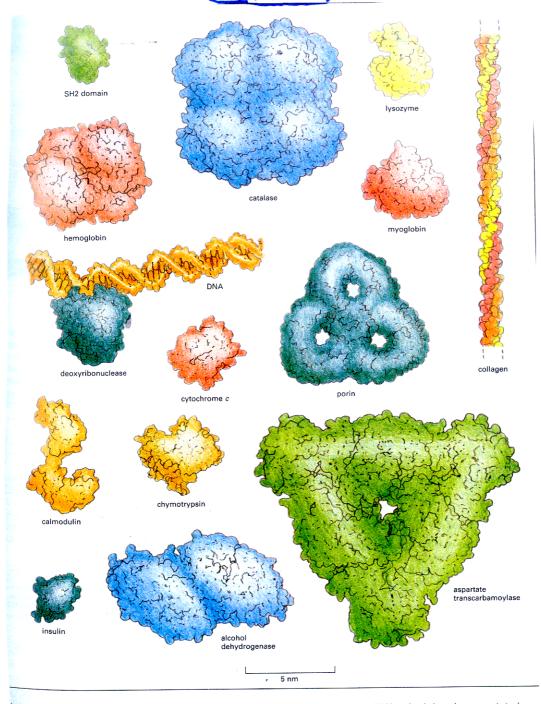
UNIQUE GENES IN A GENOME GIVE RISE TO UNIQUE, MATERIA



HOW DOES
Gene #
Relate to
Protein #?

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

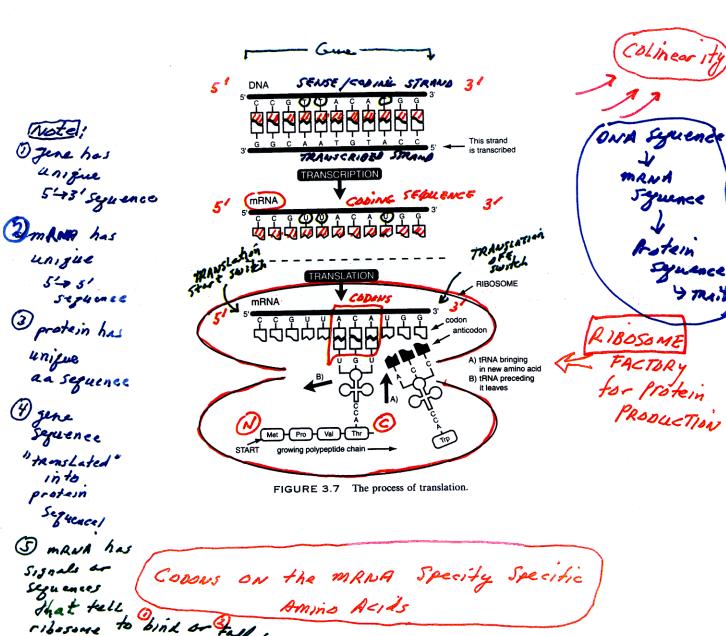
A PROTEIN is Synthesized

by TRANSLATING THE GENETIC

CODE OF the Gene/MRNA into

the Annio Acids y a specific

PROFEIR



PRotein is Nade on Rikosones Using the Genetic Code on mand for Instructions

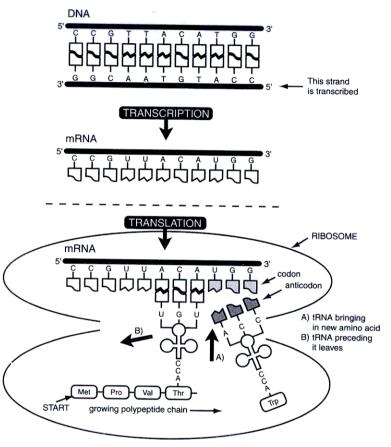
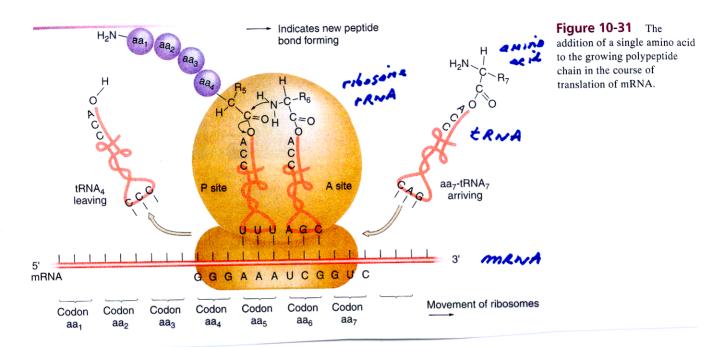


FIGURE 3.7 The process of translation.

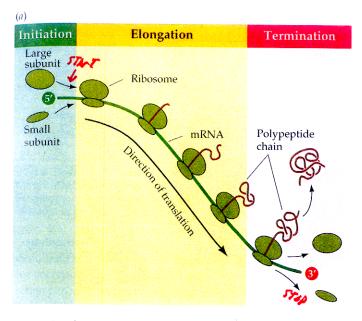
Coppus on Mana specify order & kinds

Ribosones TRANSLate the Genetic Code



colon by coon!

ONE MENA IS TRANSLATED MANY TIMES

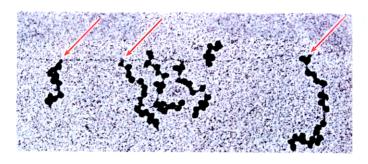


12.13 A Polysome

(a) A polysome consists of ribosomes and their growing polypeptide chains moving in single file along an mRNA molecule. (b) An electron microscopic view of a polysome.

IMPLICATIONS FOR BIOLOGY

Gene Activity CAN BE VISUALIZE!



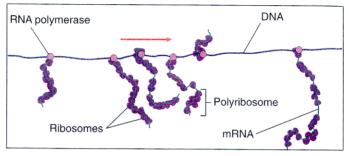


FIGURE 15.10

Translation in action. Bacteria have no nucleus and hence no membrane barrier between the DNA and the cytoplasm. In this electron micrograph of genes being transcribed in the bacterium *Escherichia coli*, you can see every stage of the process. The arrows point to RNA polymerase enzymes. From each mRNA molecule dangling from the DNA, a series of ribosomes is assembling polypeptides. These clumps of ribosomes are sometimes called "polyribosomes."

TRANSCRIPTION / TRANSLATION ARE
COUPLED in Bacteria

ELEKARYOTIC & PROKARYOTIC
Gene Expression Processes
Oitter Slightly

Gener Differ
Switches / RNA Polymereses Differ Differ
Oither Differ
Genetic Code the SAME
General Processes the SAME

but

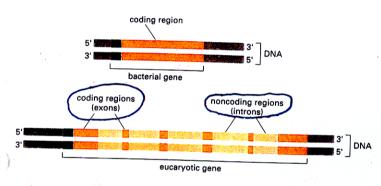


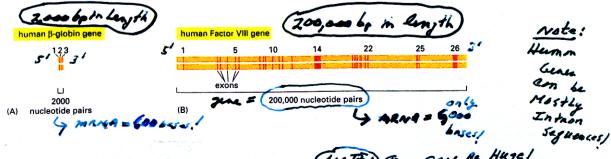
Figure 7-13 Comparison of a bacterial gene with a eucaryotic gene. The bacterial gene consists of a single stretch of unimerrupted nucleotide sequence that encodes the amino acid sequence of a protein. In contrast, the coding sequences of most eucaryotic genes (exons) are interrupted by noncoding sequences (introns). Promoters for transcription are indicated in green.

SWITCHES
WIPLE
TO
BACTERIA
TO
PLANTS JAMA

Ecchargotic Genes Thave NON-coling regions "Stuck" in coding regions

Prokaryotic Genes only have coding regions!

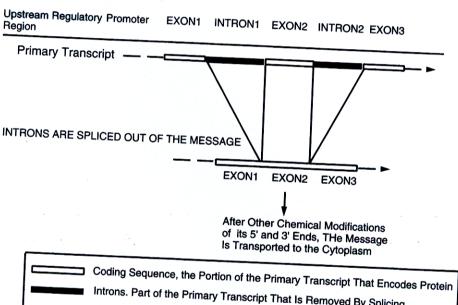
Thus: Eukargotic cells Must remove mon-coding regions in manh BEFORE genetic code com be tamslated continuously!



(VOTE) gene CAN Be Huge!

EUKARYOTIC GENES HAVE INTROVS OR NON-coomic ONA Interspersel in coding sequences or exons

> INTRONS ARE TRANSCRIBED BUT MUST BE Spliced out in nucleus to Make mand with continuous Jenete code//



Introns. Part of the Primary Transcript That Is Removed By Splicing 5' and 3' Flanking Sequences That Stay in the Mature mRNA, But Do Not Code for Protein

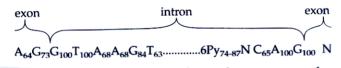
FIGURE 3.14 Exons, introns, and splicing.

BACTERIAL armes EUKARYOTE RNAS/

Yol It's in the Sequences!

intron	
exon-GT	AG-exon

The sequences shown here are for the DNA nontemplate strand (equivalent to the RNA transcript, but with T rather than U). In addition, there are short consensus sequences at the exon–intron junctions. For nuclear genes, the consensus junctions are



Specific Sequences Required FOR
RNA Splieing!

What happens it these Sequences Are Mutated in a Gene?

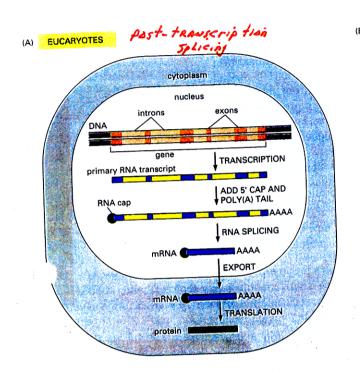


Eukaryotie Gene TRANSCRIPTS

PROCESSEL By Splicing in

the Nucleus

to FORM MRNAS



(B) PROCARYOTES DNA TRANSCRIPTION mRNA. TRANSLATION protein

NO SPLICINS

Figure 7-19 Summary of the steps leading from gene to protein. The final level of a protein in the cell depends on the efficiency of each step and on the rates of degradation of the RNA and protein molecules. (A) In eucaryotic cells, the initial RNA molecule produced by transcription (the primary transcript) contains both intron and exon sequences. Its two ends are modified, and the introns are removed by an enzymatically catalyzed RNA splicing reaction. The resulting mRNA is then transported from the nucleus to the cytoplasm, where it is translated into protein. Although these steps are depicted as occurring one at a time, in a sequence, in reality they often occur simultaneously. For example, the RNA cap is typically added and splicing typically begins before the primary transcript has been completed. (B) In procaryotes, the production of mRNA molecules is simpler. The 5' end of an mRNA molecule is produced by the initiation of transcription by RNA polymerase, and the 3' end is produced by the termination of transcription. Since procaryotic cells lack a nucleus, transcription and translation take place in a common compartment. In fact, translation of a bacterial mRNA often begins before its synthesis has been completed.

Engineer mand not june!

Modular organization y Sequences

- 1) DNA Replication
 - 2 TRANSCRIPTION

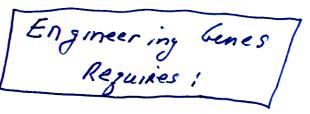
 Switch/Regulator

 Terminator
 - 3 Processing of RNA (Eukorgotes)
 Splicing Sites
- 4) Translation

Stop Genetie Code/Codons

(5) Coling Squence Genetic Code

Modules - say thing you want to so Jenets call!



- 1) The Gene & its Sequence
- 2) A Boadmap of Where Coding Sequence & All switches Iscated (What's the roal map?)
- 3 TRANSCRIPTION Start & Stap Switches
- (9) Coding Part of Gene / Genetic Code Part
- (5) TRANSLation Start x Stop Switches
- (Kingdom- Specific Switches/Signals)

Note: The general process of glas - protein is the same in all organisms but the Specific switches & enzymes (4.5., RAA polyneuse) differ in Kingdoms !!

PLANT TRANSCRIPTION ON Switch

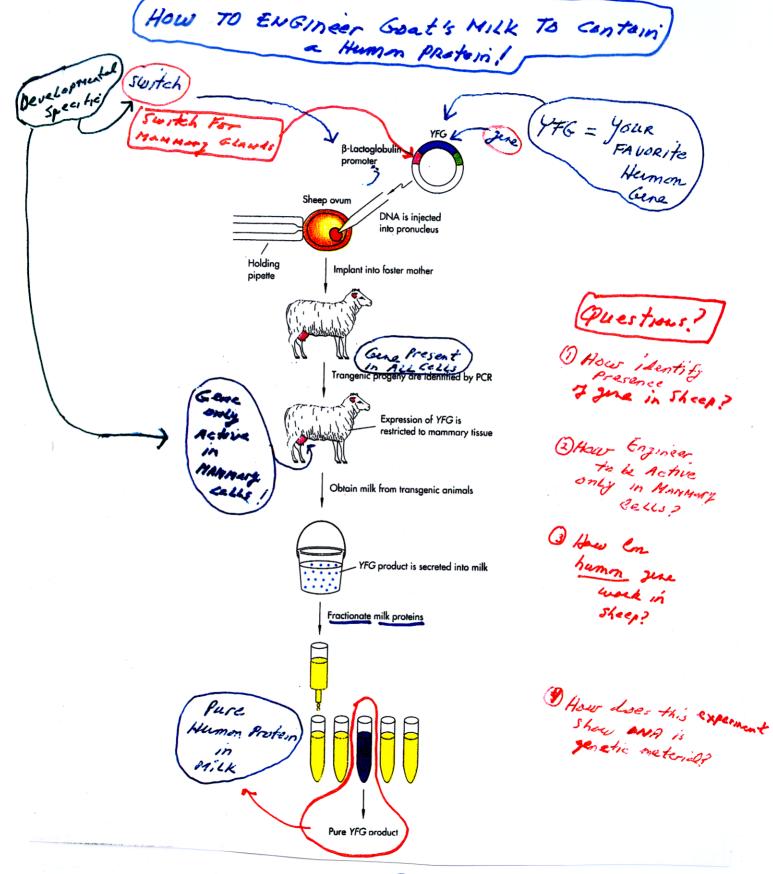
Human Coping Squence

Plant
TRANSCRIPTION
off
Switch

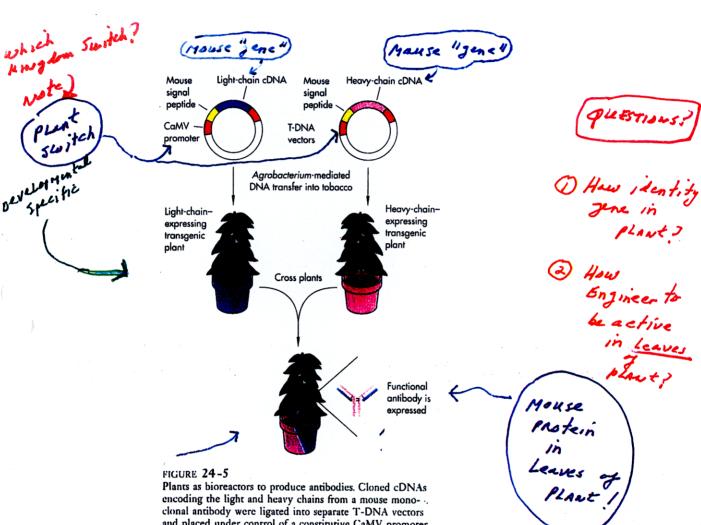
It's that easy
Because living Cells
Use SAME Overall genetic
processes.

human pritein in plant

There are NO Limits!



CAN ANIMAL GENES BE ENGINEERED TO WORK IN PLANTS?



Plants as bioreactors to produce antibodies. Cloned cDNAs encoding the light and heavy chains from a mouse monoclonal antibody were ligated into separate T-DNA vectors and placed under control of a constitutive CaMV promoter. The plasmids were transferred separately into tobacco plants by Agrobacterium infection. Transgenic plants containing the light- and heavy-chain genes were sexually crossed to produce progeny plants that contained both genes. Examination of protein extracted from leaves demonstrated the expression of functional antibody molecules in these progeny plants. Other experiments showed that the presence of a signal sequence was necessary for high-level expression. These results suggest that the plant secretion machinery can recognize the mouse signal peptide.

IMPOSTANT CONCEPT!

What does
this experiment
tell us about
fractic
processes
give.
Protein
Synthesis?

Yol It's ALL in the Sequences!

ONA, Gene, Switch, pri, mana, Pratein !

NO HOCUS POCUS/

What Does this INPLY For BIOLOGY & Genetic Engineering /