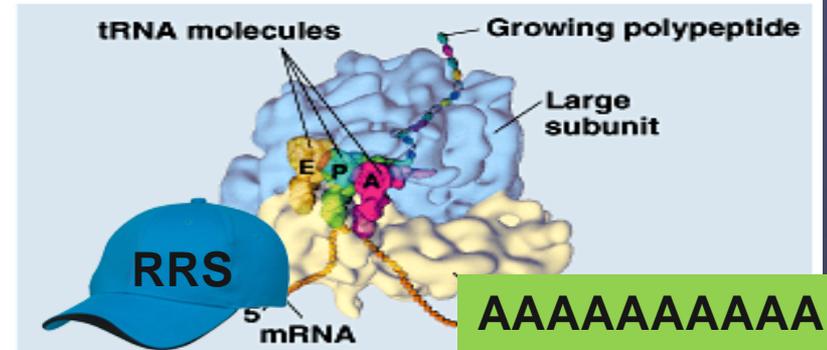


Part II: Translation

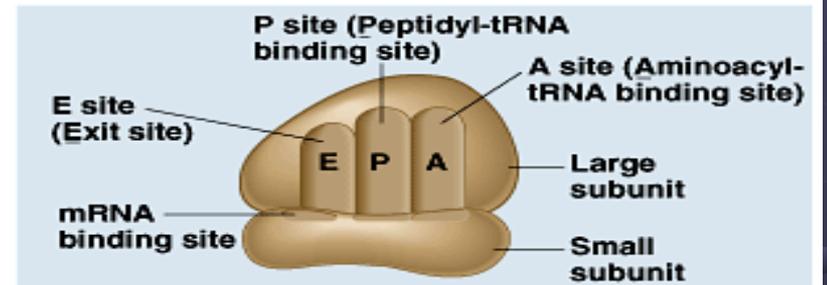
convert from mRNA language into **protein** language

Ribosomes

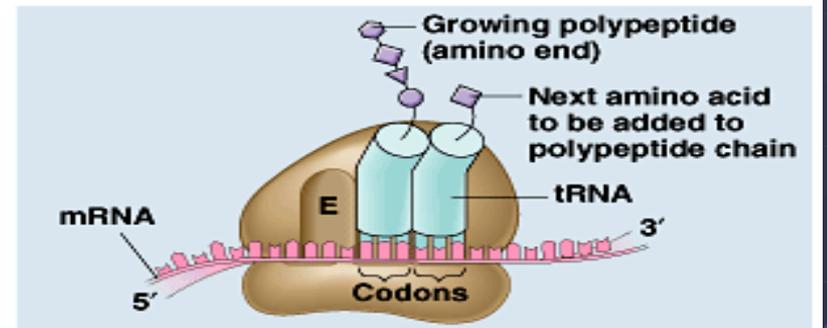
- First the mRNA binds to a ribosome in the cytoplasm or on the RER using its RRS in the CAP



(a) Computer model of functioning ribosome



(b) Schematic model showing binding sites



(c) Schematic model with mRNA and tRNA

How does mRNA code for proteins?

DNA

TACGCACATTTACGTACGCGG



mRNA

AUGCGUGUAAAUGCAUGCGCC



protein

Met Arg Val Asn Ala Cys Ala

How can you code for 20 amino acids with only 4 nucleotide bases (A,U,G,C)?

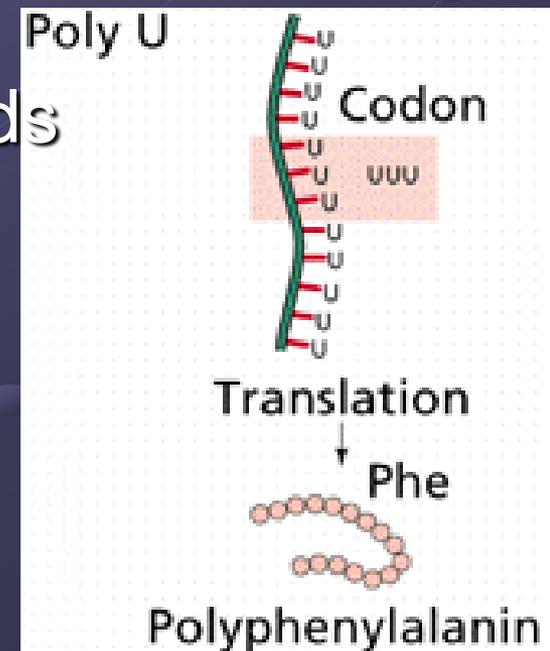
1960 | 1968

Cracking the code

● Nirenberg & Matthaei

- determined 1st codon–amino acid match
 - UUU coded for phenylalanine
- created artificial poly(U) mRNA
- added mRNA to test tube of ribosomes, tRNA & amino acids
 - mRNA synthesized single amino acid polypeptide chain

phe–phe–phe–phe–phe–phe



mRNA codes for proteins in triplets

DNA

TACGCACATTACGTACGCGG



mRNA

AUGCGUGUAAAUGCAUGCGCC

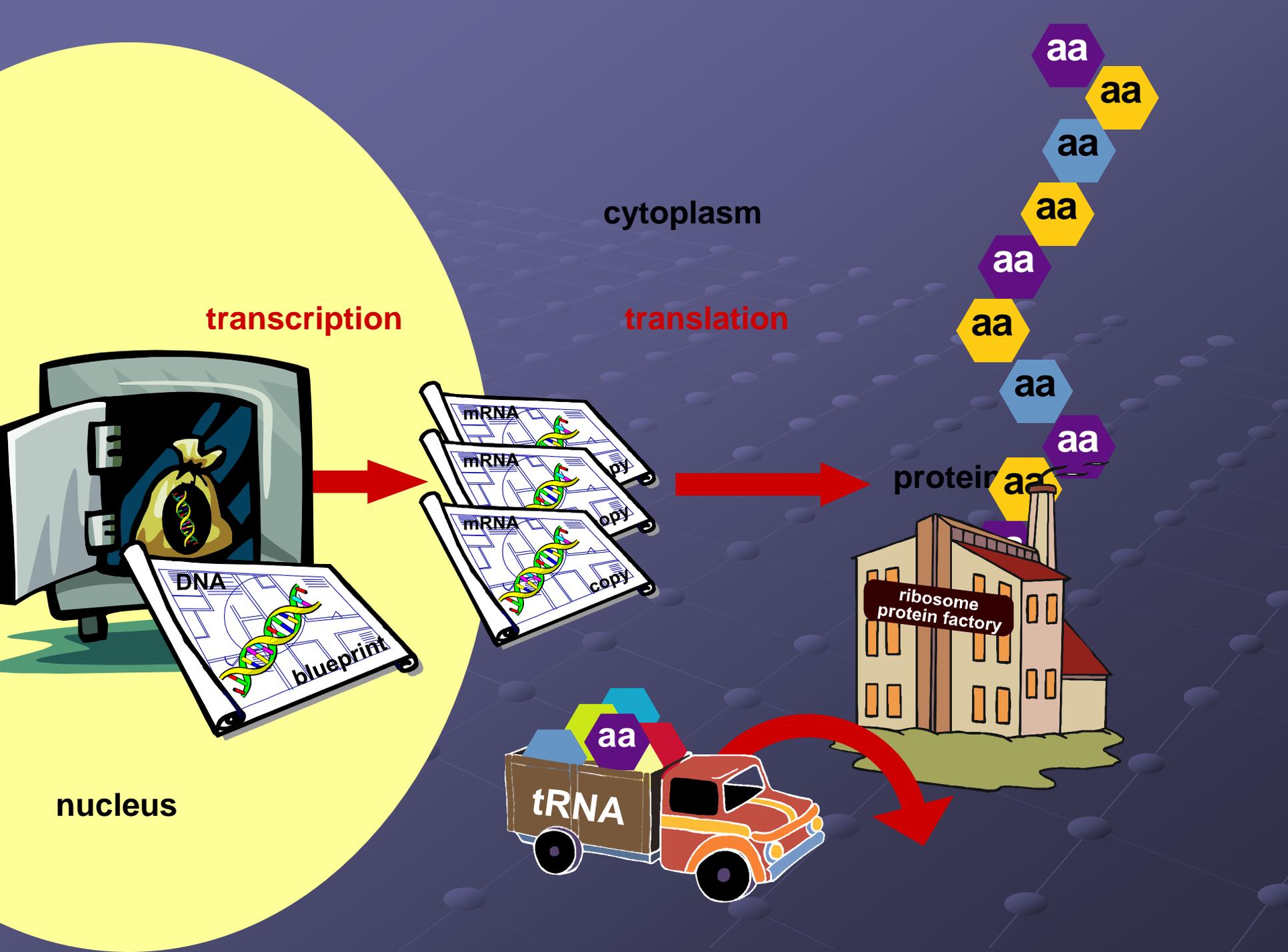


protein

Met Arg Val Asn Ala Cys Ala

- The mRNA is read 3 letters at a time (CODON)
- Each CODON represents one specific Amino Acid
 - There are 64 possible 3 letter combinations BUT only 20 amino acids....
 - SO, some Codons code for more than one amino acid

		SECOND BASE				
		U	C	A	G	
U	UUU	UCU	UAU	UGU	U	
	UUC	UCC	UAC	UGC	C	
	UUA	UCA	UAA	UGA	A	
	UUG	UCG	UAG	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	AAU	AGU	U	
	AUC	ACC	AAC	AGC	C	
	AUA	ACA	AAA	AGA	A	
	AUG	ACG	AAG	AGG	G	
G	GUU	GCU	GAU	GGU	U	
	GUC	GCC	GAC	GGC	C	
	GUA	GCA	GAA	GGA	A	
	GUG	GCG	GAG	GGG	G	



transcription

cytoplasm

translation

nucleus

protein

ribosome
protein factory

tRNA

aa

aa

aa

aa

aa

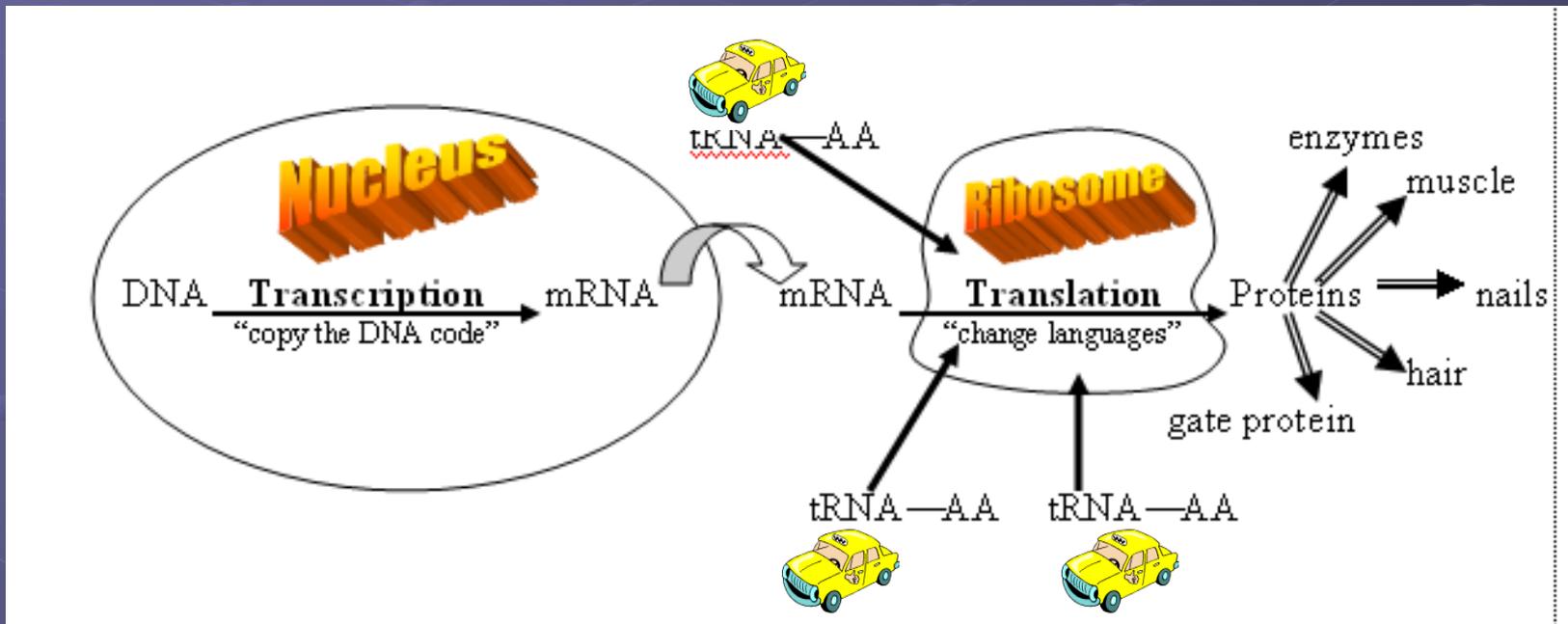
aa

aa

aa

aa

- Once the mRNA reaches the ribosome, and the CODONS are read,
- Transfer RNA (tRNA) “Taxi” service delivers the correct Amino Acid to the ribosome



- Each tRNA molecule has a triplet anticodon on one end and an amino acid attachment site on the other

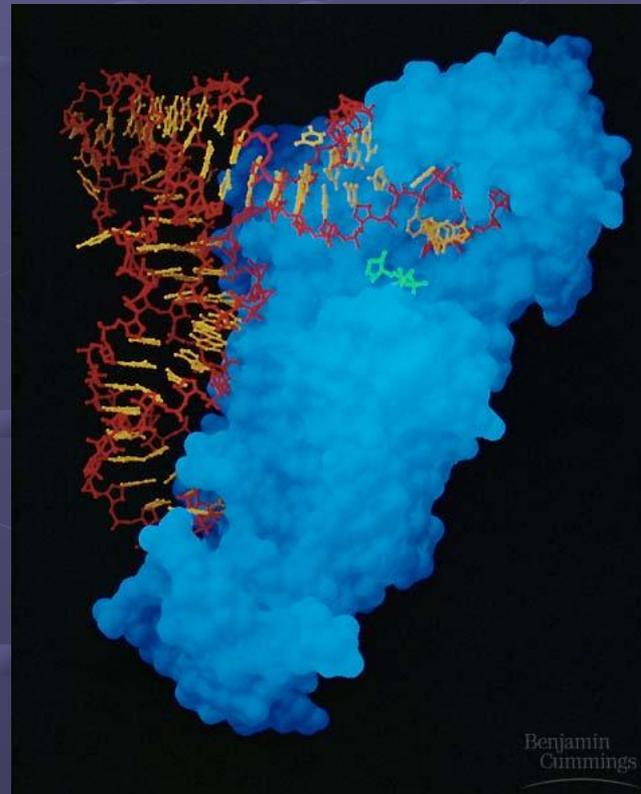
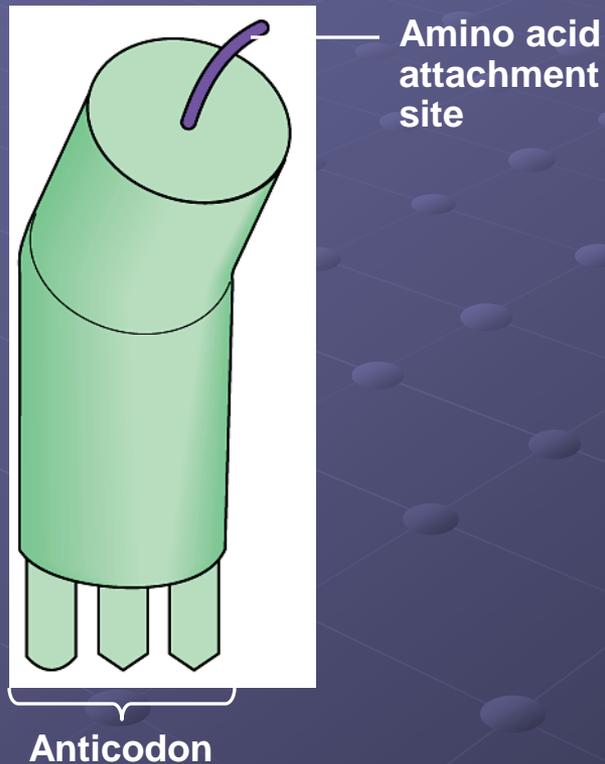
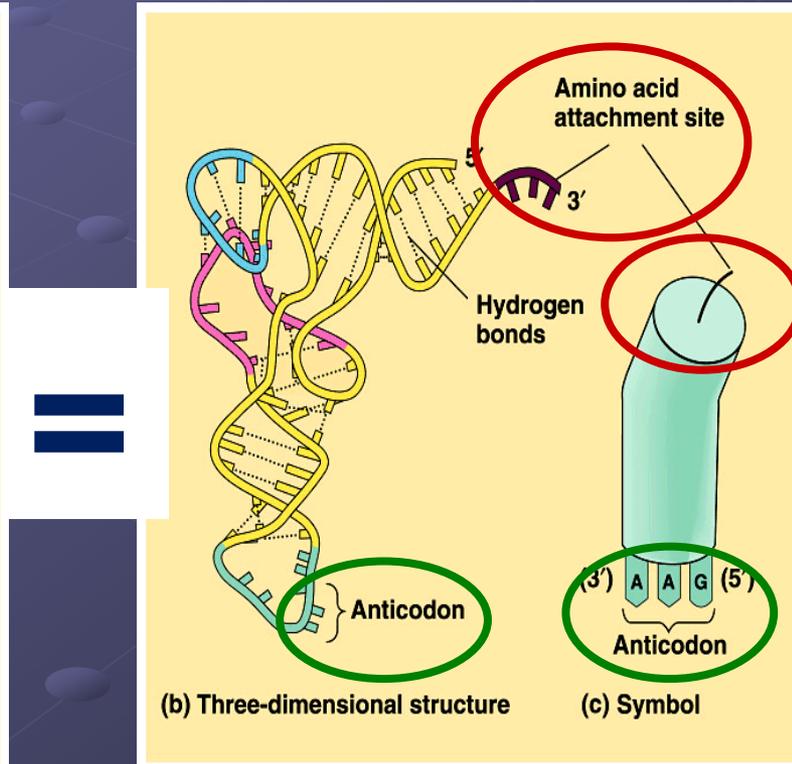
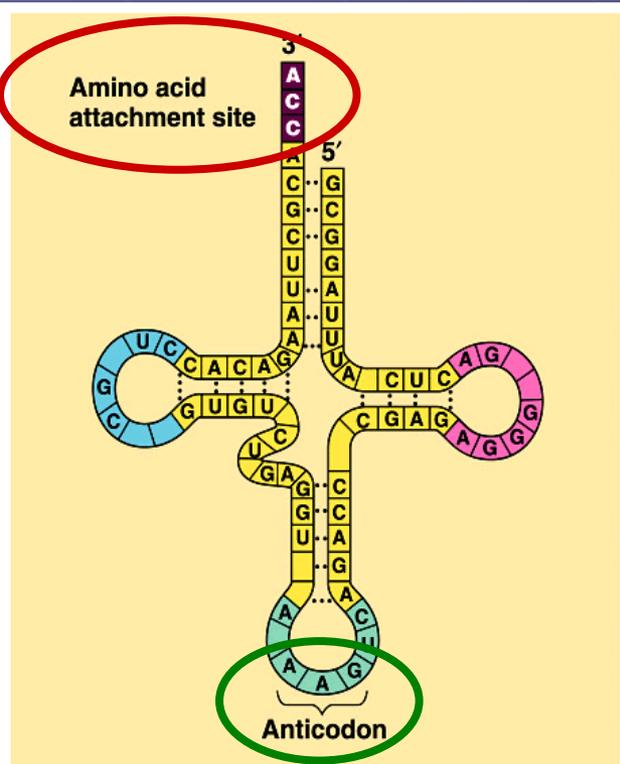


Figure 10.11B, C

tRNA delivers the correct Amino Acid

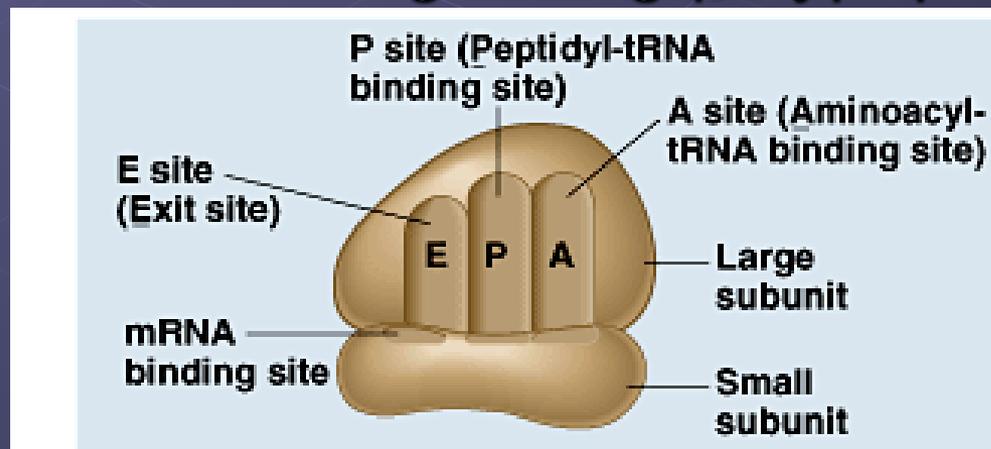
● “Clover leaf” structure

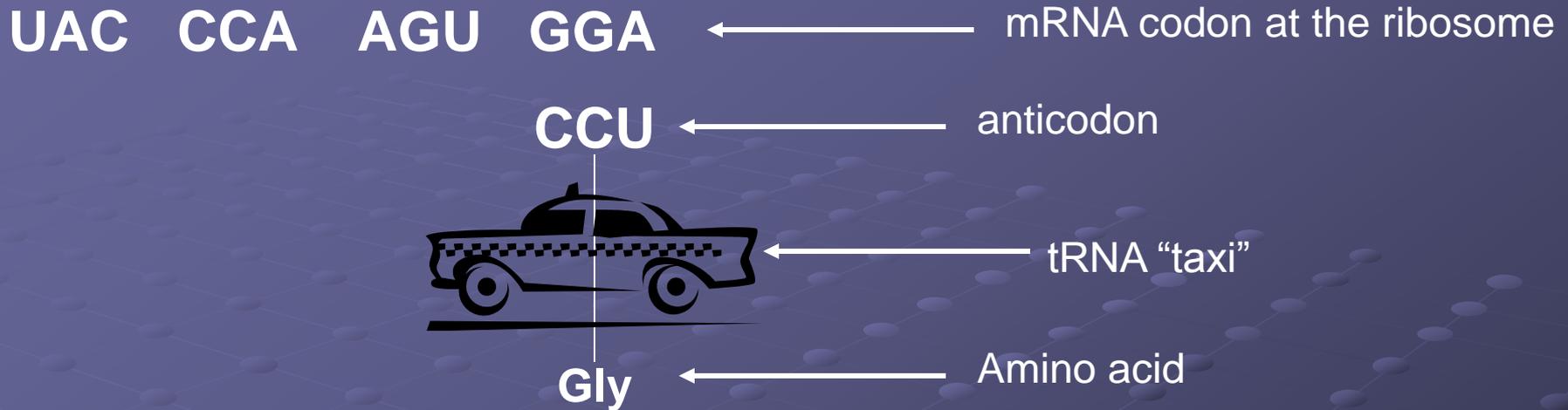
- **anticodon** on “clover leaf” end
- **amino acid** attached on 3' end



Ribosomes

- tRNA binds to the mRNA CODON with its matching 3-letter **ANTICODON**
- tRNA first Attaches to the ribosome at the **A** site (**A = Attach**)
- The ribosome **slides** down the mRNA to read the next codon
- This also causes the tRNA to slide from the A site to the **P** site (**P = Pass** the growing polypeptide)





- tRNA releases its AA "passenger" which bonds to other AA to make a polypeptide
- the empty tRNA leaves the ribosome to pick up other AA passengers
- The protein is completed when a STOP codon is read

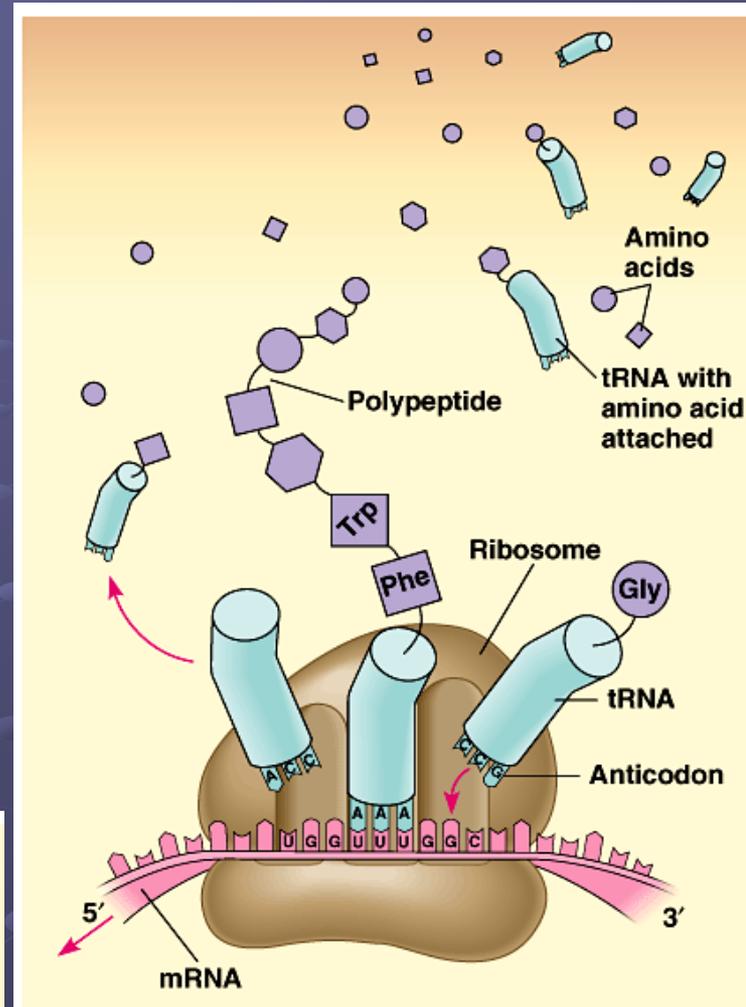
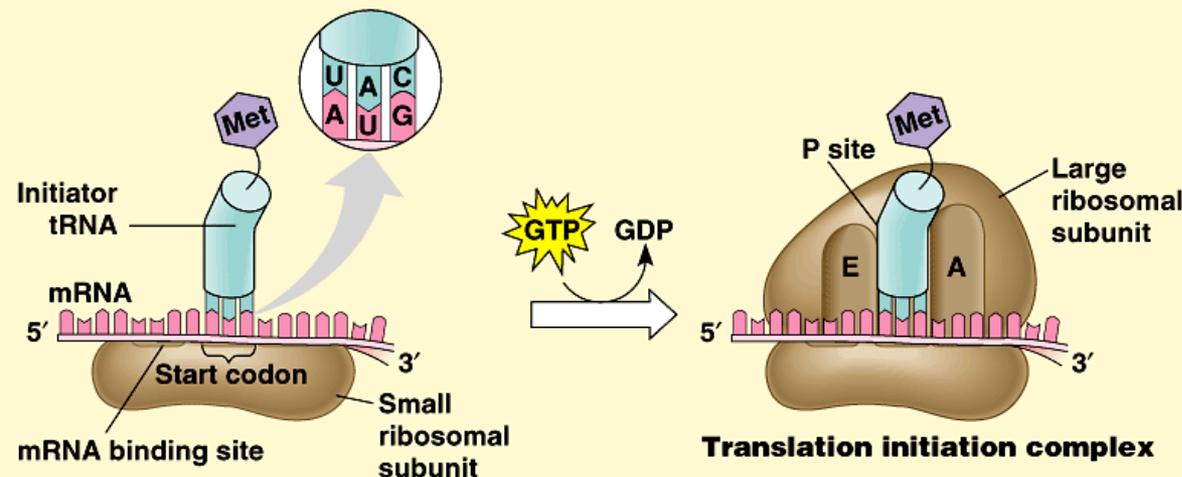
Building a polypeptide

● Initiation

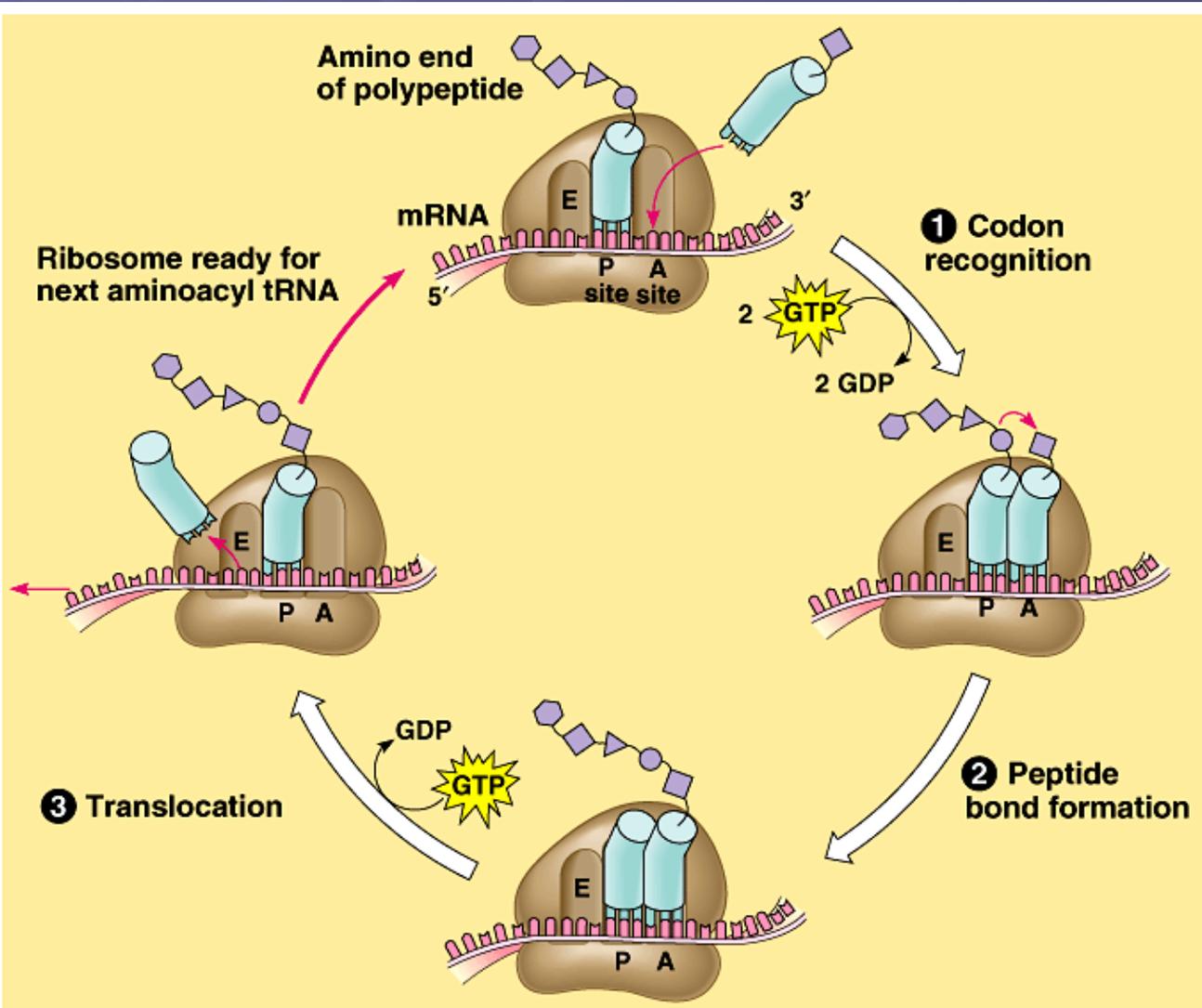
- brings together mRNA, ribosome subunits, proteins & initiator tRNA

● Elongation

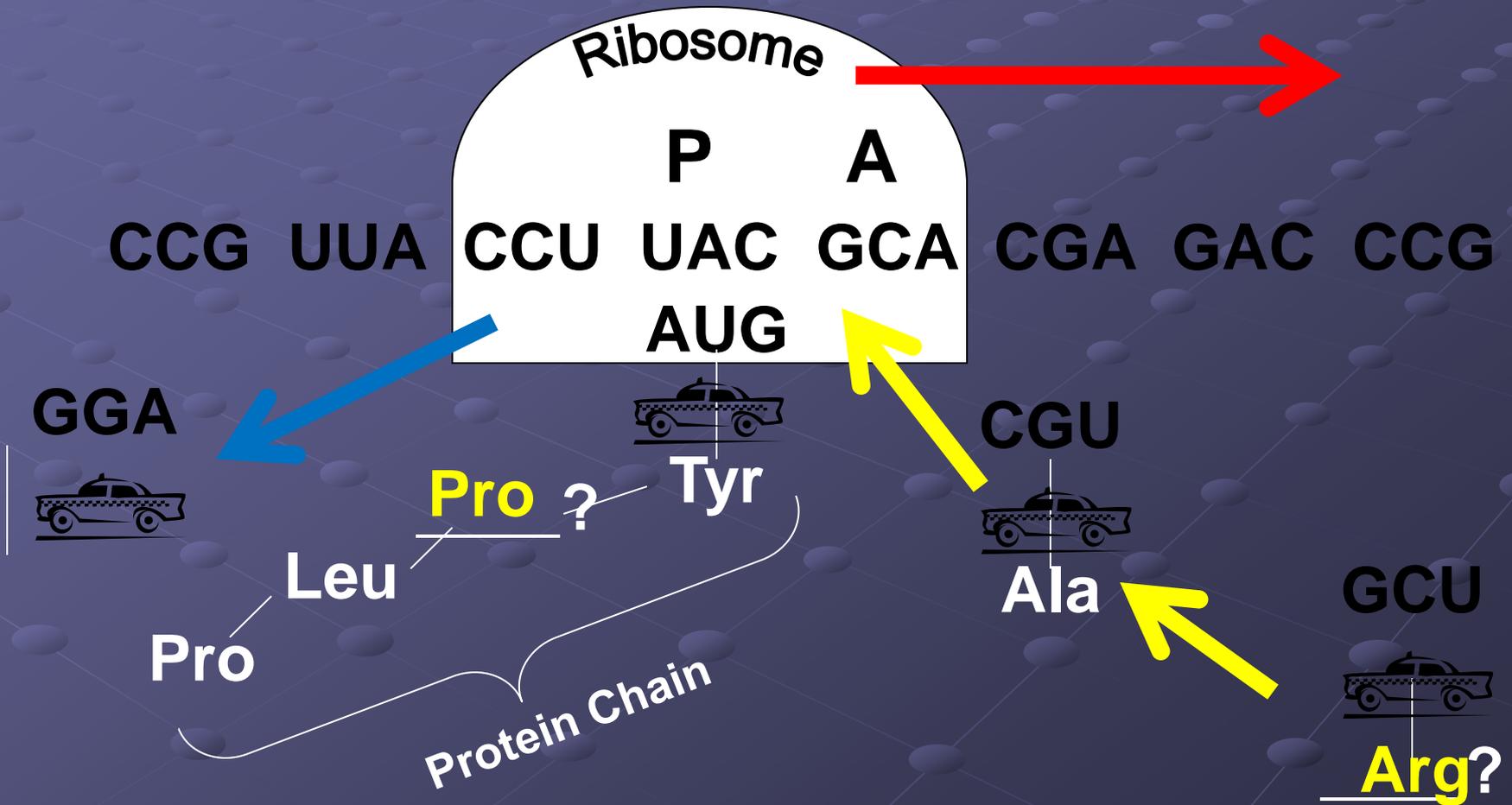
● Termination



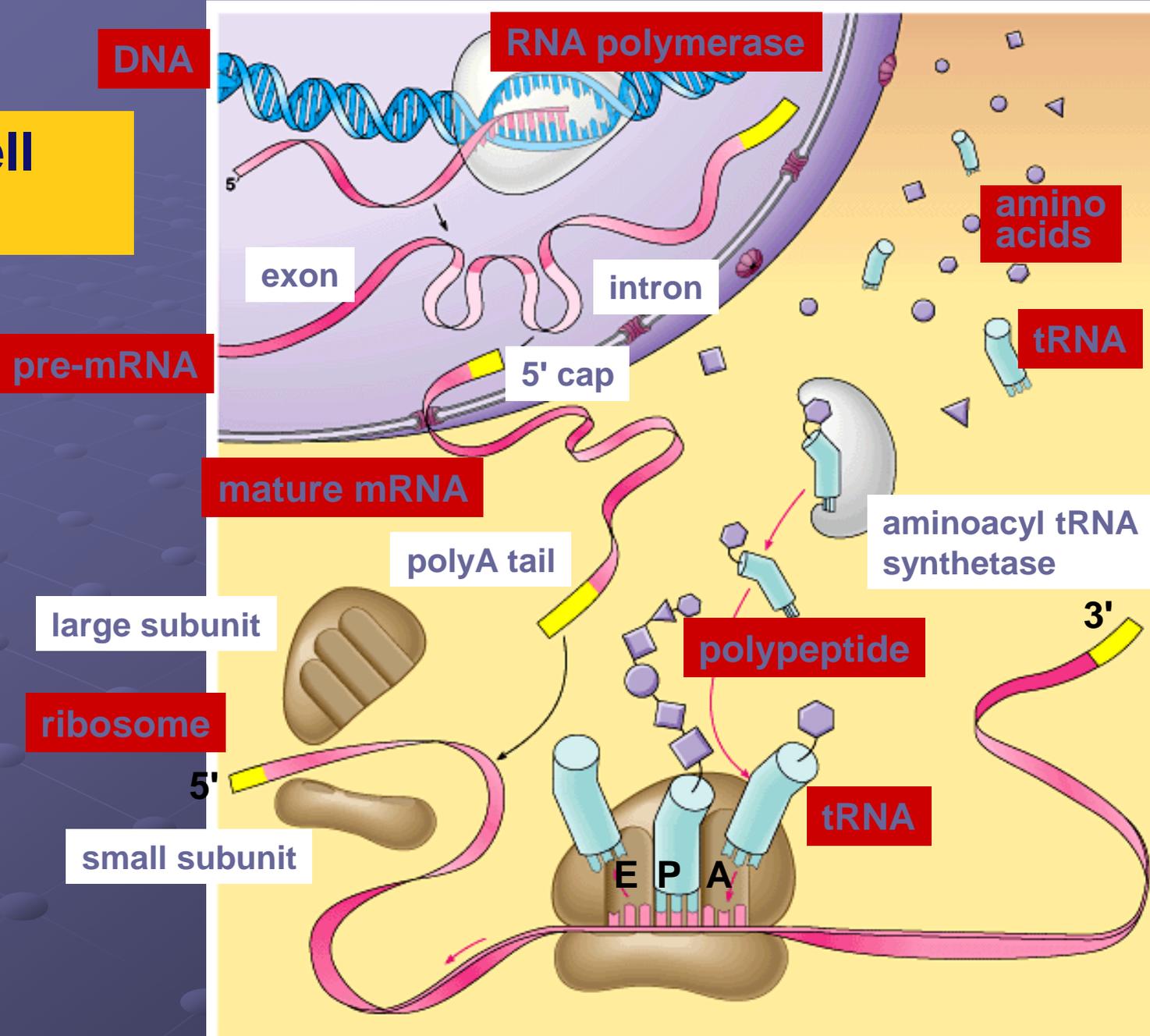
Elongation: growing a polypeptide



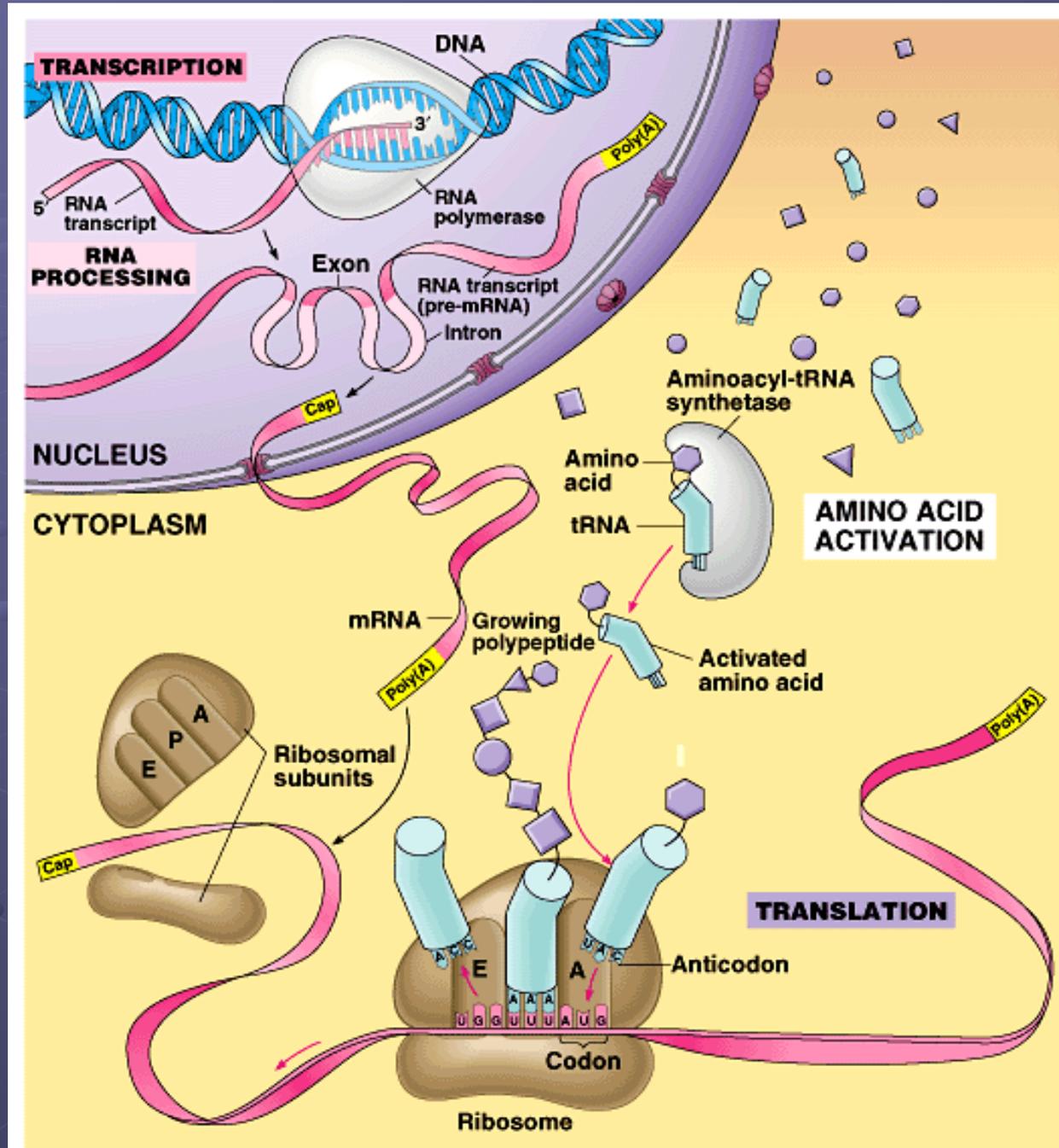
Translation



Can you tell the story?



Put it all together...



Mutation = changing the letter codes of a gene

A. Mutation **CAUSES** :

1. During DNA **Replication** :

- Base-pairing mistakes can happen but most are **fixed**

2. Exposure to powerful **Chemicals** :

- **Cigarette** smoke, smokeless tobacco, exhaust from burning petroleum fuels, pesticides, herbicides, alcohol, sawdust from CCA lumber, paints, mineral spirits & oils, asbestos, etc.

3. Exposure to powerful **Radiation** :

- **Gamma rays** (nuclear fuel & bombs)
- **X-rays** (at hospitals & dental offices)
- **UV rays** (from sunlight and tanning beds)

B. Mutation Types

- | | | Original gene | | <u>mutated</u> gene |
|----|---|---------------|---|---------------------|
| 1. | <u>Addition</u> = add 1+ letters | TAGACAT | → | TAGACCAT |
| 2. | <u>Deletion</u> = lose 1+ letters | TAGACAT | → | TGACAT |
| 3. | <u>Substitution</u> = switch 1+ letters | TAGACAT | → | TAGAGAT |

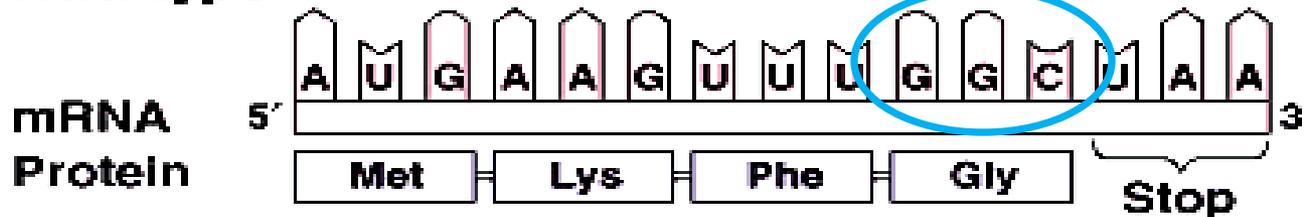
C. Mutation Effects: Was the protein produced any different shape than the original?

1. NO effect = protein remains the same → same shape
2. Small effect = protein has 1+ different AA → Small shape change
3. BIG effect = protein has many different AA → Big shape change

Mutation **Effects**: None or Small or BIG?

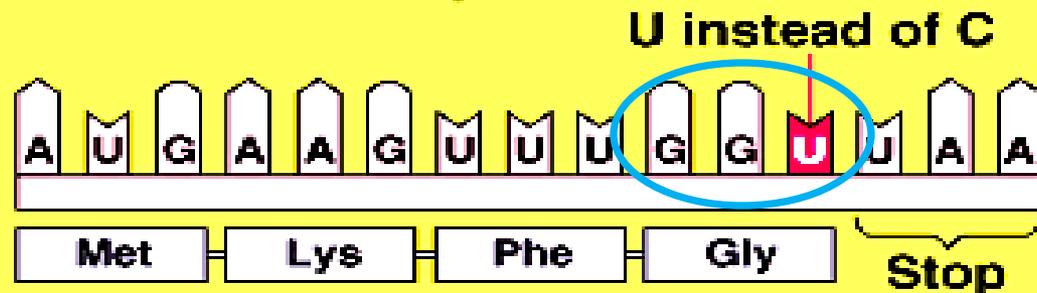
- 1) **Silent** mutation = changes a CODON but still specifies the SAME AA = NO Effect

Wild type



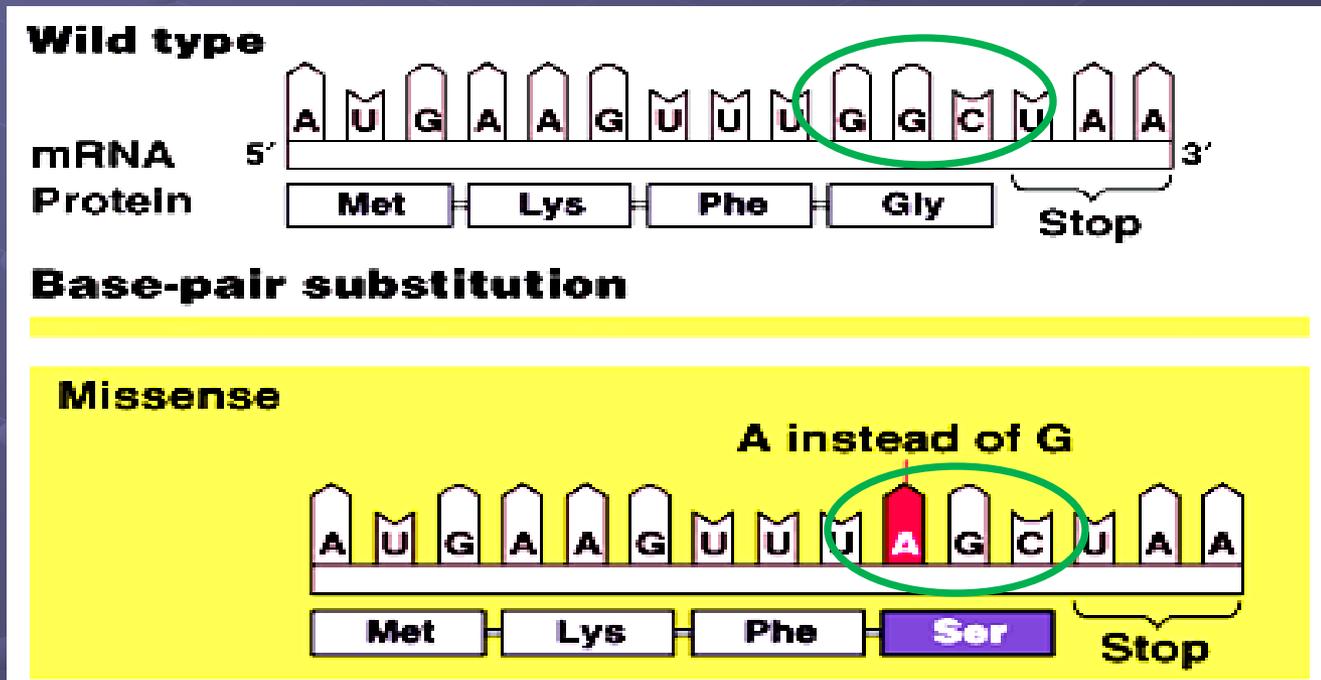
Base-pair substitution

No effect on amino acid sequence



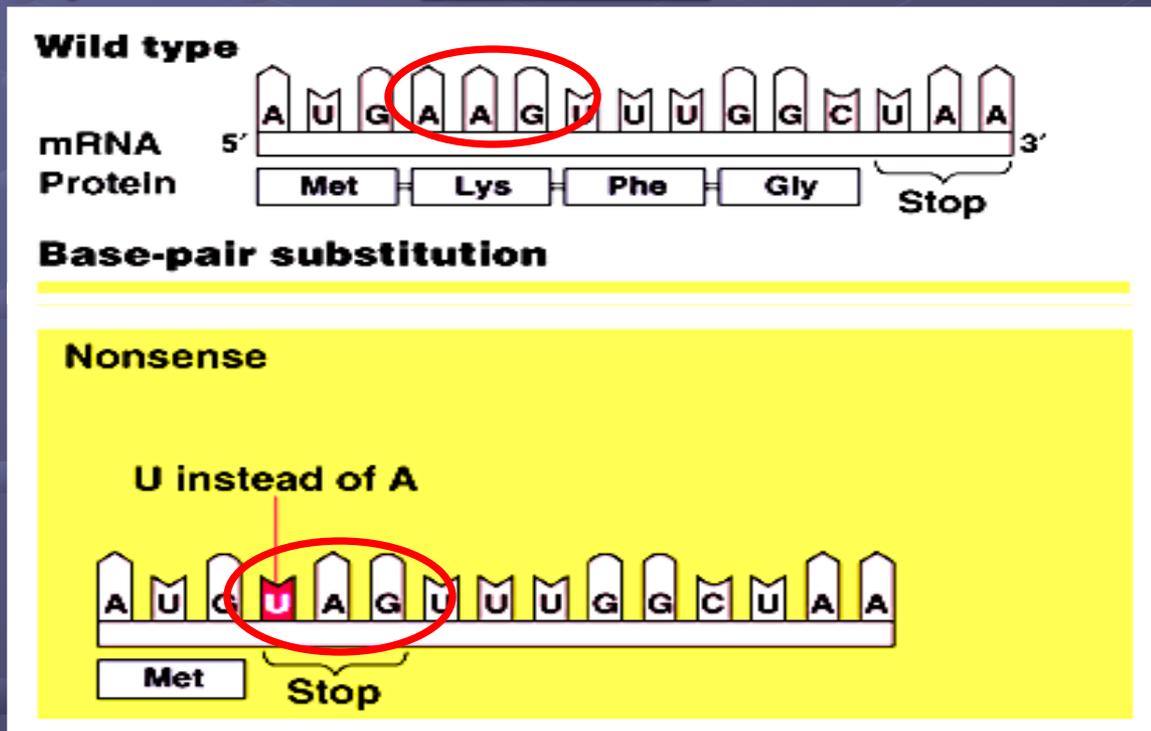
Mutation **Effects**: None or Small or BIG?

2) **Missense** mutation = changes a CODON that will specify a DIFFERENT AA = small effect



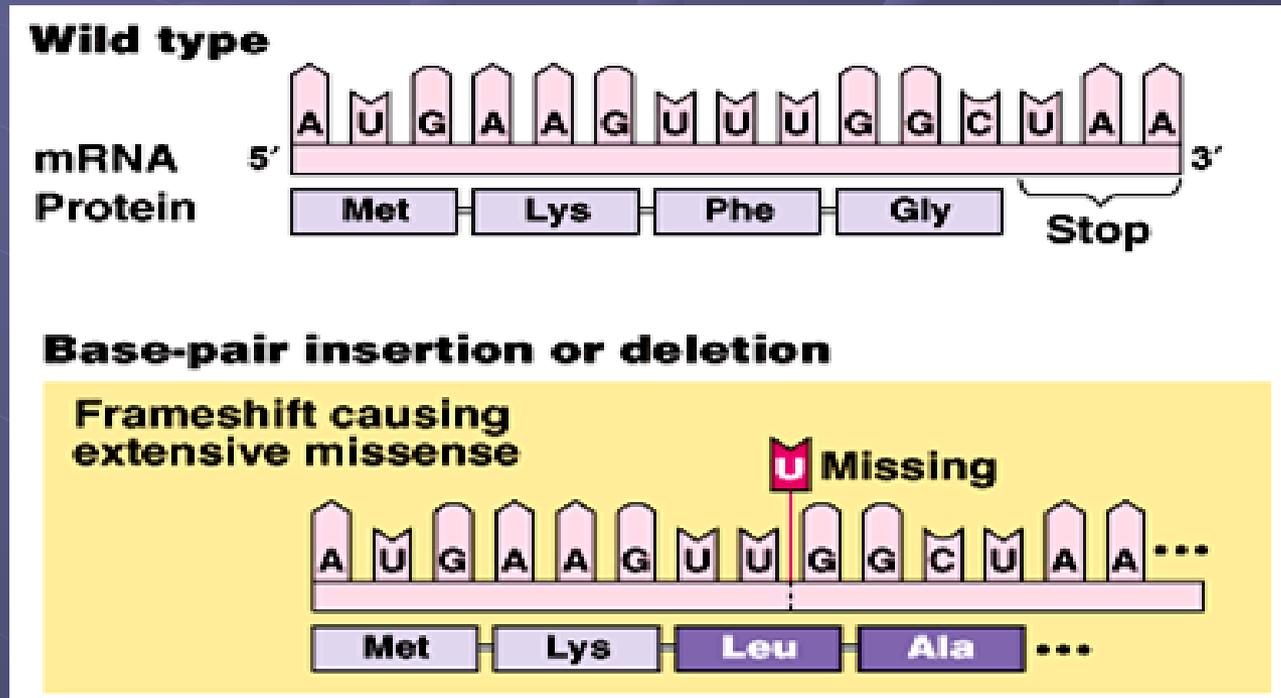
Mutation **Effects**: None or Small or BIG?

3) **Nonsense** mutation = changes a CODON to specify a **STOP** signal in the middle of the protein = **BIG** effect



Mutation **Effects**: None or Small or BIG?

4) **Frameshift** mutation = 1+ bases add or lost causes the triplet reading frame to be SHIFTED over = BIG effect



Mutation Conclusions:

- Any protein different than the original probably will NOT fold into the same 3-D SHAPE = NOT FUNCTION = cause a health problem
- Most mutations have a Negative (-) effect that lowers an organism's chance for survival
Ex.
- Some mutations have a Positive (+) effect for an organism in a certain environment that helps them better survive the challenges of life
Ex.
- Accumulating DNA mutations often lead to diseases like Cancer
- Only DNA mutations in eggs and sperm cells can be inherited
 - These inherited mutations often result in a variety of genetic diseases

Mutation Conclusions:

- Some viruses mutate quickly because mistakes during replication of their genetic material are NOT fixed by proofreading repair enzymes
- This leads to new strains of disease-causing viruses that sometimes emerge to cause major health epidemics.
- This happens because people have little immunity to the viruses' newly-shaped proteins

