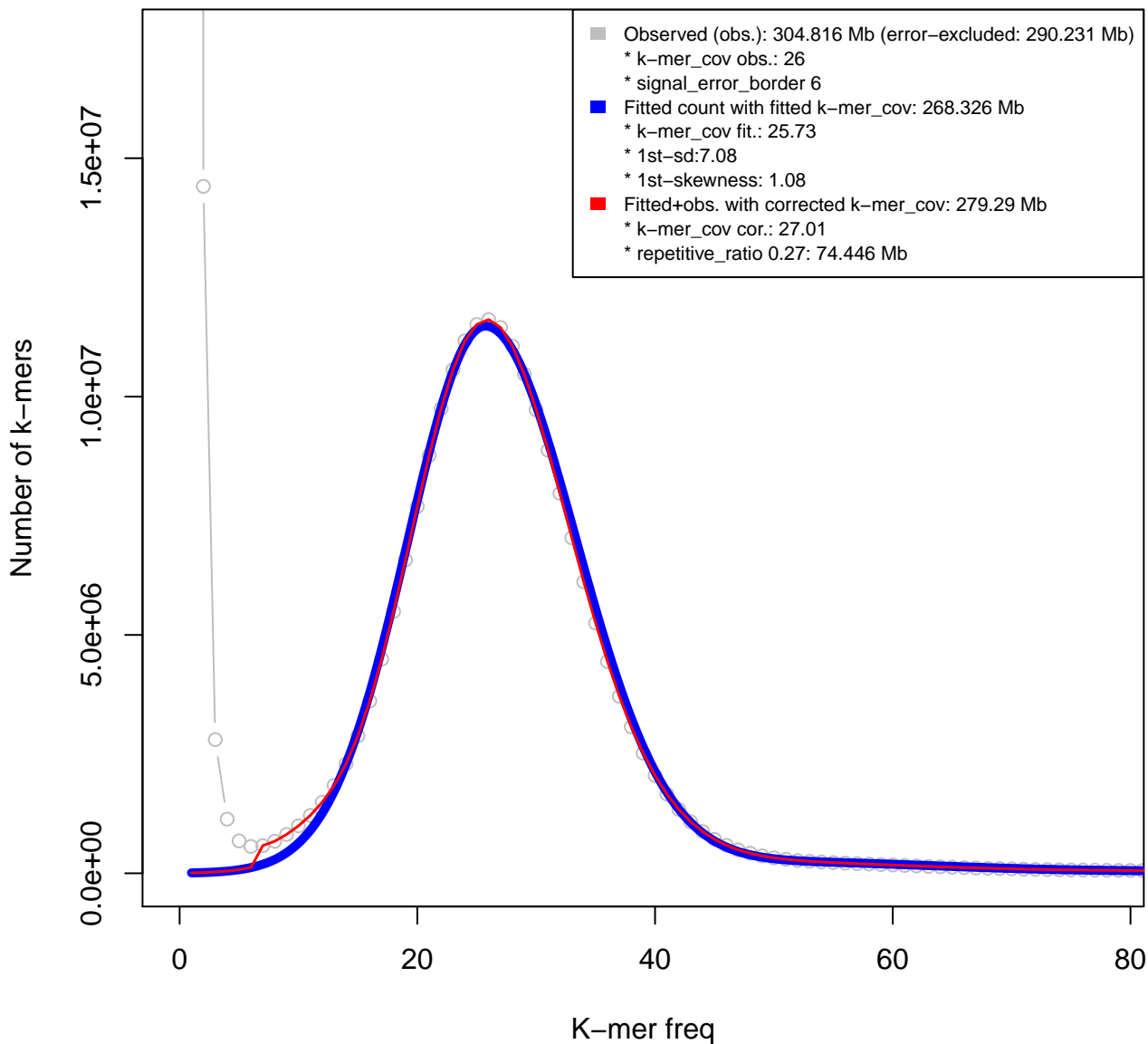


Sample HG001_novaseq_30x_combined.histo k= 21



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

