

DNA to Protein Supplemental Page

Read the article before checking out this detailed overview of how proteins are made from DNA!

1. **Central Dogma of Biology**: DNA is transcribed into RNA, which is then translated into a protein
2. The process by which DNA becomes a protein occurs in a sequence of several different phases and modifications. Here is a brief overview:
 - a. Transcription (DNA → mRNA) (occurs in the Nucleus)
 - i. Initiation
 - ii. Elongation
 - iii. Termination
 - b. Post-Transcriptional Processing (also occurs in the Nucleus)
 - i. 5' Capping
 - ii. 3' Polyadenylation
 - iii. RNA Splicing
 - c. Translation (mRNA → Protein) (Happens in the Ribosome)
 - i. Includes its own Initiation-Elongation-Termination Steps
 - d. Protein Folding and Post-Translational Modifications
 - i. Folding/Post-Translational Modifications

3. Step #1 → Initiation of Transcription

- a. RNA Polymerase identifies and binds to a specific region of DNA called the Promoter, located **upstream** in the DNA sequence.
 - i. **Upstream** = in the direction of the 5'
 - ii. Promoter = The region that the RNA polymerase identifies to start the process of transcription.
- b. The Promoter contains a specific DNA sequence that the RNA polymerase actually attaches to. It varies between Prok. and Euk.
 - i. In Prokaryotes, this is called the **Pribnow Box** (The exact sequence is TATAAT), just before the
 - ii. In Eukaryotes, this is called the **TATA Box**. (sequence is exactly what it sounds like)
- c. **Transcription Factors**
 - i. A complex variety of General Transcription Factors helps the RNA Polymerase bind to the Promoter

- d. **Enhancer** → DNA regions that connect to the Promoter using a Protein Structure called the *Mediator Complex* that helps send regulatory signals to the RNA Polymerase II as it attaches to the Promoter
- e. DNA begins unwinding into an **Open Complex** in order to create a transcription bubble, which exposes the *single-stranded template* for the base-pairs to start
- f. **Abortive Initiation:** RNA Polymerase, still attached to the Promoter at this point, *pulls in downstream DNA* in order to gain the energy needed to break the *grip on the promoter*
 - i. The enzyme may release smaller RNA Transcripts before it escapes the grip of the promoter
- g. **Promoter Escape** - The enzyme moves away from the promoter to begin the elongation step.
 - i. A *C-terminal domain* is attached to the RNA Polymerase, leading to a recruitment of the next set of proteins needed, and also releases initiation factors

4. Step #2 → Elongation (Transcription)

- a. **Elongation** occurs when the enzyme travels through the DNA strand to create a new RNA strand
 - i. 3' to 5' direction
 - ii. *Untwists the DNA Double helix* to form the transcription bubble as it does
- b. **Base Pairing and Nucleotide Addition:** Uracil replaces Thymine in mRNA when pairing with Adenine. Guanine and Cytosine continue to pair together
 - i. *Free RNA Nucleotides* are identified by the RNA Polymerase and used in the creation of the new Nucleotide
- c. Phosphodiester bonds are used to hold together the RNA nucleotides
- d. **RNA Polymerase** contains a proofreading mechanism that allows it to correct any errors in gene expression
 - i. RNA Molecule begins to pull away from the DNA template as the DNA transcription bubble closes up in order to keep less of the DNA exposed

5. Step #3 → Termination of Transcription

- a. In **Termination**, the RNA Polymerase Enzyme detaches from DNA and releases the RNA.
 - i. In Bacteria, a protein called **RHO** is used to help destabilize the interactions between DNA and RNA

- ii. In Eukaryotes, the transcription stops at a specific DNA sequence [AAUAAA]
 - 1. A group of proteins will then cut the pre-mRNA strand away and **free it for further processing**

6. Step #4 → Post Transcriptional Processing

- a. **Pre-mRNA** has been created and needs to undergo some changes before it can be translated into a protein
 - i. *Various modifications* are made in order to turn the mRNA into something **mature** and ready for export
- b. **5' Capping** → The 5 prime end is attached with a modified Guanine Nucleotide called a 7-methylguanylate.
 - i. Recognized by the cell in order to leave the nucleus
 - ii. Protects mRNA from degradation
 - iii. Helps small ribosomal subunits identify the mRNA in translation
 - iv. Interacts with spliceosome in later steps
- c. **3' Poly-A Tail**
 - i. 3' end is modified through the Poly-A Tail
 - 1. 50-250 Adenine Nucleotides are added to the 3' end
 - ii. Helps keep the mRNA stable and protected
- d. **RNA Splicing**
 - i. A Spliceosome made of proteins is responsible for cutting on the non-coding regions of the DNA (*Introns*) while keeping exons.
 - 1. Spliceosome identifies specific sequences at the ends of introns and breaks it down, degrading it as exons are connected together
 - ii. Some RNA Molecules, called **Ribozymes**, can do splicing completely on their own without a spliceosome.

7. Step #5 → Initiation of Translation

- a. The mRNA assembles at a specific "start" site to establish the correct reading frame for protein synthesis
- b. **Ribosomes**: Subunits remain separate until initiation begins
- c. **Initiator tRNA** → An initiator tRNA carries the Methionine (AUG, first amino acid)
 - i. Proteins coordinate the assembly of the initiation
- d. **Eukaryotic Initiation**:
 - i. The eIF4F Complex recognizes the 5' Cap added to the mRNA earlier, which recruits Ribosomes to the process

- ii. The Ribosomal's smaller (40S) subunit binds to the 5' end of the mRNA, forming what is called the **43S Complex** which begins searching through the mRNA for the start codon, AUG
- iii. Initiation Factors lead to the large unit being recruited in order to start the elongation.

8. Step #6 → Translation

- a. **Elongation Cycle** begins as the codons are now read by the newly formed ribosome and the amino acids are being brought into the ribosome to synthesize proteins
 - i. Codon Recognition → Specific Amino Acid enters the Ribosome carried by tRNA
 - ii. Peptide Bond Formation → Peptide bonds are created between amino acids
 - iii. Translocation → The Ribosome shifts further into the mRNA

9. Step #7 → Post Translational Modifications

- a. The now fully-created protein will continue to undergo even more changes after its synthesis is complete in the ribosome
- b. **Protein Folding:** The amino acid chain folds into a very specific three dimensional shape specific to the protein, going into a secondary structure and eventually a tertiary structure. In larger proteins, quaternary structure can form too.
 - i. **Chaperones:** While proteins fold spontaneously, others require molecular chaperones in order to fold
- c. Additional molecules may be added to the protein once it's done being transcribed
 - i. *Acetylation, Methylation, Phosphorylation, Etc.*
 - ii. Carbohydrates can be added to form Glycoproteins
- d. Unneeded proteins are marked for destruction by a polypeptide called Ubiquitin
- e. Modified Proteins travel to the Golgi Apparatus