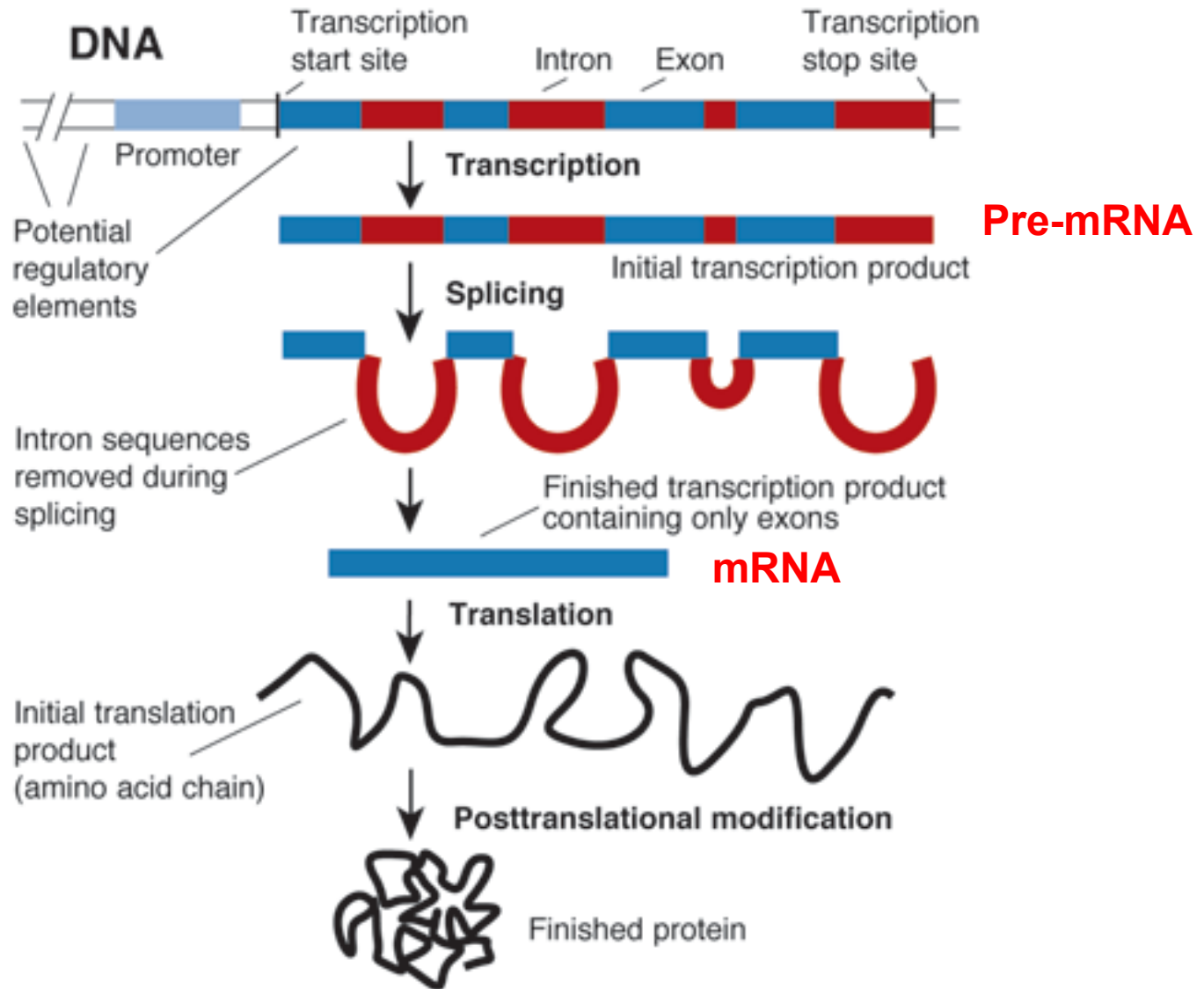


Gene Structure & Protein Synthesis



De-coding a gene sequence

5' -ATGCCTAGGTACCTATGA-3'

3' -TACGGATCCATGGATACT-5'

DNA

↓ **Transcription**

5' -AUGCCUAGGUACCUAUGA-3'

mRNA

↓ **decoded as**

5' -AUG CCU AGG UAC CUA UGA-3'



Translation

N-MET-PRO-ARG-TYR-LEU-C

Protein

The Genetic Code

- Proteins are made by joining amino acids into long chains called polypeptides. **There are 20 different amino acids.** The properties of the protein are determined by the order of amino acids.
- **The genetic code is read three letters at a time** so that each “word” is three bases long. Each three letter “word” is a **codon.**
- A **codon** consists of three consecutive nucleotides that **specify a single amino acid** that is to be added to the polypeptide.

The Genetic Code

There are **64 possible codons**. Some amino acids can be specified by **more than one codon**.

There is a **start codon, AUG** (methionine). There are three **stop codons (UAA, UAG, UGA)**.

The Genetic Code is Nearly Universal

- The process of making proteins from the information in DNA is used by nearly all cells.
- **Nearly all organisms studied to date use the same genetic code.**
- Because of this, we are able to use bacteria as factories to make massive amounts of proteins.
 - Insulin, growth factor, etc.

GCA	AGA									
GCC	AGG									
GCG	CGA						GGA			
GCU	CGC						GGC		AUA	
	CGG	GAC	AAC	UGC	GAA	CAA	GGG	CAC	AUC	
	CGU	GAU	AAU	UGU	GAG	CAG	GGU	CAU	AUU	
Ala	Arg	Asp	Asn	Cys	Glu	Gln	Gly	His	Ile	
A	R	D	N	C	E	Q	G	H	I	
UUA										
UUG										
CUA										
CUC										
CUG	AAA									
CUU	AAG	AUG	UUC	CCA	AGC	ACA				
			UUU	CCC	AGU	ACC			GUA	
				CCG	UCA	ACG			GUC	
				CCU	UCC	ACU			GUG	
					UCG				GUU	
					UCU		UGG	UAC		
								UAU		
Leu	Lys	Met	Phe	Pro	Ser	Thr	Trp	Tyr	Val	UAA
										UAG
										UGA
L	K	M	F	P	S	T	W	Y	V	stop

Figure 6–50. Molecular Biology of the Cell, 4th Edition.

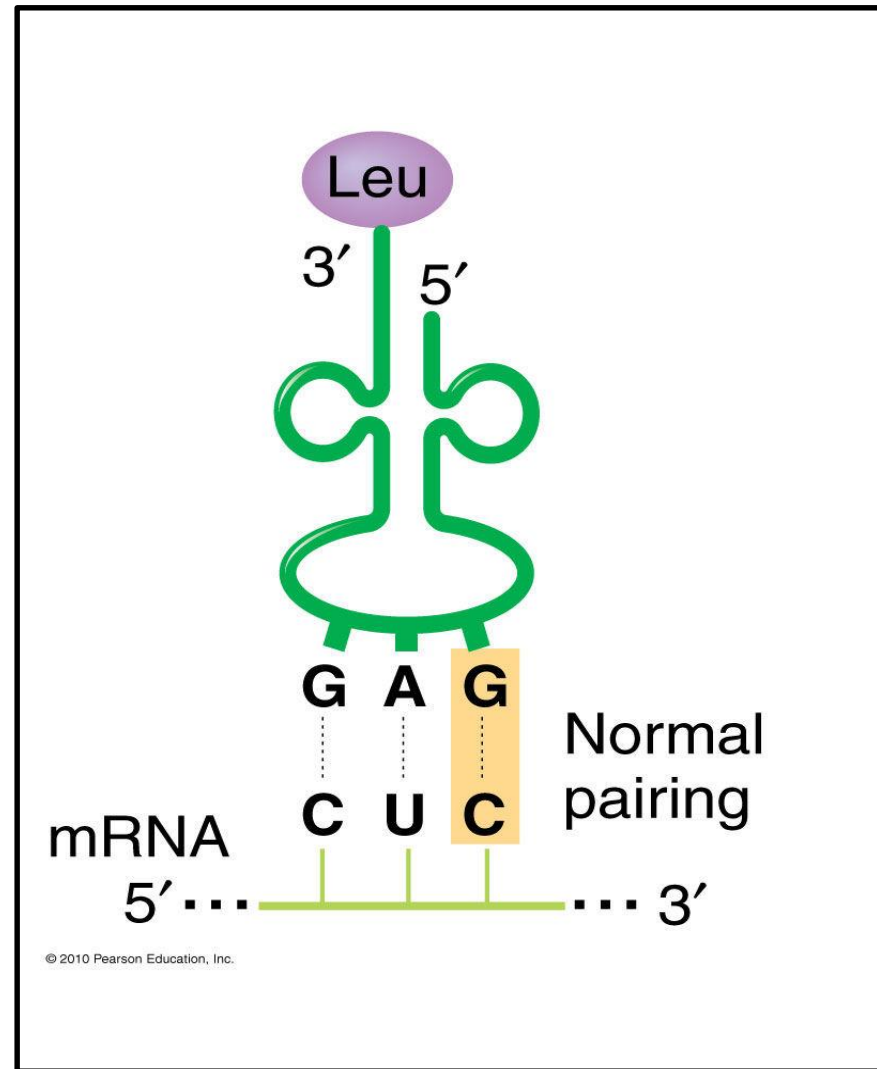
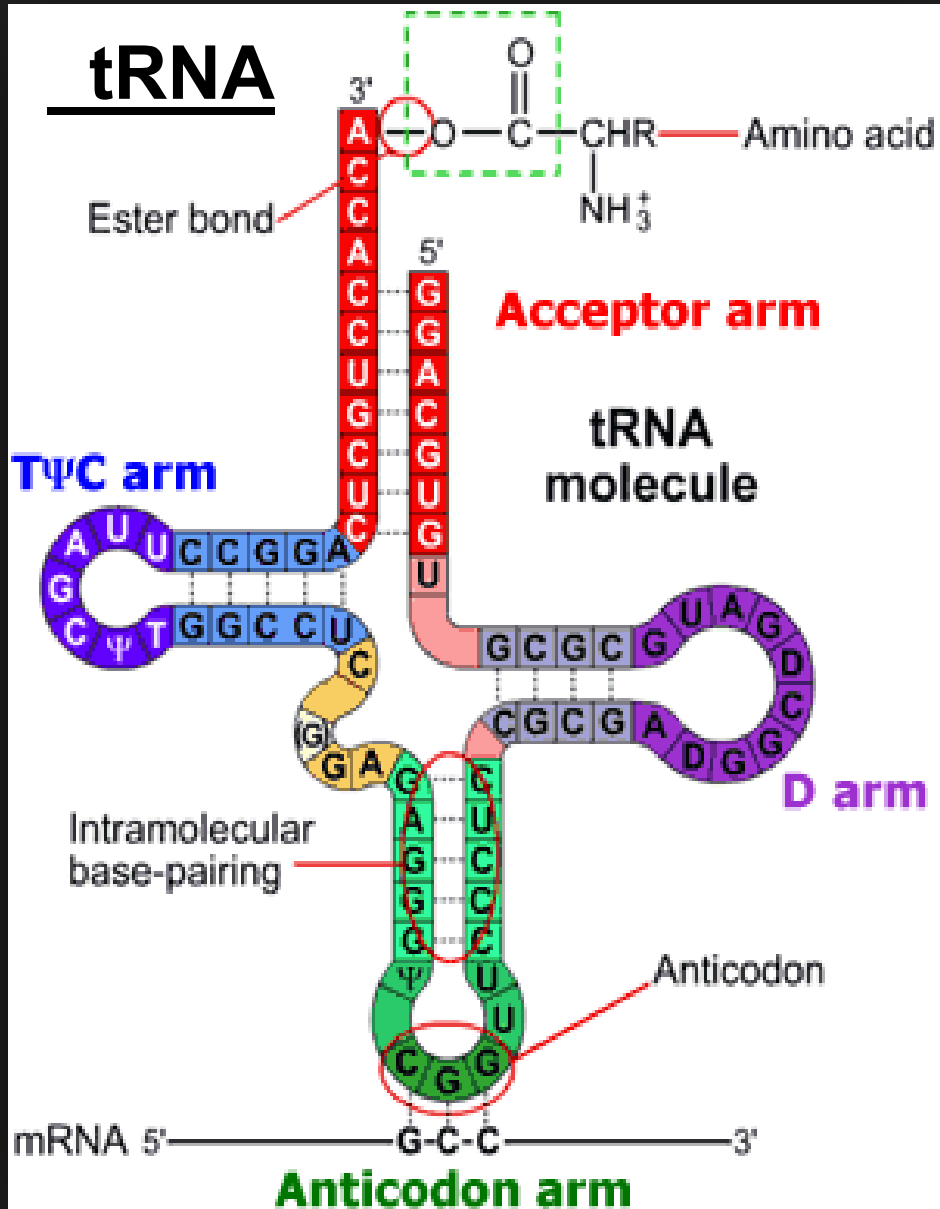
Second base in codon

		U	C	A	G	
First base in codon	U	Phe Phe Leu Leu	Ser Ser Ser Ser	Tyr Tyr STOP STOP	Cys Cys STOP Trp	U C A G
	C	Leu Leu Leu Leu	Pro Pro Pro Pro	His His Gln Gln	Arg Arg Arg Arg	U C A G
	A	Ile Ile Ile Met	Thr Thr Thr Thr	Asn Asn Lys Lys	Ser Ser Arg Arg	U C A G
	G	Val Val Val Val	Ala Ala Ala Ala	Asp Asp Glu Glu	Gly Gly Gly Gly	U C A G
		Third base in codon				

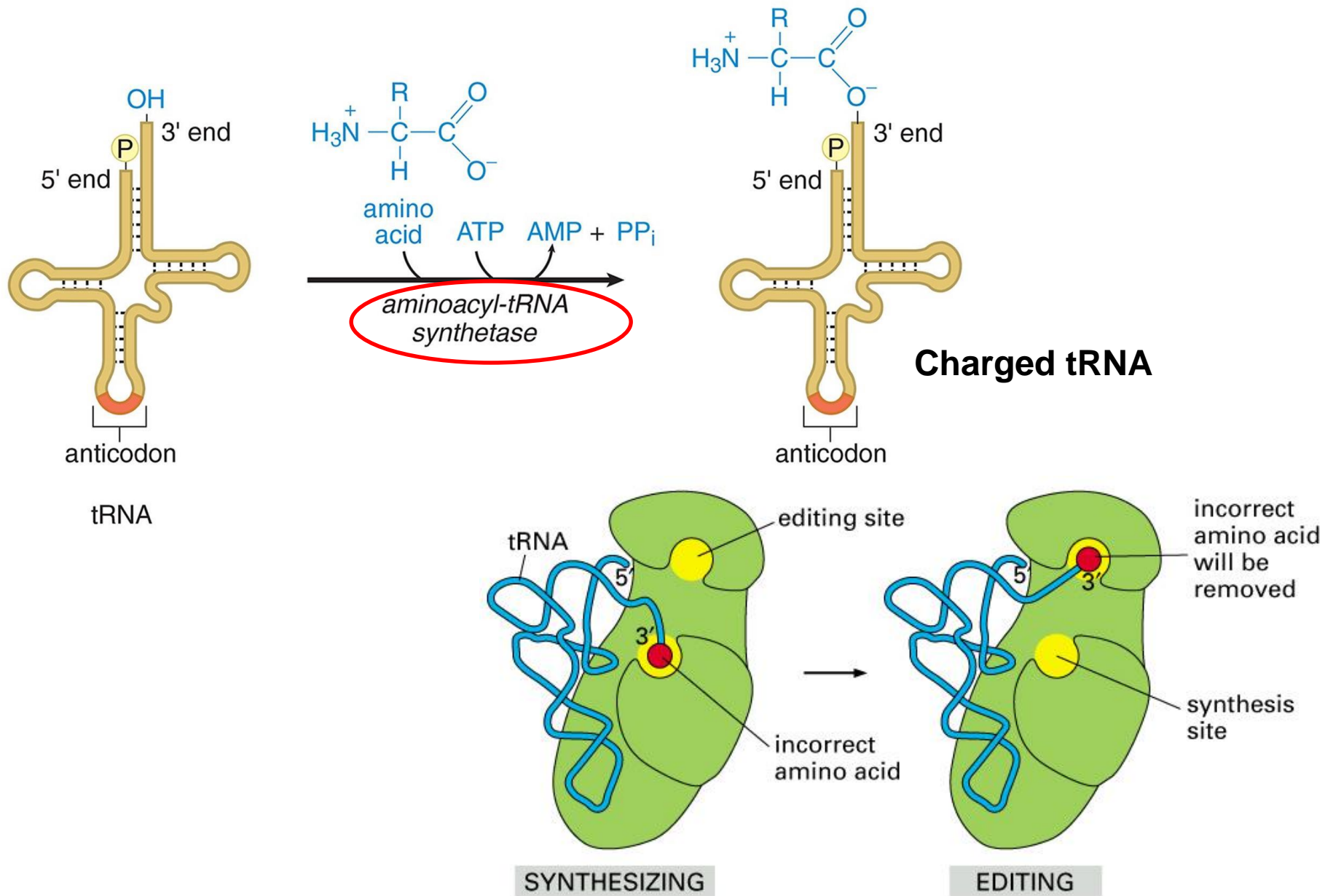
Translation

- Three types of RNA participate in translation.
 - **mRNA** carries the recipe for making the protein.
 - **tRNA and rRNA** are used to read the recipe and build the amino acid chain.
- Codons are sets of **three nucleotides** that code for specific amino acids.
 - **tRNA reads the codons** and brings the correct amino acids.

tRNA



tRNAs are activated by amino-acyl tRNA synthetases



High fidelity of amino acyl-tRNA synthetases

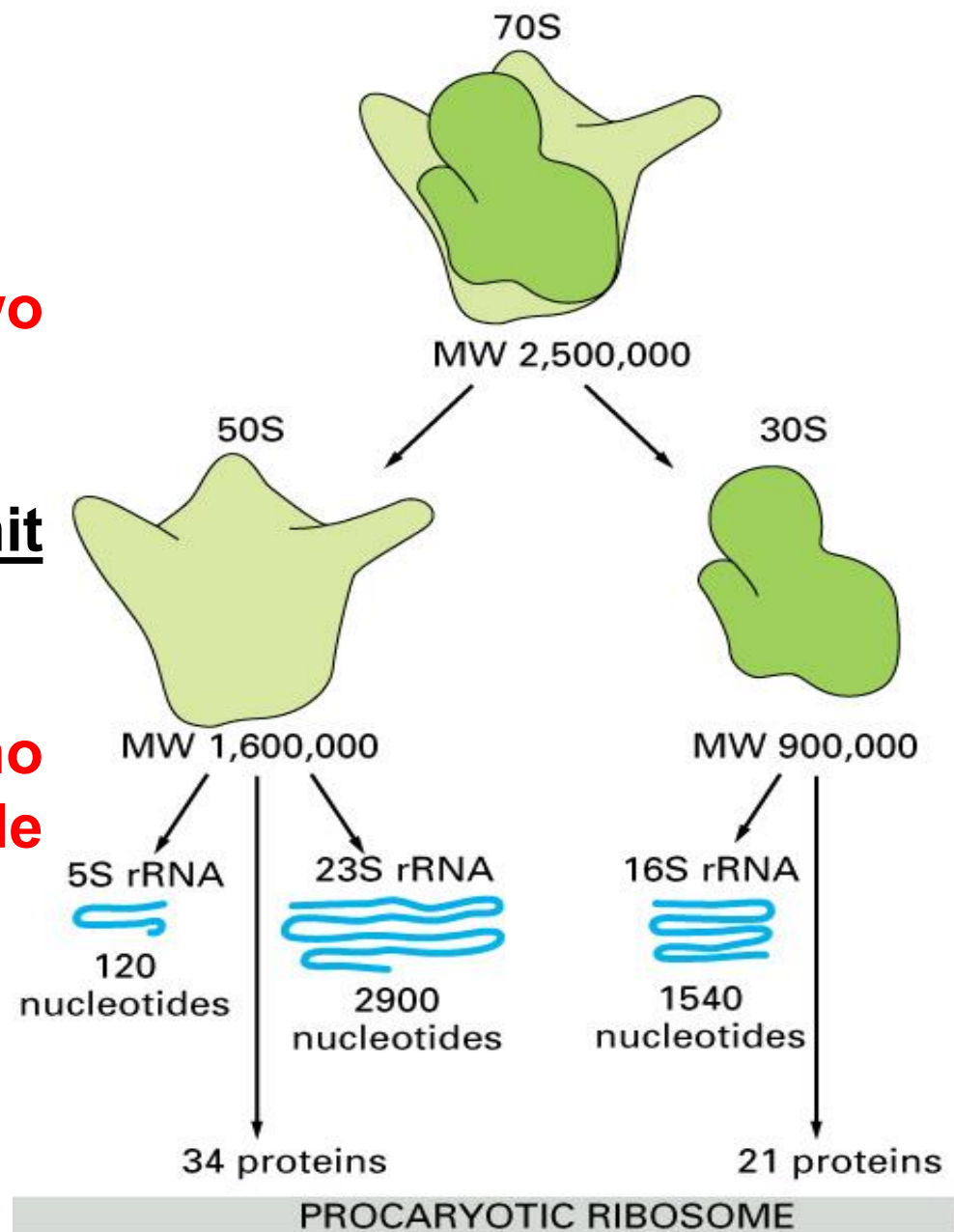
Translation

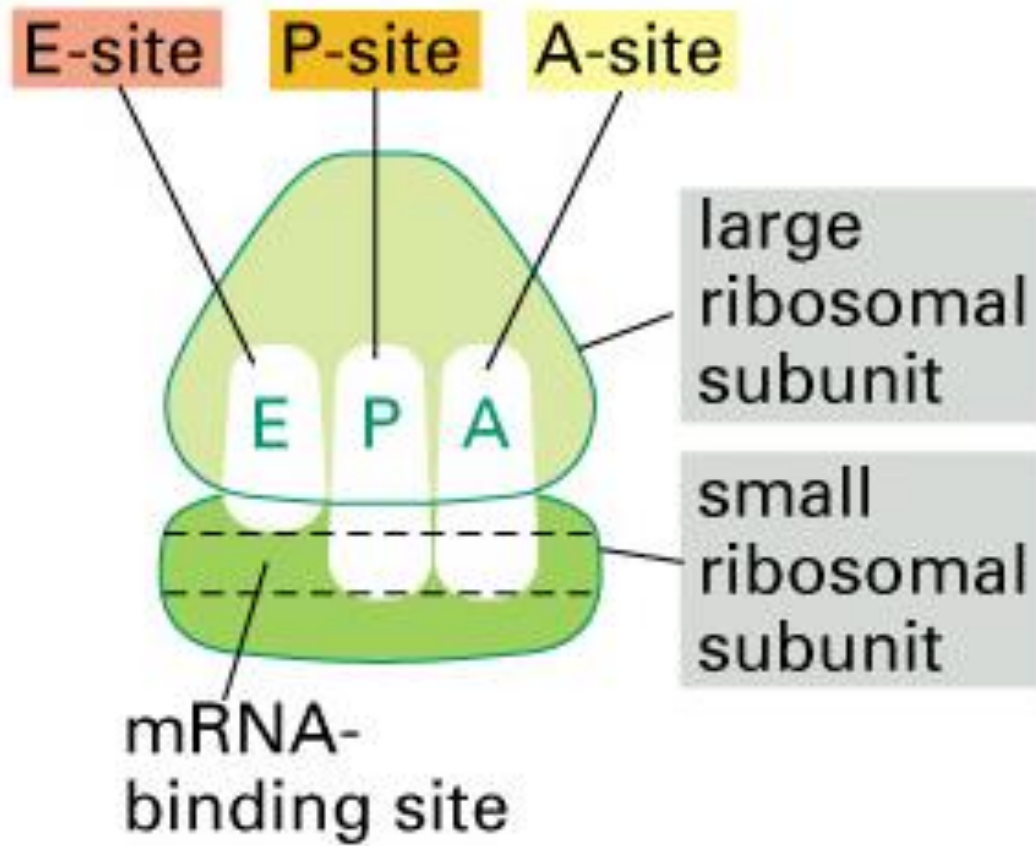
- **Ribosomes** are organelles that build proteins.
 - rRNA is found in ribosomes.
 - **mRNA is read on ribosomes.**
 - Ribosomes are found
 - Free-floating in the cytoplasm
 - Bound to the endoplasmic reticulum

Ribosomes

Ribosomes consist of **two major components**:

- small ribosomal subunit **reads the mRNA**
- large subunit **joins amino acids to form a polypeptide chain**



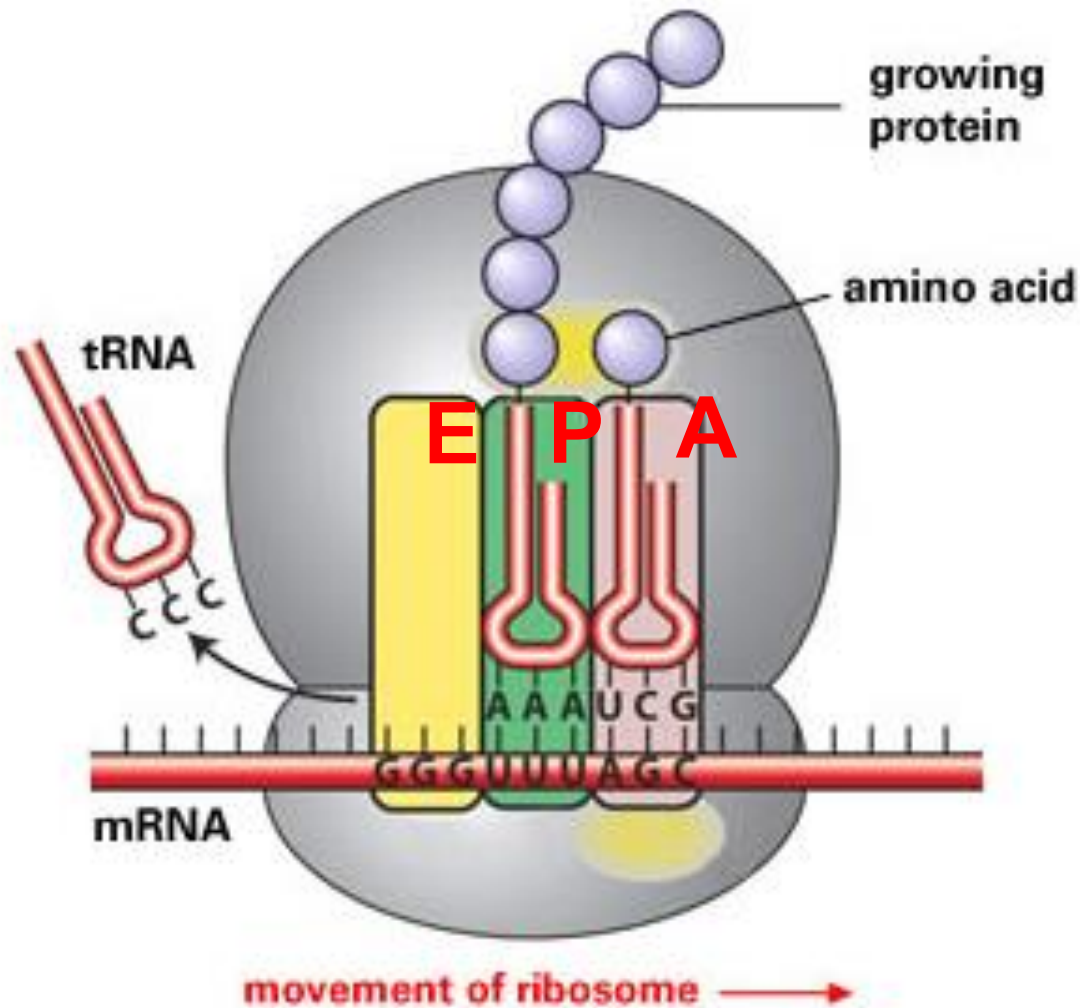


The A site is the arrival site for the aminoacyl tRNA, except for the first aminoacyl tRNA

The P site is where the peptidyl tRNA is formed in the ribosome

The E site is the exit site of the uncharged tRNA after it gives its amino acid to the growing peptide chain.

In addition to the E, P, A sites there is an mRNA binding groove that holds onto the message being translated



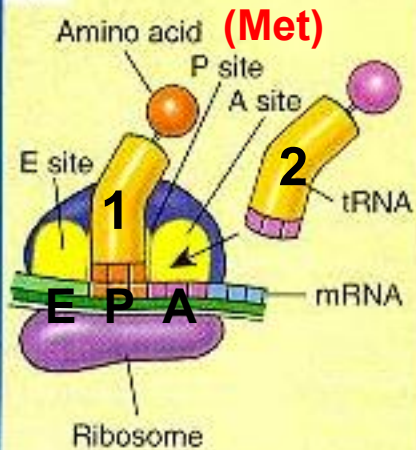
Each site of E, P and A occupies one codon each at a time, except at the time of initiation.

Translation Initiation

- Translation begins when
 - The **small ribosomal subunit** binds to the beginning of the mRNA and searches for the **AUG** start codon.
 - At this point, a tRNA brings the first amino acid.
 - The **anticodon in the tRNA matches with a codon on the mRNA.**
 - Each tRNA carries a specific amino acid based on its anticodon.
 - The start codon, **AUG**, binds to a tRNA that carries a **methionine.**
 - **Finally, the large ribosomal subunit joins the complex and the next step, translation elongation, can proceed.**

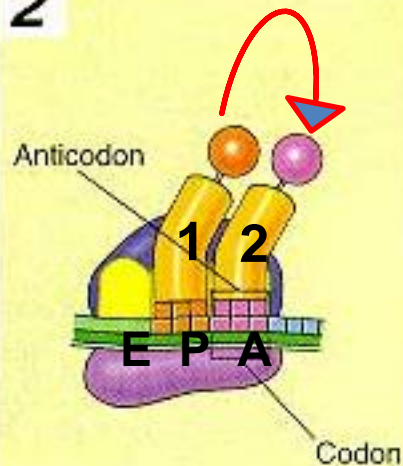
Translation

1



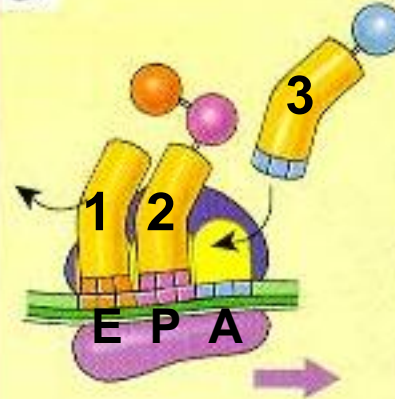
The initial tRNA occupies the P site on the ribosome. Subsequent tRNAs with bound amino acids first enter the ribosome at the A site.

2



The tRNA that binds to the A site has an anticodon complementary to the codon on the mRNA.

3



The ribosome moves three nucleotides to the right as the initial amino acid is transferred to the second amino acid at the P site.

4



The empty initiating tRNA leaves the ribosome at the E site, and the next tRNA enters at the A site.

At the time of **initiation**, **P site** occupies the **start codon**, where the **first tRNA** brings the **first amino acid (Met)**

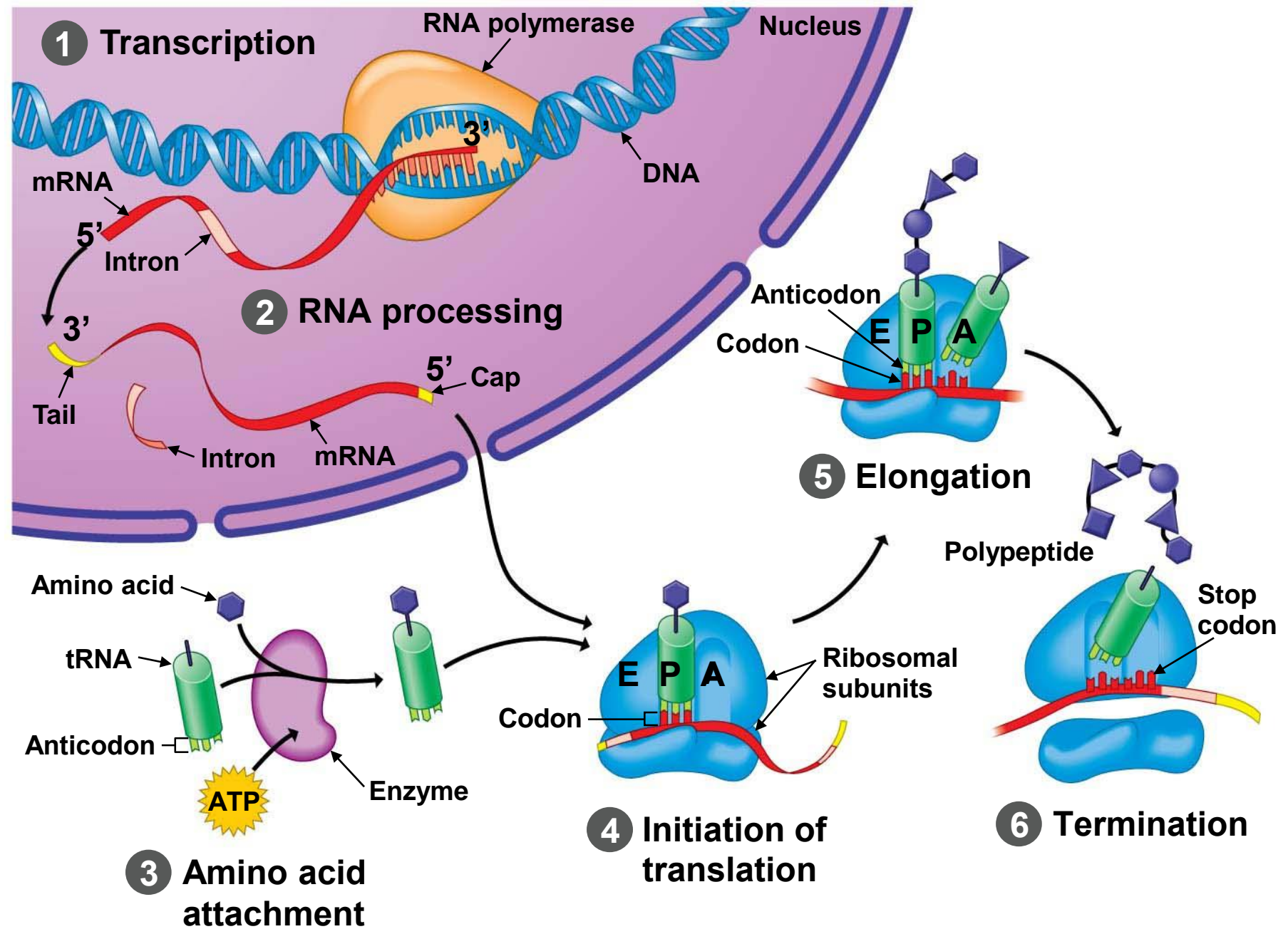
Translation Elongation

- The next tRNA binds with the next codon on the mRNA.
- The ribosome adds this amino acid to the growing polypeptide.
- **The ribosome then moves down to the next codon.**
- The process repeats itself. For each step, a new amino acid is added to the growing protein.

Translation Termination

- Elongation continues until the ribosome encounters a stop codon.
 - **UAA, UAG, UGA** are stop codons.
- **A release factor binds to the stop codon.**
 - This causes the ribosome to release the polypeptide.
 - The ribosomal subunits separate and release the mRNA.
 - **The mRNA can be translated again by another ribosome.**

Figure 10.19-s6



Control of Protein Synthesis

- **Cell types are different** from one another because they **express different sets of genes.**
 - Therefore, have different sets of proteins
- **Cells control gene expression (the decision to make a protein) in response to different environmental conditions.**
- Cells can **alter gene expression, which** controls quantity of a protein

Control of Protein Quantity

- Cells can regulate how much of a given protein has to be made by
 - Controlling *how much mRNA is available* for translation
- Cells do this in a **number of ways:**
 - 1) By regulating how tightly the chromatin is coiled in a certain region*
 - The more tightly the chromatin is coiled, the less likely a gene in that region will be transcribed.

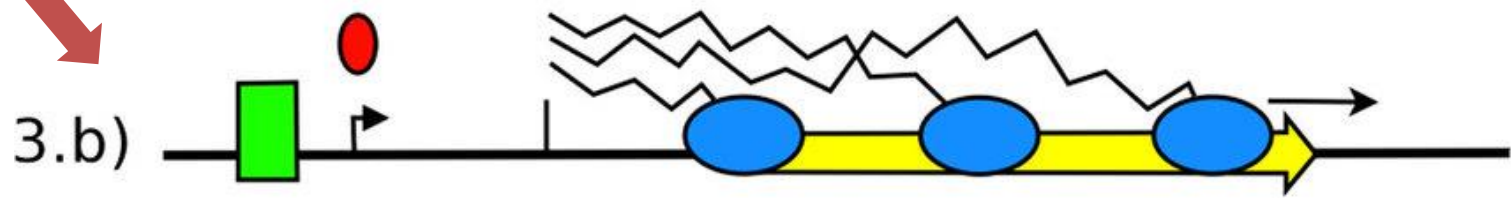
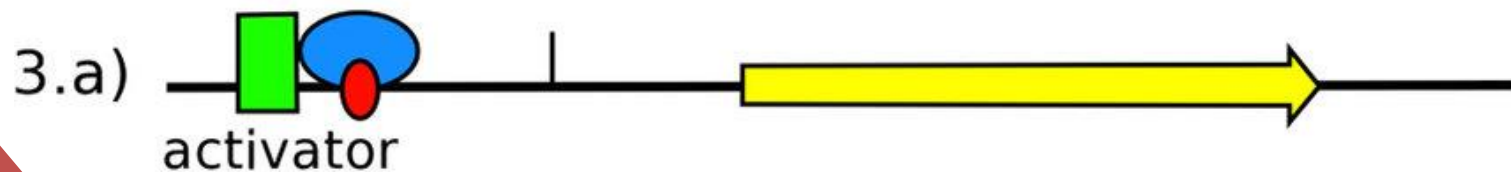
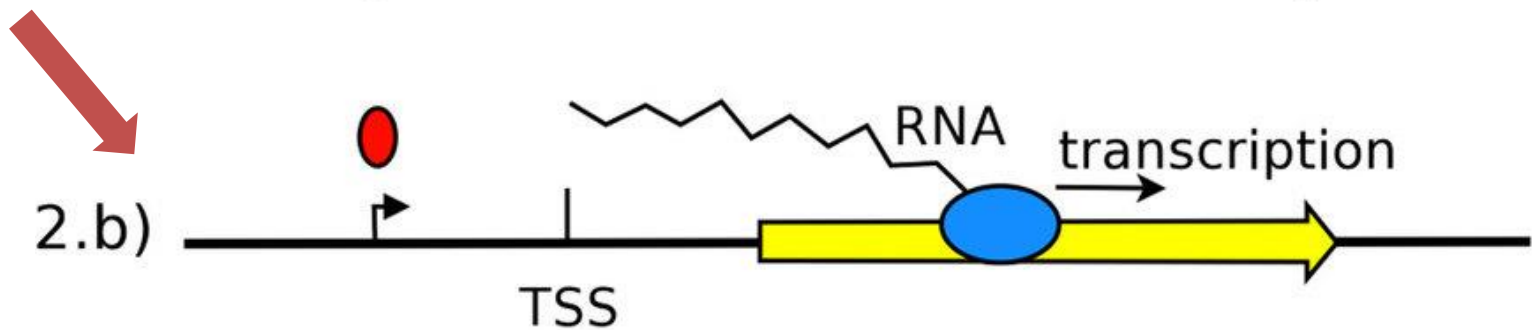
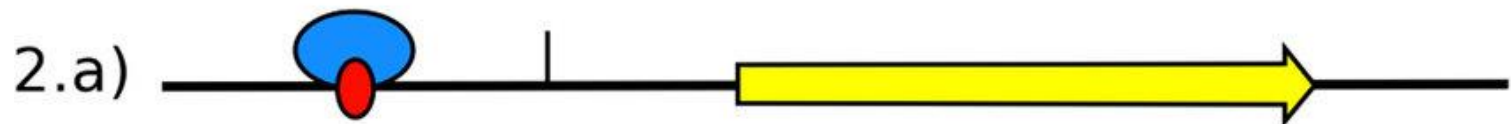
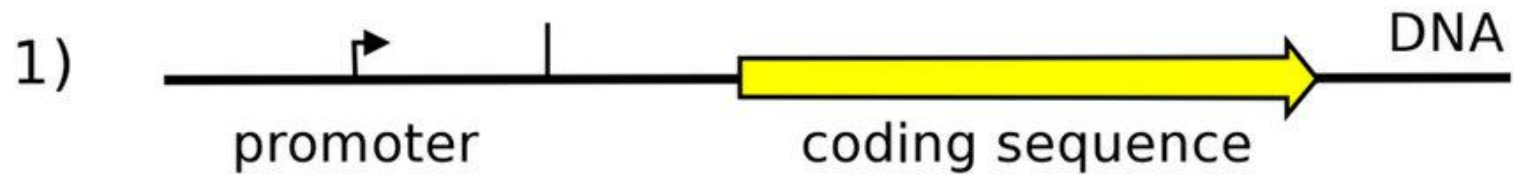
Control of Protein Quantity (Contd...)

2) By increasing or decreasing the **rate of transcription** of the gene:

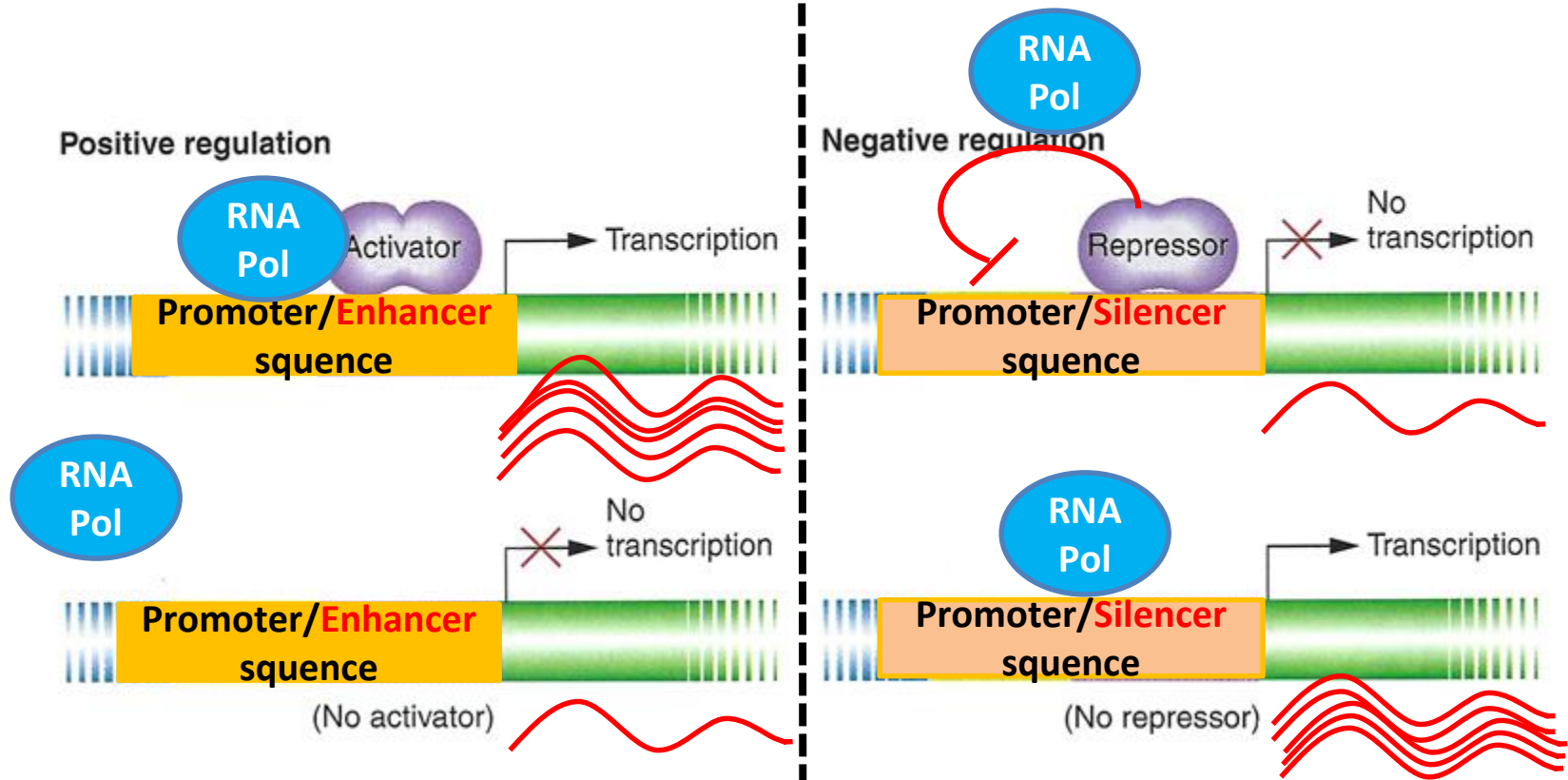
- **Activators** bound to **enhancer** regions increases transcription.
- **Repressors** bound to **silencer** regions decreases transcription.

3) By limiting the **amount of time an mRNA exists** in the cytoplasm:

- Some mRNA molecules are more stable and will exist longer in the cytoplasm, yielding more protein.



Activators and repressors of transcription



Without the activator, RNA pol cannot bind to the promoter for some of the genes, genes are turned off or transcribe at a low level

Without the repressor, RNA pol binds to the promoter, genes are turned on. If the Repressor is present, either some of the genes are turned off or transcribe at a low level

Mutations and Protein Synthesis

- A mutation is any change in the DNA sequence of an organism.
- Can be caused by **mistakes in DNA replication**
- Can be caused by **external factors**
 - **Carcinogens, radiation, drugs, viral infections**
- Only mutations in coding regions of gene will change the proteins themselves.

Point Mutations—a Change in a Single Nucleotide of the DNA Sequence

Three types:

A **nonsense mutation** changes a codon to a stop codon.

- This causes the ribosome to stop translation prematurely.
- CAA (Gln) to UAA (stop)

A **missense mutation** causes a change in the type of amino acid added to a polypeptide.

- This may change the way in which a protein functions.
- UUU (Phe) to GUU (Val)

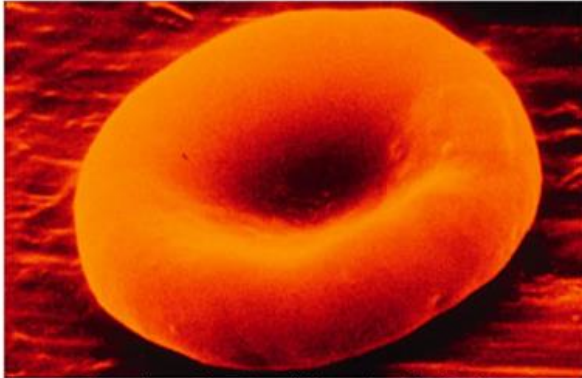
A **silent mutation** does not cause a change in the amino acid sequence.

- UUU to UUC; both code for Phe

Sickle Cell Anemia

- Results from a missense mutation in the gene for hemoglobin
 - **GAA to GUA**
 - **Glutamic acid to valine** change
 - Causes the hemoglobin protein to **change shape**
 - The red blood cells **stick together in low oxygen** conditions
 - Get stuck in blood vessels, causing the **vessels to break-apart** easily, leading to anemia
 - Also causes blood **vessels to clog, preventing oxygen delivery to tissues**, which results in tissue damage
 - Causes **weakness, brain damage, painful joints**, etc.

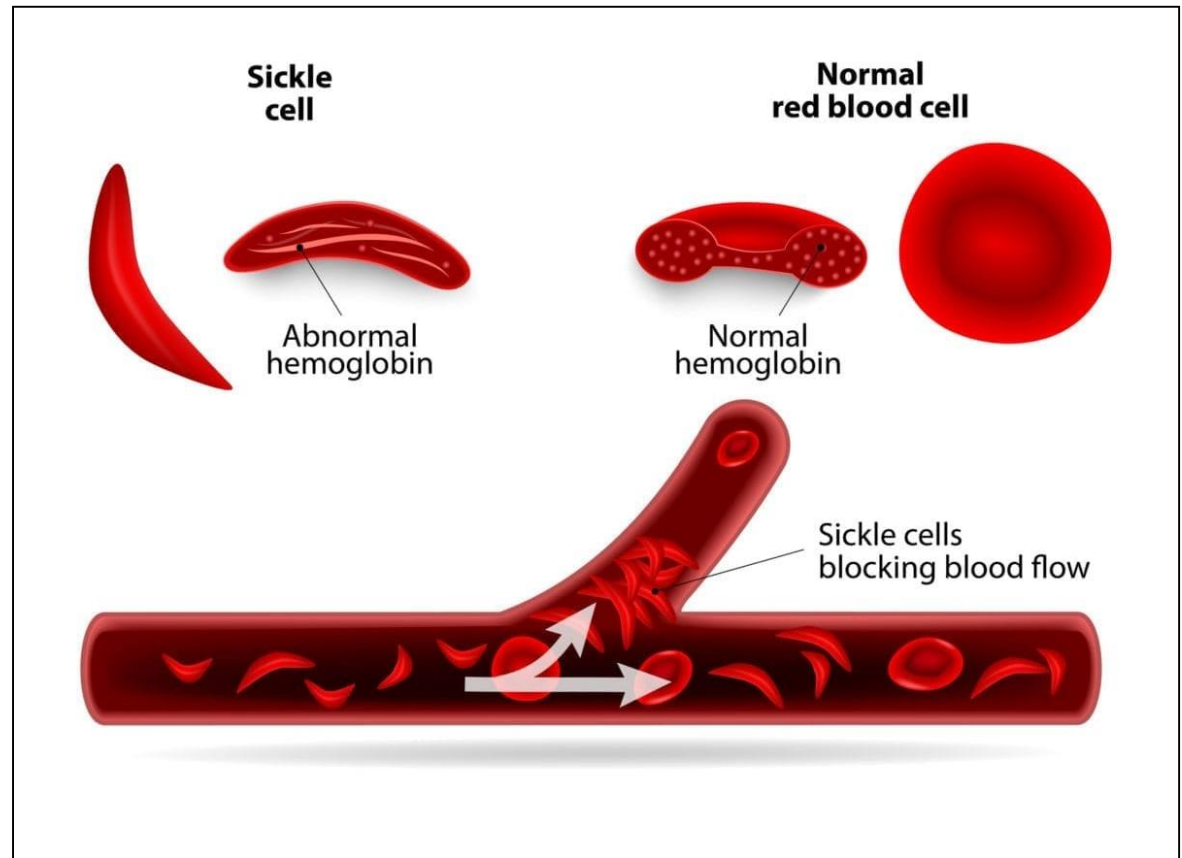
Normal and Sickled Red Blood Cells



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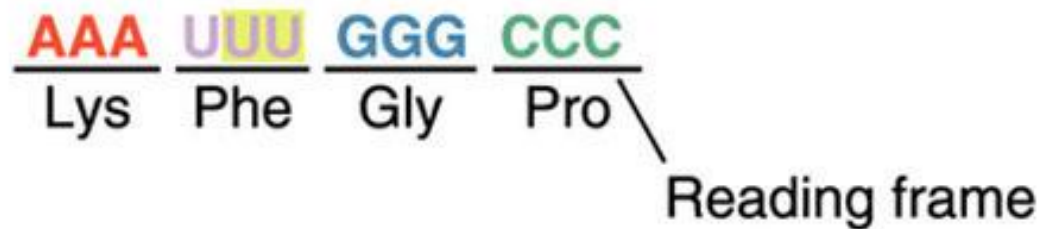
Insertions and Deletions

- An **insertion mutation** occurs when one or more nucleotides is added to the normal DNA sequence.
- A **deletion mutation** occurs when one or more nucleotides is removed from the normal DNA sequence.
- Insertions and deletions cause a **frameshift**.
 - **Ribosomes will read the wrong set of three nucleotides.**
 - **Changes the amino acid sequence dramatically**
 - **Changes the function of the protein dramatically**

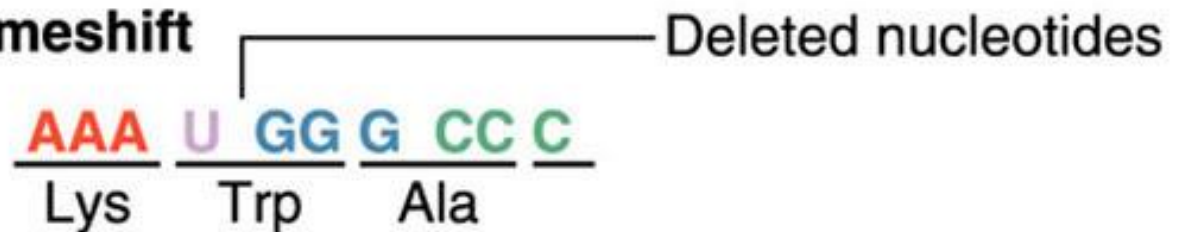
Frameshift

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Original mRNA sequence



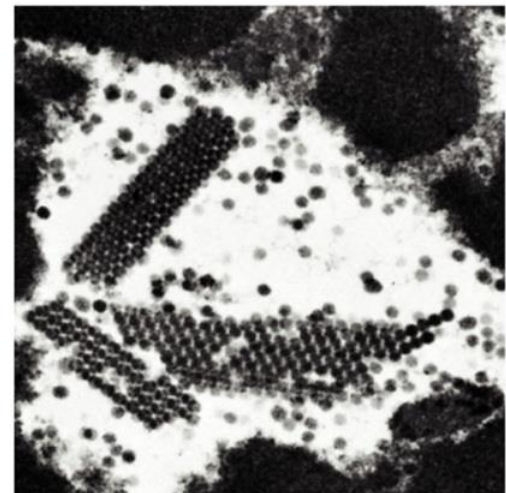
Effect of frameshift



Mutations Caused by Viruses

- **Viruses can insert their genetic material into the DNA of the host cell.**
- **The presence of the viral material may interfere with the host cell's ability to use the genetic material in that area because of this insertion.**
- **Insertion of human papillomavirus (HPV) causes an increased risk of cancer.**

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Chromosomal Aberrations

- Involves a major change in DNA at the level of the chromosome

- *Deletion*

- *Duplication*

- *Inversion*

- *Translocation*

All of these affect many genes, thus many proteins.

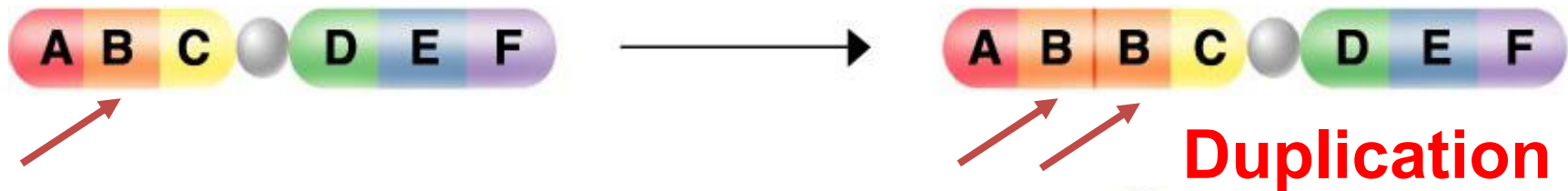
- In humans, these mutations may cause problems with fetal development.

Chromosomal Aberrations

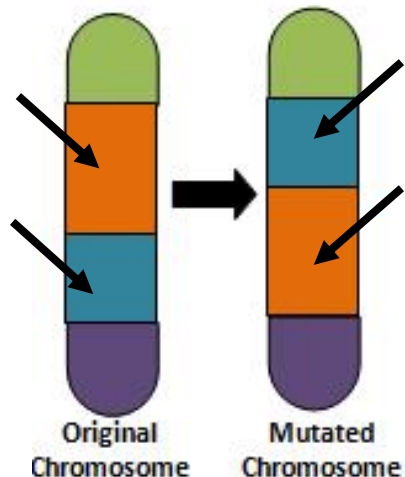
Deletion: when a broken piece of chromosome is lost or destroyed



Duplication: a chromosomal segment is replicated and reattached in sequence

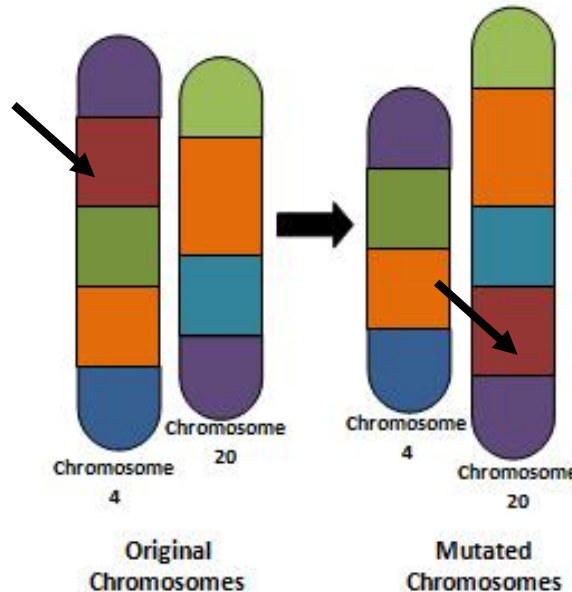


Inversions: a broken piece is reattached in the wrong orientation



Inversion

Translocation: a broken segment is integrated into a different chromosome



Translocation

5' AAGGCAAUCCCAUAAGG 3'

Given above an mRNA sequence, write the sequence of the Template strand of DNA from 5' to 3'

5' CCTTATGGGATTGCCTT 3'

5' AAGGCAAUCCCAUAAGG 3'

3' TTCCG TTAGGGTATT CC 5'

Write the coding strand sequence from the 5' to 3'

5' AAGGCAATCCCATAGG 3'

5' AAGGCAAUCCCAUAAGG 3'

Write the codons, corresponding anti codons and the amino acids of this mRNA starting from the first base from the 5' to 3'direction

Codons:	Anti-codons:	Amino acids
5' AAG 3'	5' CUU 3'	Lys
5'GCA 3'	5' UGC 3'	Ala
5' AUC 3'	5' GAU 3'	Ile
5' CCA 3'	5' UGG 3'	Pro
5' UAA 3'		

	U	C	A	G	
U	Phe	Ser	Tyr	Cys	U
	Phe	Ser	Tyr	Cys	C
	Leu	Ser	STOP	STOP	A
	Leu	Ser	STOP	Trp	G
C	Leu	Pro	His	Arg	U
	Leu	Pro	His	Arg	C
	Leu	Pro	Gln	Arg	A
	Leu	Pro	Gln	Arg	G
A	Ile	Thr	Asn	Ser	U
	Ile	Thr	Asn	Ser	C
	Ile	Thr	Lys	Arg	A
	Met	Thr	Lys	Arg	G
G	Val	Ala	Asp	Gly	U
	Val	Ala	Asp	Gly	C
	Val	Ala	Glu	Gly	A
	Val	Ala	Glu	Gly	G

5' AAGGCAAUCCCAUAAGG 3'

In the pre-mRNA, if the nucleotides drawn in the box belong to an intron, write the template strand and the matured mRNA sequence

Template strand:

5' CCTTATGGGATTGCCTT 3'

Matured mRNA:

5' AACAUAAAGG 3'

Write the codons, anti codons and the amino acids coded by the mRNA:

Codons:

5' AAC 3'

5' AUA 3'

5' AGG 3'

Anti Codons:

5' GUU 3'

5' UAU 3'

5' CCU 3'

Amino acids:

Asn

Ile

Arg

	U	C	A	G	
U	Phe	Ser	Tyr	Cys	U
	Phe	Ser	Tyr	Cys	C
	Leu	Ser	STOP	STOP	A
	Leu	Ser	STOP	Trp	G
C	Leu	Pro	His	Arg	U
	Leu	Pro	His	Arg	C
	Leu	Pro	Gln	Arg	A
	Leu	Pro	Gln	Arg	G
A	Ile	Thr	Asn	Ser	U
	Ile	Thr	Asn	Ser	C
	Ile	Thr	Lys	Arg	A
	Met	Thr	Lys	Arg	G
G	Val	Ala	Asp	Gly	U
	Val	Ala	Asp	Gly	C
	Val	Ala	Glu	Gly	A
	Val	Ala	Glu	Gly	G

Write down the polypeptide sequence of each

5' AAGGC**A**AUCCCAUAAGG 3'

5' AAGGC**U**AUCCCAUAAGG 3'

Silent Mutation, no change in polypeptide sequence

5' AAGGC**CA**AUCCCAUAAGG 3'

5' AAGG UCCCAUAAGG 3'

Lys-Val-Pro

5' AAGGC**A**AUCCCAUAAGG 3'

5' AAGGC**CU**AUCCCAUAAGG 3'

Lys-Ala

Original Amino acids:
Lys-Ala-Ile-Pro

	U	C	A	G	
U	Phe	Ser	Tyr	Cys	U
	Phe	Ser	Tyr	Cys	C
	Leu	Ser	STOP	STOP	A
	Leu	Ser	STOP	Trp	G
C	Leu	Pro	His	Arg	U
	Leu	Pro	His	Arg	C
	Leu	Pro	Gln	Arg	A
	Leu	Pro	Gln	Arg	G
A	Ile	Thr	Asn	Ser	U
	Ile	Thr	Asn	Ser	C
	Ile	Thr	Lys	Arg	A
	Met	Thr	Lys	Arg	G
G	Val	Ala	Asp	Gly	U
	Val	Ala	Asp	Gly	C
	Val	Ala	Glu	Gly	A
	Val	Ala	Glu	Gly	G

EXTRA SLIDES

Shine-Dalgarno sequence in prokaryotic translation

-Ribosomal binding site in prokaryotic mRNA

-16S rRNA present in the small subunit of ribosome binds to the mRNA, approximately 8 base pairs upstream to the start codon

-This sequence helps recruiting the ribosome to the mRNA to initiate protein synthesis

