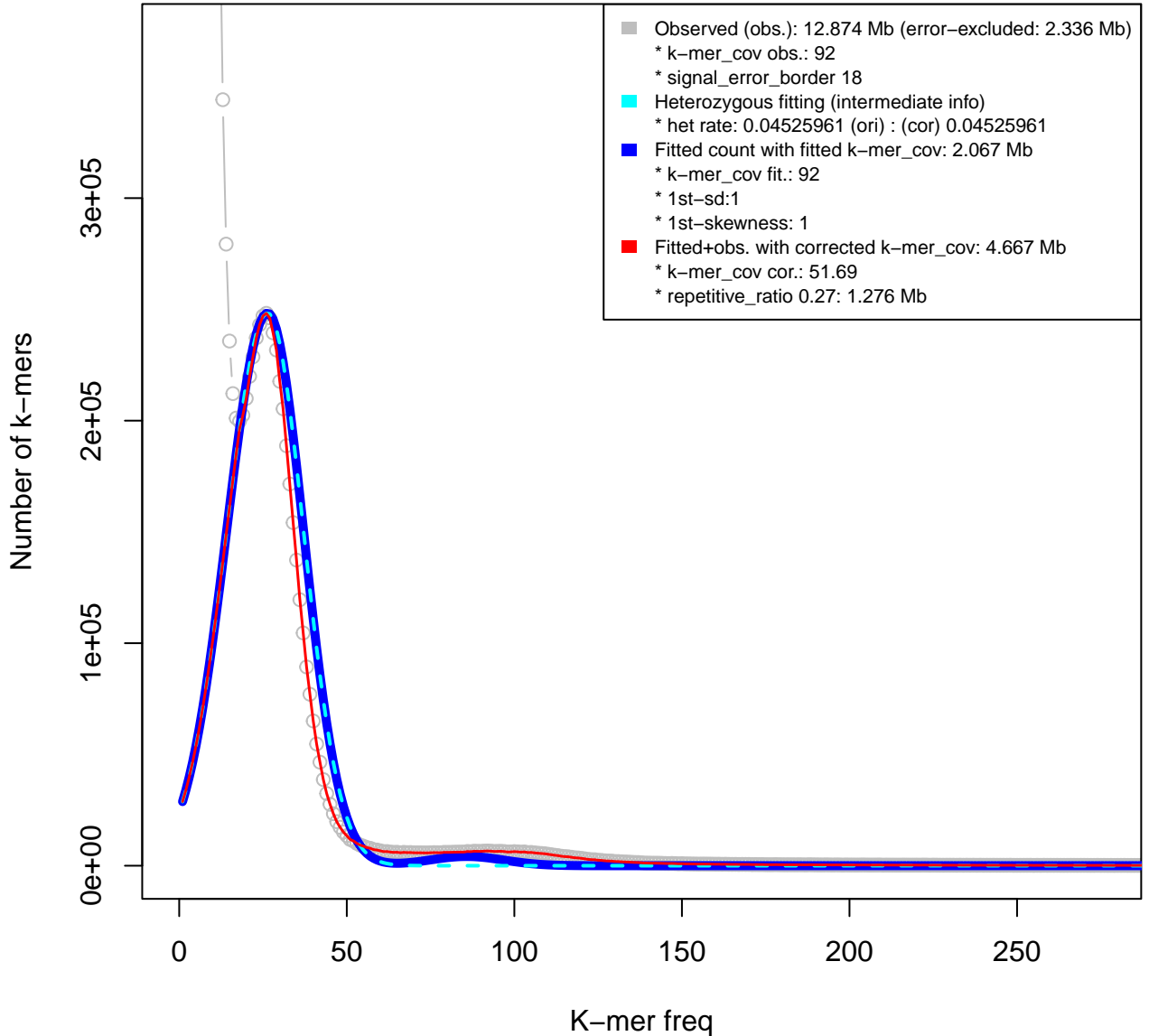


Sample 21mer_no_cut.histo k= 21



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

