

27th July, 2016

Science of Living System

BS20001 (2-0-0)

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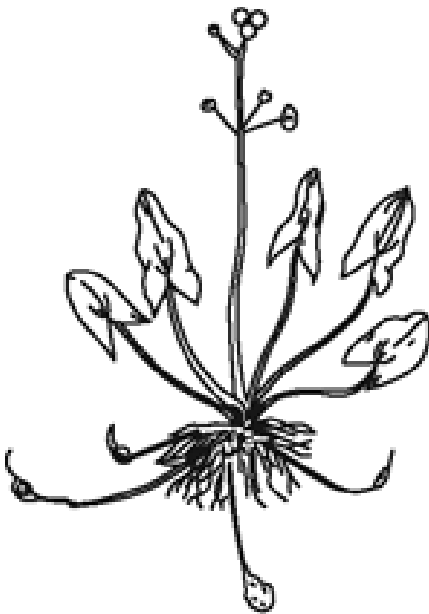
Class	Date	Schedule
1	20/7/16	L-1 (Nucleic acids: How is information stored and maintained)
2	27/7/16	L-2 (Transcription & Translation: How stored information get processed)
3	3/8/16	L-3 (Protein structure: Making sense of processed information)
4	10/8/16	L-4 (Enzymes: Working principles & concept of enzyme technology)
5	17/8/16	L-5 (Photosynthesis and Respiration: Making and breaking of sugars)



Overview of Transcription and Translation

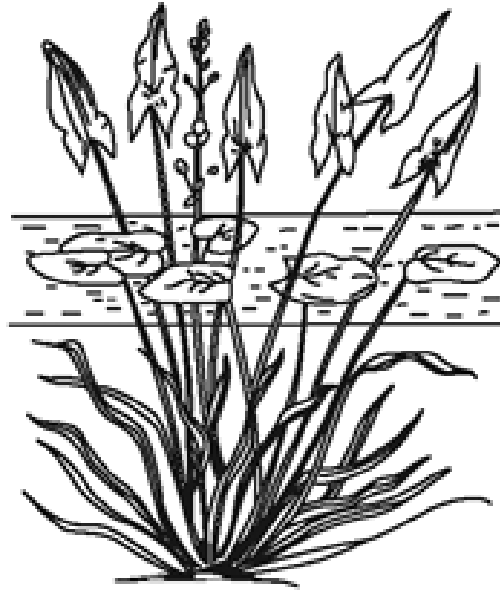
*(How stored information get
processed)*

DNA: Contains the Instruction for Life



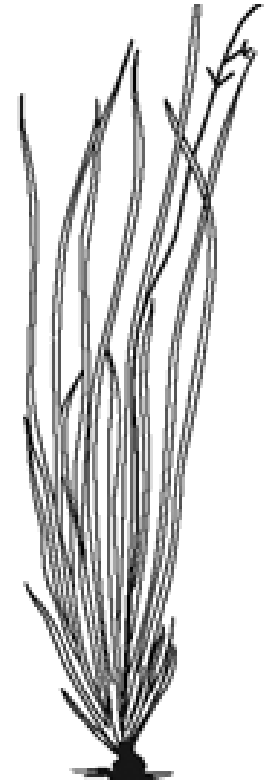
Plant-X

Completely terrestrial



Plant-Y

Partially submerged

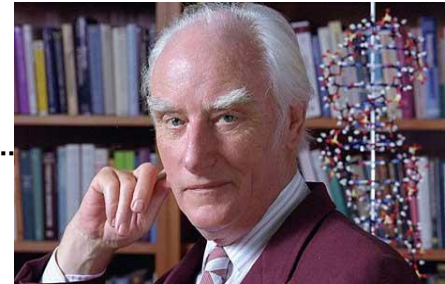
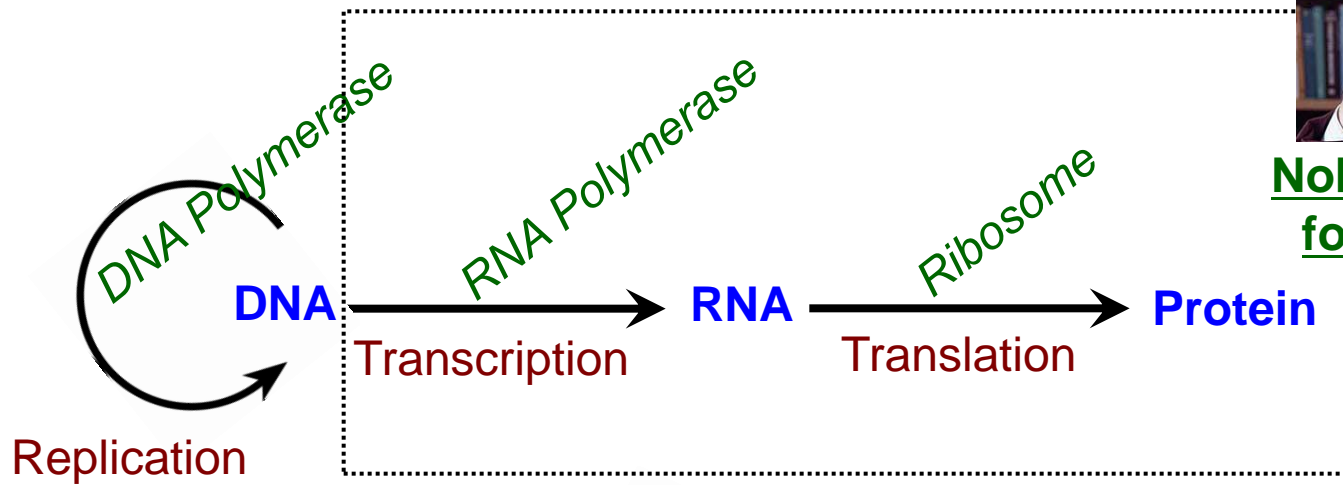


Plant-Z

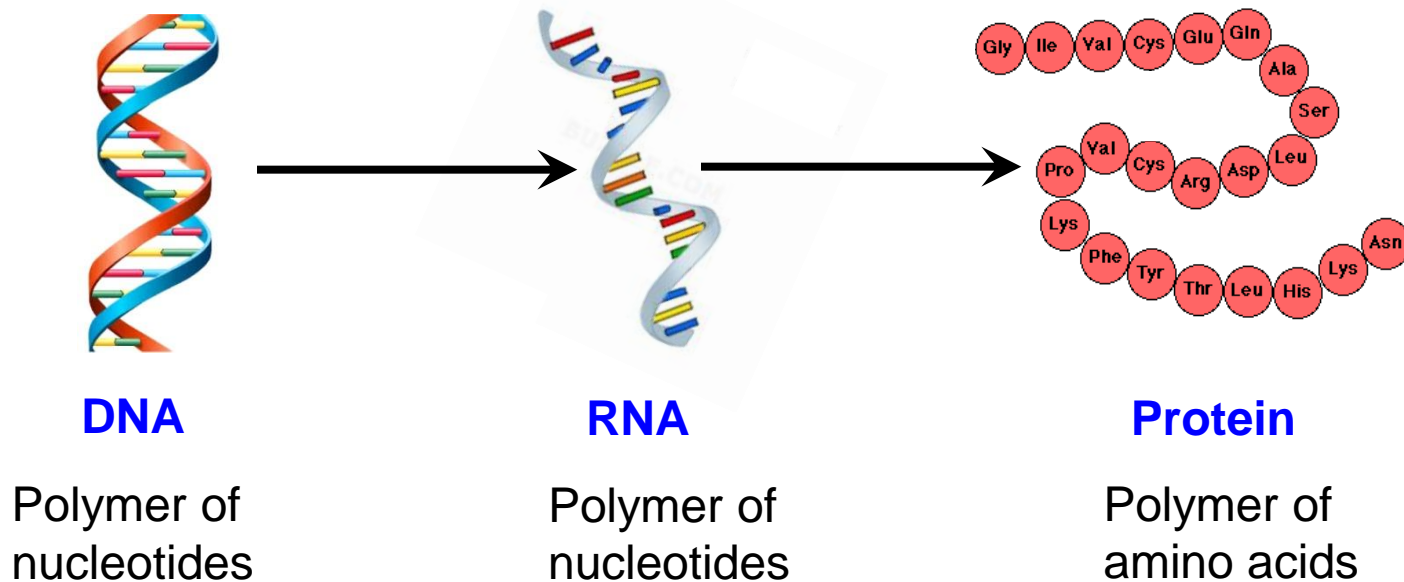
Completely submerged

The phenotype (visible configuration) of the marsh plant *Sagittaria sagittifolia* depends on its environment

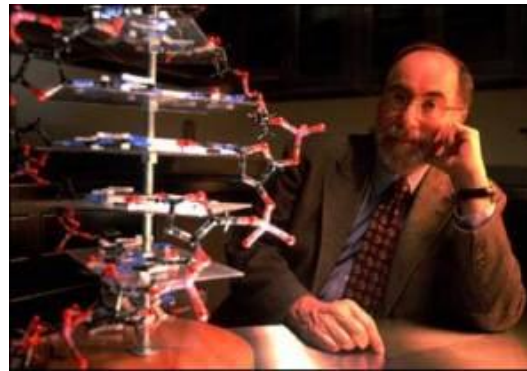
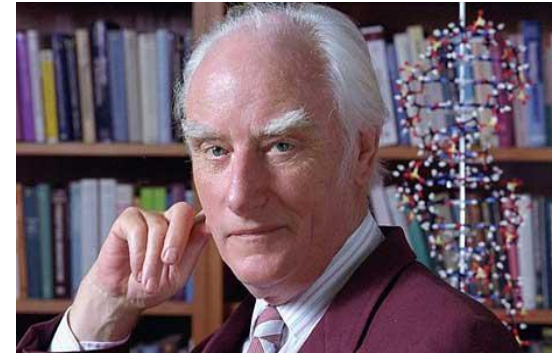
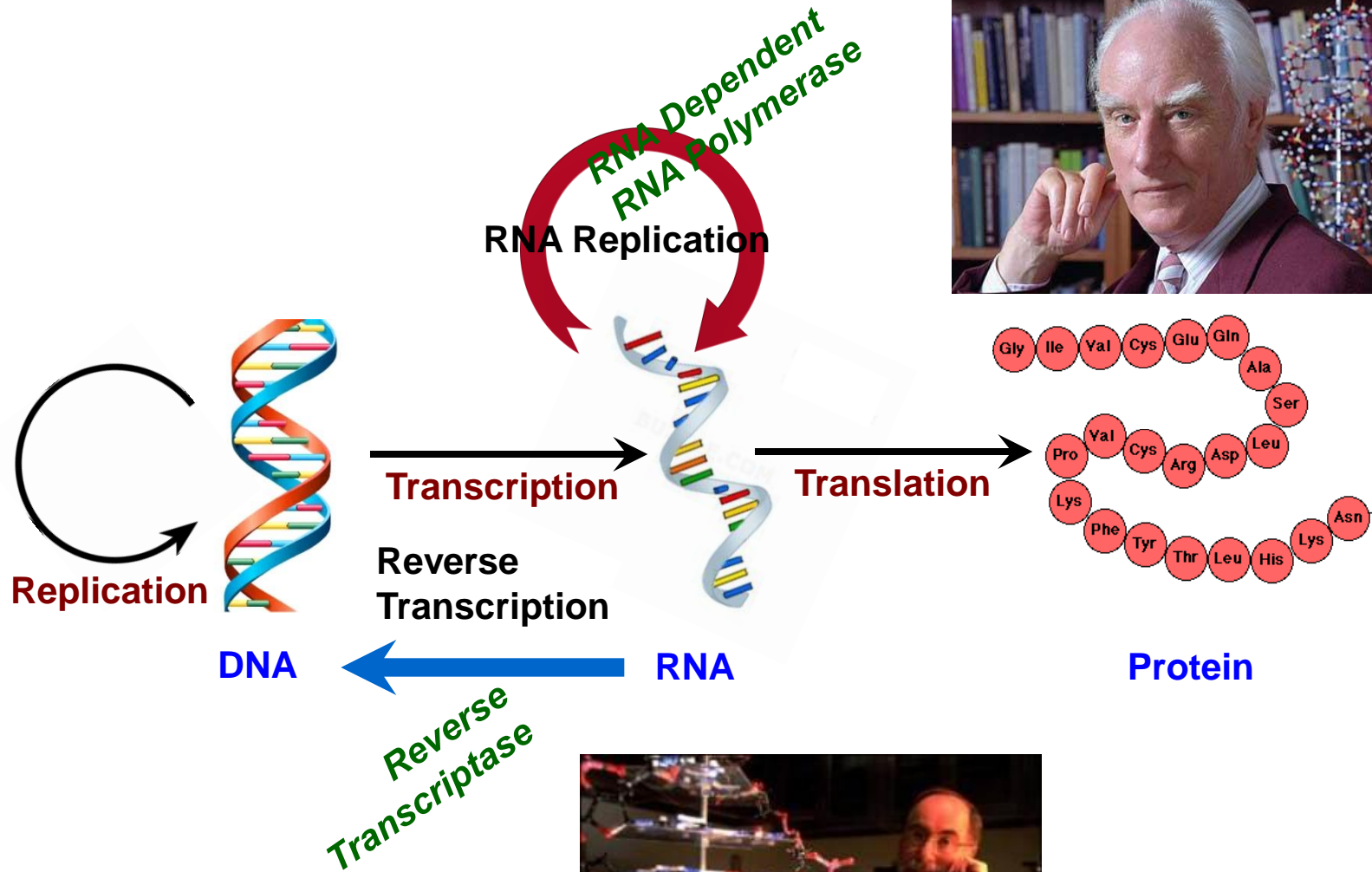
Flow of Genetic Information: The Central Dogma of Molecular Biology



Nobel Prize in 1962
for DNA Structure



Flow of Genetic Information: Updates



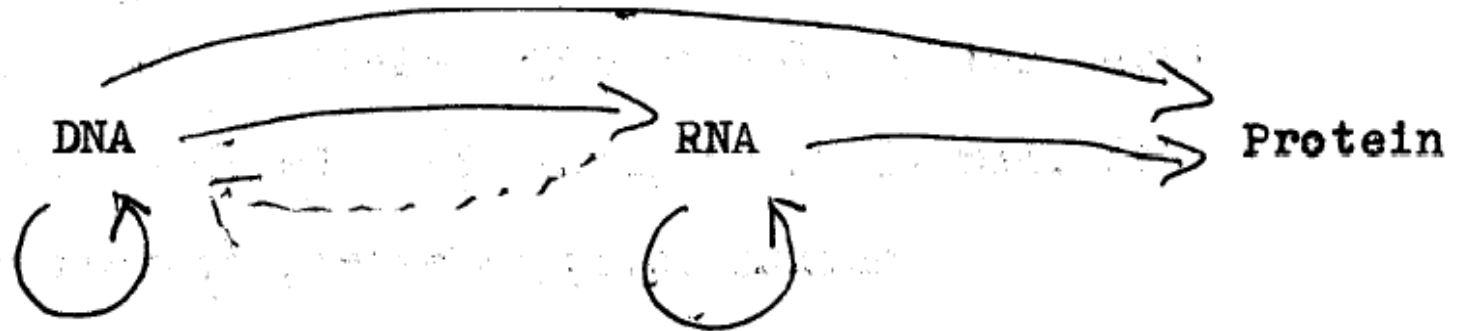
David Baltimore
Nobel Prize in 1975

Ideas on Protein Synthesis (Oct. 1956)

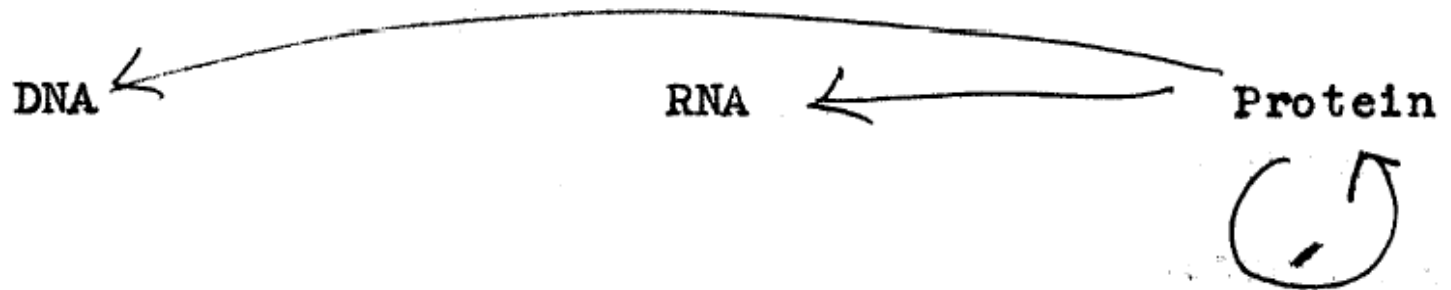
Francis Crick

The Central Dogma: "Once information has got into a protein it can't get out again". Information here means the sequence of the amino acid residues, or other sequences related to it.

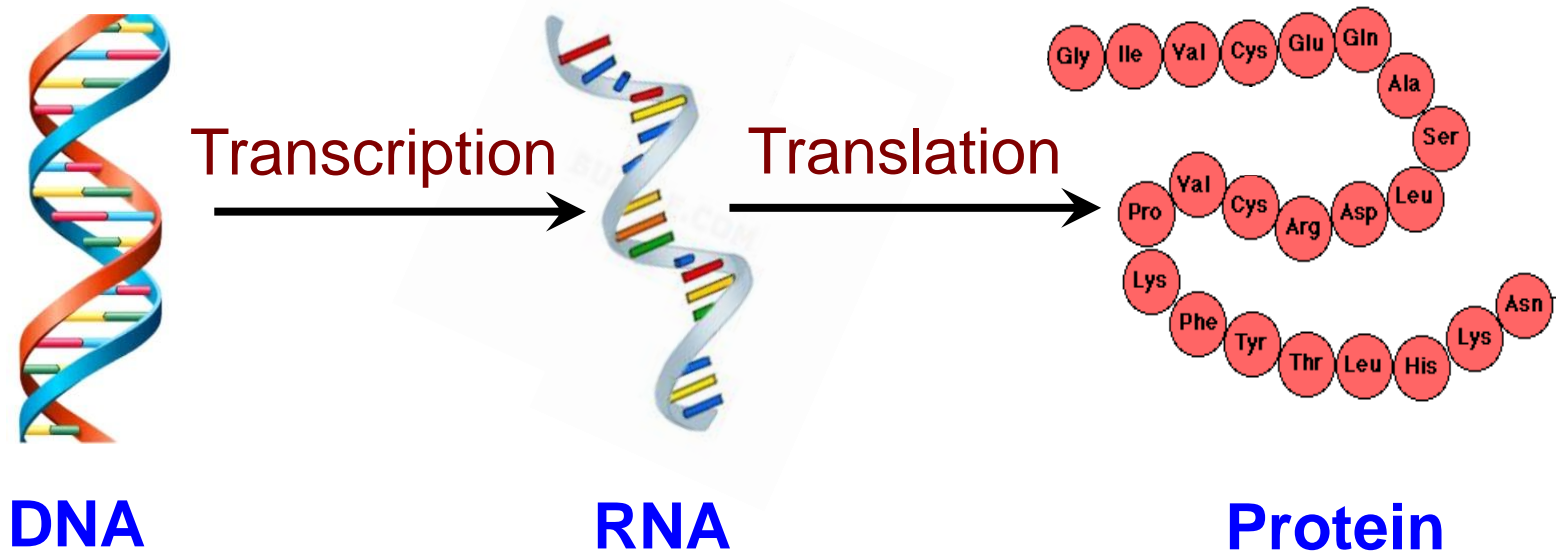
That is, we may be able to have



but never



Correlations Between DNA Content and Its Downstream Product

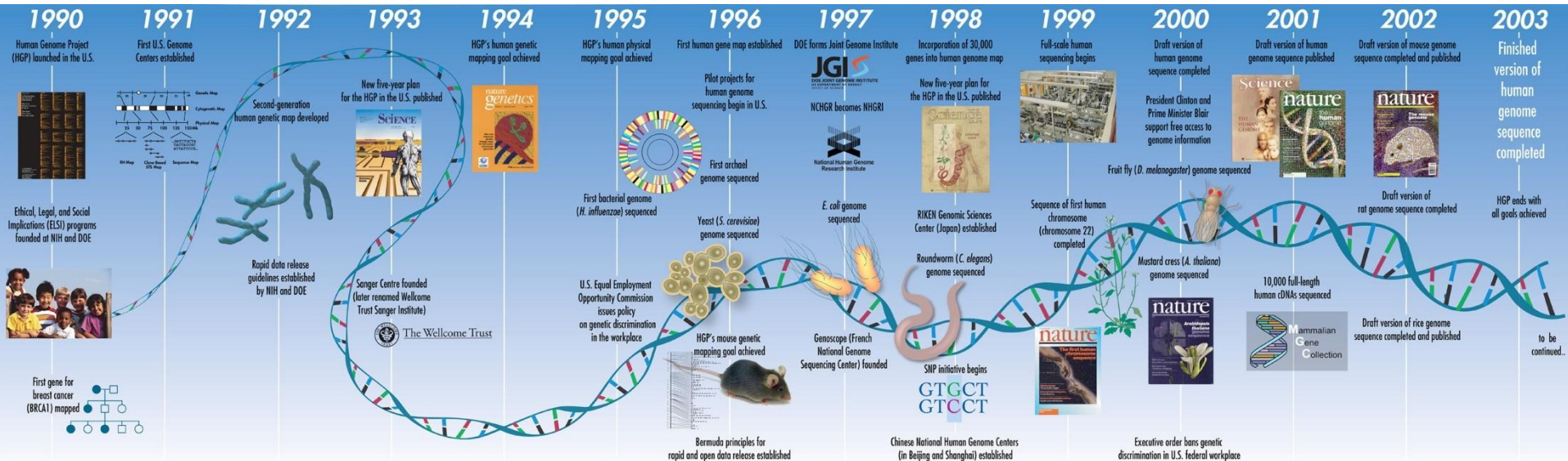


“Human Genome Project” Dramatically Enhanced Our Understanding on Gene Expression

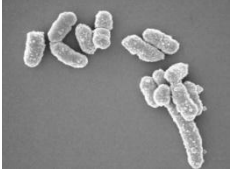
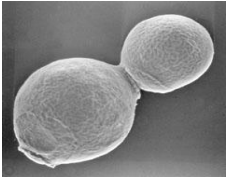



Human Genome Project



~21,000 human genes (appeared to be significantly fewer than previous estimates)



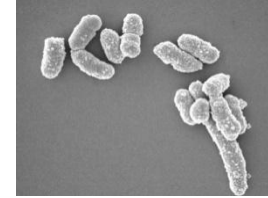
Genome Size, Gene Number, and Complexity of an Organism

	Organism	Genome size (bp)	Protein coding genes
	<i>E. coli</i>	4,600,000	4,250
	<i>S. cerevisiae</i>	12,160,000	5,616
	<i>C. elegans</i>	100,000,000	19,735
	Human	3,200,000,000	19,042
	Marbled lungfish	139,000,000,000	NA

Transcription

Total DNA content vs transcribable content

Genome size (bp)



4,600,000

- ▶ **Protein coding sequences is ~1.5% of total DNA content (human)**

Messenger RNA (mRNA)



3,200,000,000

- ▶ **Besides protein coding region, DNA can be transcribed into:**

Ribosomal RNA (rRNA)

Transfer RNA (tRNA)

Small nuclear RNA (snRNA)

Regulatory RNA

- ▶ **Most of the DNA sequences are not transcribed**

What is a Machine?

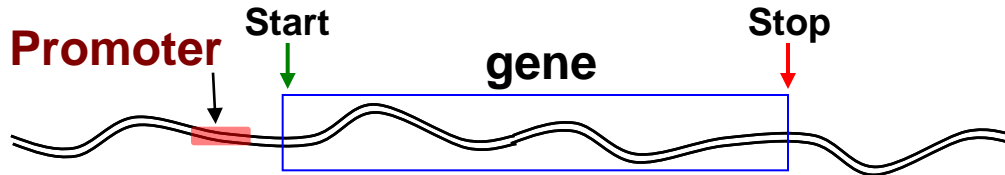
A piece of equipment with several moving parts that uses power to do a particular type of work.

- Cambridge dictionary

Biological machines:

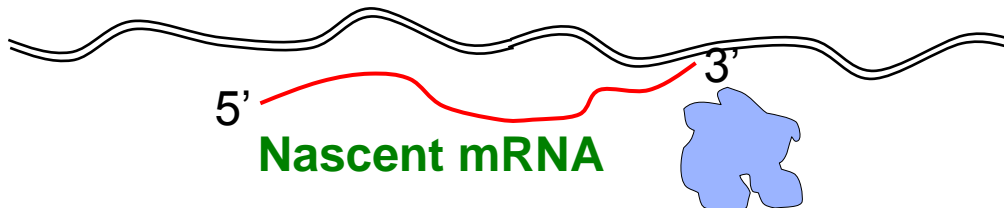
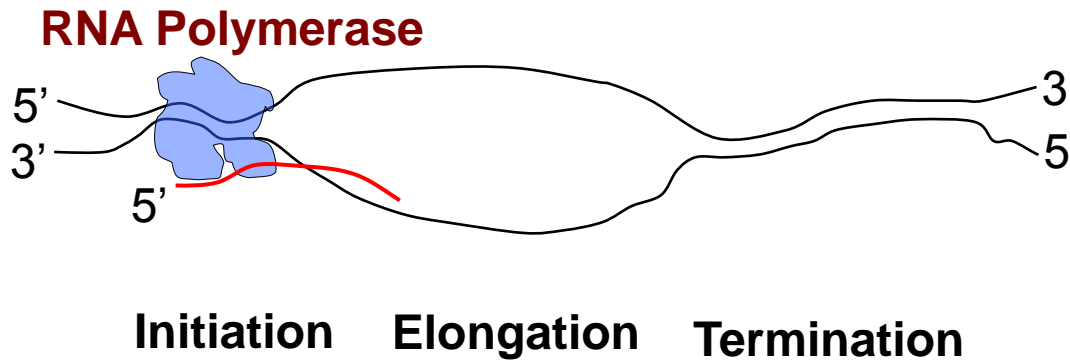
- DNA polymerase
- RNA polymerase
- Ribosome

Transcription: Involved Machineries and Processes



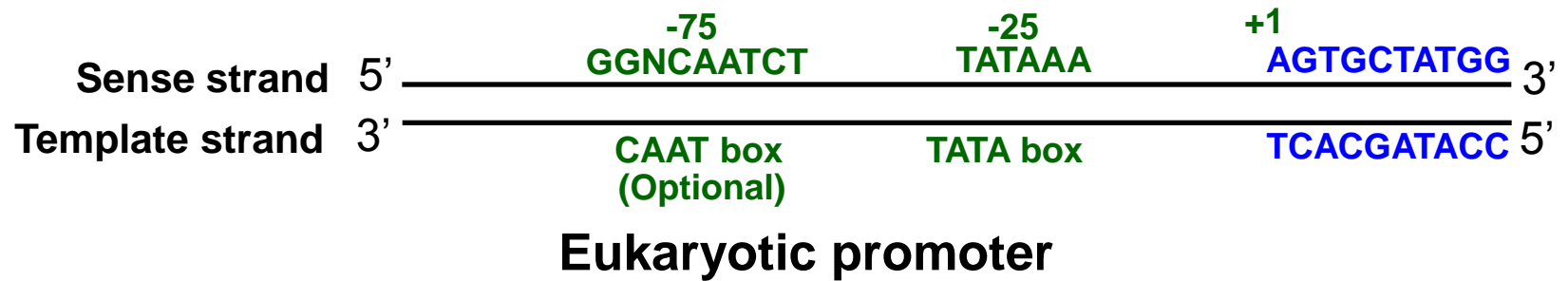
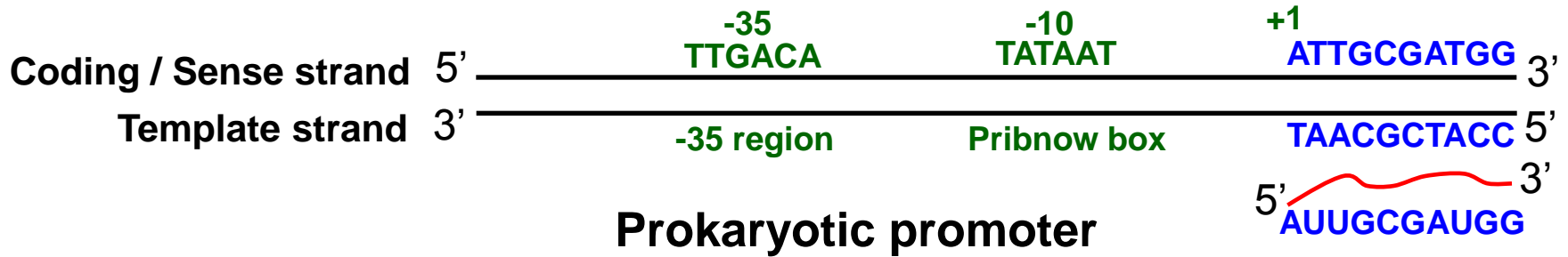
Key points to be discussed

1. Promoter
2. RNA Polymerase
3. RNA synthesis
4. Termination

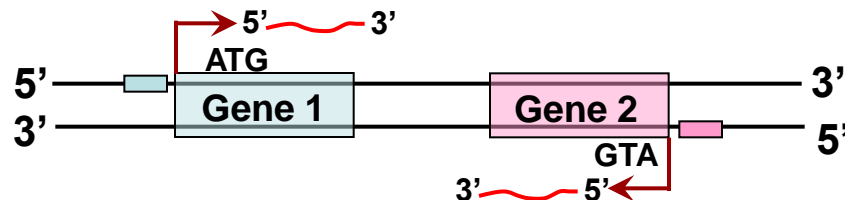


Promoter for Transcription

Promoter is just like a “start button” of an assembly line that starts the conveyor belt moving



► Transcription can start from both directions

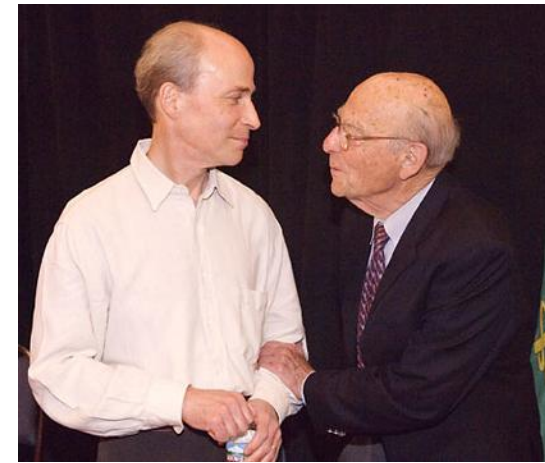
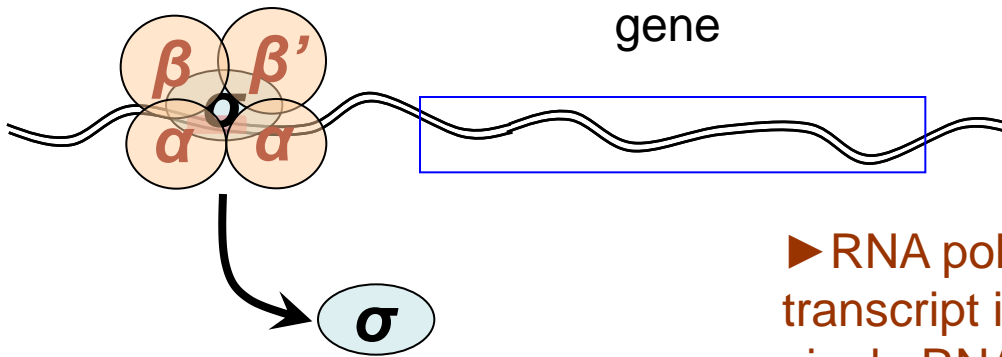


RNA Polymerase

Subunits of RNA Polymerase: α , α , β , β' and σ

Holoenzyme: α , α , β , β' and σ

Coreenzyme: α , α , β and β'



Roger Kornberg
Nobel Prize in 2006

► RNA polymerase is completely Processive: A transcript is synthesized from start to end by a single RNA polymerase molecule.

► RNA polymerase can initiate the synthesis of RNA *de-novo* (No primer required)

RNA Molecules in *E. coli*

mRNA	5%
tRNA	15%
rRNA	80%

Who transcribes this huge pool of rRNA and tRNA?

In bacteria same RNA polymerase transcribe all these three types of RNA

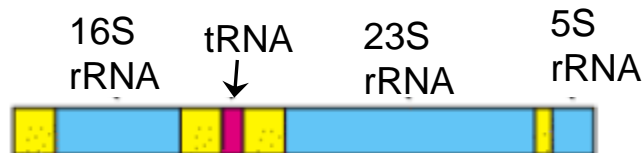
In eukaryotes different RNA polymerases are involved in transcription of mRNA, rRNA and tRNA

RNA Polymerase in Eukaryotes

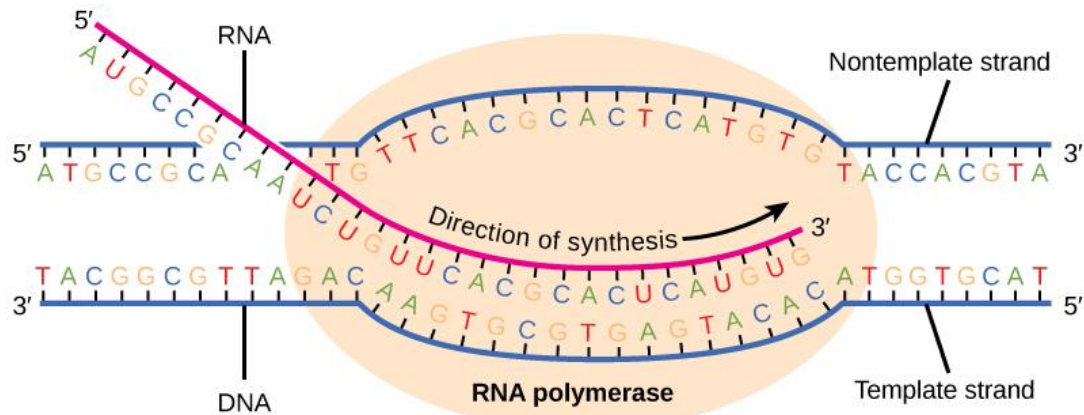
In contrast to bacteria, the nucleus of a eukaryote contains 3 types of RNA polymerase

	Template specificity	Location in the nucleus	Susceptibility to inhibitors (α -amanitin)
RNA Pol- I	18S, 5.8S, and 28S r RNA	Nucleoli	Insensitive
RNA Pol- II	mRNA	Nucleoplasm	Highly susceptible
RNA Pol- III	5S rRNA, and tRNA	Nucleoplasm	Susceptible

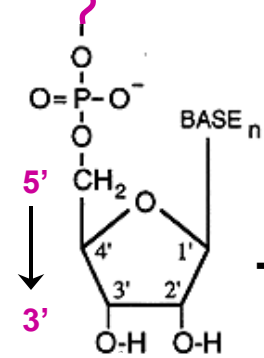
How rRNA and tRNA are generated in bacteria?



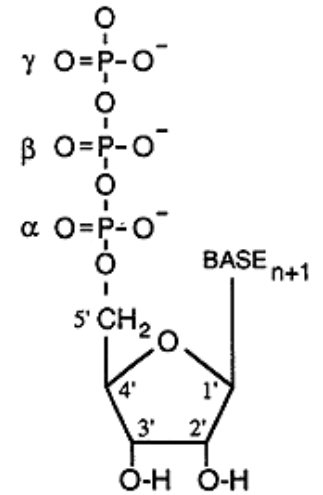
RNA Synthesis



Growing
RNA chain



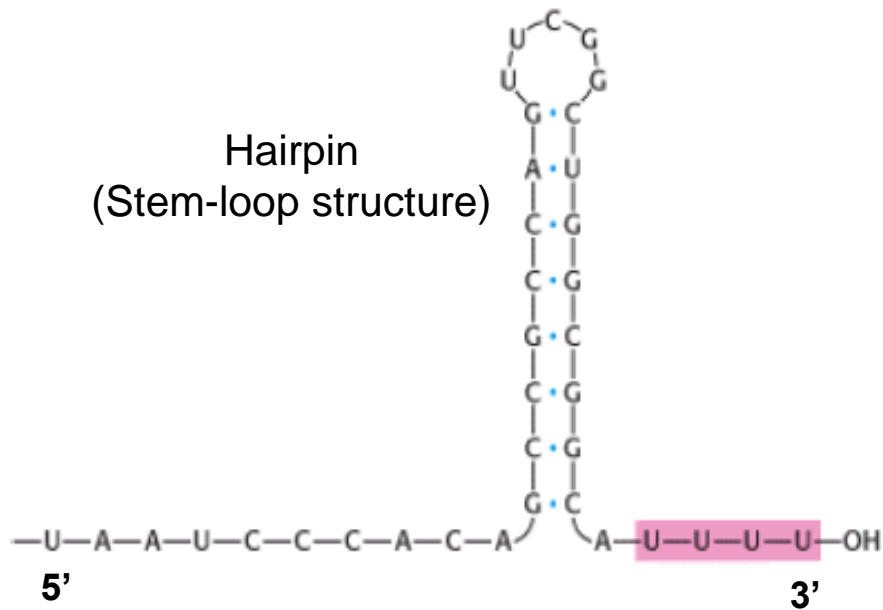
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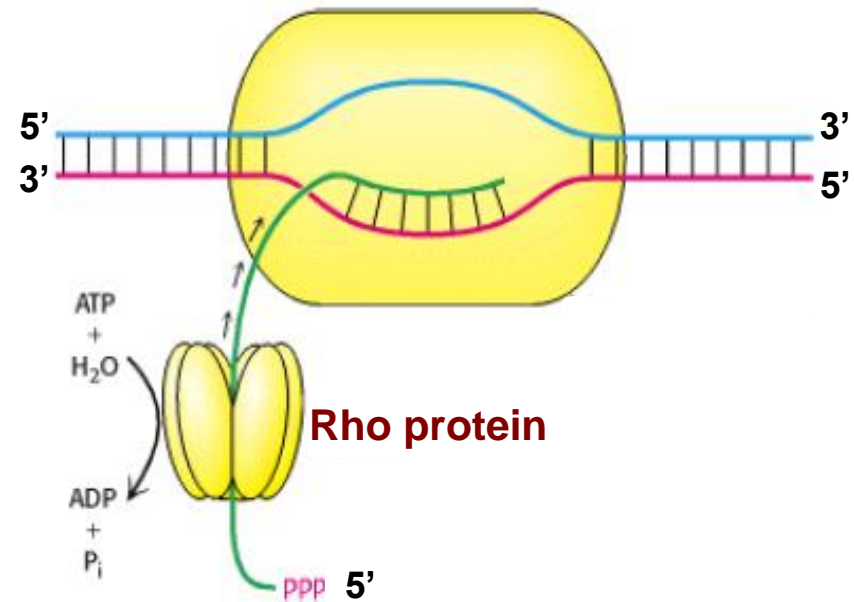
Ribonucleotide

Termination of Transcription in Prokaryotes

An RNA hairpin followed by several uracil residues terminates transcription

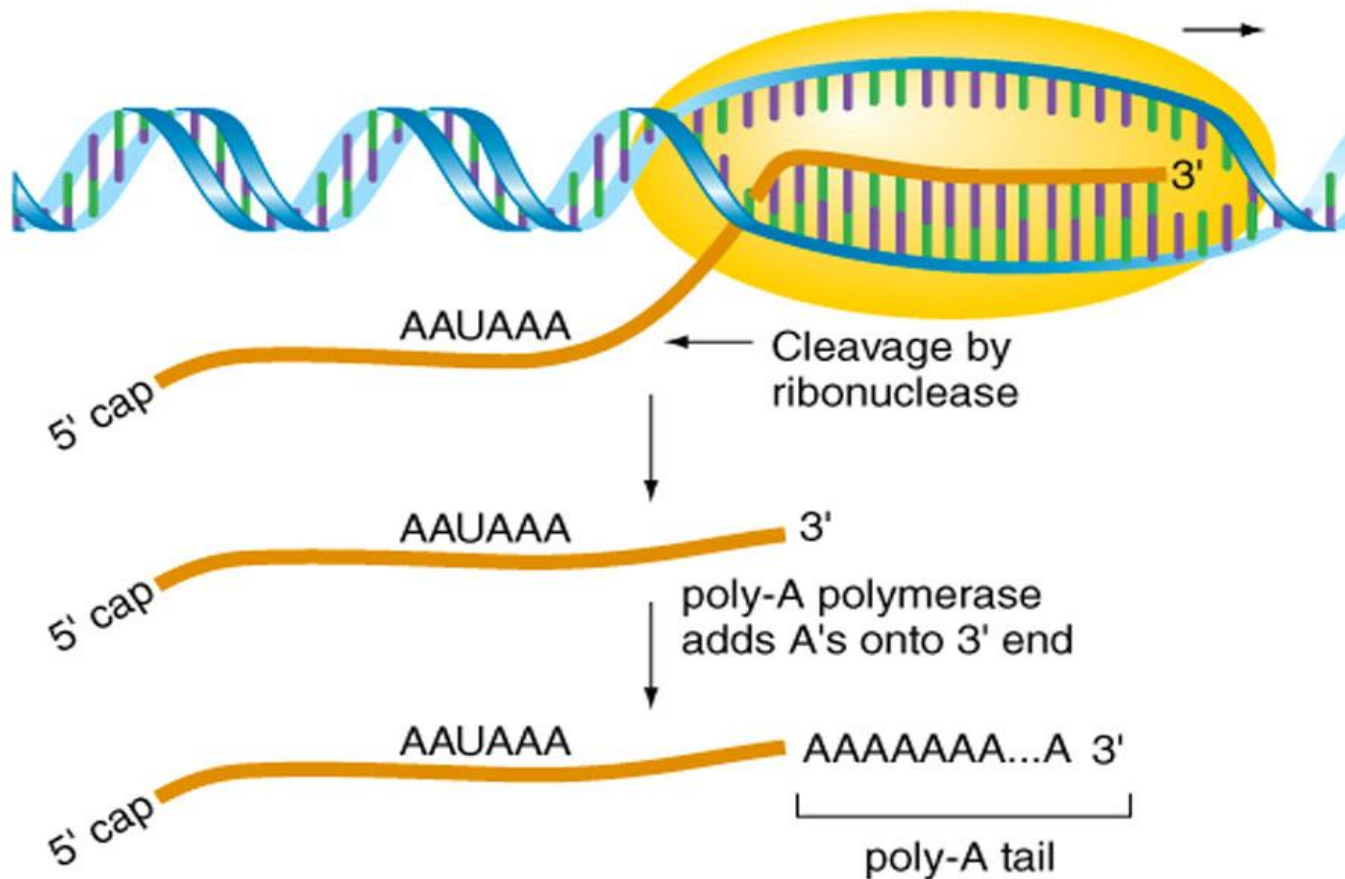


Rho binds the nascent RNA chain and pulls it away from RNA polymerase and the DNA template.



Eukaryotic Transcripts Need to be Processed

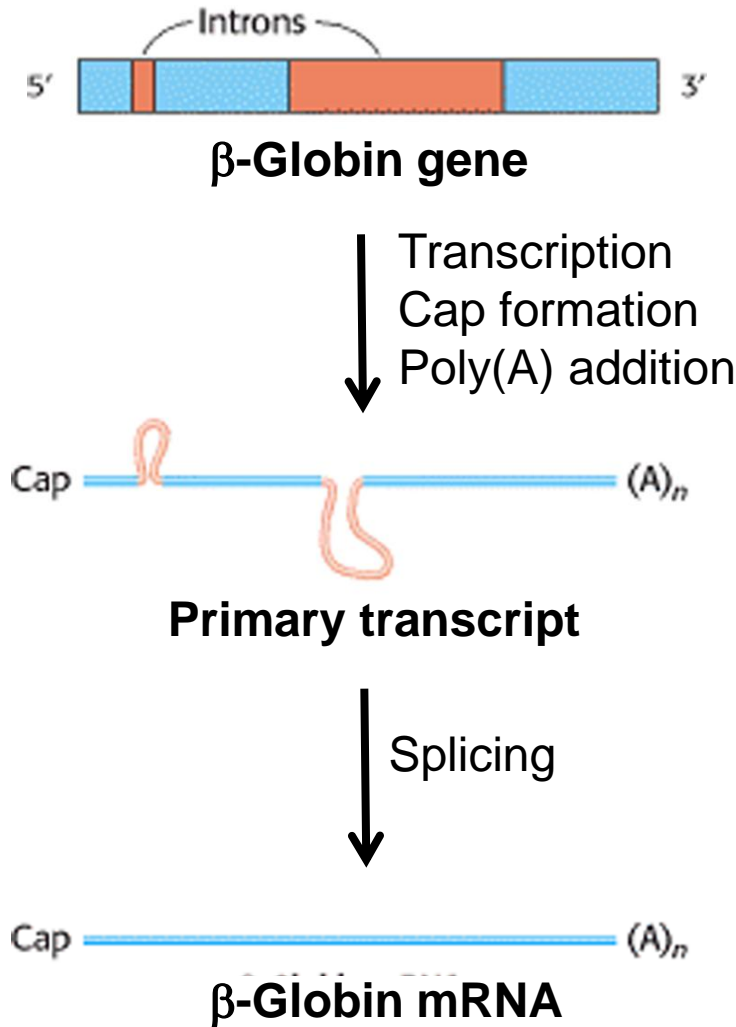
- Ends of a nascent mRNA acquire a 5' cap and a 3' poly A tail



- Increase stability of mRNA
- More effective template for translation

Eukaryotic Transcripts Need to be Processed

► **Splicing (mediated by specialized enzymatic machineries consisting of snRNAs and proteins) removes introns from nascent mRNA**



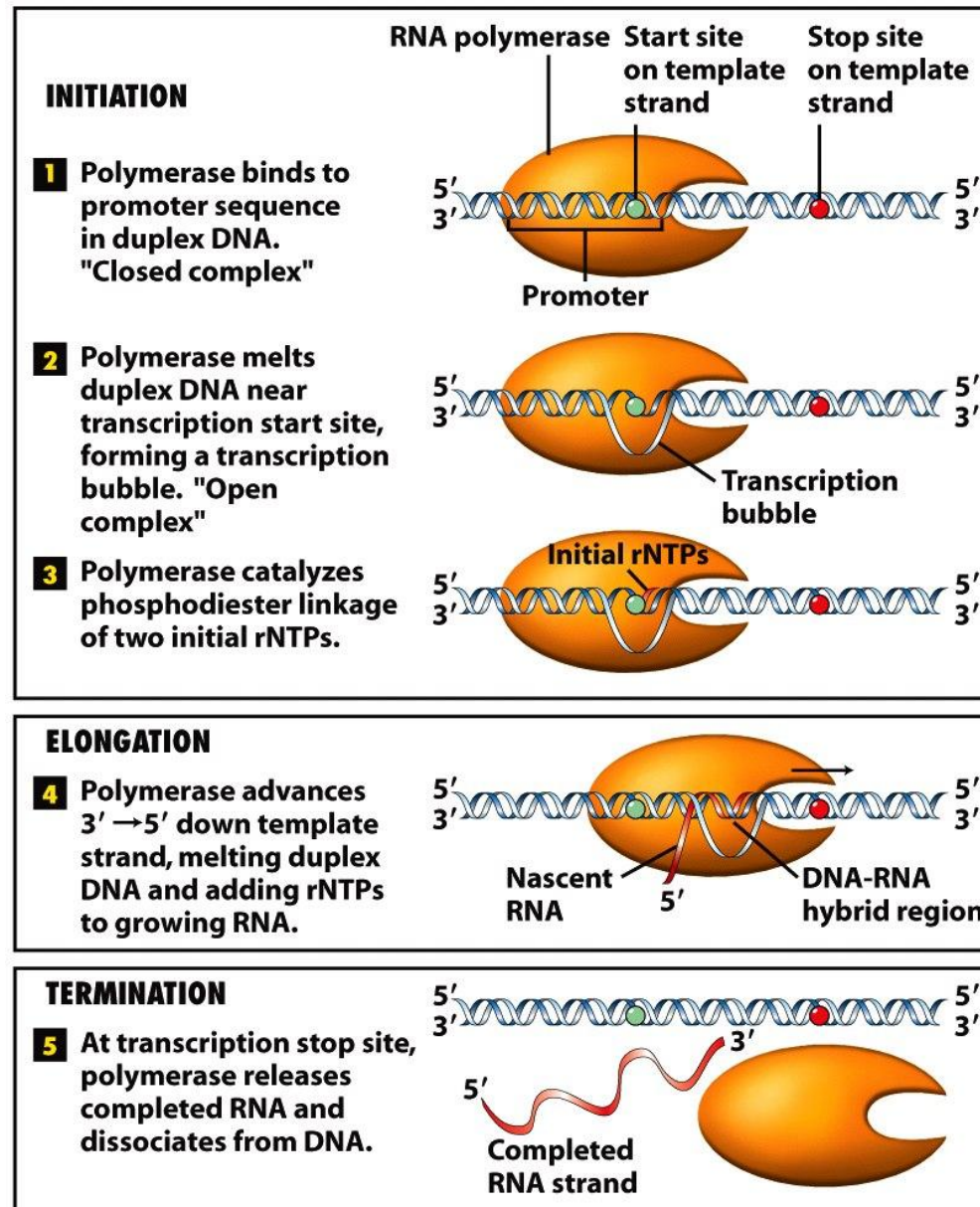
Thomas Cech
Nobel prize in 1989



****Splicing generates more variation**

Membrane bound vs soluble form of a protein

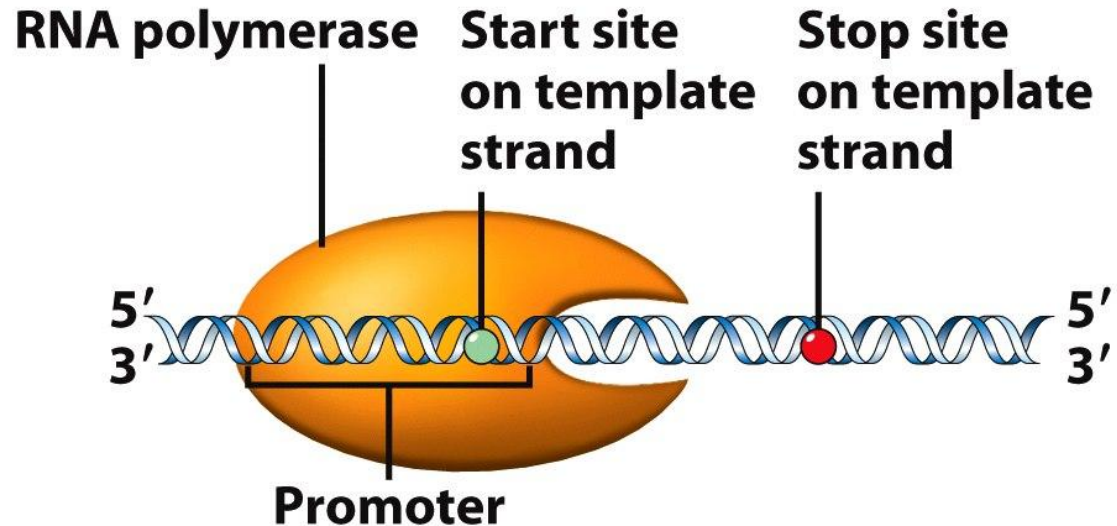
Schematic Representation of Multiple Steps Involved in Transcription



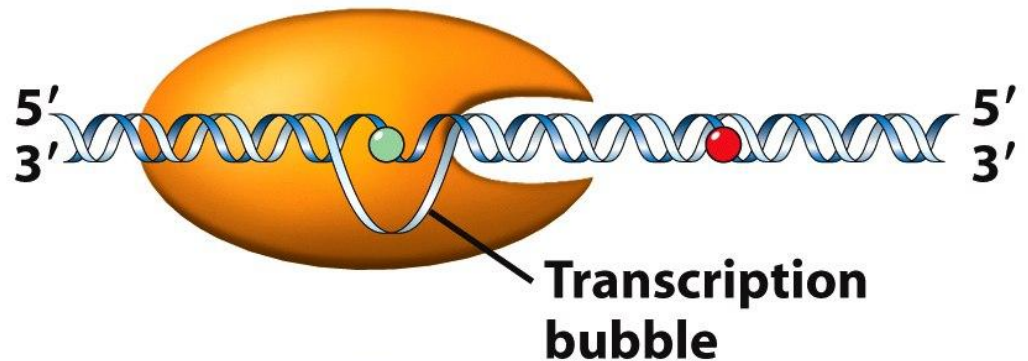
Schematic Representation of Multiple Steps Involved in Transcription

INITIATION

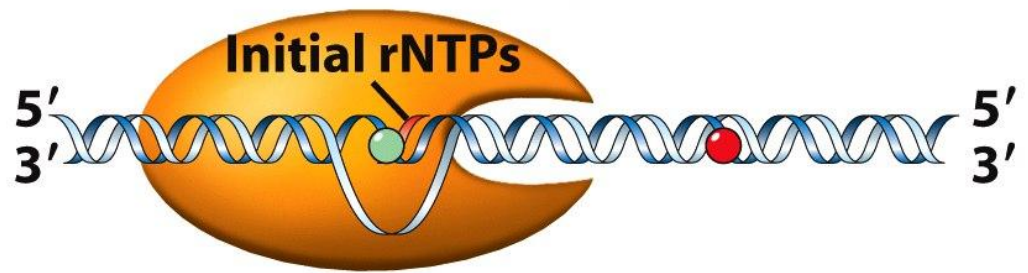
Polymerase binds to promoter sequence in duplex DNA.
"Closed complex"



Polymerase melts duplex DNA near transcription start site, forming a transcription bubble. "Open complex"



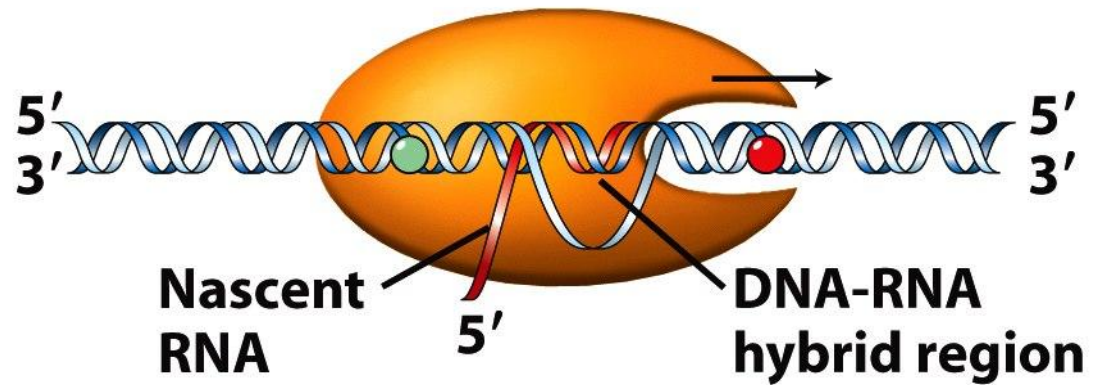
Polymerase catalyzes phosphodiester linkage of two initial rNTPs.



Schematic Representation of Multiple Steps Involved in Transcription

ELONGATION

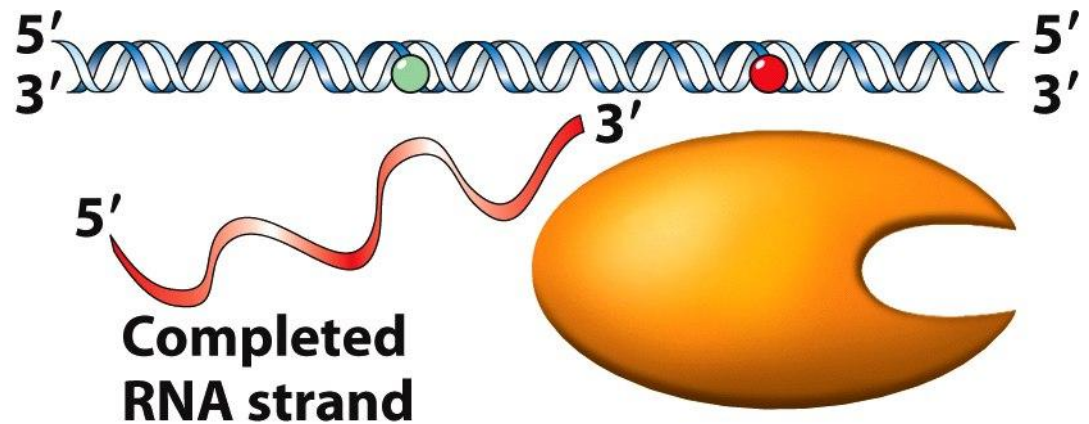
Polymerase advances 3' → 5' down template strand, melting duplex DNA and adding rNTPs to growing RNA.



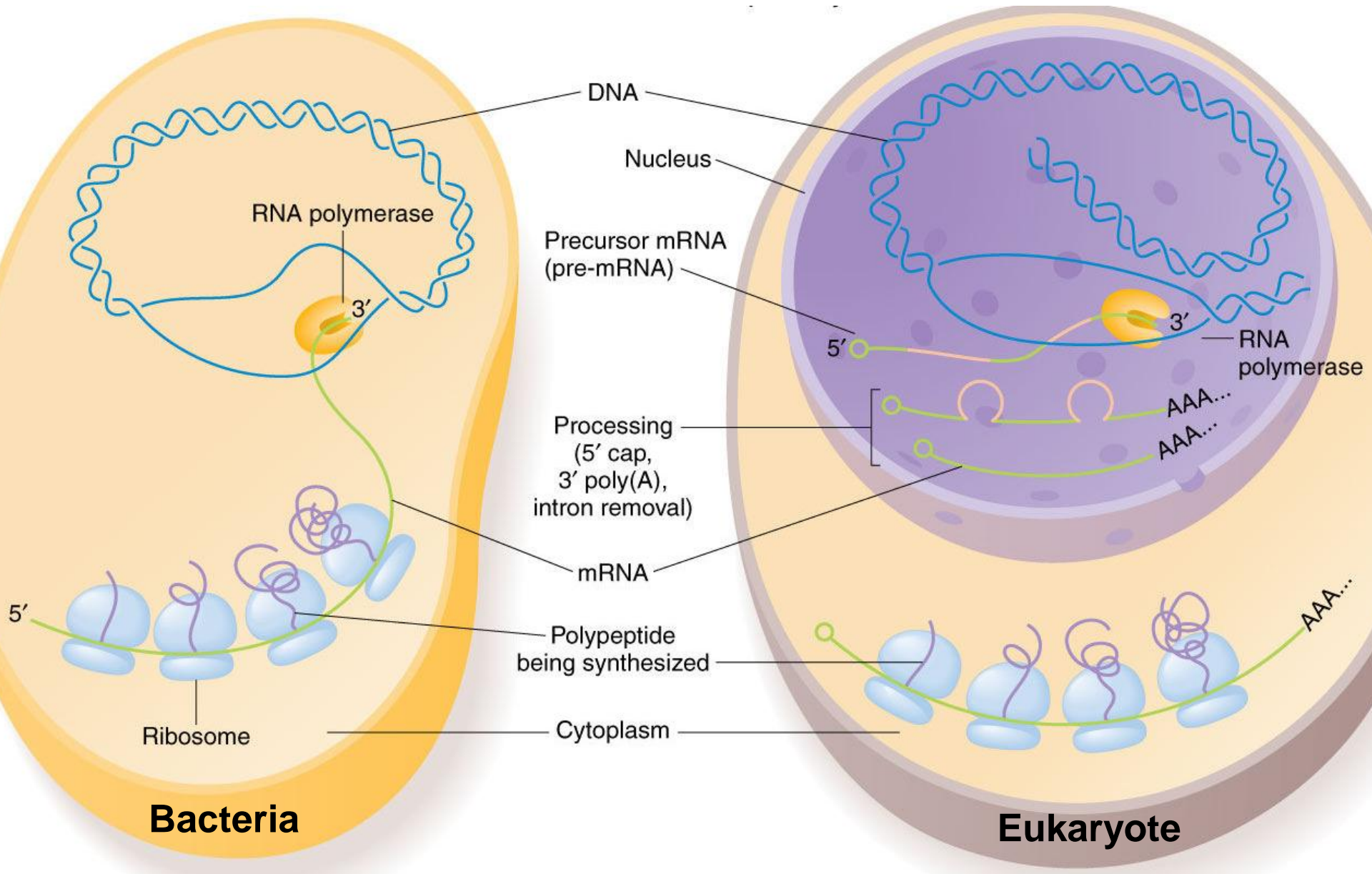
Schematic Representation of Multiple Steps Involved in Transcription

TERMINATION

At transcription stop site, polymerase releases completed RNA and dissociates from DNA.

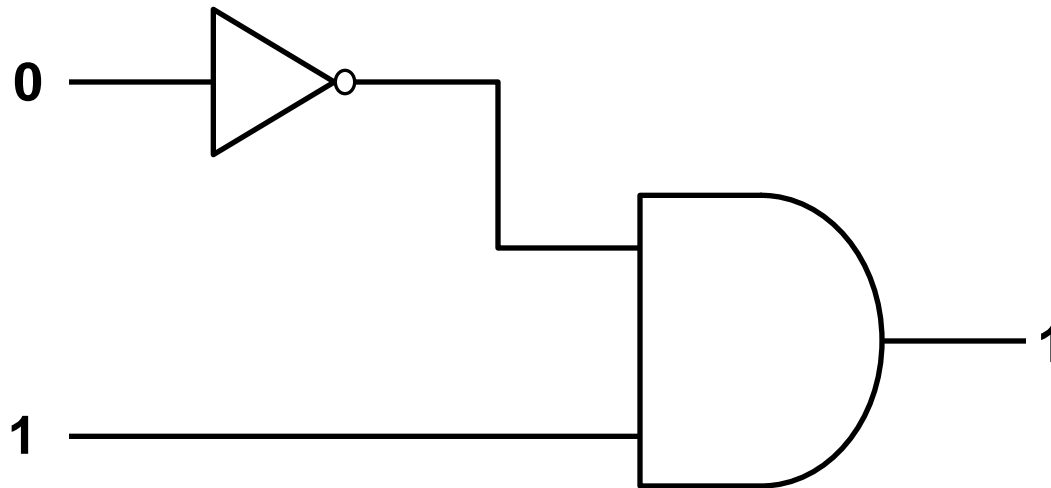


Transcription: At a Glance



Regulation of Gene Expression

(Biological circuits)



Regulation of Gene Expression

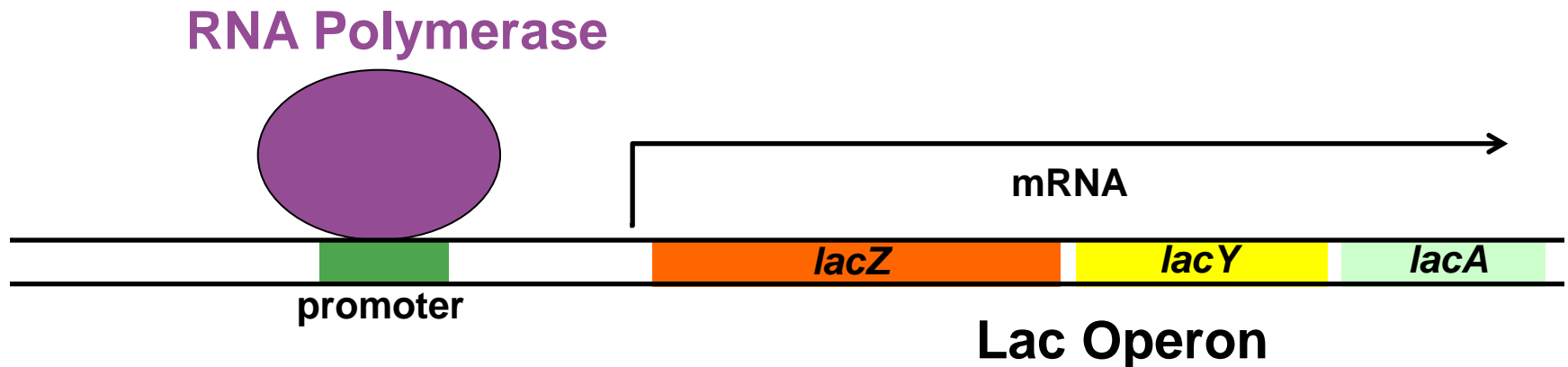
Each cell contains all the genetic material for growth and development

Some of these genes are expressed all the time

Other genes are not expressed all the time. They are switched on and off at need

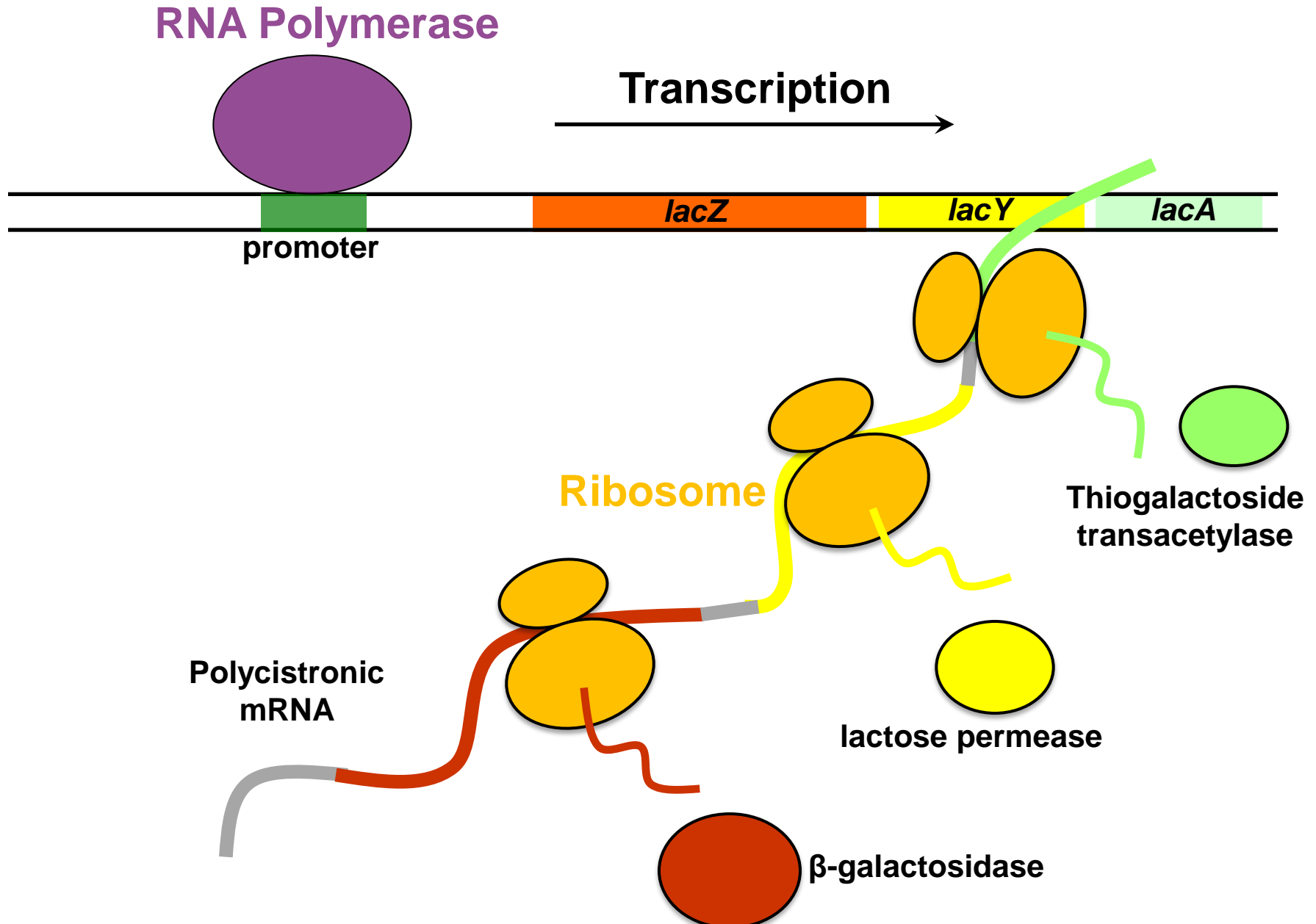
Lac Operon: A Classic Example of Bacterial Gene Expression Control

Operon: Cluster of genes, related by function, regulated by a single promoter and transcribed into one mRNA (polycistronic).

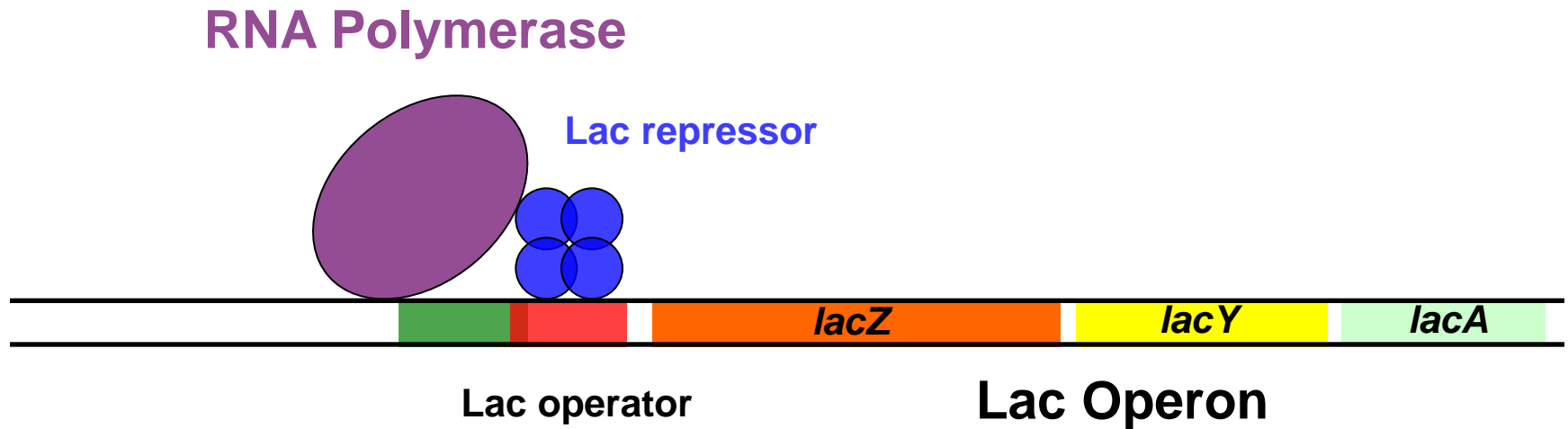


<i>lacZ</i>	β -galactosidase	Breaks lactose into galactose and glucose.
<i>lacY</i>	lactose permease	Imports lactose into the bacterial cell.
<i>lacA</i>	thiogalactoside transacetylase	Cell detoxification.

Functional Outcome of Lac Operon

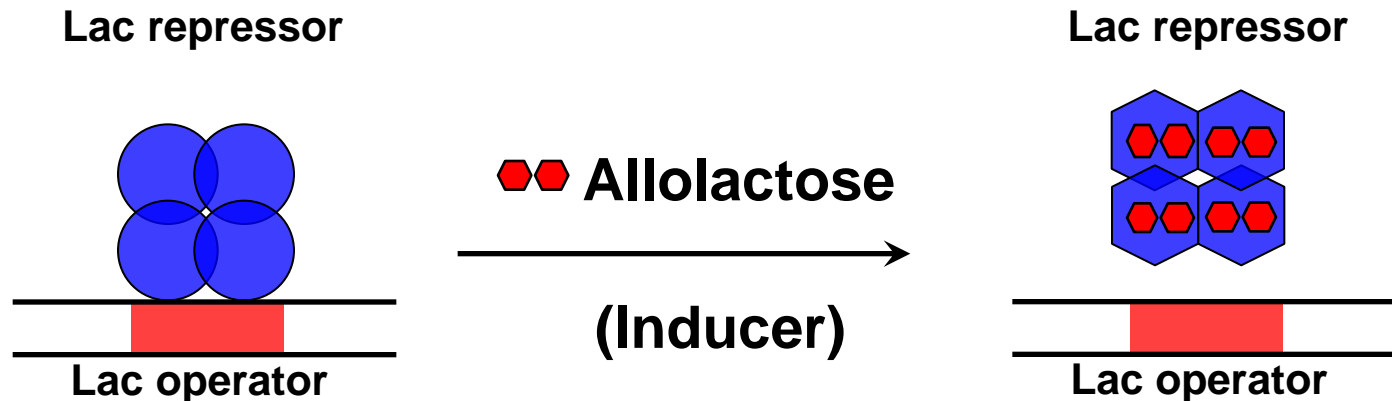
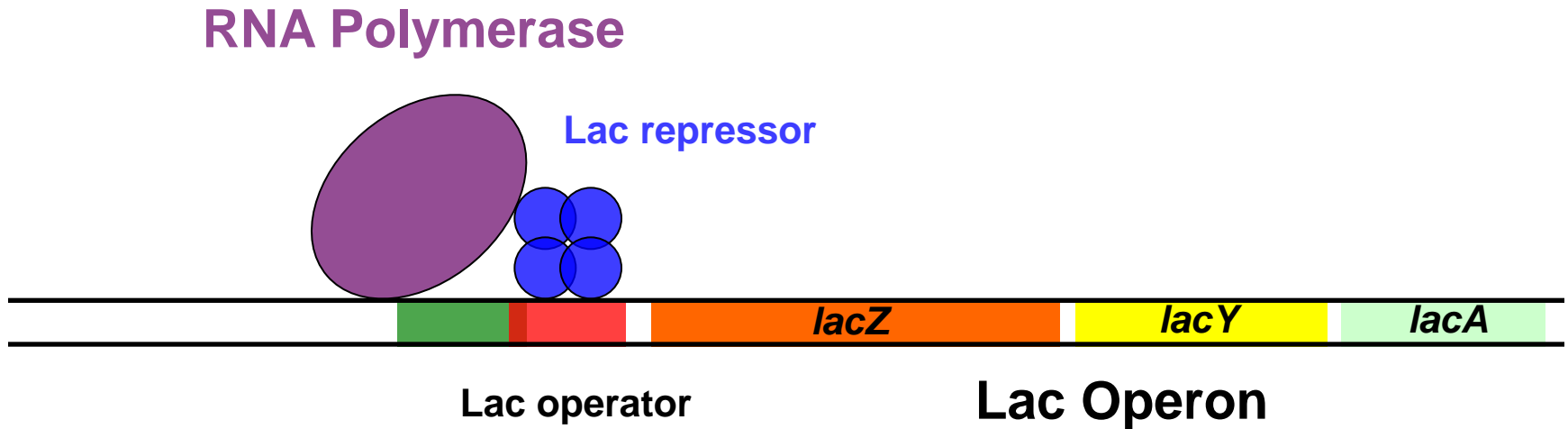


Lac repressor is a negative regulator of the Lac operon



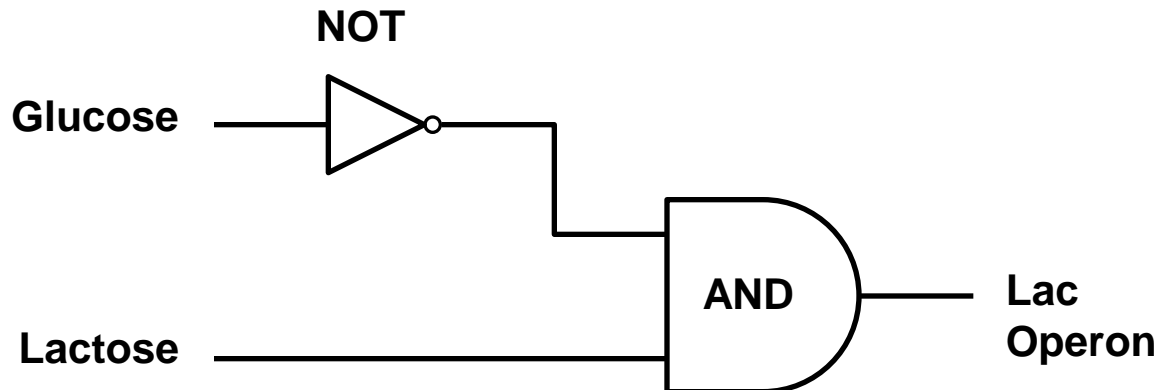
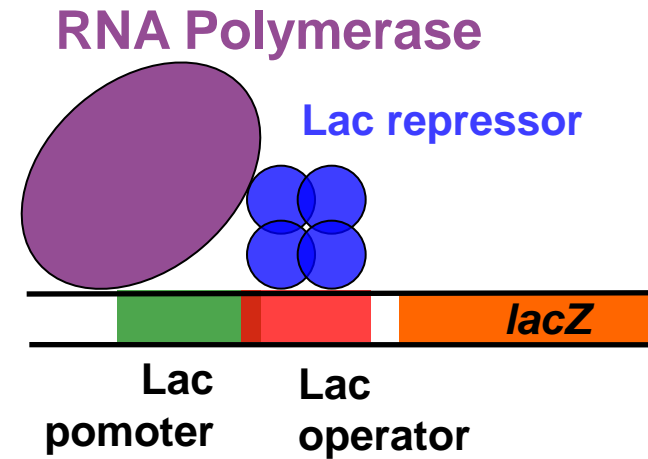
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Lactose (Allolactose) Can Displace Lac Repressor From the Operator Site

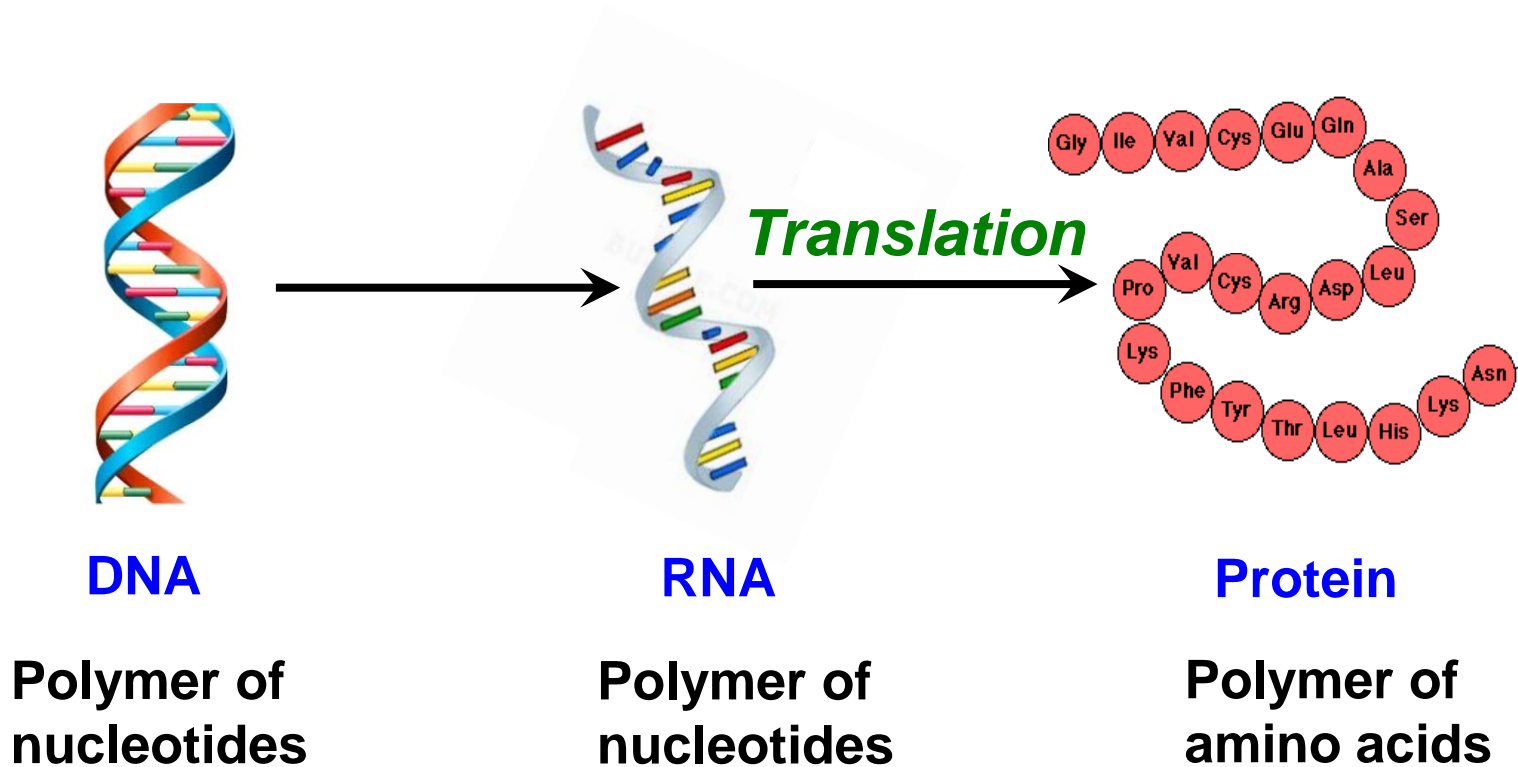


Four Possible Situations

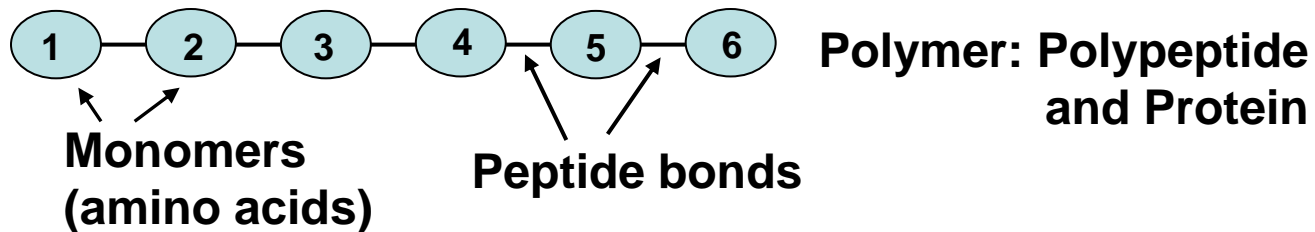
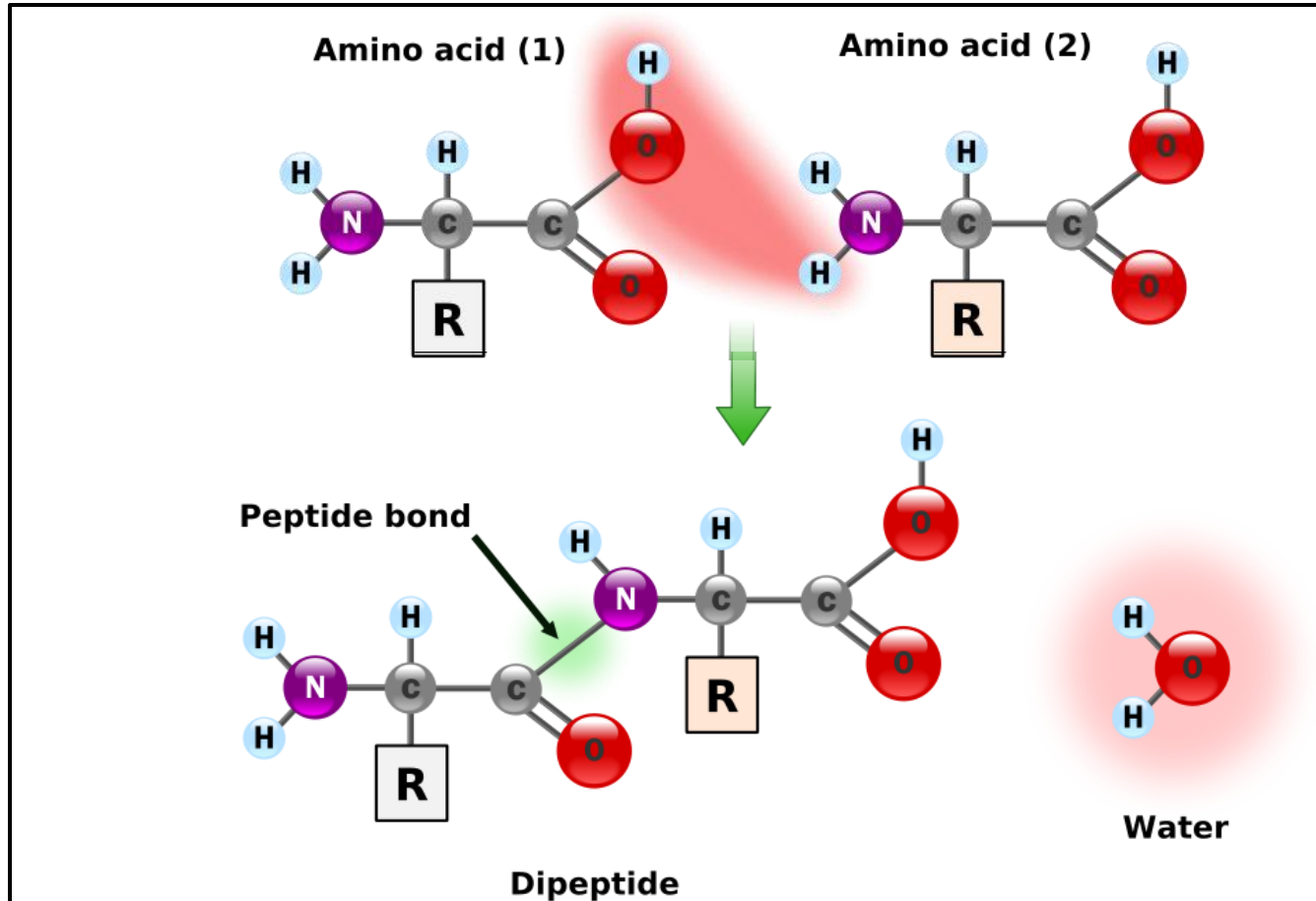
Glucose	Lactose	Lac repressor bound	Lac operon
1	0	YES	OFF (0)
1	1	YES	OFF (0)
0	1	NO	ON (1)
0	0	YES	OFF (0)



Translation



How Amino Acids are Linked Together



Translation

Template for protein synthesis

5' ————— 3' mRNA

5' UAAGGAGA AUCGUC AUG AAGAGGCC UAA UUA 3'

(RBS) Start codon Stop codon

Met—Lys—Arg—Pro.....
Polypeptide

Diagram illustrating the process of translation initiation. The ribosome is shown with the mRNA threaded through it. The mRNA sequence is 5'U AAUCGUC AUG AAGAGGCC... UAA UUA 3'. The start codon AUG is highlighted in green, and the stop codon UAA is highlighted in red. A nascent polypeptide chain is shown emerging from the ribosome.

► In Eukaryotes, 5' 7mG cap is recognized by ribosome

Protein synthesizing machinery

Ribosome: (rRNA + proteins)

30S

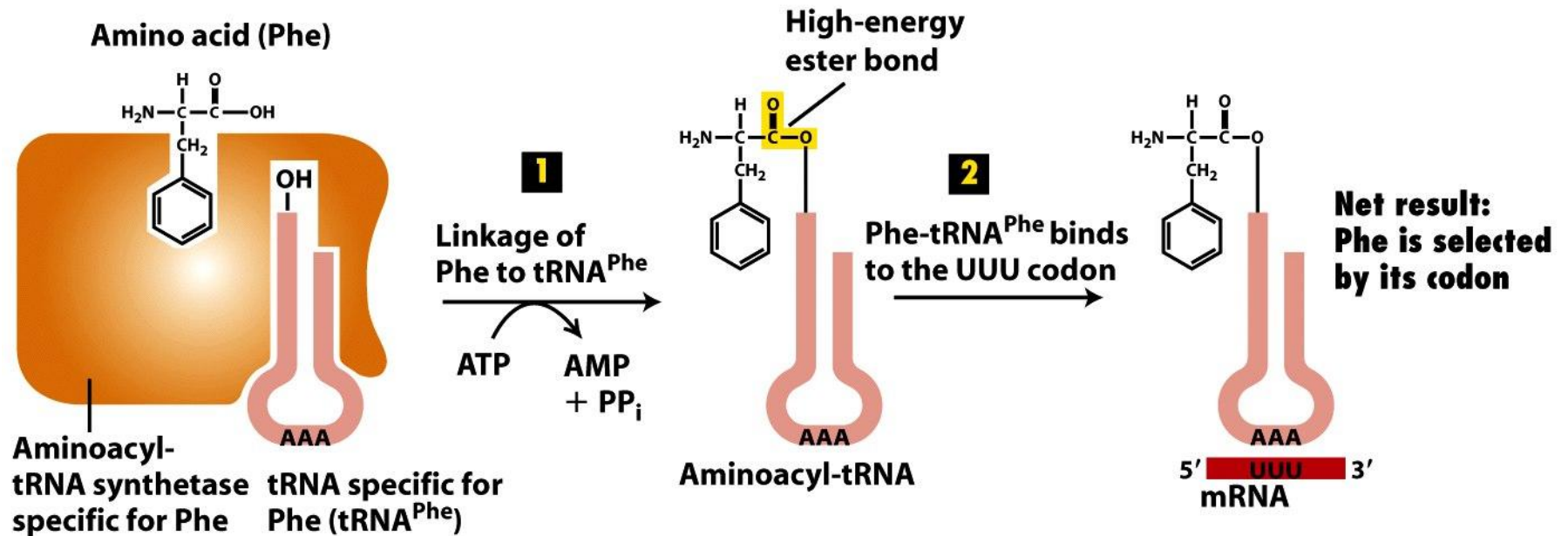
50S

16S rRNA
21 proteins

23S rRNA
5S rRNA
34 proteins

A 3D ribbon diagram of a tRNA molecule. The structure is L-shaped. The T arm is at the top left, the D arm is on the left, the V loop is on the right, the acceptor stem is at the top right, and the anticodon arm is at the bottom right. The labels are in black text.

How Correct Amino Acids are Selected During Protein Synthesis



Genetic code

Genetic code is the relation between the sequence of bases in DNA (or its RNA transcripts) and the sequence of amino acids in proteins

A codon is a set of 3 nucleotides that specifies a particular amino acid

Why three nucleotides?

64 Codons present. Three of them (UAA, UAG, UGA) can't code any amino acids, called STOP codons

AUG serves as the “initiator” or “start codon, which starts the synthesis of a protein

We have 61 codons that code for amino acids, and we have 20 amino acids. So, one codon may specify more than one amino acids

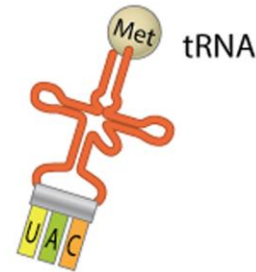


Khorana, Nirenberg, Holley
Nobel Prize in 1968

Genetic code

		Second Letter					
		U	C	A	G		
1st letter	U	UUU Phe UUC UUA Leu UUG	UCU UCC Ser UCA UCG	UAU Tyr UAC UAA Stop UAG Stop	UGU Cys UGC UGA Stop UGG Trp	U C A G	3rd letter
	C	CUU CUC Leu CUA CUG	CCU CCC Pro CCA CCG	CAU His CAC CAA Gln CAG	CGU CGC Arg CGA CGG	U C A G	
	A	AUU AUC Ile AUA AUG Met	ACU ACC Thr ACA ACG	AAU Asn AAC AAA Lys AAG	AGU Ser AGC AGA Arg AGG	U C A G	
	G	GUU GUC Val GUA GUG	GCU GCC Ala GCA GCG	GAU Asp GAC GAA Glu GAG	GGU GGC Gly GGA GGG	U C A G	

Translation: Involved Machineries and Processes

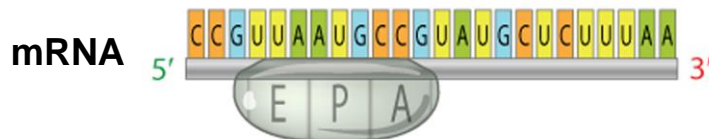


mRNA

Ribosome

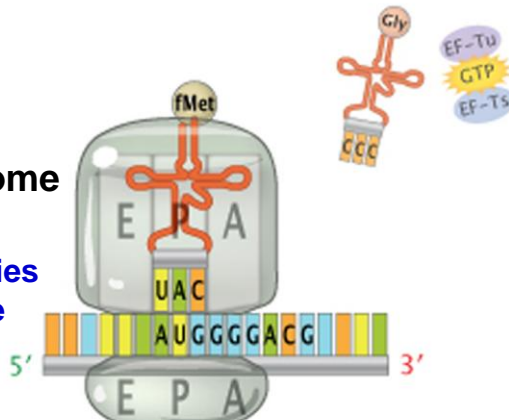
Amino acid

tRNA

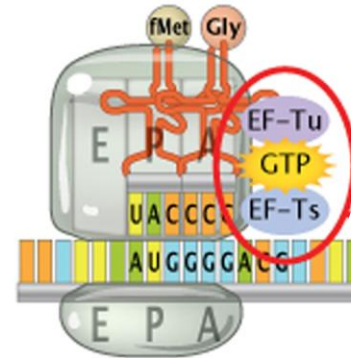


50S ribosome

tRNA^{fMet} occupies the P site of the ribosome

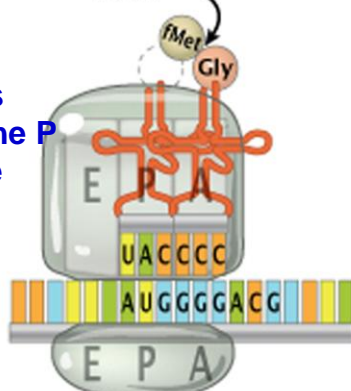


Next charged tRNA and associated translation factors enter the A site

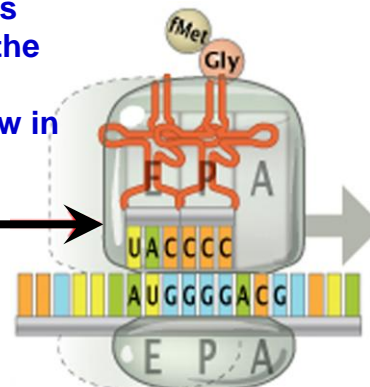


Peptide bond

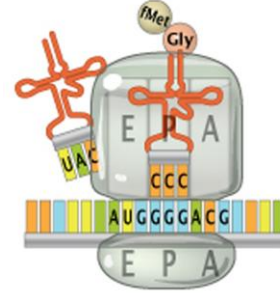
Peptide bond forms between the aa in the P and A sites, and the tRNA in the P site releases its aa



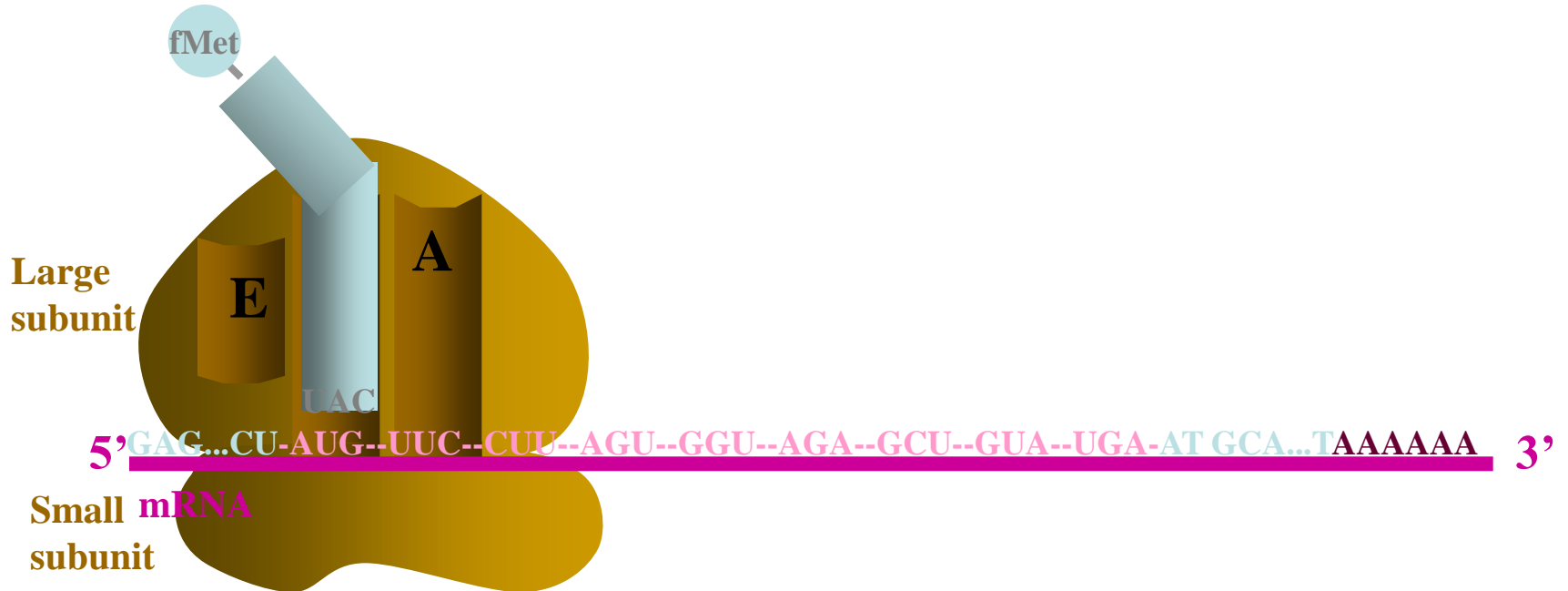
The ribosome moves down the mRNA to the next codon and the uncharged tRNA now in the E site



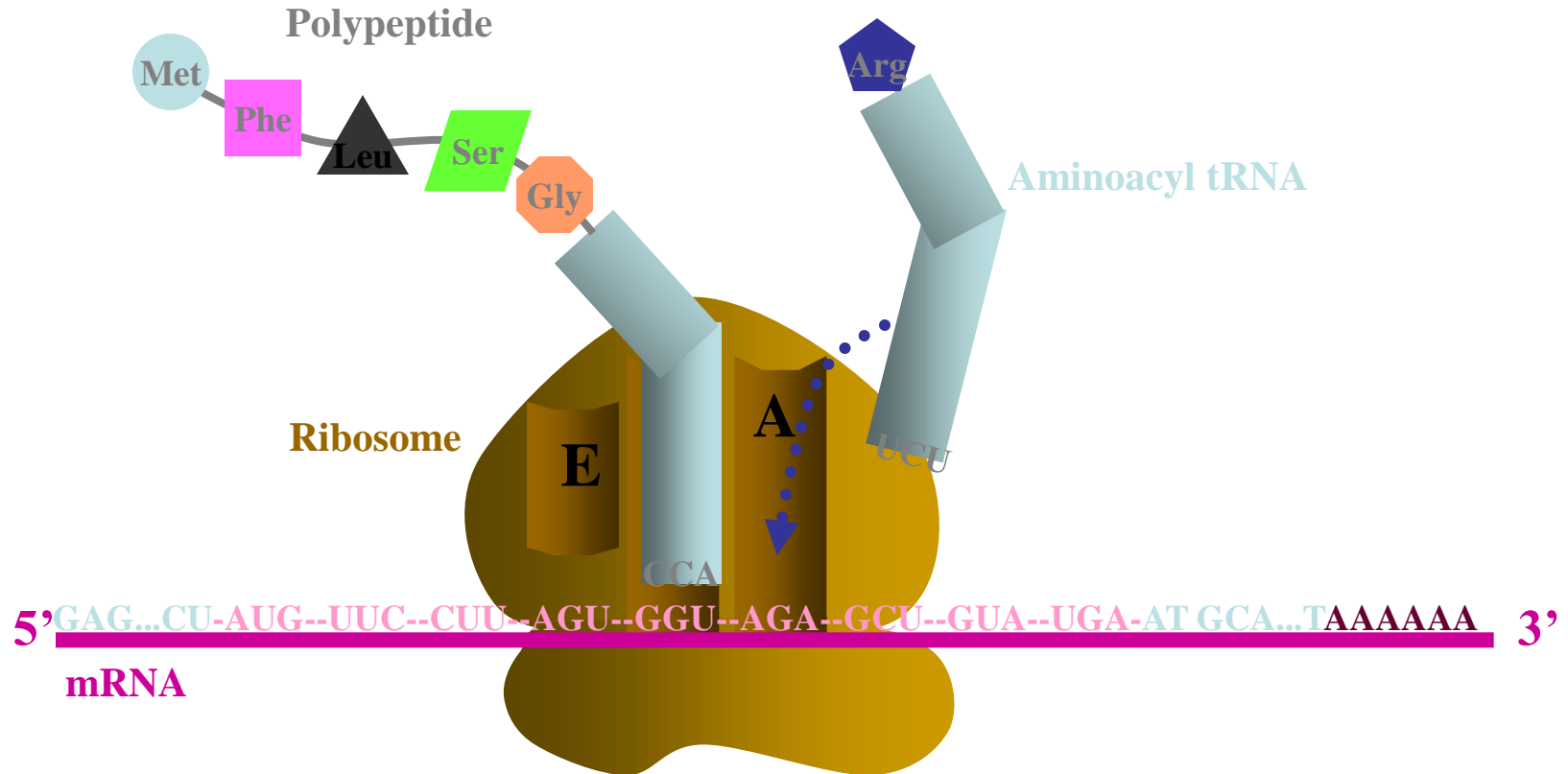
E site tRNA releases and A site is now ready to receive another tRNA



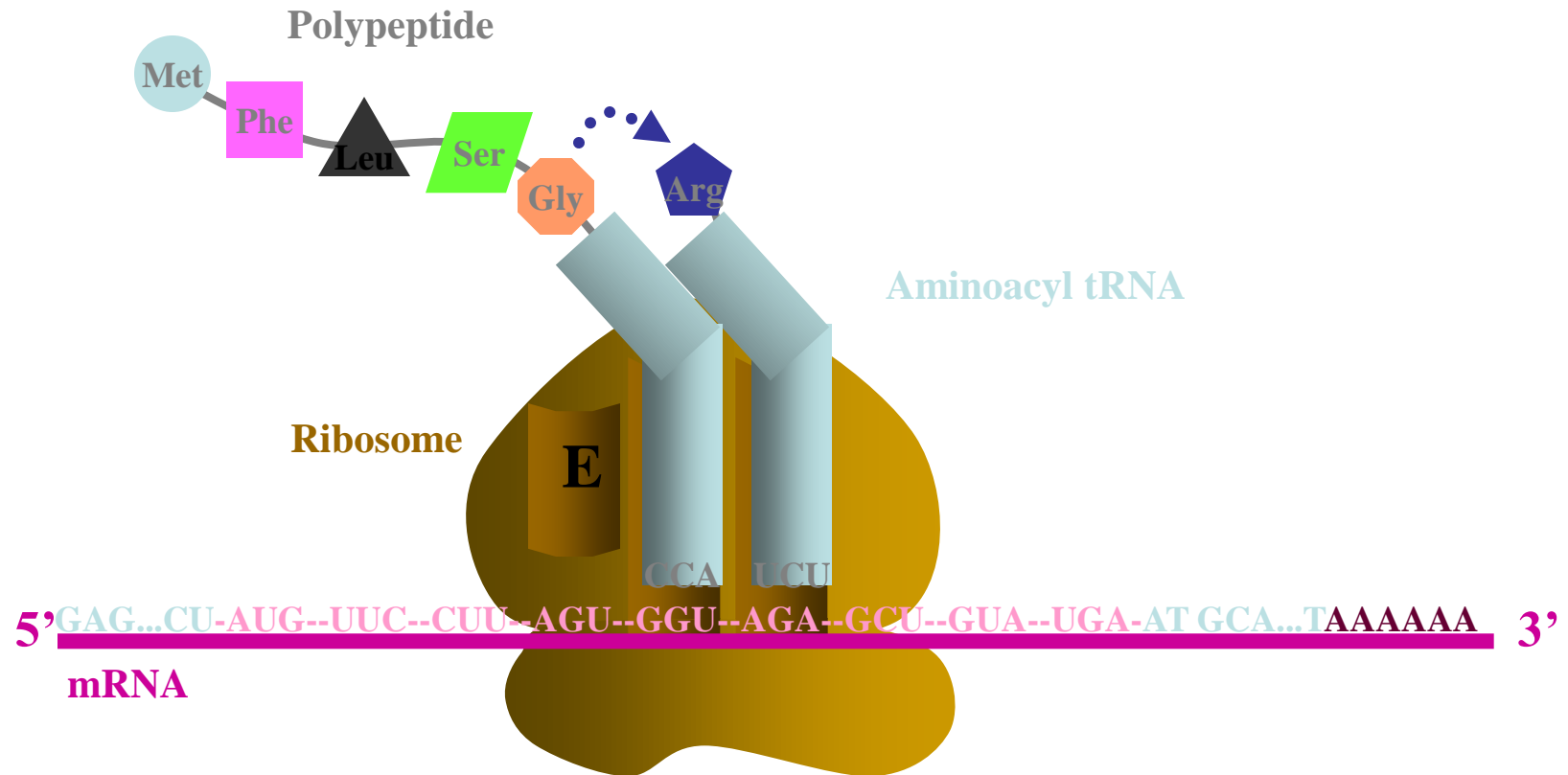
Translation - Initiation



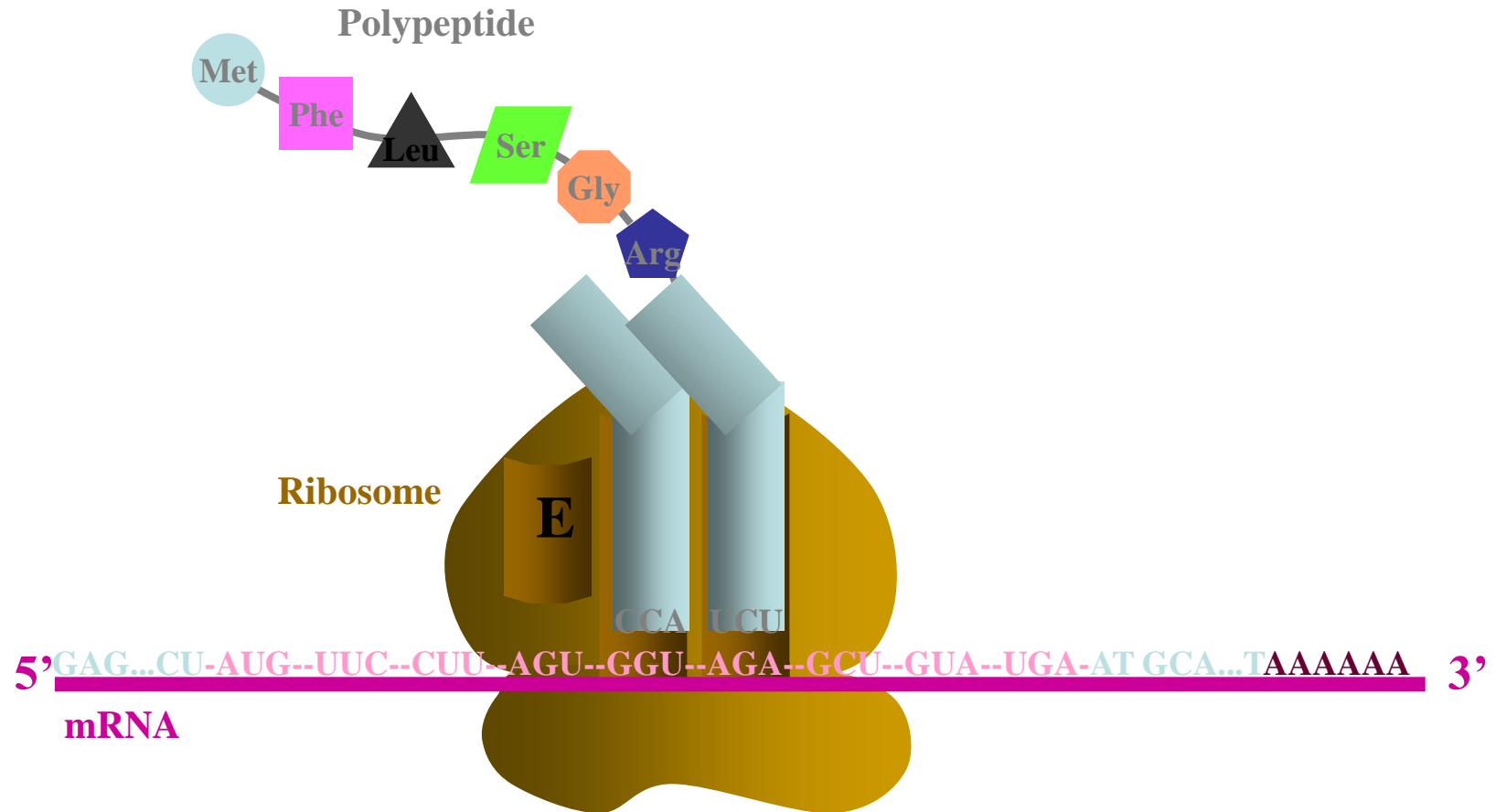
Translation - Elongation



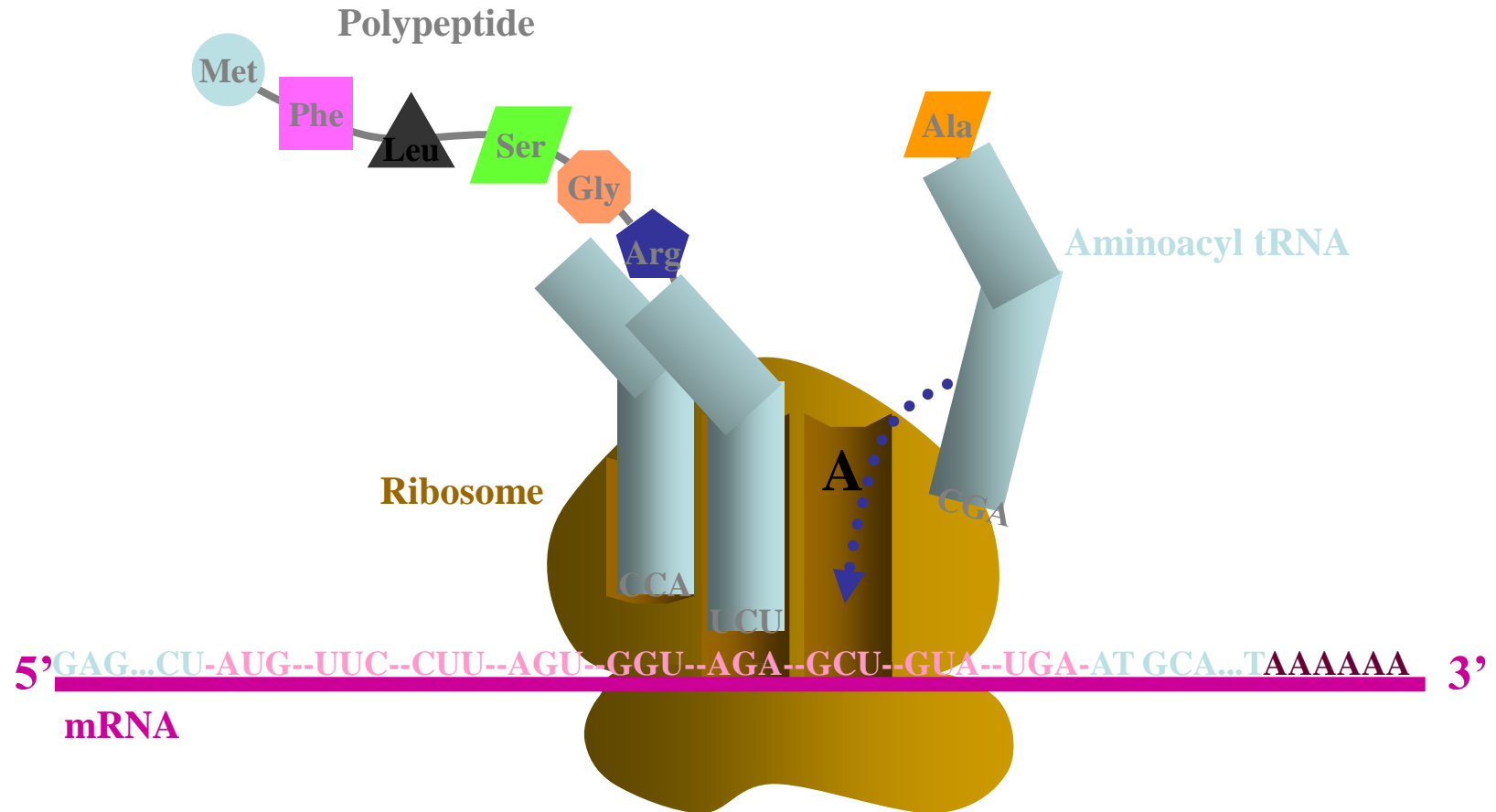
Translation - Elongation



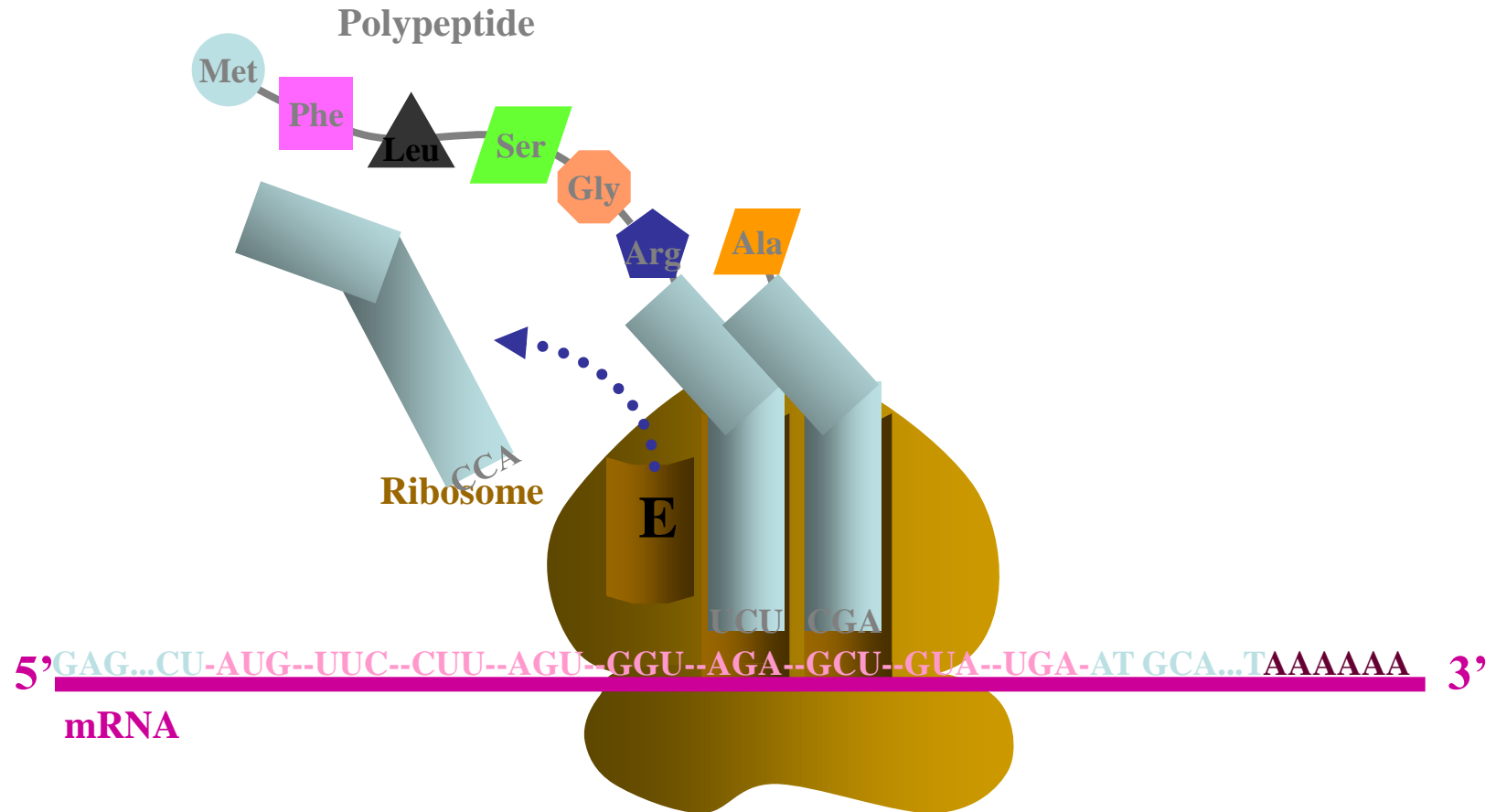
Translation - Elongation



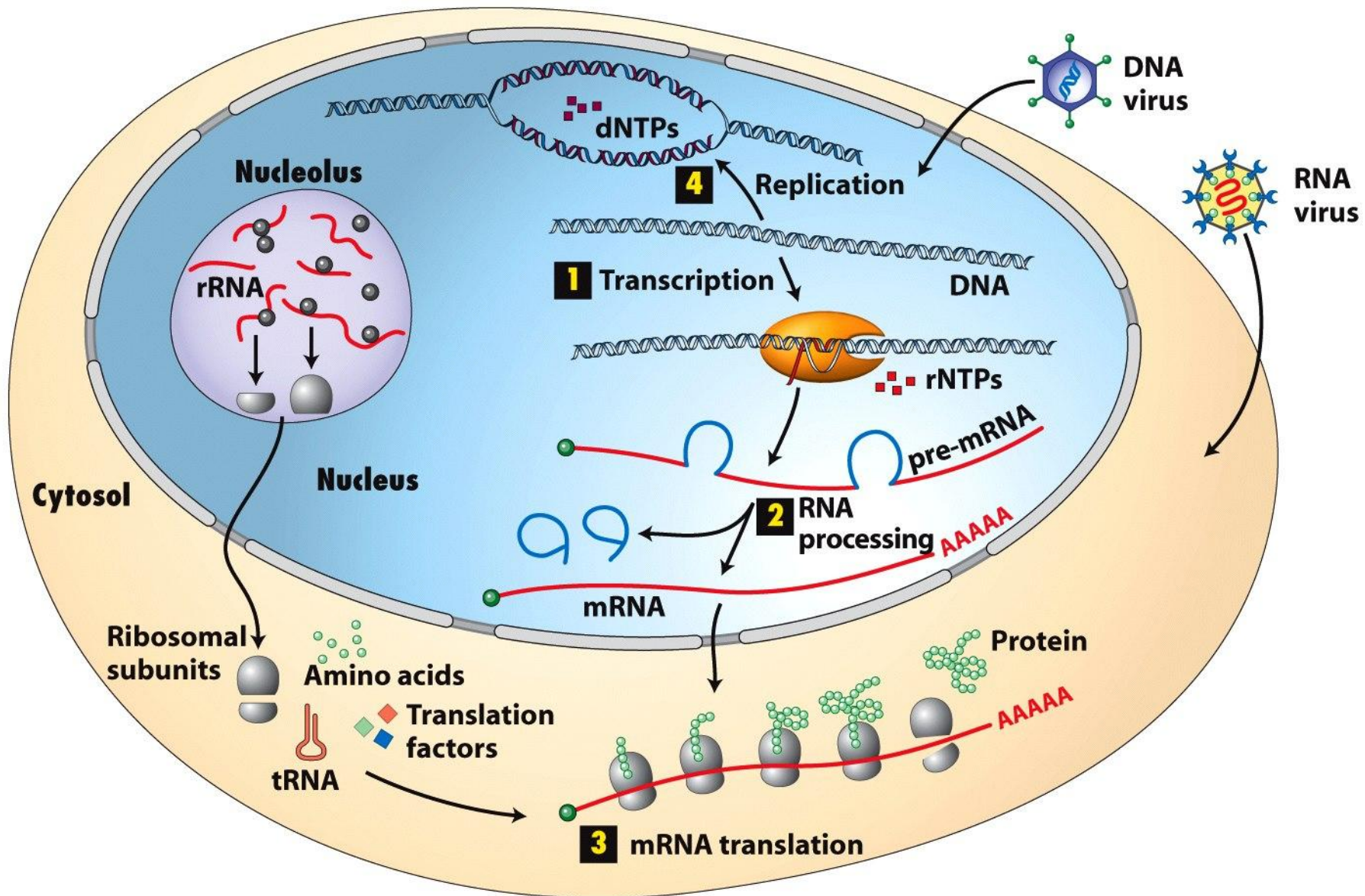
Translation - Elongation



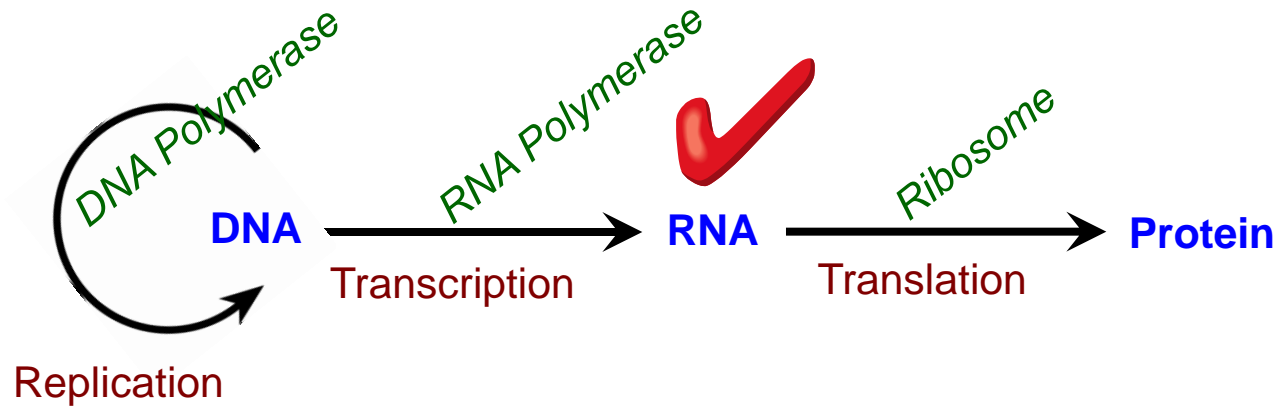
Translation - Elongation



Translation: At a Glance

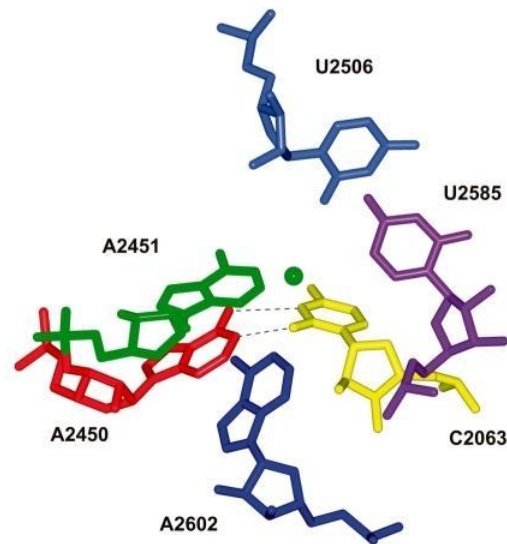
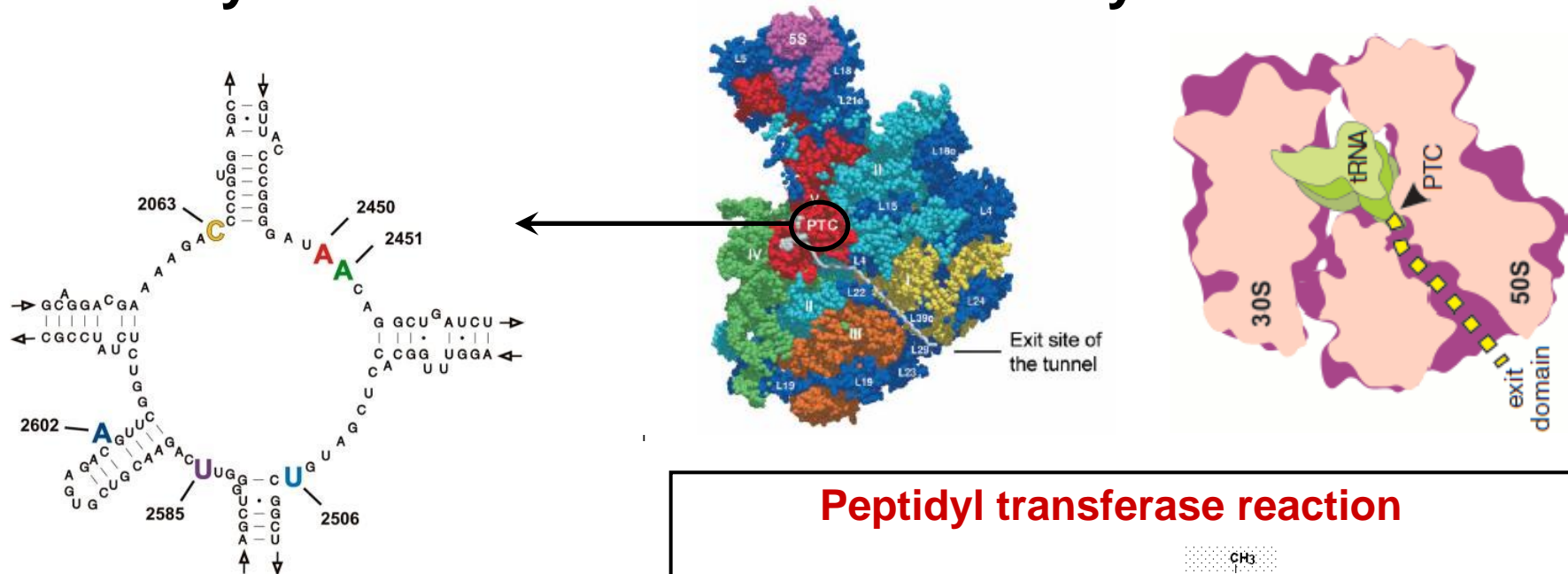


Which Came First? **Nucleic acids or Proteins**



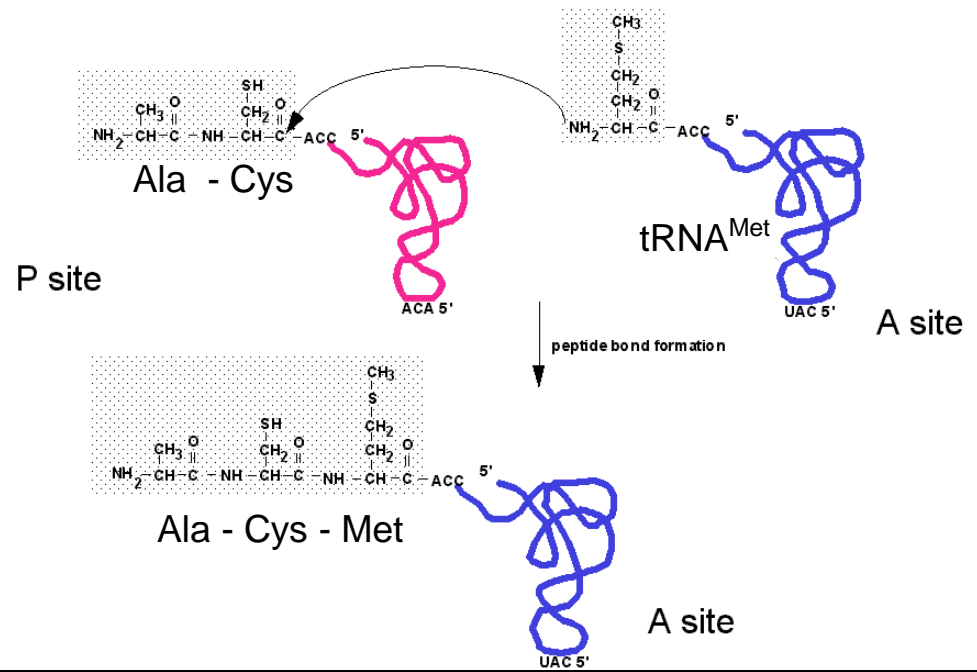
► RNA has enzymatic activity

What Happens Inside the Ribosome? Chemical and Physical Consideration of Protein Synthesis



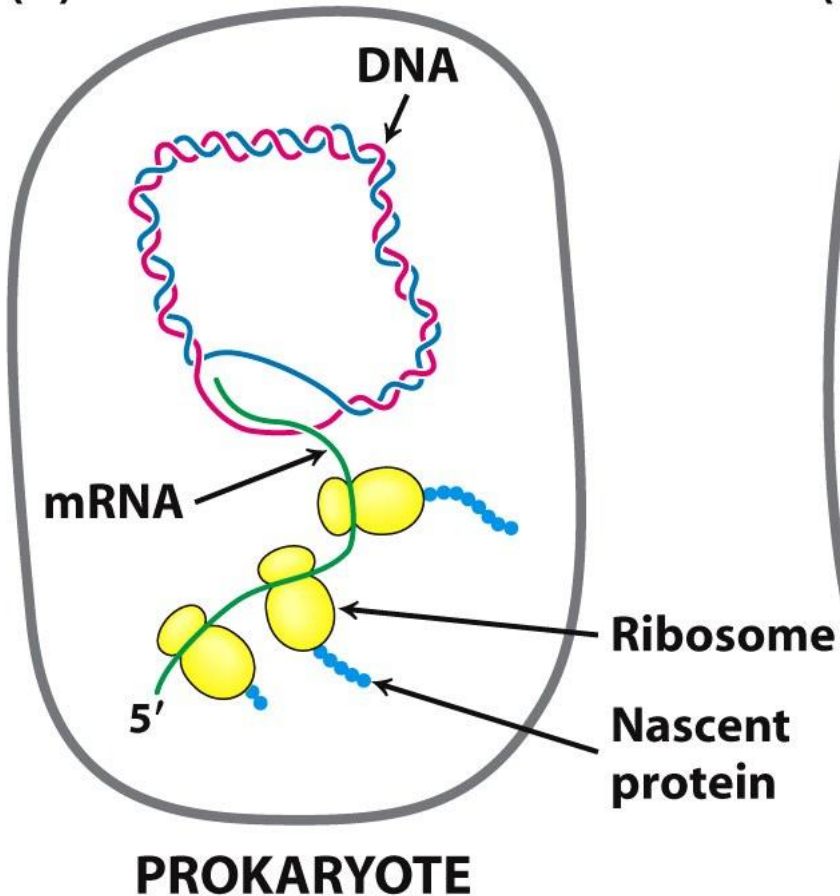
Peptidyl transferase center

Peptidyl transferase reaction

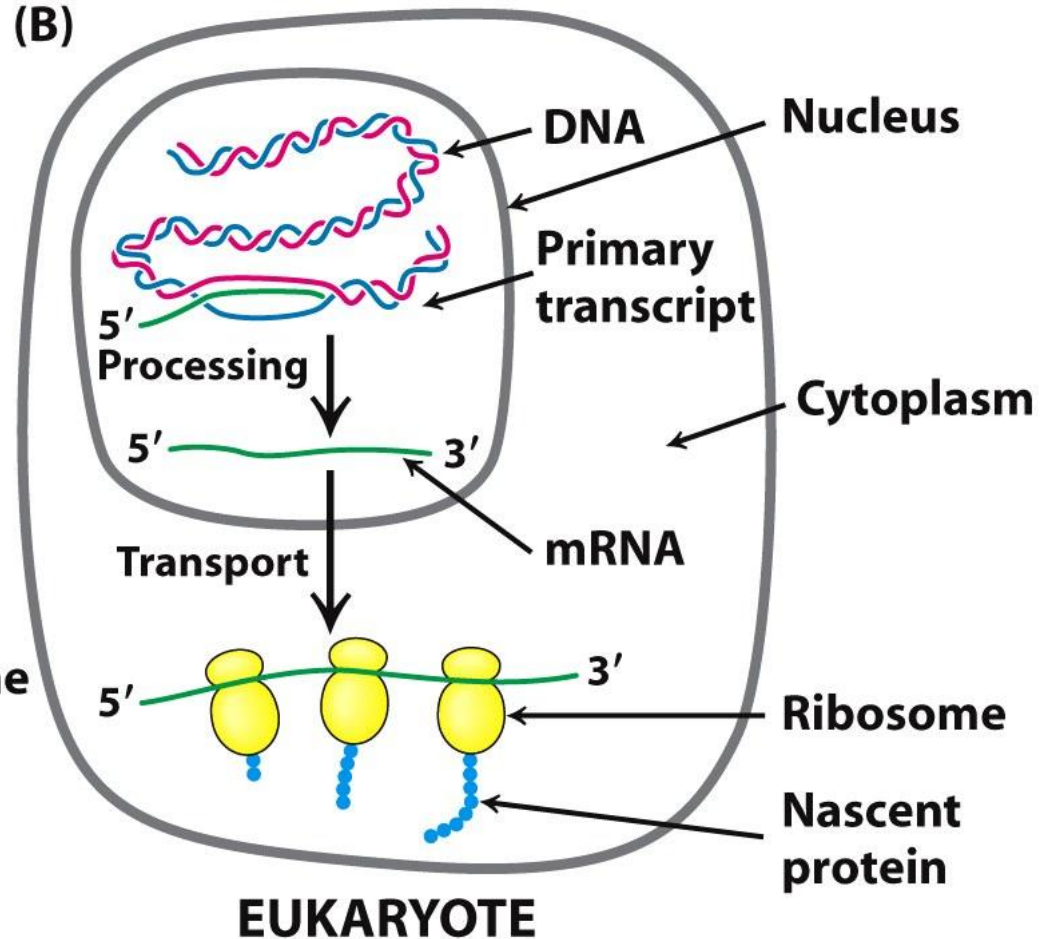


Time, Space and Correlation between Transcription and Translation

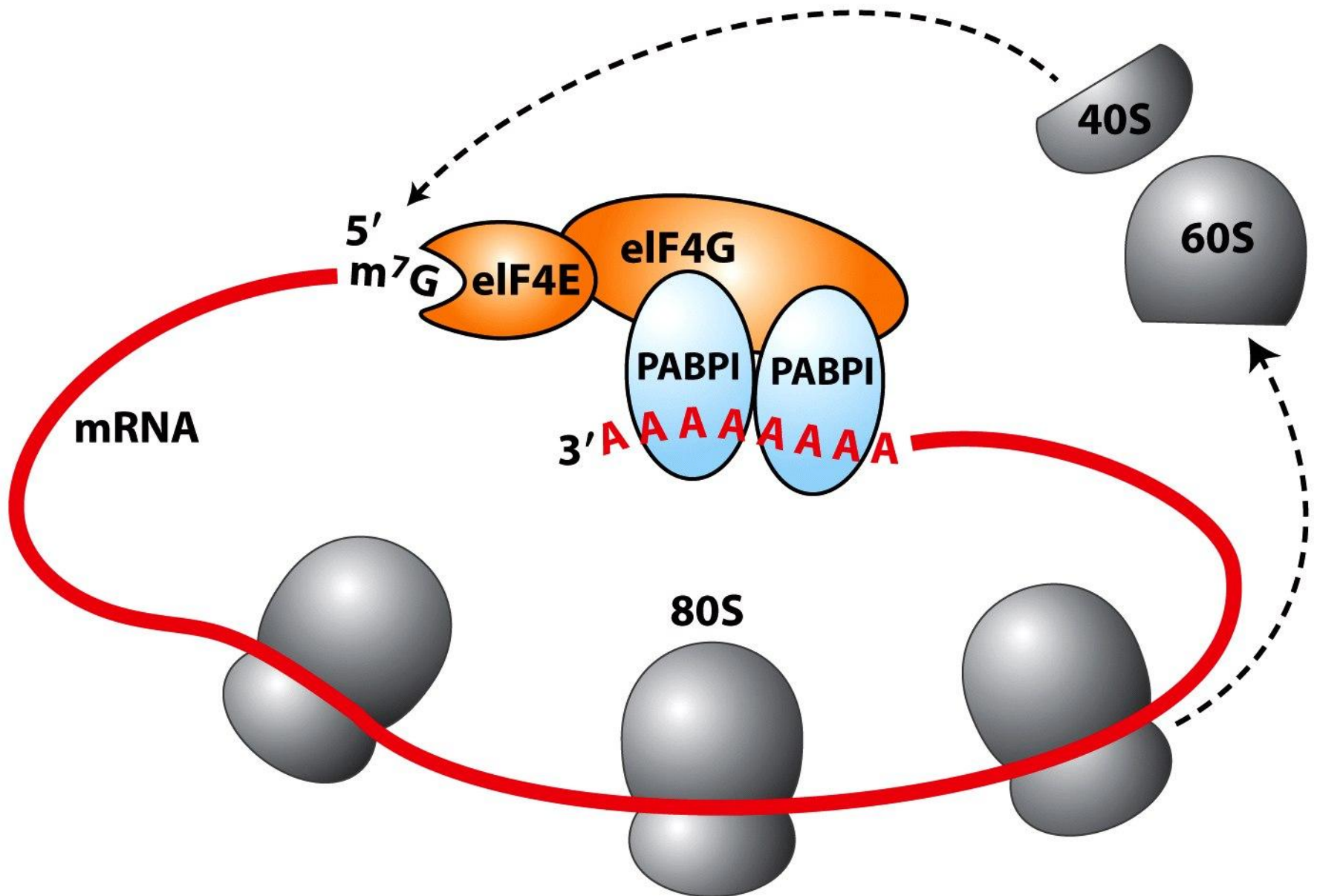
(A)



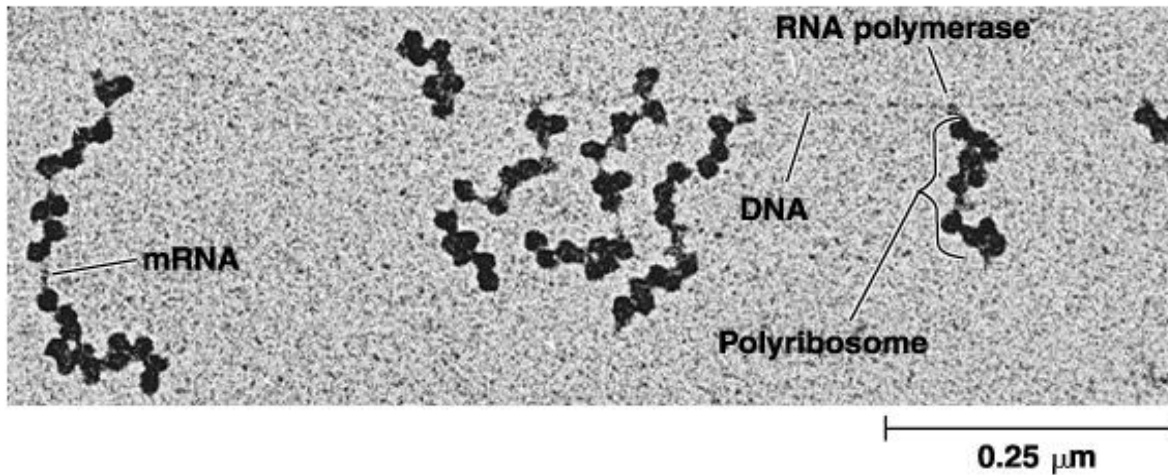
(B)



Time, Space and Correlation between Transcription and Translation



Time, Space and Correlation between Transcription and Translation



Jonathan Warner

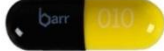
Translation Machineries: Attractive Targets For Therapeutics

Tetracycline

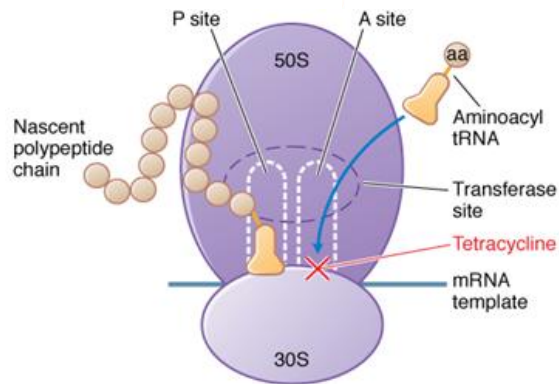
250 mg



500 mg

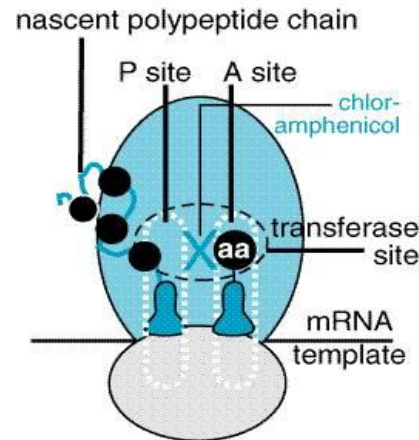


Binds to the 30S ribosome, and blocks binding of aminoacyl-tRNA to the A-site



Chloramphenicol

Blocks the peptidyl transferase reaction on 50S ribosomes



Streptomycin

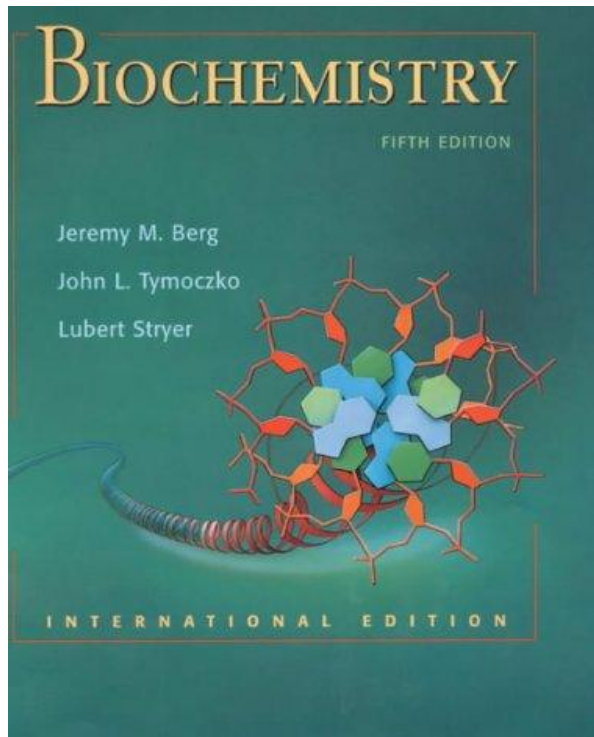
Binds to the 30S ribosome, prevents the transition from initiation to chain-elongation

Erythromycin

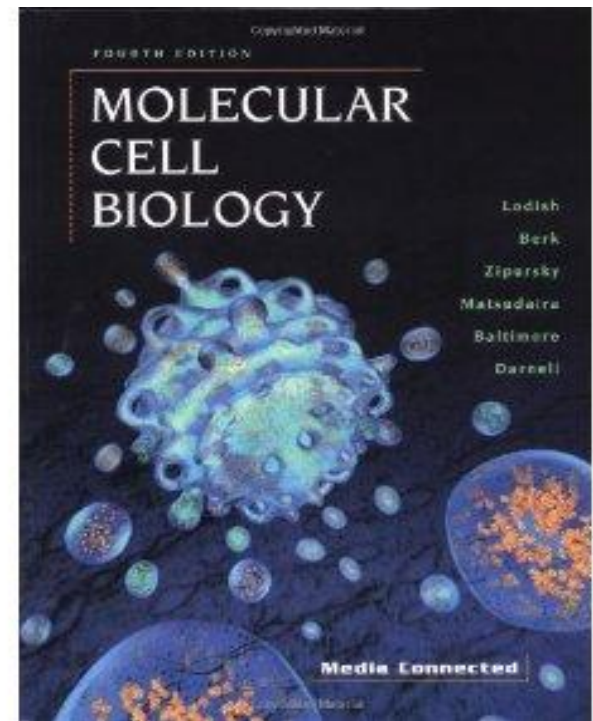
Binds to the 50S ribosome, and blocks the translocation

Why ribosome is an attractive target for the development of antibiotics?

Suggested Textbook...



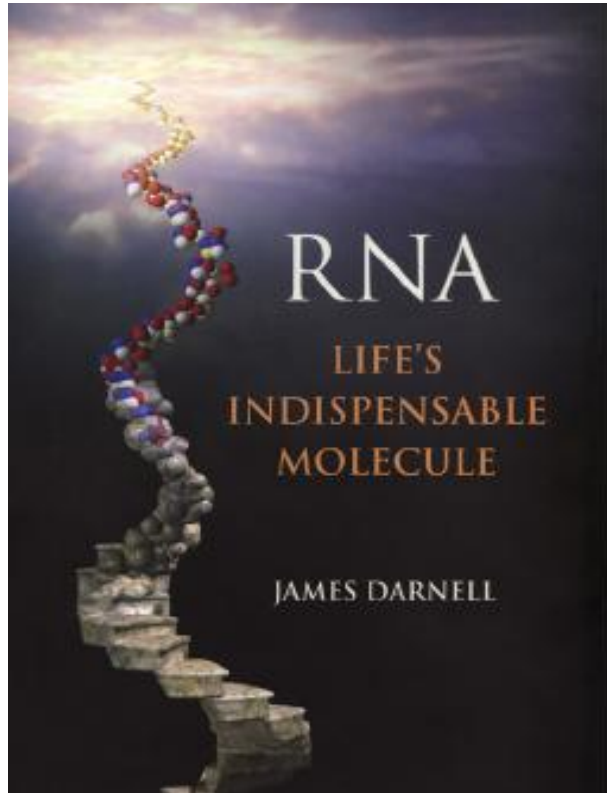
Stryer...



Baltimore, Lodish..

Extra Resources

Further Reading...



James Darnell

Videos...

mRNA synthesis (Transcription)

https://www.youtube.com/watch?v=_C9Un4dlpR4

Protein synthesis (Translation)

<https://www.youtube.com/watch?v=lkq9AcBcohA>

Overview

<https://www.youtube.com/watch?v=gG7uCskUOrA>