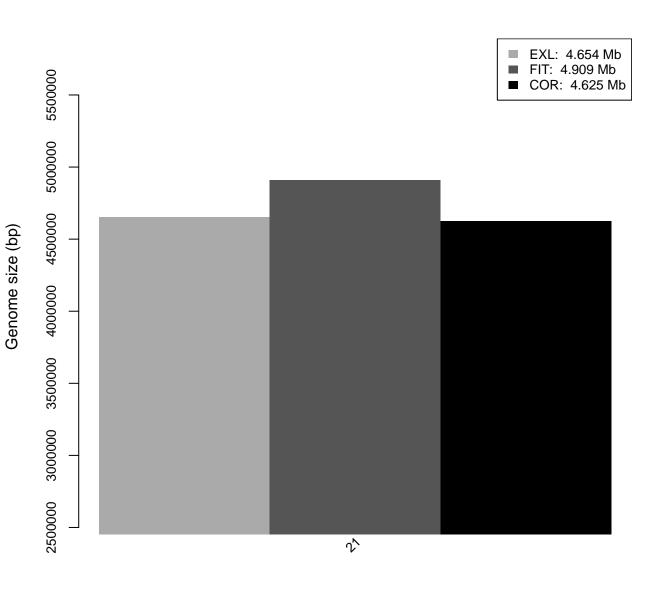


Genome size estimation by error-excluding, fitting, and correcting



Size k