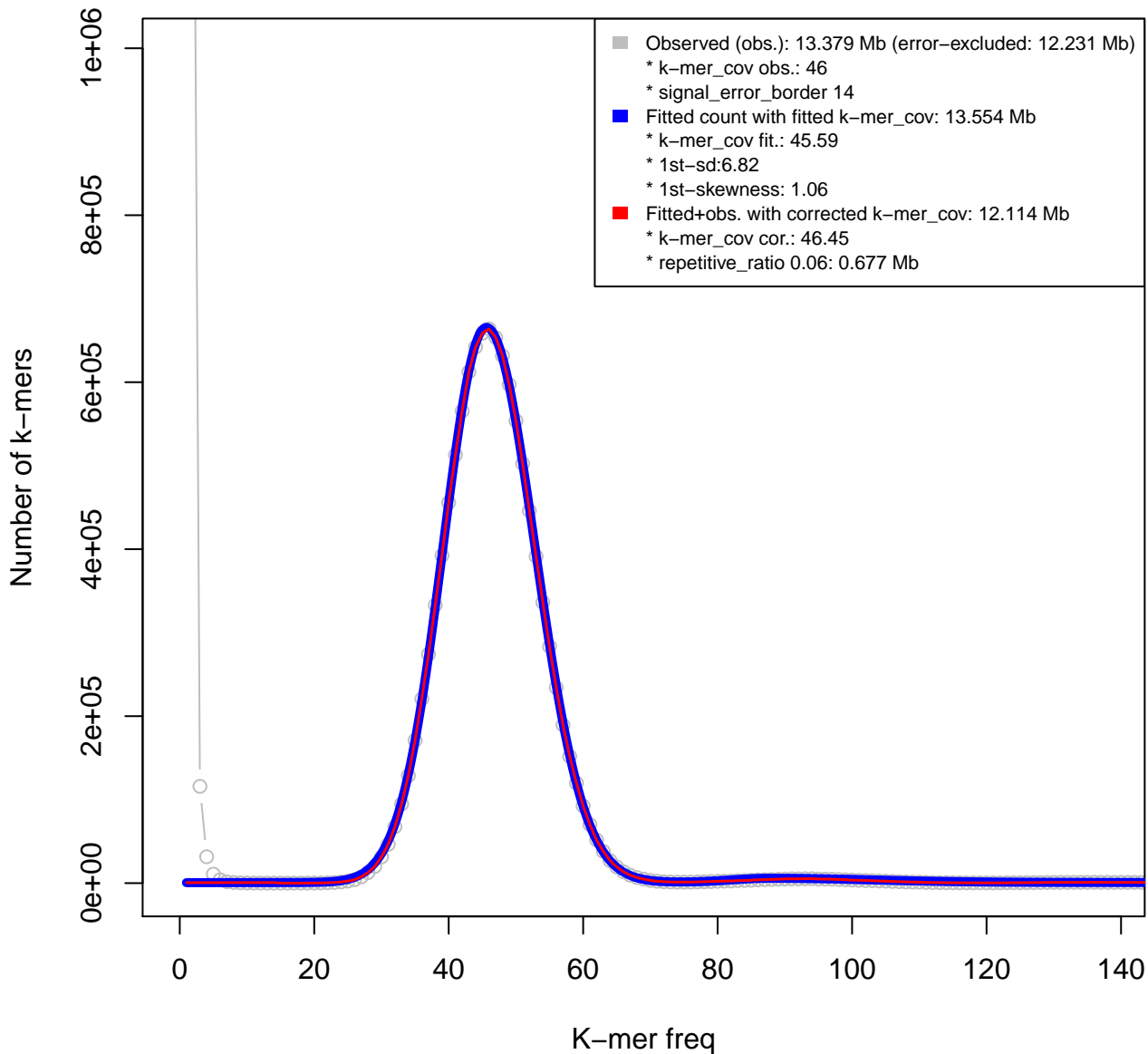


# Sample sim2.histo k= 27



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

