

Subject ? <sup>?</sup> Termination in eukaryotes ? Processing of mRNA

Lecno: 11

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\* بالمحاضرة الماضية حكينا عن ال Initiation و Elongation لل Transcription in eukaryotes هلاً رح نكمل ال Termination

#### Termination in eukaryotes:

- Leads to the dissociation of the complete transcript and the release of RNA polymerase from the template DNA. The process differs for each of the three RNA polymerases. الثلاثة الثلاثة
- As Pol II reaches the end of a gene, two protein complexes carried by the CTD (carboxy terminal domain), CPSF (cleavage and polyadenylation specificity factor) and CSTF (cleavage stimulation factor), recognize the poly-A signal (polyadenylation signal sequence AAUAAA) in the transcribed RNA.
  - The sequences that, once transcribed into RNA, trigger transfer of these factors to the RNA are called poly-A signals

- Poly-A-bound CPSF and CSTF recruit other proteins to carry out <u>RNA cleavage</u> and then <u>polyadenylation</u>. Poly-A polymerase adds approximately 200 adenines to the cleaved 3' end of the RNA without a template. The long poly-A tail is unique to transcripts made by Pol II.
- The RNA molecule made by RNA pol II is called a primary transcript, which needs extensive RNA processing in order to produce a mature mRNA for translation & protein synthesis.

لسا مش جاهن لل translation

#### اول شبى ال polymerase عمل transcription ل عشان يعطينا mRNA



# Processing of mRNA Synthesis & Processing of ribosomal RNA (rRNA) Synthesis & Processing of tRNA

Вy Dr. Walaa El Gazzar

- A) 5' capping
- splicing of exons
- عشان نكون mature RNA لازم نعمله modification ب 3 خطوات :

### B) Addition of poly (A) tail C) Removal of introns and Processing of mRNA

(Post transcription modifications)

## A. 5<sup>\</sup>-Capping :

متى بنحطله ال Cap ؟

- The RNA is capped as soon as it emerges from the RNA-exit channel of polymerase. This happens when the transcription cycle has progressed only as far as the transition from the initiation to elongation phases.
- □ The cap is a **7- methylguanosine triphosphate** attached to the  $5^{-}$  terminal end of the mRNA (which terminates at a triphosphate group).
- One of the terminal phosphate groups is removed by **<u>RNA triphosphatase</u>**, leaving a bisphosphate group بروح بشيل وحدة phosphate الى هي الغاما

رح تطلع pyrophosphate و تصير GMP

 <u>GTP</u> is added to the terminal bisphosphate by mRNA <u>guanylyltransferase</u>, losing a pyrophosphate from the GTP substrate in the process. This results in the <u>unusual 5<sup>to 5</sup></u> <u>triphosphate linkage.</u>



#### 5' end of mRNA

Methylation of this terminal guanine is catalyzed by guanine-7-methyltransferase.

- S adenosylmethionine, SAM, (<u>active</u> methionine) is the source of methyl group.
  Methylation of N-7 of guanine of the GTP cap occurs in the nucleus.
- In the cytoplasm, methylation may occur at 2<sup>\</sup>OH of ribose of some nucleotides, and at N-6 of adenine of some nucleotides

### (secondary methylations)

بس تطلع عالسيتوبلازم ممكن يصيرلها methylation بأماكن مختلفة

#### Importance of capping:

- It protects the 5<sup>\</sup> end of the mRNA from 5<sup>\</sup>
  <u>exonuclease</u> enzyme.
- It helps its recognition by the ribosome.
- It helps the initiation of protein synthesis.
- Eukaryotic mRNA lacking the cap are not efficiently translated.
- Helps transport of mRNA to the cytoplasm.

# **B. Addition of poly(A) tail:**

- ✓ The final RNA processing event, polyadenylation of the 3<sup>\</sup> end of the mRNA, is intimately linked with the termination of transcription
- ✓ It is the addition of poly- A tail at the 3<sup>\</sup> end of mRNA (100-200 A bases).
- This poly–A tail is not transcribed from DNA but added after transcription by the enzyme polyadenylate polymerase using ATP as a substrate.
- ✓ This occurs after the mRNA is cleaved 15-20 nucleotides downstream from the <u>AAUAAA</u> recognition sequence.
- ✓ The poly–A tail immediately binds several copies of a poly
  (A) binding proteins that protect mRNA against 3<sup>\</sup>
  <u>exonuclease.</u>



#### Importance of poly-A tail:

- It stabilizes the mRNA & protects it from exonucleases enzymes. The length of poly (A) tail determines the half life time of mRNA.
- Increases the efficiency of translation.
- It facilitate their exit from the nucleus .After the mRNA enters the cytosol, the poly-A tail is gradually shortened. اكتر بنلاقي ال الما بقصر (cytoplasm)

#### C. Removal of introns and splicing of exons :

It means excision of introns and joining the ends of exons to <u>leave only the functional</u> <u>mRNA molecule.</u>

This process occurs in the nucleus by the help of the small nuclear ribonucleoproteins (snRNP, or snurps) which are composed of small nuclear RNA (snRNA) and proteins.

 Snurps acting on mRNA are called <u>spliceosomes</u>. This is an example of catalytic RNAs or RNA enzymes, which are termed
 <u>ribozymes</u>. RNA وركيبها enzymes والمالي والمراجع عنوارة عن وركيبها enzymes

• The sequence of bases at the exon-intron junction determines the site of splicing.

\* ال sequence بين ال introns و ال exons هو الي بحدد لل snurps وين يقطع



تركيب الهيموغلوبين : 4 سلاسل عديد ببتيد : 2 alpha , 2 Beta

One type of β thalassemia appears to result from nucleotide change at the exon-intron junction leading to failure to remove intrones, reducing the synthesis of the β globin chain.

مرض الذئبة الحمراء Patients with systemic lupus erythematosis (SLE) produce antibodies against snRNP. خطوة ال splicing ما رح تصبر

Histone mRNAs (replication-dependent histones that are expressed during the <u>S</u>phase of the cell cycle) do not contain introns.

بكون عالي جداً بال S-phase لانه ال DNA replication بصير هناك محمد الله منيح لانه اذا ما في introns يعني ما رح يصير snRNA بكون عالي جداً بال عليها لو صار مشكلة بال

\* مهمة بال Variations

# Two advantages are suggested for having protein- coding genes organized as exons & introns:

- 1- Alternative splicing may lead to the formation of different types or new types of mRNA molecules or proteins.
- 2- Also this will <u>decreases the possibility of</u> <u>effective mutations</u>, (that result in protein abnormalities or disease), if it occurs at the regions of introns. \* وجود ال introns من حدوث الطفرات لإنه



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- Alternative splicing, or differential splicing, is a regulated process during gene expression that results in a single gene coding for multiple proteins.
- In this process, particular exons of a gene may be included within or excluded from the final, processed messenger RNA (mRNA) produced from that gene.
- Notably, alternative splicing allows the human genome to direct the synthesis of many more proteins than would be expected from its 20,000 protein-coding genes. عد البروتينات يفوق عدد الجينات