Chapter 10:

The Structure and Function of DNA

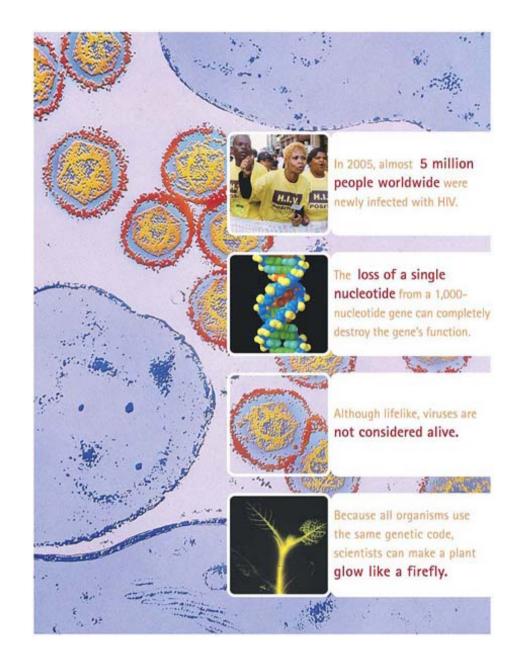
covers

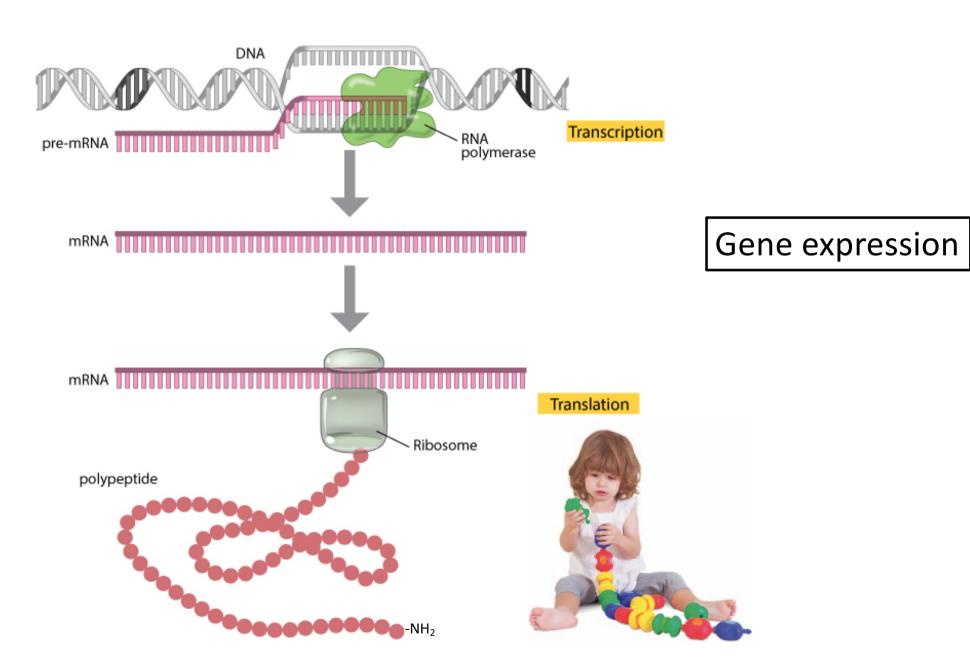
DNA replication

DNA transcription

mRNA translation

Chapter 11 covers the regulation of gene expression





What "tells" the RNA polymerase where to start transcription of a gene? → Promoter sequence

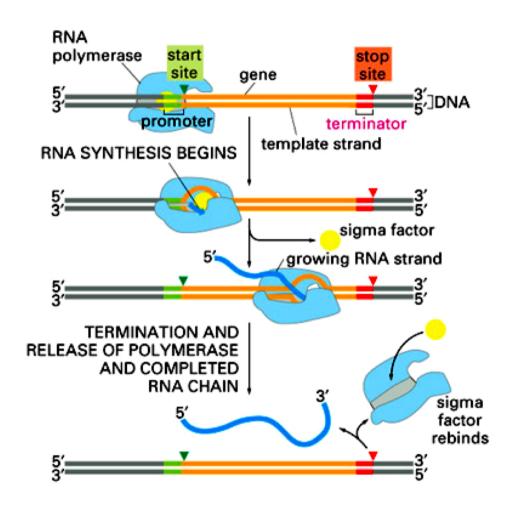
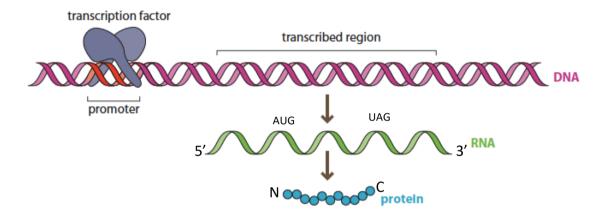


Figure shows transcription in E. coli

The enzyme RNApolymerase does not recognize promoter regions.

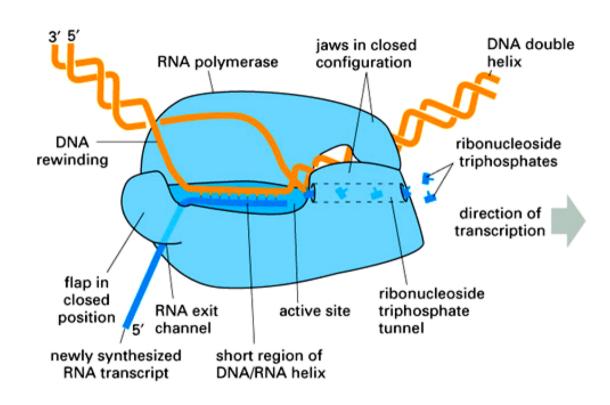


Transcription factors

- · are proteins
- · bind *specifically* to certain promoters
- · recruit the enzyme RNA polymerase
- · position the RNApolymerase (give a direction)

RNA polymerase: an enzyme made of several polypeptides

E. coli RNA polymerase



- no need for primer
- uses ATP, CTP, GTP, UTP
- in 5' -> 3' direction
- error rate: 1/10⁴ (DNA polymerase: 1/10⁷)

All eucaryotes have 3 RNA polymerases.

A protein family

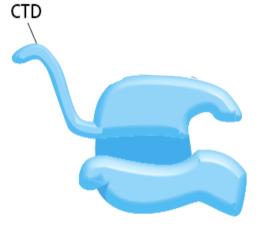


RNApol I RNA

28S rRNA

18S rRNA

5.8S rRNA



RNApol II

mRNA

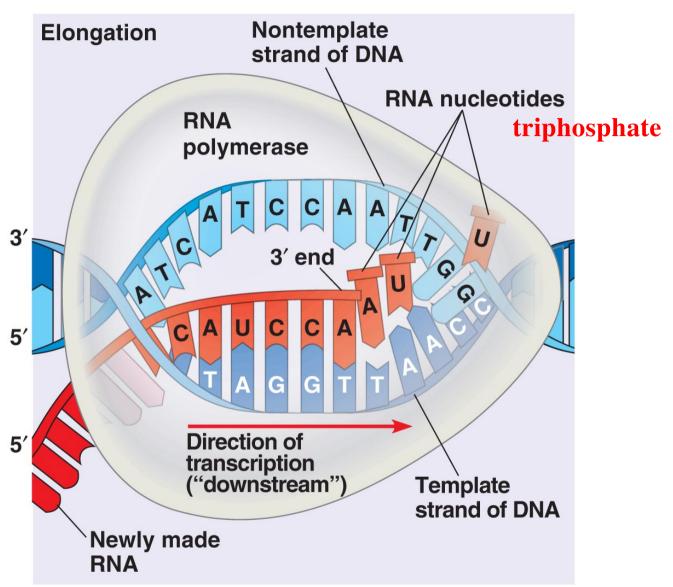
Only RNApol II has a C - Terminal Domaine (CTD)



RNApol III

5S rRNA

tRNA



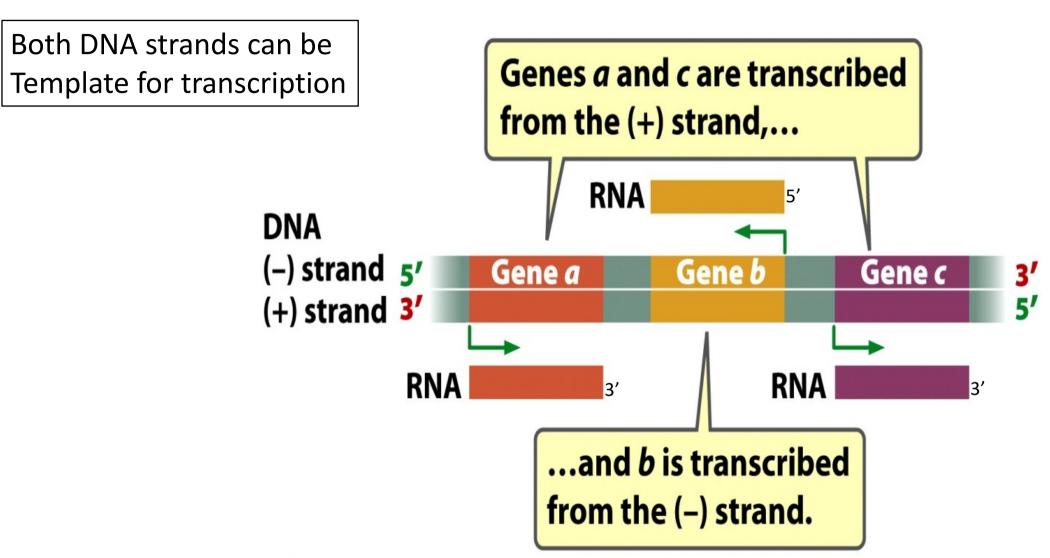
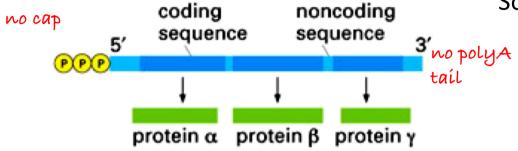


Figure 13-6

Genetics: A Conceptual Approach, Third Edition
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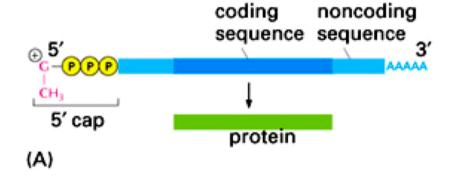
Differences between prokaryotic and eukaryotic messengers

bacterial mRNA



Some prokaryotic messengers are polycistronic.

eucaryotic mRNA



Eukaryotic messengers are always monocistronic.

No cap No introns No polyA tail

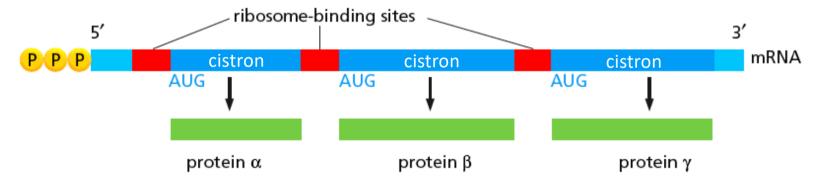


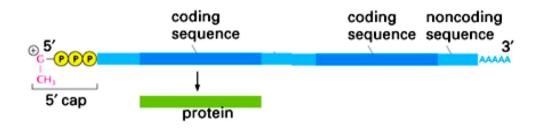
Figure 6–73 Structure of a typical bacterial mRNA molecule.

Only ~ 20% of bacterial mRNAs are *polycistronic*

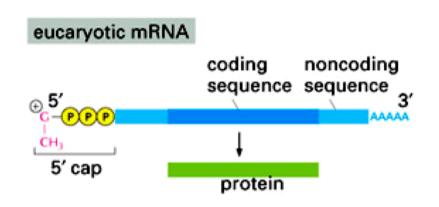
Usually proteins encoded by the same messenger work together.

Eukaryotic messengers are monocistronic.

If you engineer a bicistronic messenger only the upstream cistron is translated.



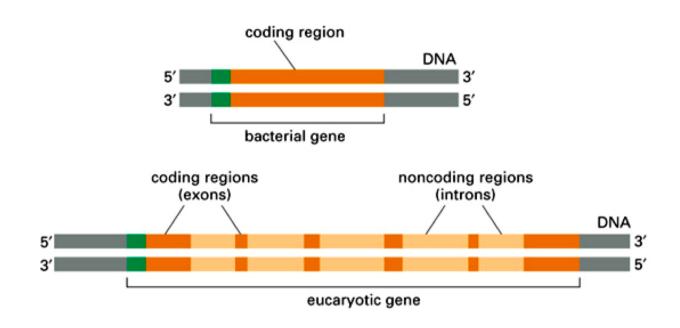
In eukaryotes: processing of primary transcripts into mature messenger RNAs.



processing includes these modifications:

- A. 5'-end: addition of "cap"
 - → protects mRNA
 - → start of translation
- B. 3'-end: addition of AAAAA-tail
 - → protects mRNA
- C. removal of introns
 - → splicing

Eukaryotic genes are composed of exons and introns



Exons: sequences encoding the protein

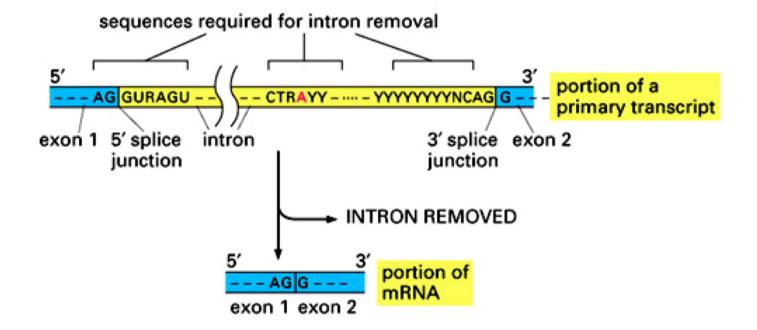
Introns: non-coding sequences

- → Exons often encode functional domains of the protein
- → Regulatory sequences can be in introns
- → Alternative splicing possible (see next slides)

Correct definition of an exon:
part of a primary transcript
not eliminated by splicing.
Coding or non coding has no
place in the definition.



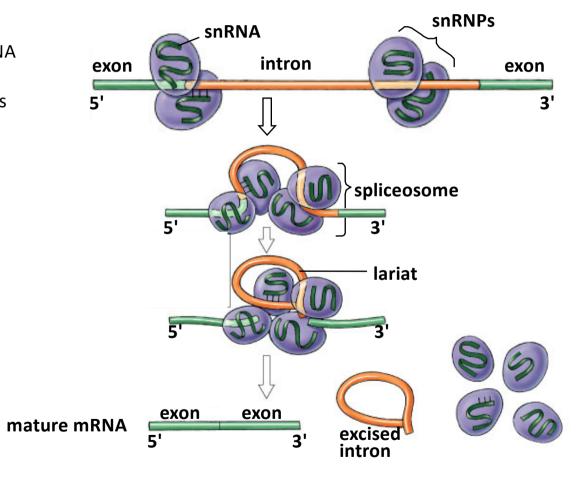
Splicing

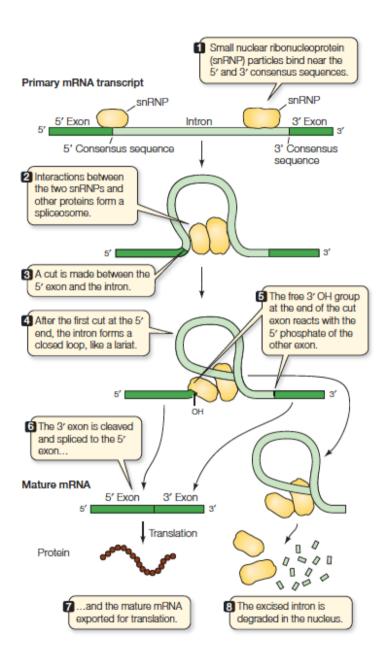


Splicing

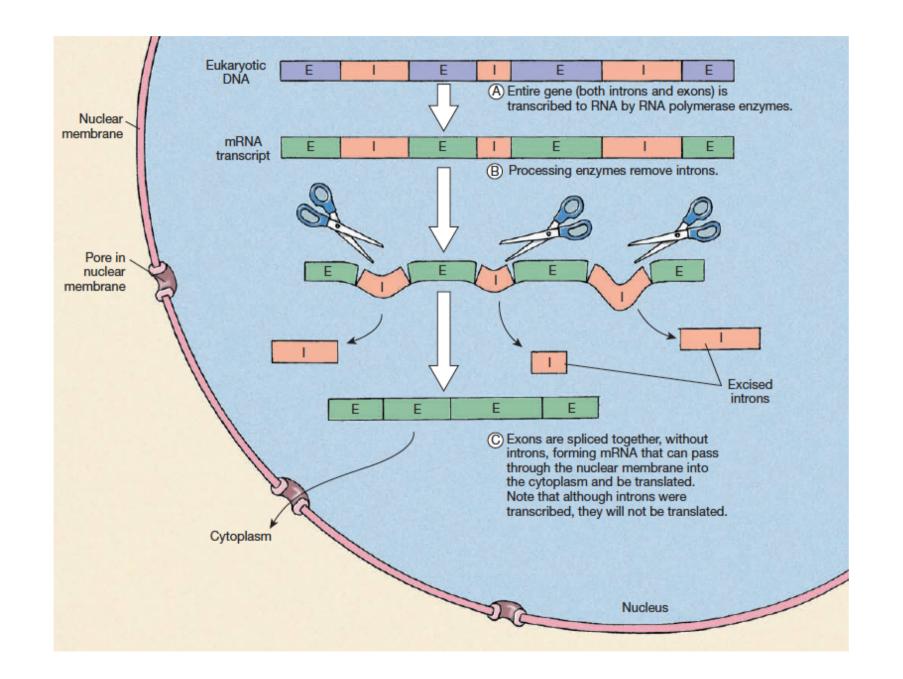
snRNA: small nuclear RNA

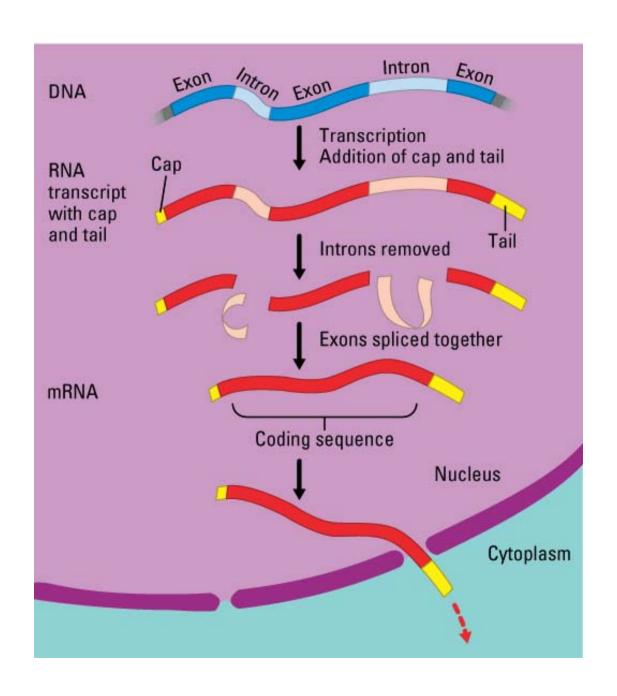
snRNP : snRNA + proteins





Introns are degraded→ nucleotids are recycled



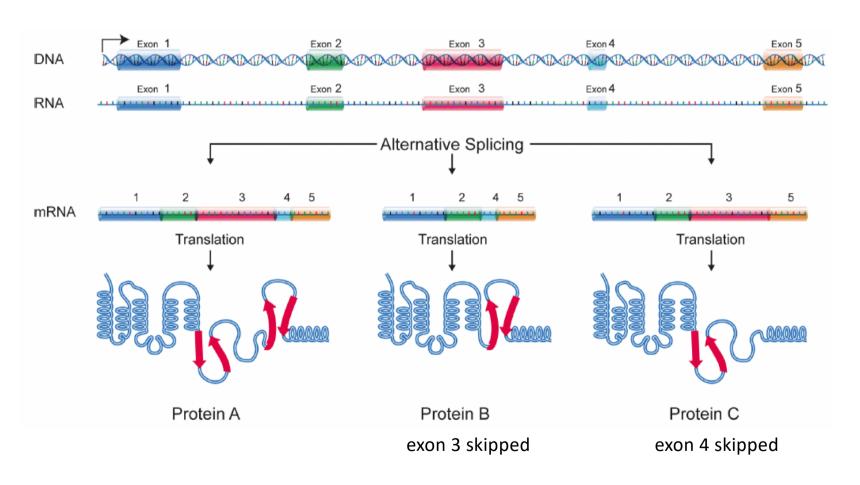


Essential Biology Fig 10.14

5'-end first

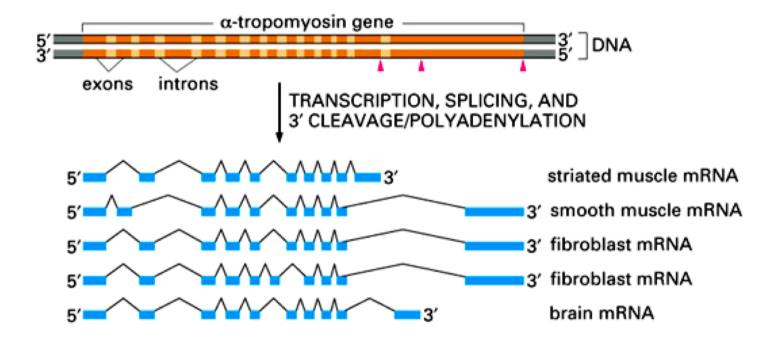
Alternative splicing

General idea



RNA splicing can create different mRNAs from 1 primary RNA

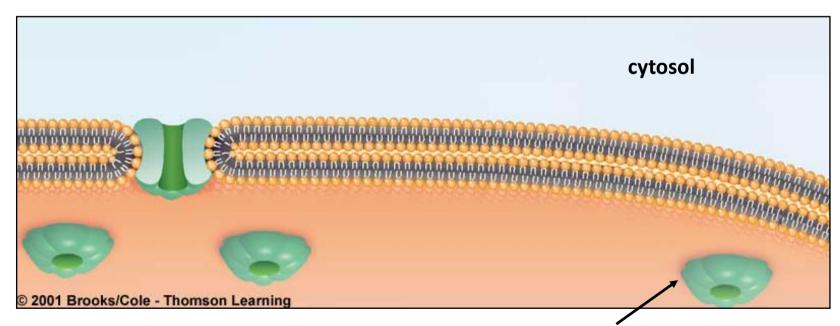
→ alternative splicing



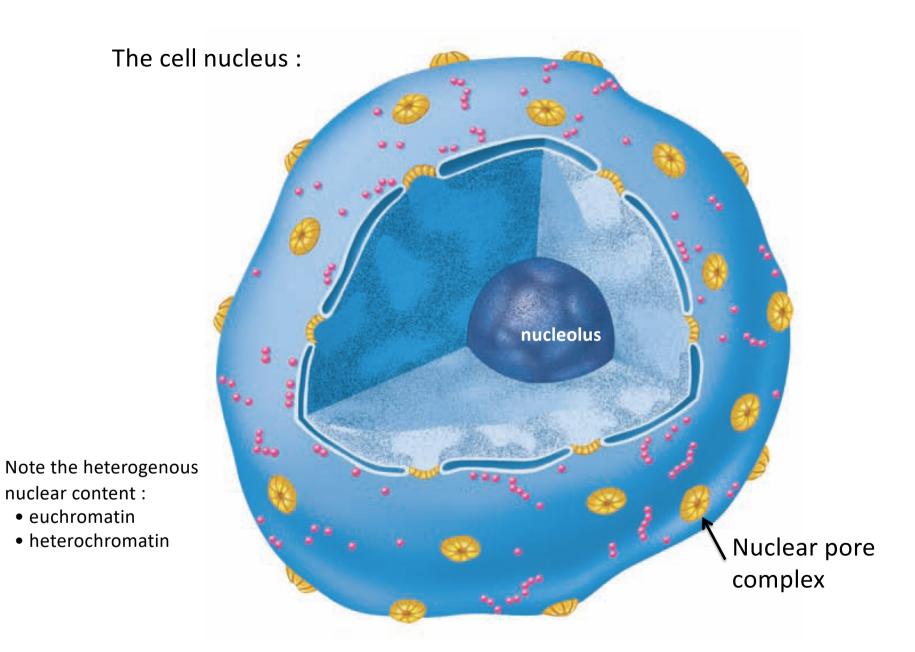
Nuclear envelope and nuclear pores.

2 membranes:

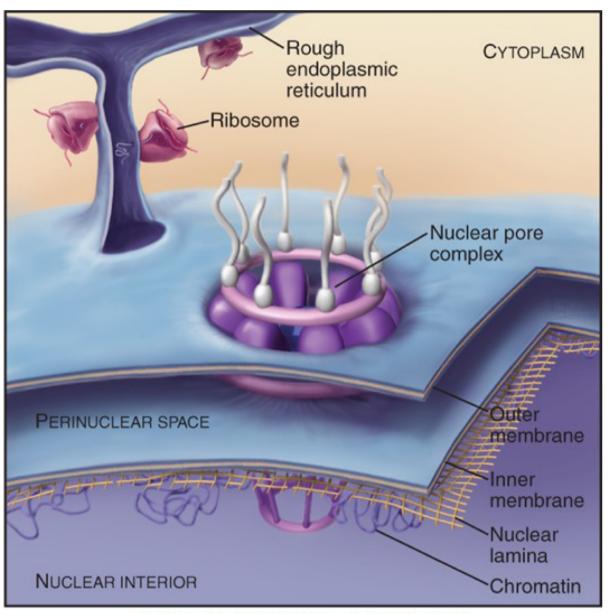
- inner nuclear membrane
- outer nuclear membrane
- perinuclear space (between the 2 membranes)



Nuclear pore made of proteins

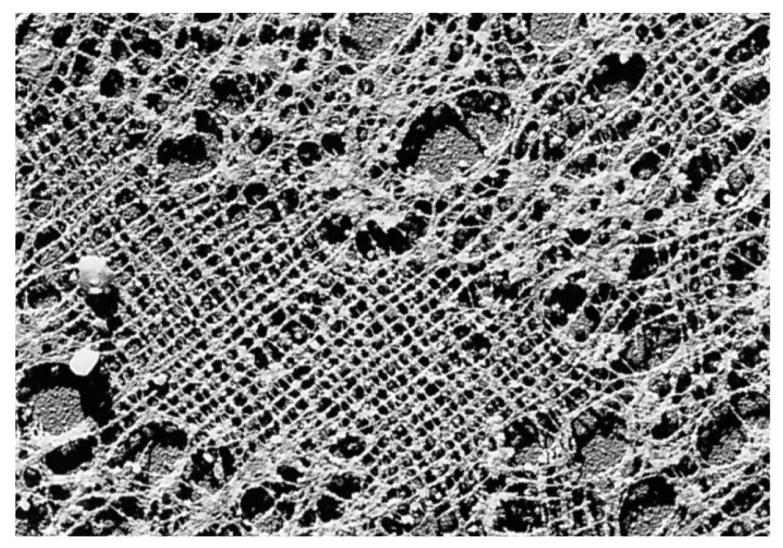


The nuclear enveloppe Inner nuclear Outer nuclear membrane membrane



Nuclear lamina

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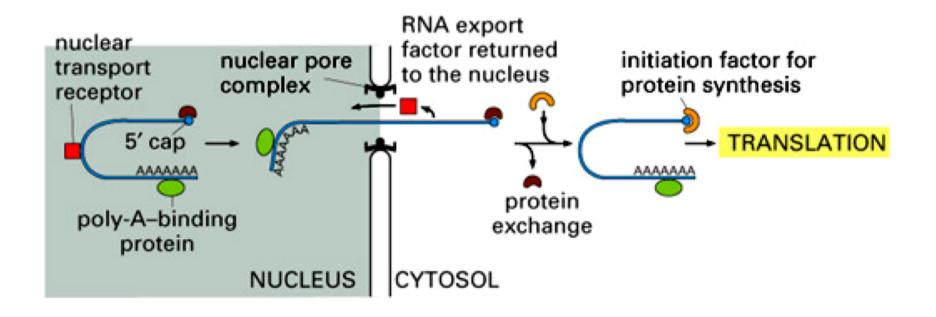


The nuclear lamina

 $1 \, \mu m$

mRNA nuclear export

a complex process



Only fully spliced RNA are exported

