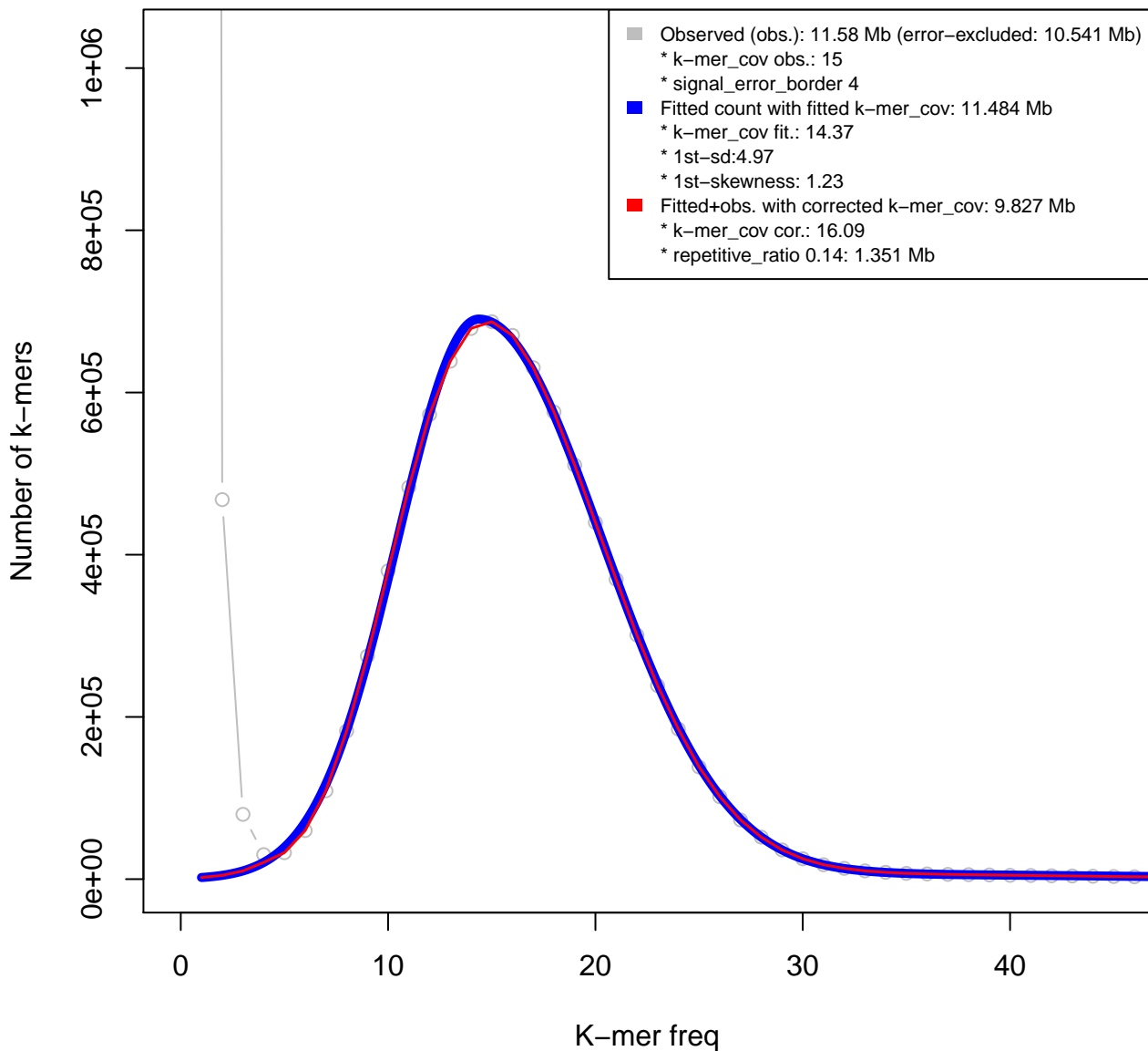


# Sample sim\_m27\_3.histo k= 27



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

