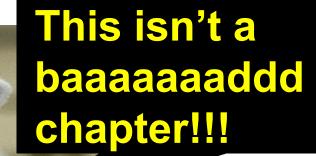
DNA Replication & Protein Synthesis



The Discovery of DNA's Structure

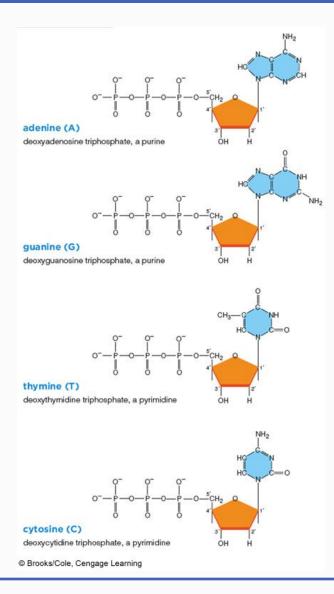
 Watson and Crick's discovery of DNA's structure was based on almost fifty years of research by other scientists

DNA's Building Blocks

Nucleotide

- A nucleic acid monomer consisting of a fivecarbon sugar (deoxyribose), three phosphate groups, and one of four nitrogen-containing bases
- DNA consists of four nucleotide building blocks
 - Two pyrimidines: **thymine** and **cytosine**
 - Two purines: adenine and guanine

Four Kinds of Nucleotides in DNA



Chargaff's Rules

- The amounts of thymine and adenine in DNA are the same, and the amounts of cytosine and guanine are the same: A = T and G = C
- The proportion of adenine and guanine differs among species

Watson and Crick's DNA Model

- A DNA molecule consists of two nucleotide chains (strands), running in opposite directions and coiled into a double helix
- Base pairs form on the inside of the helix, held together by hydrogen bonds (A-T and G-C)

Patterns of Base Pairing

- Bases in DNA strands can pair in only one way
 - A always pairs with T; G always pairs with C



- The sequence of bases is the genetic code
 - Variation in base sequences gives life diversity

Structure of DNA



Watson and Crick with their model

> 2-nanometer diameter— 0.34 nanometer between each—[base pair

3.4-nanometerlength of each full twist of the double helix

CH-

http://www.youtube.com/v/qy8dk5iS1f0

The numbers indicate the carbon of the ribose sugars (compare Figure 13.4). The 3' carbon of each sugar is joined by the phosphate group to the 5' carbon of the next sugar. These links form each strand's sugar-phosphate backbone.

The two sugar-phosphate backbones run in parallel but opposite directions (green arrows). Think of one strand as upside down compared with the other.

Key Concepts **Discovery of DNA's Structure**

- A DNA molecule consists of two long chains of nucleotides coiled into a double helix
- Four kinds of nucleotides make up the chains, which are held together along their length by hydrogen bonds

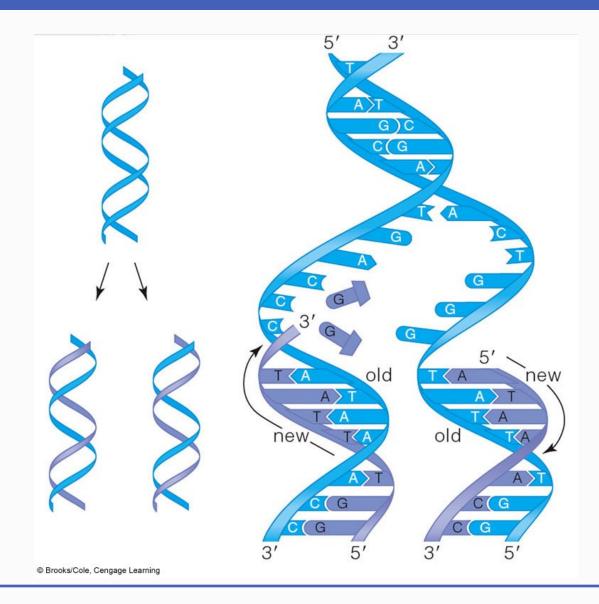
DNA Replication and Repair

- A cell copies its DNA before mitosis or meiosis I
- DNA repair mechanisms and proofreading correct most replication errors

Semiconservative DNA Replication

- Each strand of a DNA double helix is a template for synthesis of a complementary strand of DNA
- One template builds DNA continuously; the other builds DNA discontinuously, in segments
- Each new DNA molecule consist of one old strand and one new strand

Semiconservative Replication of DNA



Enzymes of DNA Replication

DNA helicase

Breaks hydrogen bonds between DNA strands

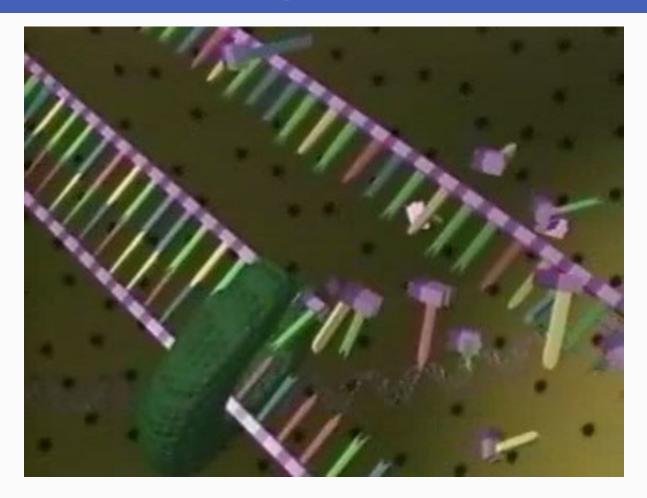
DNA polymerase

Joins free nucleotides into a new strand of DNA

DNA ligase

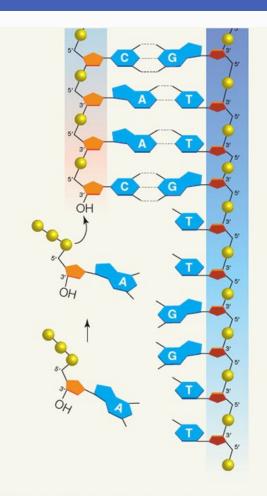
Joins DNA segments on discontinuous strand

Animation: DNA replication

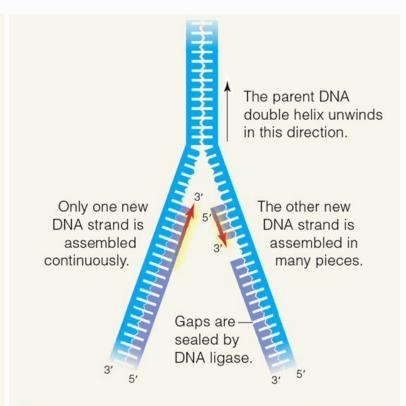


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Discontinuous Synthesis of DNA



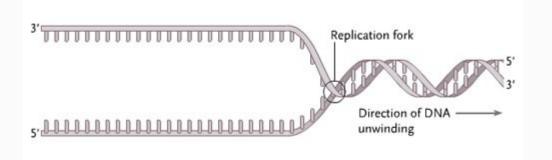
A Each DNA strand has two ends: one with a 5' carbon, and one with a 3' carbon. DNA polymerase can add nucleotides only at the 3' carbon. In other words, DNA synthesis proceeds only in the 5' to 3' direction. © Brooks/Cole, Cengage Learning



B Because DNA synthesis proceeds only in the 5' to 3' direction, only one of the two new DNA strands can be assembled in a single piece.

The other new DNA strand forms in short segments, which are called Okazaki fragments after the two scientists who discovered them. DNA ligase joins the fragtransfer back of the strand of the strand of the strand brock of the strand of t

Animation: Semidiscontinuous DNA replication



http://www.youtube.com/v/teV62zrm2P0

Checking for Mistakes

Because so much DNA is being replicated in the many cells of the body, there is a potential for errors to occur!

DNA repair mechanisms

- DNA polymerases proofread DNA sequences during DNA replication and repair damaged DNA
- When proofreading and repair mechanisms fail, an error becomes a mutation – a permanent change in the DNA sequence.
 - Can alter the genetic message and affect protein synthesis

Mutations

- Mutations in germ cells
 - Passed to future generations
 - Important for evolutionary change
- Mutations in somatic cells
 - Not passed to future generations but passed to all other somatic cells derived from it

Key Concepts How Cells Duplicate Their DNA

- Before a cell begins mitosis or meiosis, enzymes and other proteins replicate its chromosome(s)
- Newly forming DNA strands are monitored for errors
- Uncorrected errors may become mutations

From DNA to Protein – Transcription and Translation

The Nature of Genetic Information

- Each strand of DNA consists of a chain of four kinds of nucleotides: A, T, G and C
- The sequence of the four bases in the strand is the genetic information
- Transcription and translation are used to turn the DNA strand's base sequence into a protein

Converting a Gene to an RNA

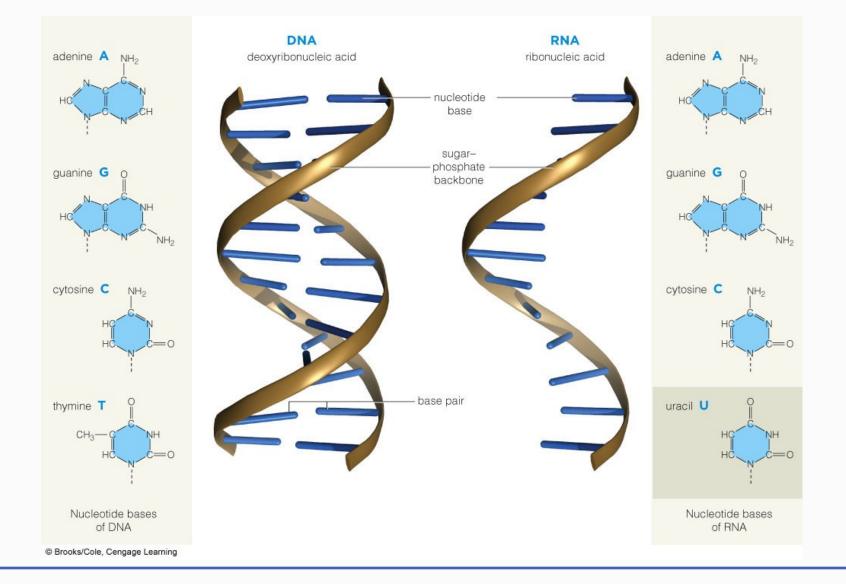
Transcription

• Enzymes use the nucleotide sequence of a gene to synthesize a complementary strand of RNA

DNA is transcribed to RNA

- Most RNA is single stranded
- RNA uses uracil in place of thymine
- RNA uses ribose in place of deoxyribose

DNA and RNA



RNA in Protein Synthesis

Messenger RNA (mRNA)

Contains information transcribed from DNA

Ribosomal RNA (rRNA)

 Main component of ribosomes, where polypeptide chains are built

Transfer RNA (tRNA)

Delivers amino acids to ribosomes

Converting mRNA to Protein

Translation

- The information carried by mRNA is decoded into a sequence of amino acids, resulting in a polypeptide chain that folds into a protein
- mRNA is translated to protein
 - rRNA and tRNA translate the sequence of base triplets in mRNA into a sequence of amino acids

Key Concepts **DNA to RNA to Protein**

- Proteins consist of polypeptide chains
- The chains are sequences of amino acids that correspond to sequences of nucleotide bases in DNA called genes
- The path leading from genes to proteins has two steps: transcription and translation

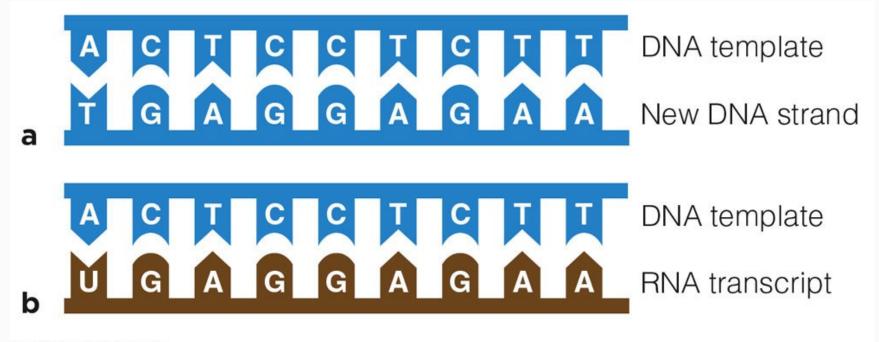
Transcription: DNA to RNA

- RNA polymerase assembles RNA by linking RNA nucleotides into a chain, in the order dictated by the base sequence of a gene
- A new RNA strand is complementary in sequence to the DNA strand from which it was transcribed

DNA Replication and Transcription

- DNA replication and transcription both synthesize new molecules by base-pairing
- In transcription, a strand of mRNA is assembled on a DNA template using RNA nucleotides
 - Uracil (U) nucleotides pair with A nucleotides
 - RNA polymerase adds nucleotides to the transcript

Base-Pairing in **DNA Synthesis and Transcription**

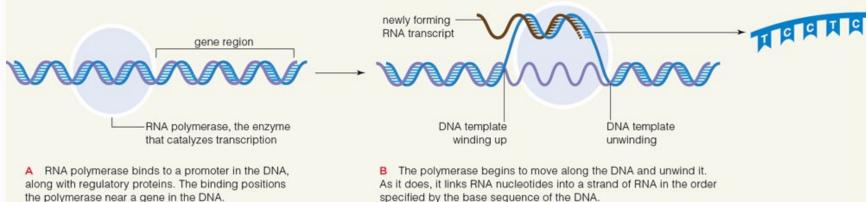


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The Process of Transcription

- RNA polymerase and regulatory proteins attach to a promoter (a specific binding site in DNA close to the start of a gene)
- RNA polymerase moves over the gene in a 5' to 3' direction, unwinds the DNA helix, reads the base sequence, and joins free RNA nucleotides into a complementary strand of mRNA

Transcription



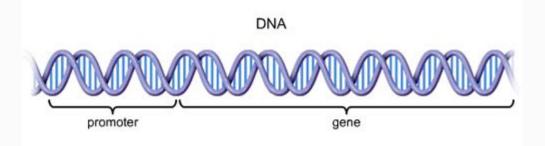
In most cases, the nucleotide sequence of the gene occurs on only one of the two strands of DNA. Only the complementary strand will be translated into RNA.

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specified by the base sequence of the DNA.

The DNA double helix winds up again after the polymerase passes. The structure of the "opened" DNA molecule at the transcription site is called a transcription bubble, after its appearance.

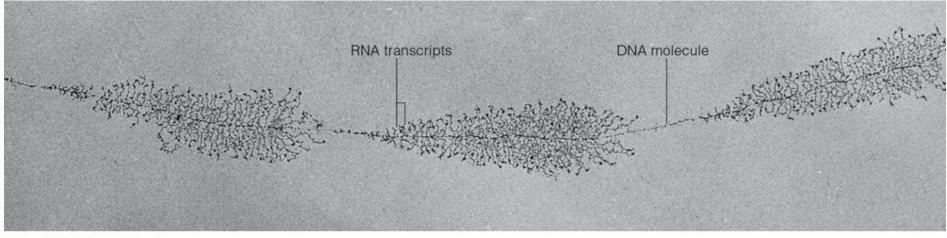
Animation: Gene transcription details



http://www.youtube.com/v/vJSmZ3DsntU

Transcription

 Many RNA polymerases can transcribe a gene at the same time



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Animation: Transcription



http://www.youtube.com/v/OtYz_3rkvPk

Key Concepts DNA to RNA: Transcription

- During transcription, one strand of a DNA double helix is a template for assembling a single, complementary strand of RNA (a transcript)
- Each transcript is an RNA copy of a gene

RNA and the Genetic Code

- Base triplets in an mRNA are words in a proteinbuilding message
- Two other classes of RNA (rRNA and tRNA) translate those words into a polypeptide chain

Post-Transcriptional Modifications

 In eukaryotes, RNA is modified before it leaves the nucleus as a mature mRNA

Introns

 Nucleotide sequences that are removed from a new RNA

Exons

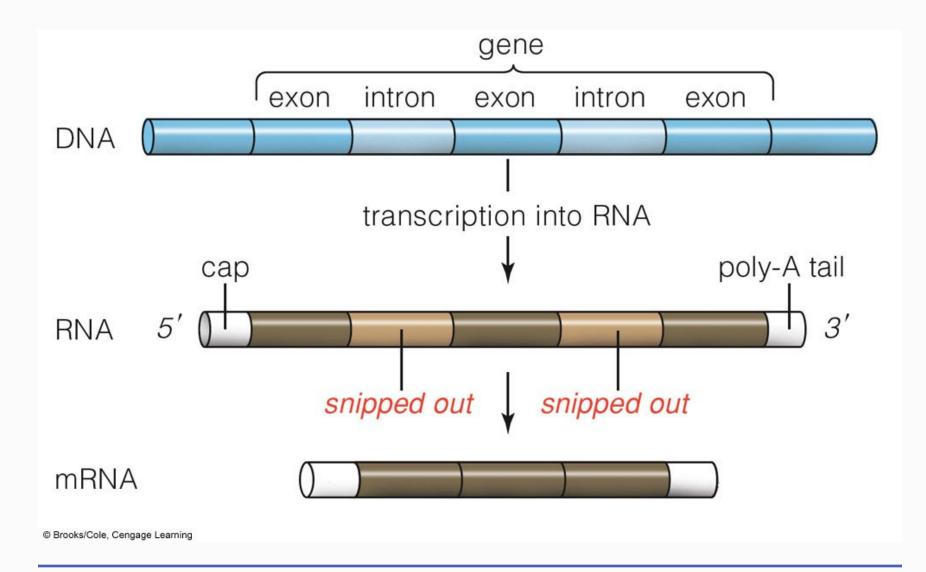
Sequences that stay in the RNA

Alternative Splicing

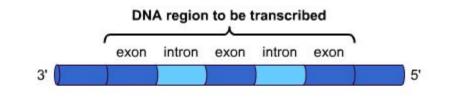
Alternative splicing

- Allows one gene to encode different proteins
- Some exons are removed from RNA and others are spliced together in various combinations

Post-Transcriptional Modifications



Animation: Pre-mRNA transcript processing





http://www.youtube.com/v/BpL4dgVGnCk

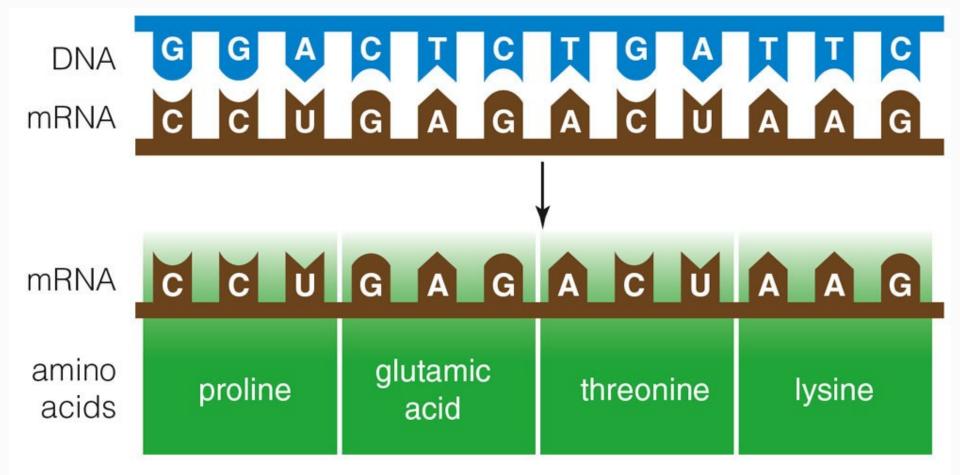
 mRNA carries protein-building information to ribosomes and tRNA for translation

Codon

- A sequence of three mRNA nucleotides that codes for a specific amino acid
- The order of codons in mRNA determines the order of amino acids in a polypeptide chain

Genetic Information

From DNA to mRNA to amino acid sequence



Genetic Code

Genetic code

- Consists of 64 mRNA codons (triplets)
- Some amino acids can be coded by more than one codon
- Some codons signal the start or end of a gene
 - AUG (methionine) is a start codon
 - UAA, UAG, and UGA are stop codons

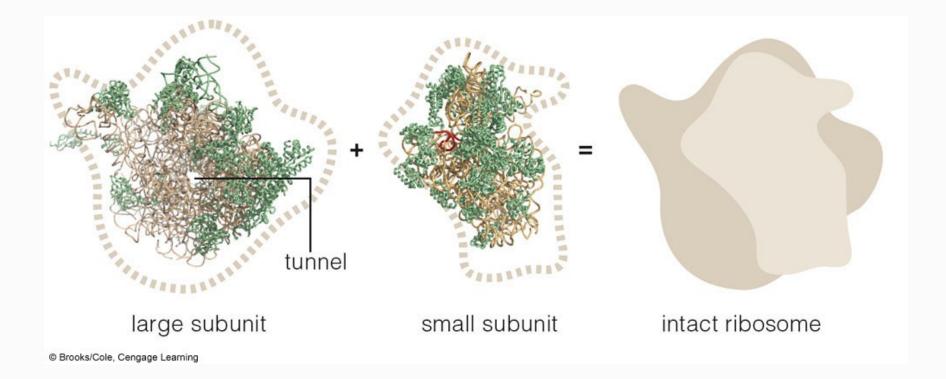
The Genetic code

		U	С	Α	G
Base Pool	2.3	phenylalanine	serine	tyrosine	cysteine
	11	phenylalanine	serine	tyrosine	cysteine
UCAG	0	leucine	serine	stop	stop
		leucine	serine	stop	tryptophan
	6	leucine	proline	histidine	arginine
Resulting Codon	С	leucine	proline	histidine	arginine
		leucine	proline	glutamine	arginine
		leucine	proline	glutamine	arginine
1st 2nd 3rd		isoleucine	threonine	asparagine	serine
1st 2nd 3rd Base Base Base	Α	isoleucine	threonine	asparagine	serine
2400 2400 2400	A	isoleucine	threonine	lysine	arginine
		methionine (start)	threonine	lysine	arginine
Clean Chart		valine	alanine	aspartate	glycine
Clear Chart	G	valine	alanine	aspartate	glycine
	G	valine	alanine	glutamate	glycine
	2 2	valine	alanine	glutamate	glycine

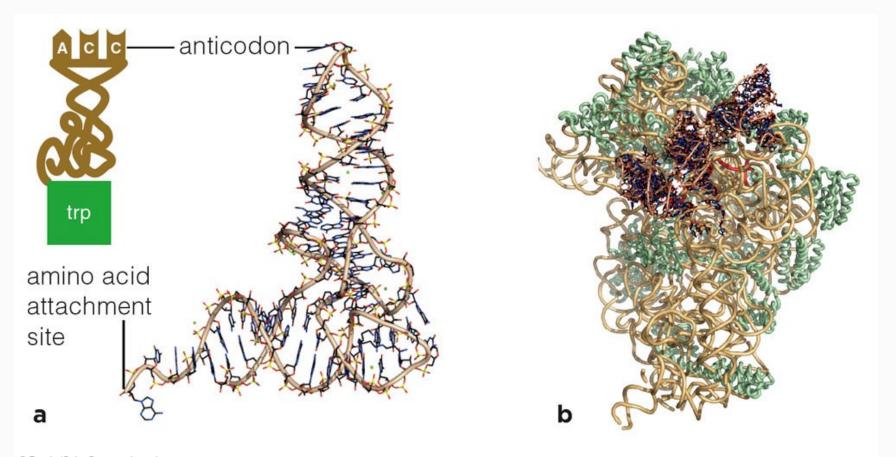
rRNA and tRNA – The Translators

- tRNAs deliver amino acids to ribosomes
 - tRNA has an anticodon complementary to an mRNA codon, and a binding site for the amino acid specified by that codon
- Ribosomes, which link amino acids into polypeptide chains, consist of two subunits of rRNA and proteins

Ribosomes



tRNA



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Key Concepts RNA

- Messenger RNA carries DNA's protein-building instructions
- Its nucleotide sequence is read three bases at a time
- Sixty-four mRNA base triplets—codons represent the genetic code
- Two other types of RNA interact with mRNA during translation of that code

Translation: RNA to Protein

- Translation converts genetic information carried by an mRNA into a new polypeptide chain
- The order of the codons in the mRNA determines the order of the amino acids in the polypeptide chain



- Translation occurs in the cytoplasm of cells
- Translation occurs in three stages
 - Initiation
 - Elongation
 - Termination

Initiation

- An initiation complex is formed
 - A small ribosomal subunit binds to mRNA
 - The anticodon of initiator tRNA base-pairs with the start codon (AUG) of mRNA
 - A large ribosomal subunit joins the small ribosomal subunit

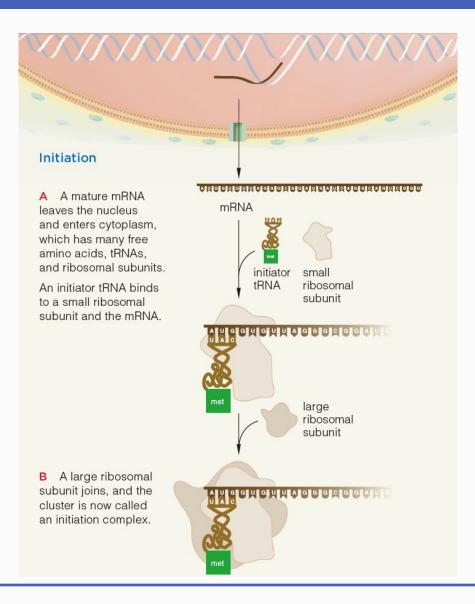
Elongation

- The ribosome assembles a polypeptide chain as it moves along the mRNA
 - Initiator tRNA carries methionine, the first amino acid of the chain
 - The ribosome joins each amino acid to the polypeptide chain with a peptide bond

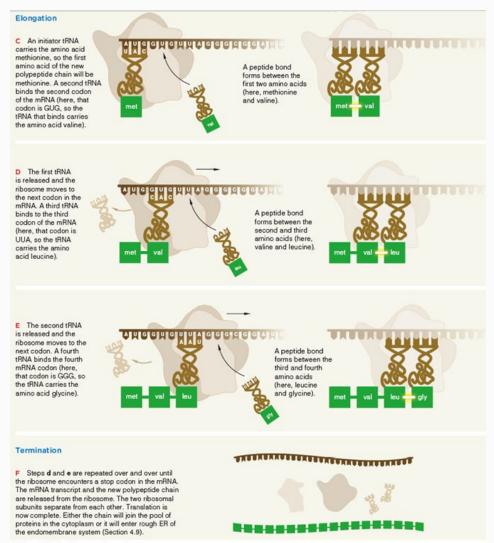
Termination

- When the ribosome encounters a stop codon, polypeptide synthesis ends
 - Release factors bind to the ribosome
 - Enzymes detach the mRNA and polypeptide chain from the ribosome

Translation in Eukaryotes

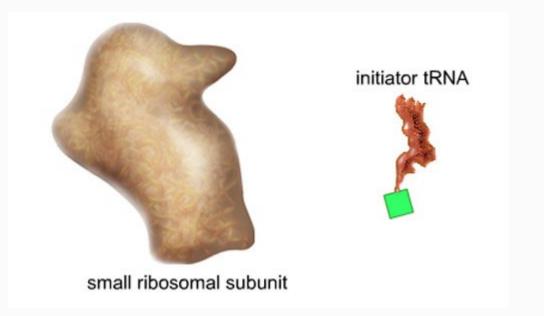


Translation in Eukaryotes



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Animation: Translation



http://www.youtube.com/watch?v=D5vH4Q_tAkY

Key Concepts RNA to Protein: Translation

 Translation is an energy-intensive process by which a sequence of codons in mRNA is converted to a sequence of amino acids in a polypeptide chain

Mutated Genes and Their Protein Products

 If the nucleotide sequence of a gene changes, it may result in an altered gene product, with harmful effects

Mutations

 Small-scale changes in the nucleotide sequence of a cell's DNA that alter the genetic code

Common Mutations

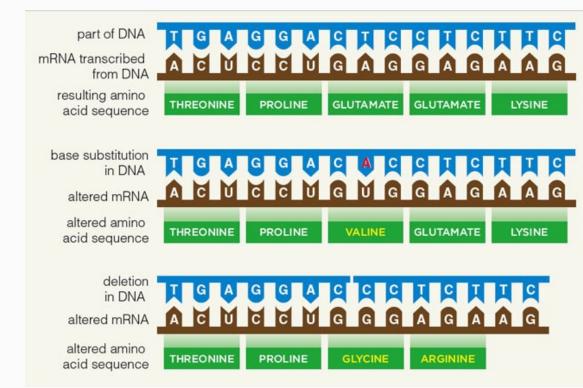
Base-pair-substitution

- May result in a premature stop codon or a different amino acid in a protein product
- *Example:* sickle-cell anemia

Deletion or insertion

- Can cause the reading frame of mRNA codons to shift, changing the genetic message
- *Example:* Huntington's disease

Common Mutations



A Part of the DNA, mRNA, and amino acid sequence of the beta chain of a normal hemoglobin molecule.

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B A base-pair substitution in DNA replaces a thymine with an adenine. When the altered mRNA is translated, valine replaces glutamate as the sixth amino acid of the new polypeptide chain. Hemoglobin with this chain is HbS—sickle hemoglobin (Section 3.6).

C Deletion of the same thymine causes a frameshift. The reading frame for the rest of the mRNA shifts, and a different protein product forms. This mutation results in a defective hemoglobin molecule. The outcome is thalassemia, a type of anemia.

http://www.youtube.com/v/kp0esidDr-c

What Causes Mutations?

Transposable elements

- Segments of DNA that can insert themselves anywhere in a chromosomes
- Spontaneous mutations
 - Uncorrected errors in DNA replication
- Harmful environmental agents
 - Ionizing radiation, UV radiation, chemicals

Key Concepts **Mutations**

- Small-scale, permanent changes in the nucleotide sequence of DNA may result from replication errors, the activity of transposable elements, or exposure to environmental hazards
- Such mutation can change a gene's product

Summary: Protein Synthesis in Eukaryotic Cells

