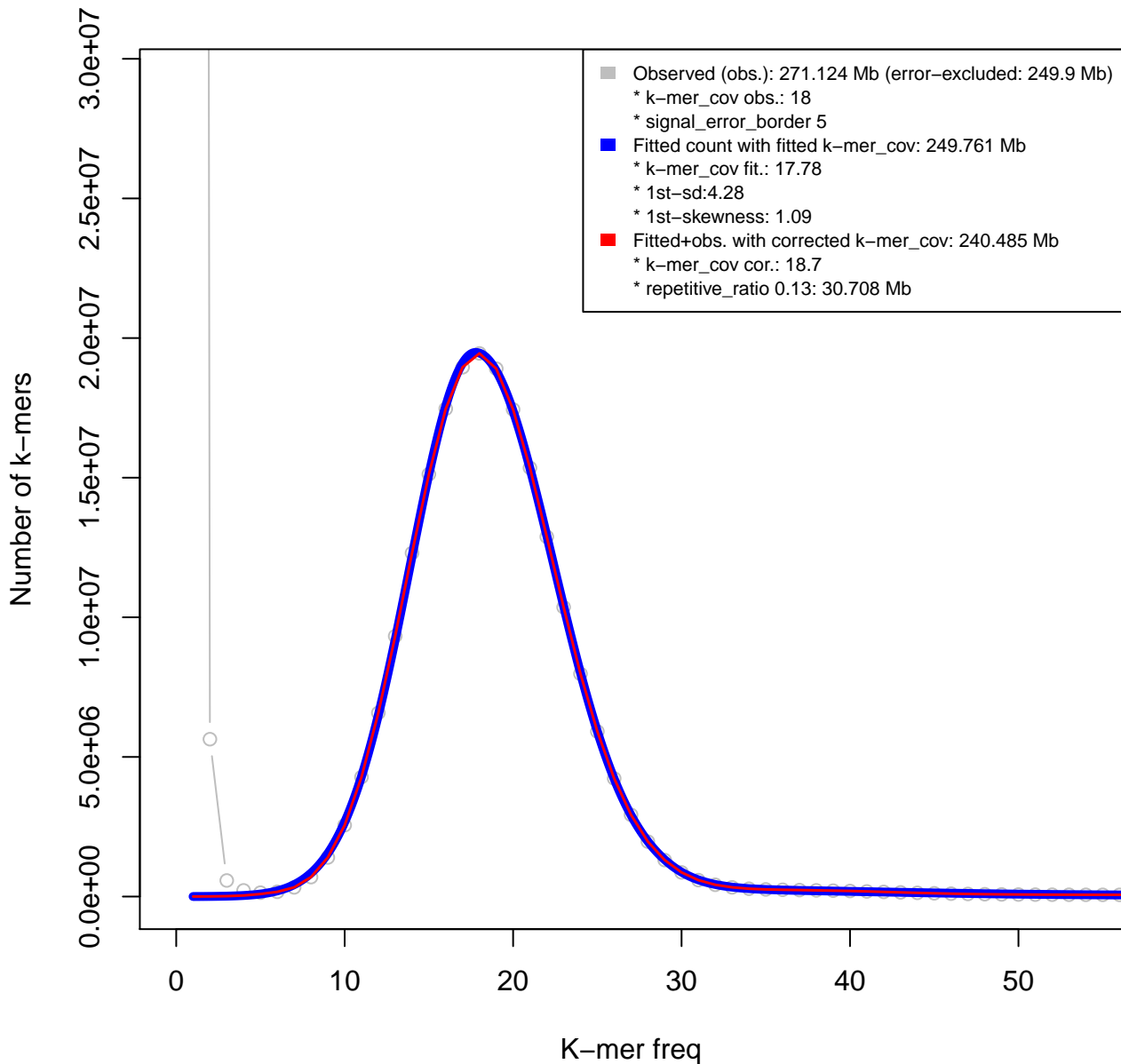


Sample K_27_2.histo k=27



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

