Chapter 20

Biotechnology

PowerPoint® Lecture Presentations for

Biology

Eighth Edition Neil Campbell and Jane Reece

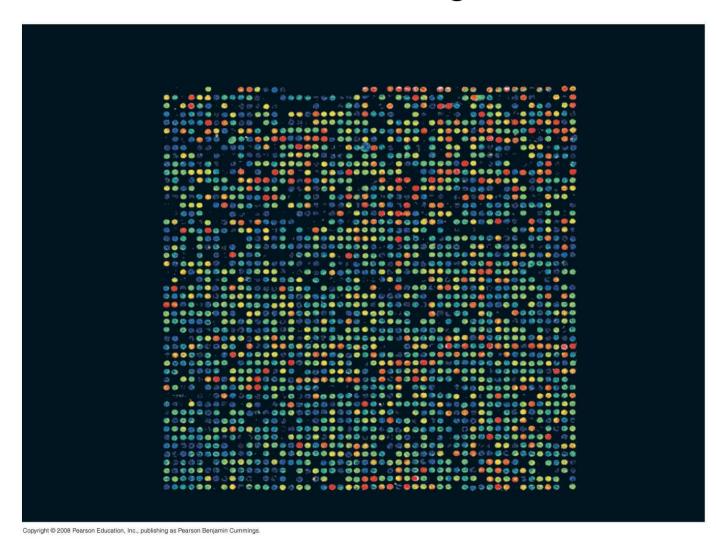
Lectures by Chris Romero, updated by Erin Barley with contributions from Joan Sharp

Overview: The DNA Toolbox

- One of the greatest achievements of modern science has been the sequencing of the human genome (2003)--- depended on advances in recombinant DNA technology
- In recombinant DNA, nucleotide sequences from two different sources, often two species, are combined in vitro into the same DNA molecule

- Methods for making recombinant DNA are central to genetic engineering, the direct manipulation of genes for practical purposes
- DNA technology has revolutionized biotechnology
 - the manipulation of <u>organisms or their</u> genetic components to make useful products

An example of DNA technology is the **microarray**, a measurement of gene expression of tens of thousands of different genes



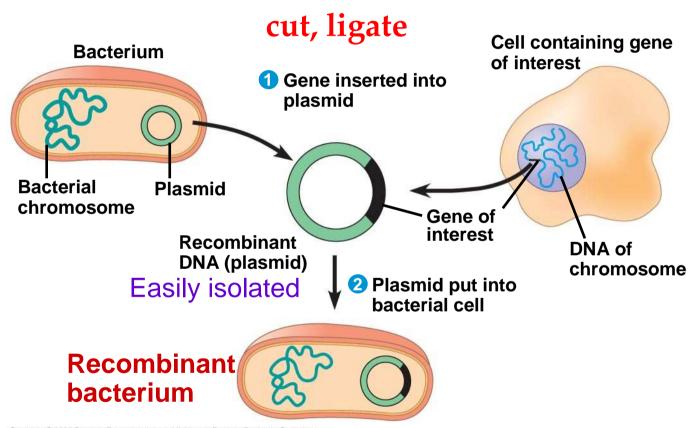
Concept 20.1: DNA cloning yields multiple copies of a gene or other DNA segment

- The first step in recombinant DNA technology
 - DNA cloning
- It is the production of multiple identical copies of a <u>specific gene</u> (gene cloning) or other <u>DNA segment</u>

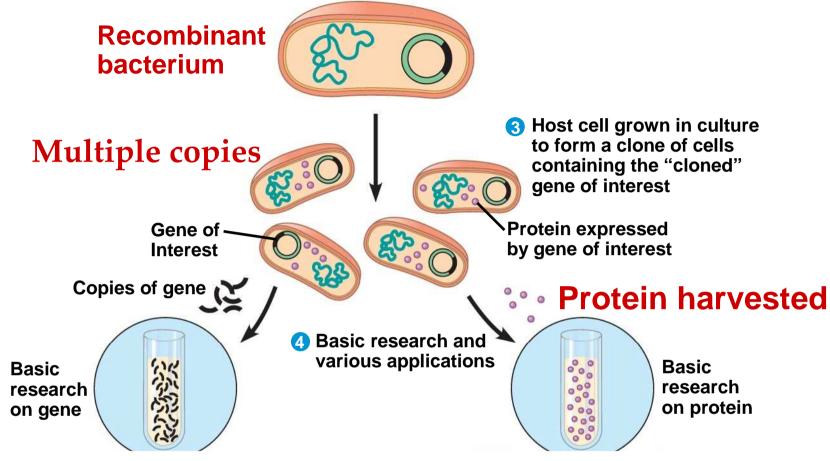
DNA Cloning and Its Applications: A Preview

- How to do DNA cloning:
 - (1) 目標:DNA of interest
 - (2) 載體: Plasmids (cloning vector):
 replication origin, multiple cloning sites,
 selection marker (ampicillin resistance gene;
 LacZ gene: b-galactosidae digests X-gal into
 blue product)
 - (3) 宿主:Host: bacteria (E. Coli), yeast
- Cloned genes are useful for making copies of a particular gene and producing a protein product

A preview of gene cloning and some uses of cloned genes

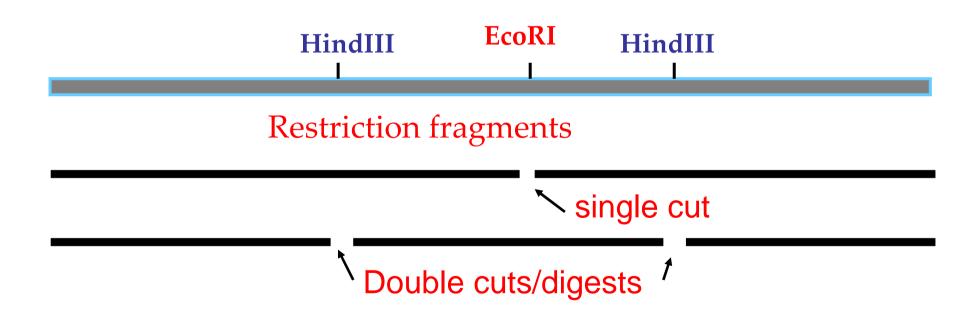


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How to Make Recombinant DNA: Restriction Enzymes

(1) Bacterial **restriction enzymes** -- cut DNA molecules at a limited number of **specific** DNA sequences (4-8 nucleotides), called **restriction sites** – GAATTC (EcoRI),



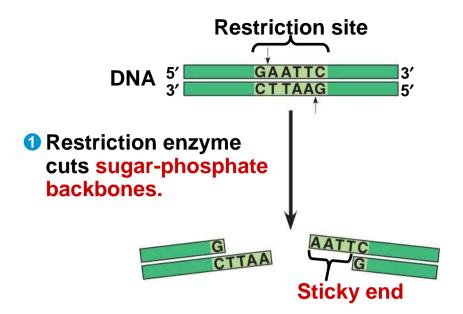
Using Restriction Enzymes to Make Recombinant DNA

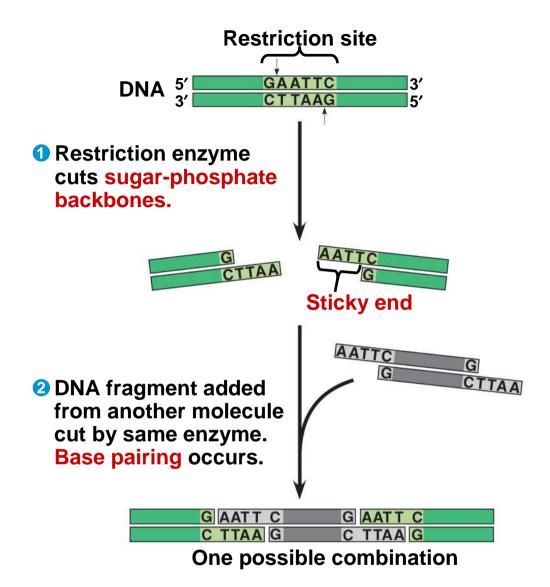
 The most useful restriction enzymes cut DNA in a staggered way, producing fragments with "sticky ends" that bond with complementary sticky ends of other fragments

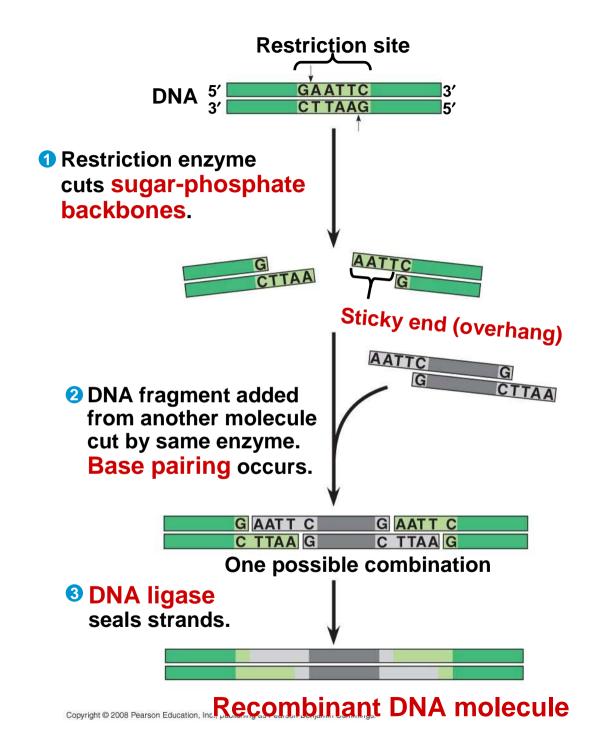
PLAY

Animation: Restriction Enzymes

 DNA ligase is an enzyme that seals the bonds between restriction fragments







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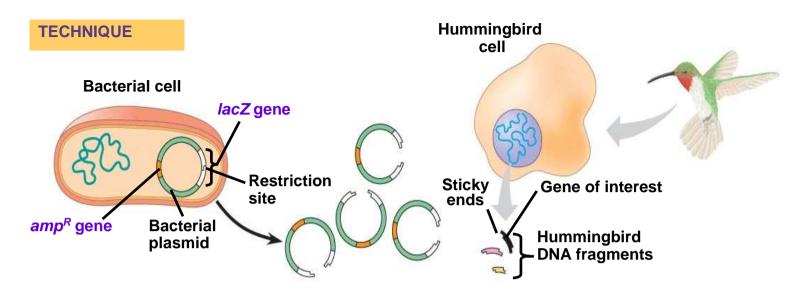
Cloning a Eukaryotic Gene in a Bacterial Plasmid

- In gene cloning, the original plasmid is called a cloning vector
- A cloning vector is a DNA molecule that can carry foreign DNA into a host cell and replicate there

Producing Clones of Cells Carrying Recombinant Plasmids

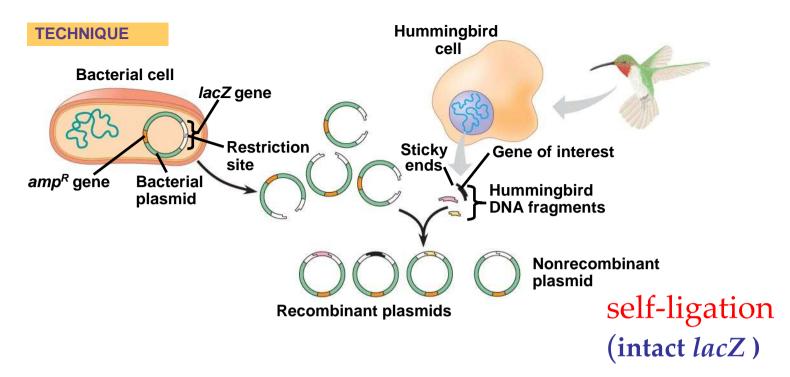
- Several steps are required to clone the hummingbird β-globin gene in a bacterial plasmid:
 - The hummingbird genomic DNA and a bacterial plasmid are isolated
 - Both are digested with the same restriction enzyme
 - The fragments are mixed, and DNA ligase is added to bond the fragment sticky ends

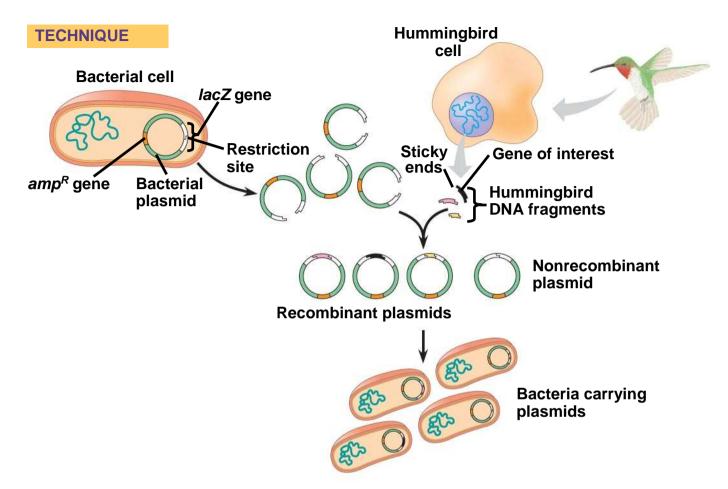
- Some recombinant plasmids now contain hummingbird DNA
- The DNA mixture is added to bacteria that have been genetically engineered to accept it
- The bacteria are plated on a specific type of agar that selects for the bacteria with recombinant plasmids
- This results in the cloning of many hummingbird DNA fragments, including the β-globin gene

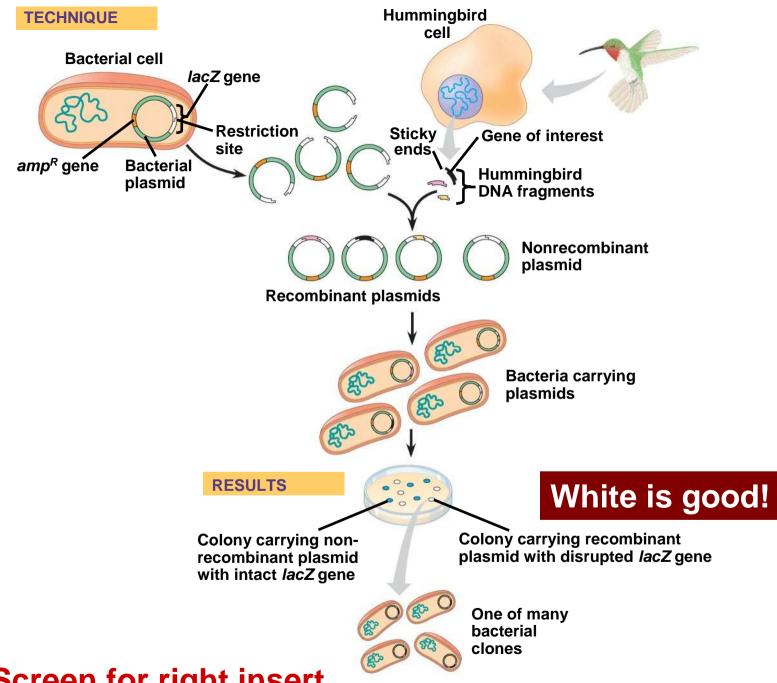


amp^R gene: makes *E. coli* cells resistant to the antibiotic ampicillin.

lacZ gene: encodes β-galactosidase. This enzyme hydrolyzes a molecular mimic of lactose (X-gal) to form a blue product. (藍白箭)







Screen for right insert

Summay of cloning a gene

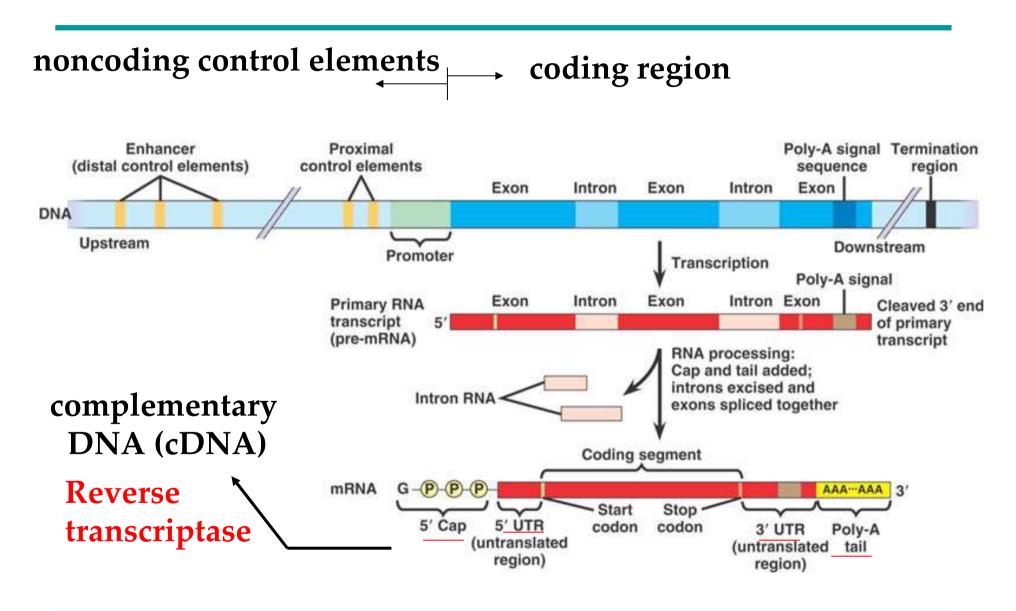
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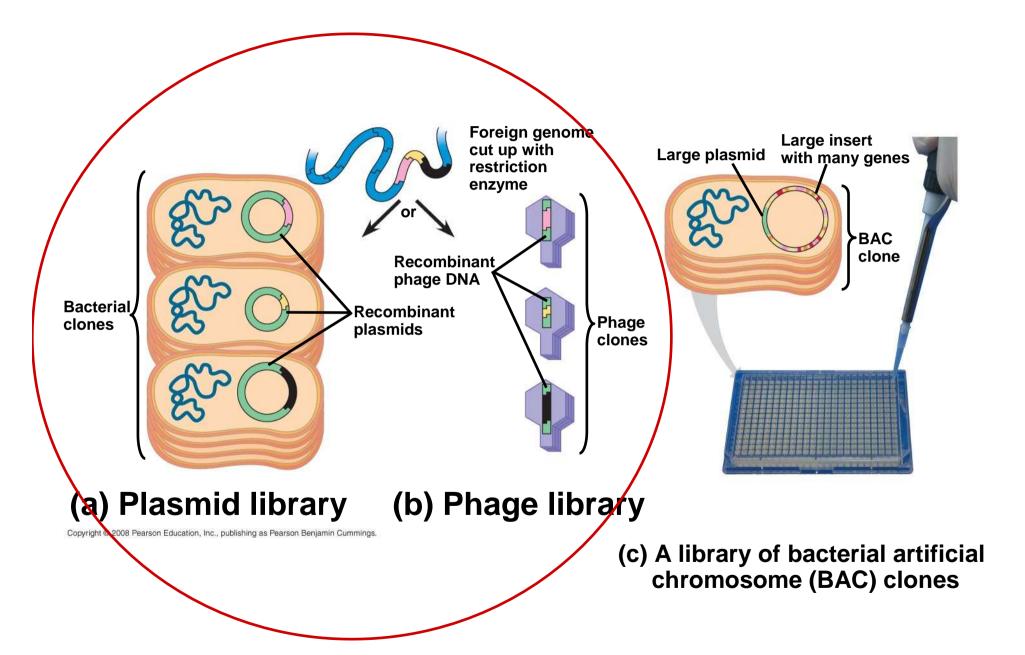
Animation: Cloning a Gene

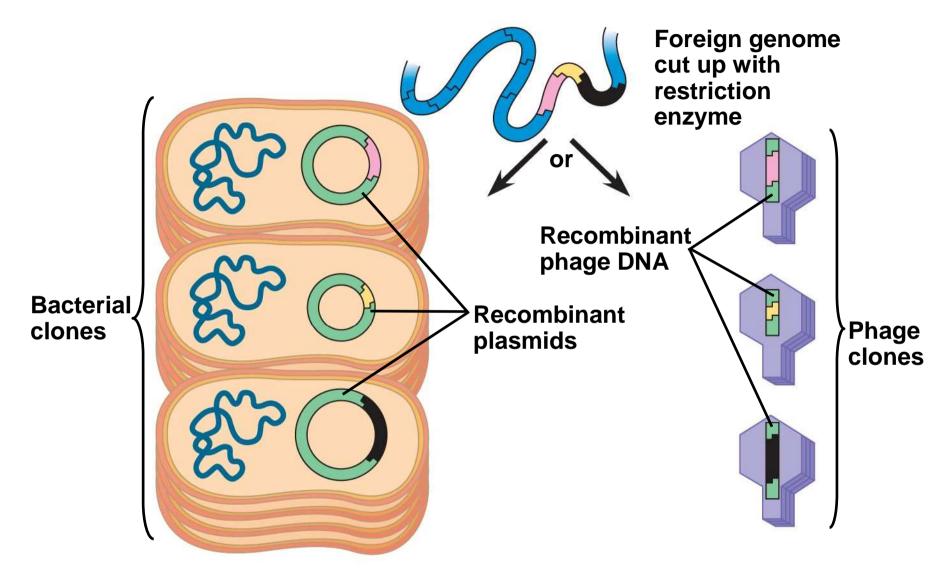
Storing Cloned Genes in DNA Libraries

- A genomic library that is made using bacteria is the collection of recombinant vector clones produced by cloning DNA fragments from an entire genome - complete sequence with regulatory region (enhancers), exon, introns, nonconding region, etc.
- A genomic library that is made using bacteriophages is stored as a collection of phage clones

Organization of eukaryotic genes: with multiple control elements





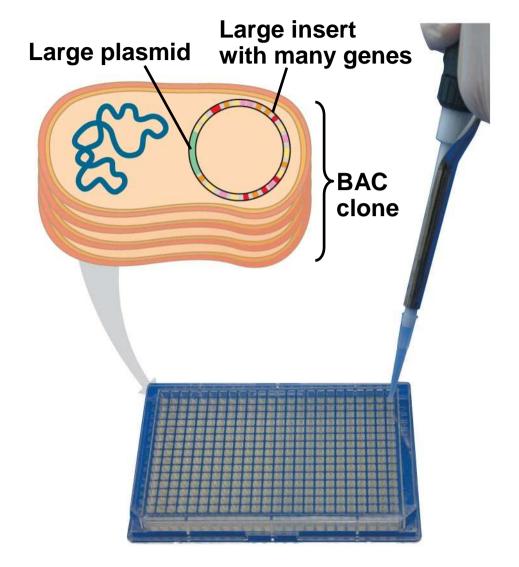


(a) Plasmid library

(b) Phage library

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- A bacterial artificial chromosome (BAC) is a large plasmid that has been trimmed down and can carry a large DNA insert
- BACs are another type of vector used in DNA library construction

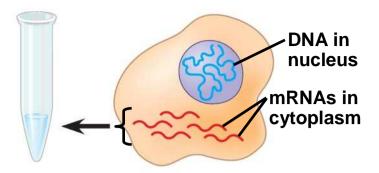


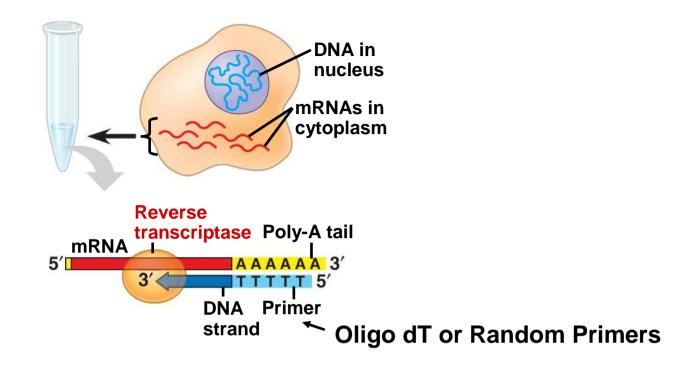
(c) A library of bacterial artificial chromosome (BAC) clones

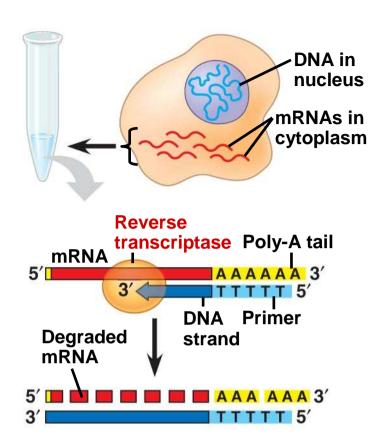
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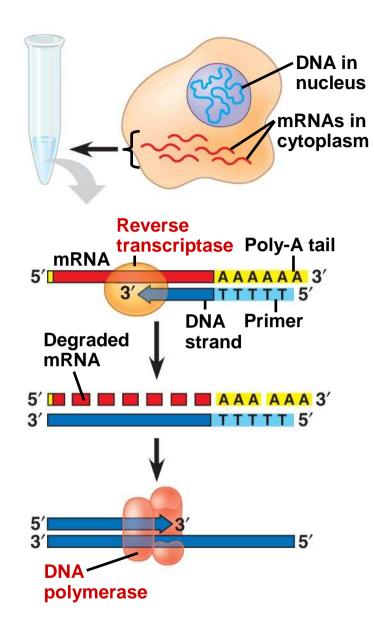
cDNA library

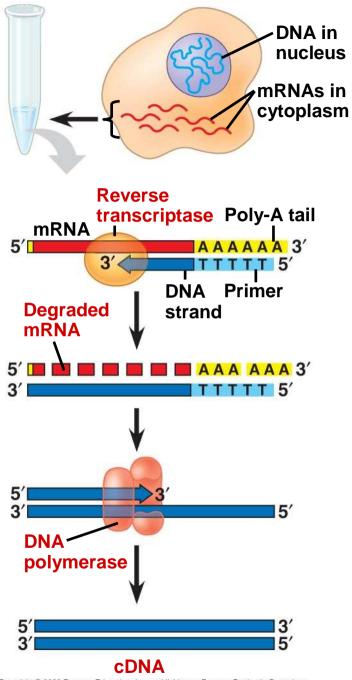
- A complementary DNA (cDNA) library is made by cloning DNA made in vitro by reverse transcription of all the mRNA produced by a particular cell or tissue
- A cDNA library represents only part of the genome — only the <u>subset of genes</u> transcribed into mRNA in the original cells











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Screening a Library for Clones Carrying a Gene of Interest

- A clone carrying the gene of interest can be identified with a nucleic acid probe having a sequence complementary to the gene
- This process is called nucleic acid hybridization

- A probe can be synthesized that is complementary to the gene of interest
- For example, if the desired gene is

Then we would synthesize this probe



DNA denature and probe hybridization (colony hybridization)

complementary

Template 5'...GGCTAACTTAGC....3'
probe 3'...CCGATTGAATCG...5'

3'...CCGATTGAATCG...5'

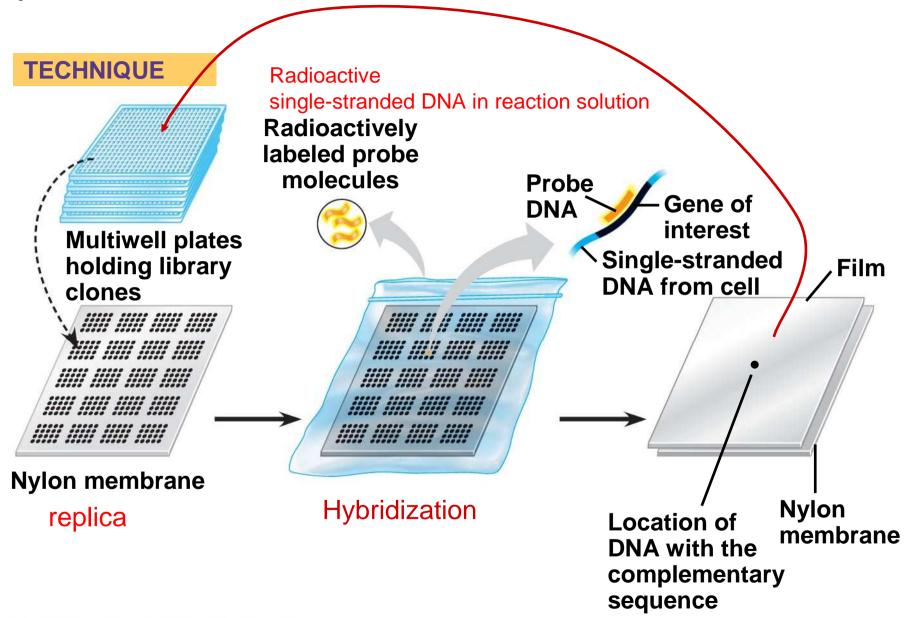
Sense vs. Anti-sense strand

Template 5'...CCATGGATACCG....3'

probe 5'...CCGATTGAATCG...3'
3'...GGTACCTATGGC...5'

- The DNA probe can be used to <u>screen a large</u> number of clones <u>simultaneously</u> for the gene of interest
- Once identified, the clone carrying the gene of interest can be cultured

Fig. 20-7



Expressing Cloned Eukaryotic Genes

- After a gene has been cloned, its protein product can be produced in larger amounts for research
- Cloned genes can be expressed as protein in either bacterial or eukaryotic cells

Bacterial Expression Systems

- Most popular methods to deliver plamids into E.coli – heat shock (42°C for 45 seconds)
- Several technical difficulties hinder expression of cloned eukaryotic genes in bacterial host cells: no expression, not soluble, not proper folded, etc.
- To overcome differences in promoters and other DNA control sequences, scientists usually employ an expression vector, a cloning vector that contains a highly active prokaryotic promoter

Eukaryotic Cloning and Expression Systems

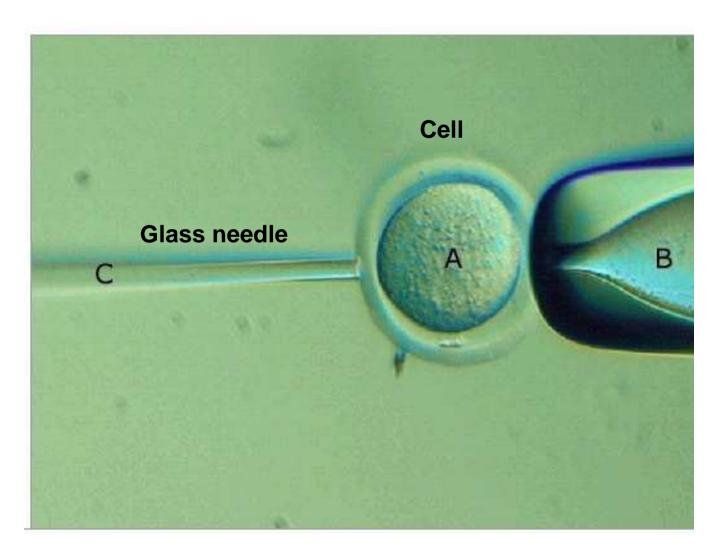
- The use of cultured eukaryotic cells as host cells and yeast artificial chromosomes (YACs) as vectors helps avoid gene expression problems
- YACs behave normally in mitosis and can carry more DNA than a plasmid
- Eukaryotic hosts can provide the <u>post-translational modifications</u> that many proteins require

Delivery gene into the eukaryotic cells

- One method of introducing recombinant DNA into <u>eukaryotic cells</u> is <u>electroporation</u>, applying a <u>brief electrical pulse</u> to create temporary holes in plasma membranes
- Alternatively, scientists can inject DNA into cells using microscopically thin needles (microinjection)
- Once inside the cell, the DNA is incorporated into the cell's DNA by natural genetic recombination

Microinjector

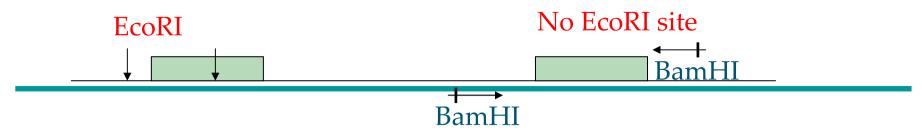




Suction

Amplifying DNA *in Vitro*: The Polymerase Chain Reaction (PCR)

- The polymerase chain reaction, PCR, can produce many copies of a specific target segment of DNA
- A three-step cycle heating, cooling, and replication — brings about a chain reaction that produces an exponentially growing population of identical DNA molecules

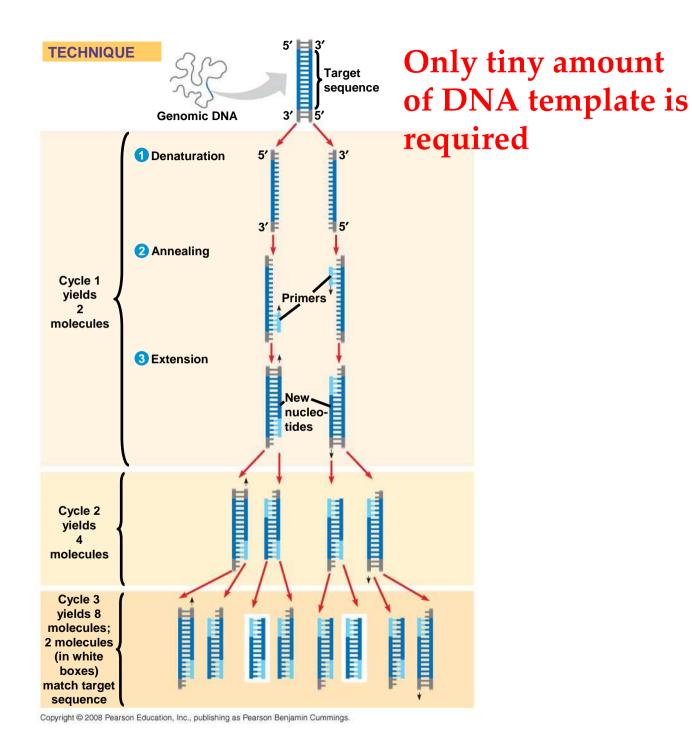


With polymerase chain reaction (PCR), any specific segment (with too many or no restriction enzymes available) of the target sequence within a DNA sample can be copied many times (amplified) completely *in vitro*.

Key components in a PCR reaction:

- --Uses <u>primers</u> (with unique restriction site) that bracket the desired sequence
- --Uses a heat-resistant DNA polymerase (active at high temperature) isolated from prokaryotes living in hot spring: extend primers in $5' \rightarrow 3'$ direction

Fig. 20-8



The polymerase chain reaction (PCR)

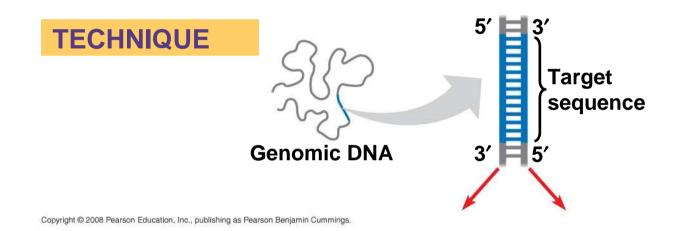
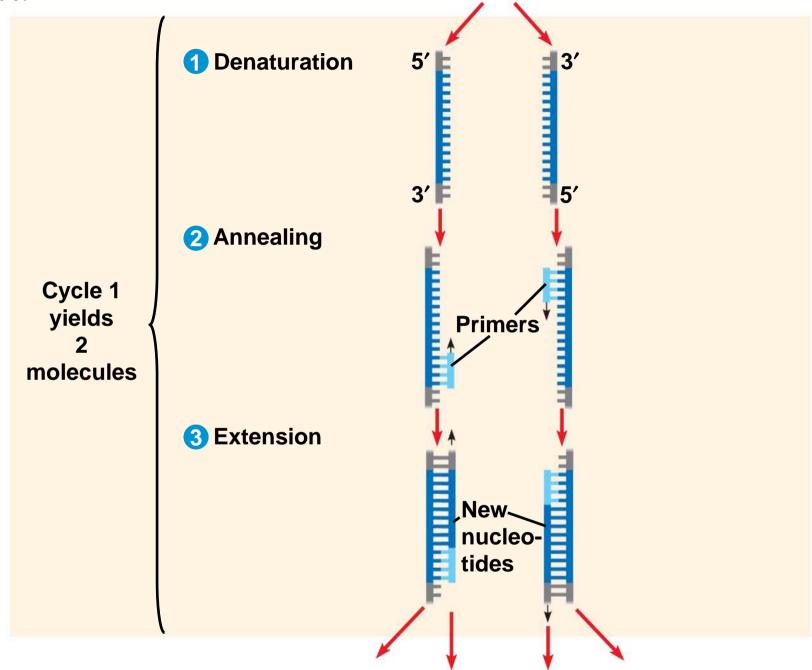
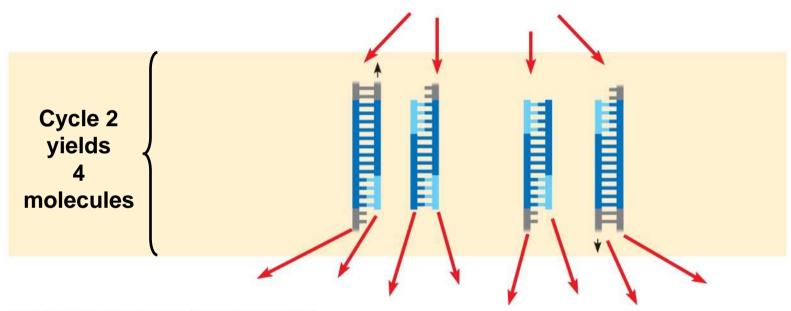
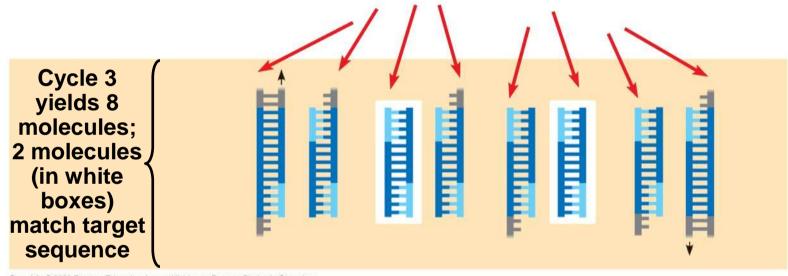


Fig. 20-8b





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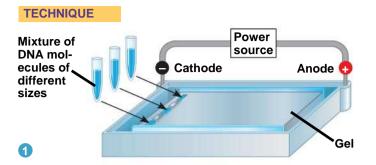
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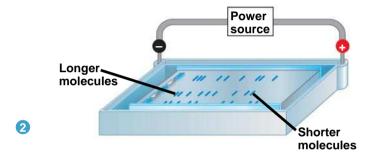
Concept 20.2: DNA technology allows us to study the sequence, expression, and function of a gene

- DNA cloning allows researchers to
 - Compare genes and alleles between individuals
 - Locate gene expression in a body
 - Determine the role of a gene in an organism
- Several techniques are used to analyze the DNA of genes

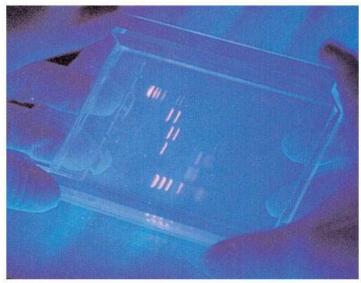
Gel Electrophoresis and Southern Blotting

- One indirect method of rapidly analyzing and comparing genomes is gel electrophoresis
- This technique uses a gel as a molecular sieve to separate nucleic acids or proteins by size
- A electrical current is applied that causes charged molecules to move through the gel
- Molecules are sorted into "bands" by their size

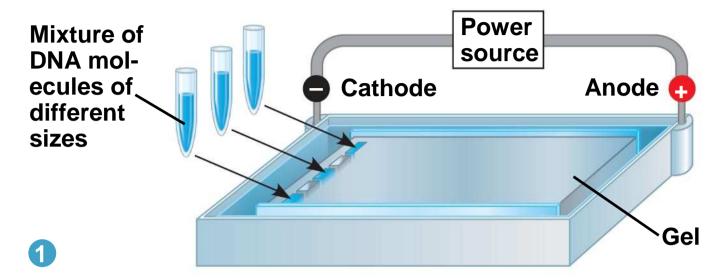


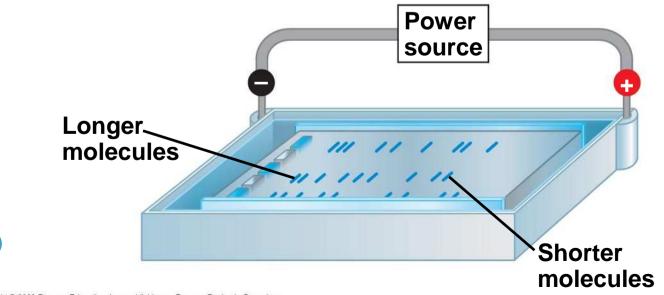


RESULTS



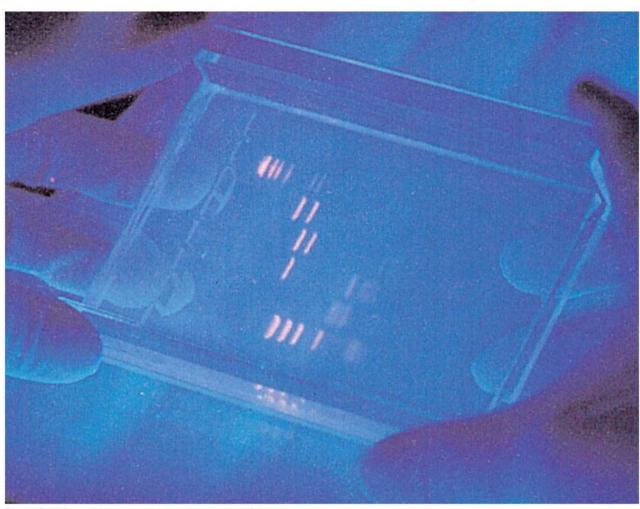
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Gel electrophoresis

RESULTS



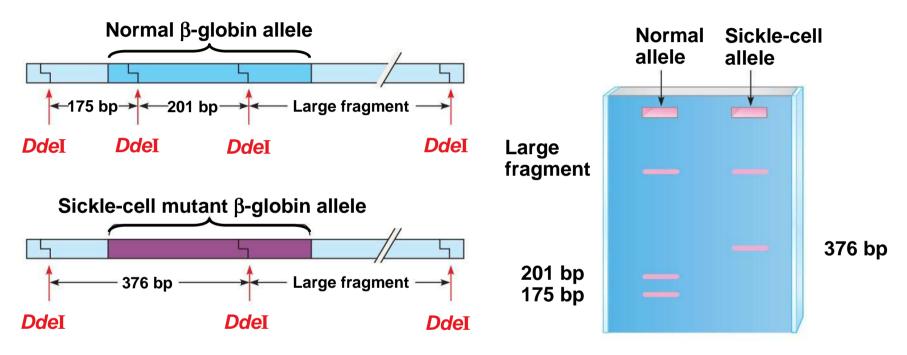
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Visualization by EtBr staining

Restriction fragment analysis

- In restriction fragment analysis, DNA fragments produced by restriction enzyme digestion of a DNA molecule are sorted by gel electrophoresis
 - A restriction enzyme will usually make many cuts in a DNA molecule
 - Yielding a set of restriction fragments
 - Gel electrophoresis to Separates DNA restriction fragments of different lengths:
 - Small DNA molecules from virus, plasmid— discrete bands
 - Large DNA molecules from eukaryotic chromosome --- smear
- Restriction fragment analysis is useful for comparing two different DNA molecules, such as two alleles for a gene
- The procedure is also used to prepare pure samples of individual fragments

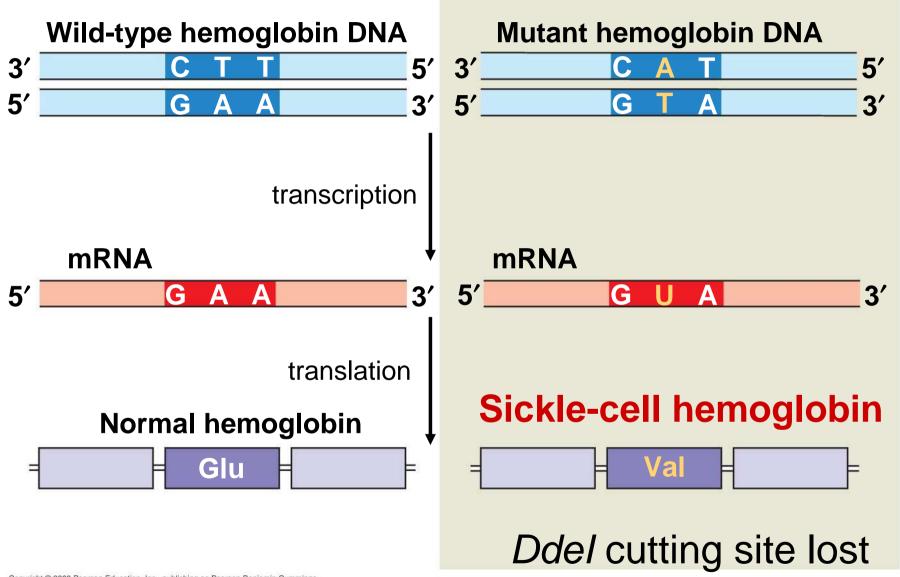
Using restriction fragment analysis to distinguish the normal and sickle-cell alleles of the β -globin gene



(a) *DdeI* restriction sites in normal and sickle-cell alleles of β-globin gene

(b) Electrophoresis of restriction fragments from normal and sickle-cell alleles

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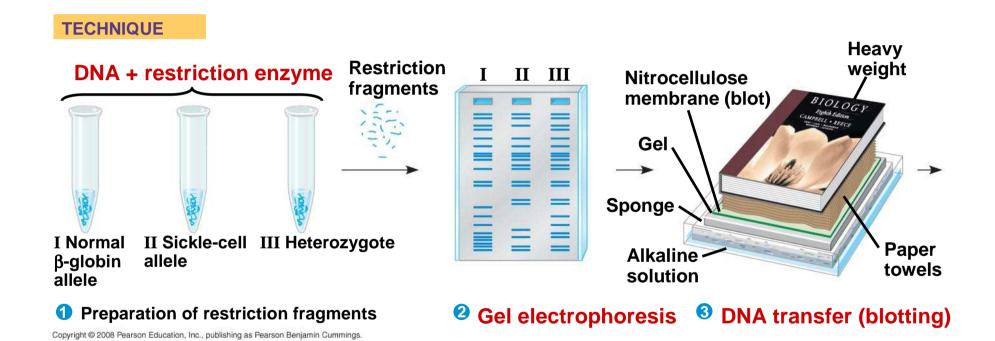


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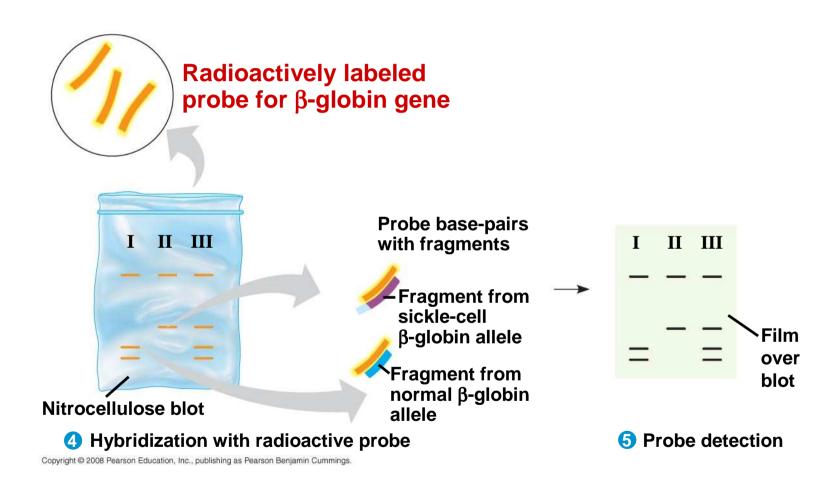
Southern blotting (DNA – DNA)

- A technique called Southern blotting combines gel electrophoresis of DNA fragments with nucleic acid hybridization
- Specific DNA fragments can be identified by Southern blotting, using labeled probes that hybridize to the DNA immobilized on a "blot" of gel

Southern blotting of DNA fragments

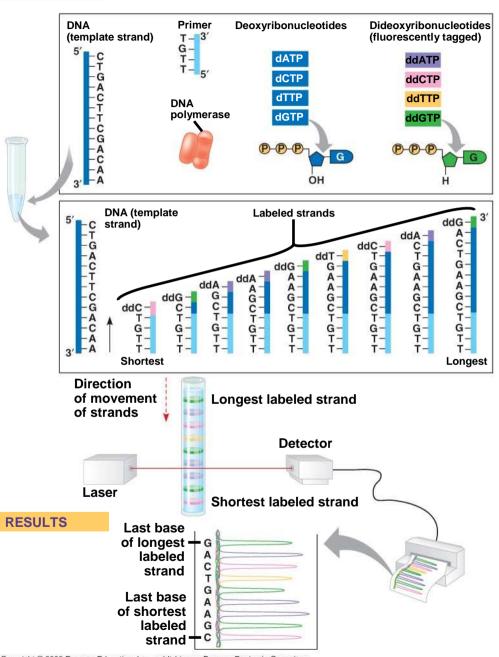


Southern blotting of DNA fragments

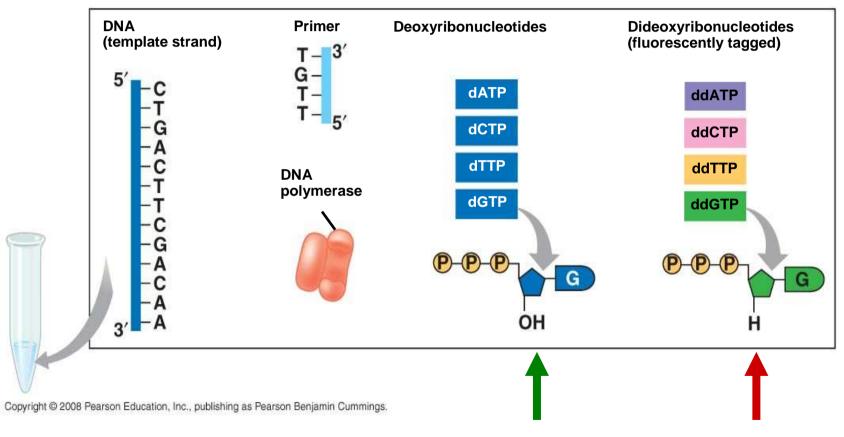


DNA Sequencing

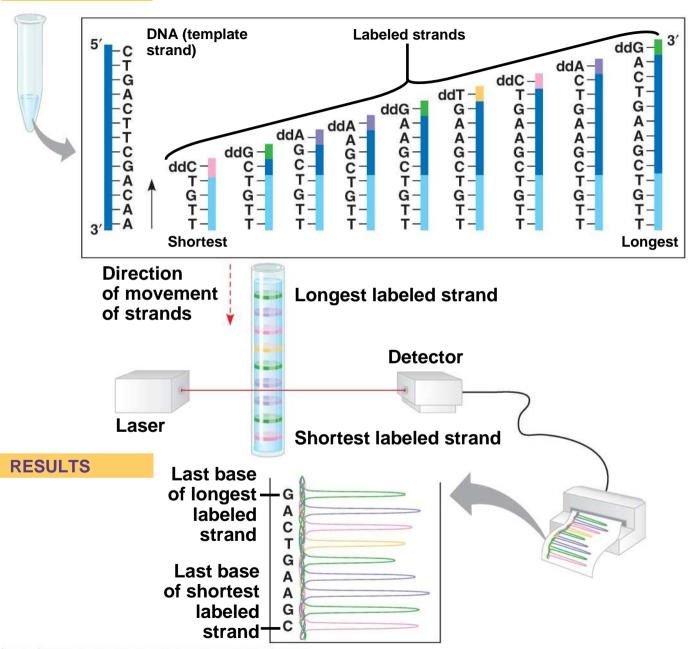
- Relatively short DNA fragments can be sequenced by the dideoxy chain termination method
- Modified nucleotides called dideoxyribonucleotides (ddNTP) attach to synthesized DNA strands of different lengths
- Each type of ddNTP is tagged with a distinct fluorescent label that identifies the nucleotide at the end of each DNA fragment
- The DNA sequence can be read from the resulting spectrogram



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Chain reaction terminated



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Latest Development on

Massive & Rapid whole genome DNA Sequencing



Getting personal

The commercialization of personal genomics is moving with dizzying speed and scientists need to find innovative ways of discussing the implications with consumers.

s the first conference on personal genomes opened earlier this month at Cold Spring Harbor Laboratory in New York, some present were wondering whether the event was a little premature. After all, only four people's genomes have so far been fully sequenced and assembled, and it's still quite difficult to interpret the genetic variation found in them (see page 1014). But the participants soon began to realize that, in one sense, the meeting was overdue. Increasingly, private companies are offering personal genome scans and genetic tests for sale — and consumers are buying them. Meanwhile, some scientists earlier this week made public parts of their genetic and medical data through the Personal Genome Project, spearheaded by George Church, a geneticist at Harvard University. In this context, the ethical, legal and social issues usually sidelined at such gatherings kept intruding with uncommon urgency.

The day before the meeting began, for instance, deCODE Genetics of Reykjavik, Iceland, began selling a US\$1,625 risk-assessment test for breast cancer, which surveys seven of the single-point genetic variations known as single nucleotide polymorphisms (SNPs). The company says that the test will help identify women with a significantly elevated risk of the disease.

But the test is worrying some oncologists and geneticists. Mary-Claire King of the University of Washington in Seattle told the meeting that the SNPs included in the test are present at slightly different background frequencies in different populations across Europe. She raised the possibility that rather than predicting breast-cancer risk, the test might instead be detecting population differences between the control and test groups included in the studies that linked the SNPs to disease.

Regulation of such tests is patchy at best, so women have little official

guidance when it comes to balancing the potentially confusing scientific and medical facts about a product against their fear of breast cancer. But, as *New York Times* reporter Amy Harmon told the meeting, the public desperately wants help in making such decisions.

Part of the problem is the information overload provided by the Internet. Consumers can point their browsers to a slew of content, which can come from both trusted sources and charlatans. And in the age of e-mail whisper campaigns, lies "Scientists cannot

can proliferate, opinions can replace facts, and experts are no longer trusted to know the truth.

"Scientists cannot put the genie back in the bottle."

Scientists can and should help the public sift through this information, interacting through blogs, newspaper articles or science cafes, for example. And if researchers are surprised at how quickly genomewide association studies have become consumer products, they need to realize that things will only move faster in the future, with findings moving from the lab to Internet chat rooms and people's lives with astonishing rapidity.

The issue was articulated at the meeting by Robert Cook-Deegan, an ethicist at Duke University in Durham, North Carolina, who pointed out that scientists cannot put the genie back in the bottle. Anyone can now access his or her genome information through a personal genomics scan. And if people can get that information, they will — with or without the advice of genetic counsellors, doctors or expert scientists. It is impossible to guess what issues this will raise as the science matures, although new discoveries will no doubt trigger a fresh and more complicated set of societal discussions. Scientists need to get creative about how they participate in these discussions, because they won't have the luxury of opting out.

NGS

Analyzing Gene Expression

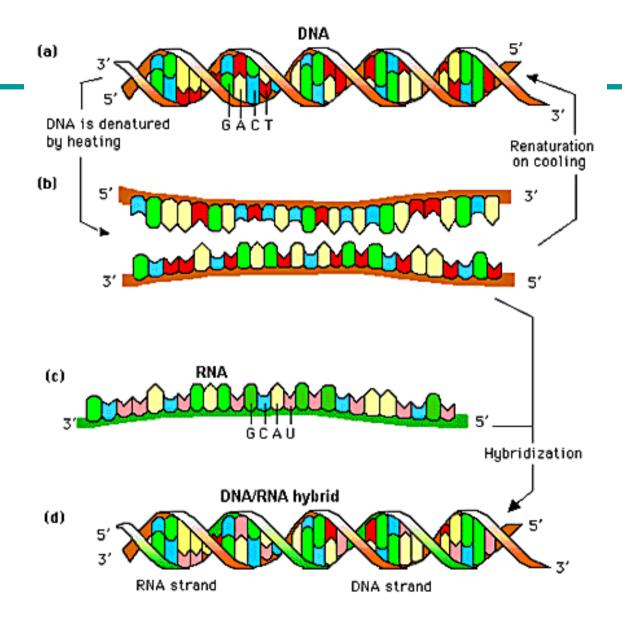
- Nucleic acid probes can hybridize with mRNAs transcribed from a gene
- Probes can be used to identify where or when a gene is transcribed in an organism

Studying the Expression of Single Genes

- Changes in the expression of a gene during embryonic development can be tested using
 - Northern blotting
 - Reverse transcriptase-polymerase chain reaction
- Both methods are used to compare mRNA from different developmental stages

Northern blotting (DNA-RNA)

- Northern blotting combines gel electrophoresis of mRNA followed by hybridization with a probe on a membrane
- Identification of mRNA at a particular developmental stage suggests protein function at that stage



Nucleic Acid Hybridization

RT-PCR

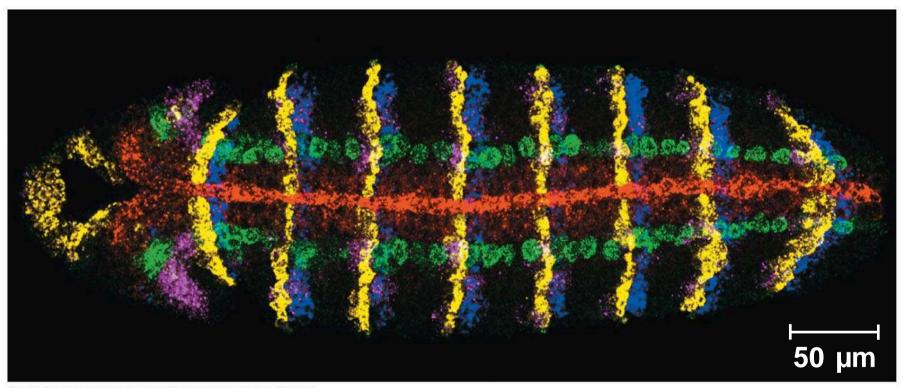
- Reverse transcriptase-polymerase chain reaction (RT-PCR) is quicker and more sensitive
- Reverse transcriptase is added to mRNA to make cDNA, which serves as a template for PCR amplification of the gene of interest
- The products are run on a gel and the mRNA of interest identified

TECHNIQUE 1 cDNA synthesis mRNAs 1 **cDNAs Primers 2** PCR amplification β-globin gene **3** Gel electrophoresis **Embryonic stages RESULTS** 6

In situ hybridization

 In situ hybridization uses fluorescent dyes attached to probes to identify the location of specific mRNAs in place in the intact organism

Determining where genes are expressed by *in* situ hybridization analysis



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Studying the Expression of Interacting Groups of Genes

- Automation has allowed scientists to measure expression of thousands of genes at one time using DNA microarray assays
- DNA microarray assays compare patterns of gene expression in <u>different tissues</u>, at <u>different</u> <u>times</u>, or under <u>different conditions</u>

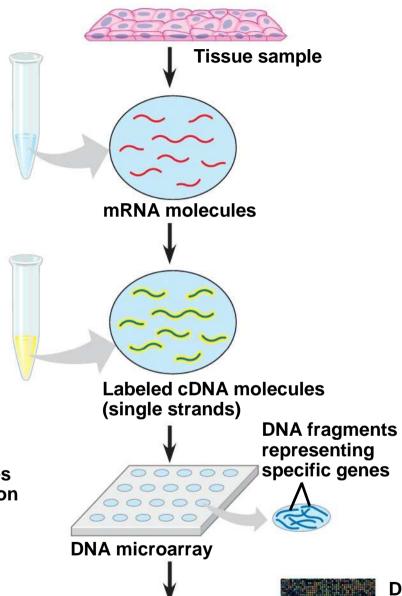
TECHNIQUE

1 Isolate mRNA.

Make cDNA by reverse transcription, using fluorescently labeled nucleotides.

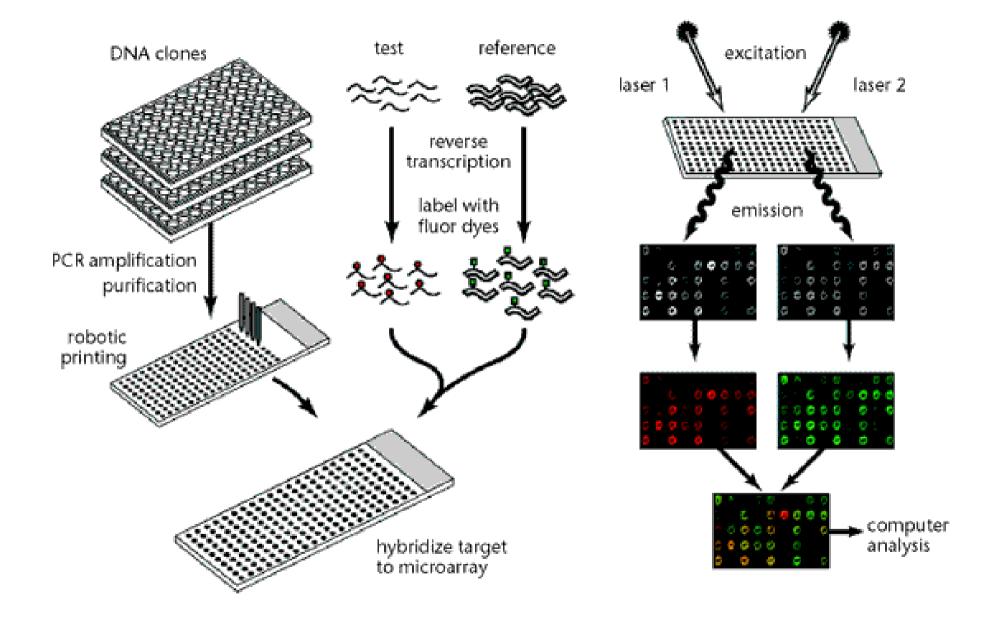
3 Apply the cDNA mixture to a microarray, a different gene in each spot. The cDNA hybridizes with any complementary DNA on the microarray.

4 Rinse off excess cDNA; scan microarray for fluorescence. Each fluorescent spot represents a gene expressed in the tissue sample.

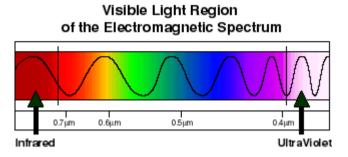




DNA microarray with 2,400 human genes



Fluor Dyes Cy3 and Cy5



In a typical cDNA microarray experiments, the samples will be visibly pink (Cy3) or blue (Cy5) in color after successful labeling.

If one looks at the visible light spectrum, the Cy3 emission filter (550-605 nM) covers the **blue-green** region, while the Cy5 emission filter (650-725 nM) covers the **orange-red** region.

Note:

- Cy3: Three-Tree- Tree is Green!
- Cy5: Five-Fire- Fire is Red!

Determining Gene Function

- One way to determine function is to disable the gene and observe the consequences – loss of function
- Using in vitro mutagenesis, mutations are introduced into a cloned gene, altering or destroying its function
- When the mutated gene is returned to the cell, the normal gene's function might be determined by examining the mutant's phenotype

- Gene expression can also be silenced using RNA interference (RNAi)
- Synthetic double-stranded RNA molecules matching the sequence of a particular gene are used to break down or block the gene's mRNA

Concept 20.3: Cloning organisms may lead to production of stem cells for research and other applications

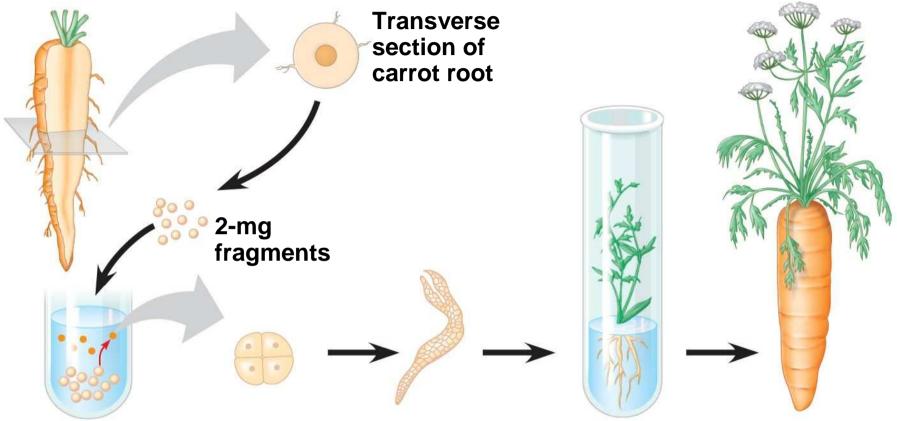
 Organismal cloning produces one or more organisms genetically identical to the "parent" that donated the single cell

Cloning Plants: Single-Cell Cultures

- One experimental approach for testing genomic equivalence is to see whether a differentiated cell can generate a whole organism
- A totipotent cell is one that can generate a complete new organism

EXPERIMENT

RESULTS



Fragments were cultured in nutrient medium; stirring caused single cells to shear off into the liquid.

Single cells free in began to divide.

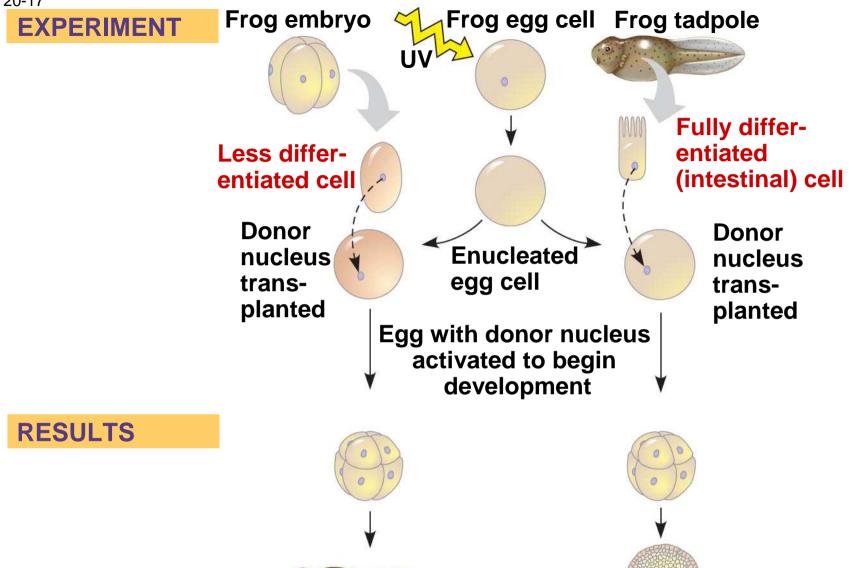
Embryonic plant developed from a cultured suspension single cell.

Plantlet was cultured on agar medium. Later it was planted in soil.

A single somatic carrot cell developed into a mature carrot plant.

Cloning Animals: Nuclear Transplantation

- In nuclear transplantation, the nucleus of an unfertilized egg cell or zygote is replaced with the nucleus of a differentiated cell
- Experiments with frog embryos have shown that a transplanted nucleus can often support normal development of the egg
- However, the older the donor nucleus, the lower the percentage of normally developing tadpoles



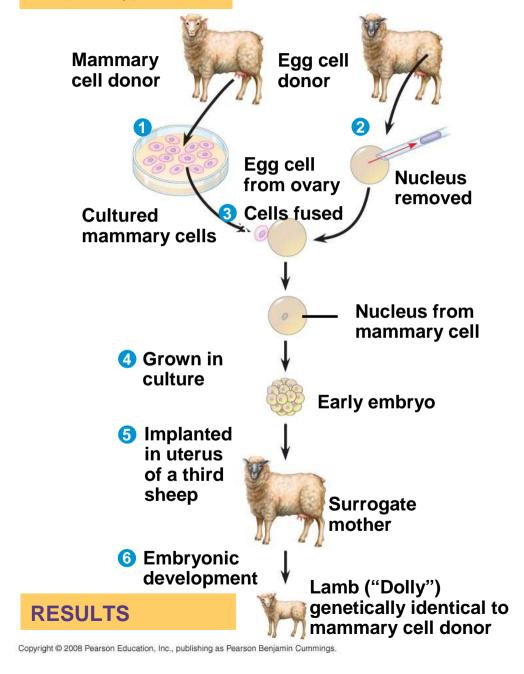
Most develop into tadpoles

Most stop developing before tadpole stage

Reproductive Cloning of Mammals

- In 1997, Scottish researchers announced the birth of Dolly, a lamb cloned from an adult sheep by nuclear transplantation from a differentiated mammary cell
- Dolly's premature death in 2003, as well as her arthritis, led to speculation that her cells were not as healthy as those of a normal sheep, possibly reflecting incomplete reprogramming of the original transplanted nucleus

TECHNIQUE



 Since 1997, cloning has been demonstrated in many mammals, including mice, cats, cows, horses, mules, pigs, and dogs

 CC (for Carbon Copy) was the first cat cloned; however, CC differed somewhat from her

female "parent"

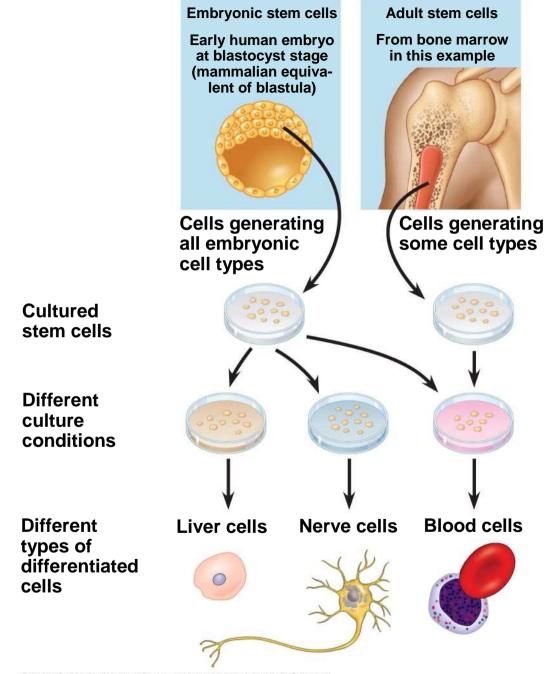


Problems Associated with Animal Cloning

- In most nuclear transplantation studies, only a small percentage of cloned embryos have developed normally to birth
- Many epigenetic changes, such as acetylation of histones or methylation of DNA, must be reversed in the nucleus from a donor animal in order for genes to be expressed or repressed appropriately for early stages of development

Stem Cells of Animals

- A stem cell is a relatively unspecialized cell that can reproduce itself indefinitely and differentiate into specialized cells of one or more types
- Stem cells isolated from early embryos at the blastocyst stage are called embryonic stem cells; these are able to differentiate into all cell types
- The adult body also has stem cells, which replace nonreproducing specialized cells



 The aim of stem cell research is to supply cells for the repair of damaged or diseased organs

Concept 20.4: The practical applications of DNA technology affect our lives in many ways

Many fields benefit from DNA technology and genetic engineering

Medical Applications

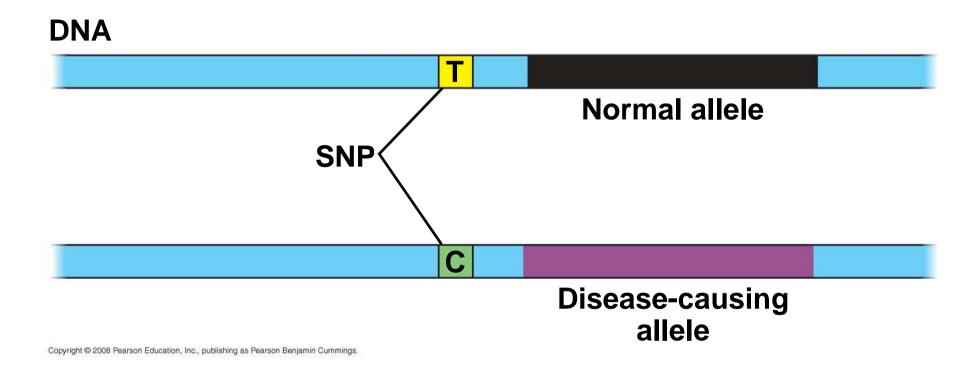
 One benefit of DNA technology is identification of human genes in which mutation plays a role in genetic diseases

Diagnosis of Diseases

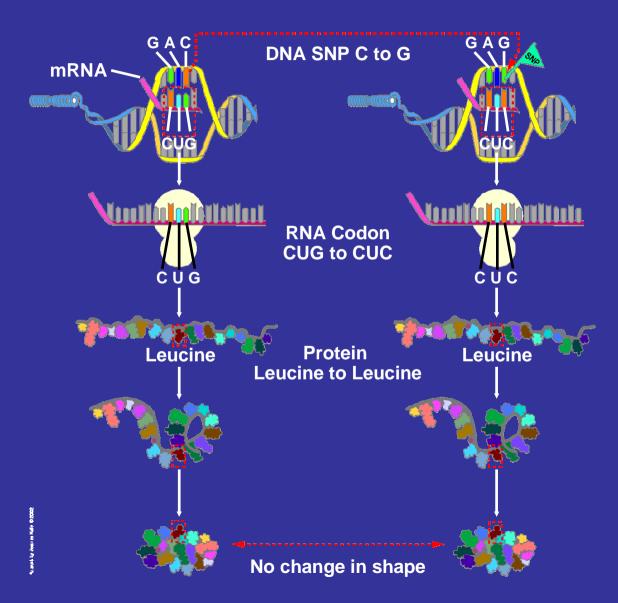
- Scientists can diagnose many human genetic disorders by using PCR and primers corresponding to cloned disease genes, then sequencing the amplified product to look for the disease-causing mutation
- Genetic disorders can also be tested for using genetic markers that are linked to the disease-causing allele

- Single nucleotide polymorphisms (SNPs) are useful genetic markers
- These are single base-pair sites that vary in a population
- When a restriction enzyme is added, SNPs result in DNA fragments with different lengths, or restriction fragment length polymorphism (RFLP)

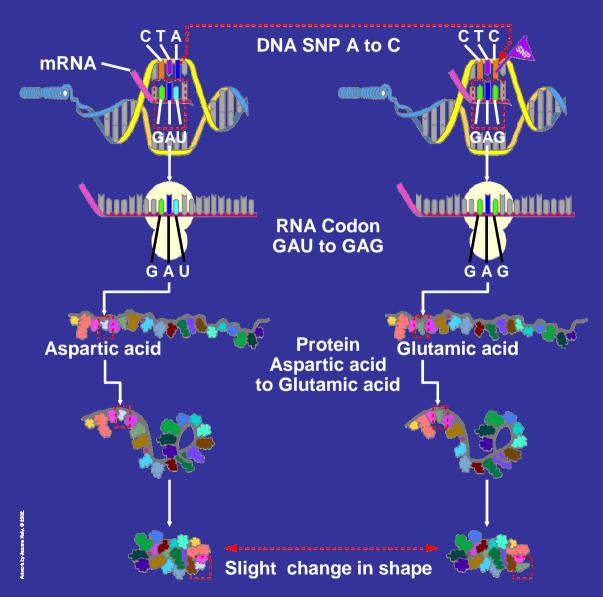
Single nucleotide polymorphisms (SNPs) as genetic markers for disease-causing alleles



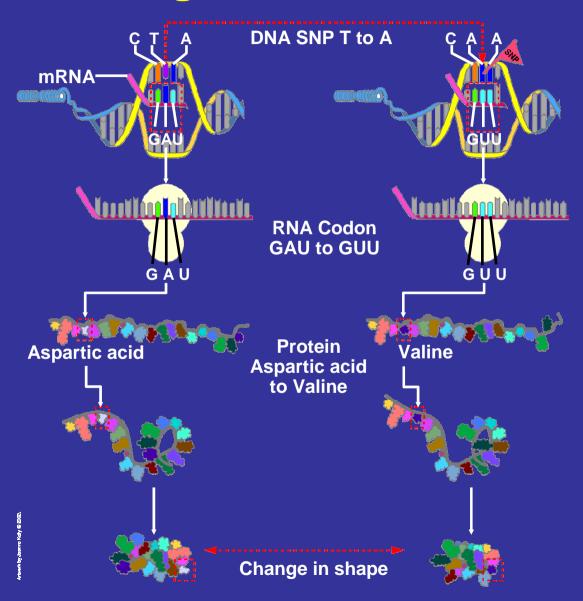
SNPs in Coding Regions – No Changes in Protein



SNPs in Coding Regions – Subtle, Harmless Changes in Protein

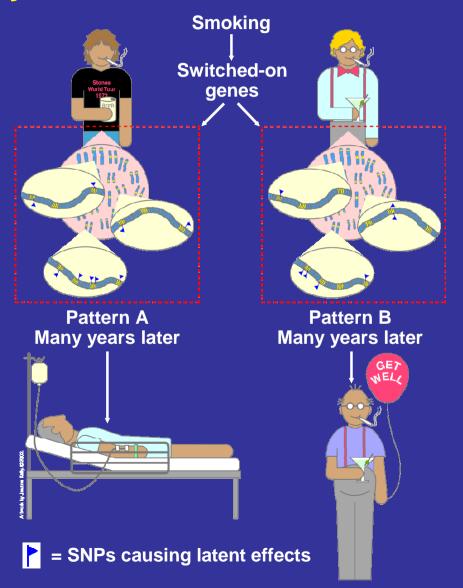


SNPs in Coding Regions – Harmful Changes in Protein – Mutations



SNPs in Coding Regions –

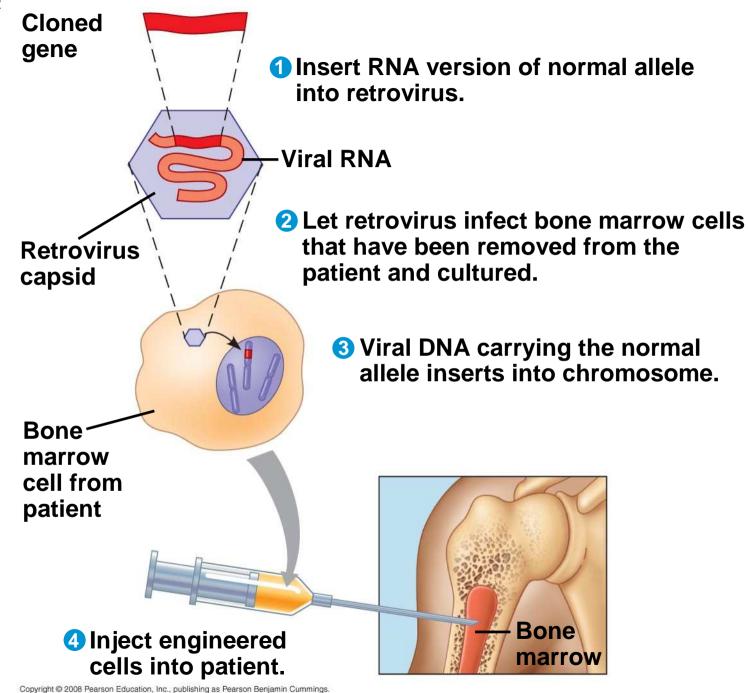
Subtle Changes in Proteins
That Only Switch on Under Certain Conditions



Human Gene Therapy

- Gene therapy is the alteration of an afflicted individual's genes
- Gene therapy holds great potential for treating disorders traceable to a single defective gene
- Vectors are used for delivery of genes into specific types of cells, for example bone marrow
- Gene therapy raises ethical questions, such as whether human germ-line cells should be treated to correct the defect in future generations

Fig. 20-22



Pharmaceutical Products

 Advances in DNA technology and genetic research are important to the development of new drugs to treat diseases

Synthesis of Small Molecules for Use as Drugs

- The drug imatinib is a small molecule that inhibits overexpression of a specific leukemiacausing receptor
- Pharmaceutical products that are proteins can be synthesized on a large scale

Protein Production in Cell Cultures

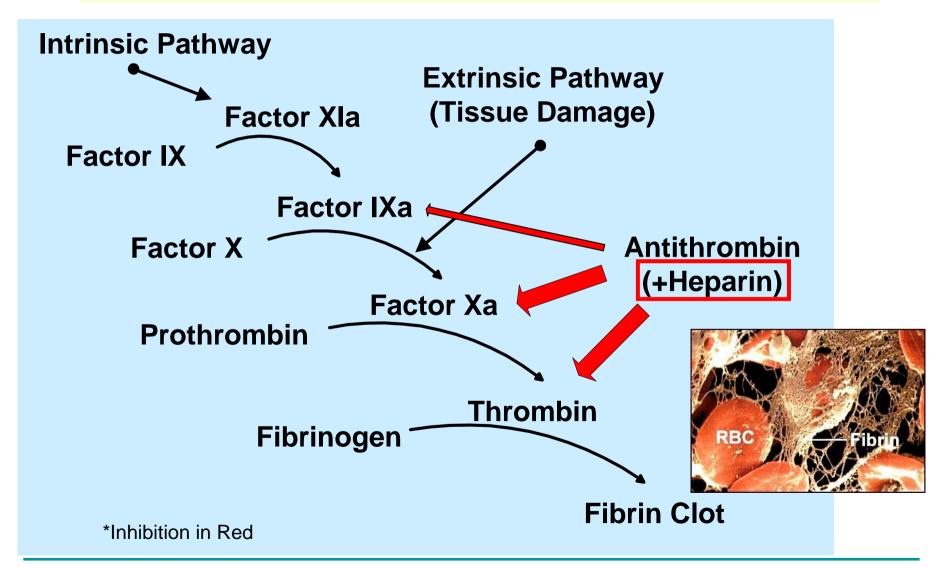
- Host cells in culture can be engineered to secrete a protein as it is made
- This is useful for the production of insulin, human growth hormones, and vaccines

"Pharm" Animals and Plants

Protein Production by "Pharm" Animals and Plants

- Transgenic animals are made by introducing genes from one species into the genome of another animal
 - Transgenic animals are pharmaceutical "factories," producers of large amounts of otherwise rare substances for medical use
- "Pharm" plants are also being developed to make human proteins for medical use

Regulation of the Blood Clotting Cascade By Antithrombin



Goats as "pharm" animals







Forensic Evidence and Genetic Profiles

- An individual's unique DNA sequence, or genetic profile, can be obtained by analysis of tissue or body fluids
- Genetic profiles can be used to provide evidence in criminal and paternity cases and to identify human remains
- Genetic profiles can be analyzed using RFLP analysis by Southern blotting

Short tandem repeats (STRs)

- Even more sensitive is the use of genetic markers called short tandem repeats (STRs), which are variations in the number of repeats of specific DNA sequences
- PCR and gel electrophoresis are used to amplify and then identify STRs of different lengths
- The probability that two people who are not identical twins have the same STR markers is exceptionally small

(a) This photo shows Earl Washington just before his release in 2001, after 17 years in prison.



Source of sample	STR marker 1	STR marker 2	STR marker 3
Semen on victim	17, 19	13, 16	12, 12
Earl Washington	16, 18	14, 15	11, 12
Kenneth Tinsley	17, 19	13, 16	12, 12

(b) These and other STR data exonerated Washington and led Tinsley to plead guilty to the murder.

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Environmental Cleanup

- Genetic engineering can be used to modify the metabolism of microorganisms
 - Some modified microorganisms can be used to extract minerals from the environment or degrade potentially toxic waste materials
- Biofuels make use of crops such as corn, soybeans, and cassava to replace fossil fuels

Agricultural Applications

 DNA technology is being used to improve agricultural productivity and food quality

Animal Husbandry

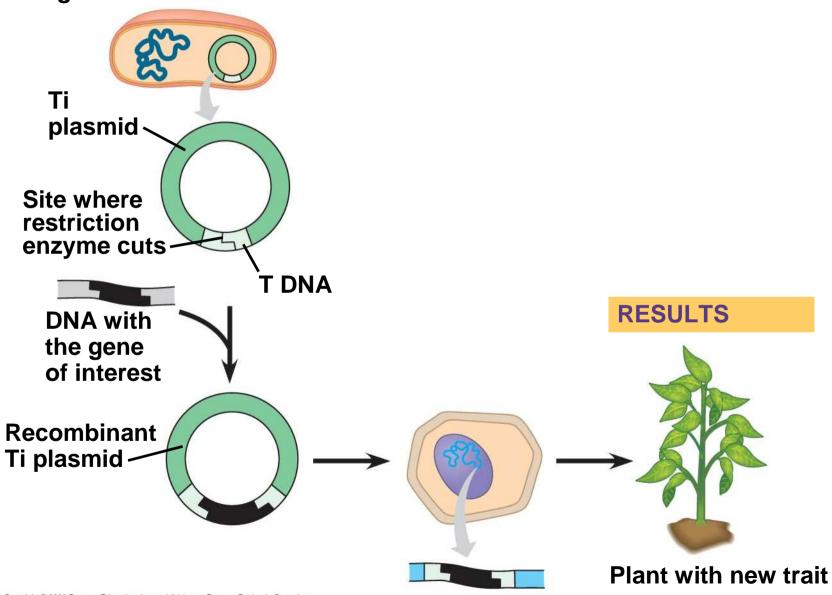
- Genetic engineering of transgenic animals speeds up the selective breeding process
- Beneficial genes can be transferred between varieties or species

Genetic Engineering in Plants

- Agricultural scientists have endowed a number of crop plants with genes for desirable traits
- The Ti plasmid is the most commonly used vector for introducing new genes into plant cells
- Genetic engineering in plants has been used to transfer many useful genes including those for herbicide resistance, increased resistance to pests, increased resistance to salinity, and improved nutritional value of crops

TECHNIQUE

Agrobacterium tumefaciens



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Safety and Ethical Questions Raised by DNA Technology

- Potential benefits of genetic engineering must be weighed against potential hazards of creating harmful products or procedures
- Guidelines are in place in the United States and other countries to ensure safe practices for recombinant DNA technology

GMO

- Most public concern about possible hazards centers on genetically modified (GM) organisms used as food
- Some are concerned about the creation of "super weeds" from the transfer of genes from GM crops to their wild relatives

Ethical guidelines

- As biotechnology continues to change, so does its use in agriculture, industry, and medicine
- National agencies and international organizations strive to set guidelines for safe and ethical practices in the use of biotechnology

You should now be able to:

- Describe the natural function of restriction enzymes and explain how they are used in recombinant DNA technology
- 2. Outline the procedures for cloning a eukaryotic gene in a bacterial plasmid
- 3. Define and distinguish between genomic libraries using plasmids, phages, and cDNA
- Describe the polymerase chain reaction (PCR) and explain the advantages and limitations of this procedure

- 5. Explain how gel electrophoresis is used to analyze nucleic acids and to distinguish between two alleles of a gene
- Describe and distinguish between the Southern blotting procedure, Northern blotting procedure, and RT-PCR
- Distinguish between gene cloning, cell cloning, and organismal cloning
- 8. Describe how nuclear transplantation was used to produce Dolly, the first cloned sheep

- Describe the application of DNA technology to the diagnosis of genetic disease, the development of gene therapy, vaccine production, and the development of pharmaceutical products
- 10.Define a SNP and explain how it may produce a RFLP
- 11.Explain how DNA technology is used in the forensic sciences

12. Discuss the safety and ethical questions related to recombinant DNA studies and the biotechnology industry