

DNA Sequencing

ENCODE: ENCyclopedia Of DNA Elements

Objective:
To identify all functional elements in the human genome sequence

ENCODE By the Numbers

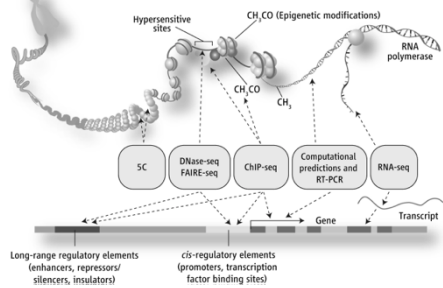
147 cell types studied
80% functional portion of human genome
20,687 protein-coding genes
18,400 RNA genes
1640 data sets
30 papers published this week
442 researchers
\$288 million funding for pilot, technology, model organism, and current project

E Pennisi Science 2012;337:1159-1161



Published by AAAS

Zooming in. A diagram of DNA in ever-greater detail shows how ENCODE's various tests (gray boxes) translate DNA's features into functional elements along a chromosome.



E Pennisi Science 2012;337:1159-1161



Published by AAAS

DNA Sequencing

Restriction enzymes (1973; Boyer & Cohen) cleave the polynucleotide to smaller fragments. These smaller fragments (100-200 base pairs) are sequenced. The two strands are separated.

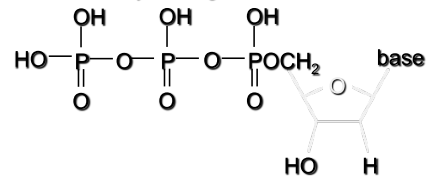
DNA Sequencing

Restriction enzymes cleave the polynucleotide to smaller fragments in specific patterns.

<div> <div>C</div> <div>G</div> <div>T</div> <div>A</div> <div>G</div> <div>C</div> </div>	<div> <div>T</div> <div>A</div> <div>C</div> <div>G</div> <div>A</div> <div>T</div> </div>	<div> <div>C</div> <div>G</div> <div>T</div> <div>A</div> <div>G</div> <div>C</div> </div>
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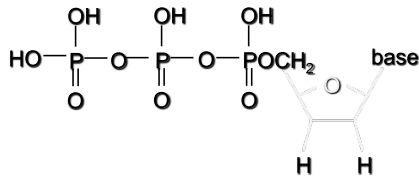
DNA Sequencing

Single stranded DNA divided in four portions. Each tube contains adenosine, thymidine, guanosine, and cytidine plus the triphosphates of their 2'-deoxy analogs.



DNA Sequencing

The first tube also contains the 2,3'-dideoxy analog of adenosine triphosphate (ddATP); the second tube the 2,3'-dideoxy analog of thymidine triphosphate (ddTTP), the third contains ddGTP, and the fourth ddCTP.



DNA Sequencing

Each tube also contains a "primer," a short section of the complementary DNA strand, labeled with radioactive phosphorus (^{32}P).

DNA synthesis takes place, producing a complementary strand of the DNA strand used as a template.

DNA synthesis stops when a dideoxynucleotide is incorporated into the growing chain.

DNA Sequencing

The contents of each tube are separated by electrophoresis and analyzed by autoradiography.

There are four lanes on the electrophoresis gel.

Each DNA fragment will be one nucleotide longer than the previous one.

DNA Profiling

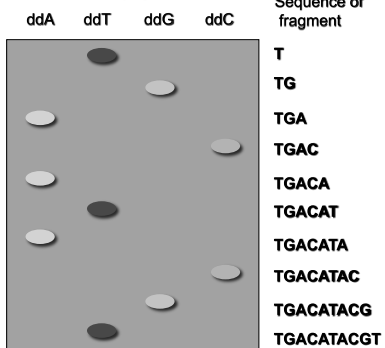
DNA sequencing involves determining the nucleotide sequence in DNA.

The nucleotide sequence in regions of DNA that code for proteins varies little from one individual to another, because the proteins are the same.

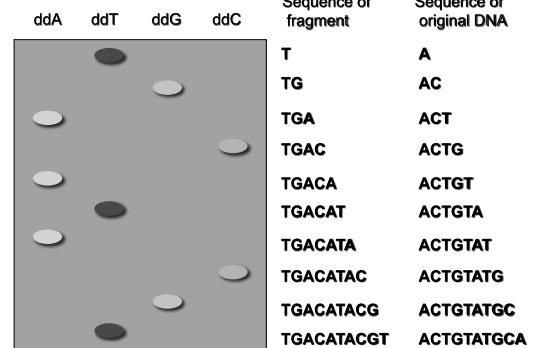
Most of the nucleotides in DNA are in "noncoding" regions and vary significantly among individuals.

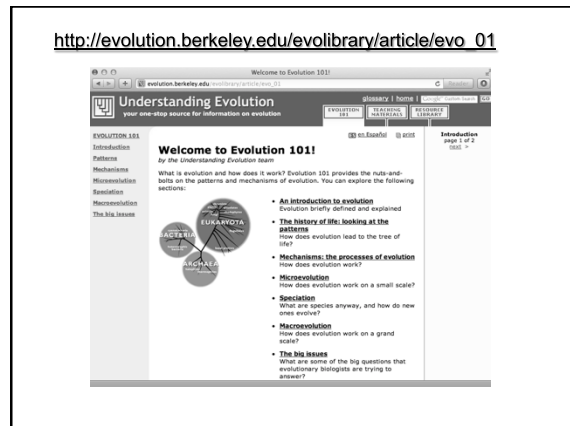
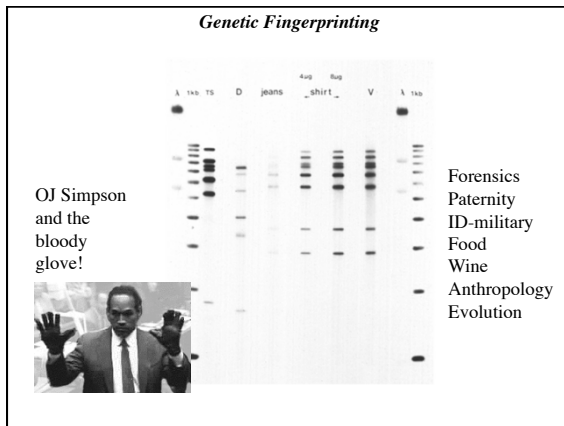
Enzymatic cleavage of DNA give a mixture of polynucleotides that can be separated by electrophoresis to give a "profile" characteristic of a single individual.

The bloody glove?



A bloody glove?





PCR: Polymerase Chain Reaction

PCR

When a sample of DNA is too small to be sequenced or profiled, the *polymerase chain reaction* (PCR) is used to make copies ("amplify") portions of it.

PCR amplifies DNA by repetitive cycles of the following steps.

1. Denaturation
2. Annealing ("priming")
3. Synthesis ("extension" or "elongation")

PCR

(a) Consider double-stranded DNA containing a polynucleotide sequence (the target region) that you wish to amplify.

Target region

(b) Heating the DNA to about 95°C causes the strands to separate. This is the denaturation step.

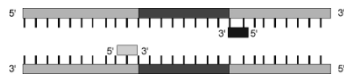
PCR

(c) Cooling the sample to ~60°C causes one primer oligonucleotide to bind to one strand and the other primer to the other strand. This is the annealing step.

(b) Heating the DNA to about 95°C causes the strands to separate. This is the denaturation step.

PCR

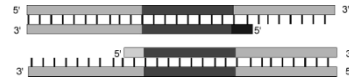
(c) Cooling the sample to $\sim 60^{\circ}\text{C}$ causes one primer oligonucleotide to bind to one strand and the other primer to the other strand. This is the annealing step.



(d) In the presence of four DNA nucleotides and the enzyme DNA polymerase, the primer is extended in its 3' direction. This is the synthesis step and is carried out at 72°C .

PCR

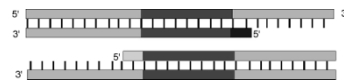
This completes one cycle of PCR.



(d) In the presence of four DNA nucleotides and the enzyme DNA polymerase, the primer is extended in its 3' direction. This is the synthesis step and is carried out at 72°C .

PCR

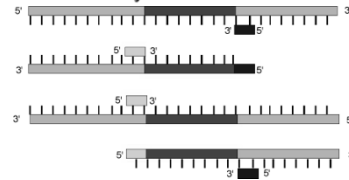
This completes one cycle of PCR.



(e) The next cycle begins with the denaturation of the two DNA molecules shown. Both are then primed as before.

PCR

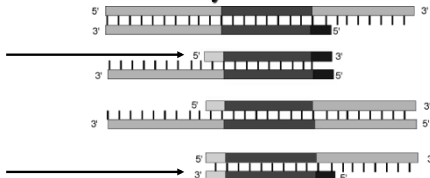
(f) Elongation of the primed fragments completes the second PCR cycle.



(e) The next cycle begins with the denaturation of the two DNA molecules shown. Both are then primed as before.

PCR

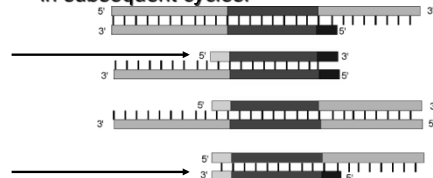
(f) Elongation of the primed fragments completes the second PCR cycle.



(g) Among the 8 DNAs formed in the second cycle are two having the structure shown.

PCR

The two contain only the target region and are the ones that increase disproportionately in subsequent cycles.



(g) Among the 8 DNAs formed in the second cycle are two having the structure shown.

PCR

Cycle	Total DNAs	Contain only target
0 (start)	1	0
1	2	0
2	4	0
3	8	2
4	16	8
5	32	22
10	1,024	1,004
20	1,048,566	1,048,526
30	1,073,741,824	1,073,741,764

Recombinant Methods

Recombinant DNA : GMOs

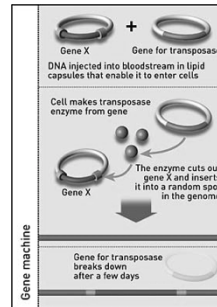
Restriction enzymes (1973), plasmids, promoters, recombinant DNA (rDNA) -> New Organisms -> Genentech et. al. (1976)

ABSTRACT

The Cohen-Boyer licensing program, by any variety of metrics, was widely successful. Recombinant DNA (rDNA) products provided a new technology platform for a range of industries, resulting in over US\$55 billion in sales for an estimated 2,442 new products. Over the duration of the life of the patents (they expired in December 1997), the technology was licensed to 468 companies, many of them fledgling biotech companies who used the licenses to establish their legitimacy. Over the 25 years of the licensing program, Stanford and the University of California system accrued US\$255 million in licensing revenues (to the end of 2001), much of which was subsequently invested in research and research infrastructure. In many ways, Stanford's management of the Cohen-Boyer patents has become the gold standard for university technology licensing. Stanford made pragmatic decisions and was flexible, adapting its licensing strategies as circumstances changed.

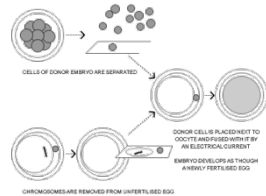
<http://www.nytimes.com/1999/12/07/business/robert-a-swanson-52-co-founder-of-genentech.html>

Transgenic Crops Monsanto Syngenta Luis?



CLONING

Hello Dolly, and Lassie, and Tabby



DNA Mutations

Understanding Evolution : Personal response question

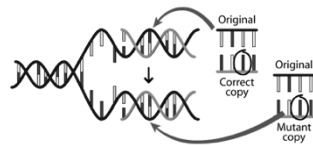
Mutation is a random process.

- a. agree
- b. disagree

DNA Mutations

Substitution	CTGGAG CTGGGG
Insertion	CTGGAG CTGGTGGAG
Deletion	CTGGAG CTAG
Frameshift	Xhe fat cat sat hef atc ats at

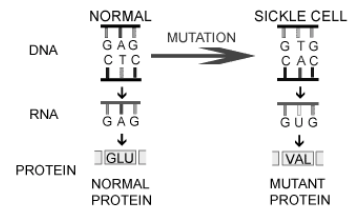
DNA Mutations



Enzymatic corrections are common
But, they are not always done

DNA Mutations

Sickle Cell Anemia



RNA

ENCODE: ENCyclopedia Of DNA Elements

Objective:
To identify all functional elements in the human genome sequence

ENCODE By the Numbers

- 147 cell types studied
- 80% functional portion of human genome
- 20,687 protein-coding genes
- 18,400 RNA genes
- 1640 data sets
- 30 papers published this week
- 442 researchers
- \$288 million funding for pilot, technology, model organism, and current project

COMPARING DIFFERENT KINDS OF SMALL RNAs			
	siRNA	microRNA	piRNA
Length	21–24 nucleotides	20–25 nucleotides	21–31 nucleotides
Organization	Double-stranded	Single-stranded	Single-stranded
Requires Dicer for maturation?	Yes	Yes	No
Found in	Animals, plants, fungi, protists	Animals, plants, protists	Only animals
Function	Controlling gene expression, blocking transposons	Controlling gene expression	Blocking transposons

M Leslie Science 2013;339:25-27

Published by AAAS

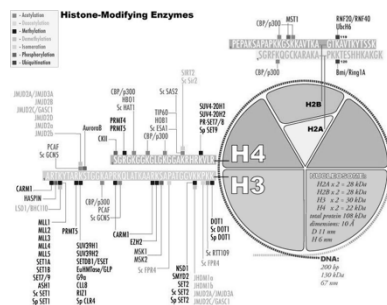
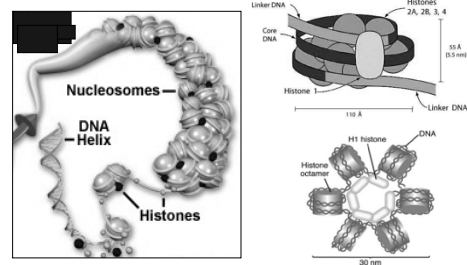
Science
AAAS

Epigenetics

Epigenetics

- Chemical reactions switch parts of the genome off and on at strategic times and locations.
- Epigenetics is the study of these reactions and the factors that influence them.
- View video:
<http://learn.genetics.utah.edu/content/epigenetics/intro/>
- <http://learn.genetics.utah.edu/content/epigenetics/control/>

Nucleosomes & Histones



Neoplasia is characterized by "methylation imbalance"