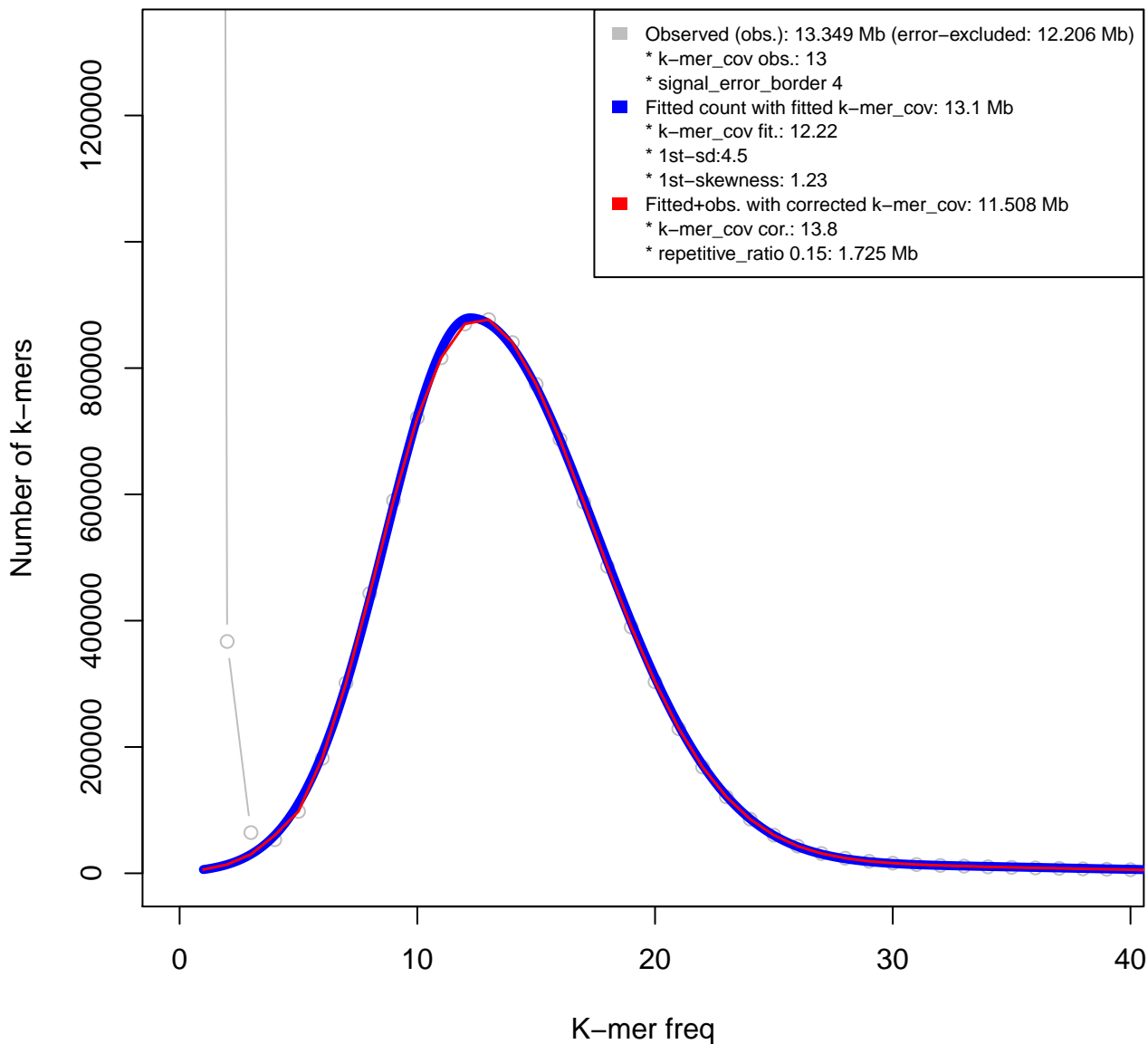


# Sample M\_27\_2.histo k= 27



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

