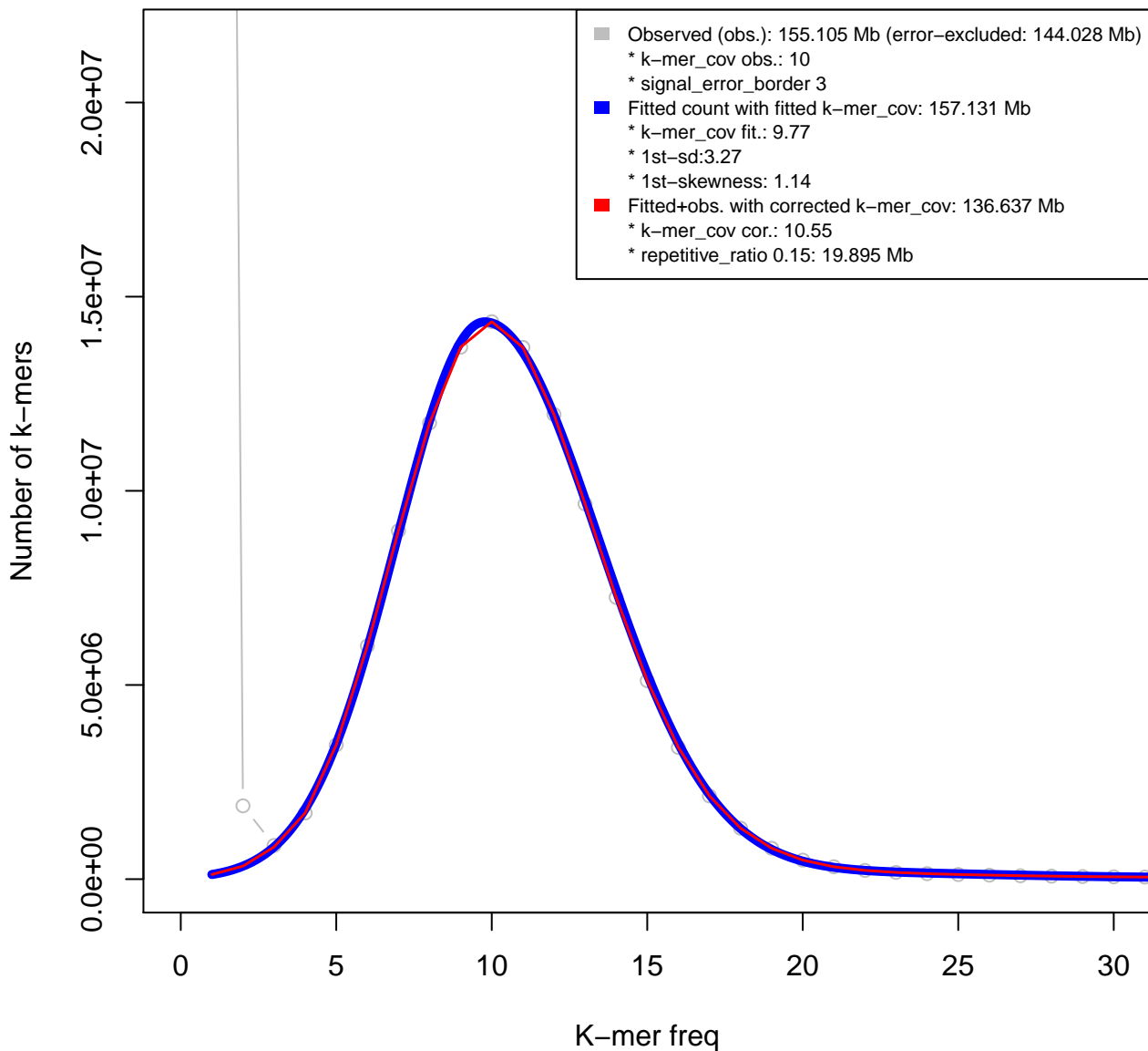


Sample sim_k23_2.histo k= 23



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

