



Genetics Review

[PubMed](#)
[Entrez](#)
[BLAST](#)
[OMIM](#)
[Taxonomy](#)
[Structure](#)
[NCBI Home](#)
[NCBI Site Map](#)
[brief/complete](#)

A few more terms...

[Course](#)
[Description](#)
[Schedule](#)
[Introduction](#)

- [acrocentric chromosome](#)
- [euchromatin](#)
- [heterochromatin](#)
- [draft sequence](#)
- [finished sequence](#)

[Genetics Review](#)

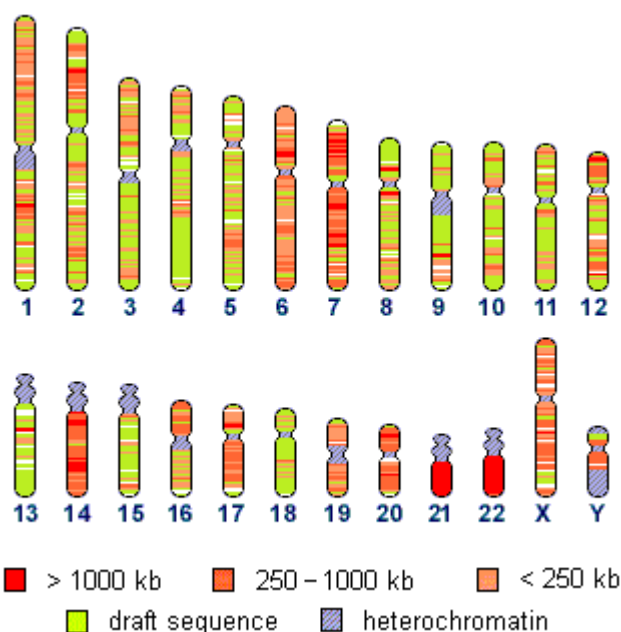
Acrocentric chromosome


[Types of Databases](#)

A chromosome in which the centromere is very close to one end.

[Format of Sequence Record](#)

For example, the centromere of chromosomes 13, 14, 15, 21, and 22 is very close to the p telomere, making the p arm very short. The p arm in those chromosomes contains [heterochromatin](#), colored in blue.

[Entrez](#)
[BLAST](#)
[3-D Structures](#)
[Genomes and Maps](#)
[Librarian Roles](#)
[WWW Sites](#)
[Glossaries and Dictionaries](#)


The Human Genome Project focuses primarily on sequencing the [euchromatic](#) regions of the chromosomes. The illustration above uses a color code to show the sequencing progress (as of June 25, 2000).

The [Human Genome Sequencing](#) page shows an illustration that is updated as new sequence data are generated, and includes a link to progress statistics.

Source:

definition based on:

Smith, A.D., et al., eds. 1997. *Oxford Dictionary of*

Biochemistry and Molecular Biology. New York: Oxford University Press.

illustration from:
NCBI Human Genome Sequencing web page, July 20, 2000
<http://www.ncbi.nlm.nih.gov/genome/seq/>

Euchromatin



"The fraction of the nuclear genome which contains transcriptionally active DNA and which, unlike [heterochromatin](#), adopts a relatively extended conformation."

Source: Strachan, T. and Read, A.P. 1999. Glossary in *Human Molecular Genetics, 2nd ed.* New York: John Wiley & Sons.

Heterochromatin



"A chromosomal region that remains highly condensed throughout the cell cycle and shows little or no evidence of active gene expression."

Source: Strachan, T. and Read, A.P. 1999. Glossary in *Human Molecular Genetics, 2nd ed.* New York: John Wiley & Sons.

Draft Sequence



A region of sequence which still contains gaps, and in which sections of DNA might still be of unknown order and orientation.

The sections of DNA are grouped together into a single GenBank submission because they have been sequenced from the same clone. Once the order and orientation of the sections is determined, and as the gaps are filled, the sequence will move into the finished phase.

The draft sequence in the [illustration](#) above is shown in green, while [finished sequence](#) is shown in shades of orange or red (depending on the length of the finished segment of sequence).

The phases of sequencing are described on the [High-Throughput Genomic Sequences \(HTG\)](#) page. Draft sequence can be either phase 1 or phase 2.

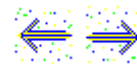
Finished Sequence



A region of sequence which has been completely sequenced. It contains no gaps, and the order and orientation of all the sequence subsections are known.

The finished sequence in the [illustration](#) above is shown in shades of orange or red (depending on the length of the finished segment of sequence), while the [draft sequence](#) is shown in green.

The phases of sequencing are described on the [High-Throughput Genomic Sequences \(HTG\)](#) page. Finished sequence is phase 3.



Help Desk

NCBI

NLM

NIH

Credits

Revised February 13, 2001

Comments/questions about course to Renata Geer

renata@ncbi.nlm.nih.gov

Questions about NCBI resources to info@ncbi.nlm.nih.gov