



Molecular Biology 2

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Notes and explanations

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highlights



Objectives:

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• 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

1- it replicates faithfully. (without mutations or mistakes)

2-has the coding ability (ability to transfer information) to produce proteins for all cellular functions.

Because it carries all the important genetic information; we should be able to inherent it without mistakes, and we should have a correction system if there were mistakes.



Features of Eukaryotic DNA:

Semiconservative	Daughter DNA molecule contains one parental strand and one newly-replicated strand		
Bidirectional goes in both directions with multiple origins of replication	Origin of replication: is a particular sequence • in a genome at which replication is initiated. The opening point of DNA	Fork movement 35 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	Fork movement
Primed by short stretches of RNA.	a short segment of RNA, 10 nucleotides long) is required to • start a new chain; removed later on during replication		
Semi- discontinuous	The leading strand is synthesized continuously and the lagging strand is discontinuous (in fragments) The directions will always be 5' -> 3' (either away or into the of fork)	DNA: Reading : 3' > 5' Synthesis :5' > 3'	

Proteins	
involved in	
DNA	
replication	
	,

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Name	Types	Functions
DNA Helicase		-An enzyme that binds to DNA sequences called Origins and unwinds DNA strands.
Single-stranded DNA binding proteins		Prevents single strands from winding.
DNA Primase		An enzyme that makes a short segment of RNA primer which is complementary to the DNA
DNA polymerases	α (Alpha), β (Beta), γ (Gamma), δ (Delta), ϵ (Epsilon).	An enzyme that adds DNA nucleotides to the RNA primerProofreads bases added and replaces incorrect nucleotides.
DNA ligase		An enzyme that forms bonds between the sugar-phosphate backbone.
Topoisomerases	Topoisomerases I	(cut one of the DNA strands)
	Topoisomerases II	(cut both of the DNA strands)
Telomerases		An enzyme that adds nucleotides to telomeres (a reign at the end of a chromosome)

Steps in DNA replication

Helicase	Binds at the origin point , Untwists DNA strands, making a fork-like structure \rightarrow called replication fork
Single-stranded binding proteins.	Prevents single strands from twisting again. (keeps DNA strands opened)
Primase	Makes RNA primer. (a short segment of RNA complementary to the DNA)
DNA polymerase.	 1-Adds nucleotides to the primer (from 5' to 3') and checks if the bases added are correct or not, if not → removes incorrect nucleotides.(proofreads and corrects) 2- Removes the RNA primers by exonuclease activity (which creates gaps) 3-Fills the gaps that we got after removing the RNA primers (adding form 3' to 5')
Ligase	Connects the okazaki fragments by forming bonds between the sugar-phosphate backbone.(from 5' to 3')

DNA replication



Leading strand : continuous synthesis from 5' to 3' direction. Lagging strand : discontinuous synthesis produce 5' to 3' DNA segments (okazaki fragments)

The central dogma of Molecular Biology



Transcription (mRNA synthesis)

• A portion of **DNA** (a **gene**) is transcribed into messenger RNA (**mRNA**).

ملاحظة: Pay attention to the

number of the protein (RNA polymerase II)

• Only one of the DNA strands is transcribed (antisense strand).

• The **RNA polymerase II** is responsible for this process.

• The direction of transcription is $5' \rightarrow 3'$.

the template strand on which the mRNA is synthesized

Dr.note: The mRNA chain is complementary to the mother DNA chain, whereas the amino acid chain has the same sequence of the mother DNA and a complementary sequence to the mRNA

Steps of mRNA synthesis

• 1-Chain initiation:

• RNA polymerase II binds to *promoter region* of DNA to start transcription.



Steps of mRNA synthesis

Chain elongation:

- A portion of DNA template unwinds (opens) at the point of RNA synthesis. (called a bubble)
- This forms a short length of RNA-DNA hybrid.

Chain termination:

• DNA contains specific sites which stop transcription (at a sequence of 4-10 AT base pairs).

After the termination we get an immature RNA, because this RNA has to travel to the cytoplasm, it needs some modifications to protect it and help the ribosomes recognize it. (post transcriptional modification)



Post-transcriptional modification

• **Capping:** Addition of a methylated guanine nucleotide at

5' end of mRNA

Function:

- To prevent mRNA degradation by exonucleases.
- It helps the transcript bind to the ribosome during protein synthesis.
- **Polyadenylation:** Addition of a poly(A) tail (a highly conserved

AAUAA sequence) at 3' end of mRNA.

Functions:

- To protect the mRNA from degradation
- For ribosomal RNA recognition



• Intron removal for releasing mature mRNA from nucleus.

Translation (Protein synthesis)

It 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.
- Each individual word in the code is composed of three nucleotide bases (codons).

الكودون هو ثلاثة " "نيوكليتايد تكون شفرة لحمض أميني معين

> 61 coding codons (including the start codon) 3 non coding stop codons



جملة لتسهيل حفظ كودونات التوقف:

" يُقال من يواعد يُواجه" UGAl mn UAAed UAGeh

You only need to memorize the stop codons and the start codon (written in the previous slide)

second base in codon



linst base in codon

Components required for Translation

- Amino acids
- Transfer RNA (tRNA)
- Aminoacyl tRNA synthetase
- mRNA
- Functionally competent ribosomes
- Protien factors
- ATP and GTP



1. Initiation: in general it means {making complements}

Puts (ribosoms +tRNA+ mRNA} together and make it ready to start

Produces an initiation complex made of : 1- small ribosomal subunit 2- large ribosomal subunit 3-mRNA 4- methionine tRNA



Elongation: it happens until the termination (stop codon)

* P =peptide side, A=incoming amino acids, E= Exit site

Important enzyme in elongation : peptidyl transferase



Termination : the release factor binds to the A site > everything disassembles

(يتفكك الريبوسـوم*ر* بمجرد ارتباط عامل فك الارتباط في الموقع أ)









Mentioned in males lecture:

- All prokaryotes have 70S ribosomes. 70S ribosome is made up of a large subunit (50S) and small subunit (30S)
- Eukaryotes have larger ribosomes (80S) made up of 60S and 40S subunits.
- S= Svedberg units to measure the size of the subunits
- In humans the start codon is **Methionine.**
- In bacteria the start codon might code with different codons.



Quiz

• Onlyof the DNA strands is transcribed (antisense strand)

One	Two	Three	Four	
Protein makes	a short segment of RNA			
Single stranded	Primase	DNA Helicase	Ligase	
• Addition of a me	ethylated guanine nucleotide at 5	5' end of mRNA		
Polyadenylationcc	intron	axone	capping	
• Polyadenylation	and capping function as			
helps the transcript	bind to the ribosome during prot	ein synthesis		
ribosomal RNA rec	ognition			1-A
a protect the mRNA	from degradation			2-B 3-D
None				4- C

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