



DNA  
Genetic Code of Life



Entire Genetic Code  
of a Bacteria



DNA Fingerprinting



Cloning: Ethical Issues  
and Future Consequences



Plants of Tomorrow

# HC70A Winter 2008 Genetic Engineering in Medicine, Agriculture, and Law Professor Bob Goldberg

## Lecture 3 What Are Genes & How Do They Work-Part Two



DNA  
Genetic Code of Life



Entire Genetic Code  
of a Bacteria



DNA Fingerprinting



Cloning: Ethical Issues  
and Future Consequences



Plants of Tomorrow

## THEMES

1. What is the Function of a Gene-Review?
2. How Does DNA Replication Occur?
3. What is the Polymerase Chain Reaction (PCR) and How is PCR Used?
4. How Do Mutations Occur?
5. How Can Pedigrees Be Used To Follow the Inheritance of Mutant Genes?
6. How Do Mutations Change Phenotypes?
7. What is the Colinearity Between Genes & Proteins (how does DNA→protein)?
8. What is the Genetic Code?
9. How Do Gene Expression Processes Differ in Eukaryotes & Prokaryotes?
10. How Can Splicing Cause One Gene To Specify Several Different Proteins?
11. Yo!-It's in the DNA Sequences-What Are the Implications For Genetic Engineering?

**The Beginning of a Gene is its:**

- a. 5' End**
- b. 3' End**

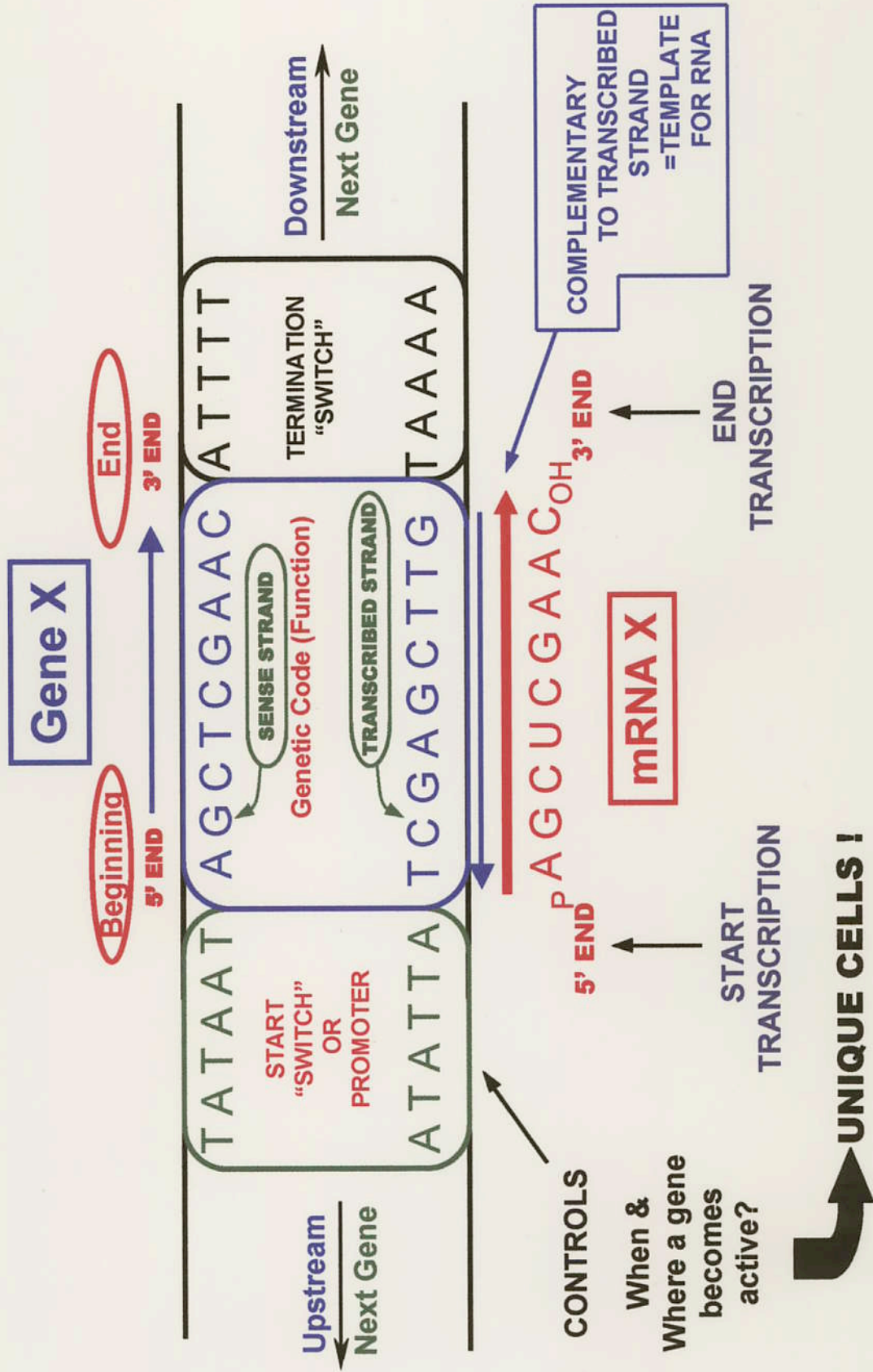
**The Sense Strand of a Gene Is Transcribed?**

- a. Yes**
- b. No**

The Sequence of a mRNA is the Same As the Sequence of the:

- a. Sense Strand
- b. Antisense Strand

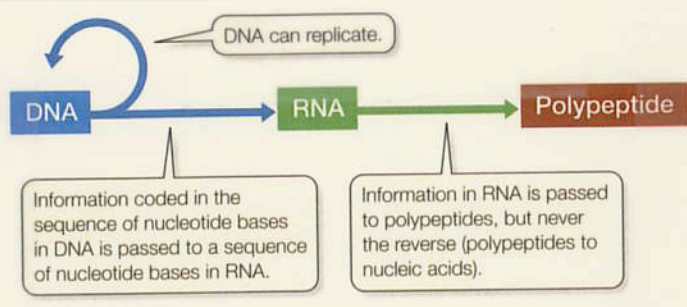
A Gene is a Specific DNA Sequence That Directs the Expression of a Unique Trait



Note: mRNA Sequence = Sense Strand Sequence

# HOW DO GENES WORK?

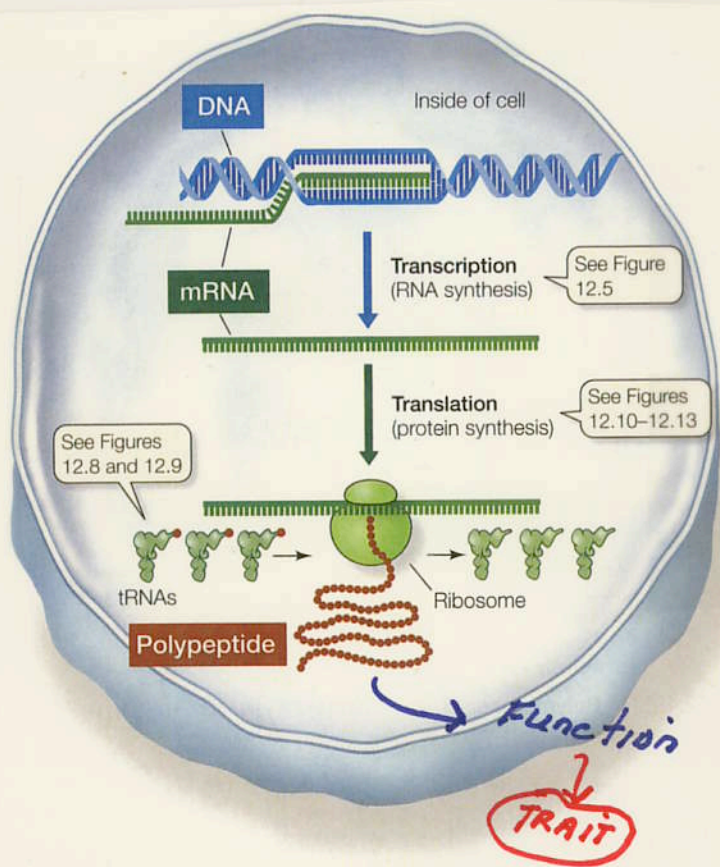
① Replication



② Gene Activity to Function

12.2 The Central Dogma Information flows from DNA to RNA to polypeptide, as indicated by the arrows.

Gene Activity  
↓  
Protein  
↓  
Function  
↓  
Phenotype (Trait)

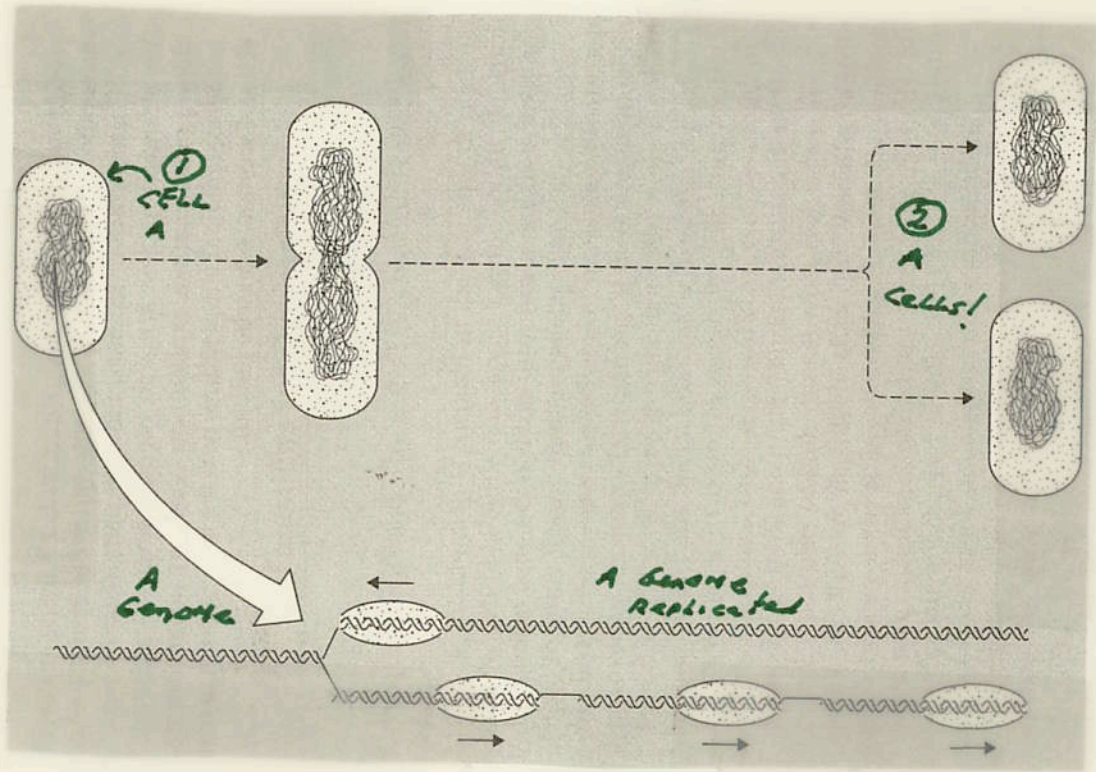


12.3 From Gene to Protein This diagram summarizes the processes of gene expression in prokaryotes. In eukaryotes, the processes are somewhat more complex.

A GENE IS NOT EXPRESSED UNLESS A FUNCTIONAL PROTEIN IS PRODUCED!

①

HOW ARE GENES REPLICATED EACH CELL GENERATION?



HOW IS THE DNA SEQUENCE COPIED / REPLICATED EACH CELL DIVISION?

PASS ON GENES TO NEXT GENERATION Precisely?

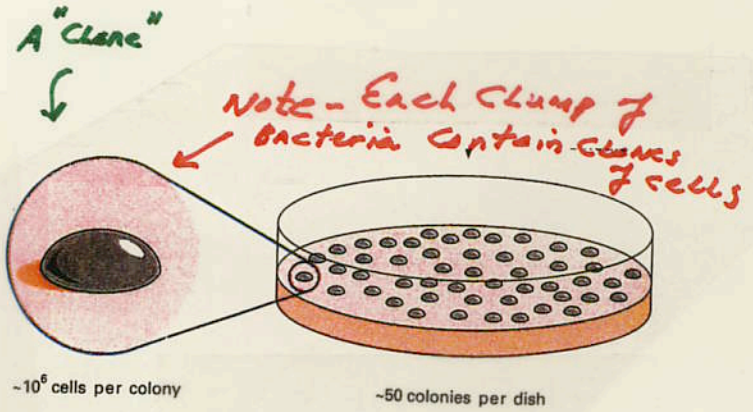
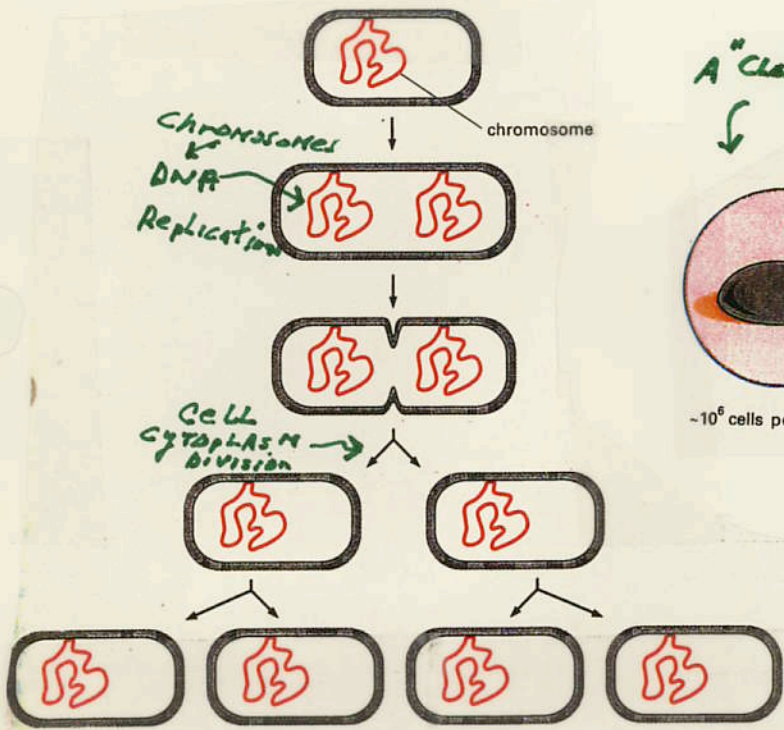
BASIS OF LIFE!



**The Structure of DNA "Allows" the Genotype of a Cell To Be Passed on From Generation to Generation?**

- a. Yes**
- b. No**

**GENES ARE REPLICATED DURING EACH CELL DIVISION**



A BACTERIAL colony contains many copies of SAME CELL or CLONES which are genetically identical!

**EACH DAUGHTER CELL CONTAINS THE SAME COLLECTION OF GENES**

CLONES!

MAJOR PROPERTIES OF GENETIC MATERIAL  
Replication + Stability

THE SEQUENCE OF EACH DNA STRAND MUST BE MAINTAINED DIVISION AFTER DIVISION

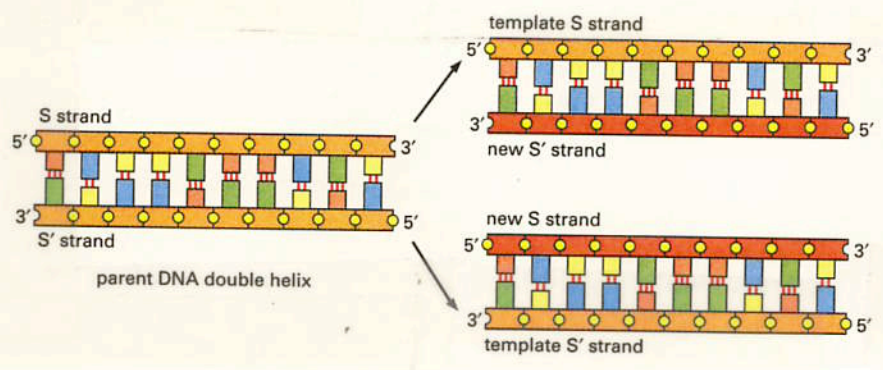


Figure 5-2 The DNA double helix acts as a template for its own duplication. Because the nucleotide A will successfully pair only with T, and G only with C, each strand of DNA can serve as a template to specify the sequence of nucleotides in its complementary strand by DNA base-pairing. In this way, a double-helical DNA molecule can be copied precisely.

HOW DOES THAT OCCUR?  
PROPERTY OF THE DNA MOLECULE

NOTE → SEQUENCE & POLARITY

DNA REPLICATION OCCURS SEMI-CONSERVATIVELY

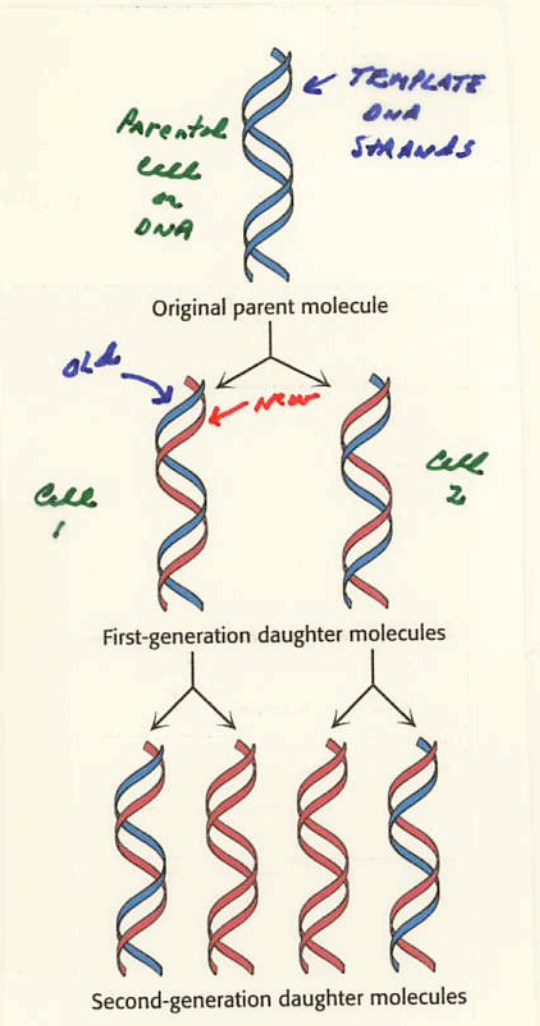


FIGURE 5.16 Diagram of semiconservative replication.

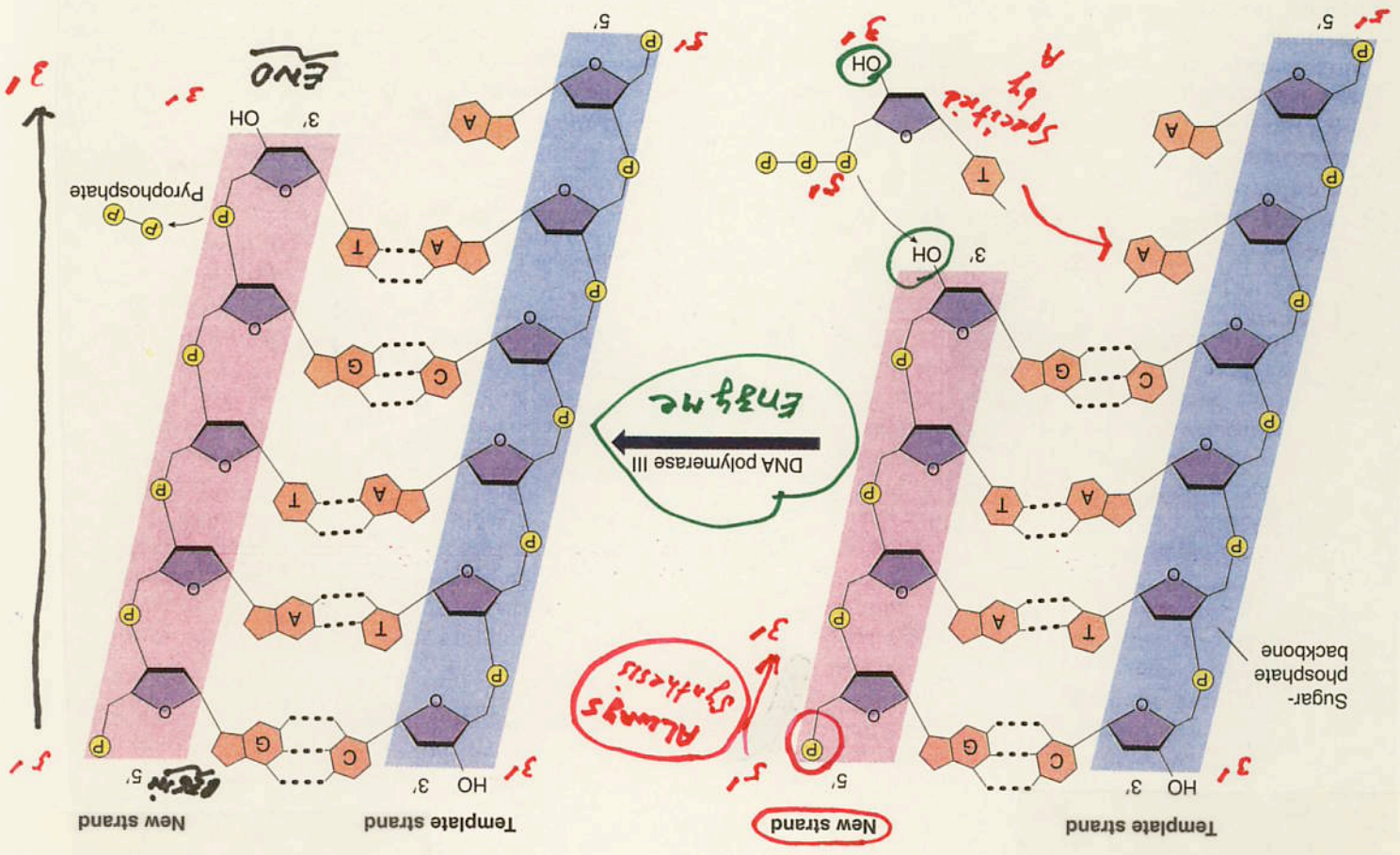
- ① DNA STRUCTURE ALLOWS DNA sequence to be Maintained  
Complementary bases
- ② Each STRAND serves as a Template for the synthesis of a complementary strand of DNA
- ③ NEW MOLECULES of DNA ARE PRECISE COPIES OF PARENTAL DNA - ONE TEMPLATE STRAND + ONE NEWLY SYNTHESIZED COMPLEMENTARY STRAND!

Meselson & Stahl  
1957

②  
 NOTE 5'P \* 3'OH  
 SEQUENCE IS SPECIFIED BY  
 COMPLEMENTARY  
 BASES

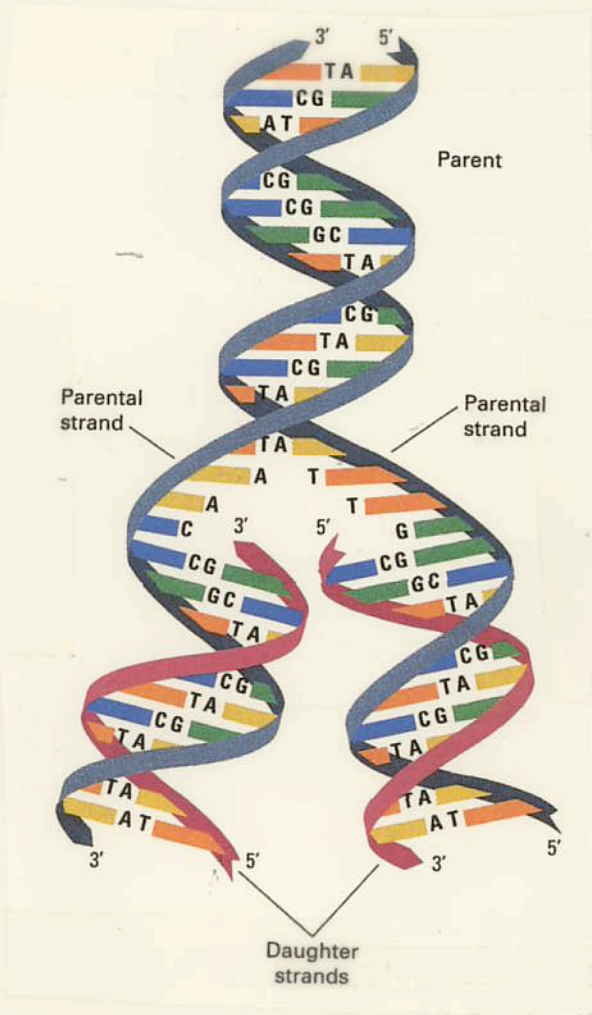
POLARITY  
 ||  
 SEQUENCE

FIGURE 14.14 How nucleotides are added in DNA replication. DNA polymerase III, along with other enzymes, catalyzes the addition of nucleotides to the growing complementary strand of DNA. When a nucleotide is added, two of its phosphates are lost as pyrophosphate.



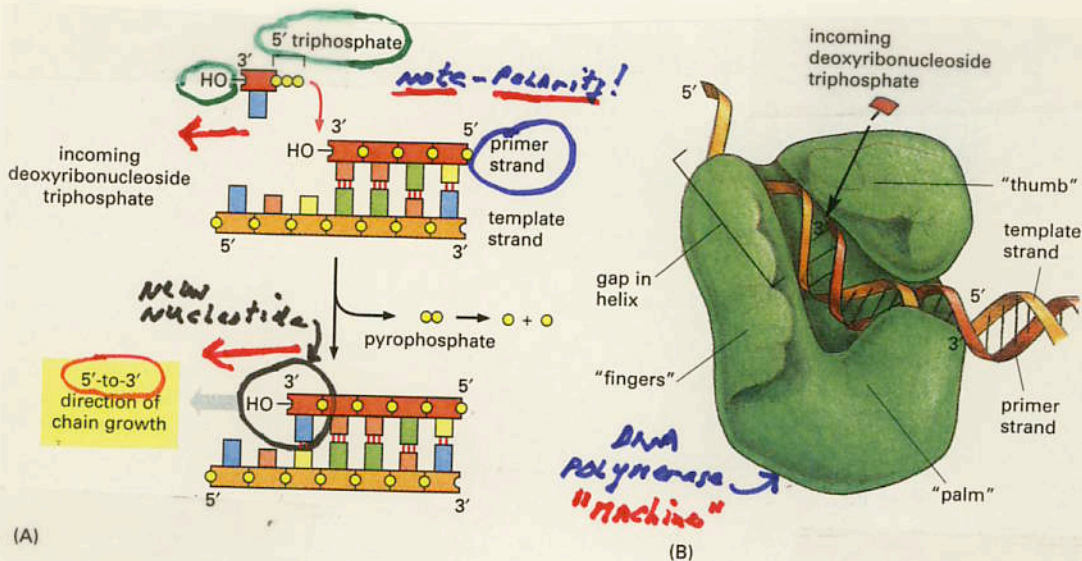
DNA SEQUENCE OF ONE STRAND IS A  
 TEMPLATE FOR THE NEW STRAND

THE DNA SEQUENCE IS MAINTAINED  
GENERATION TO GENERATION



THE DNA SEQUENCE "LIVES"  
FOREVER!

**DNA REPLICATION REQUIRES AN ENZYME - DNA POLYMERASE**

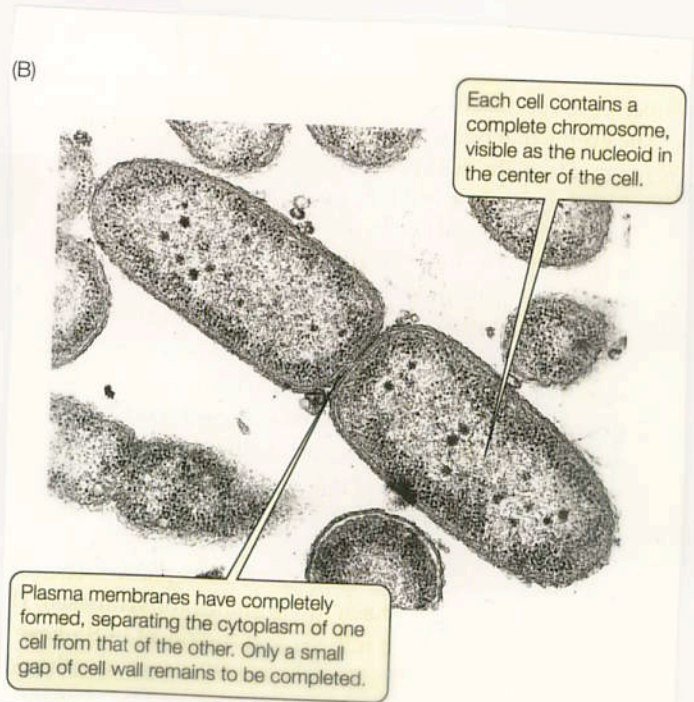
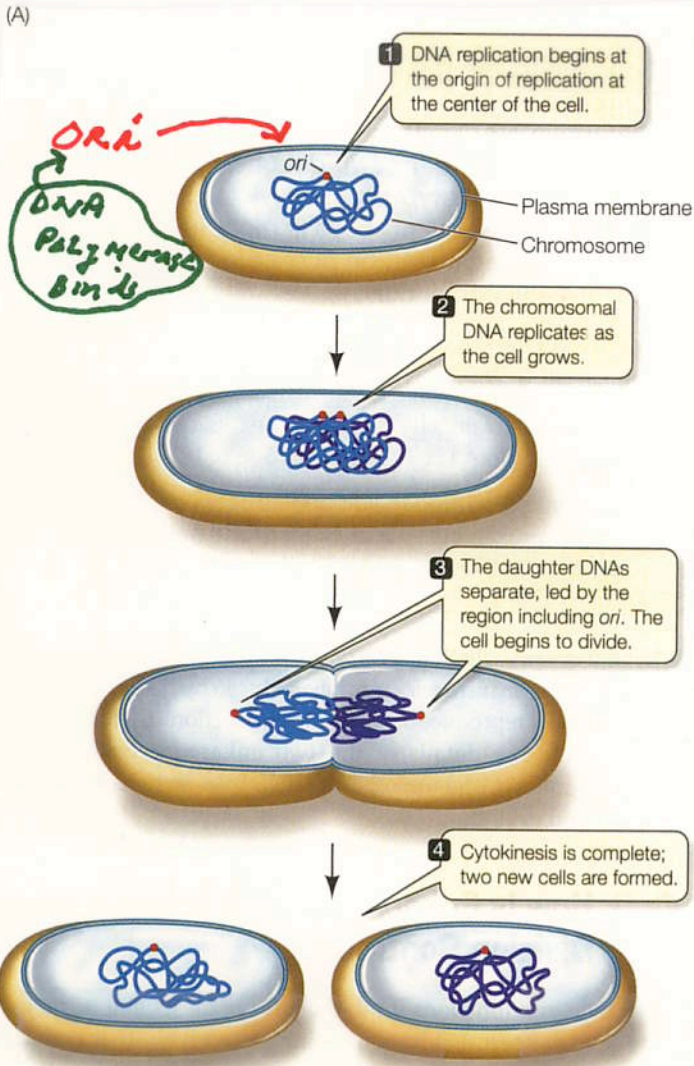


**DNA POLYMERASE CATALYZES PHOSPHODIESTER BONDS AND "COPIES" THE TEMPLATE**

**NUCLEOTIDES ARE ALSO NEEDED**

**NEED A PRIMER + TEMPLATE + DNA POLYMERASE + NTS**

# DNA Replication Requires AN ORIGIN of Replication



9.2 Prokaryotic Cell Division (A) The process of cell division in a bacterium. (B) These two cells of the bacterium *Pseudomonas aeruginosa* have almost completed cytokinesis.

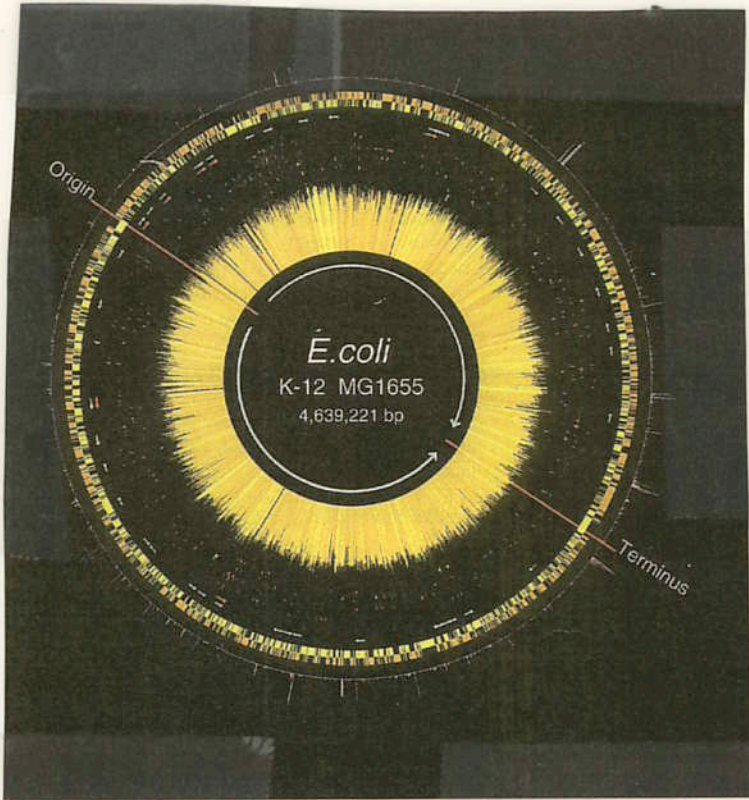
## DNA Replication Also Requires:

- ① Template
- ② Nucleotides
- ③ DNA Polymerase (Machine)
- ④ "Primer" to Start Replication



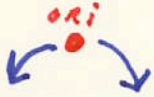
ORI

DNA Replication Starts at the ORIGIN OF REPLICATION



Key: ● tRNA genes; ● rRNA genes; — origin and terminus of replication

DNA Replication is bidirectional from the ori!!

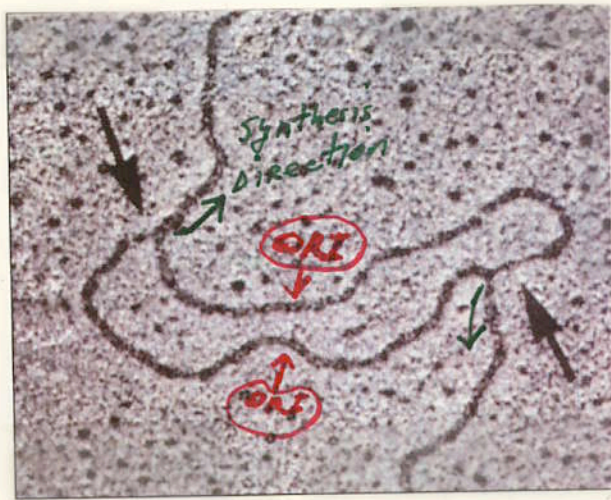


Hypothesis for Two Direction Synthesis?

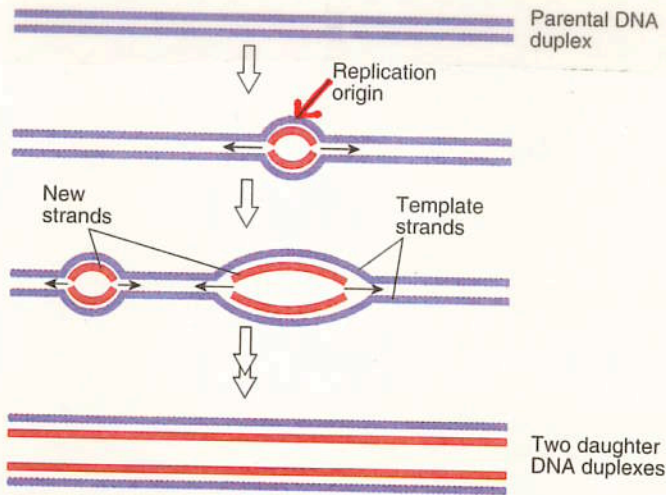
DNA Polymerase Binds to the ORIGIN OF REPLICATION (ORI) TO Begin DNA Synthesis

CONTROL DIVISION?

DNA in The PROCESS OF BEING REPLICATED



Replication Moves Bidirectionally FROM ORIGIN



**FIGURE 14.13**  
**Origins of replication.** At a site called the replication origin, the DNA duplex opens to create two separate strands, each of which can be used as a template for a new strand. Eukaryotic DNA has multiple origins of replication.

CONCEPT!

Foreign DNA segments use ori of chromosomes/DNA they are inserted into

e.g., bacteria insect<sup>R</sup> gene  
 ↳ use plant ori

The ORIGIN OF REPLICATION IS A SPECIFIC SEQUENCE

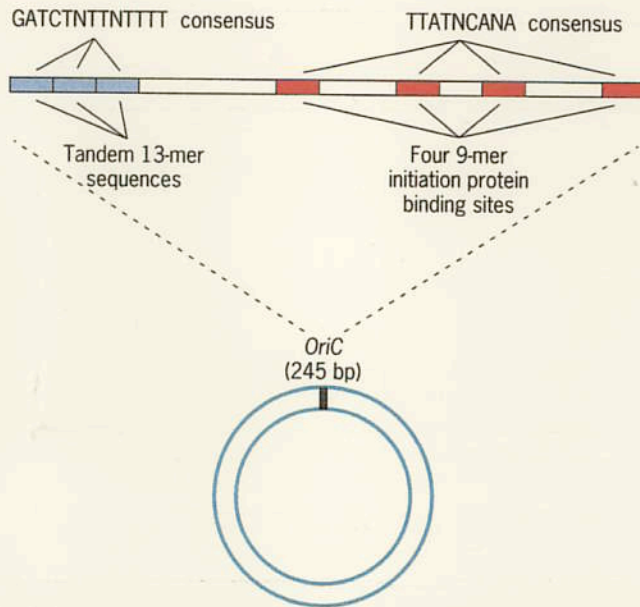


Figure 11.6 Structure of OriC, the single origin of replication in the *E. coli* chromosome.

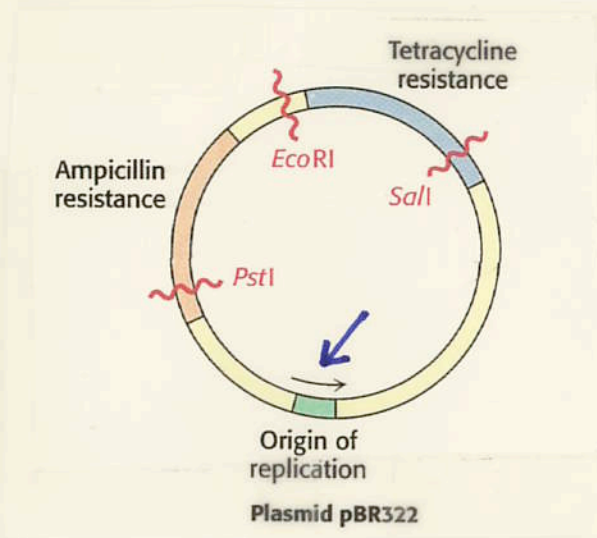
How CLONE AN ORIGIN OF REPLICATION?

Specific Sequence  
What does this mean for Genetic Engineering?

What is the significance for Genetic Engineering?

CAN Replicating "Chromosomes" Be Made?

VECTORS ARE NEEDED TO REPLICATE GENES IN SPECIFIC CELLS



- ① ORI is a specific sequence
  - ② ORI is genome & organism specific
  - ③ DNA POLYMERASES ARE SPECIFIC FOR EACH ORGANISM
- ∴ need correct ORI to replicate gene in a specific organism!

Note → need bacterial ori to clone human gene in bacteria. Need human ori to replicate a bacterial gene in human cell.

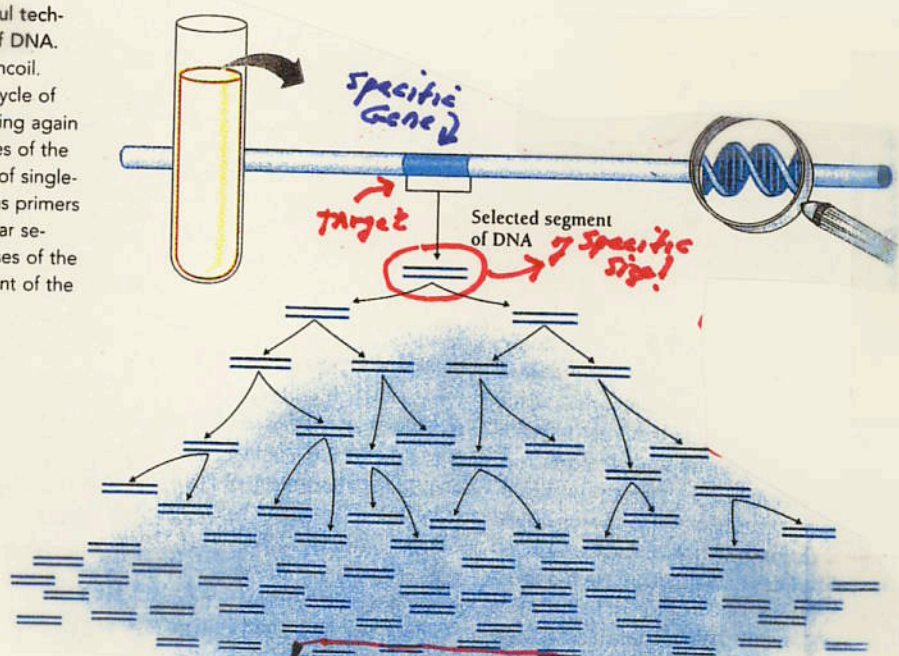
Yo! It's in the sequence = function

∴  
 Vectors CAN BE ENGINEERED!  
 ORIs CAN BE CLONED/SYNTHEZISED!

MODULAR

The Polymerase Chain Reaction or PCR  
is a Molecular  
Xerox Machine

Figure 13-2 PCR is a simple, powerful technique for multiplying specific sequences of DNA. A. When DNA is heated, the two strands uncoil. They are then cooled and replicated. The cycle of heating, cooling, replicating, and then heating again is repeated until millions or billions of copies of the sequence are obtained. B. Short segments of single-stranded DNA called oligonucleotides act as primers and allow researchers to replicate a particular sequence, not just any DNA. The 20 or so bases of the oligonucleotide pair with the correct segment of the DNA and initiate replication.



How many copies  
after 10 replication  
cycles?

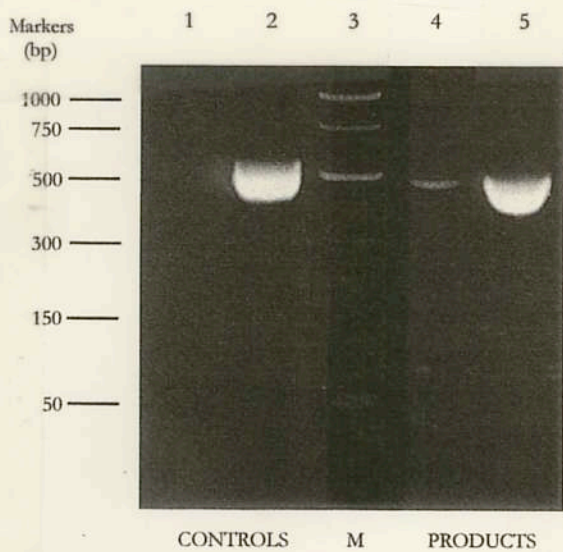
PCR HAS REVOLUTIONIZED DNA ANALYSIS!  
SPECIFIC DNA SEQUENCES / GENES CAN BE  
"COPIED" DIRECTLY FROM "TINY" AMOUNTS OF DNA!

NO CLONING NEEDED!

but need sequence!  
∴ have to clone "gene" first

DNA Polymerase  
↑ ↑ ↑

USING GEL ELECTROPHORESIS TO VISUALIZE PCR PRODUCTS



TARGET SPECIFIC BAND

Specific Diagnostic DNA Band Unique to DNA Sequence Being Amplified

Fig. 7.8. Visualisation of PCR products of ornithine decarboxylase on an agarose gel. Lane 1 – negative control (no DNA); lane 2 – positive control (cloned ornithine decarboxylase fragment, 460 bp); lane 3 – PCR size markers; lanes 4 and 5 – PCR product using rat liver genomic DNA and the ornithine decarboxylase primers used in lane 2. Lane 4 shown product after 15 cycles, lane 5 after 30 cycles of PCR. Photograph courtesy of Dr F. McKenzie.

CAN AMPLIFY ONE DNA SEQUENCE IN A WHOLE GENOME

## PCR Has MANY Uses That HAVE Changed MANY Fields

- ① Amplify any DNA Sequence or Gene from "TINY" Amounts of DNA. *NO Need For Bacteria or Vector!*
- ② Study DNA from Limited Sources: *a single hair, an ancient insect/plant, a bone fragment, cheek cell*
- ③ Used in: *Forensics, DNA Fingerprinting, Law, Evolution, Disease diagnosis, Identification, Pathogen Identification, Basic Molecular Biology, Evolution Studies, mRNA Detection*

Need as little as ONE molecule of DNA!

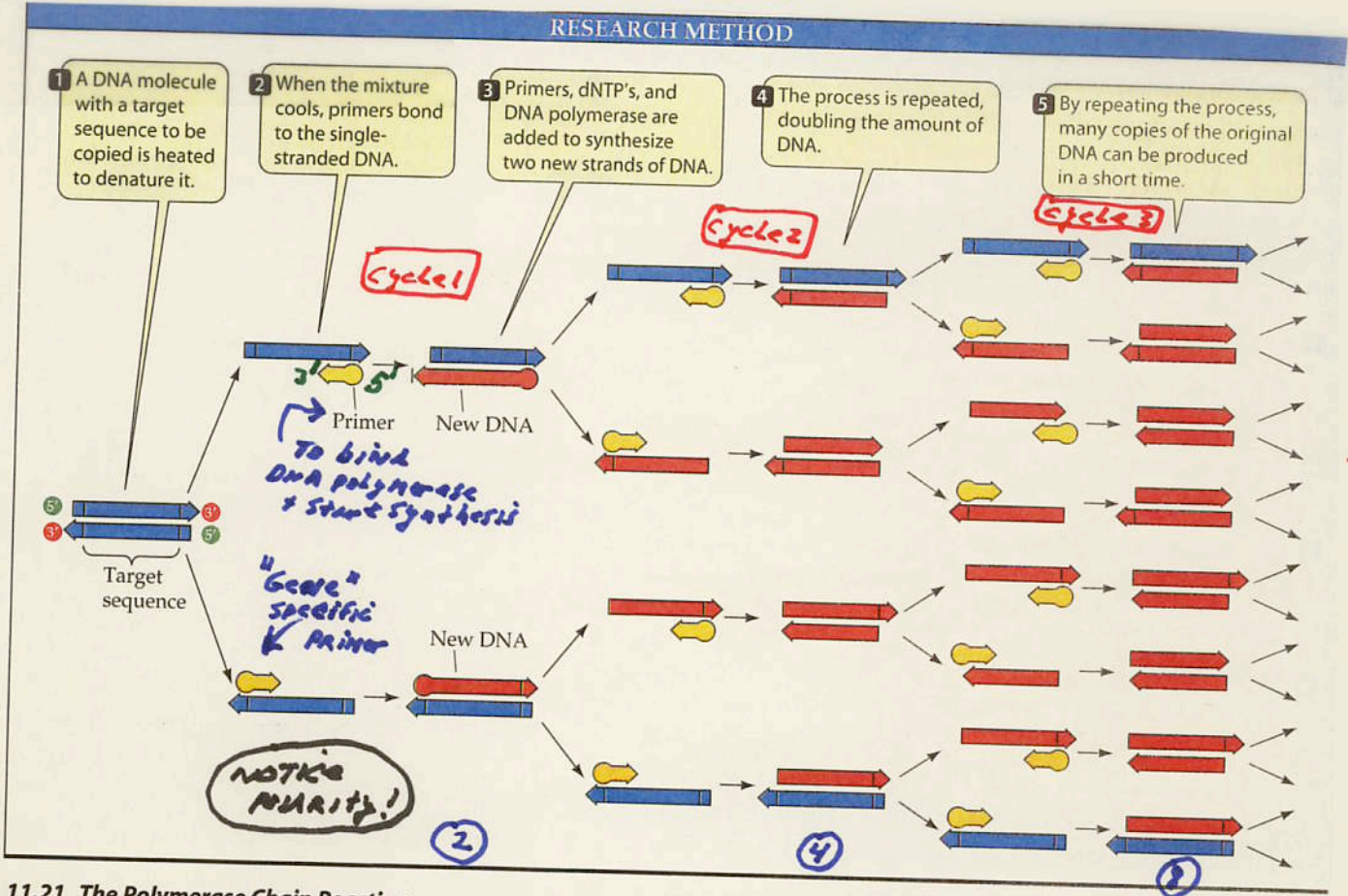
*CAN Make an ∞ amount of DNA!*

BUT - *Need sequence of DNA segment to be used for PCR! is have to clone a sequence first!*

REVOLUTIONIZED MANIPULATING DNA



# PCR IS A CYCLIC PROCESS OF DNA REPLICATION



## 11.21 The Polymerase Chain Reaction

The steps in this cyclic process are repeated many times to produce multiple copies of a DNA sequence.

$$2^D \text{ molecules of DNA}$$

where  $D = \# \text{ cycles}$

## PCR Requirements

- ① Knowledge of DNA Sequence :: Must clone DNA the "old fashioned way" first
- ② DNA Polymerase - Heat stable to  $\sim 100^{\circ}\text{C}$ !!  
Where isolated?
- ③ Thermo programmer / Cycler to Heat + Cool DNA in cycles - Separate strands + Allow new strands to form
- ④ Primers - Recognize specific DNA sequences + Initiate DNA synthesis + Binding of DNA polymerase Note! one not needed in test tube!

It's ALL in the DNA Sequences  
Know Sequence - can "make" an  
infinite amount of DNA sequence!

In 1 hour can do what took months before PCR!

# PCR IS A CYCLICAL PROCESS OF DNA SYNTHESIS

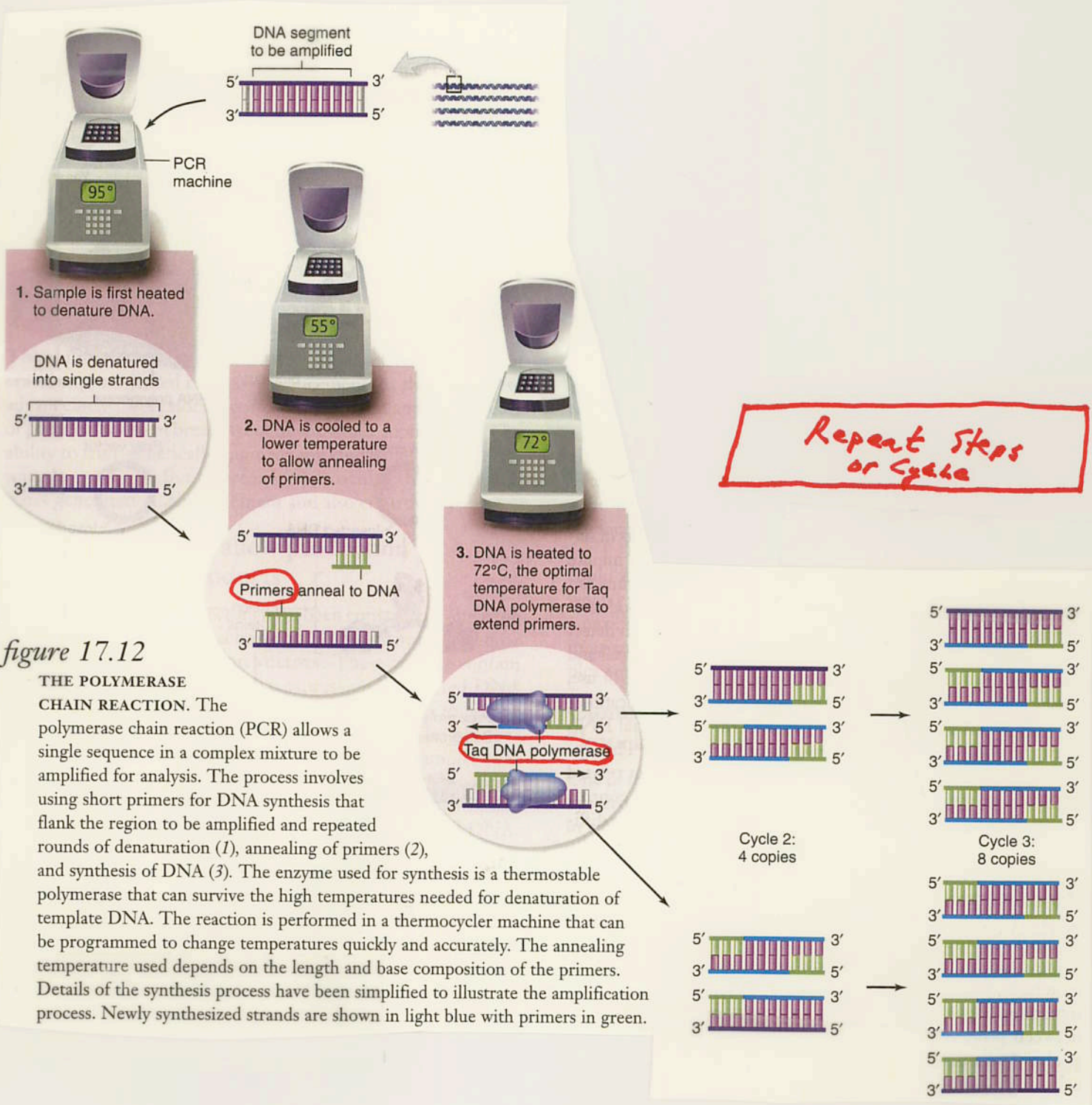


figure 17.12

**THE POLYMERASE CHAIN REACTION.** The polymerase chain reaction (PCR) allows a single sequence in a complex mixture to be amplified for analysis. The process involves using short primers for DNA synthesis that flank the region to be amplified and repeated rounds of denaturation (1), annealing of primers (2), and synthesis of DNA (3). The enzyme used for synthesis is a thermostable polymerase that can survive the high temperatures needed for denaturation of template DNA. The reaction is performed in a thermocycler machine that can be programmed to change temperatures quickly and accurately. The annealing temperature used depends on the length and base composition of the primers. Details of the synthesis process have been simplified to illustrate the amplification process. Newly synthesized strands are shown in light blue with primers in green.

Diagnostic for Amplified DNA Sequence

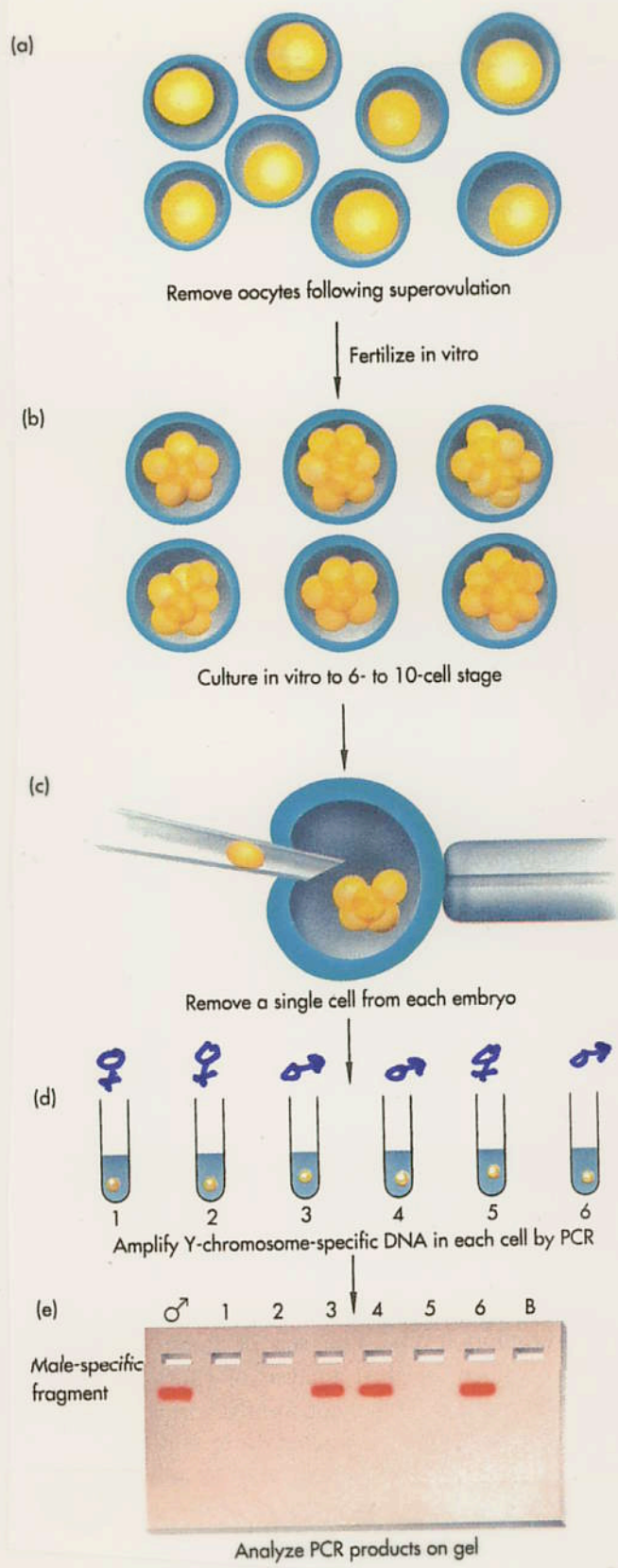
DNA FRAGMENTS ALL THE SAME SIZE  
Primer Sequence Primer

# PCR APPLICATIONS

- ① Amplify known DNA sequence for study  
+ / or Engineering (e.g., with another DNA)  
*Specific gene, specific switch, ori*
- ② Genetic diagnosis to determine genotype  
(e.g., disease genes)  
*PGD or pre-implantation genetic testing  
diagnosis*
- ③ Ancient DNA (e.g., sources where DNA present  
in small amounts + / or degraded)  
*bones, fossils (Neanderthal), mummies*
- ④ DNA forensics (i.e., DNA fingerprints)  
*crime scenes, mass graves, criminal suspects,  
innocent people wrongly convicted*
- ⑤ Identity testing (i.e., DNA fingerprints)  
*paternity, family relationships, immigration,  
individuals (tracing)*
- ⑥ Pathogen identity (e.g., medicine / agriculture)

PCR CAN BE USED TO ANALYZE GENES IN A SINGLE EMBRYO CELL

**PGD**  
pre-implantation  
genetic  
diagnosis



WHAT IS THE IMPLICATION OF THIS PROCEDURE CONSIDERING THAT THE HUMAN GENOME HAS BEEN SEQUENCED!

SEX DETERMINATION IN 8-CELL EMBRYOS!

Parents Should Be Allowed to Use PGD To Test  
Their Embryos For Any Gene and Select Those  
With the Gene Combination They Want To  
Become Their Child?

- a. Yes
- b. No

**Parents Should Be Allowed to Use PGD To Test  
Their Embryos For Any Gene and Select the  
"Best" One To Become Their Child?**

- a. Yes**
- b. No**

A STEVEN SPIELBERG FILM



An Adventure  
65 Million Years In The Making.



USING PCR TO DETECT GENES IN ANCIENT DNA

Ancient DNA Milestones

These extinct organisms have yielded meaningful genetic sequences.



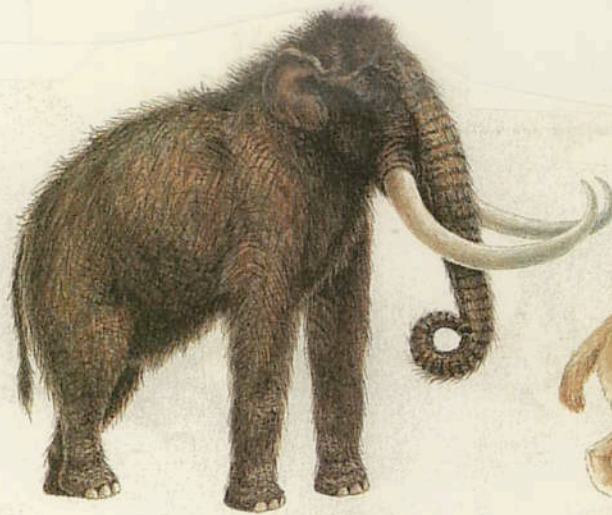
AMBER INSECTS

40 MILLION YEARS OLD



FOSSIL LEAVES

17 MILLION



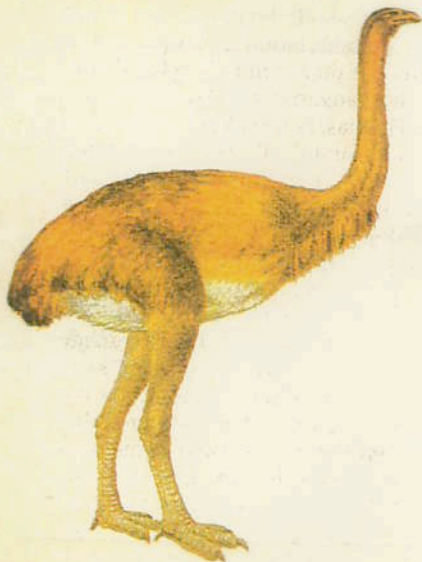
MAMMOTH

40,000



SMILODON

13,000



MOA

4,300



QUAGGA

140



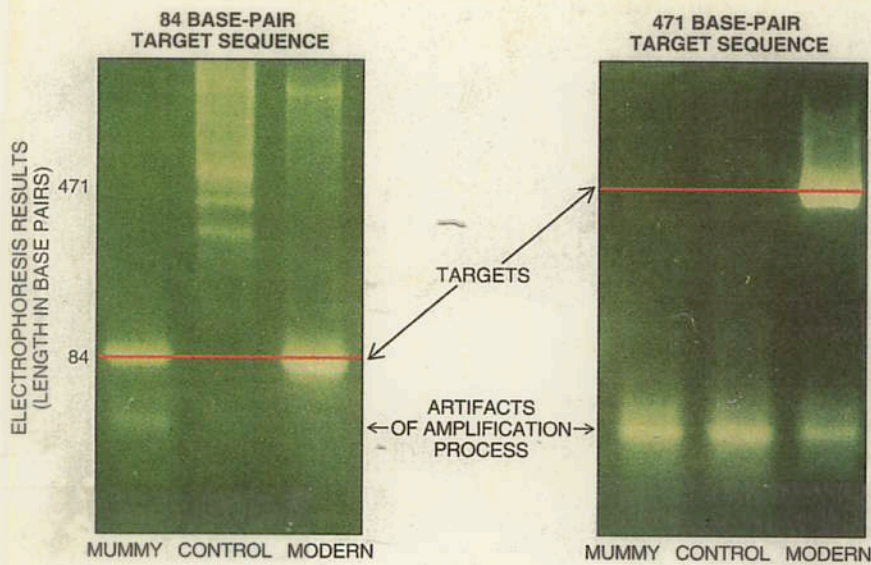
THYLACINE

80

PRESENT

→ JUST NEED ONE MOLECULE OF DNA ←

Using PCR TO DETECT GENES  
IN MUMMY DNA



Using PCR TO AMPLIFY NEANDERTHAL BONE  
DNA + SEQUENCE THE ENTIRE GENOME!

Neanderthal DNA vs. Homo Sapien (Human)  
DNA

SEQUENCE TO DETERMINE  
RELATIONSHIPS

# USING PCR IN CRIME SCENES

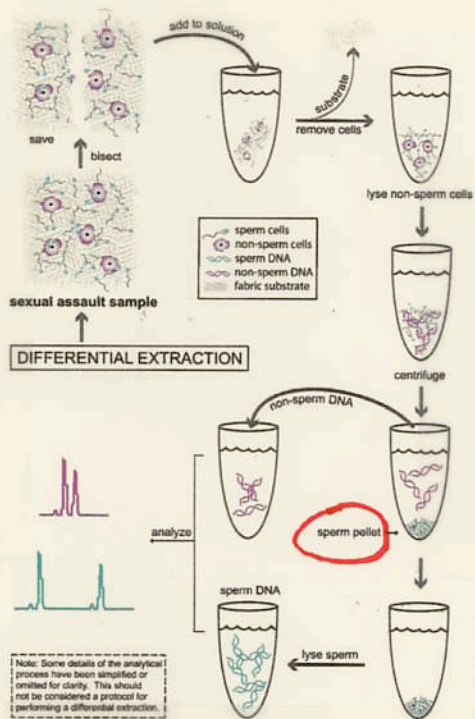
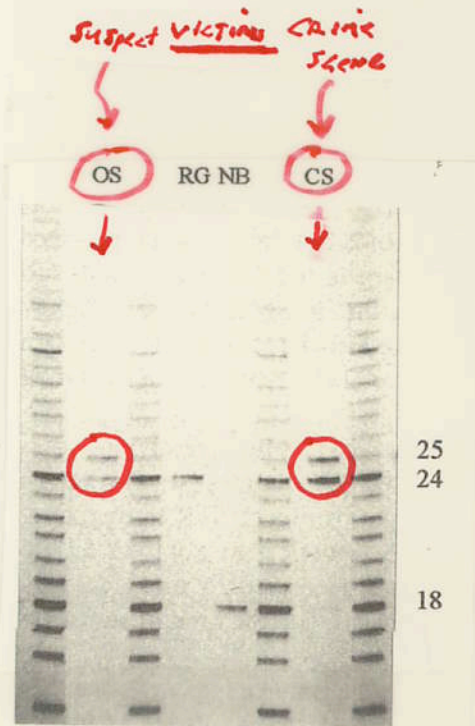


Plate 4 Differential extraction.

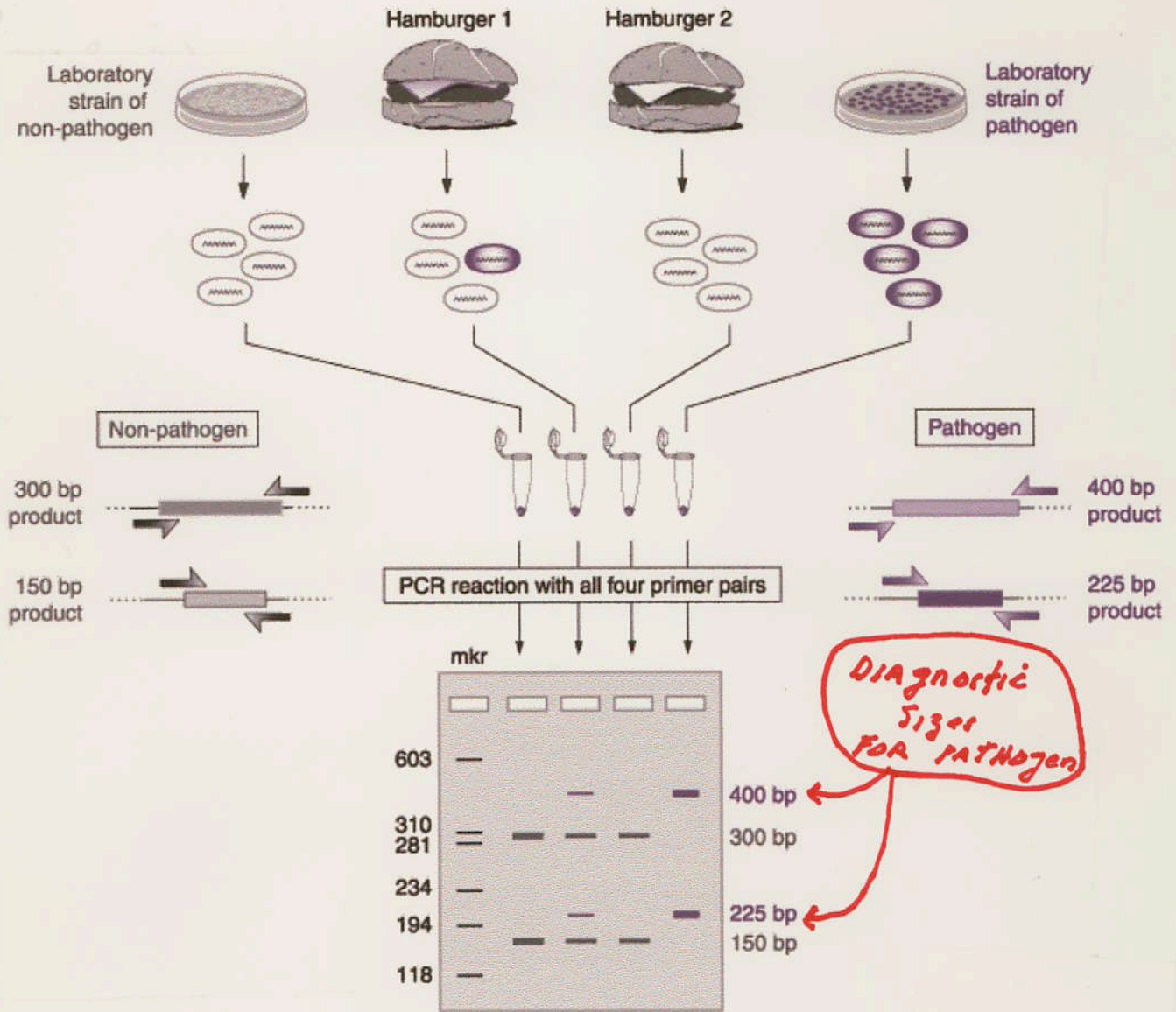


"Match"  
 What is probability  
 that this  
 will  
 occur by  
 chance?

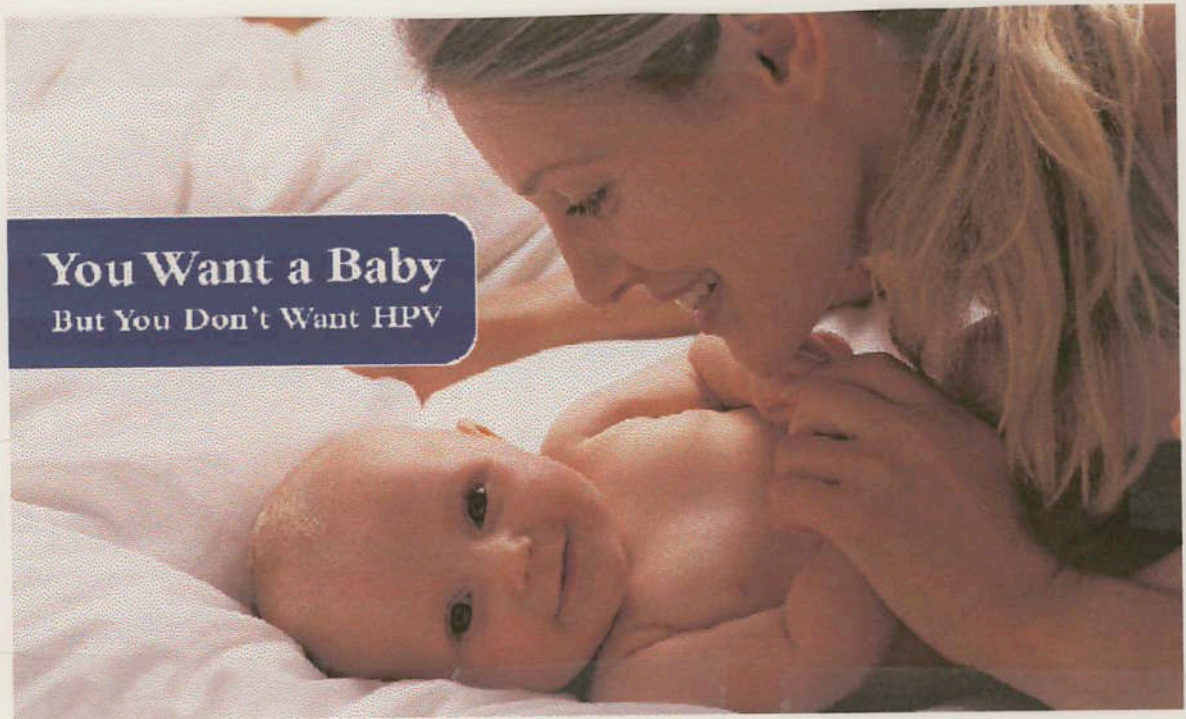
OS = Suspect  
 CS = CRIME SCENE  
 RG + NB = VICTIMS

DNA doesn't "lie"!!

# USING PCR TO DETECT FOOD PATHOGENS



USING PCR TO TEST FOR  
HUMAN PATHOGENS

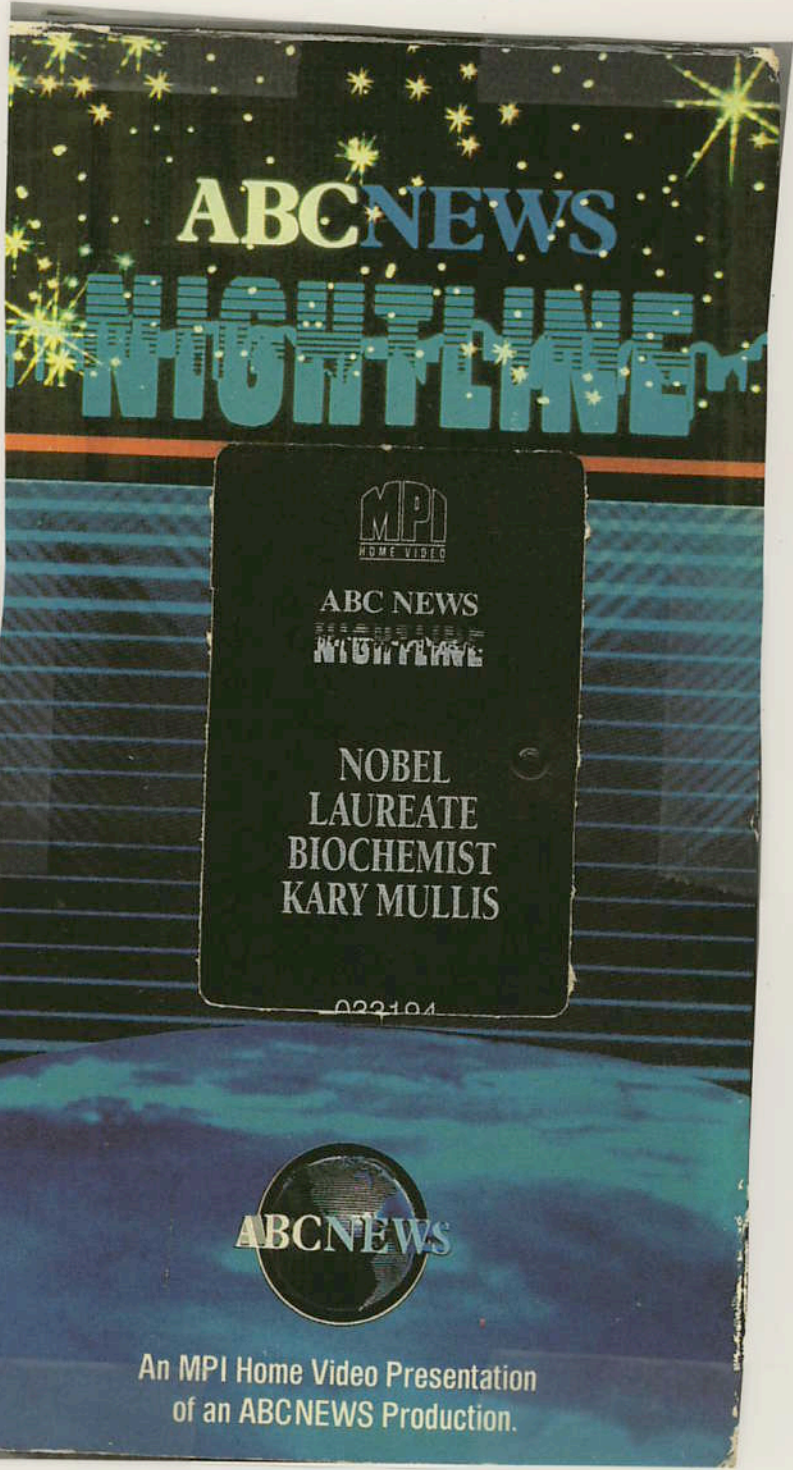


Identify Specific Pathogenic  
Bacteria, Viruses, Fungi

Each Genome Has Specific  
DNA Sequences + therefore  
DIAGNOSTIC DNA FRAGMENTS

**The Polymerase Chain Reaction Has Made DNA Cloning and Recombinant DNA Obsolete?**

- a. Yes**
- b. No**



ABC NEWS

NIGHTLINE



ABC NEWS  
NIGHTLINE

NOBEL  
LAUREATE  
BIOCHEMIST  
KARY MULLIS

022104



An MPI Home Video Presentation  
of an ABC NEWS Production.

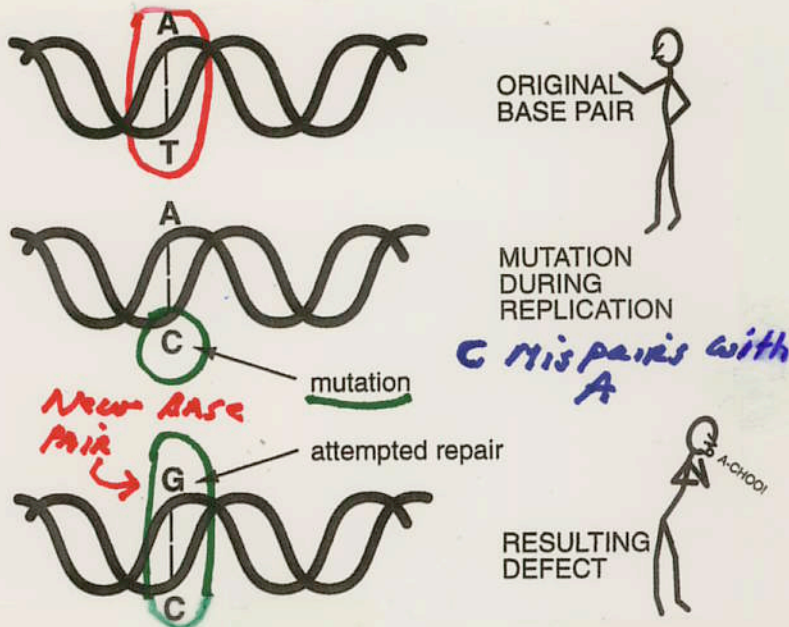
DNA REPLICATION IS PRECISE  
BUT MISTAKES OR **MUTATIONS**  
CAN OCCUR!

GENE A

Replication ①

Replication ②

GENE A'  
ALLELIC  
VARIANT



RARE  
BASE  
MISMATCH

SEE  
MUTATION  
AS  
CHANGE  
in  
Phenotype

FIGURE 1.3 A mutated gene makes a defective protein that leads to an illness in an individual.

CHANGE DNA SEQUENCE FROM  
A=T TO G=C

∴ CHANGE PROTEIN AMINO ACID  
SEQUENCE → Alter Function!



Big TOMATO to  
SMALL TOMATO



MUTATIONS IN GENES ARE RARE BUT ARE INHERITED

**ONE**  
gene per  
gamete  
 $\sigma + \text{♀}$

**TWO**  
genes per  
somatic  
cells

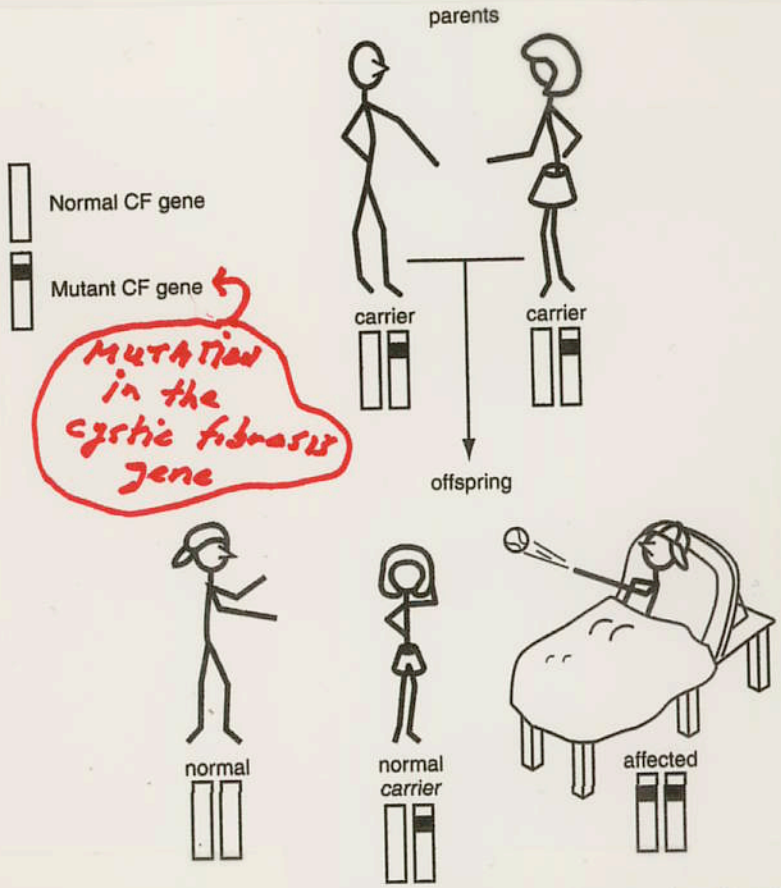


FIGURE 1.4 Two parents heterozygous for CF and producing normal or affected kids.

HOW FOLLOW  
INHERITANCE?  
WHAT ALLOW DISEASE  
TO BE FOLLOWED?

**MARKER!**

**Alleles Are Different Forms of the Same Gene?**

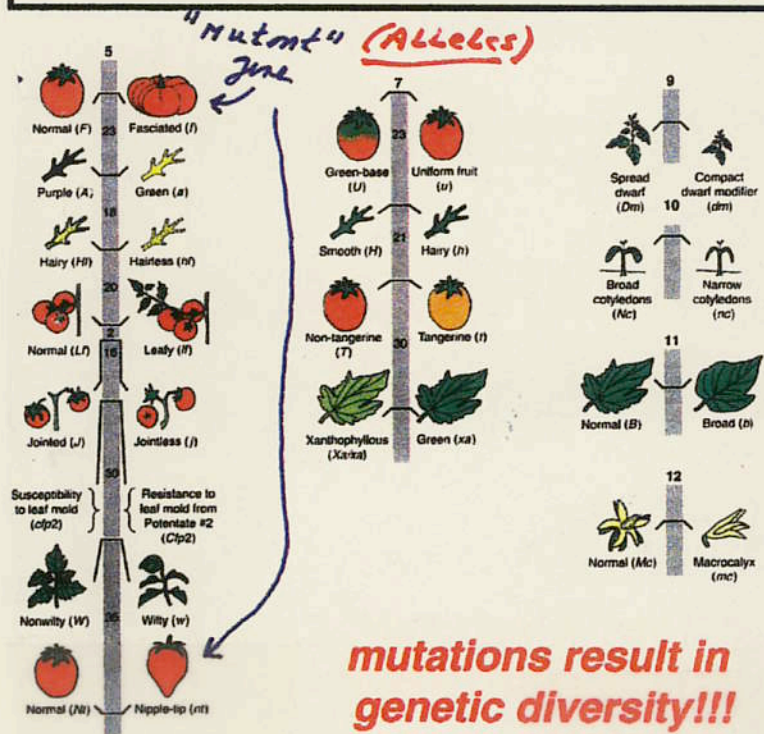
- a. Yes**
- b. No**

**Alleles Are Generated By Mutations and Lead to Genetic Diversity in a Population?**

- a. Yes**
- b. No**

MUTATIONS GIVE RISE TO GENETIC DIVERSITY

Alternative Forms of the Same Gene Lead to Genetic Diversity



"Mutant" (ALLELES)  
gene

Change in DNA Sequence!

How know Sequence changed?

Different forms of SAME Gene

SNP - Single nucleotide polymorphism!







mutations result in genetic diversity!!!

VARIABILITY Acted on & used by "Nature" & by our Ancestors / Early Gene Engineers!

SAME PROCESS → Diversity of HUMAN Genes!  
How know MUTATION? ARE MUTATIONS Good, Bad, Neutral?

# MUTATIONS CAN OCCUR DIFFERENT WAYS

Table 18.1 Types of Mutation

Mutation	Example result
<b>NO MUTATION</b>	
	Normal B protein is produced by the B gene.
<b>POINT MUTATION</b>	
<b>Base substitution</b> Substitution of one or a few bases 	B protein is inactive because changed amino acid disrupts function.
<b>Insertion</b> Addition of one or a few bases 	B protein is inactive because inserted material disrupts proper shape.
<b>Deletion</b> Loss of one or a few bases 	B protein is inactive because portion of protein is missing.
<b>CHANGES IN GENE POSITION</b>	
<b>Transposition</b> 	B gene or B protein may be regulated differently because of change in gene position.
<b>Chromosomal rearrangement</b> 	B gene may be inactivated or regulated differently in its new location on chromosome.

① BASE-PAIR CHANGE

② Add/delete BASE PAIRS

BASE Sequence of Gene Change!

③ Move Gene or PART of Gene to NEW Location!

Switches Change!

LOSS OF FUNCTIONAL PROTEIN

MUTATIONS CAN OCCUR BY  
 CHANGING, ADDING TO, OR DELETING  
 SPECIFIC NUCLEOTIDES

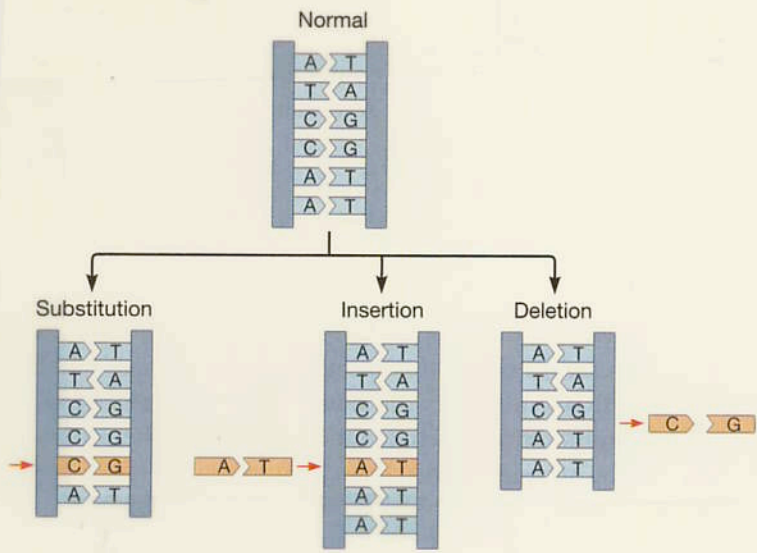


FIGURE 2-10

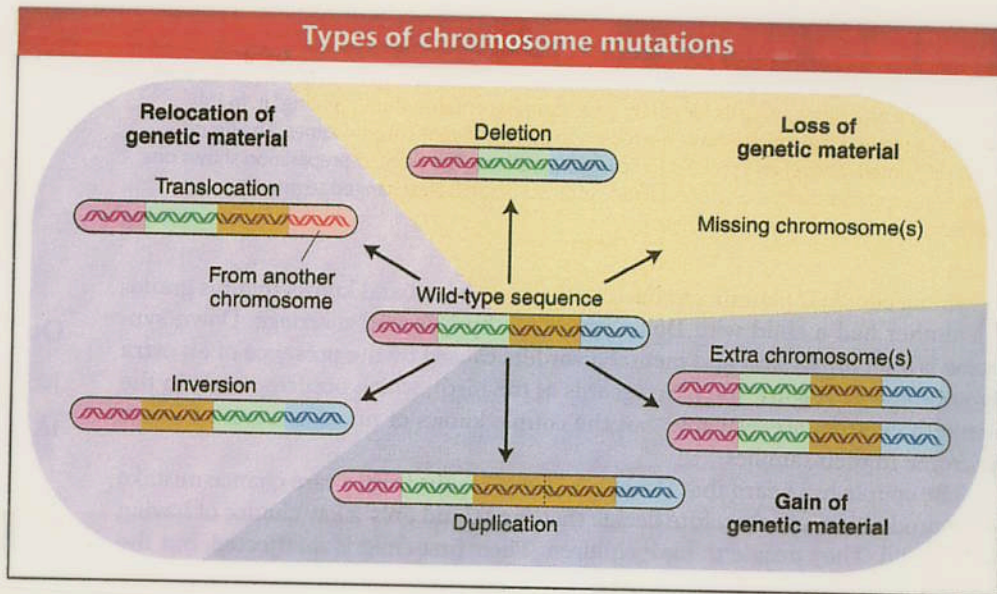
The three types of mutation. They are substitution, addition, and deletion of a base pair in a DNA strand.

CHANGE

ADD

DELETE

"MUTATIONS" CAN ALSO OCCUR  
BY LARGE CHROMOSOMAL  
CHANGES

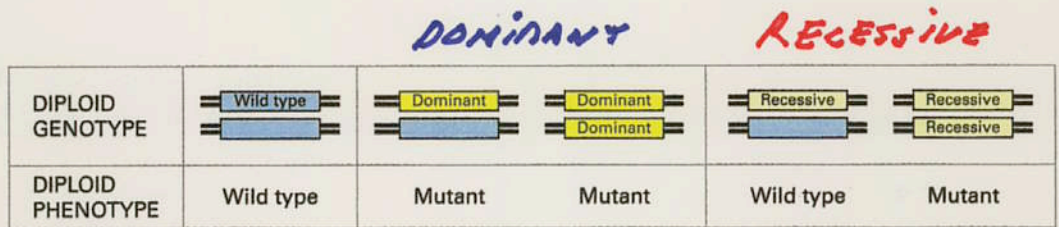


These changes affect MANY Genes!

e.g. Down's Syndrome (3 Chromosome 21s)

# HUMAN GENETIC DISORDERS OCCUR AS A RESULT OF MUTATIONS

Disorder	Symptom	Defect	Dominant/ Recessive	Frequency Among Human Births
Cystic fibrosis	Mucus clogs lungs, liver, and pancreas	Failure of chloride ion transport mechanism	Recessive	1/2500 (Caucasians)
Sickle cell anemia	Blood circulation is poor	Abnormal hemoglobin molecules	Recessive	1/600 (African Americans)
Tay-Sachs disease	Central nervous system deteriorates in infancy	Defective enzyme (hexosaminidase A)	Recessive	1/3500 (Ashkenazi Jews)
Phenylketonuria	Brain fails to develop in infancy	Defective enzyme (phenylalanine hydroxylase)	Recessive	1/12,000
Hemophilia	Blood fails to clot	Defective blood-clotting factor VIII	X-linked recessive	1/10,000 (Caucasian males)
Huntington disease	Brain tissue gradually deteriorates in middle age	Production of an inhibitor of brain cell metabolism	Dominant	1/24,000
Muscular dystrophy (Duchenne)	Muscles waste away	Degradation of myelin coating of nerves stimulating muscles	X-linked recessive	1/3700 (males)
Hypercholesterolemia	Excessive cholesterol levels in blood lead to heart disease	Abnormal form of cholesterol cell surface receptor	Dominant	1/500

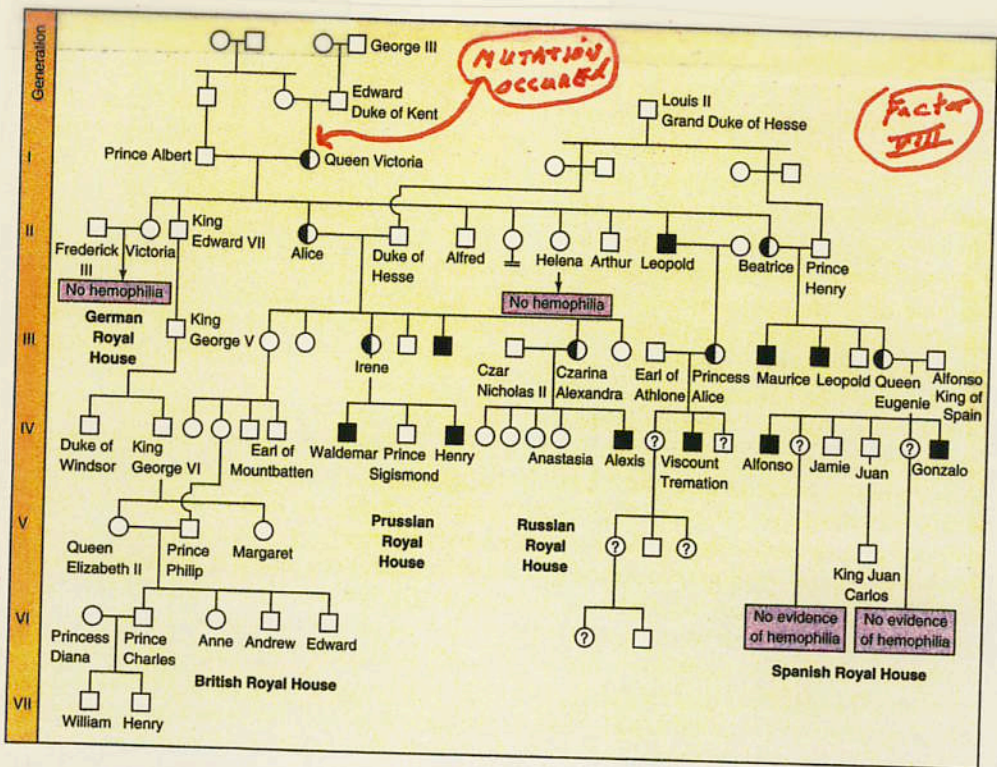


**▲ FIGURE 5-2 Effects of dominant and recessive mutant alleles on phenotype in diploid organisms.** A single copy of a dominant allele is sufficient to produce a mutant phenotype, whereas both copies

of a recessive allele must be present to cause a mutant phenotype. Recessive mutations usually cause a loss of function; dominant mutations usually cause a gain of function or an altered function.



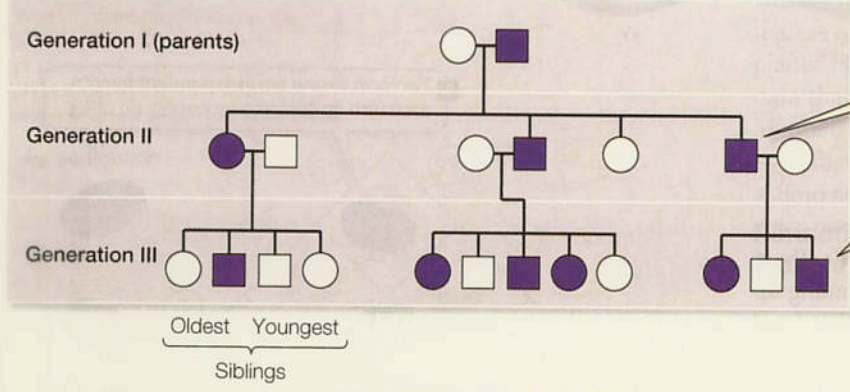
PEDIGREES CAN BE USED TO FOLLOW DISEASE GENES IN HUMAN FAMILIES



**FIGURE 13.26**  
**The Royal hemophilia pedigree.** Queen Victoria's daughter Alice introduced hemophilia into the Russian and Austrian royal houses, and Victoria's daughter Beatrice introduced it into the Spanish royal house. Victoria's son Leopold, himself a victim, also transmitted the disorder in a third line of descent. Half-shaded symbols represent carriers with one normal allele and one defective allele; fully shaded symbols represent affected individuals.

**PEDIGREES CAN BE USED TO DETERMINE IF TRAIT IS DOMINANT OR RECESSIVE**

(A) Dominant inheritance

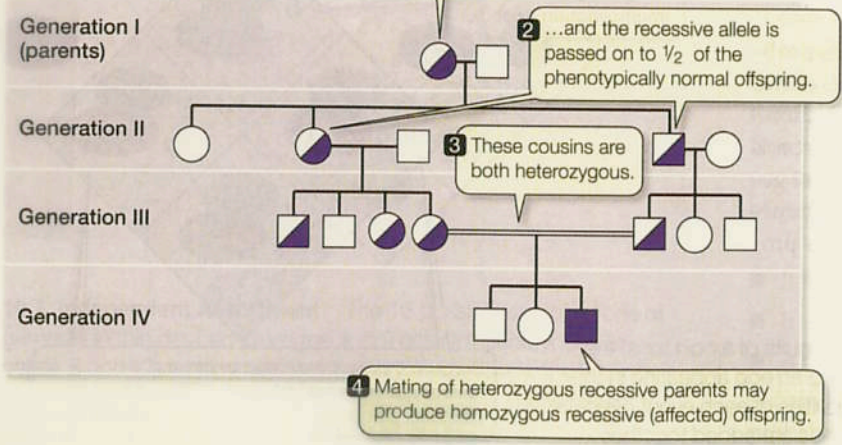


Every affected individual has an affected parent.

*MUSCULAR DYSTROPHY*  
*Huntington disease*

About 1/2 of the offspring (of both sexes) of an affected parent are affected.

(B) Recessive inheritance

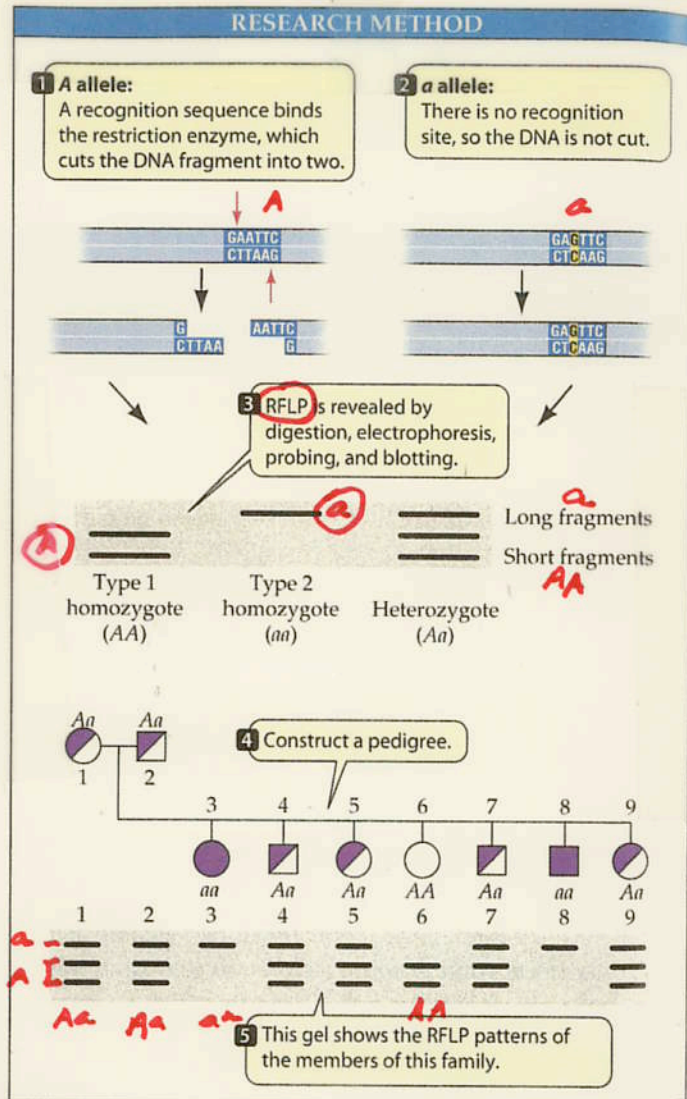


*cystic fibrosis*  
*hemophilia*  
*Tay-Sachs*

	Unaffected	Affected	Heterozygote (unaffected phenotype)
Female	○	●	◐
Male	□	■	◑
Mating	○—□	●—■	◐—◑
Mating between relatives	○—○	■—■	◐—◐

**EACH TYPE OF INHERITANCE PREDICTS SPECIFIC RESULTS IN EACH GENERATION**

OR FOLLOW BY DNA TESTS USING MOLECULAR METHODS (e.g., PCR)



CUT - A

a - NOT CUT

USE PCR + for STR on Methods

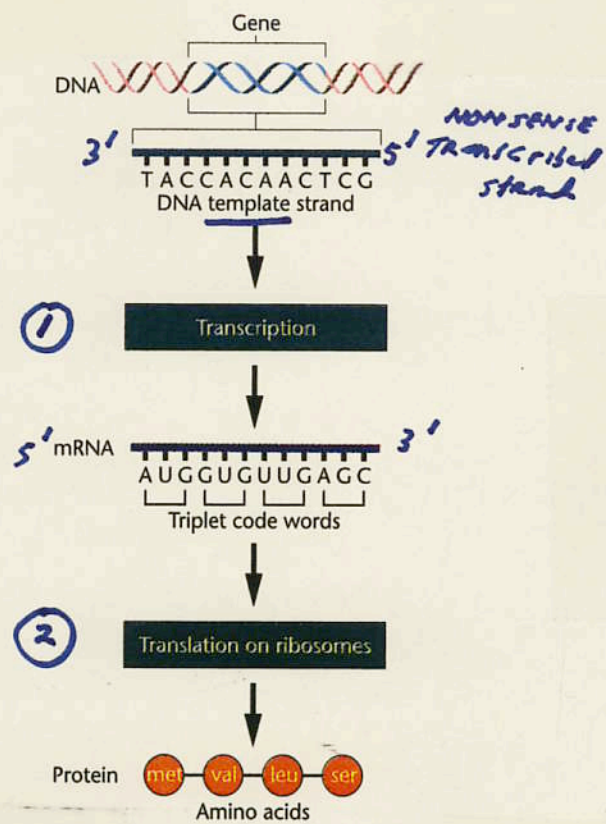
DNA Fingerprint  
RFLP

Follow in Family

**17.7 RFLP Mapping** Restriction fragment length polymorphisms are differences in DNA sequences that serve as genetic markers. Thousands of such markers have been described for the human genome.

Implications? Combined with Sequence of Human Genome, Embryo testing (PGD), + PCR ?!!!

**HOW DOES A GENE LEAD TO A PHENOTYPE?**



**FIGURE 13.1** An overview of the concept of the flow of genetic information encoded in DNA to messenger RNA to protein.

① mRNA Synthesized by Transcription

Complementary to Transcribed, Nonsense strand

Same sequence as sense strand

② mRNA Translated into Protein by Translation of the Genetic Code

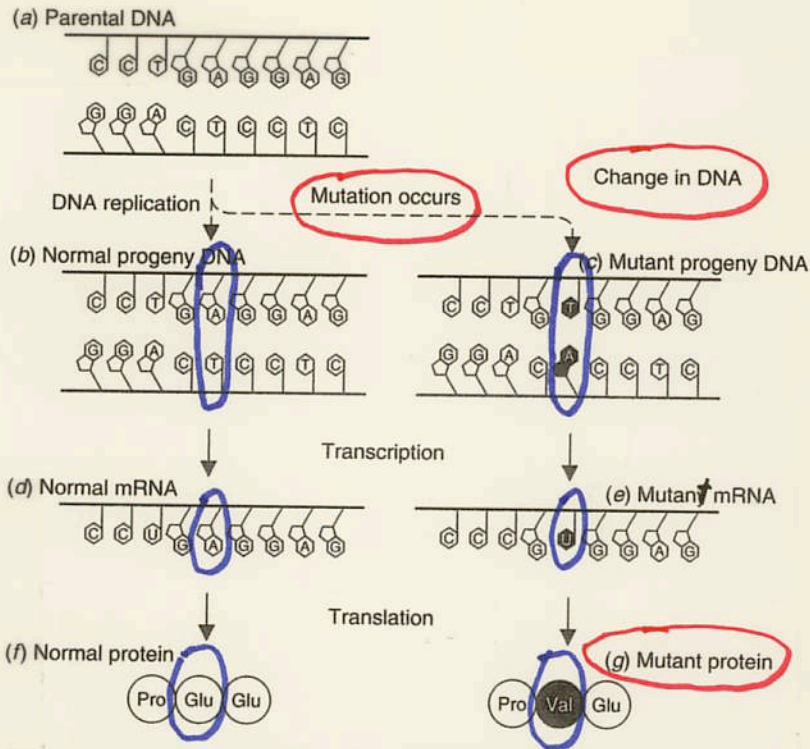
Genetic Code on mRNA Translated to Protein in sequence

∴ Sequence of Gene  
 ↳ Sequence of mRNA  
 ↳ Sequence of protein

KNOW SEQUENCE  
 KNOW PROTEIN

↳ ENGINEER NEW PROTEINS

**MUTATIONS - CHANGE DNA SEQUENCE  
 ↓ CHANGE CORRESPONDING PROTEIN  
 SEQUENCE !!**



Change in  
DNA  
↓  
Change in  
mRNA  
↓  
Change in  
Protein  
↓  
Change in  
Phenotype

**Figure 3-5. A Point Mutation Changes the Sequence of Amino Acids in a Protein.** DNA replication is very accurate, so the nucleotide sequence in the progeny DNA (b) is usually identical to that of normal parental DNA (a). Occasionally an error is made. In this example, a particular A · T base pair in parental DNA changes to a T · A pair in the mutant, progeny DNA (c). During transcription, the information in DNA is converted into messenger RNA. The mutation in DNA results in a conversion of particular GAG codon in normal messenger RNA (d) into a GUG codon in mutant messenger RNA (e). During translation of the information into protein, GAG codes for the amino acid glutamic acid (Glu) (f), while GUG codes for valine (Val) (g) (see Figure 2-6). The two amino acids have very different chemical properties. Since the structure of the resulting protein is determined by the precise order of the amino acids, the mutant protein will differ significantly from the normal protein. The differences between the normal and mutant molecules shown are identical to those found between healthy people and patients suffering from sickle-cell disease.

# GENE MUTATION LEADING TO SICKLE CELL ANEMIA

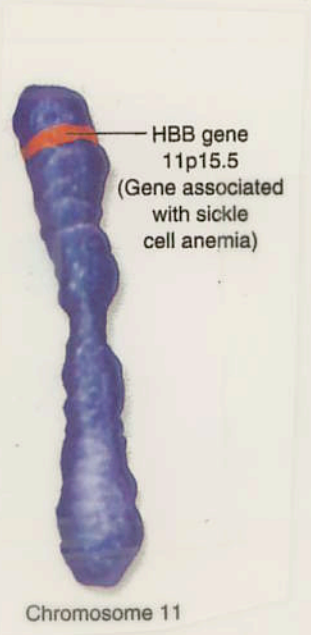
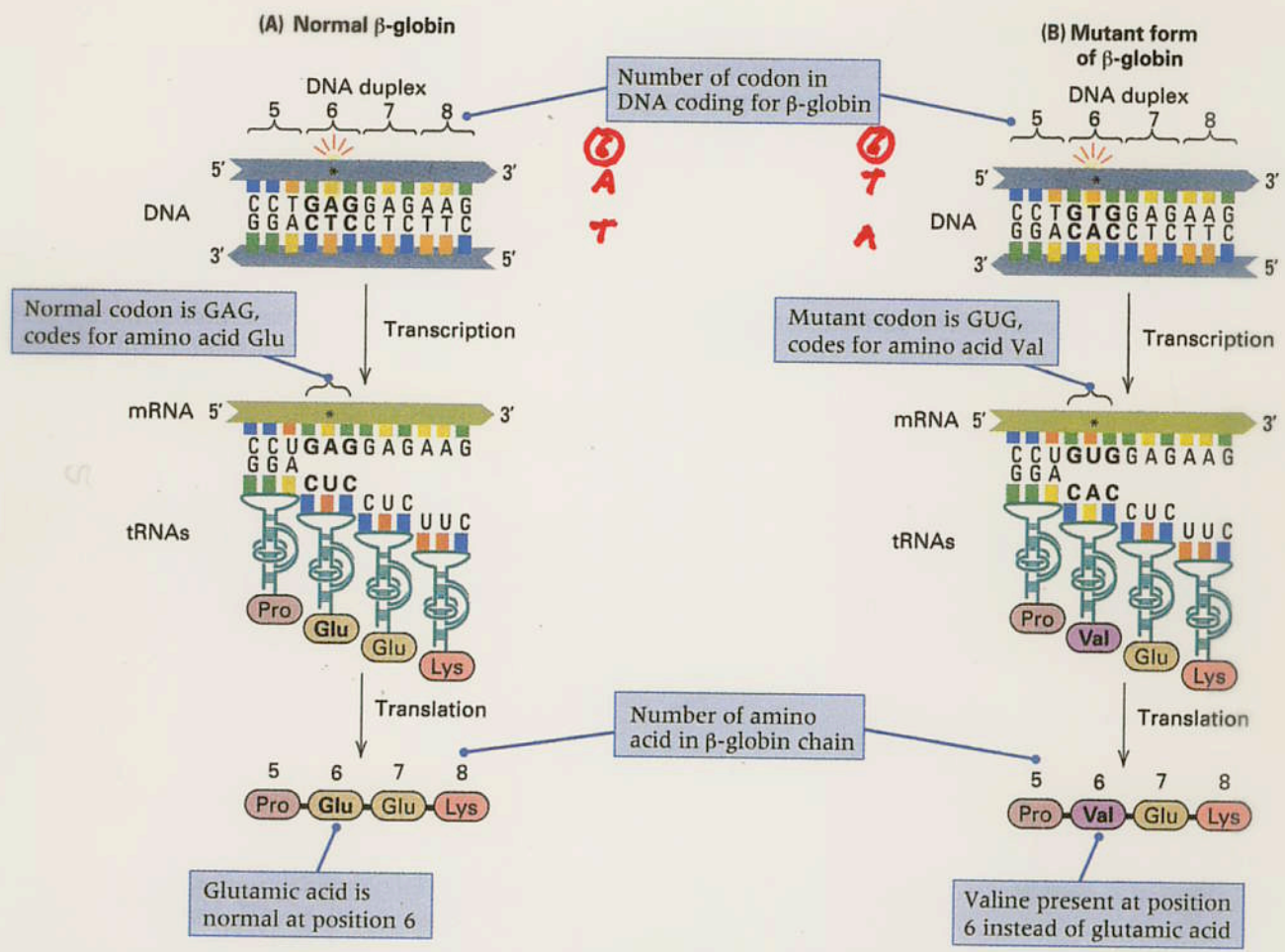


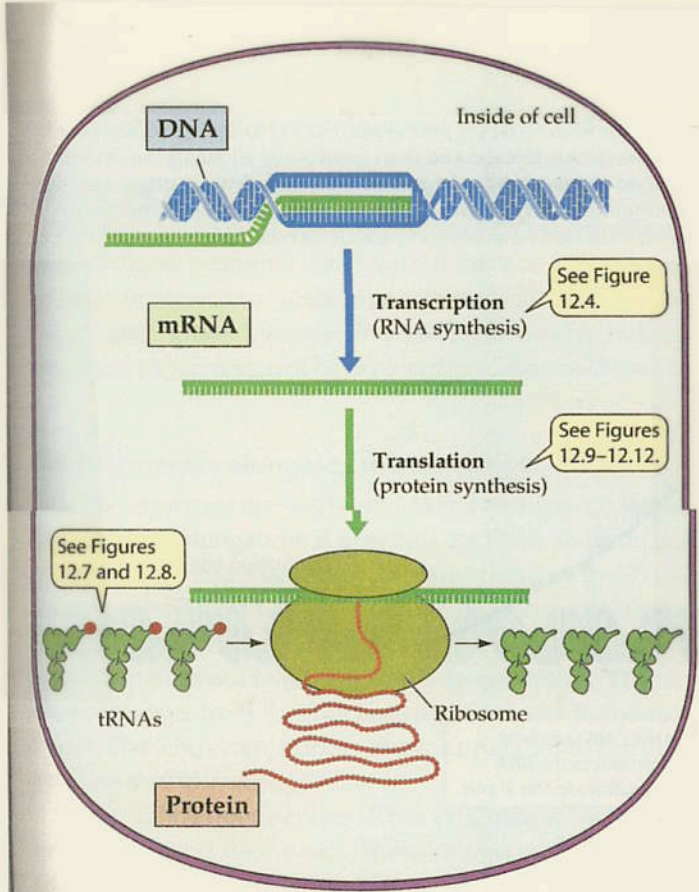
figure 13.11

SICKLE CELL ANEMIA. In individuals homozygous for the sickle cell trait, many of the red blood cells have sickled or irregular shapes, such as the cell on the far right.



**Figure 7.3** Molecular basis of sickle-cell anemia. (A) Part of the DNA in the normal β-globin gene is transcribed into a messenger RNA coding for the amino acid sequence Pro-Glu-Glu-Lys. The T in the marked A-T base pair is transcribed as the A in the GAG codon for Glu (glutamic acid). (B) Mutation of the normal A-T base pair to a T-A base pair results in the codon GUG instead of GAG. The codon GUG codes for Val (valine), so the polypeptide sequence in this part of the molecule is Pro-Val-Glu-Lys. The resulting hemoglobin is defective and tends to polymerize at low oxygen concentration.

AN ELABORATE CELLULAR MACHINERY  
 REQUIRING THOUSANDS OF GENES  
 IS REQUIRED TO PRODUCE  
 PROTEINS ENCODED BY SPECIFIC  
 GENES!



IT TAKES GENES  
 TO EXPRESS  
 (one replicate)  
 A GENE!!

12.3 From Gene to Protein This diagram summarizes the processes of gene expression in prokaryotes. In eukaryotes, the processes are somewhat more complex.

GENETIC CODE ALLOWS THE SEQUENCE OF NUCLEOTIDES in mRNA / sense strand of Gene to be translated into sequence of amino acids in proteins

mRNA  
protein

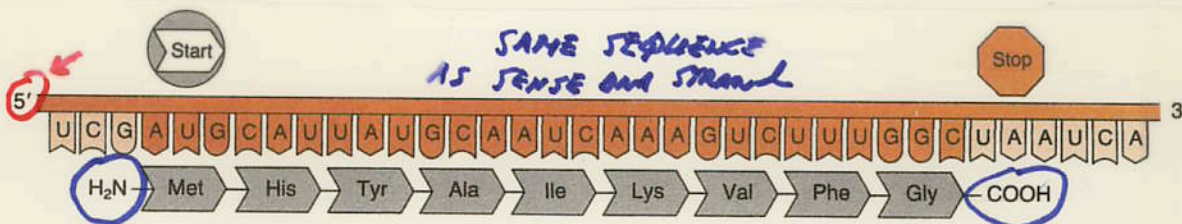


Figure 3.4 Decoding a messenger RNA sequence into a polypeptide.

NOTE: SEQUENCE in mRNA (= sense Gene strand) is TRANSLATED 5' → 3' (= beginning of sense strand to end) \* protein made in N → C direction ∴ order nts in gene = order aa in protein!



The Genetic Code is Universal!

How Know?

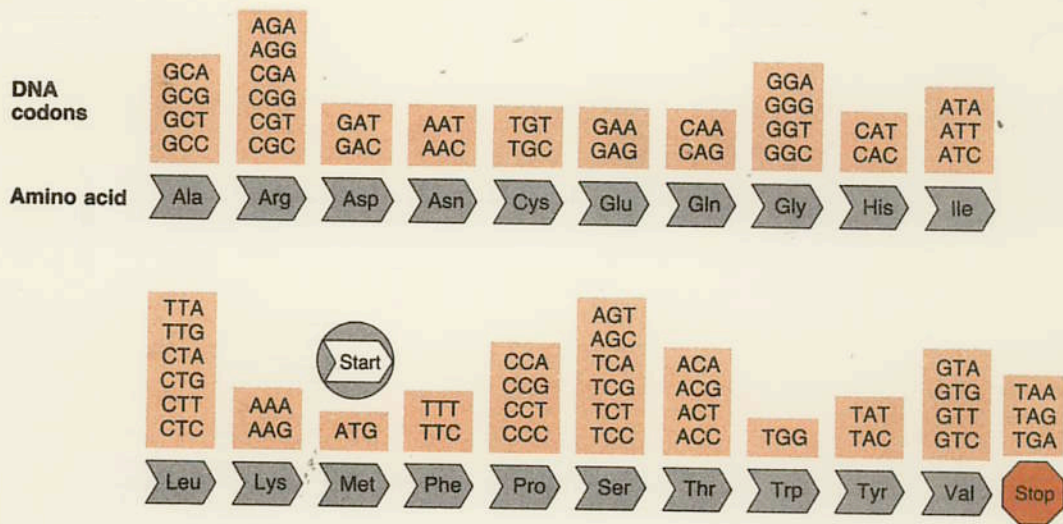


Figure 3.3 The genetic code. The codons shown for each amino acid are those for DNA. For RNA, the Ts are replaced by Us.

- ① Universal
- ② Triplet
- ③ Punctuation
- ④ Degenerate

KNOW SEQUENCE OF GENE - KNOW SEQUENCE OF PROTEIN USING GENETIC CODE

Big Implication For Genetic Engineering! CAN MAKE GENES, GENOMES & SPECIFY PROTEINS WANTED! CAN EXPRESS GENES FROM ONE ORGANISM IN ANOTHER!

DESIGN AN EXPERIMENT TO SHOW UNIVERSAL!

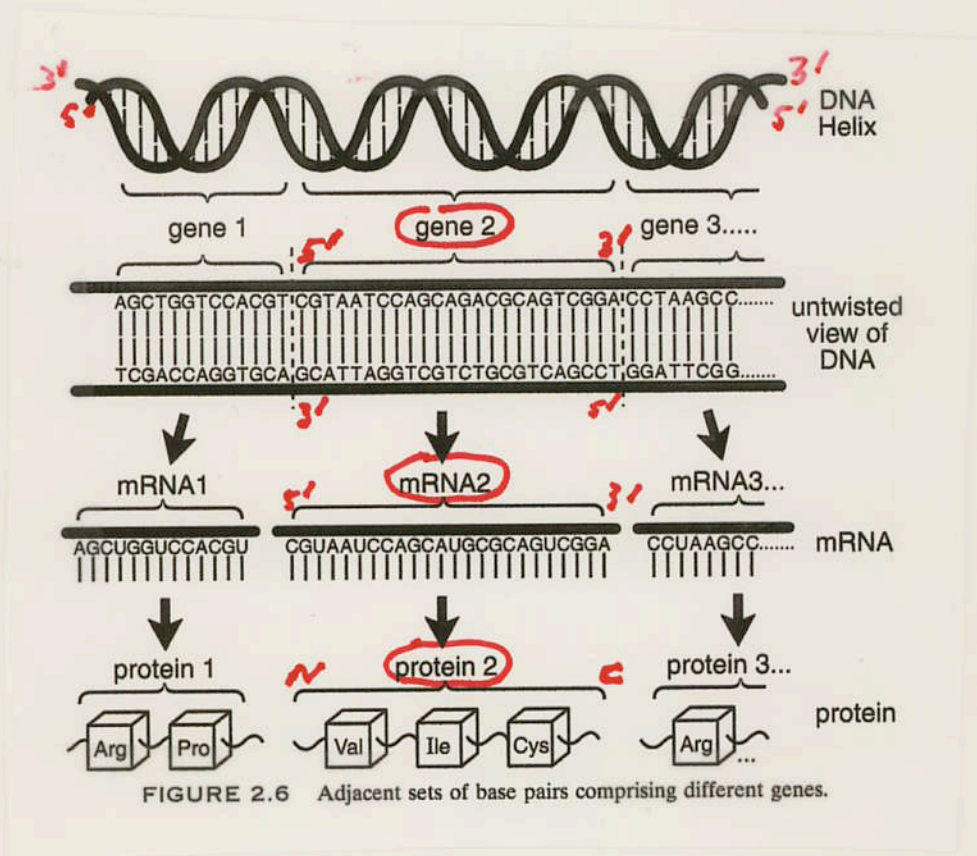
EXPRESSION OF JELLYFISH GREEN  
FLUORESCENCE PROTEIN (GFP)  
IN PIGS SHOWS THAT  
GENETIC CODE IS UNIVERSAL!



*figure 15.3*

TRANSGENIC PIG. The piglet on the right is a conventional piglet. The piglet on the left was engineered to express a gene from jellyfish that encodes green fluorescent protein. The color of this piglet's nose is due to expression of this introduced gene. Such transgenic animals indicate the universal nature of the genetic code.

THERE IS A COLINEARITY BETWEEN THE DNA SEQUENCE OF A GENE & THE AMINO ACID SEQUENCE OF A PROTEIN



FUNCTION 2  
 ↳ SPECIFIC TRAITS

GENES FUNCTION AS INDIVIDUAL UNITS!

PROTEINS ARE MADE OF AMINO ACIDS AND.....

AMINO ACIDS ARE JOINED BY PEPTIDE BONDS

ORDER!

Corresponds to 5' & 3' ends of mRNA/sense strand of gene!

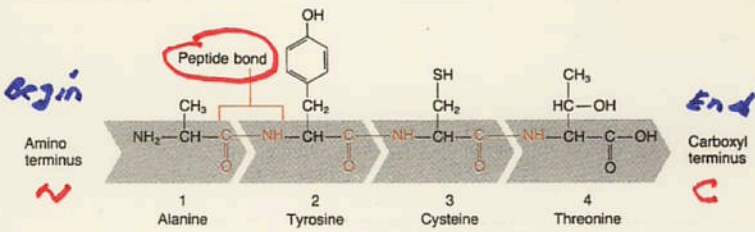
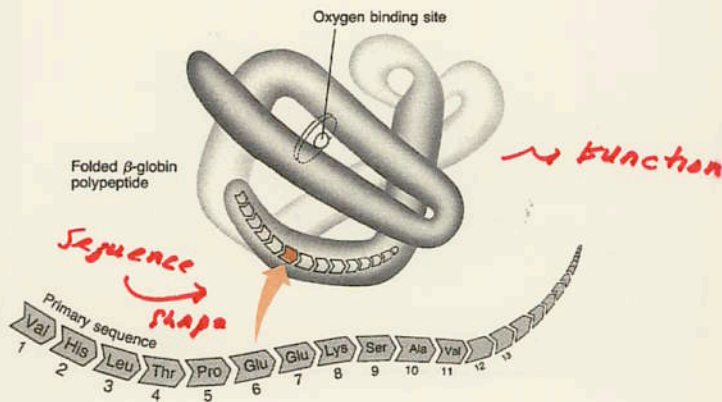


Figure 2.16 Peptide bonds between amino acids in a tetrapeptide (four amino acids).

Mutations!

Change Shape  
Change Function  
Change Phenotype



In sickle-cell hemoglobin, the Glu at position 6 is replaced by Val

Figure 2.17 A portion of the primary structure of the  $\beta$ -globin polypeptide and its location in the folded, complete polypeptide. Also shown is the amino acid that is altered in the  $\beta$ -globin polypeptide in sickle-cell disease.

CHEMISTRY  
↳ BIOLOGY!

ORDER OF AMINO ACIDS → Specific Protein Shape & Function

SHAPE → FUNCTION → Phenotype → Specific

UNIQUE PROTEINS HAVE A UNIQUE COMPOSITION & ORDER OF AMINO ACIDS & HAVE UNIQUE SIZES, SHAPES, & FUNCTIONS

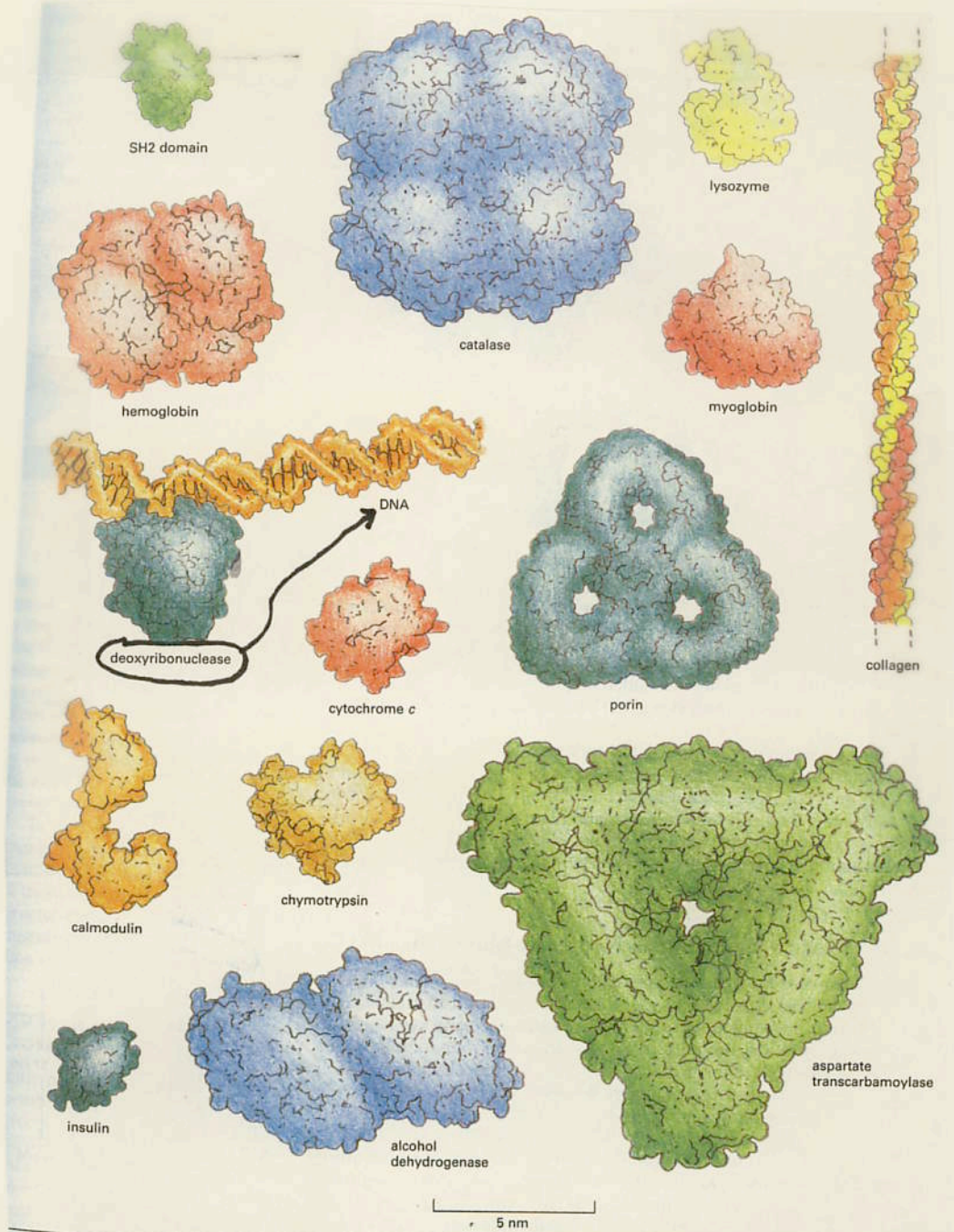
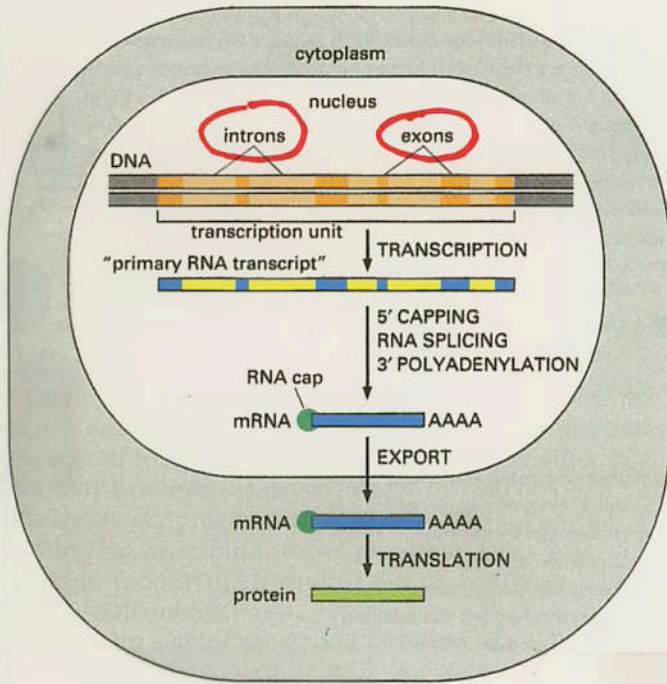


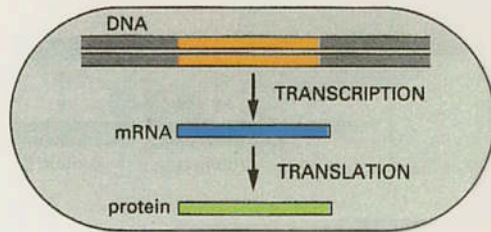
Figure 3-24 A collection of protein molecules, shown at the same scale. For comparison, a DNA molecule bound to a protein is also illustrated. These space-filling models represent a range of sizes and shapes. Hemoglobin, catalase, porin, alcohol dehydrogenase, and aspartate transcarbamoylase are formed from multiple copies of subunits. The SH2 domain (top left) is presented in detail in Panel 3-2 (pp. 138-139). (After David S. Goodsell, Our Molecular Nature. New York: Springer-Verlag, 1996.)

# EUKARYOTIC AND PROKARYOTIC GENE EXPRESSION PROCESSES DIFFER SLIGHTLY

(A) EUKARYOTES

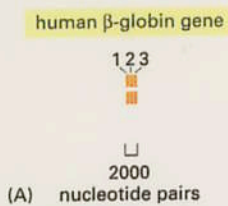


(B) PROCARYOTES

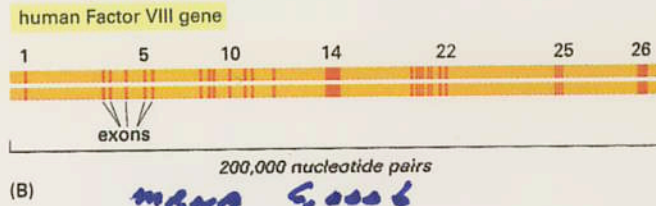


*Genes differ  
Switches differ  
Genetic code the same  
General processes same  
Eukaryotic genes have  
introns - noncoding regions  
in gene!*

## GENES CAN VARY IN SIZE!



*mRNA 600b*



*mRNA 6,000b*

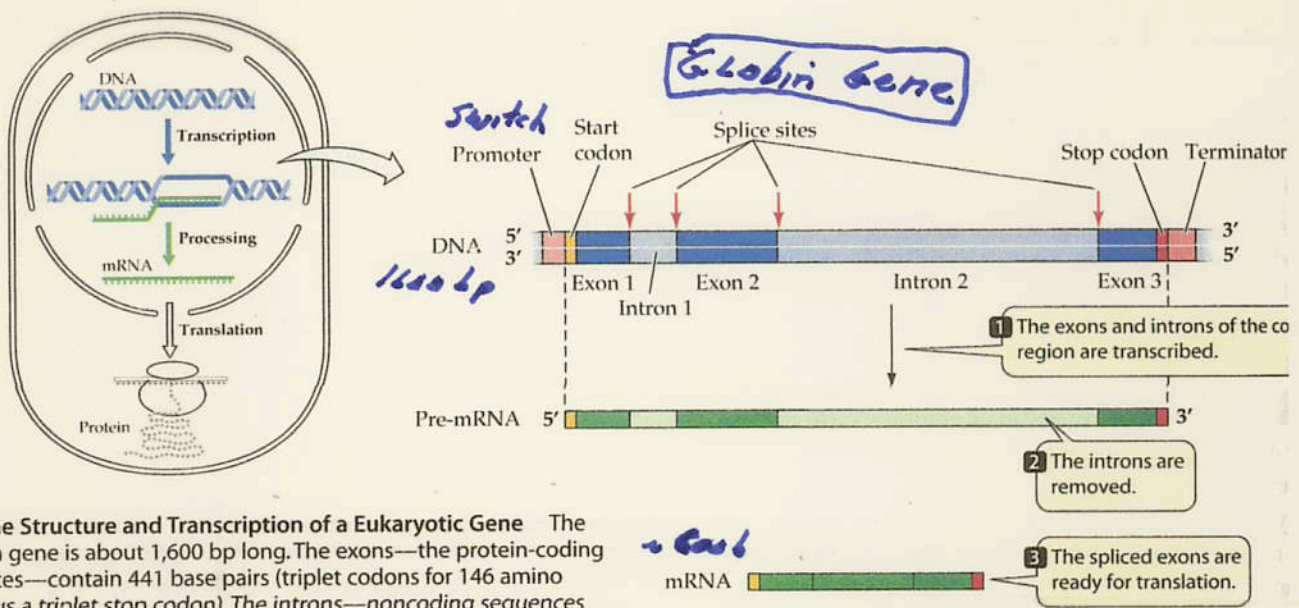
*HUMAN GENES mostly introns!*

**EUKARYOTIC CELLS MUST REMOVE NON-CODING REGIONS OF RNA BEFORE genetic CODE CAN BE TRANSLATED CONTINUOUSLY!**

**The Modular Organization of Genes and Gene Function Implies That There is No Limit to How Genes Can be Functionally Rearranged and Recombined Using Genetic Engineering?**

- a. Yes**
- b. No**

RNA Splicing - Removing Non-Coding Sequences from Primary Transcripts & Generating Functional mRNAs



14.4 The Structure and Transcription of a Eukaryotic Gene The β-globin gene is about 1,600 bp long. The exons—the protein-coding sequences—contain 441 base pairs (triplet codons for 146 amino acids plus a triplet stop codon). The introns—noncoding sequences of DNA—between codons 30 and 31 (130 bp long) and 104 and 105 (850 bp long), are initially transcribed, but are spliced out of the initial mRNA transcript.

Mutations → Blood Disorders  
Where can these occur?

MUTATIONS CAN OCCUR IN CODING REGION,  
SWITCH, & RNA SPLICING SITES

→ MUTANT PHENOTYPE

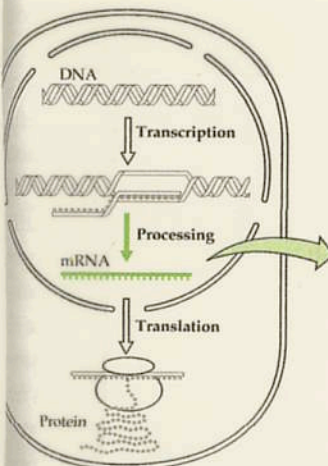
IMPLICATIONS FOR ENGINEERING EUKARYOTIC  
GENE IN BACTERIAL CELL FOR EXPRESSION?



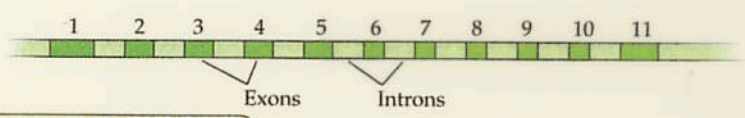


# ALternative Splicing - one Gene ↳ Several RNAs & Proteins!

Gene Active in variety of Cells  
But...!!!

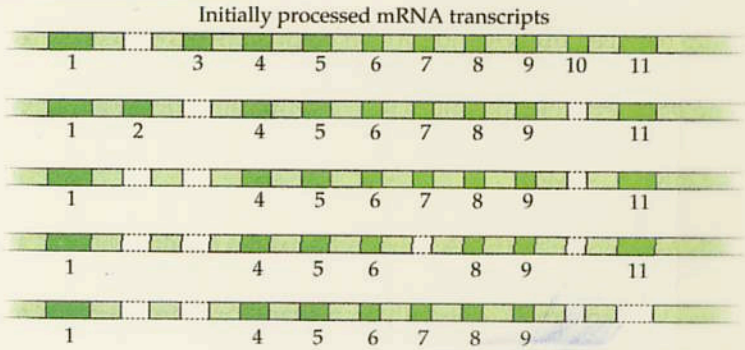


Primary RNA transcript for tropomyosin: 11 exons



Different splicing patterns in different tissues result in a unique collection of exons in mRNA for each tissue.

- Skeletal muscle: missing exon 2
- Smooth muscle: missing exons 3 and 10
- Fibroblast: missing exons 2, 3, and 10
- Liver: missing exons 2, 3, 7, and 10
- Brain: missing exons 2, 3, 10, and 11



5 different mRNAs!

Different mRNAs = different proteins = different functions!

4.20 Alternative Splicing Results in Different mRNAs and proteins In mammals, the protein tropomyosin is encoded by a gene that has 11 exons. Tropomyosin pre-mRNA is spliced differently in different tissues, resulting in five different forms of the protein.

IMPLICATION - HUMAN genome has only 30,000 genes but can give rise to many more proteins which are responsible for producing the phenotype!  
↳ human gene ≅ 3 transcripts

∴ Reason why human genome can contain same # of genes as the fly & mouse genomes!  
Implications for Genetic Engineering? use specific cDNA!

# IMPLICATIONS FOR "0-IT's in the DNA!!"

## Modular organization of Sequences

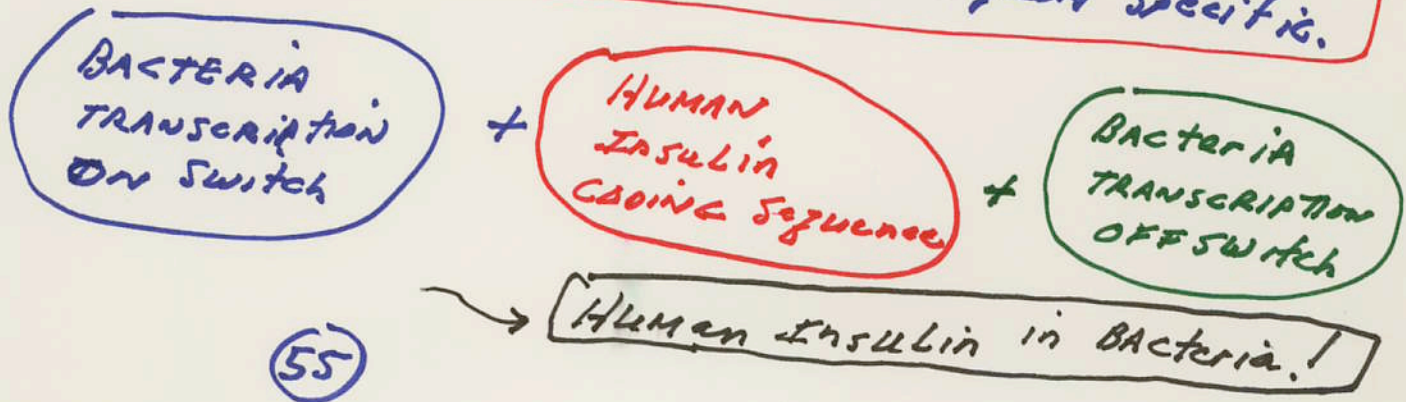
- ① DNA Replication  
ORI
- ② TRANSCRIPTION  
Switch/Regulator  
Terminator
- ③ Processing of RNA (Eukaryotes)  
Splicing Sites
- ④ Translation  
Start  
Stop  
Genetic Code / Codons
- ⑤ Coding Sequence  
Genetic Code

Modules → anything you  
want to do first only!

# ENGINEERING GENES REQUIRES:

- ① The **GENE** & its DNA Sequence
- ② A **ROADMAP** of where coding SEQUENCE & ALL SWITCHES LOCATED (Sequence, Restriction site Map)
- ③ **TRANSCRIPTION START AND STOP SWITCHES**
- ④ **CODING** Region of Gene (Genetic code part)
- ⑤ **TRANSLATION START AND STOP SWITCHES**
- ⑥ Kingdom-Specific Switches / Signals

NOTE: The General process of Gene → protein is the SAME in ALL organisms, but the Specific Switches & Enzymes (e.g. RNA Polymerase) are Kingdom Specific.



HOW DO GENES WORK & What  
ARE GENES IN CONTEXT OF....



THINKING ABOUT THE CONSEQUENCES  
of GMS

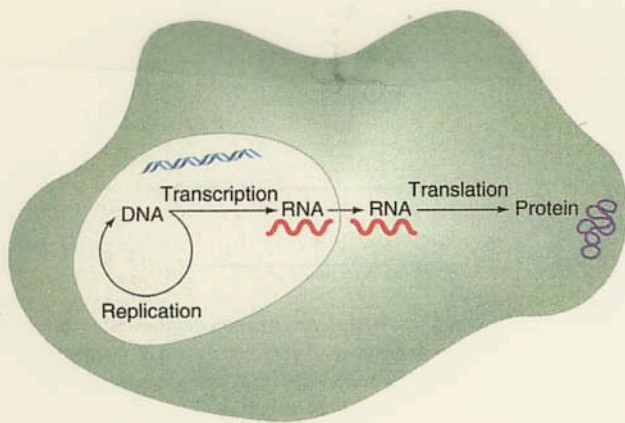


Figure 10-2 The three processes of information transfer: replication, transcription, and translation.

Need Science-based  
Questions & Science-based  
solutions

There's NO HOCUS FOCUS  
all hypotheses are  
testable!!

- ① What is a Gene?
- ② What is the Anatomy of a Gene?
- ③ How does the Gene Replicate?
- ④ How does the Gene direct Synthesis of a Protein?
- ⑤ Does the Gene work independently of other Genes?
- ⑥ What is the Sequence & Structure of the Protein?
- ⑦ How does it work in Cell?
- ⑧ Does the Protein Structure imply any potential "Harm"?
- ⑨ Does the Gene Change the organism? Fitness?

"Behind" ALL TRAITS!

SAME PROCESSES!