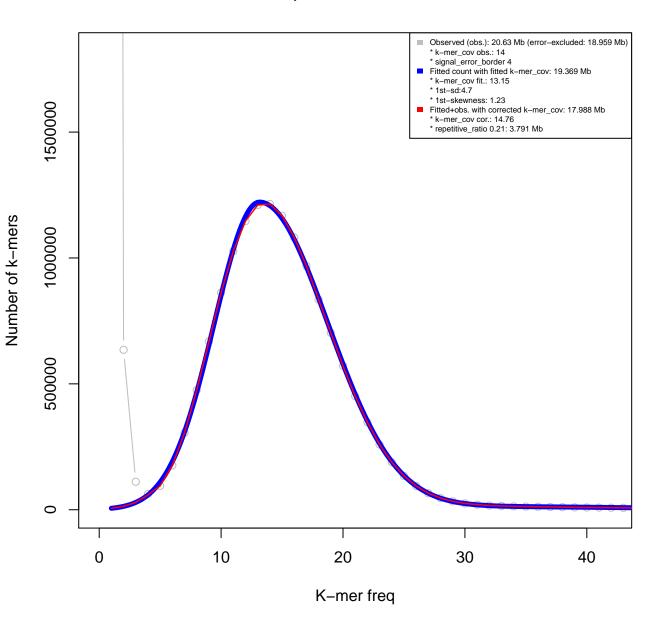
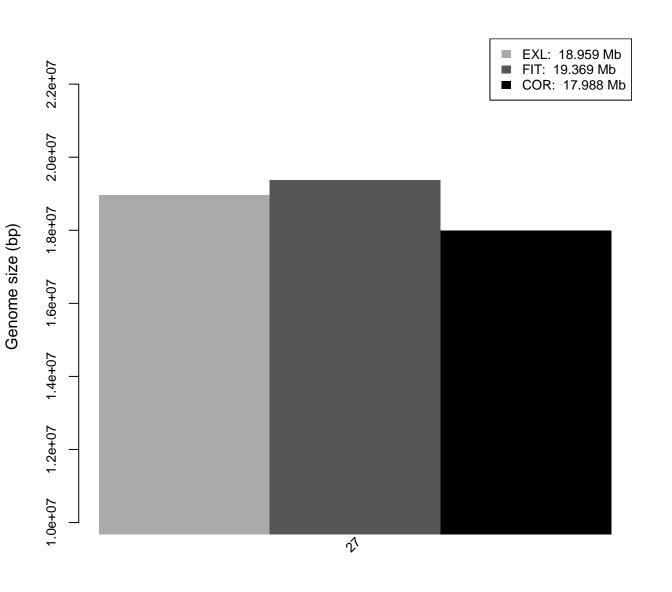
## Sample M\_27\_2.histo k= 27



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



Size k