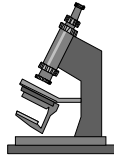
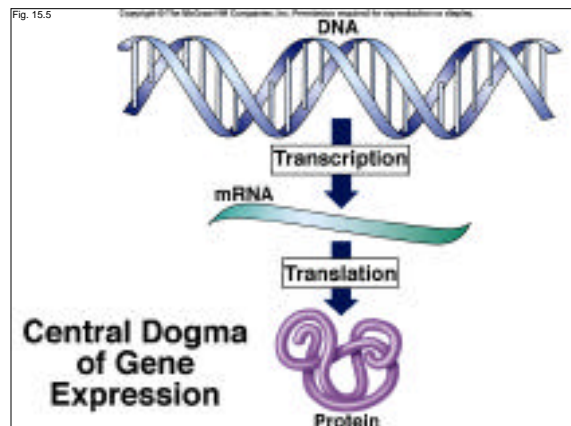
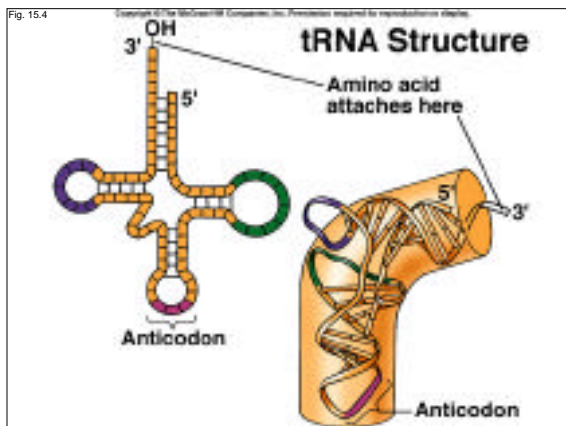
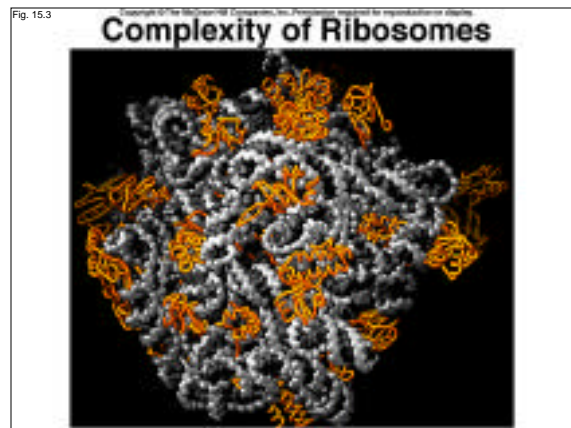
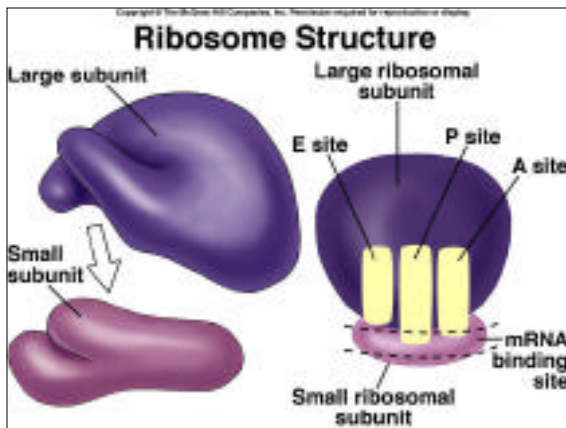
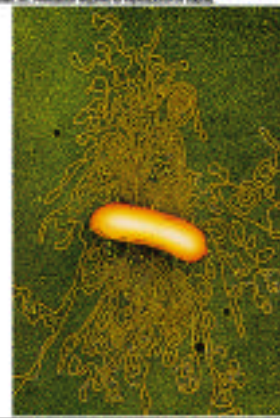
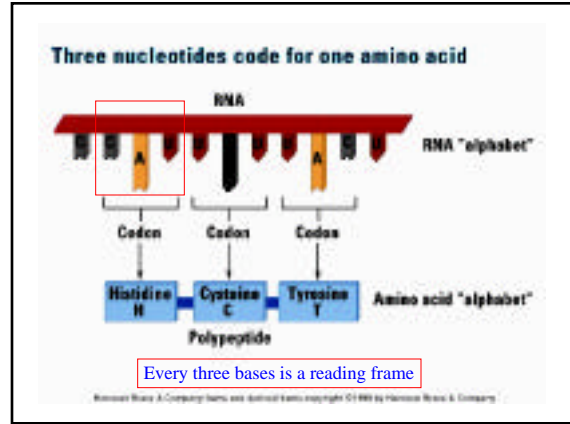
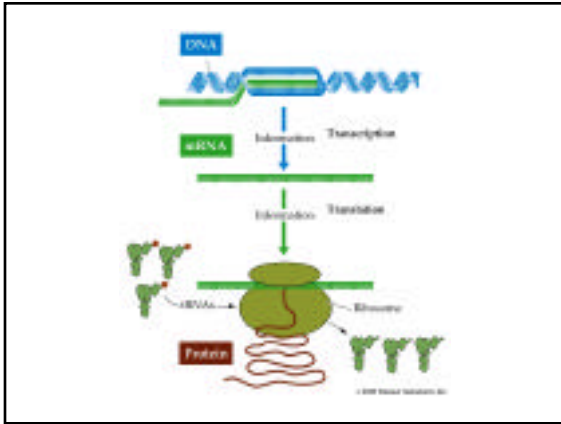


Transcription and Translation



Unraveled Chromosome of *E. coli* Bacterium





12.1 Signals that Start and Stop Transcription and Translation

	TRANSCRIPTION	TRANSLATION
Initiation	Promoter sequence in mRNA	AUG start codon
Termination	Terminator sequence in DNA	UAA, UAG, or UGA stop codon in mRNA

Reading-Frame Alterations

Delete 1 base

Hypothesis A: unpunctuated
 WHY DID HER EDB ATE ATT HEF ATR AT?
 (Nonsense)

Hypothesis B: punctuated
 WHY DID HE O EDO ATO ATO HE O ATO AT?
 (Nonsense)

Delete 3 bases

Hypothesis A: unpunctuated
 WHY DID THE RED B ATE AT THE FAT RAT?
 (Sense)

Hypothesis B: punctuated
 WHY DID HE O DOB OEA OTH OFA ORA?
 (Nonsense)

Breaking the genetic code

EXPERIMENT

Question: What are the amino acids specified by the triplet codons UUU, AAA, and CCC?

METHOD

Three test tubes containing yellow liquid are shown. The first tube is labeled 'UUUUUUUU' and points to three brown circles labeled 'Phe'. The second tube is labeled 'AAAAAAA' and points to three red circles labeled 'Lys'. The third tube is labeled 'CCCCCCC' and points to three blue circles labeled 'Pro'.

RESULTS

Conclusion: UUU is an mRNA codon for phenylalanine. AAA is an mRNA codon for lysine. CCC is an mRNA codon for proline.

Genetic code

First letter (5' end)	Second letter			Third letter (3' end)
	U	C	A	
U	UUU Phe UUC Phe UUA Leu UUG Leu	UCU Phe UCC Ser UCA Ser UCG Ser	UAU Tyr UAC Tyr UAA Stop UAG Stop	UGU Cys UGC Cys UGA Stop UGG Trp
C	CUU Leu CUC Leu CUA Leu CUG Leu	CCU Pro CCC Pro CCA Pro CCG Pro	CAU His CAC His CAA Gln CAG Gln	CGU Arg CGC Arg CGA Arg CGG Arg
A	AUU Ile AUA Ile AUG Met	ACU Thr ACC Thr ACA Thr ACG Thr	AAG Lys AAA Lys AAG Lys AAA Lys	AGU Ser AGC Ser AGA Arg AGG Arg
G	GUC Val GUA Val GUG Val	GCC Ala GCA Ala GCG Ala	GAU Asp GAC Asp GAA Gly GAG Gly	GUU Gly GUC Gly GUA Gly GUG Gly

The genetic code is redundant
 2 bases per codon $4 \times 4 = 16$
 3 bases per codon $4 \times 4 \times 4 = 64$

Diagram showing four different codons (UCU, UCC, UCA, UCG) all coding for the amino acid Serine.

Fig. 15.6b

The Genetic Code

First Letter	Second Letter				Third Letter	
	U	C	A	G		
U	UUU	UCU	UAU	UGU	U	
	UUC	UCC	UAC	UGC		C
	UUA	UCA	UUA	UGA		A
	UUG	UCG	UUG	UGG		G
C	CUU	CCU	CAU	CGU	U	
	CUC	CCC	CAC	CGC		C
	CUA	CCA	CAA	CGA		A
	CUG	CCG	CAG	CGG		G
A	AUU	ACU	AUU	AUU	U	
	AUC	ACC	AAC	AAC		C
	AUA	ACA	AAA	AAA		A
	AUG	ACG	AAU	AAG		G
G	GUU	UCU	GAU	GGU	U	
	GUC	UCC	GAC	GGC		C
	GUA	UCA	GAA	GGA		A
	GUG	UCG	GAG	GGG		G

Table 15.1 The Genetic Code

First Letter	Second Letter				Third Letter	
	U	C	A	G		
U	UUU	UCU	UAU	UGU	U	
	UUC	UCC	UAC	UGC		C
	UUA	UCA	UUA	UGA		A
	UUG	UCG	UUG	UGG		G
C	CUU	CCU	CAU	CGU	U	
	CUC	CCC	CAC	CGC		C
	CUA	CCA	CAA	CGA		A
	CUG	CCG	CAG	CGG		G
A	AUU	ACU	AUU	AUU	U	
	AUC	ACC	AAC	AAC		C
	AUA	ACA	AAA	AAA		A
	AUG	ACG	AAG	AAG		G
G	GUU	UCU	GAU	GGU	U	
	GUC	UCC	GAC	GGC		C
	GUA	UCA	GAA	GGA		A
	GUG	UCG	GAG	GGG		G

Fig. 15.7

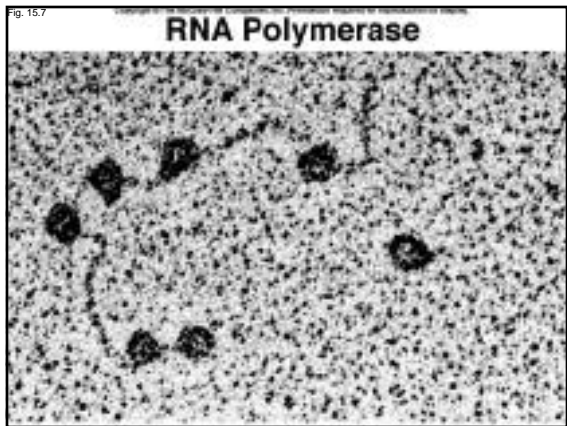


Fig. 15.8

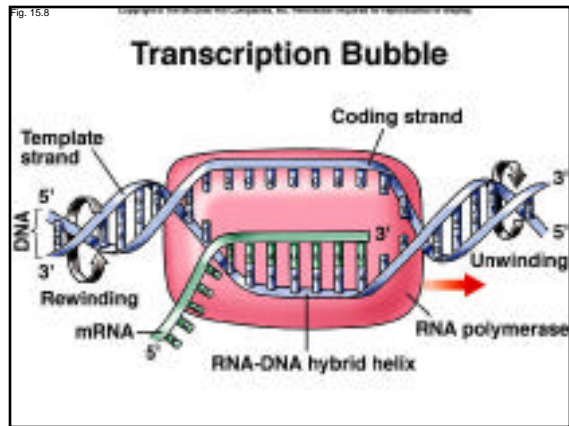


Fig. 15.9

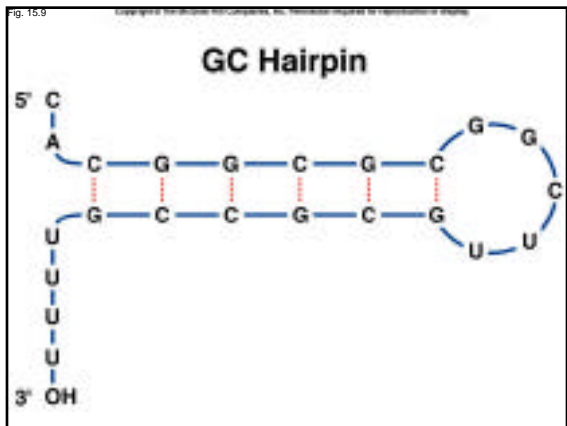
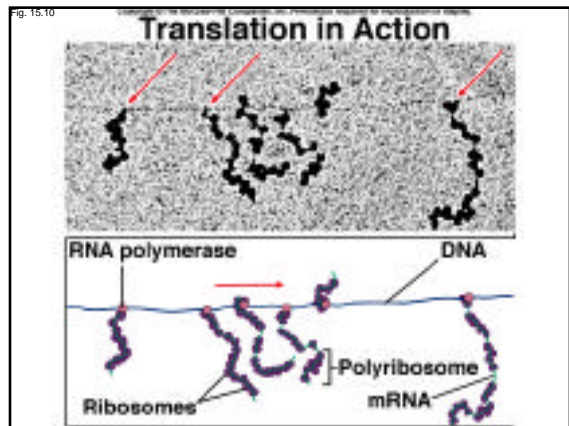
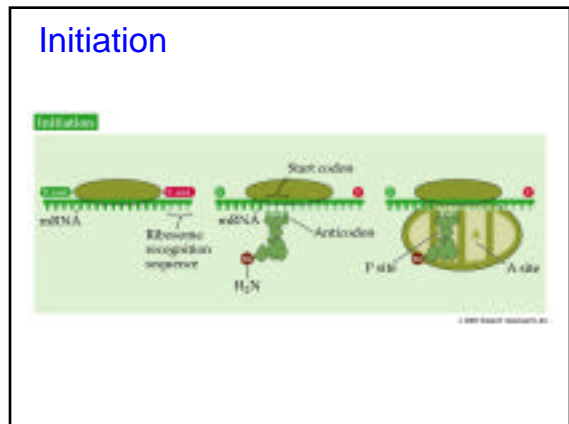
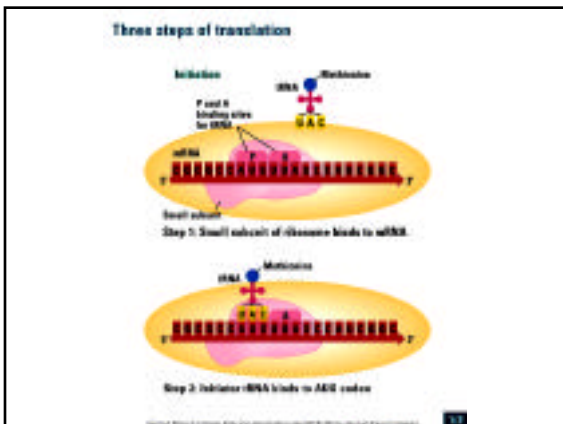
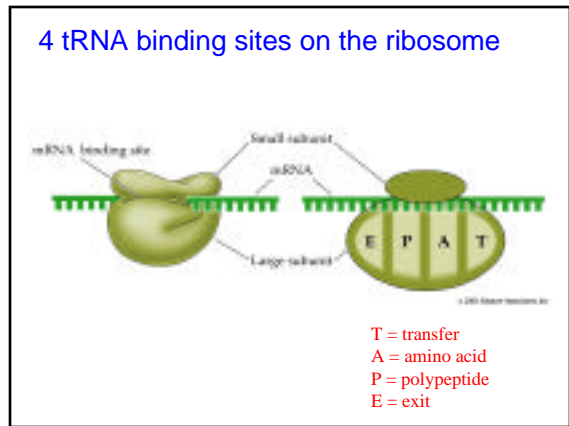
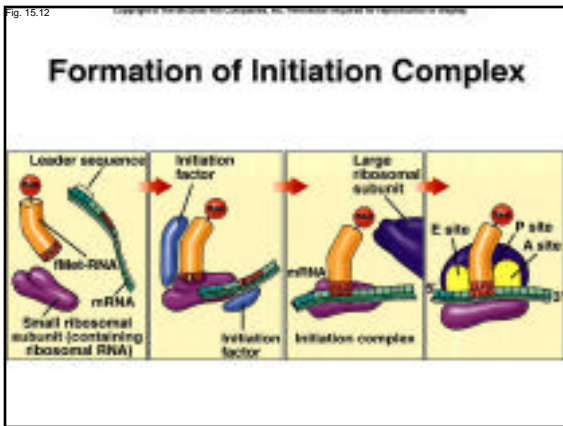
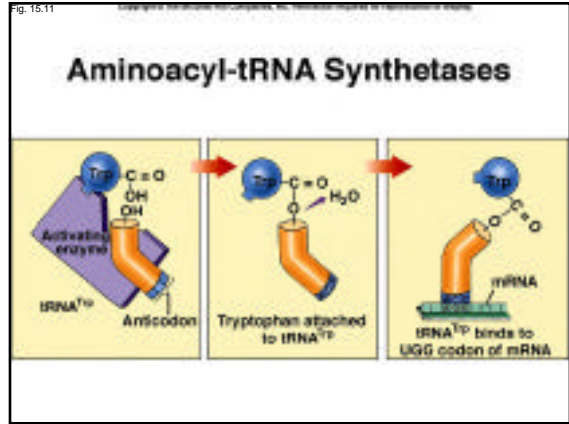
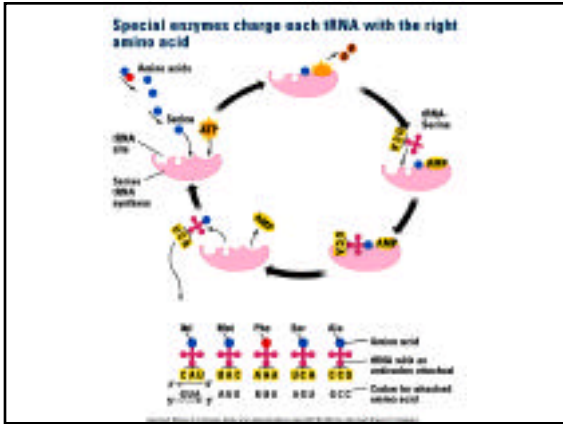
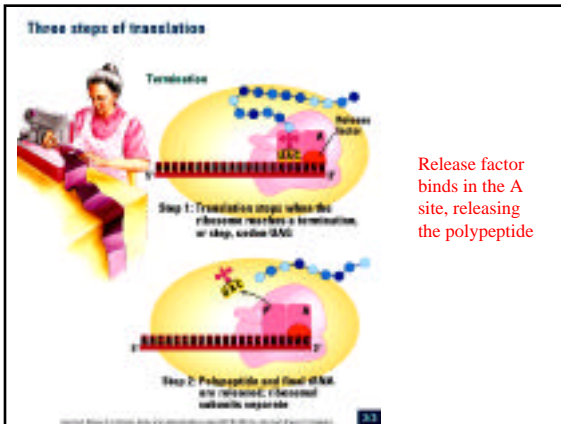
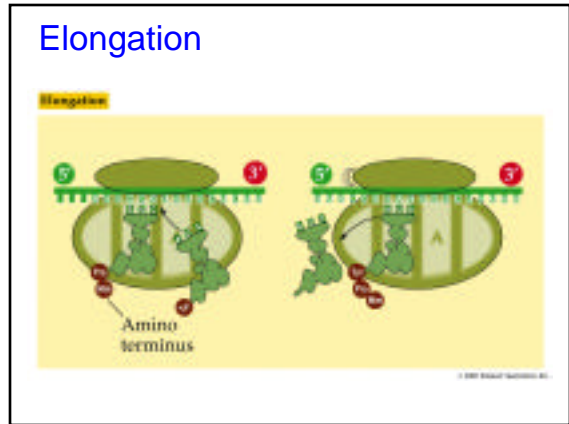
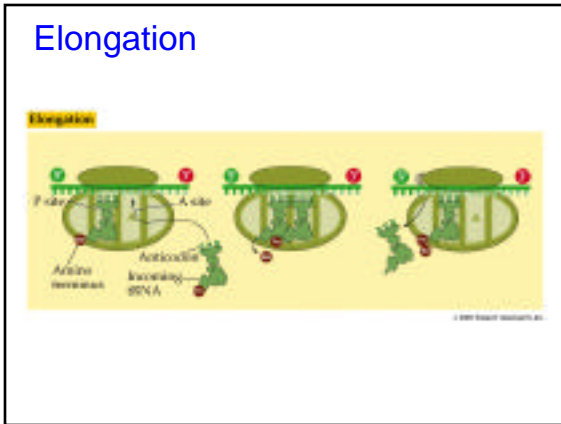
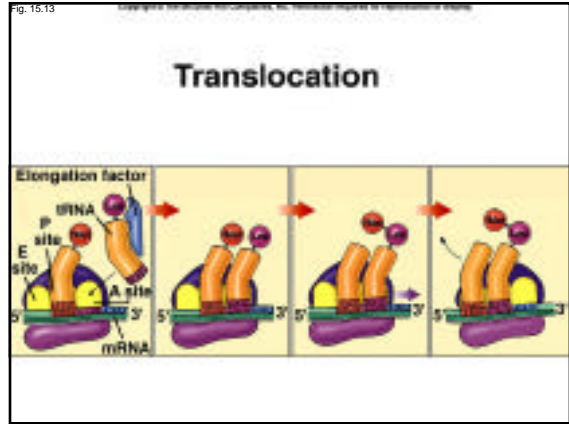
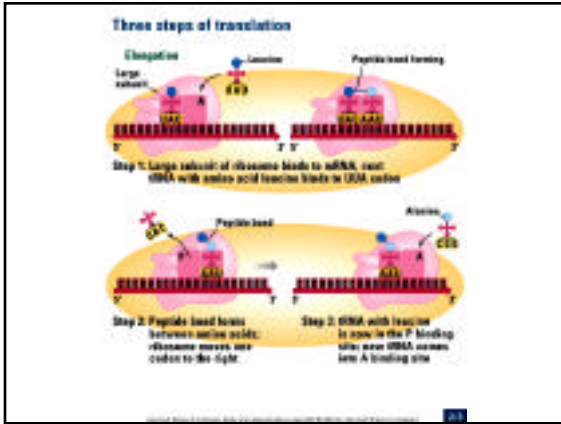


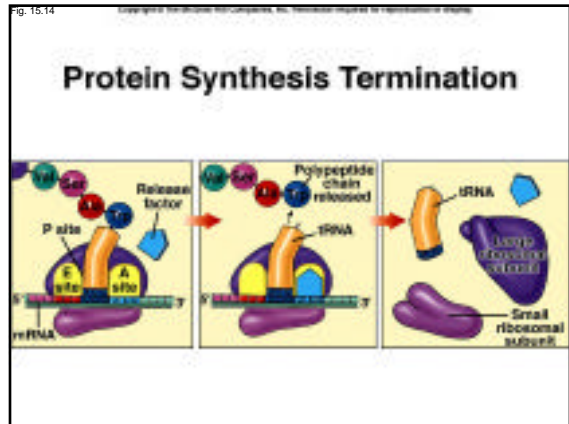
Fig. 15.10







Release factor binds in the A site, releasing the polypeptide



Termination



12.2 Antibiotics that Inhibit Bacterial Protein Synthesis

ANTIBIOTIC	STEP INHIBITED
Chloramphenicol	Formation of peptide bonds
Erythromycin	Translocation of mRNA along ribosome
Neomycin	Interactions between tRNA and mRNA
Streptomycin	Initiation of translation
Tetracycline	Binding of tRNA to ribosome

Translation summary

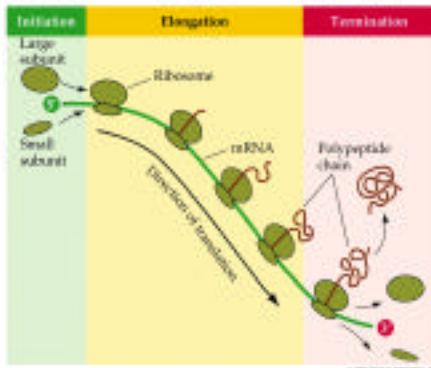


Fig. 15.15a

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Introns and Exons (1)

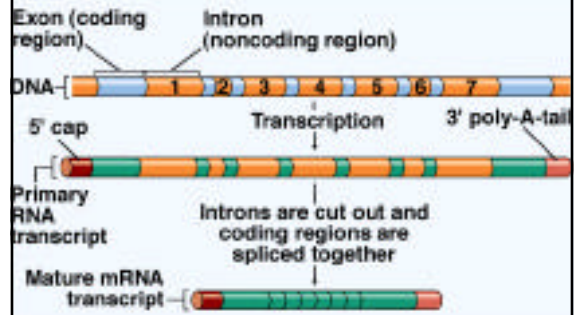


Fig. 15.15b

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Introns and Exons (2)

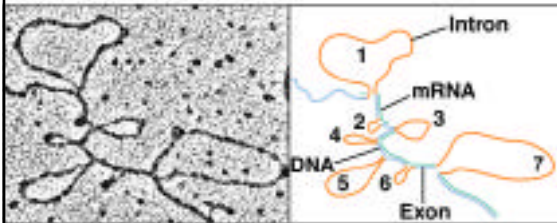
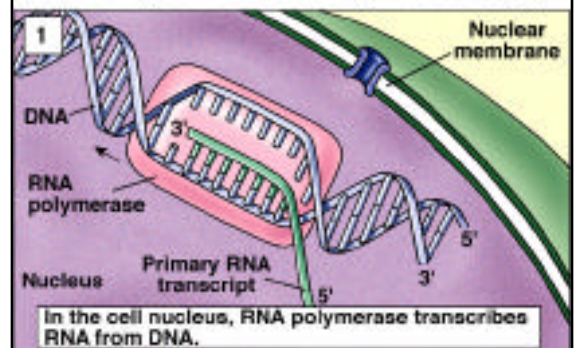
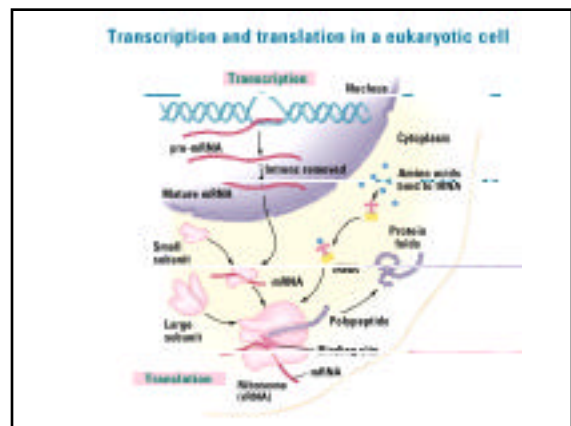
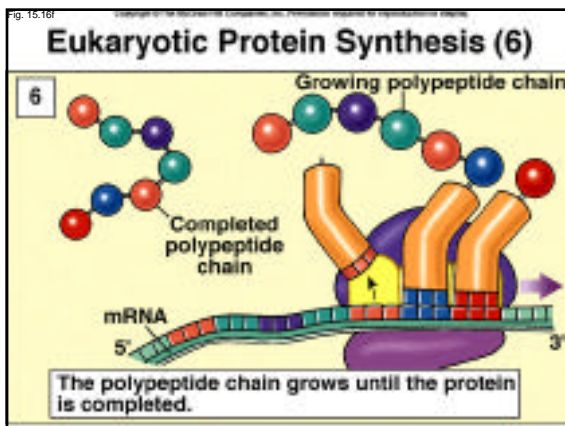
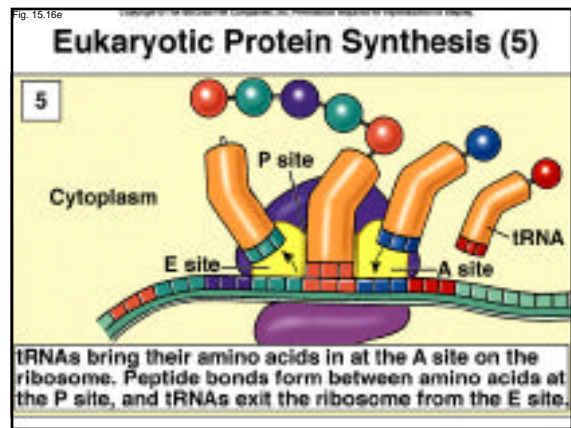
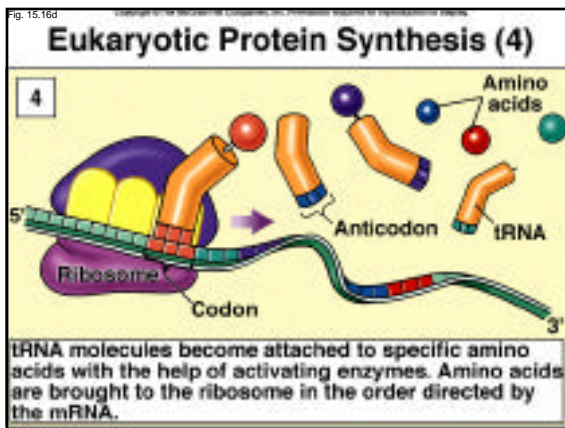
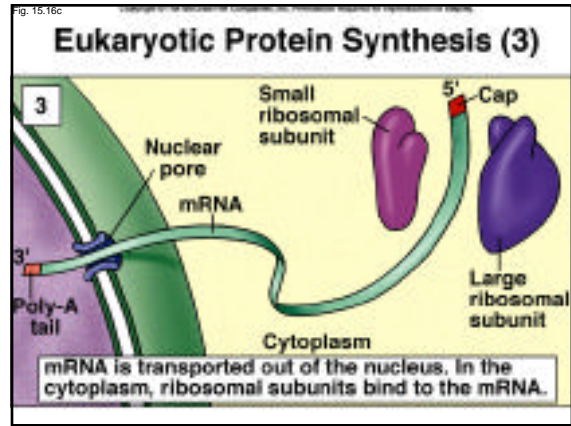
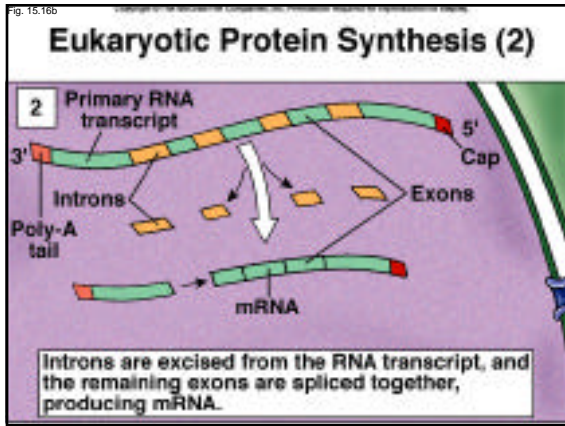


Fig. 15.15a

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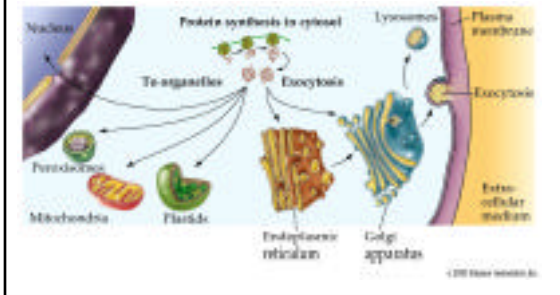
Eukaryotic Protein Synthesis (1)





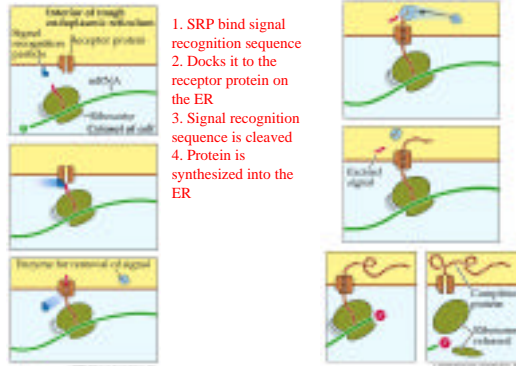
Signal sequences direct proteins

Signal sequences are short sequences at the N-terminus or in the middle that direct the proteins

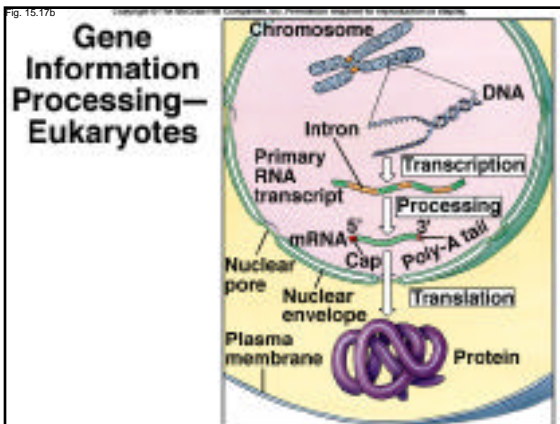
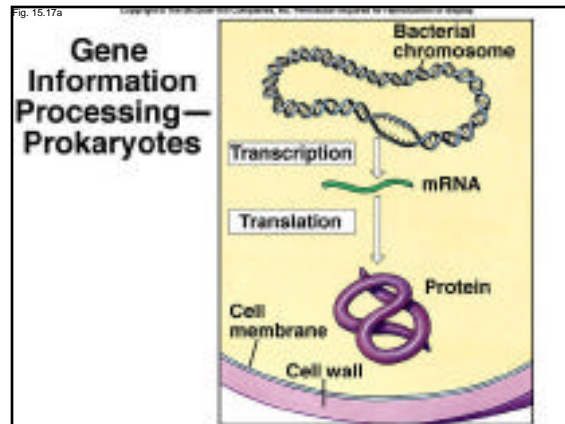
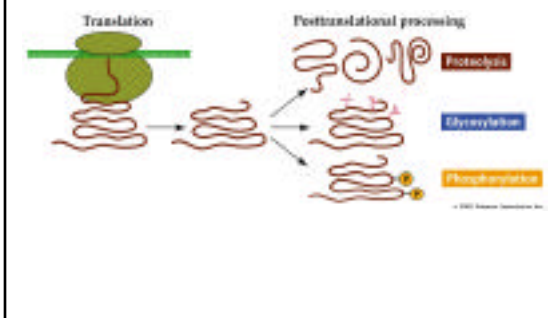


ER targeting of proteins

1. SRP bind signal recognition sequence
2. Docks it to the receptor protein on the ER
3. Signal recognition sequence is cleaved
4. Protein is synthesized into the ER



Post-translational modifications



Mutations

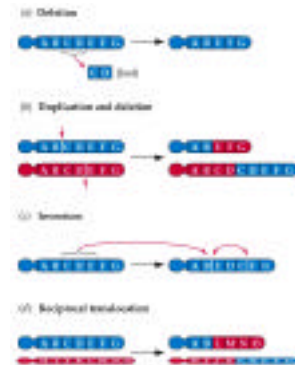
Permanent changes in the DNA sequence

Somatic - non gamete cells
Germ-line mutations

Point mutations - substitution or deletion/addition of a nucleotide

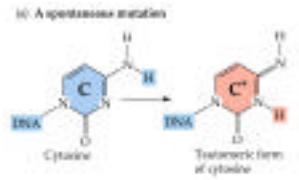
- Silent (no change)
- missense (sickle cell)
- Nonsense (Stop)
- Frame shift

Chromosome mutation - loss or change in a DNA segment



Spontaneous and induced mutations

Tautomeric C pairs with A



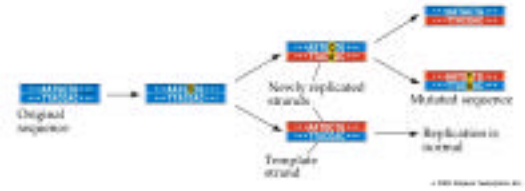
Base modifications by chemicals or radiation

benzpyrene
Nitrates
X-rays
UV light



Mutation propagation

(i) The consequences of either mutation



The End.