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Sequencing of the human genome has provided a great deal of information about each of the chromosomes. Below is a table compiling statistics for the chromosomes, based on the Sanger Institute's human genome information in the Vertebrate Genome Annotation (VEGA) database. Number of genes is an estimate, as it is in part based on gene predictions. Total chromosome length is an estimate as well ...

[12.1 Viruses – Concepts of Biology – 1st Canadian Edition](#)

The provisions of this Chapter 4 issued under the Public School Code of 1949 (24 P.S. § § 1-101—27-2702), unless otherwise noted. Source. The provisions of this Chapter 4 adopted January 15, 1999, effective January 16, 1999, 29 Pa.B. 399, unless otherwise noted. Cross References

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