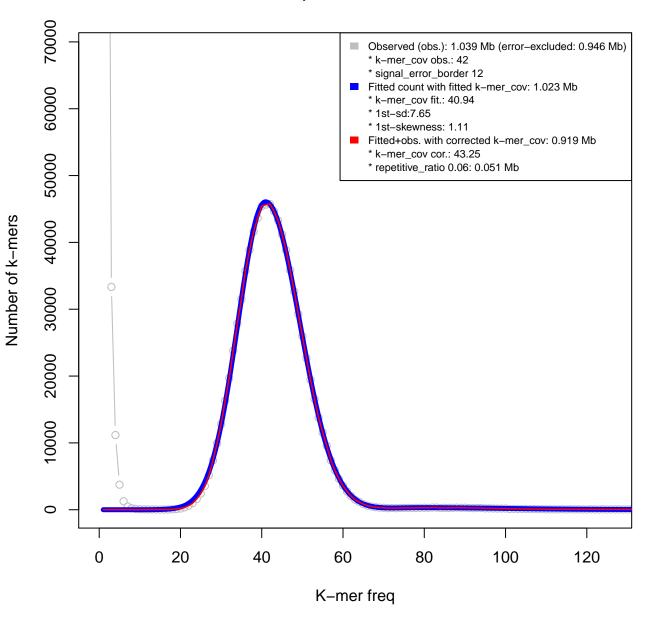
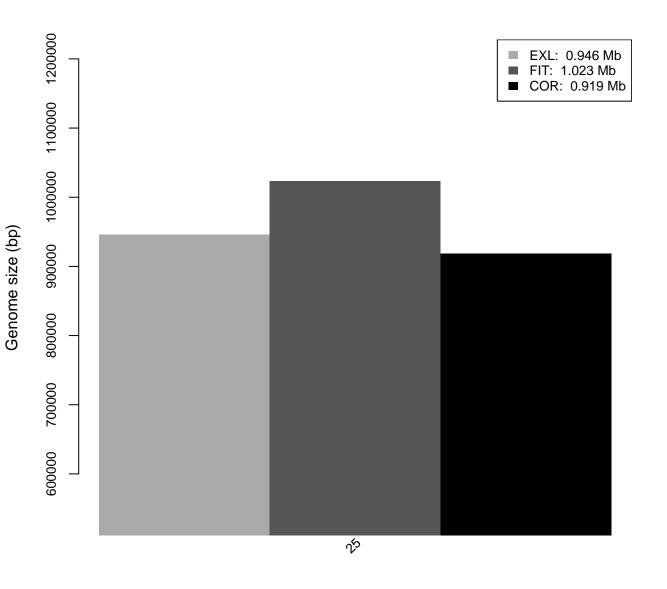
## Sample sim2.histo k= 25



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



Size k