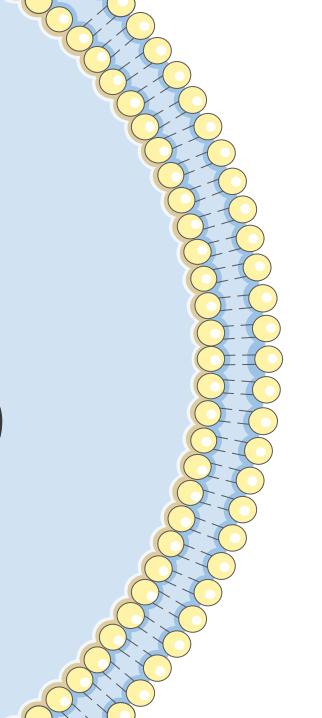


Molecular Biology (2)





Color Index:

- Main text
- **Important**
- Notes
- Boys slides' Girls slides'
- Extra

Editing File

Objectives

- ✓ To understand DNA replication
- ✓ To know the transcription of genetic material into messenger RNA
- ✓ To get an idea about the translation of mRNA into a functional protein.



DNA is the genetic material

Therefore it must:

- Replicate faithfully (precisely).
- Have the coding ability to produce proteins for all cellular functions.

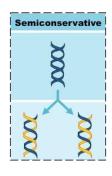


Features of Eukaryotic DNA Replication



Semiconservative with respect to parental strand:

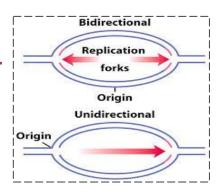
Daughter DNA molecules contain one parental strand and one newly-replicated strand.



02

Bidirectional with multiple origins of replication.

Bidirectional: it goes both ways. Origins of replication: specific sequence where the parent DNA strands separate



03

Primed by short stretches of RNA.

It's a primer (initial fragment) that help DNA polymerase to know where to add the DNA nucleotide. The primer≈10 nucleotides.

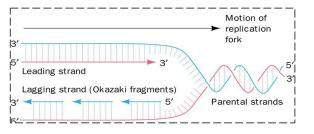
** 'semi' means half, so *semiconservative* : a half is from parent DNA *semi-discontinous* : a half has intervals 04

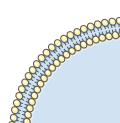
Semi-discontinous.

In DNA replication, both daughter strands (leading strand red, lagging strand blue) are synthesized in their $5' \rightarrow 3'$ directions.

Leading strand: a new strand that synthesized continuously

Lagging strand: a new strand that synthesized in fragment (Okazaki-fragments)







Proteins involved in DNA Replication

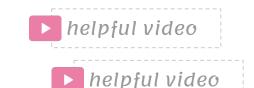




| Protein name | Function Unwinds the DNA at the site of origin by breaking the hydrogen bond between the bases | | |
|--|---|--|--|
| DNA Helicase. | | | |
| Single-stranded DNA binding proteins. | Prevent rebinding of the DNA helix (prevent hydrogen bond formation) | | |
| (DNA) Primase. | Make short segments of RNA prime complementary to the DNA. | | |
| DNA polymerases (5 types: α ; β ; γ ; δ ; ϵ) (each with different job but we are not going that deep). | Add nucleotides to RNA primer. Proofreads bases added and replaces incorrect nucleotides. (Remover RNA primers / fill the gaps / spell check) | | |
| DNA ligase. | Join the segments that DNA polymerase made instead of RNA nucleotides, to form bonds between sugar-phosphate beckbone. | | |
| Topoisomerases: Topoisomerase I + Topoisomerase II. | prevent supercoiling of the chromosome. | | |
| Telomerases | prevent the shorting of the chromosome (Maintain the chromosome length). | | |



Steps in DNA Replication



01

Helicase protein. (pink)

binds to DNA sequences called <u>origins</u> and **unwinds DNA strands**.

-open the DNA strands and form **replication fork**

02

Single-stranded DNA binding proteins. (purple)

prevent single strands from rewinding.

03

DNA Primase protein. (red)

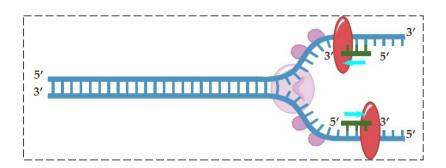
makes a short segment of RNA primer complementary to the DNA

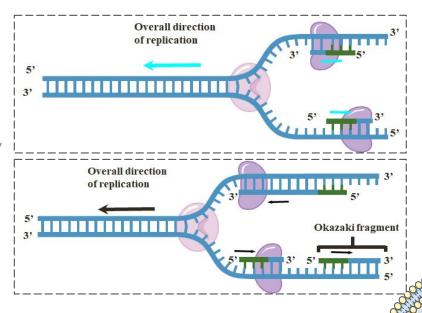
- -always the synthesis direction $5' \rightarrow 3'$
- -made by DNA polymerase.

04

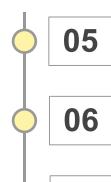
DNA polymerases

- Adds DNA nucleotides to the RNA primer.
 - -start adding complementary nucleotide. **leading strand** is $(5' \rightarrow 3')$ toward the fork. but the **lagging strand** is $(5' \rightarrow 3')$ in the opposite direction of the fork.
- Proofreads bases added and replaces incorrect nucleotides.
 - if a wrong nucleotide is added that is <u>not</u> complementary to the parent strand, DNA polymerase will **recognize** it and **replace** it
- Leading strand synthesis **continues** in a 5' to 3' direction.
- Discontinuous synthesis produces 5' to 3' DNA segments (Okazaki fragments).









DNA polymerases. (yellow)

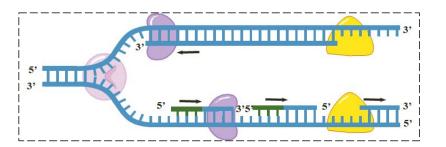
Exonuclease activity of DNA polymerase **removes RNA primers Exonucleases** are enzymes that remove nucleotides.

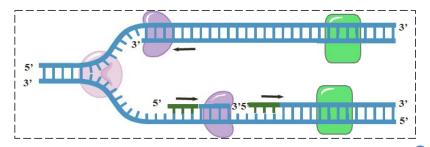
DNA polymerases

DNA polymerase fills the gaps.

07 Ligase. (green)

Ligase forms bonds between sugar-phosphate backbone (phosphodiester bond)
The DNA is consist of *sugar-phosphate* chains on the <u>sides</u> and *nitrogenous bases* in the <u>center</u>





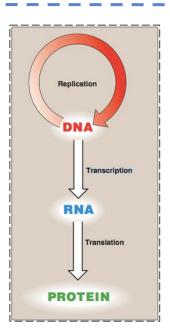


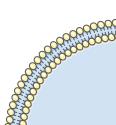
The central dogma of Molecular Biology

A portion of DNA, called a **gene**, is transcribed into RNA.

RNA is translated into proteins.

only 5% of DNA wil transcribed into mRNA





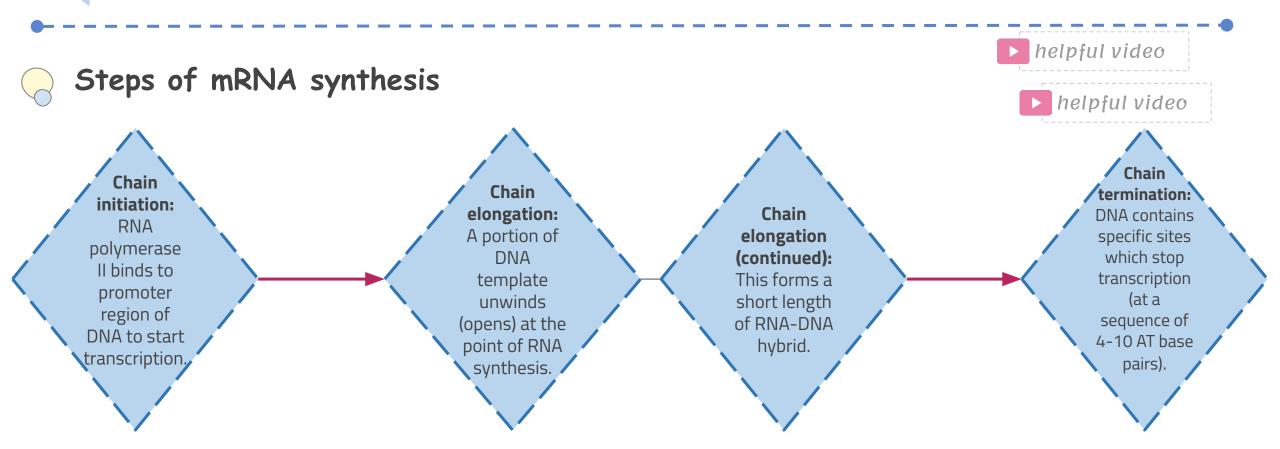


Transcription (mRNA synthesis)

- A portion of DNA (a gene) is transcribed into messenger RNA (mRNA).
- The direction of transcription is 5'--->3'.

- Only one of the DNA strands is transcribed (antisense strand).
 - Antisense strand: It is the strand that contains the opposite codon of the required mRNA to be transcribed.

The RNA polymerase II is responsible for this process.

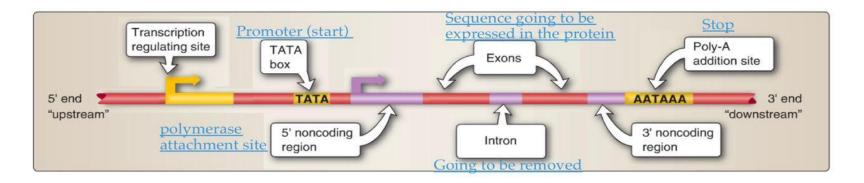




Steps of mRNA synthesis:

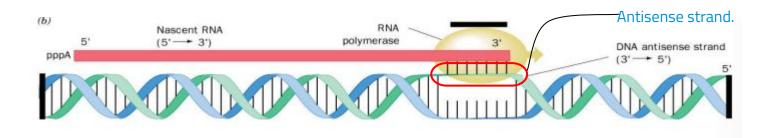
shows:

- Initiation (first step), where RNA polymerase II binds to promoter region of DNA to start the transcription process. (always at 5' side)
- Termination (last step), where RNA polymerase II stops transcription. (always at the 3' side)



shows:

 Elongation (second step), where a portion of DNA unwinds (opens) at the point of RNA synthesis, which forms a short length of RNA-DNA hybrid.





Post-transcriptional modification



*The pre-mRNA has to go through some modifications to become a mature mRNA, the modifications occur in the cell nucleus before the RNA is translated.

Post-transcriptional modification (3 steps)

2- Polyadenylation:

Addition of a poly(A) tail

(a highly conserved

end of mRNA.

AAUAA sequence) at

1- Capping:

addition of a methylated guanine nucleotide at 5' end of mRNA

Function:

To **prevent** mRNA degradation by exonucleases.

It helps the transcript during protein

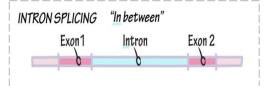
bind to the ribosome synthesis.

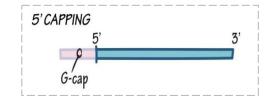
To **protect** the mRNA from **degradation**

Function:

For ribosomal RNA recognition

3- Intron removal for releasing mature mRNA from nucleus.





Exonucleases are enzymes that remove nucleotides.





Translation (Protein synthesis)

- Translation is a process of protein synthesis from mRNA.
- mRNA has genetic codes for amino acids present in proteins.
- The genetic code is a dictionary that identifies the correspondence between a sequence of nucleotide bases and a sequence of amino acids. *it is a code that connect the nucleotide bases in mRNA and the amino acids.
- Each individual word in the code is composed of three nucleotide bases (codons).
 *each codon specifies a particular amino acid
- **⋖** 64 possible codons:
 - 1. 61 codons specify 20 amino acids. there are different codes give the same amino acid.
 - 2. One start codon (AUG). AUG code is for Methionine, every translation start with AUG.
 - 3. Although methionine (Met) is the first amino acid incorporated into any new protein, it is not always the first amino acid in mature proteins, it may be removed after translation.
 - 4. 3 stop codons: UAA, UAG and UGA

| First position (5' end) | Second position | | | | Third position (3' end) |
|-----------------------------|---|---------------------------------|--|------------------------------|-------------------------|
| | U | C | A | G | |
| UUU Phe UUC Phe UUA UUG Leu | Phe | UCU UCC | UAU Tyr UAC | UGU Cys | U C |
| | UCA Ser UCG | | UGA Stop UGG Trp | A G | |
| c | CUU CUC CUA CUG | CCU CCC CCA Pro | CAU CAC His CAC CAA Gln | CGU CGC CGA Arg CGG | U C A G |
| A | AUU AUC Ile AUA AUG Met ^b | ACU ACC ACA Thr ACG | AAU AAC AAA AAG Lys | AGU Ser AGC AGA Arg | U C A G |
| G | GUU GUC GUA Val GUG | GCU GCC GCA Ala GCG | GAU GAC Asp GAA GAA GAG | GGU GGC GGA GGG | U C A G |

Just memorize the **start** + **stop** codons



Components required for Translation

01

Amino acids.

02

Transfer RNA (tRNA).

read the code and bring the amino acids

Aminoacyl-tRNA synthetases.

Aminoacyl-tRNA: (tRNA + amino acid)

synthetases: enzyme that make Aminoacyl-tRNA



04

mRNA



05

Functionally competent ribosomes.

Functional Ribosomes consists of 2 subunits:

- Small ribosomal subunit has mRNA binding site.
- Large ribosomal subunit



06

Protein factors

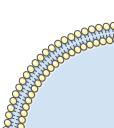
Catalytic or stabilizer for the synthetic machinery.



7

ATP and GTP

Energy is needed.





Steps in Protein Translation

helpful video: start at 3:40

*Important to know that:

- ◀ tRNA has two important areas:
 - 1- the **anticodon** which forms base pairs with its <u>complementary</u> <u>sequence</u> on mRNA.
 - 2- a region for attaching a specific amino acid.
- Ribosome has 2 subunits that join to form a functional ribosome:
 - 1- small subunit: where the mRNA come and site
 - 2- **large subunit:** has three sites for tRNA:

A site: Acceptor site, that binds to tRNA which holds the new amino acid to be added to the polypeptide chain.

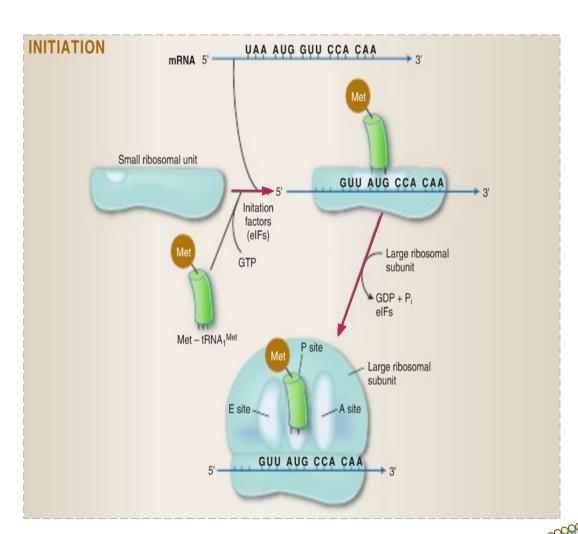
P site: Peptidyl site, hold tRNA carrying the growing polypeptide chain.

E site: Exit site, that discharge tRNA to leave the ribosome

1- Initiation:

It requires <u>ribosomal subunits</u>, <u>mRNA</u>, <u>aminoacyl-tRNA</u> for methionine, initiation <u>factors</u> and <u>energy</u>. These all join to form the **initiation complex**.

- **1-** small ribosomal subunit bind to the aminoacyl-tRNA and mRNA which carries the start codon.
- **2-** the large ribosomal subunit binds to the small ribosomal subunit to complete the initiation complex.





Steps in Protein Translation

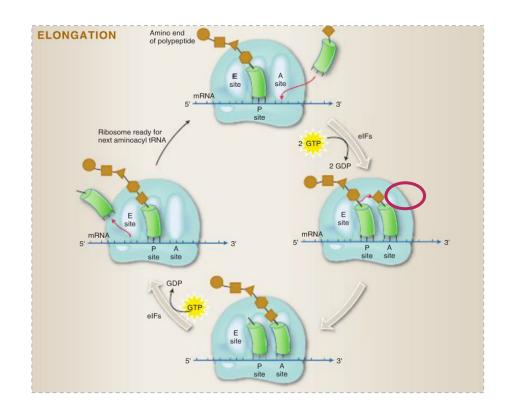
2- Elongation:

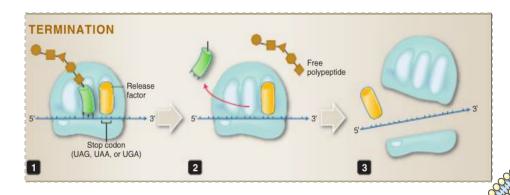
- **1-** An aminoacyl-tRNA will attach to **A site**. the attachment will be facilitated by **elongation factors**.
- **2- peptidyl transferase**: it is an enzyme that **separates** the growing chain bond with tRNA in the **P site**, and **transfers** the growing polypeptide chain to tRNA at **A site**
- **3-**then the ribosome will move 3 nucleotides. So the **empty tRNA** at P site moves to **E site** and **discharge**, and the **tRNA** with the **growing chain** moves to the **P site**.
- **4-** the **A site** will be ready to **receive another aminoacyl-tRNA** and repeat the steps.

3- Termination:

Occurs when one of the three **stop codons (UAA,UAG,UGA)** reaches the **A site**.

- 1- **Release factor** binds to the **stop codon** and cut the bond between the polypeptide and its tRNA in the **P site**.
- 2- This frees the polypeptide and the translation complex **disassembles.**





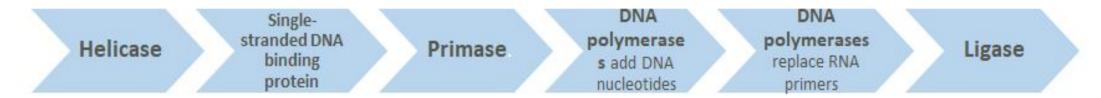




Features of Eukaryotic DNA Replication

1- Semiconservative 2-Bidirectional with multiple origins of replication 3-Primed RNA. 4-Semi-discontinous

DNA Replication



transcription: mRNA synthesis



Protein Translation





- ✓ DNA is the genetic material, so it must replicate faithfully and have the coding ability to produce proteins for all cellular functions.
- ✓ Only one strand of DNA (antisense strand) is transcribed into mRNA.
- ✓ The synthesized mRNA is protected from the destruction and prepared for translation through post-transcriptional modification.
- ✓ mRNA transcription and protein synthesis processes are the same in both prokaryotic eukaryotic cells with some differences.



Q1: DNA polymerases is responsible for? Leading strand Adds DNA A Proofreads bases All nucleotides synthesis Q2:In post-transcriptional modification, where does capping have place? at the promoter B at the 3' end C at introns A at the 5' end region Q3: How many proteins are involved in DNA replication? D 7 A 4 11 Q4: What is DNA polymerases function Prevent single Forms Unwinds DNA phosphodiester Fills the gaps strands from strands bonds rewinding Q5:What is the function of A site in ribosomes? holds tRNA empty tRNA leaves binds incoming dissociation of the ribosome carrying the aminoacyl-tRNA. ribosome. through it. polypeptide chain.

Answer Key:

8(9

3)**D**

A(S

a(r

Q6:Mention all stop codons?

Q7:What is the codon of methionine?

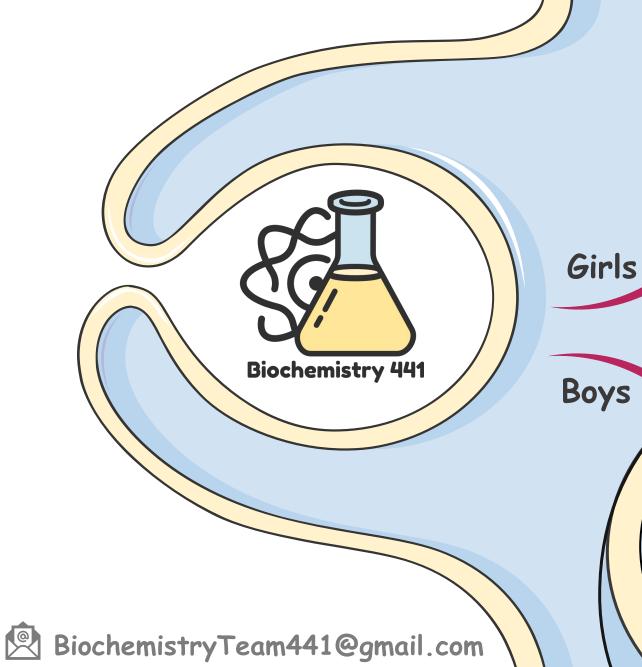
Q8:What is the function of DNA primase?

Q9:What is the function of Polyadenylation

Answer:

6-Three stop codons: UAA, UAG and UGA 7-AUG.

8-makes a short segment of RNA primer complementary to the DNA.
9-To protect the mRNA from degradation + for ribosomal RNA recognition



Leader: Ghadah Alarify

Members:

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Members:

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