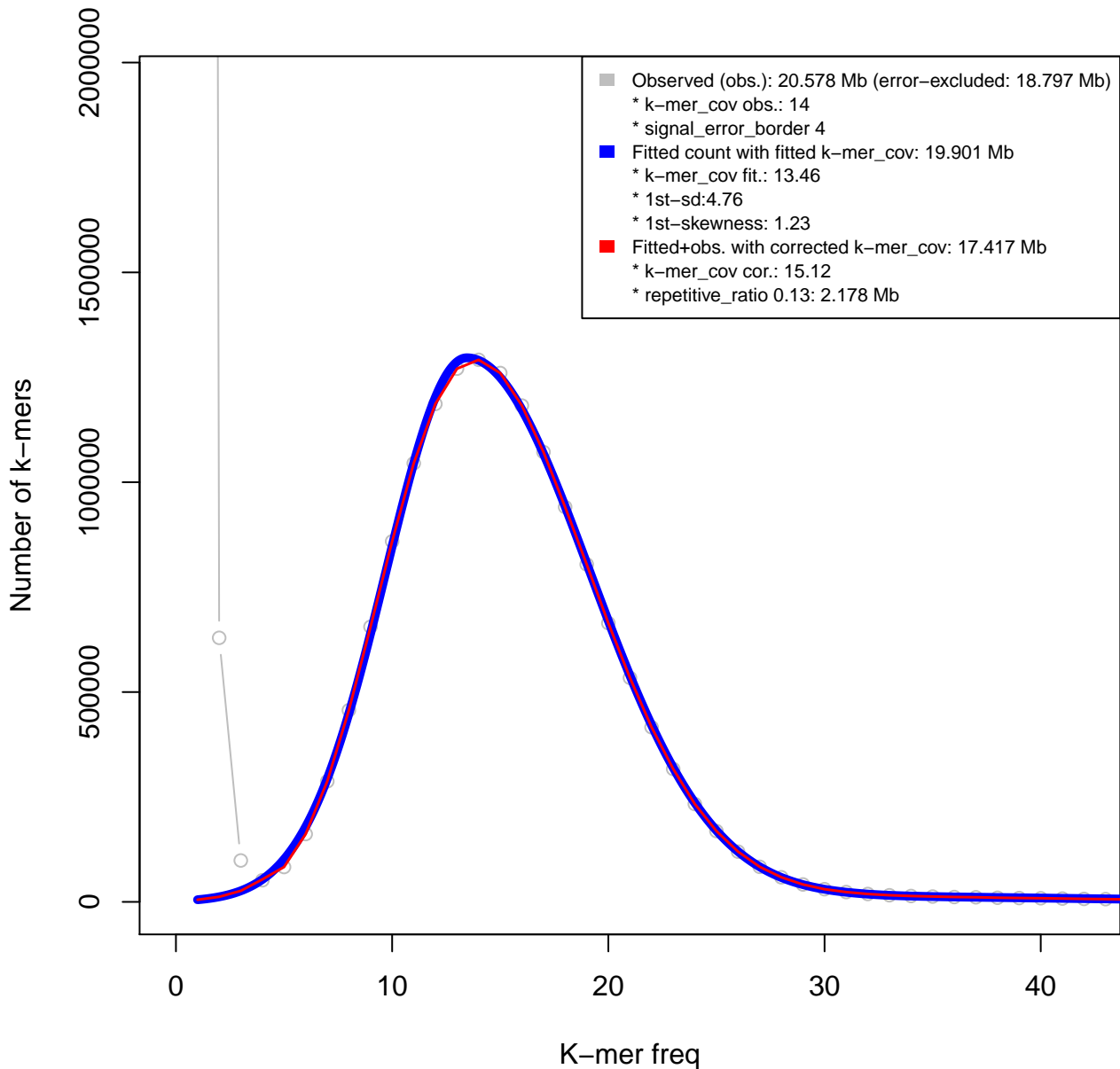


Sample M_27_2.histo k= 27



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

