

# General Biology

## Ch 10 and 11: The Structure and Function of DNA and How Genes are controlled



**“Translation and Mutation”**

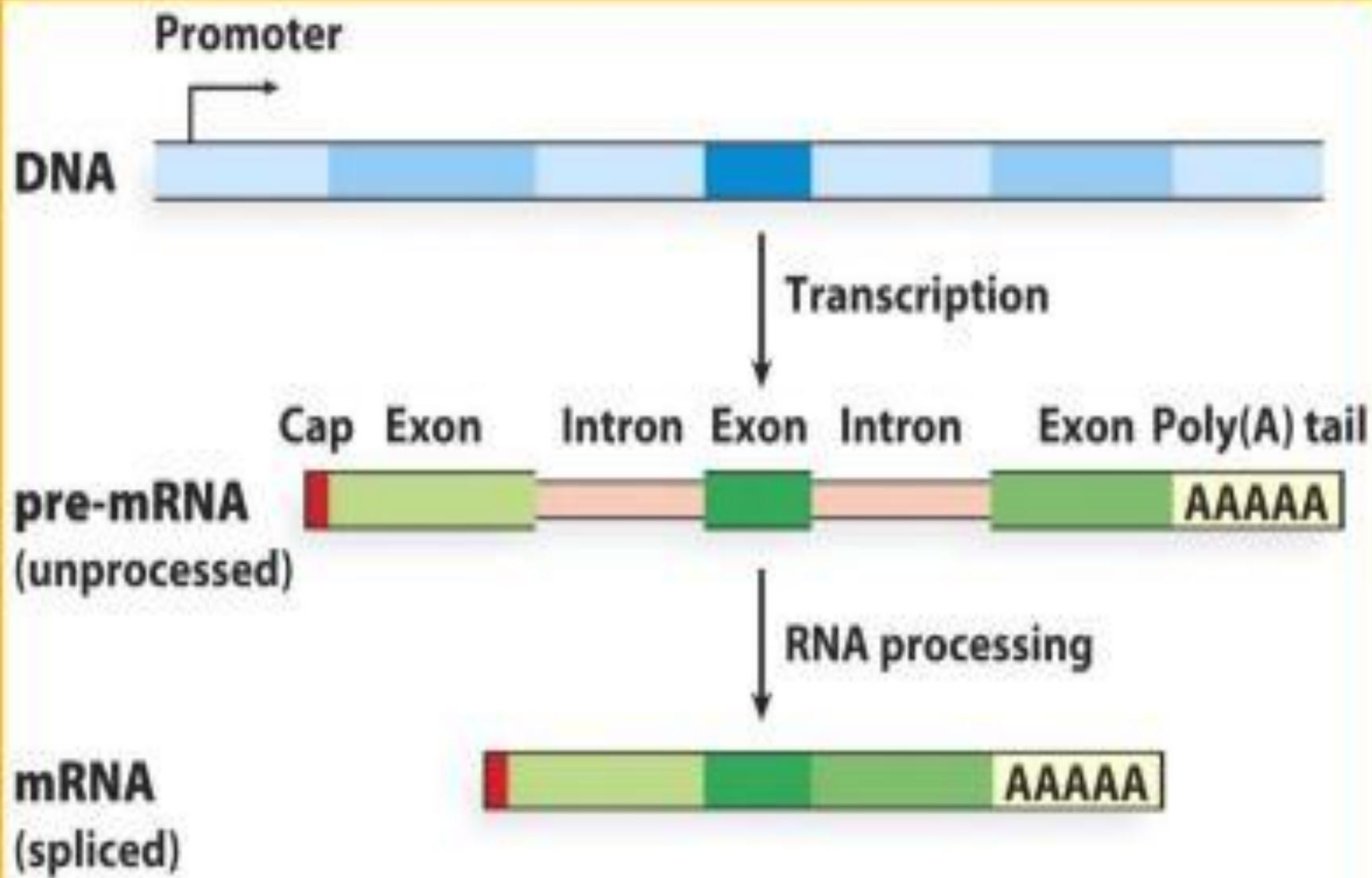


Figure 16-7  
*Molecular Biology: Principles and Practice*  
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# **Food for thought!!!!**

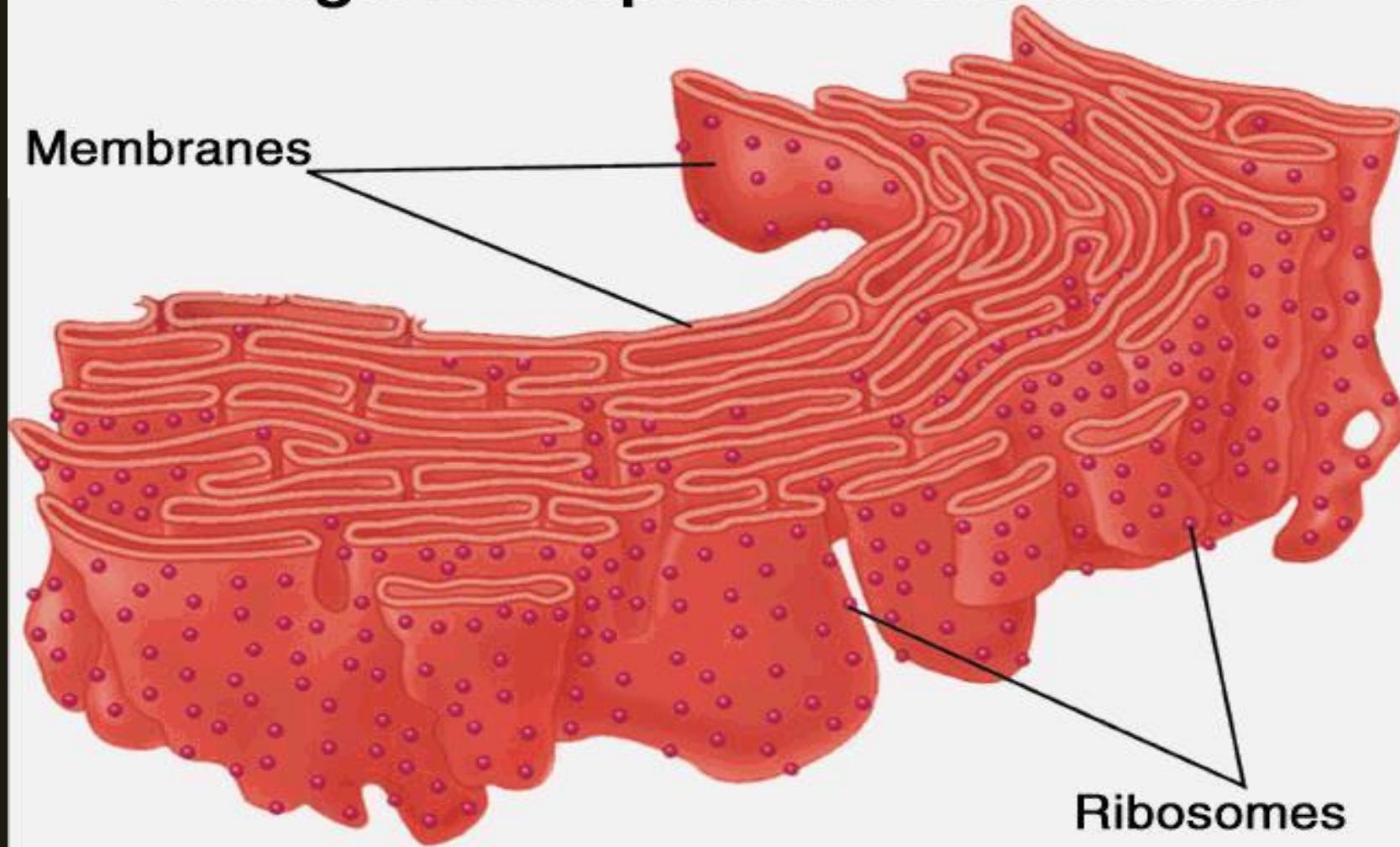
- What is the advantage of having introns?
- What happens to the introns after they are spliced off?
- Why do cells contain nucleases that chop off nucleotides from 5' or 3' end?
- Why do you think prokaryotic RNAs don't have a cap or tail?
- What are the so called 'junk' sequences in DNA?
- How can you regulate transcription?

# Eukaryotic or Prokaryotic cell?

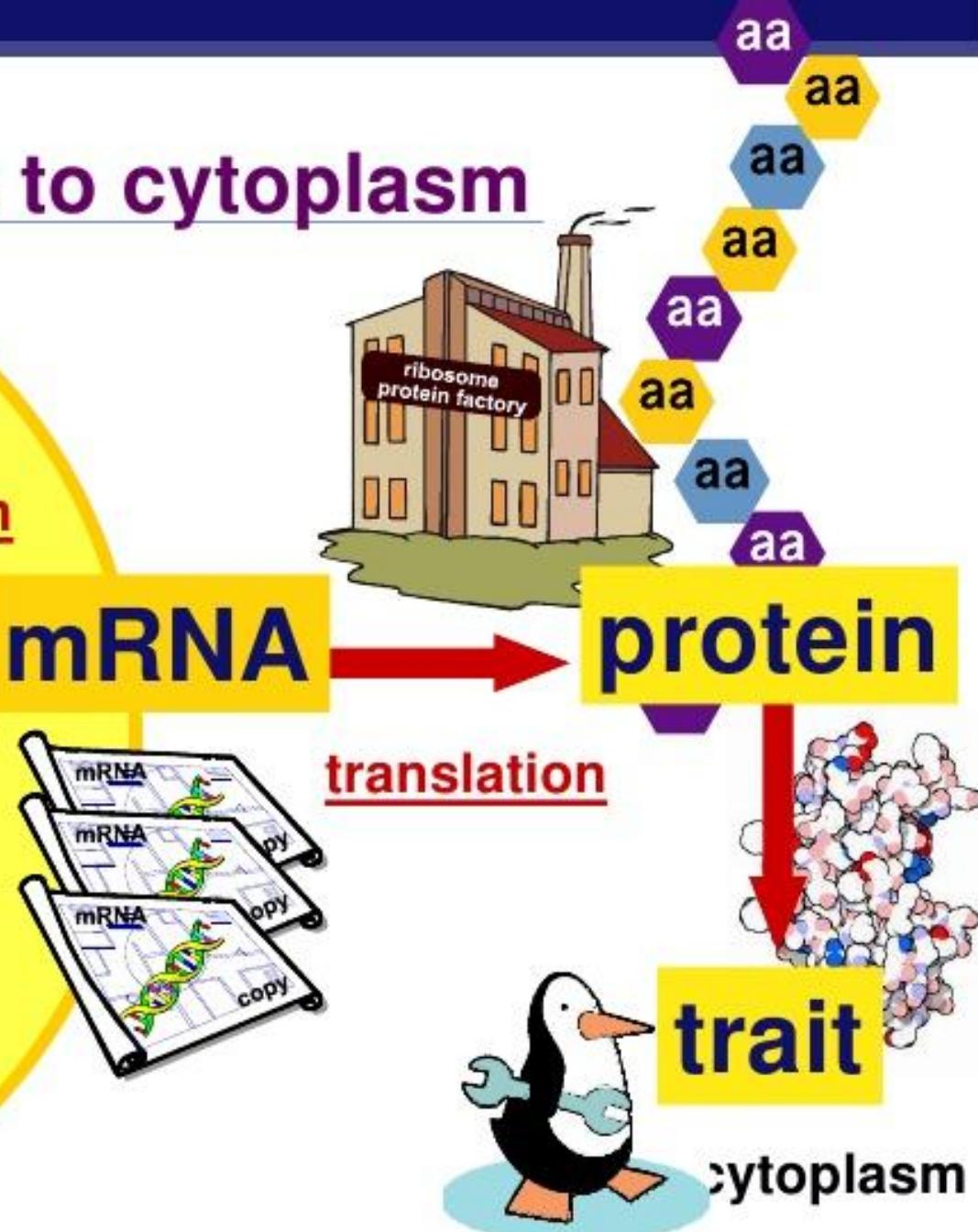
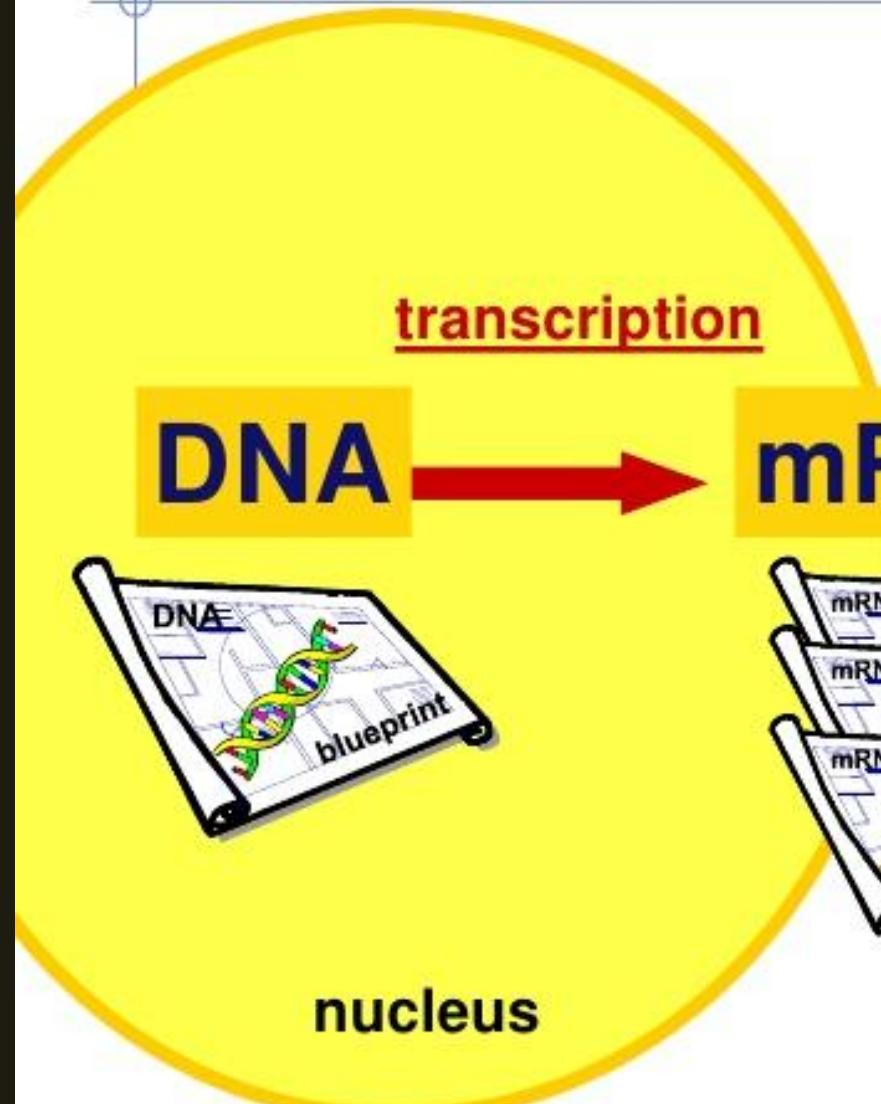


# The Destination

## Rough Endoplasmic Reticulum



# From nucleus to cytoplasm

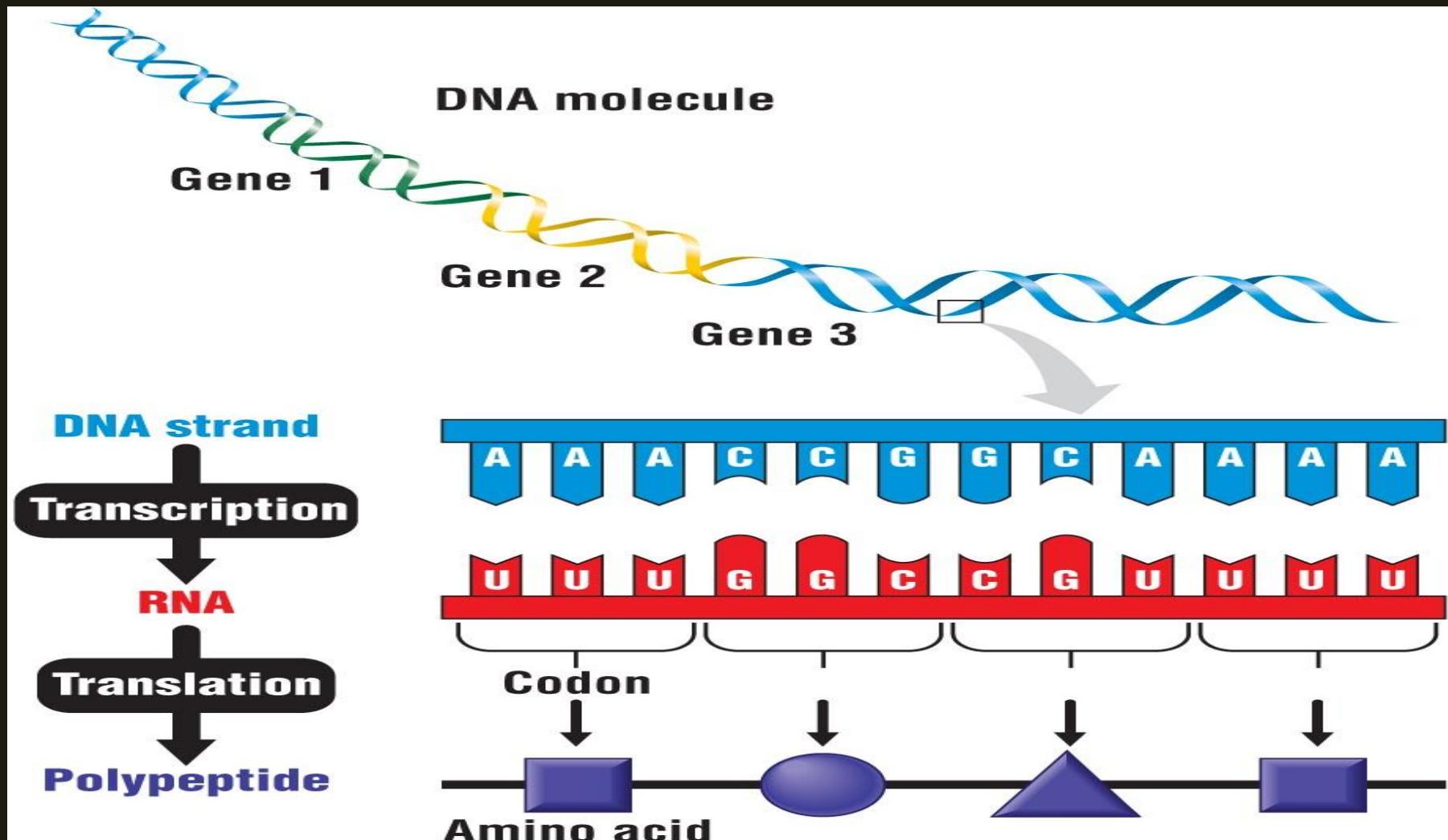


for what is considered  
to be best in any related  
point of view.

## **Translation**

another language; in  
language foreign to the  
written communication  
that is thought

# Protein Translation



**Codons are triplets**

DNA

5' — ATG GGC TAC CCC TGC CTG — 3'

3' — TAC CCG ATG GGG ACG GAC — 5'

↓  
Transcription

mRNA

5' — |AUG|GGC|UAC|CCC|UGC|CUG| — 3'

↓  
Translation

Protein N-terminus

met      gly      tyr      pro      cys      leu  
M          G          Y          P          C          L

C-terminus

# Deciphering the genetic code...



## Second base

	U	C	A	G		
U	UUU UUC UUA UUG	UCU UCC UCA UCG	UAU UAC UAA UAG	Tyrosine (Tyr) Stop Stop	UGU UGC UGA UGG	Cysteine (Cys) Stop Tryptophan (Trp)
C	CUU CUC CUA CUG	CCU CCC CCA CCG	CAU CAC CAA CAG	Histidine (His) Glutamine (Gln)	CGU CGC CGA CGG	
A	AUU AUC AUA AUG Met or start	ACU ACC ACA ACG	AAU AAC AAA AAG	Asparagine (Asn) Lysine (Lys)	AGU AGC AGA AGG	Serine (Ser) Arginine (Arg)
G	GUU GUC GUA GUG	GCU GCC GCA GCG	GAU GAC GAA GAG	Aspartic acid (Asp) Glutamic acid (Glu)	GGU GGC GGA GGG	Glycine (Gly)

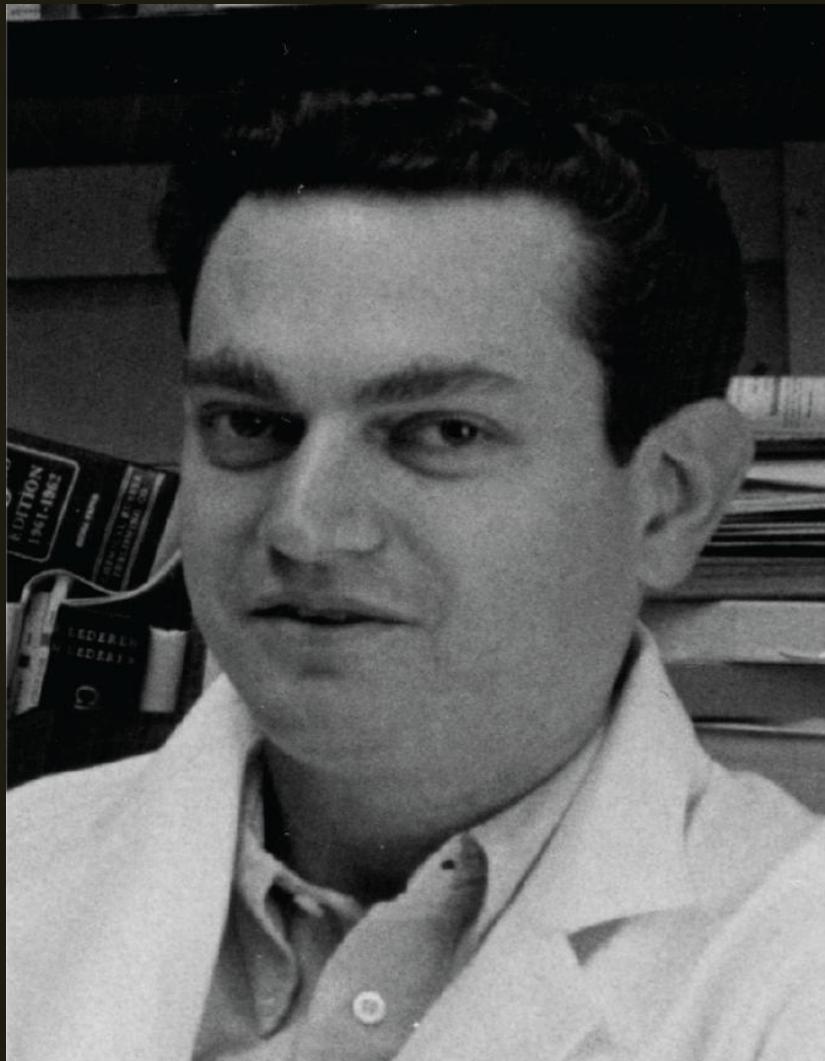
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**Note the start and stop codon**  
**(Nonsense codon)**

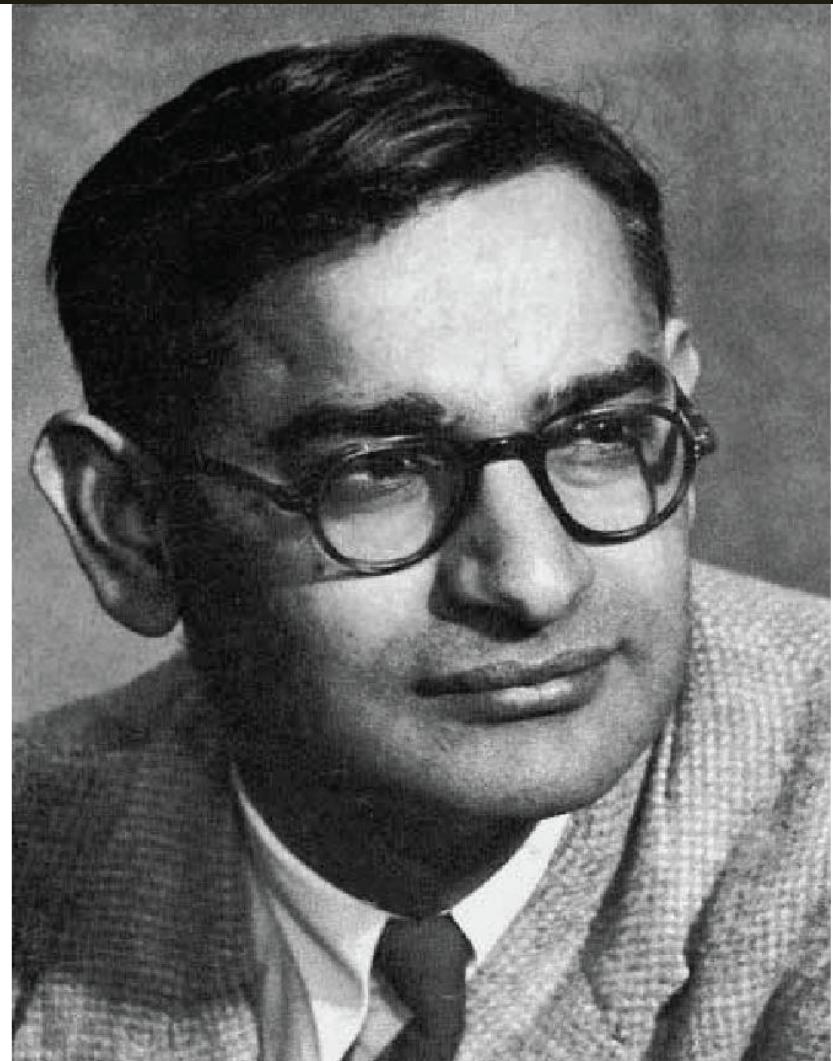


**Why codons constitute “3” nucleotides and not one or two?**

# The *Nobel* Prize in Physiology or Medicine 1968



Marshall Nirenberg



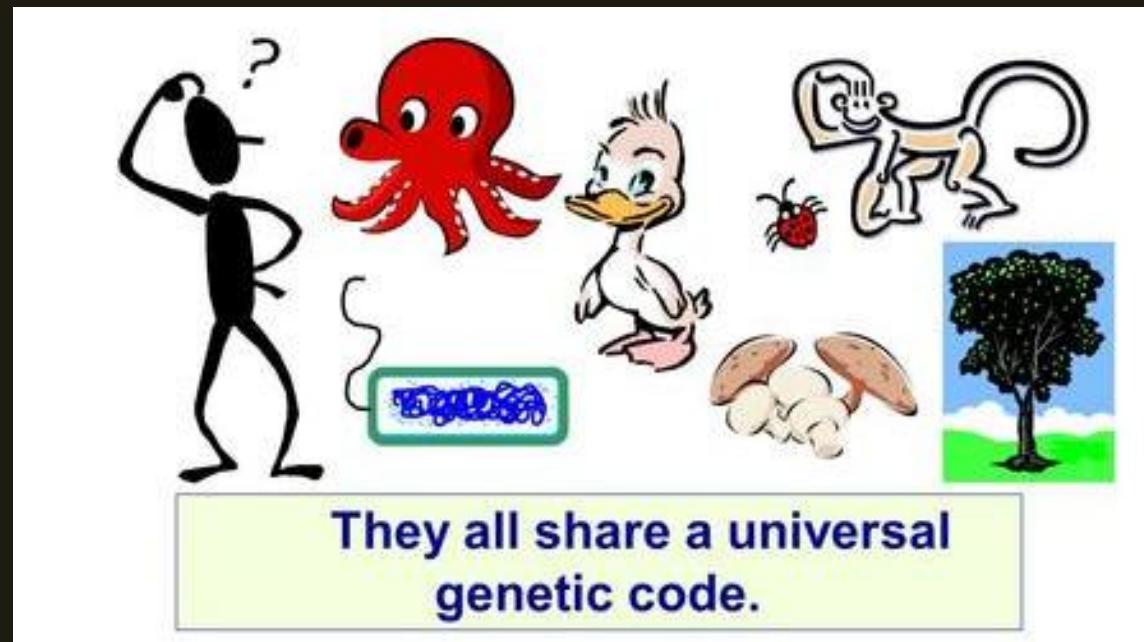
Har Gobind Khorana

# **Features of Genetic Code**

# The Genetic Code

- How many codons combinations can you make from four bases?
- 64
- Of the 64 triplets,
  - 61 code for amino acids and
  - 3 are stop codons, instructing the ribosomes to end the polypeptide.
- A given RNA triplet always specifies a given amino acid.

The genetic code is universal,  
shared by organisms from the  
simplest bacteria to the most  
complex plants and animals!!!!



What can be its advantage to us?

# Degeneracy of the Genetic Code

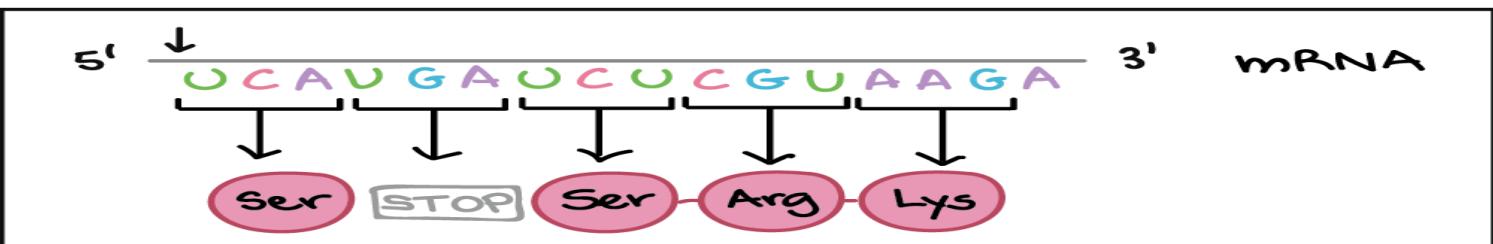
		Second base						
		U	C	A	G			
First base	U	UUU UUC UUA UUG	UCU UCC UCA UCG	UAU UAC UAA UAG	UGU UGC UGA UGG	Cysteine (Cys) Stop Tryptophan (Trp)		
	C	CUU CUC CUA CUG	CCU CCC CCA CCG	CAU CAC CAA CAG	CGU CGC CGA CGG	Histidine (His) Proline (Pro) Glutamine (Gln)		
	A	AUU AUC AUA AUG	ACU ACC ACA ACG	AAU AAC AAA AAG	AGU AGC AGA AGG	Arginine (Arg) Serine (Ser) Arginine (Arg)		
	G	GUU GUC GUA GUG	GCU GCC GCA GCG	GAU GAC GAA GAG	GGU GGC GGA GGG	Asparagine (Asn) Threonine (Thr) Lysine (Lys) Alanine (Ala) Aspartic acid (Asp) Glutamic acid (Glu)		
						Glycine (Gly)		
Third base								

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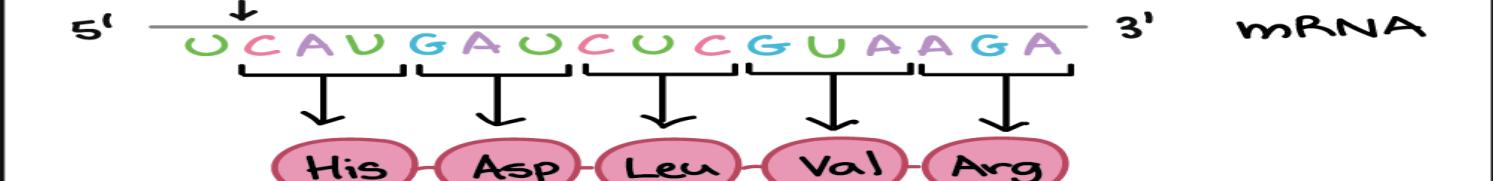
Although each codon is specific for only one amino acid (or one stop signal), the genetic code is described as degenerate, or redundant, because a single amino acid may be coded for by more than one codon.

# The Reading Frame: Genetic is non-overlapping

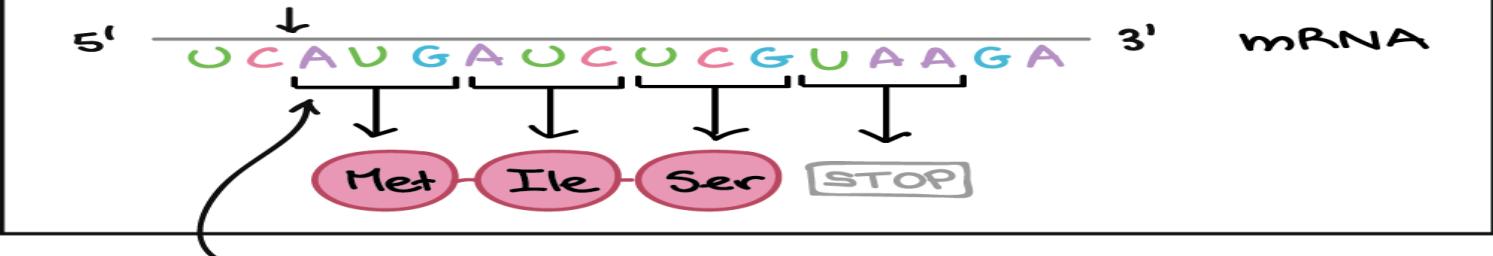
FRAME 1



FRAME 2



FRAME 3



Start codon's position ensures that this frame is chosen

How does a cell know which of these protein to make?

# How do you know where to start off.....

genomic DNA sequence:

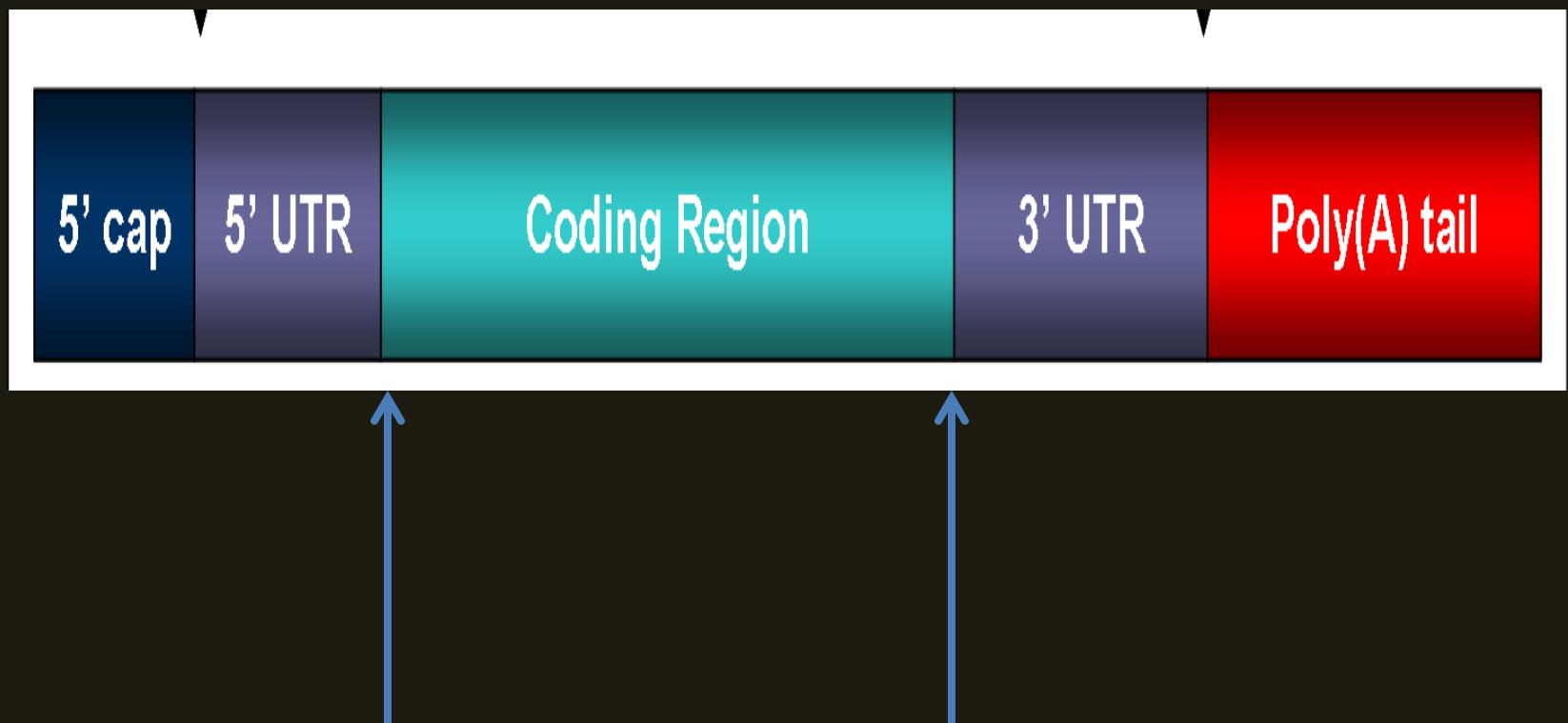
5'-AGCTCATGTGCGAGTCCTGACGCTGACTAGG-3' top strand  
3'-TCGAGTACACGCTCAGGACTGCGACTGATCC-5' bottom strand



mature mRNA sequence:

5'-7-methylGcap\*UCAUGUGCGAACGCUGACUAGGAAAAAAA....-3'

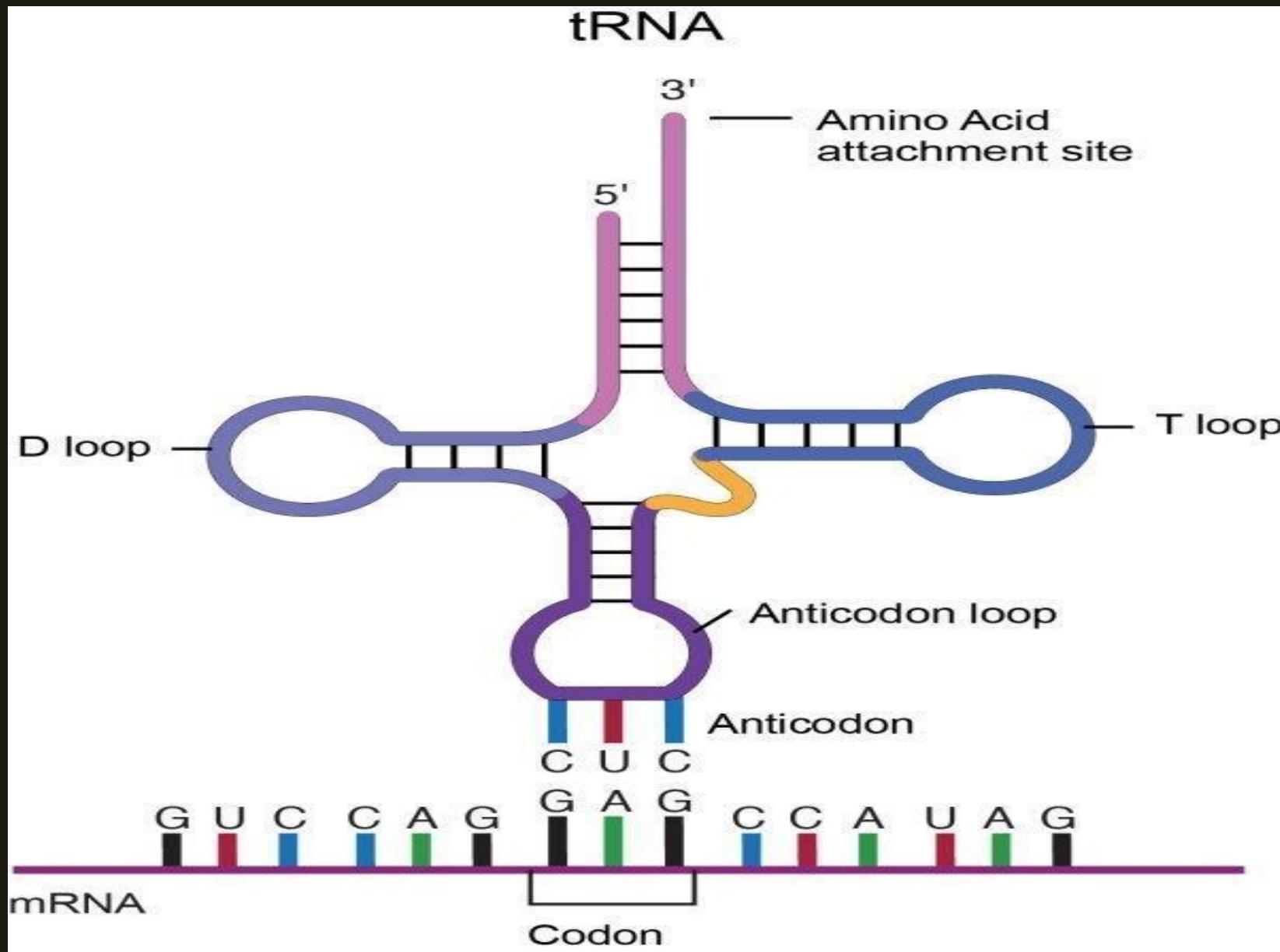
# The Concept of UTR- Un-translated Region in mRNA



# Translation: The Players

- Translation is a process of encoding protein from an mRNA.
- The ingredient required for translation
  - mRNA produced by transcription.
  - ribosomes and
  - transfer RNA
  - enzymes and
  - sources of chemical energy, such as ATP

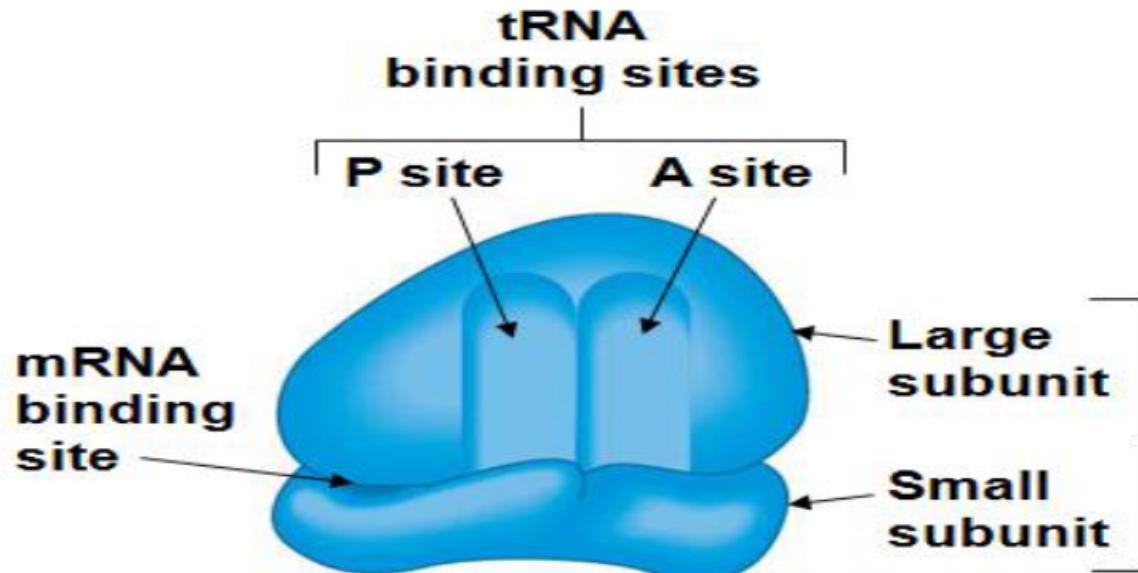
# Player-1 (tRNA)

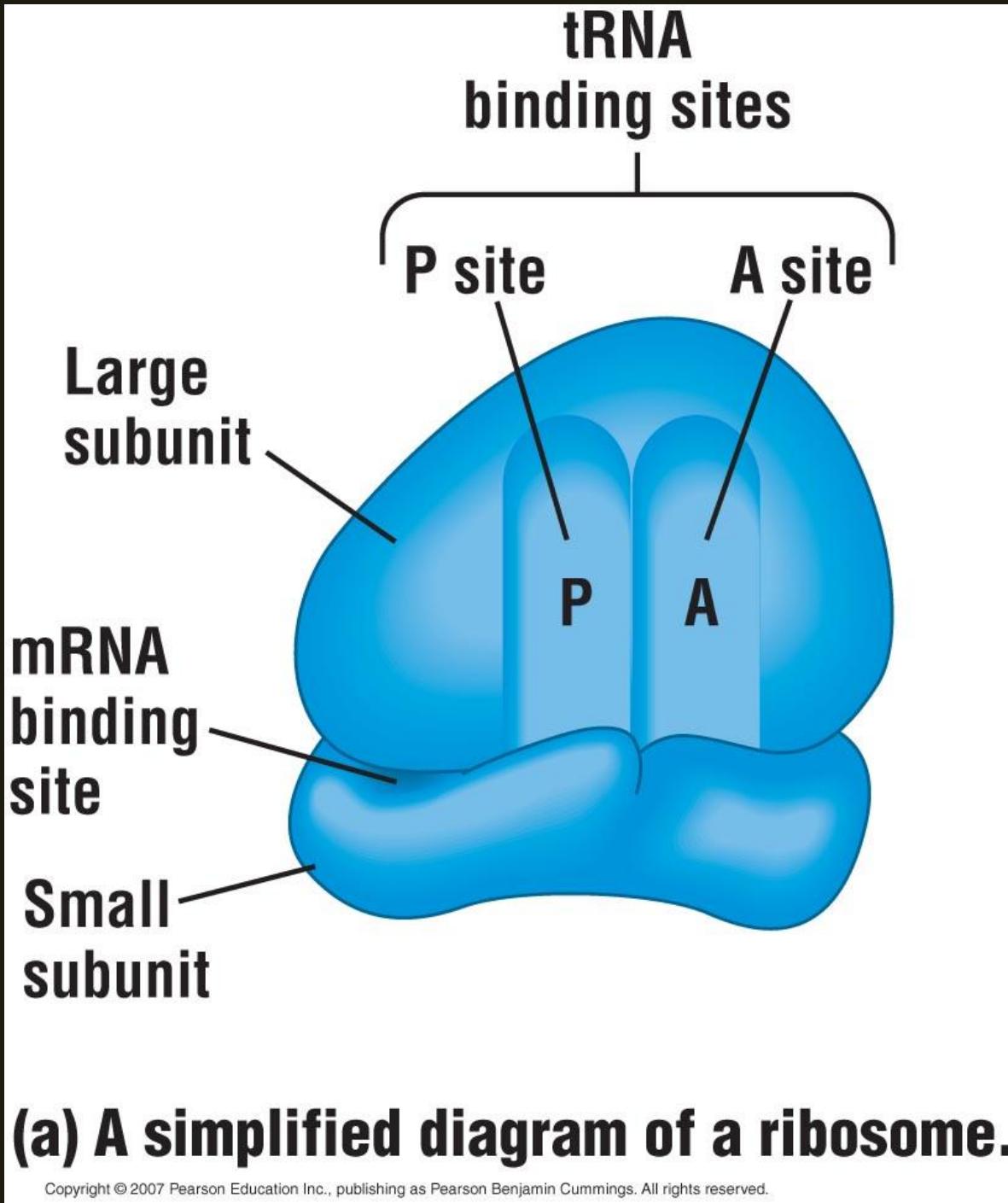


# Player-2 (Ribosomes)

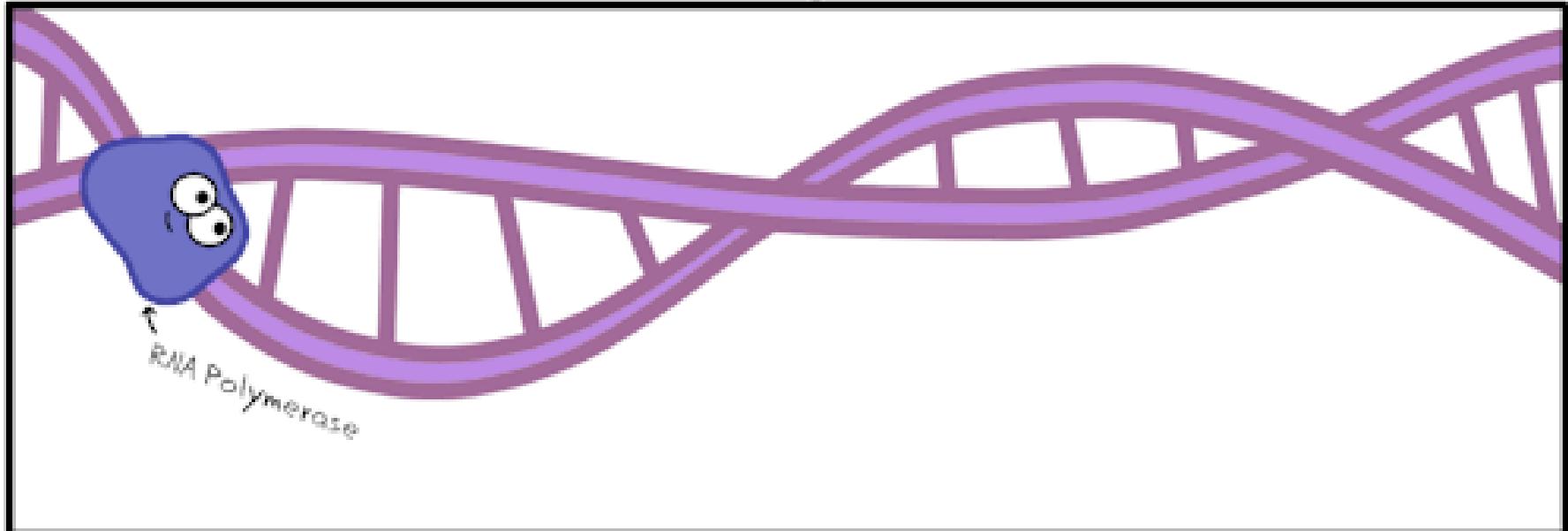
## Ribosomes

- Ribosomes are the organelles in the cytoplasm that
  - coordinate the functioning of mRNA and tRNA and
  - actually make polypeptides.
- A ribosome consists of two subunits – large & small.
- Each subunit is made up of
  - proteins and
  - **ribosomal RNA (rRNA)**.

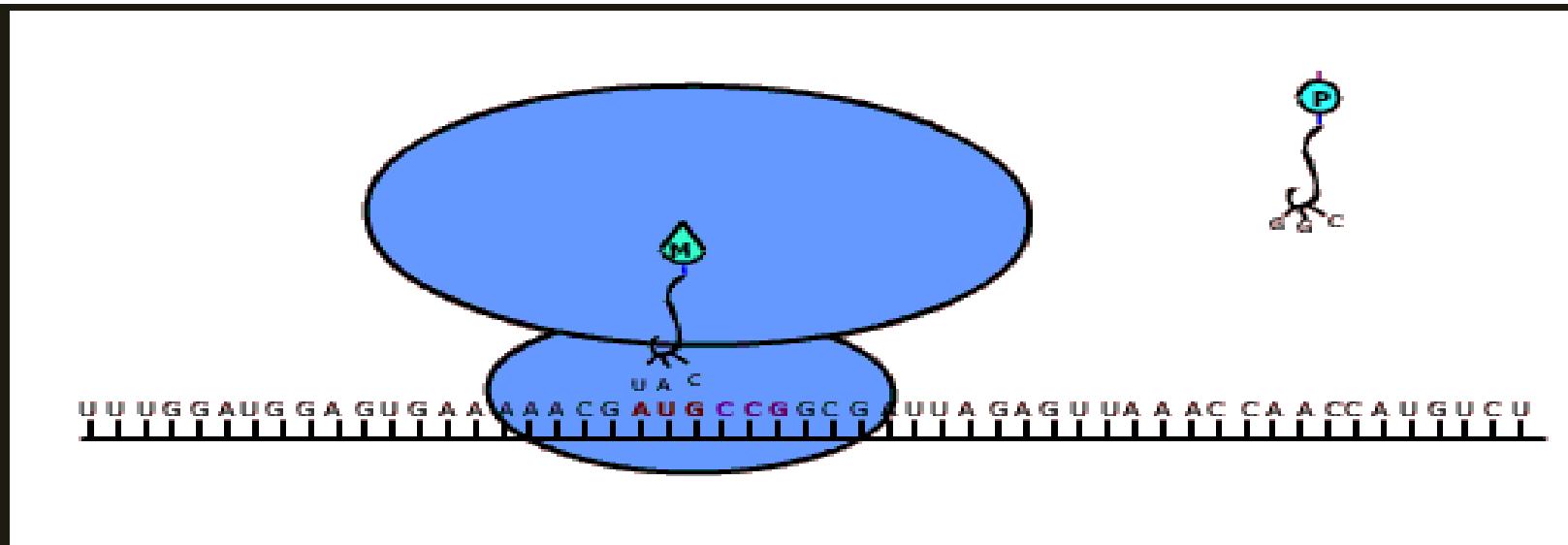


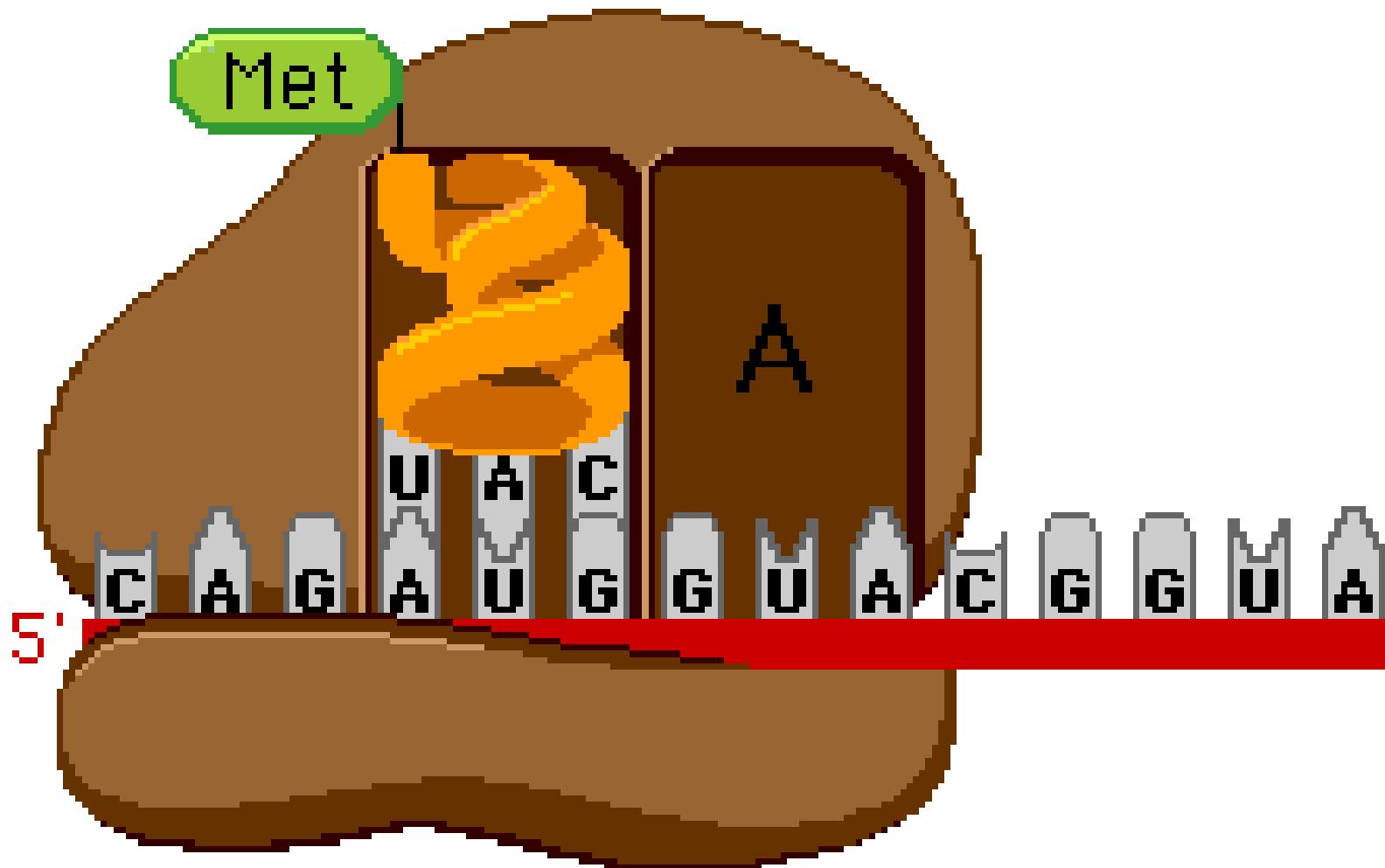


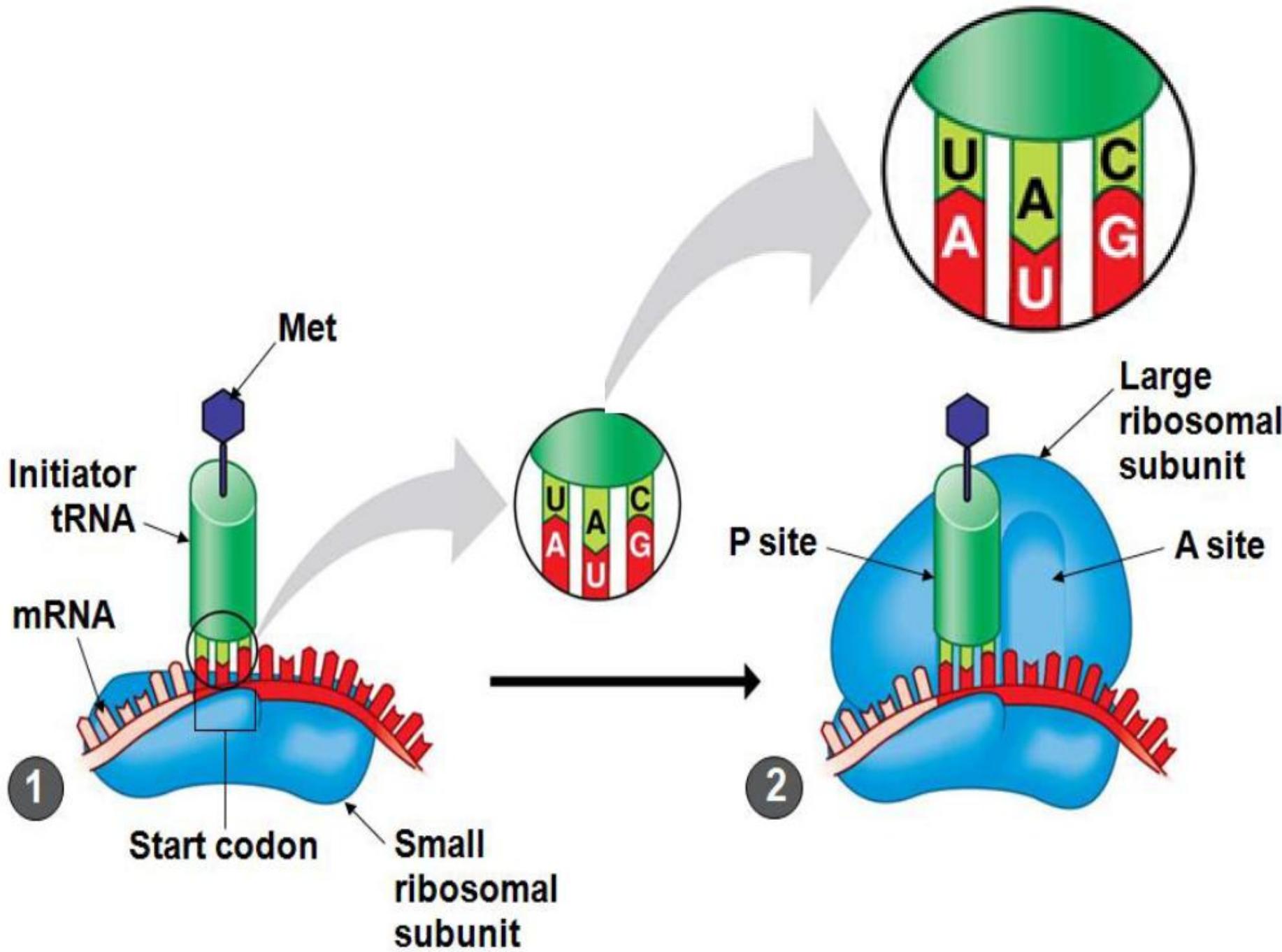
# Protein Synthesis



## Step 1: Transcription

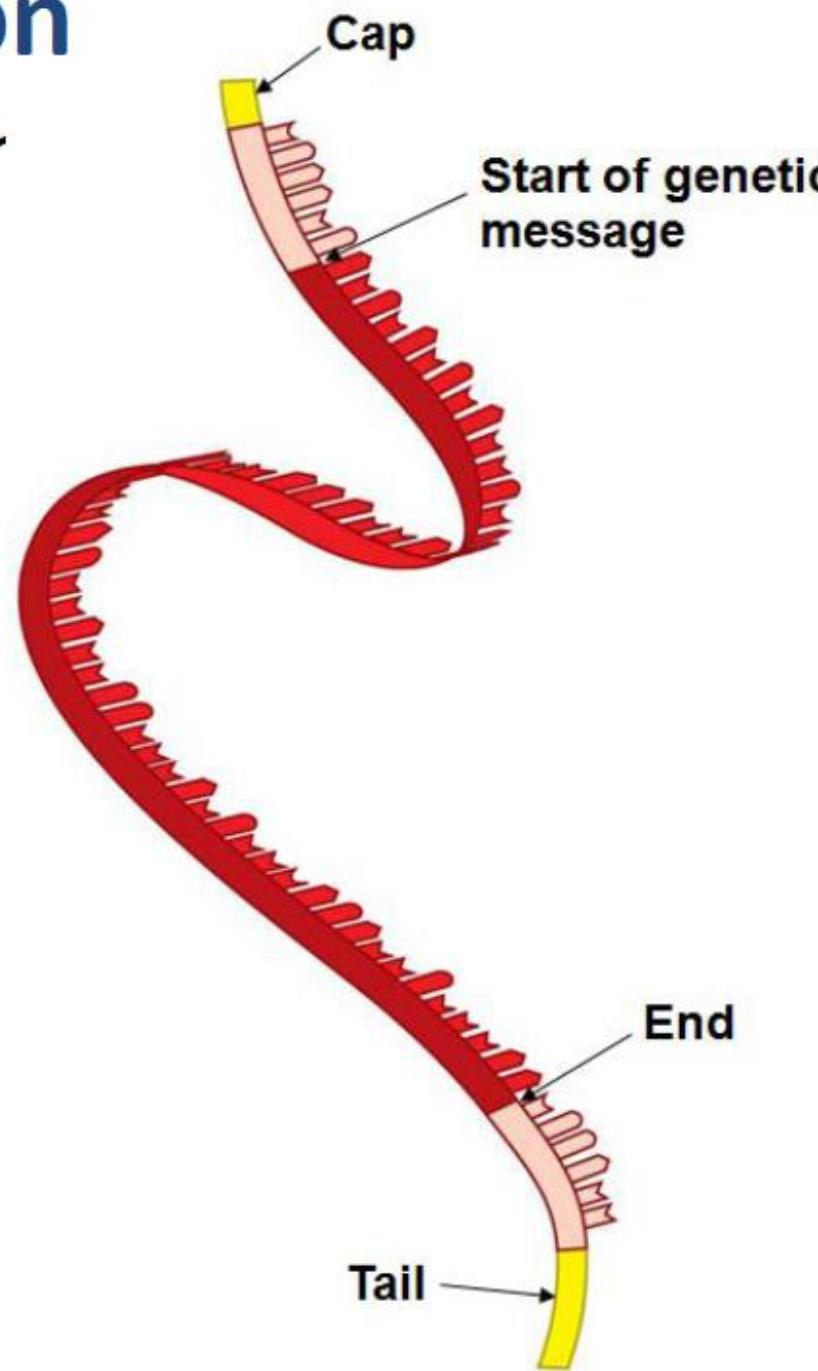






# Initiation

- Initiation brings together
  - mRNA - binds to a small ribosomal subunit,
  - the tRNA carrying first amino acid attaches to **start codon** and
  - large subunit of the ribosome.
- cap and tail help the mRNA bind to the ribosome.



# Elongation

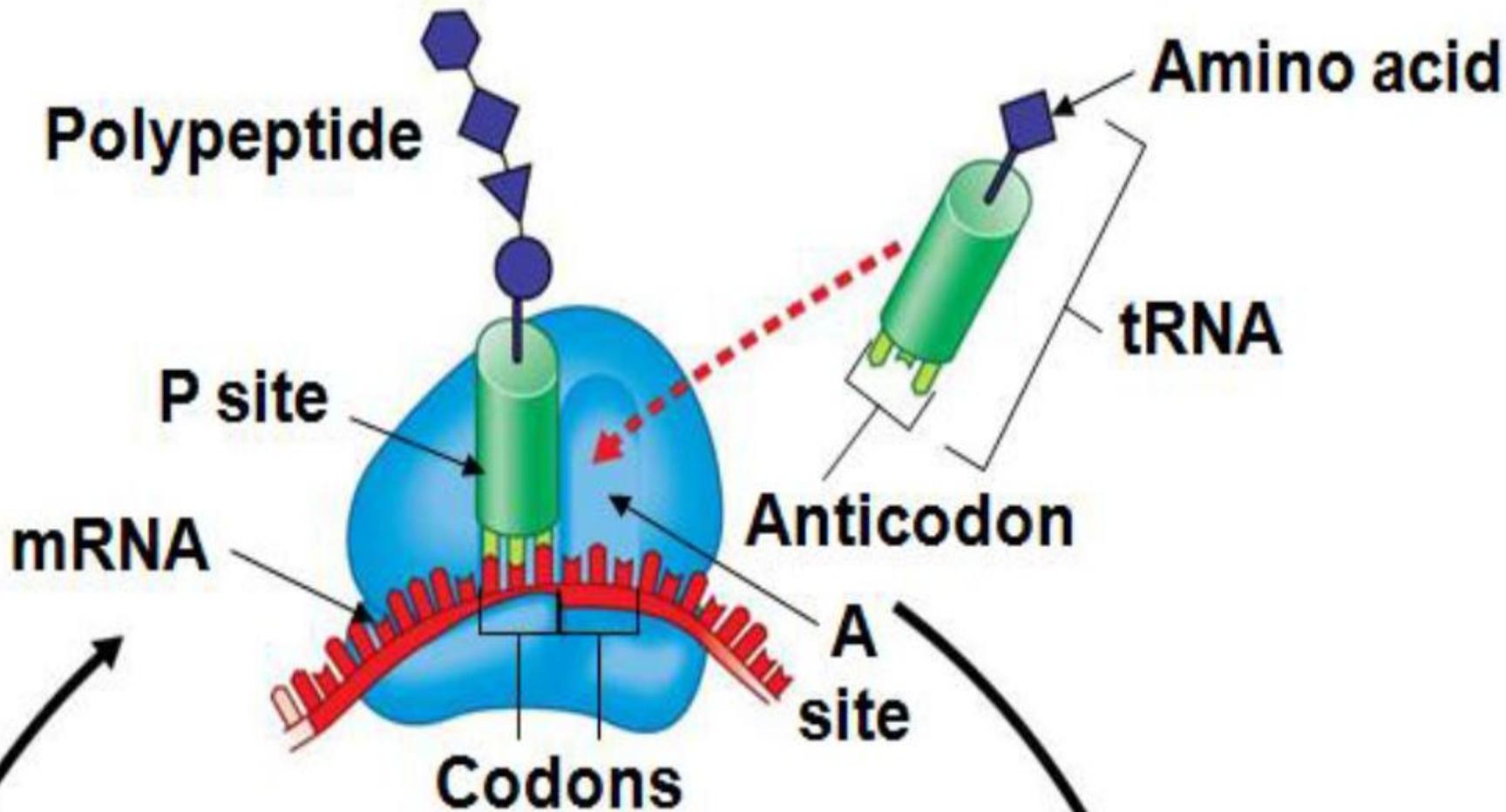
- After initiation, amino acids are added one by one to the first amino acid.
- Each addition occurs in the three-step elongation process.
  - **Step 1: Codon recognition.** The anticodon of an **incoming tRNA** molecule, carrying its amino acid, pairs with the mRNA codon in the **A site** of the ribosome.

# Elongation

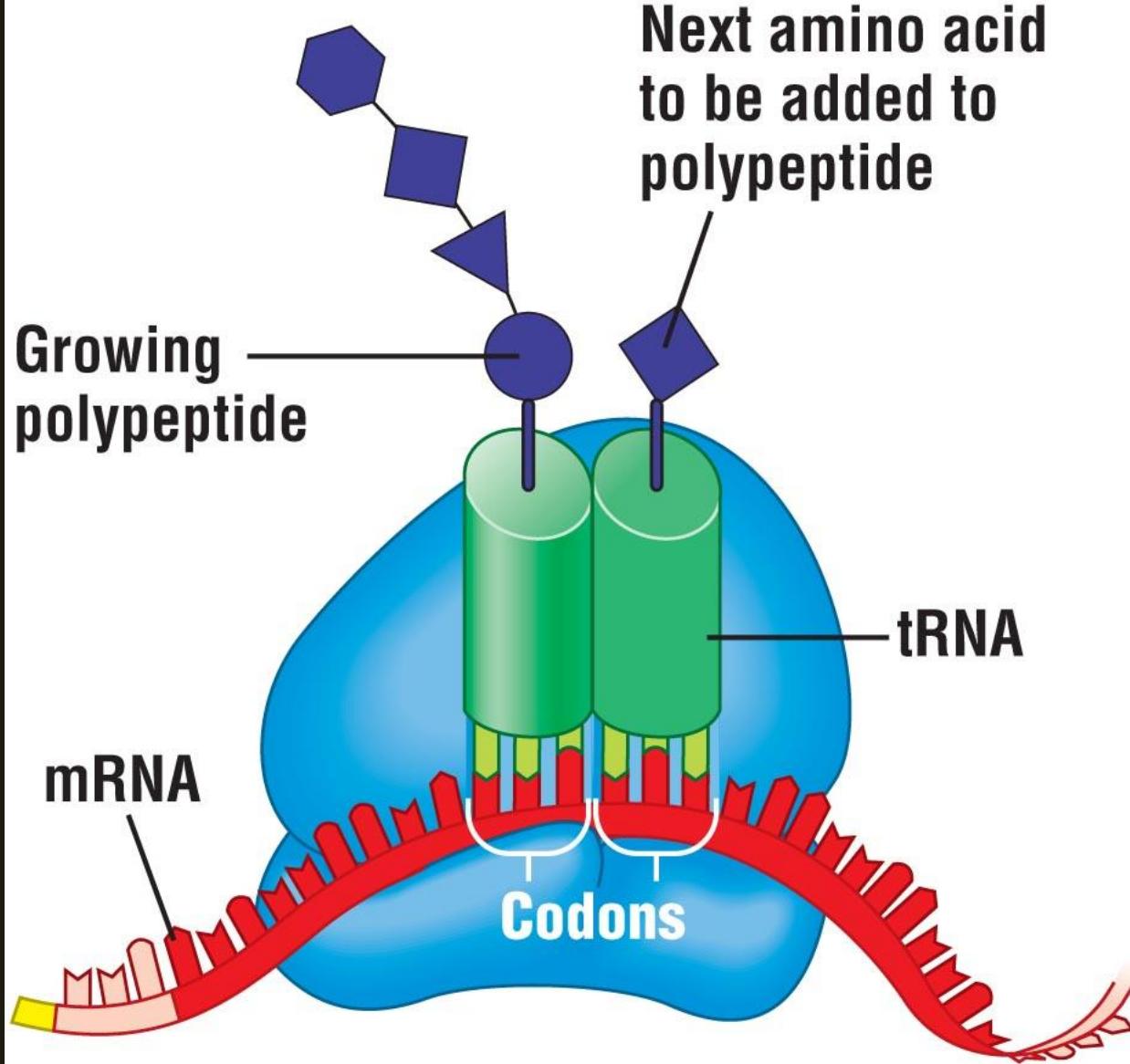
- Step 2: Peptide bond formation.
  - The polypeptide leaves the tRNA in the **P site** and attaches to the amino acid on the tRNA in the **A site**.
  - The ribosome creates a new peptide bond.
  - Now the chain has one more amino acid.

# Elongation

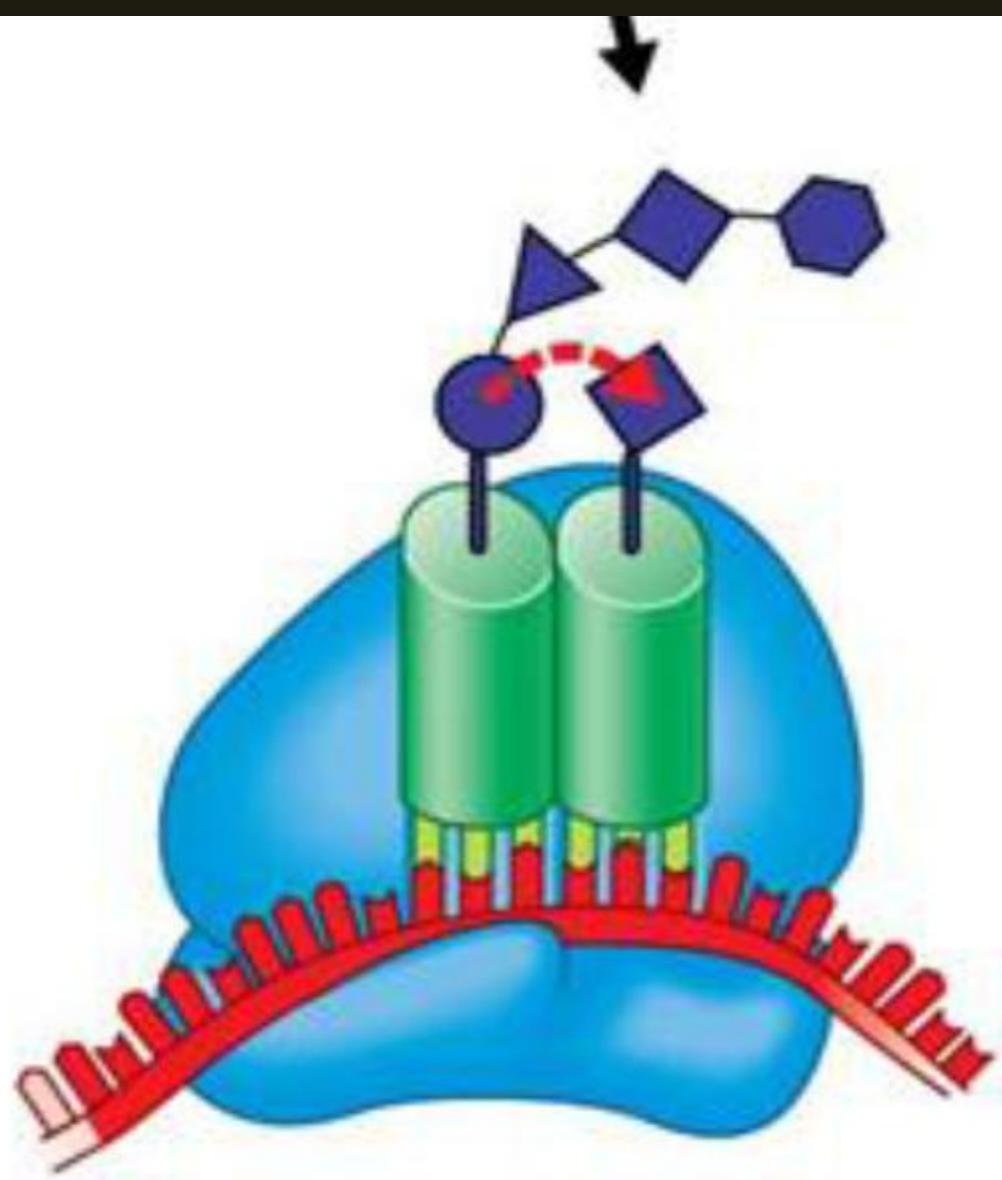
- **Step 3: Translocation.**
  - The P site tRNA now leaves the ribosome, and the ribosome moves the remaining tRNA, carrying the growing polypeptide, to the P site.
  - The mRNA and tRNA move as a unit.
  - This movement brings into the A site the next mRNA codon to be translated, and the process can start again with step 1.



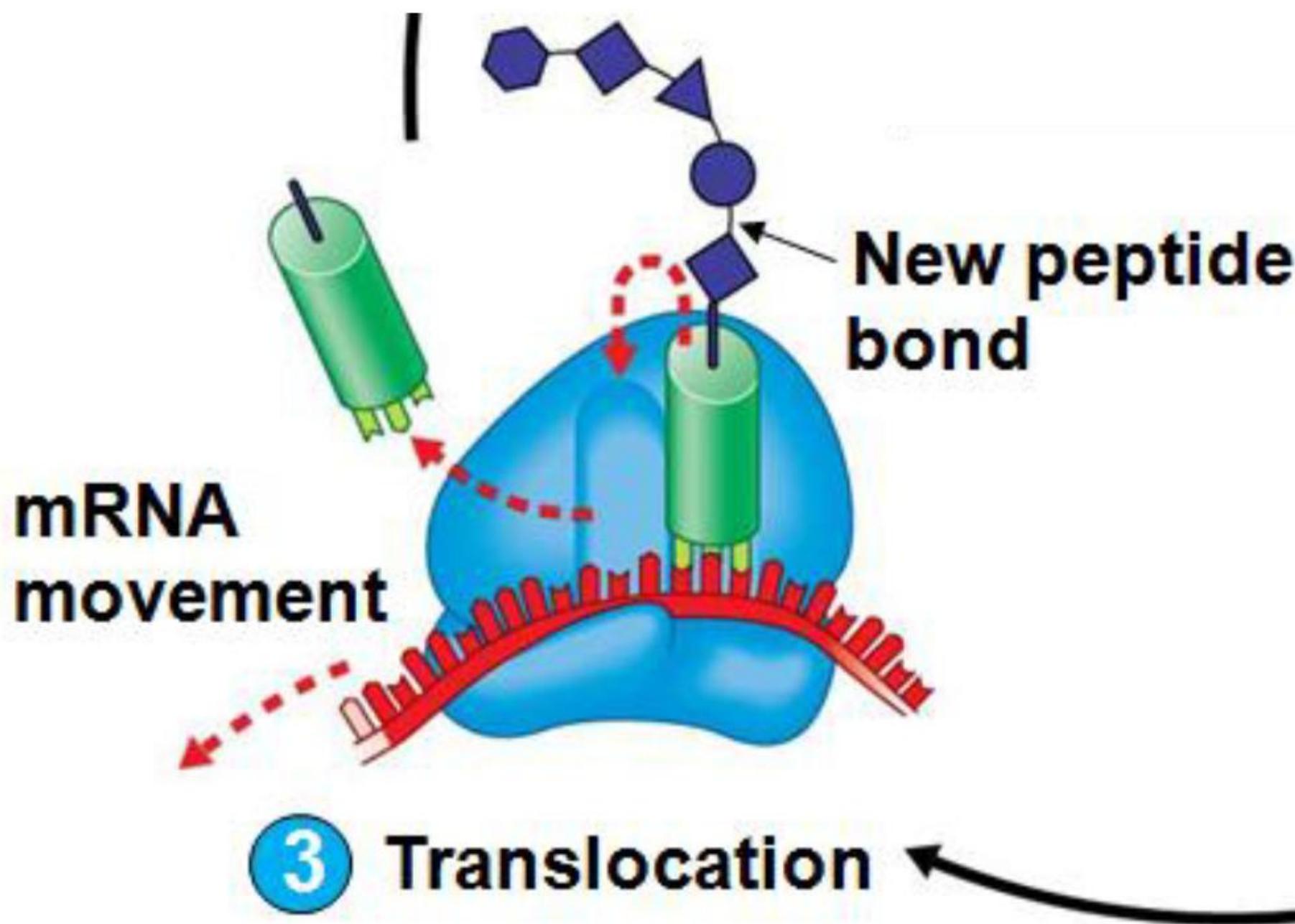
① Codon recognition

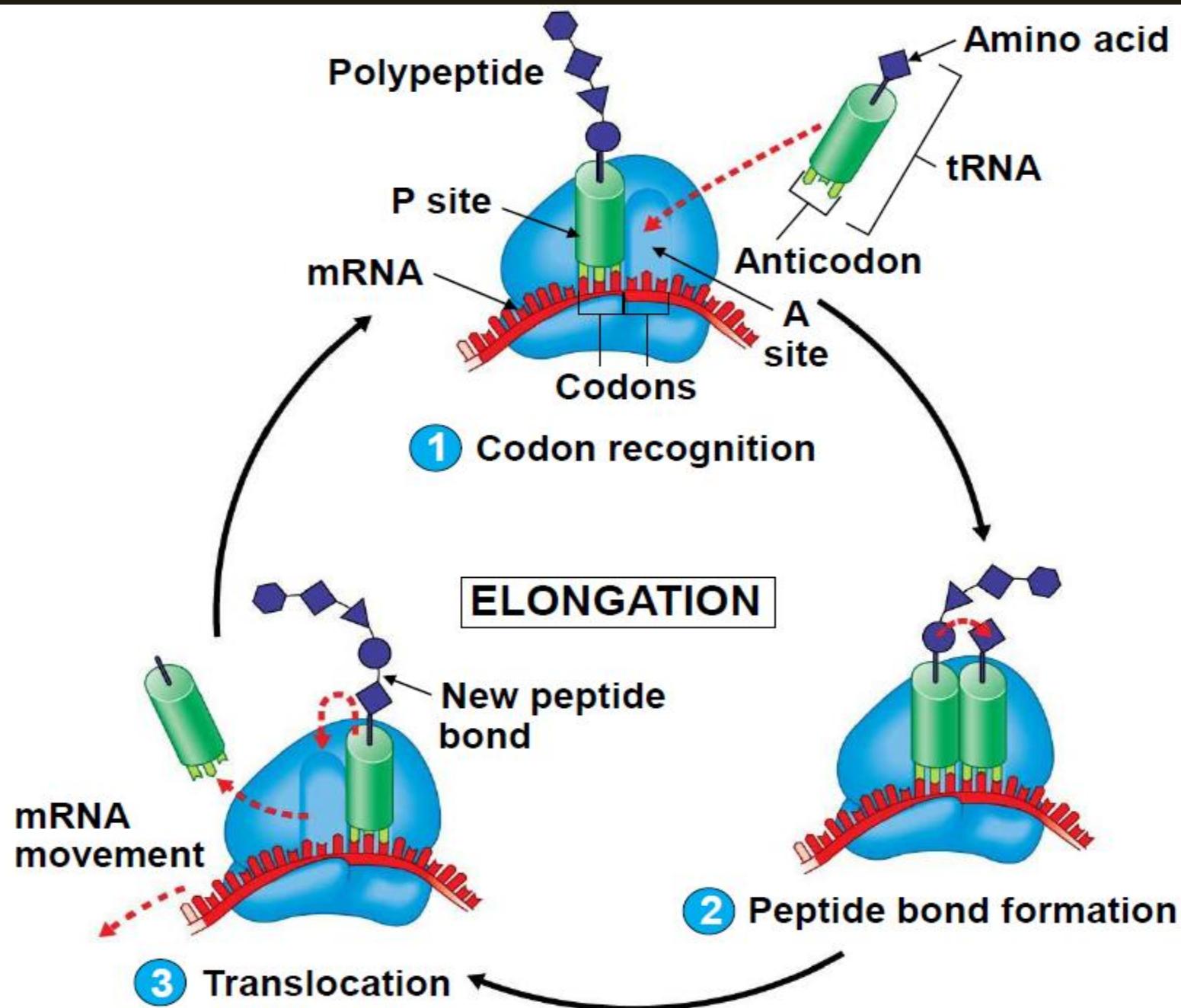


**(b) The “players” of translation.**



## ② Peptide bond formation





# Termination

- Elongation continues until
  - a **stop codon (UAA/UGA/UAG)** reaches the ribosome's A site,
  - the completed polypeptide is freed, and
  - the ribosome splits back into its subunits.