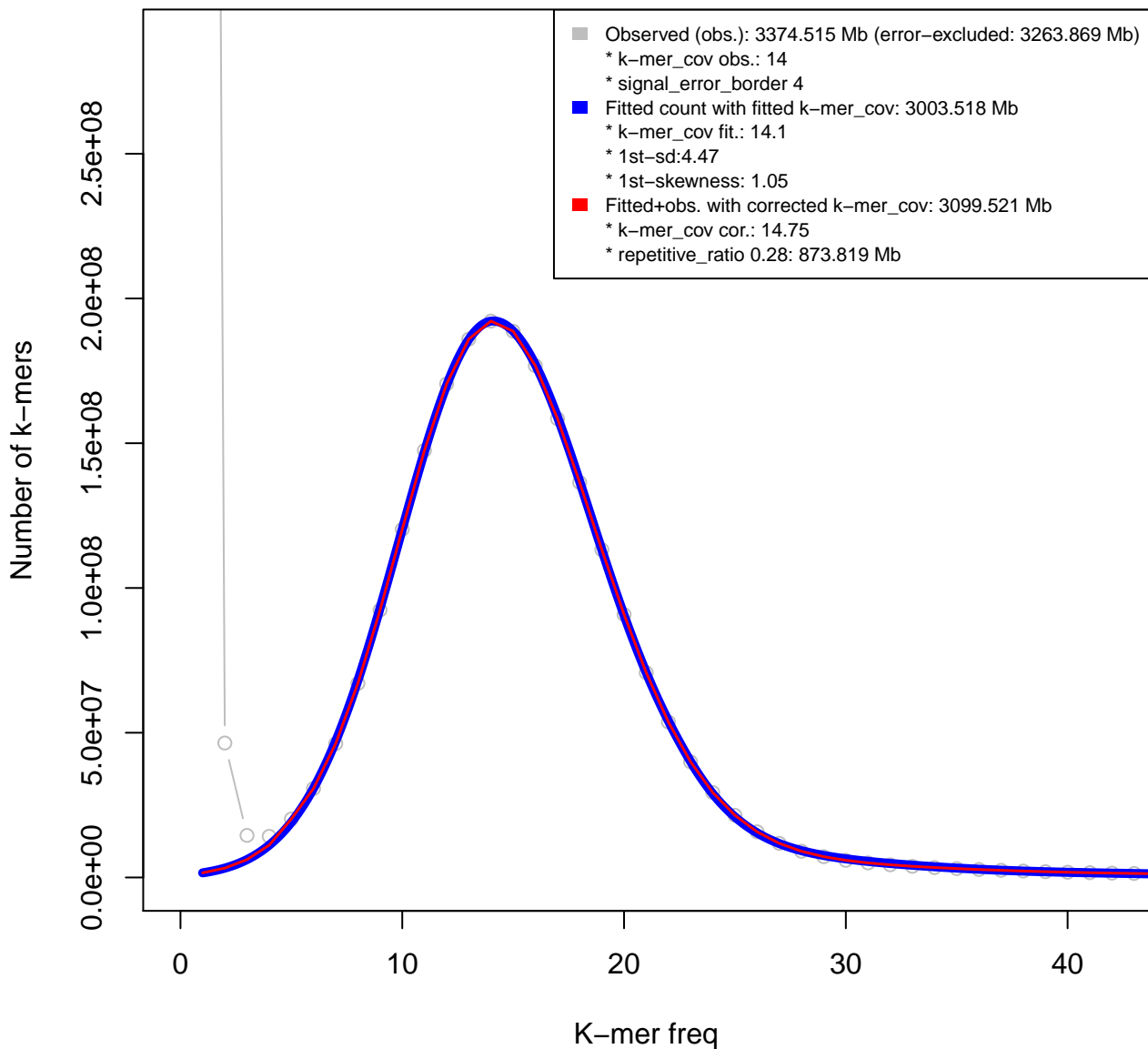


Sample human_30x_21m.histo k= 21



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

