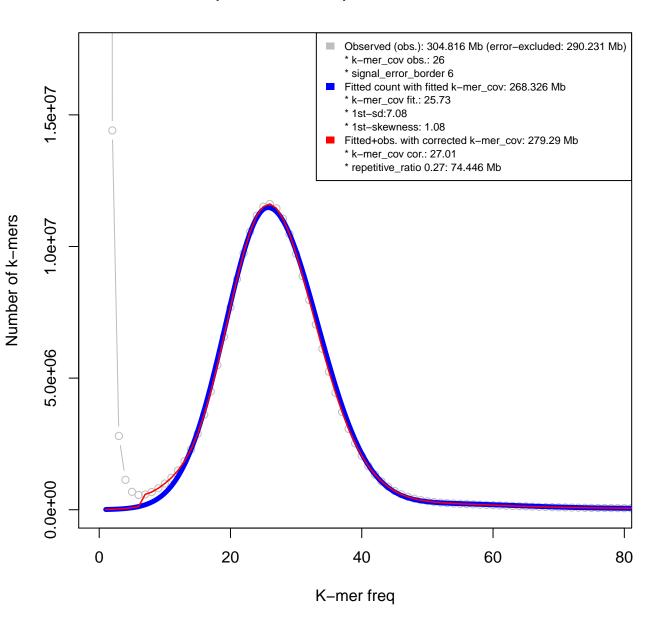
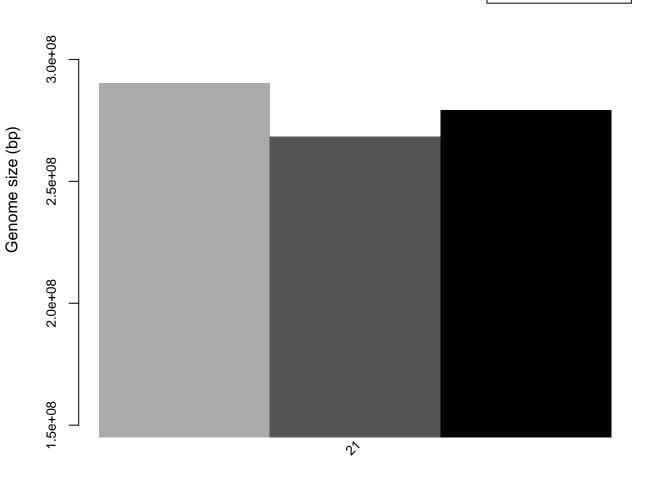
Sample HG001_novaseq_30x_combined.histo k= 21



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

EXL: 290.231 MbFIT: 268.326 MbCOR: 279.29 Mb



Size k