University of Anbar Department of Biotechnology

College of Science

Dr. Huda Musleh Biotechnology 1

# Lecture 3 GeneTranscription and Translation

# In this lecture...

- Central Dogma
- RNA and proteins
- Codons
- Transcription
  - Initiation
  - Elongation
  - Termination
- Translation
- Post-translational modification
- Mutations

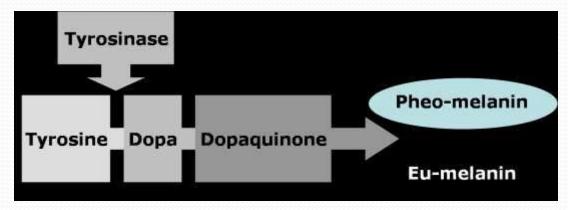
# Proteins link genotype and

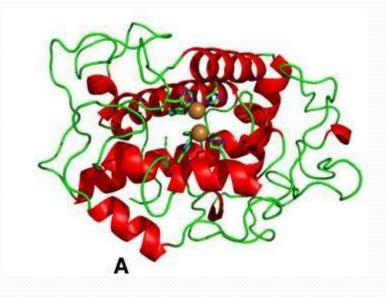
# phenotype



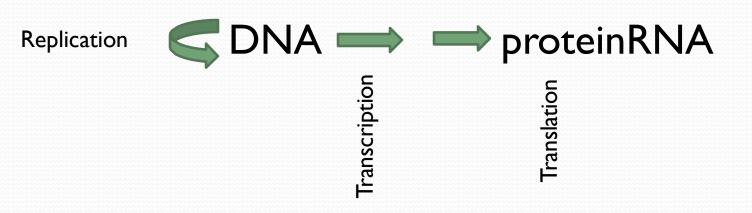
- Albinism occurs when the protein tyrosinase is defective
- Tyrosinase directs the synthesis of melanin, a pigment found in skin and eyes
- Heterozygotes with one functional copy of tyrosinase produce enough melanin to not display the phenotype
- However, recessive homozygotes will lack all pigment

# Melanin synthesis pathway and GPR413 structure







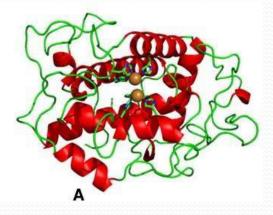


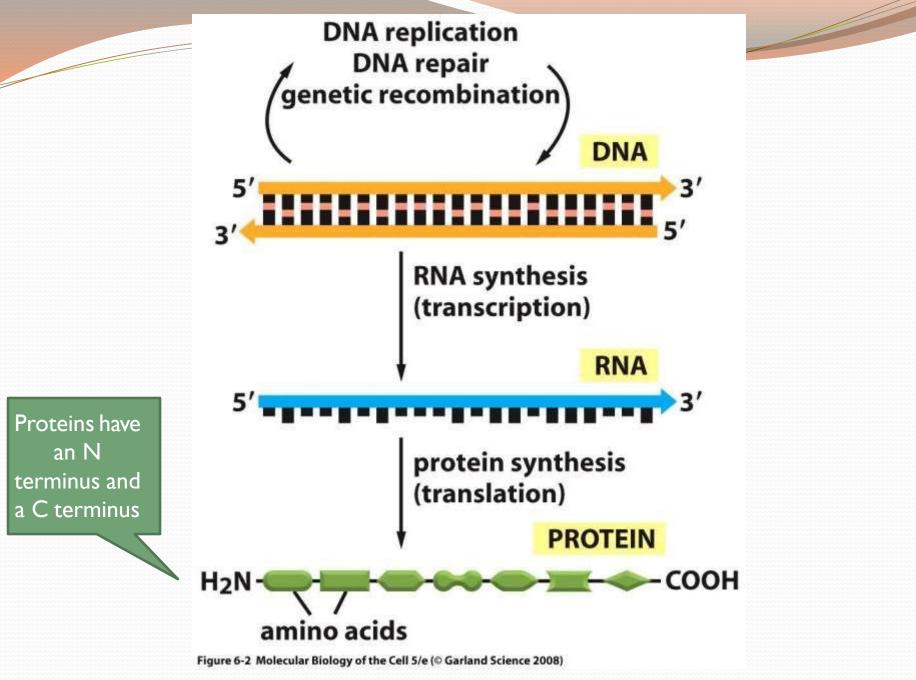
Primary structure of tyrosinase protein

gi|209571475:5001-122888 Homo sapiens tyrosinase (oculocutaneous albinism IA) (TYR), RefSegGene on chromosome [] ATCACTGTAGTAGTAGCTGGAAAGAGAAATCTGTGA CTCCAATTAGCCAGTTCCTGCAGACCTTGTGAGG ACTAGAGGAAGAATGCTCCTGGCTGTTTTGTACTGC CTGCTGTGGAGTTTCCAGACCTCCGCTGGCCATT TCCCTAGAGCCTGTGTCTCCTCTAAGAACCTGATGG AGAAGGAATGCTGTCCACCGTGGAGCGGGGGACAG GAGTCCCTGTGGCCAGCTTTCAGGCAGAGGTTCCT GTCAGAATATCCTTCTGTCCAATGCACCACTTGGG CCTCAATTTCCCTTCACAGGGGTGGATGACCGGGA GTCGTGGCCTTCCGTCTTTTATAATAGGACCTGCC AGTGCTCTGGCAACTTCATGGGATTCAACTGTGGAA ACTGCAAGTTTGGCTTTTGGGGGACCAAACTGCAC AGAGAGACGACTCTTGGTGAGAAGAACATCTTCG ATTTGAGTGCCCCAGAGAAGGACAAATTTTTGCC TACCTCACTTTAGCAAAGCATACCATCAGCTCAGAC TATGTCATCCCCATAGGGACCTATGGCCAAATGA AAAATGGATCAACACCCATGTTTAACGACATCAATA TTTATGACCTCTTTGTCTGGATGCATTATTATGT GTCAATGGATGCACTGCTTGGGGGGATCTGAAATCT GGAGAGACATTGATTTTGCCCATGAAGCACCAGCT TTTCTGCCTTGGCATAGACTCTTCTTGTTGCGGTGG GAACAAGAAATCCAGAAGCTGACAGGAGATGAAA ACTTCACTATTCCATATTGGGACTGGCGGGGATGCAG AAAAGTGTGACATTTGCACAGATGAGTACATGGG AGGTCAGCACCCCACAAATCCTAACTTACTCAGCC CAGCATCATTCTTCTCCTCTTGGCAGGTAAGATAT GCTAGATATACGATGTCAGAGTAGGGAGGAACCTT AACAATCACTTCTTCAGGCAGGGTATAAACTTCTC ACCTGAACACTCATTGCAGCCCCCATCAAGGACAG AAATGGTGCCCTGTTAAGAACTCTCAATGTATCTT

>gi|403422|gb|AAB60319.1| tyrosinase [Homo sapiens] MLLAVLYCLLWSFQTSAGHFPRACVSSKNLMEKECCPP WSGDRSPCGQLSGRGSCQNILLSNAPLGPQFP FTGVDDRESWPSVFYNRTCOCSGNFMGFNCGNCKF GFWGPNCTERRLLVRRNIFDLSAPEKDKFFAYLTL AKHTISSDYVIPIGTYGQMKNGSTPMFNDINIYDLFVW MHYYVSMDALLGGSEIWRDIDFAHEAPAFLPW HRLFLLRWEQEIQKLTGDENFTIPYWDWRDAEKCDIC TDEYMGGOHPTNPNLLSPASFFSSWOIVCSRLE EYNSHQSLCNGTPEGPLRRNPGNHDKSRTPRLPSSADV **EFCLSLTQYESGSMDKAANFSFRNTLEGFASP LTGIADASOSSMHNALHIYMNGTMSOVOGSANDPIFLL** HHAFVDSIFEQWLQRHRPLQEVYPEANAPIGH NRESYMVPFIPLYRNGDFFISSKDLGYDYSYLQDSDPDSF **QDYIKSYLEQASRIWSWLLGAAMVGAVLTA** LLAGLVSLLCRHKRKQLPEEKOPLLMEKEDYHSLYOSHL

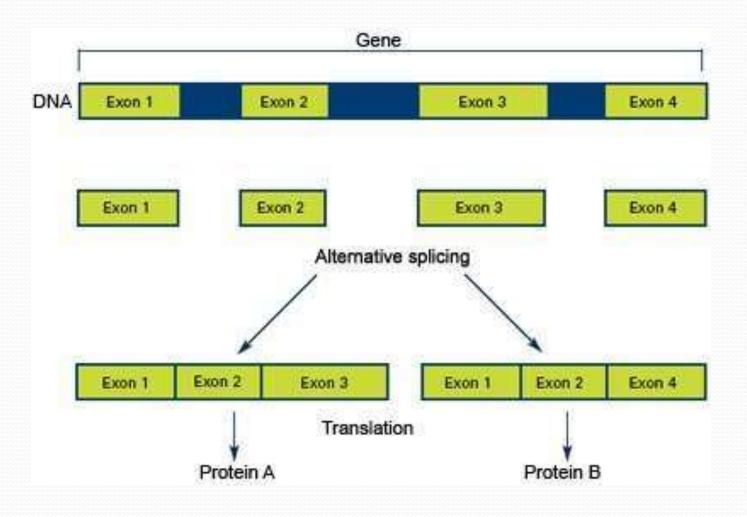
#### Quaternary structure of tyrosinase protein





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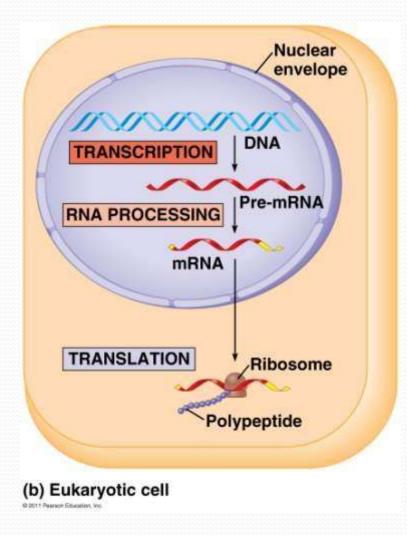
# One gene, one protein hypothesis



# How do genes produce proteins?

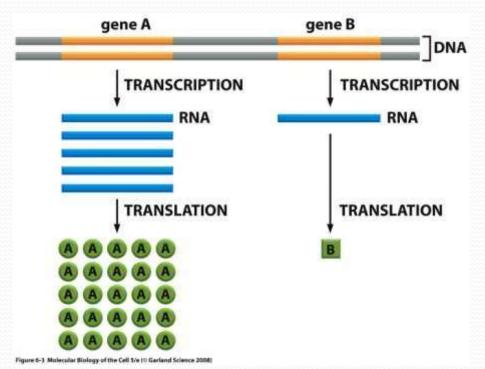
Gene expression: the process by which genes produce proteins

- Two stages:
- Transcription
- Translation



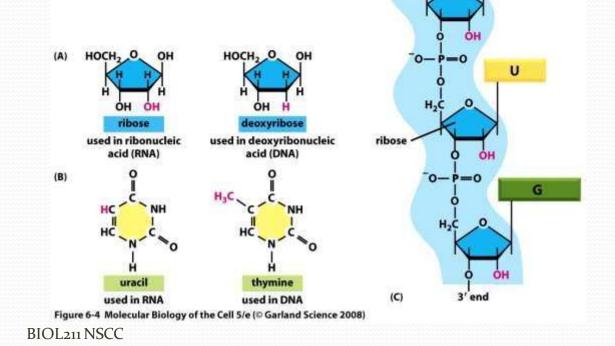
#### Genes can be expressed at different efficiencies

- Gene A is transcribed much more efficiently than gene B
- This allows the amount of protein A in the cell to be greater than protein B
- The lower expression of gene B is a reason behind incomplete dominance



# A reminder: what is RNA?

 RNA is the bridge between genes and the proteins for which they code



5' end

0-P=0

0-P=0

C

bases

A

Type of RNA	Functions
Messenger RNA (mRNA)	Carries information specifying amino acid sequences of proteins from DNA to ribosomes
Transfer RNA (tRNA)	Serves as translator molecule in protein synthesis; translates mRNA codons into amino acids
Ribosomal RNA (rRNA)	Plays catalytic (ribozyme) roles and Structural roles in ribosomes
Primary transcript	Is a precursor to mRNA, rRNA, or tRNA, before being processed; some intron RNA acts as a ribozyme, catalyzing its own splicing
Small nuclear RNA (snRNA)	Plays structural and catalytic roles in spliceosomes, the complexes of protein and RNA that splice pre-mRNA

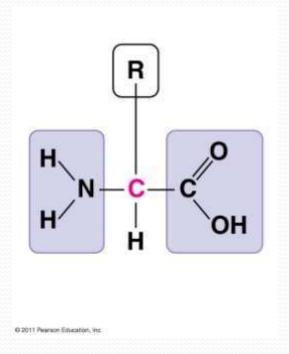
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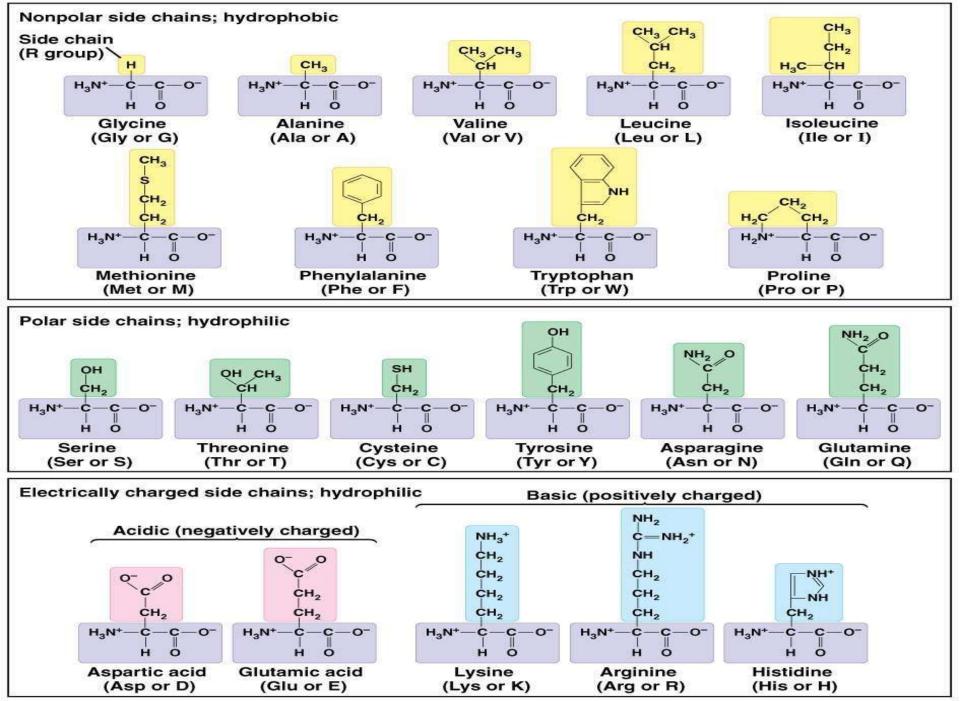
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# A reminder: what are proteins made of?

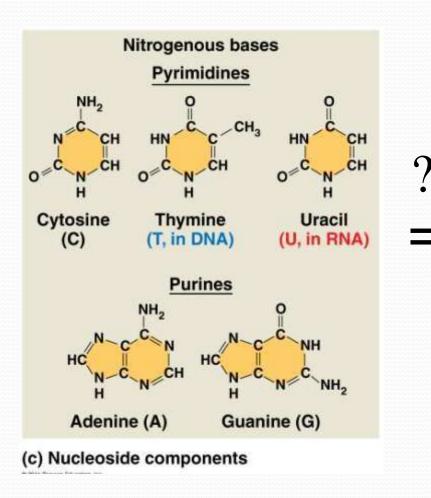
• Monomers of proteins are amino acids

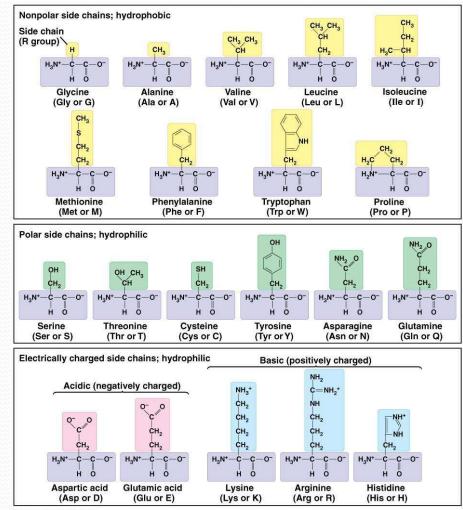
There are 20 amino acids Each has a different property depending on its R group/side chain





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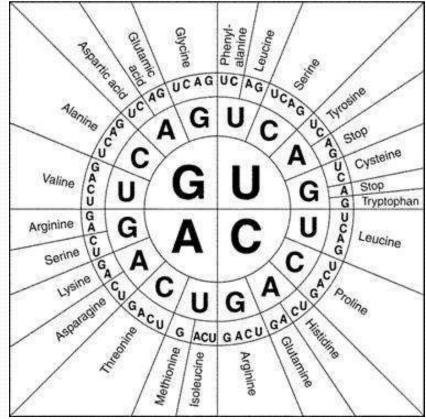




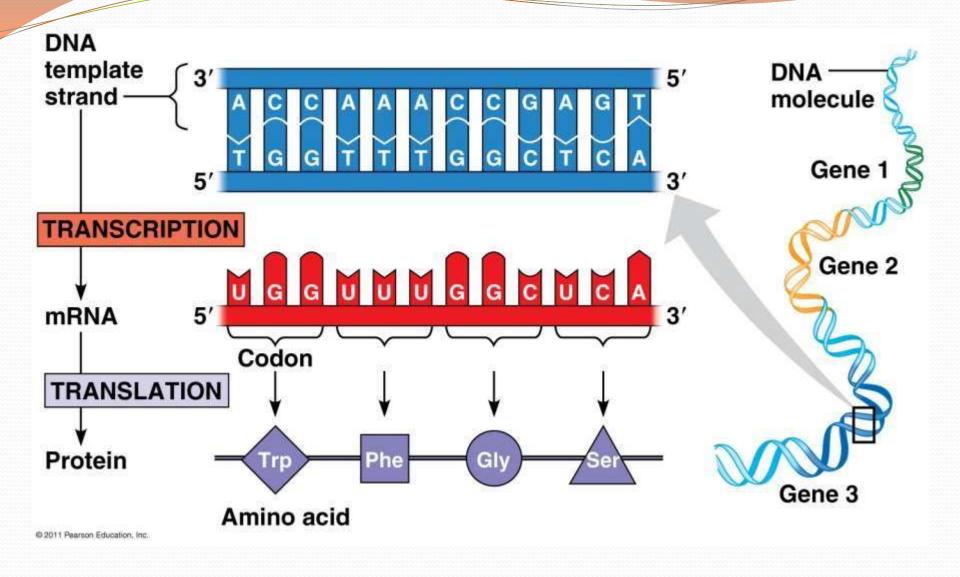
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Codons

- A **codon** is three nucleotides in a row on an RNA molecule that codes for a single amino acid
- A specific three-nucleotide sequence encodes for each amino acid



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#### **Template Strand**

- During transcription, one of the two DNA strands, called the template strand, provides a template for ordering the sequence of complementary nucleotides in an RNA transcript
  - The template strand is always the same strand for a given gene
  - However, different genes may be on opposite strands

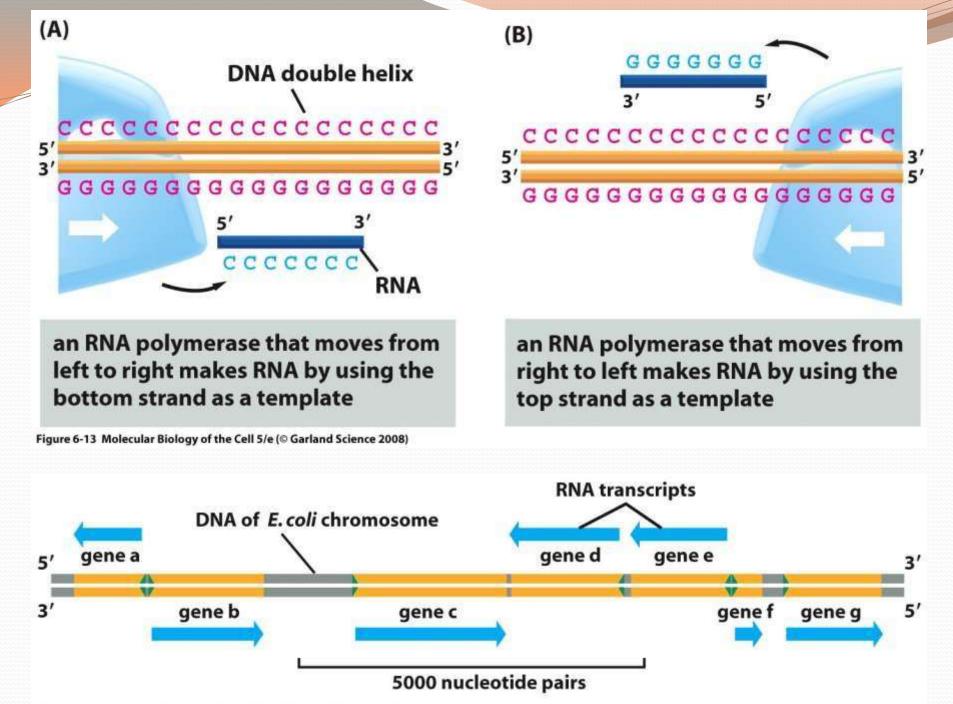


Figure 6-14 Molecular Biology of the Cell 5/e (© Garland Science 2008)

### Degeneracy of the code

- There are 64 codons, but only 20 amino acids
- One amino acid will have multiple codons
- The genetic code is said to be **degenerate** for this reason
- Each codon specifies the amino acid (one of 20) to be placed at the corresponding position along a polypeptide

Codons along an mRNA molecule are read by translation machinery in the 5' to 3' direction

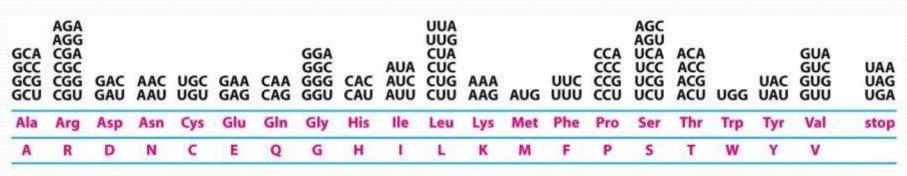
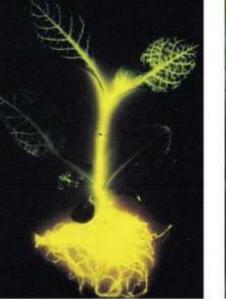


Figure 6-50 Molecular Biology of the Cell 5/e (© Garland Science 2008)

# **Evolution of the code**

- The genetic code is nearly universal, shared by the simplest bacteria to the most complex animals
  - Some species prefer certain codons (codon bias)
- Genes can be transcribed and translated after being transplanted from one species to another





(a) Tobacco plant expressing a firefly gene

(b) Pig expressing a jellyfish gene

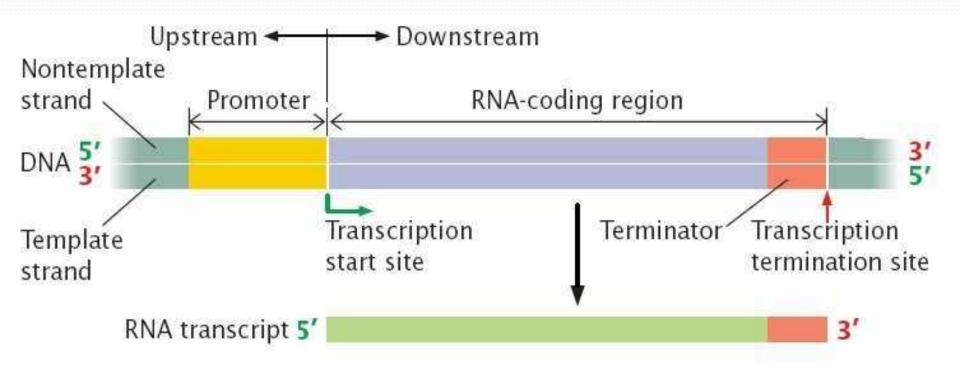


# Transcription DNA → RNA

- DNA is transcribed into a special type of RNA called messenger RNA, or mRNA
- mRNA synthesis is catalyzed by RNA polymerase, which pries the DNA strands apart and hooks together the RNA nucleotides
- The RNA is complementary to the DNA template strand
  - RNA synthesis follows the same base-pairing rules as DNA, except that uracil substitutes for thymine

# The Transcription Unit

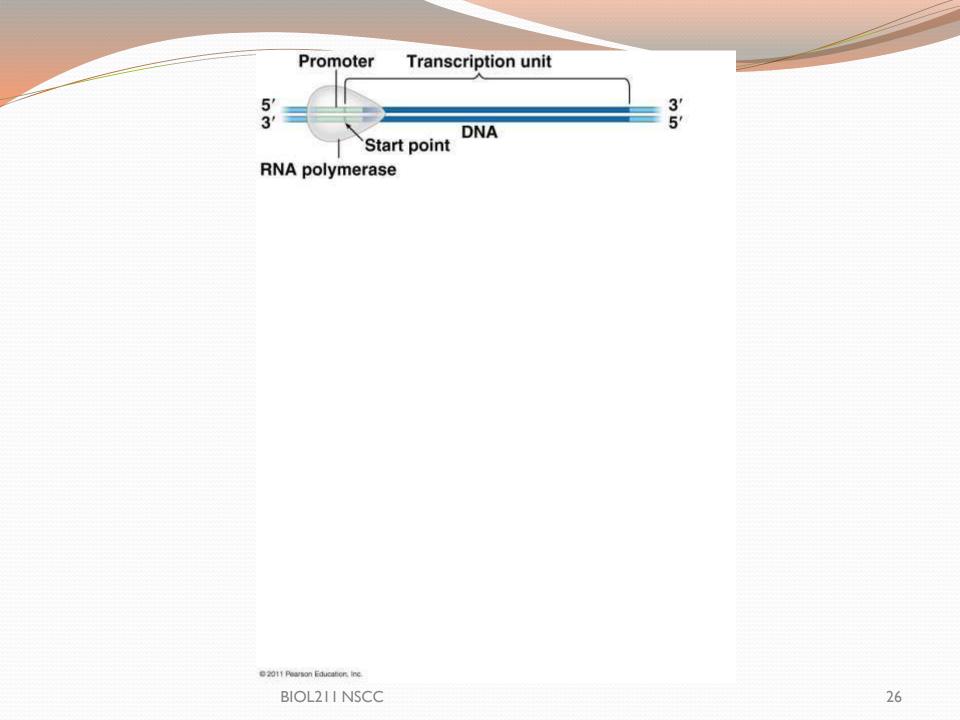
• The stretch of DNA that is used in transcription

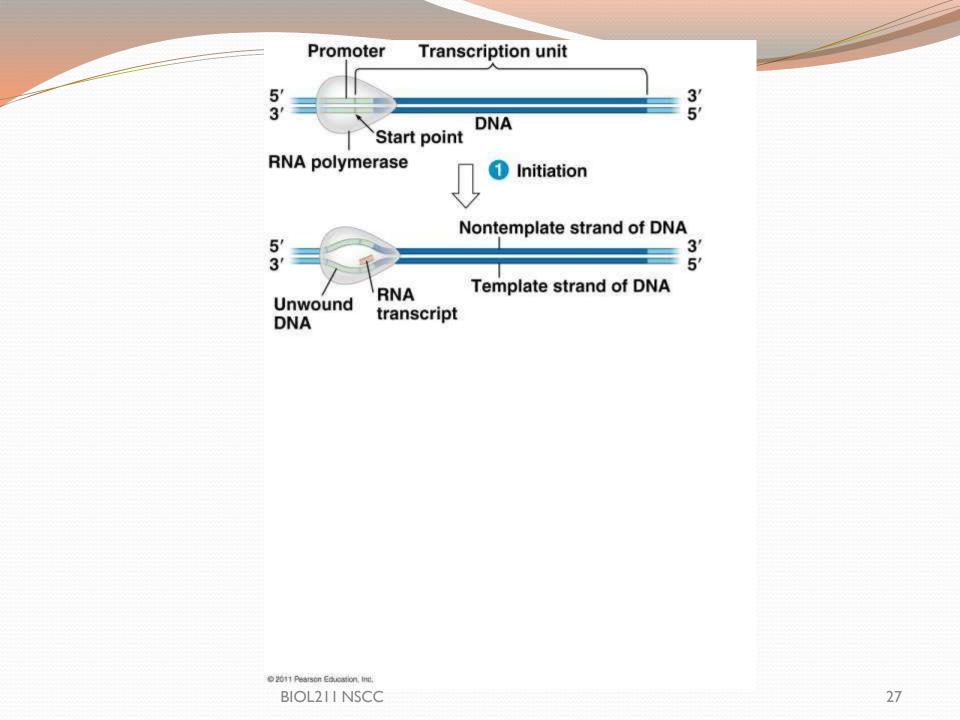


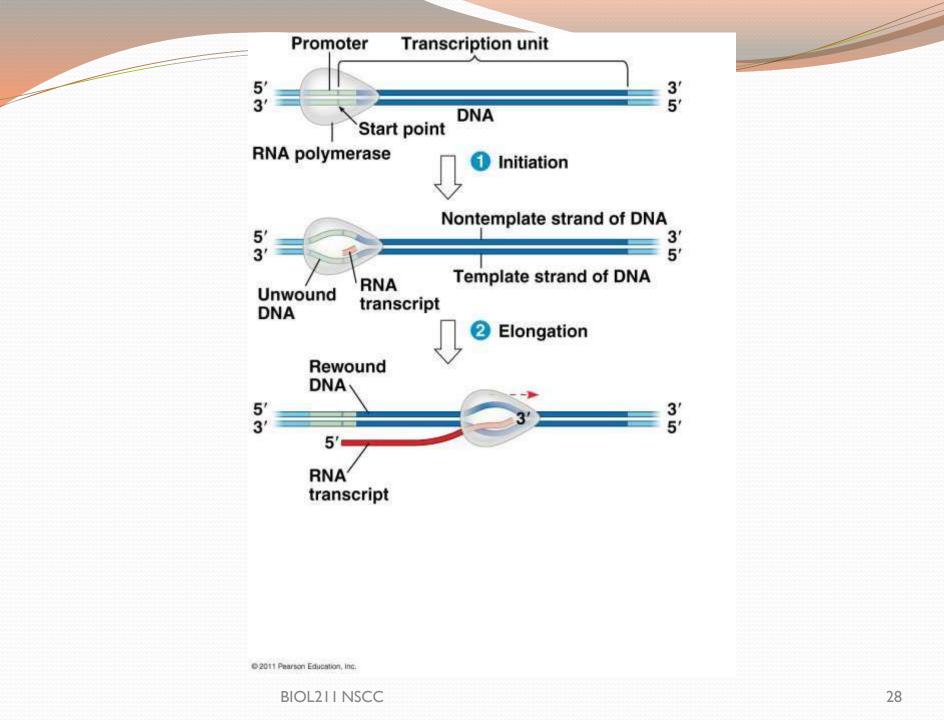
#### Transcription

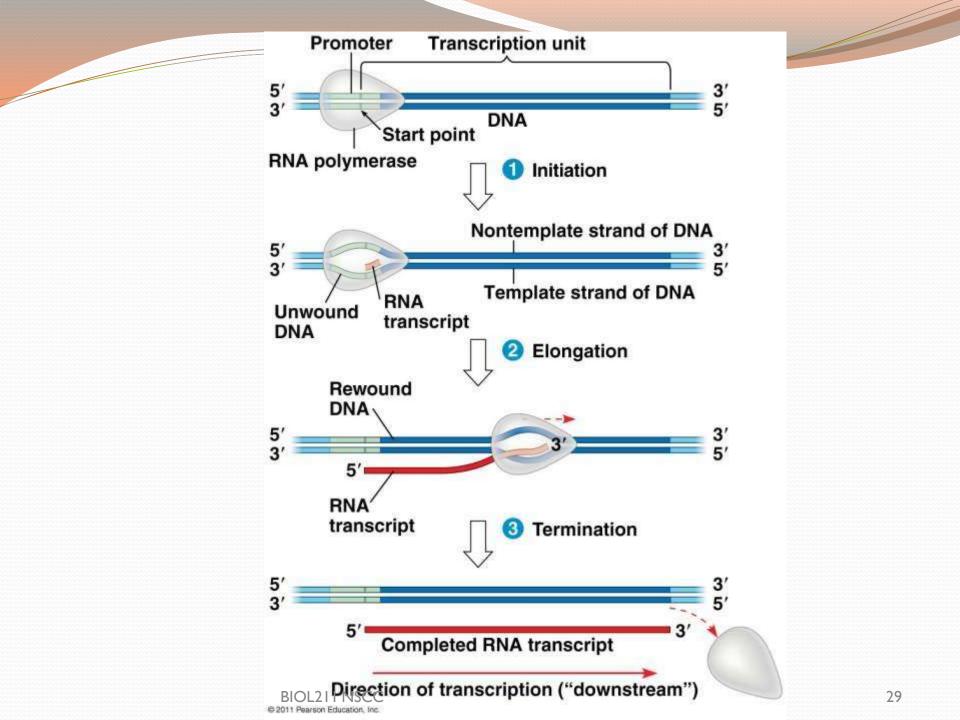
Generic process for both prokaryotes and eukaryotes

- Initiation
  - RNA polymerase binds to the promoter of a gene and begins to unwind the DNA
- Elongation
  - RNA pol 'reads' the template strand in 3' to 5' direction and adds complementary ribonucleotides
- Termination
  - RNA pol hits a stop signal (prokaryotes) or falls of (eukaryotes)









# **Transcription: Initiation**

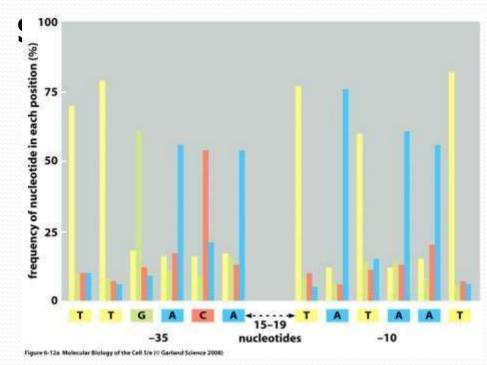
- **Promoters** control when, how, and at what level a gene is transcribed
  - Composed of a distinct sequence of nucleotides
- **Transcription factors** help RNA polymerase bind to the promoter
  - Transcription factors bind to areas in and around the promoter and provide a "landing pad" for RNA pol

# **Regulators of gene transcription**

- Promoters
- Enhancers
  - Upstream sequences in eukaryotes that help to control the expression of genes
  - Can be thousands of nucleotides away from the protein-coding region
- Silencers
  - When transcription factors bind, they prevent a gene from being transcribed

### **Promoters in prokaryotes**

- Prokaryotes
  - Highly conserved nucleotide sequences at -10 and
     -35 base pairs upstream of the start of the gene
    - -10 sequence is TATA, called "TATA box"



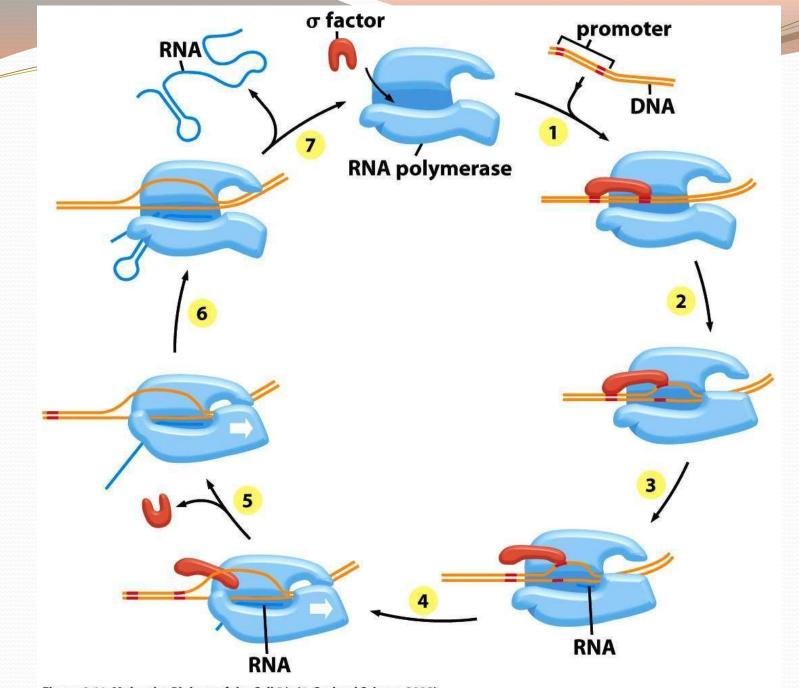


Figure 6-11 Molecular Biology of the Cell 5/e (© Garland Science 2008)

#### Promoters in eukaryotes

#### • Eukaryotes

- A "basal promoter" made of a TATA box and other elements ~25bp upstream
- A conserved region around the start of the gene called "Inr" for initiation
- Many different transcription factors

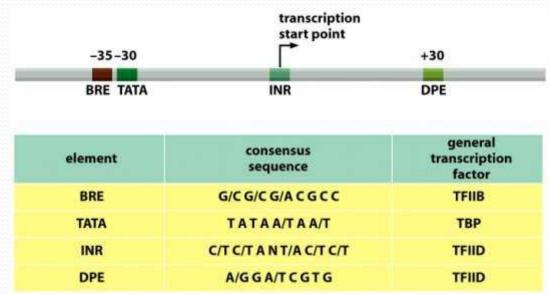
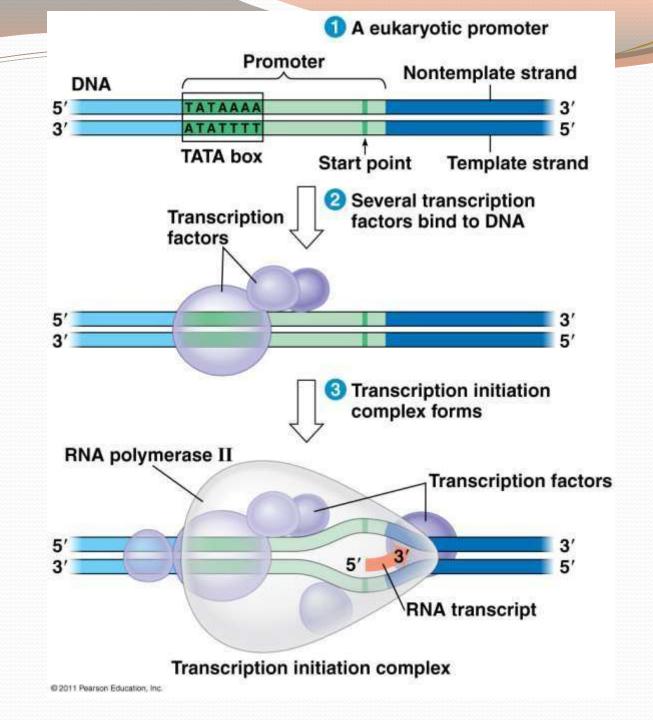
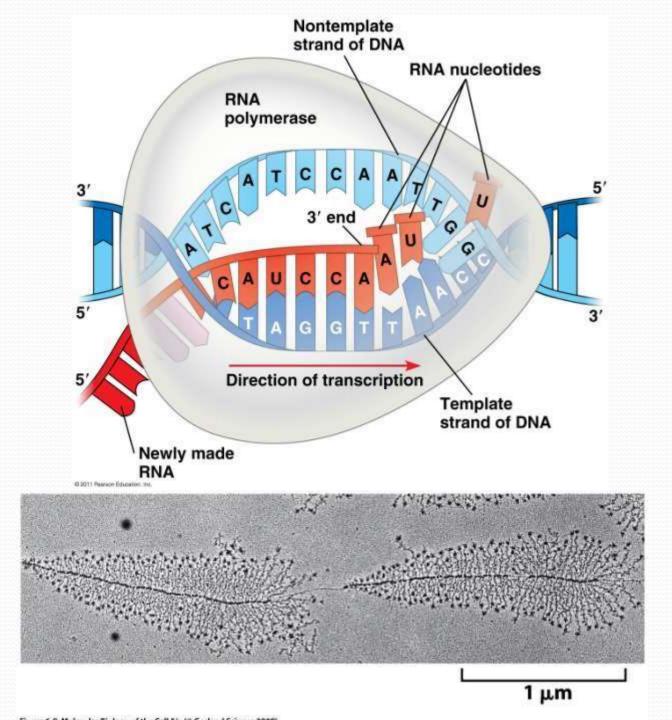


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### **Transcription: Elongation**

- As RNA polymerase moves along the DNA, it untwists the double helix, 10 to 20 bases at a time
- Transcription progresses at a rate of 40 nucleotides per second in eukaryotes
  - Actually 24/s in eukaryotes and 60/s in prokaryotes
- A gene can be transcribed simultaneously by several RNA polymerases
- Nucleotides are added to the 3' end of the growing RNA molecule



## **Transcription:Termination**

- The mechanisms of termination are different in bacteria and eukaryotes
  - In bacteria, the polymerase stops transcription at the terminator signal and the mRNA can then be translated without further modification

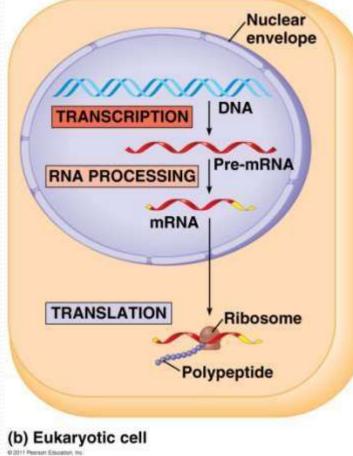
## Transcription:Termination

- In eukaryotes, RNA polymerase II transcribes the **polyadenylation signal sequence**
- The RNA transcript is released 10–35 nucleotides past this polyadenylation sequence
  - What exactly boots RNA pol off after the poly-A tail is unknown

## **RNA** Processing

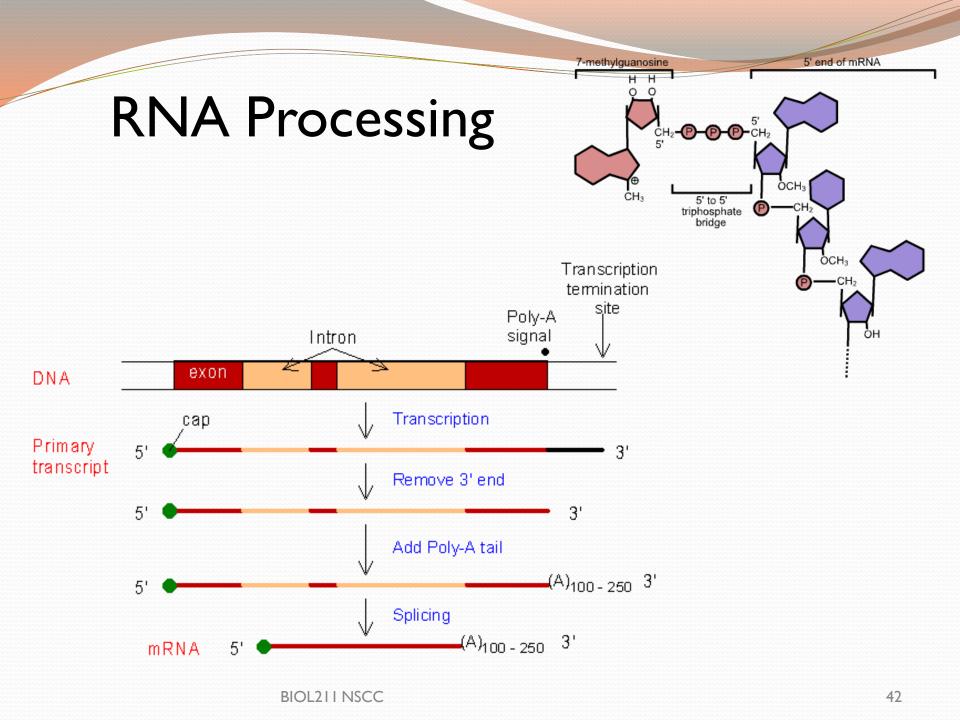
- In eukaryotes, before mRNA can be translated it has to first be processed
  - Prokaryotes don't need processing
- Pre-mRNA is renamed to mRNA after being processed

DNA pre-mRNA mRNA protein



## **RNA** Processing

- A pre-mRNA is processed in three particular ways:
  - The 5' end receives a modified nucleotide 5' cap
  - The 3' end gets a poly-A tail
  - Exons are spliced out
- These modifications share several functions
  - They seem to facilitate the export of mRNA
  - They protect mRNA from hydrolytic enzymes
  - They help ribosomes attach to the 5' end



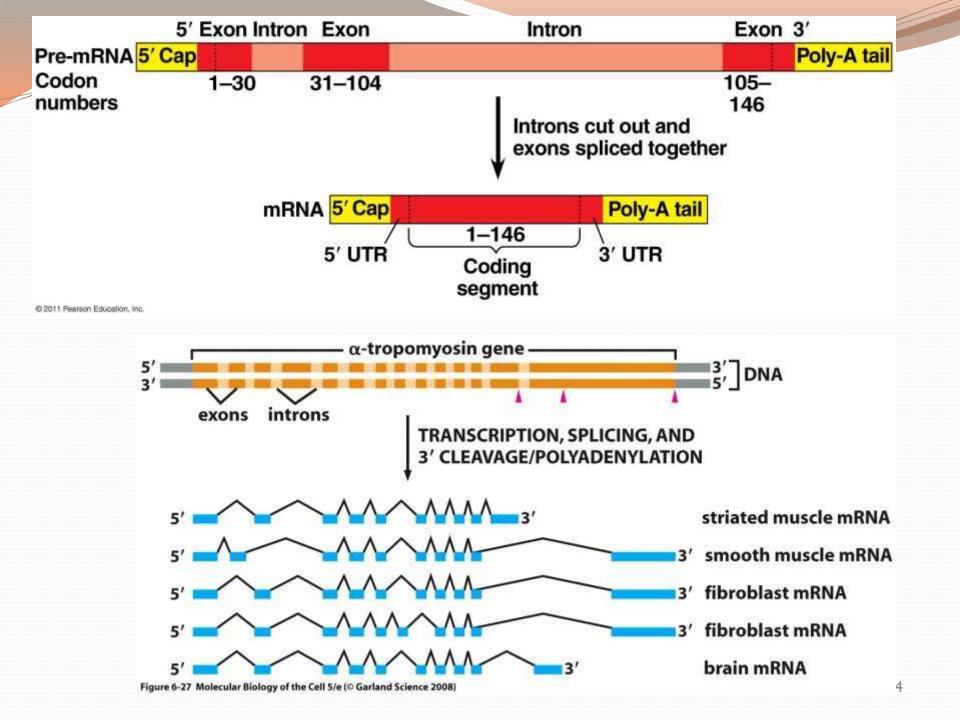
## mRNA splicing

Most eukaryotic genes and their RNA transcripts • have long noncoding stretches of nucleotides that lie between coding regions

These noncoding regions are called intervening • sequences, or **introns** 

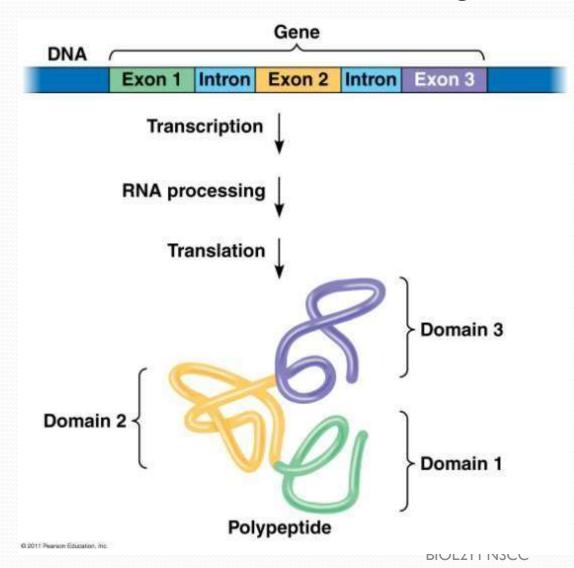
The other regions are called **exons** because they • are eventually expressed, usually translated into amino acid sequences

**RNA splicing** removes introns and joins exons, • creating an mRNA molecule with a continuous coding sequence



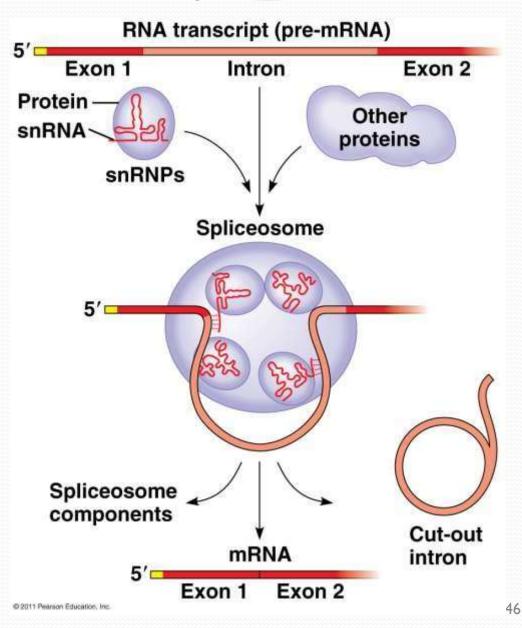
## **Alternative Splicing**

The rearrangement of exons during splicing •



#### How do we splice?

- Protein complexes called
   spliceosomes
   recognize and splice
   out introns
- Spliceosomes
   consist of a variety
   of proteins and
   several small nuclear
   ribonucleoproteins
   (snRNPs) that
   recognize the splice
   sites



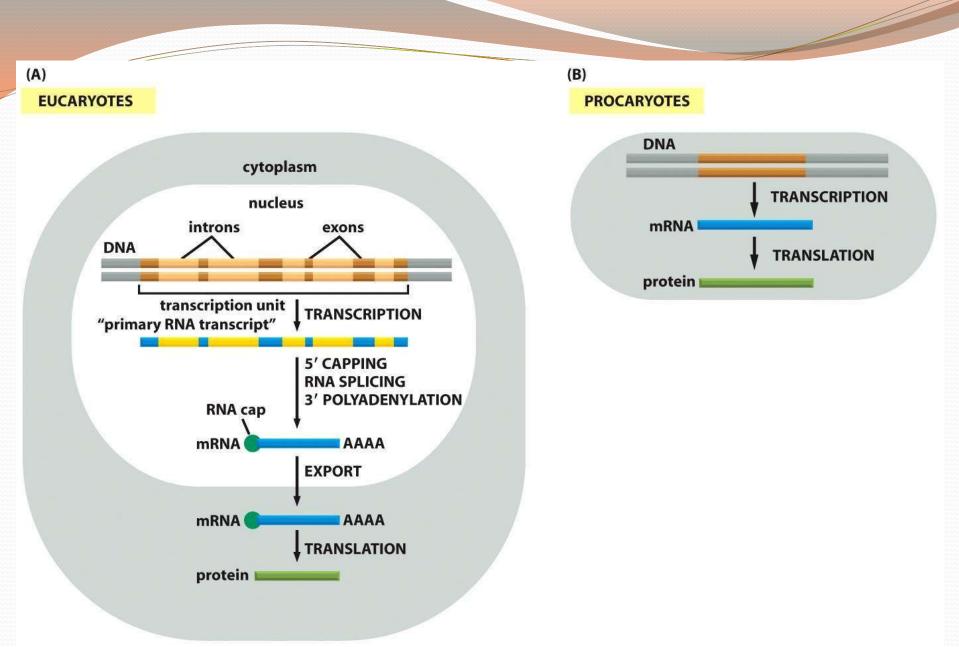
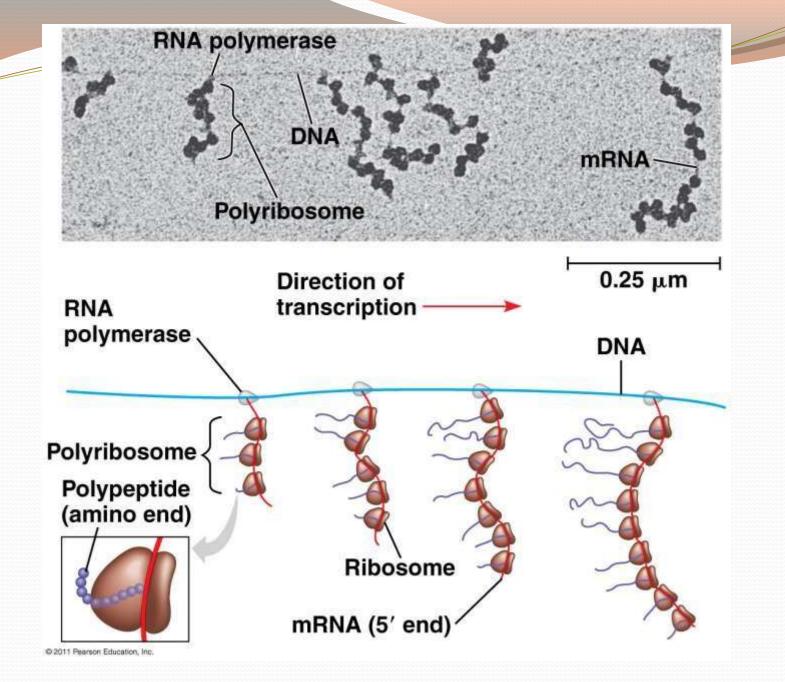


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- Bacteria and eukarya differ in their RNA polymerases, termination of transcription, and ribosomes; archaea tend to resemble eukarya in these respects
  - Bacteria can simultaneously transcribe and translate the same gene
  - In eukarya, transcription and translation are separated by the nuclear envelope
  - In archaea, transcription and translation are likely coupled



#### After processing, mRNA moves through nuclear

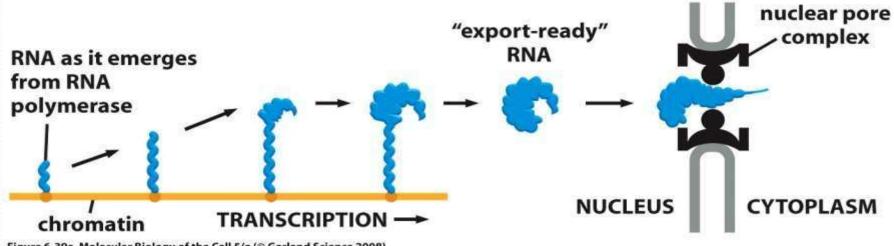
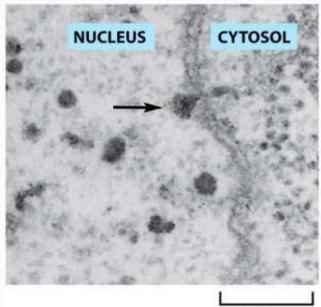


Figure 6-39a Molecular Biology of the Cell 5/e (© Garland Science 2008)



# Reference

• - Molecular biology of the cell, Garland Science , 2008.