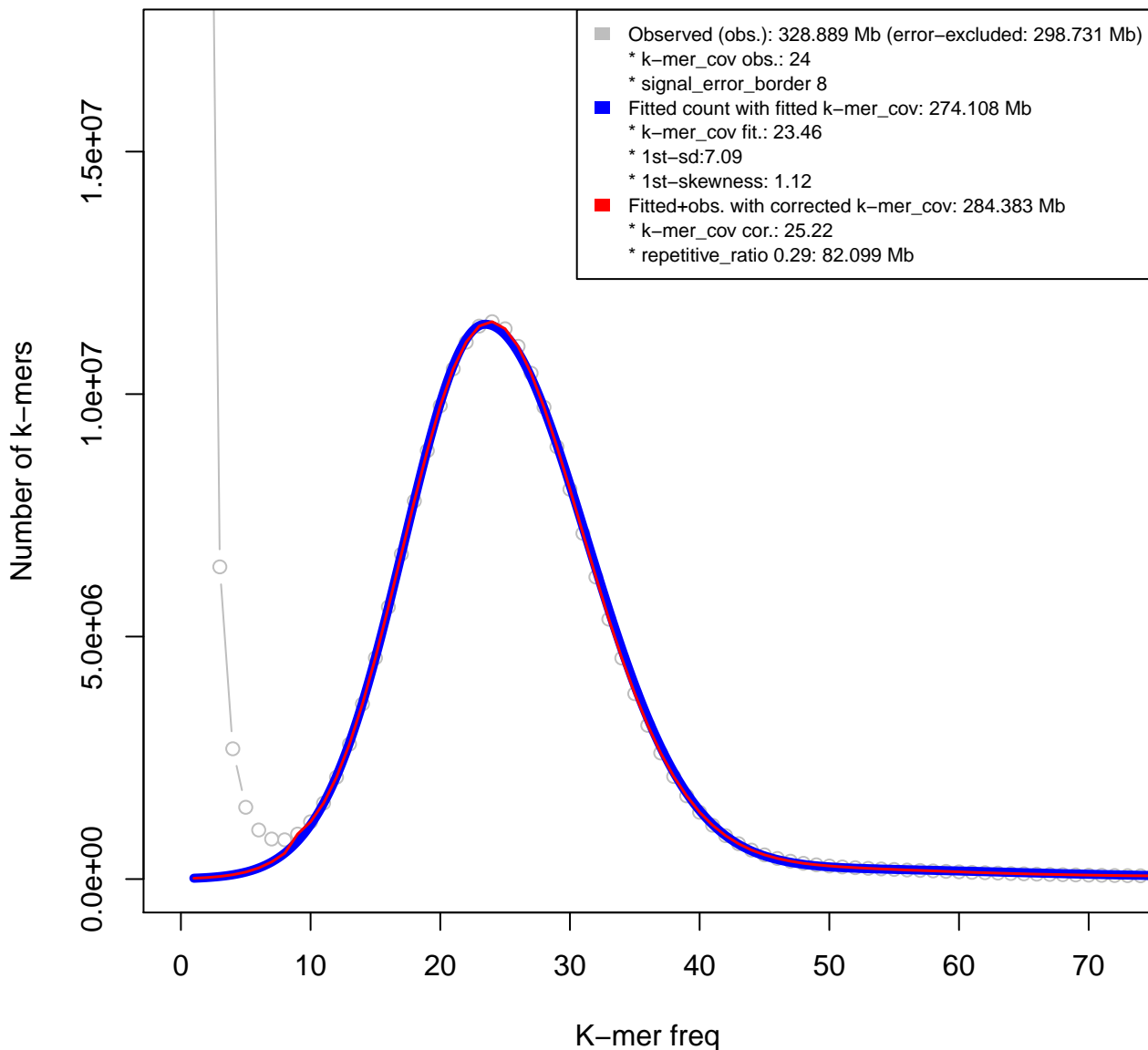


Sample list2.21.25x.minimizer.histo k= 21



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

