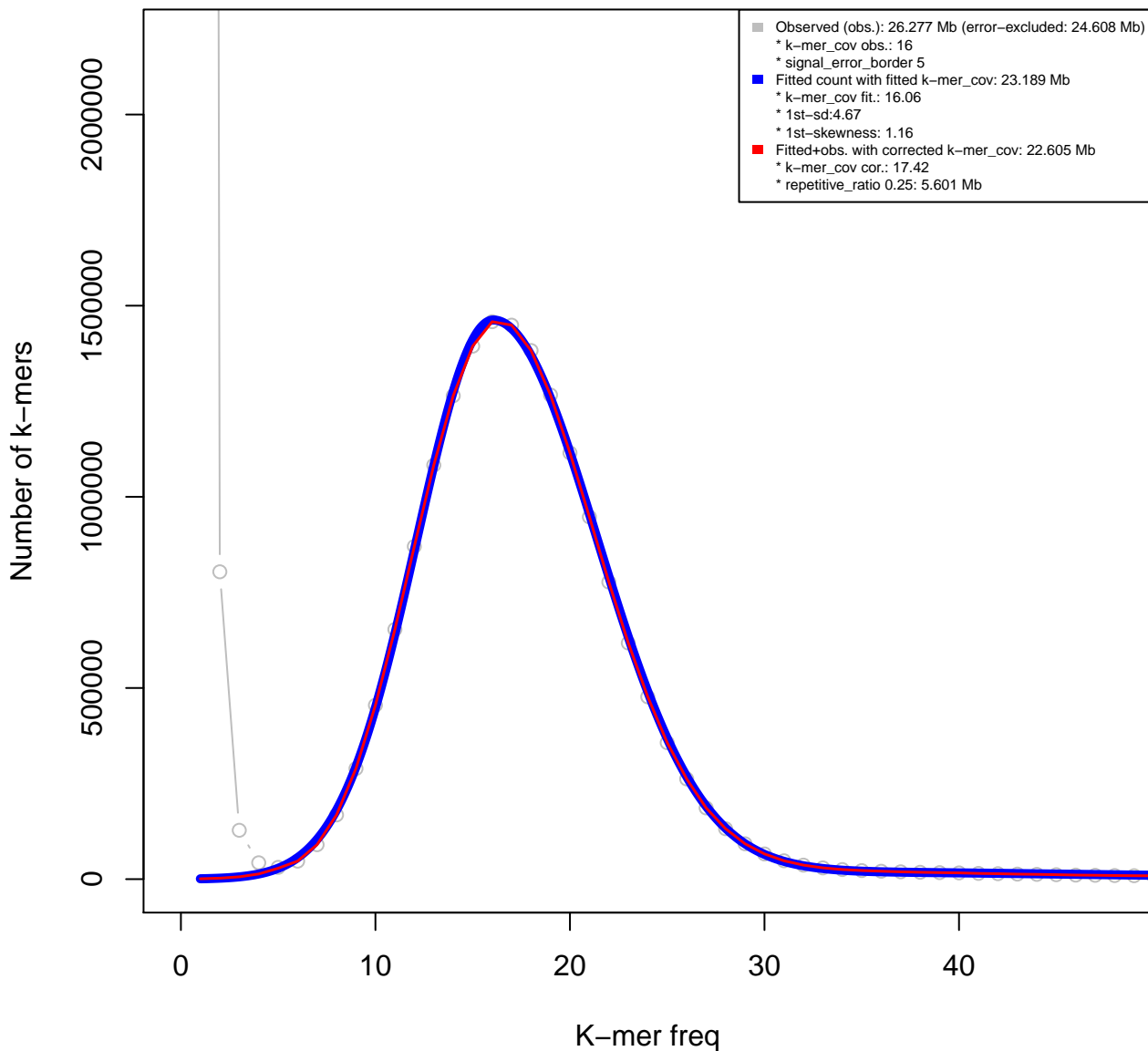


Sample M_21_2.histo k= 21



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

