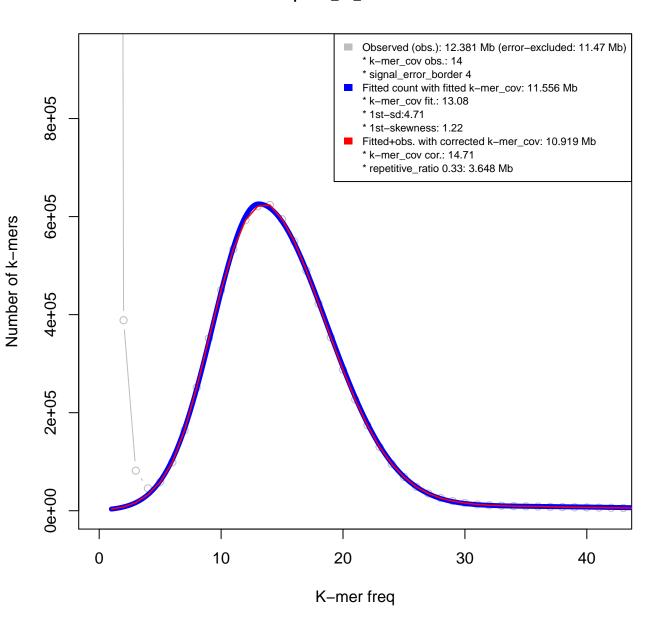
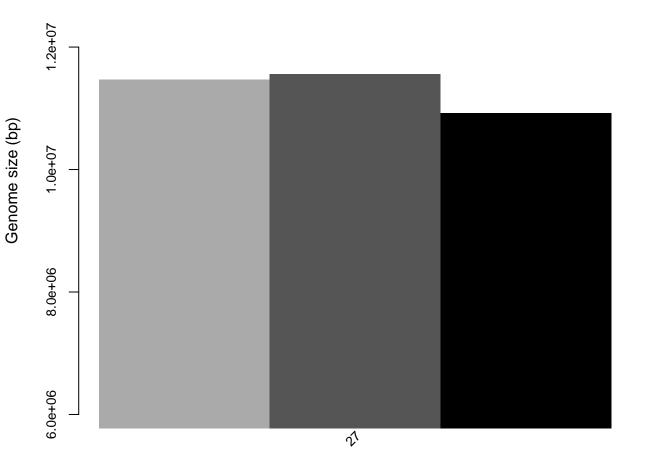
## **Sample M\_27\_2.histo k= 27**



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





Size k