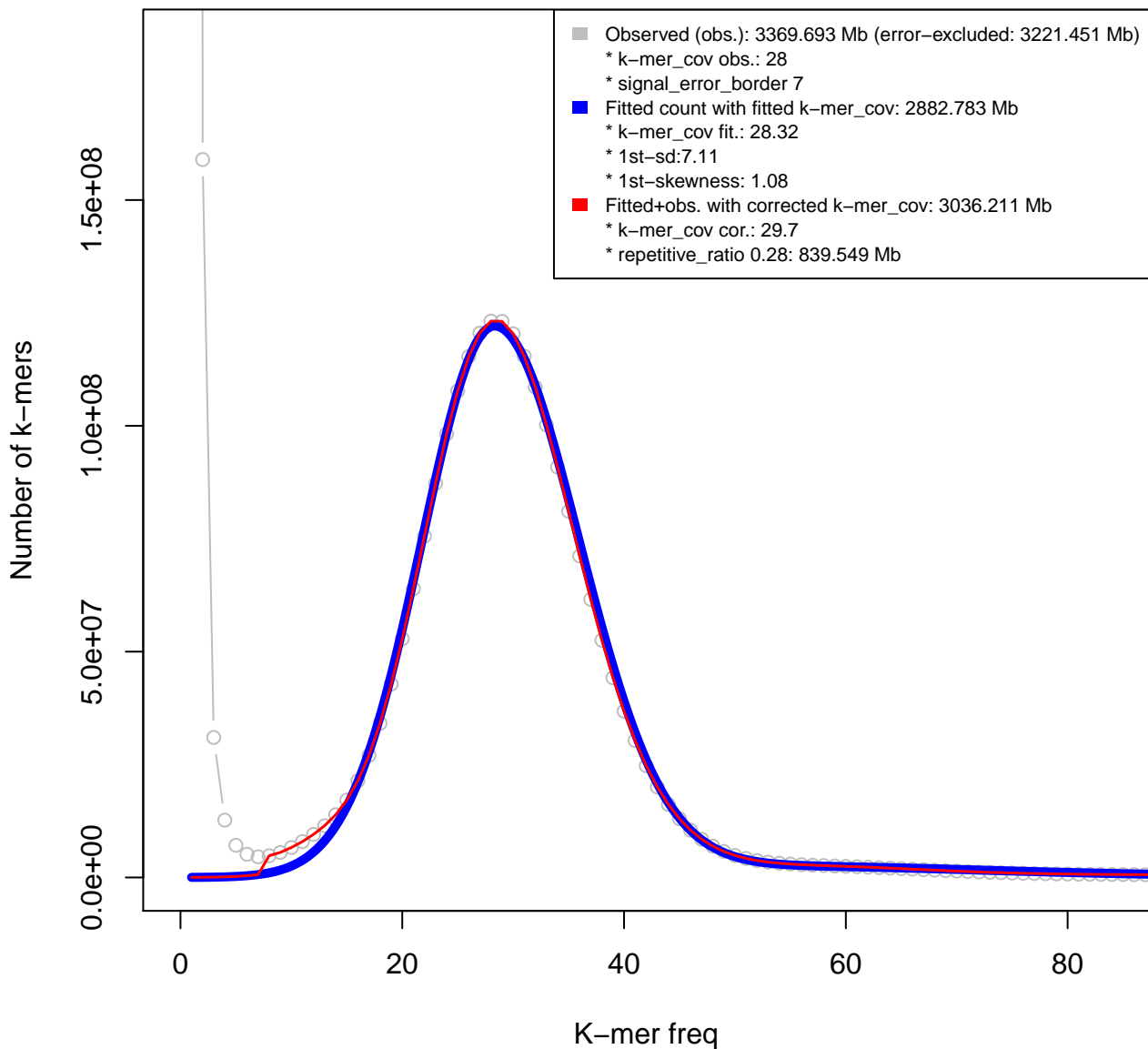


# Sample HG001\_novaseq\_30x\_original\_combined.histo k= 21



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

