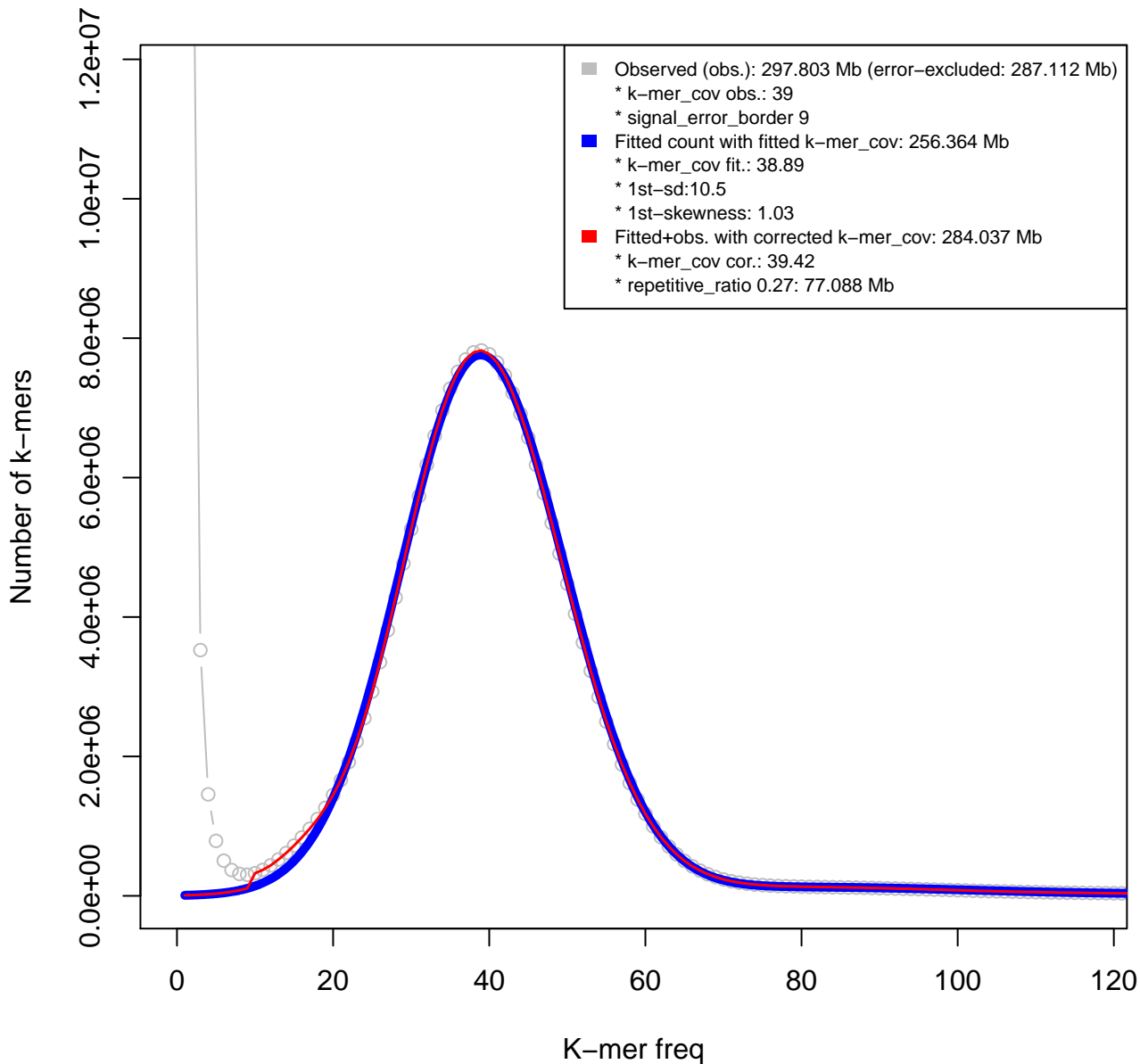


Sample HG001-NA12878-50x-combined.histo k= 21



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

