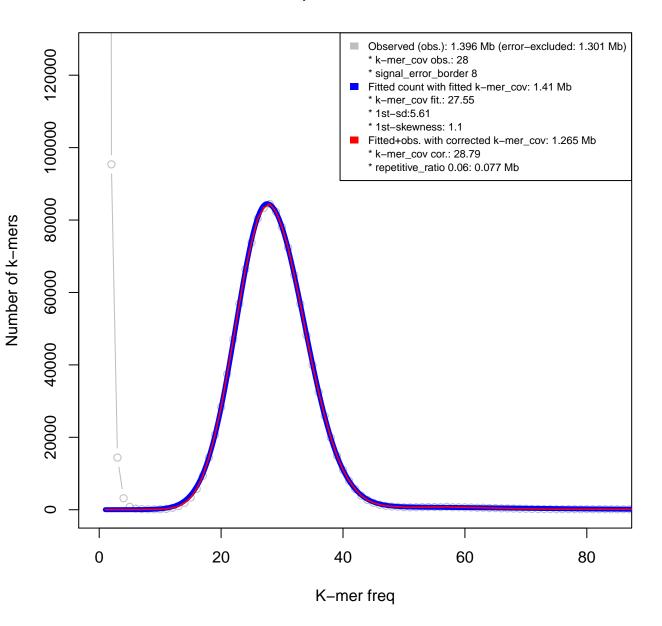
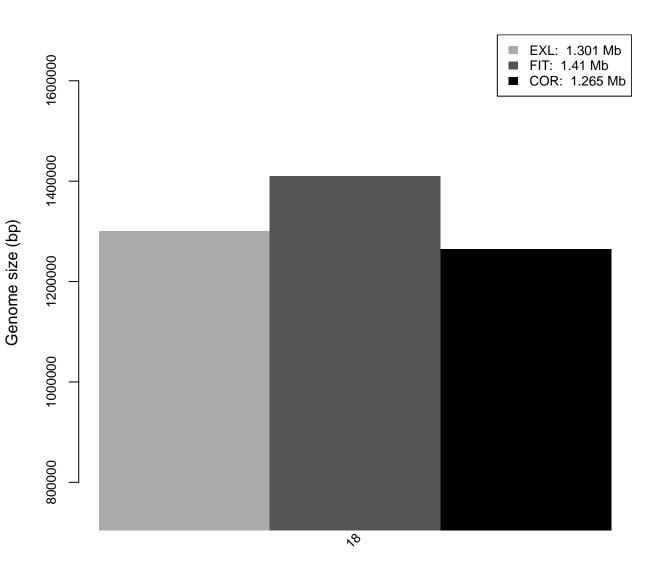
## Sample sim5.histo k= 18



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



Size k