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[Chapter 14 The Human Genome](#)

Human genome project (HGP) was an international scientific research project which got successfully completed in the year 2003 by sequencing the entire human genome of 3.3 billion base pairs. The HGP led to the growth of bioinformatics which is a vast field of research.

[The Human Genome Project \(Stanford Encyclopedia of Philosophy\)](#)

If a non-human gene were introduced into the human genome then the person would no

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longer be fully human but would by definition become a hybrid. The ancient name for a creature that is composed of different kinds is known as a chimera and that is also the name given to hybrid creatures by scientists today.

[Whole genome sequencing - Wikipedia](#)

Genome size refers to the amount of DNA contained in a haploid genome expressed either in terms of the number of base pairs, kilobases (1 kb = 1000 bp), or megabases (1 Mb = 1 000 000 bp), or as the mass of DNA in picograms (1 pg = 10⁻¹² g). Genome sizes of bacteriophages and viruses range from about 2 kb to over 1 Mb.

[CRISPR'd babies: human germline genome editing in the 'He ...](#)

For example, since the release of the draft sequence of the human reference genome in 2001 (Lander et al., 2001; Venter et al., 2001), thousands of individual human genomes have been sequenced, including such comparative genome-sequencing projects as: a deep catalog of human variation of thousands of individuals, 7 normal versus tumor cells ...

[Chromosome - Wikipedia](#)

*by Junying Yu * and James A. Thomson **. Human embryonic stem (ES) cells capture the imagination because they are immortal and have an almost unlimited developmental potential (Fig. 1.1: How hESCs are derived). After many months of growth in culture dishes, these remarkable cells maintain the ability to form cells ranging from muscle to nerve to blood—potentially any cell type that makes up ...*

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