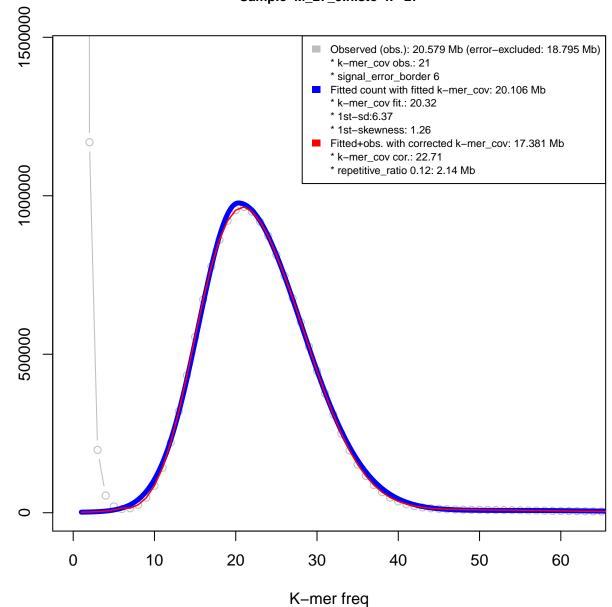
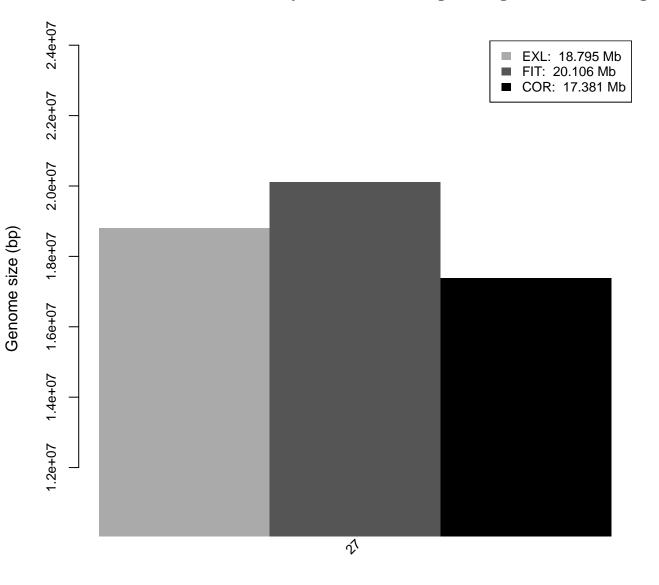
Sample M_27_3.histo k= 27



Number of k-mers

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



Size k