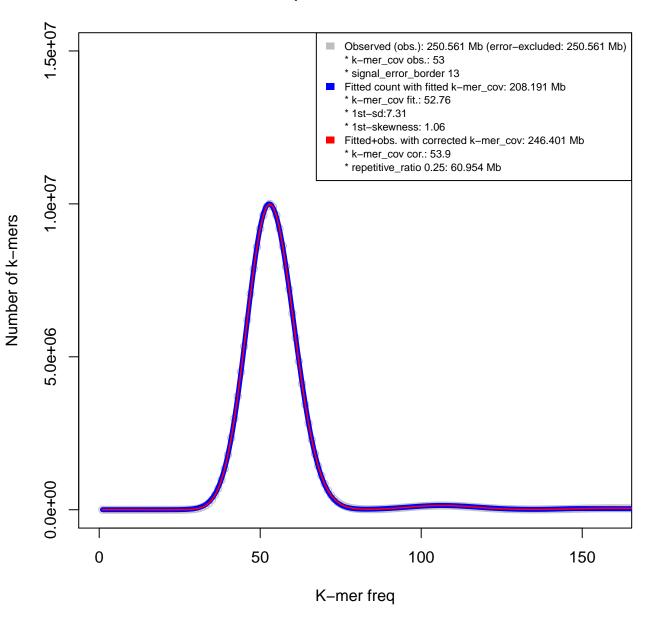
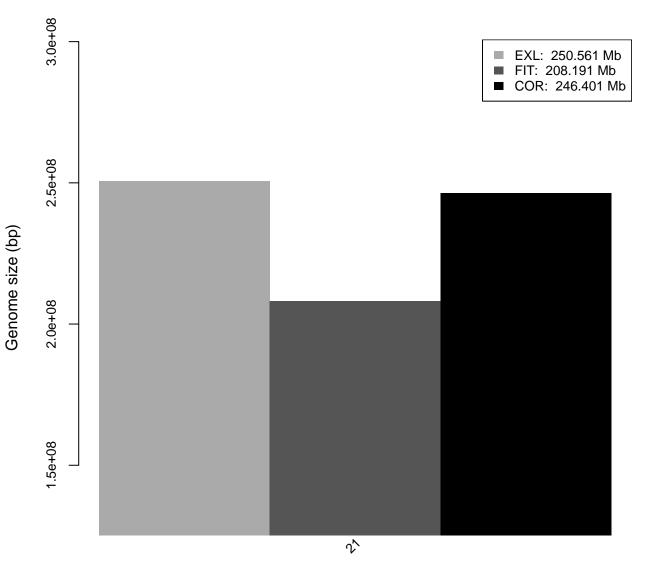
## Sample $K_21_5$ .histo k=21



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



Size k