



## Activity 17.1 Modeling Transcription and Translation: What Processes Produce RNA from DNA and Protein from mRNA?

Create a model of the processes of transcription and translation. Your model should be a dynamic (working or active) representation of the events that occur first in transcription in the nucleus and then in translation in the cytoplasm.

When developing and explaining your model, be sure to include definitions or descriptions of the following terms and structures:

gene	start codon (methionine)
DNA	aminoacyl-tRNA synthetase
nucleotides: A, T, G, and C versus A, U, G, and C	amino acids (see Figure 17.4, page 313, in <i>Biology</i> , 7th edition)
RNA modification(s) after transcription	peptidyl transferase
mRNA	polypeptide
RNA polymerase	energy
poly(A) tail	codons
5' cap	stop codons
translation	anticodons
protein synthesis	initiation
ribosome (large versus small subunit)	elongation
A, P, and E sites	termination
tRNA	polypeptide
rRNA	

For the purposes of this activity, assume there are *no introns* in the mRNA transcript.

## Building the Model

- Use chalk on a tabletop or a marker on a large sheet of paper to draw a cell's plasma membrane and nuclear membrane. The nucleus should have a diameter of about 12 inches.
- Draw a DNA molecule in the nucleus that contains the following DNA sequence:  
Template strand      3' TAC TTT AAA GCG ATT 5'  
Nontemplate strand    5' ATG AAA TTT CGC TAA 3'
- Use playdough or cutout pieces of paper to represent the various enzymes, ribosome subunits, amino acids, and other components.
- Use the pieces you assembled to build a dynamic (claymation-type) model of the processes of transcription and translation.
- When you feel you have developed a good working model, use it to explain the processes of transcription and translation to another student or to your instructor.

### Use your model of transcription and translation to answer the questions.

1. How would you need to modify your model to include intron removal? Your explanation should contain definitions or descriptions of the following terms and structures:

pre-mRNA  
RNA splicing  
introns

exons  
spliceosome

2. If 20% of the DNA in a guinea pig cell is adenine, what percentage is cytosine? Explain your answer.

3. A number of different types of RNA exist in prokaryotic and eukaryotic cells. List the three main types of RNA involved in transcription and translation. Answer the questions to complete the chart.

a. Types of RNA:	b. Where are they produced?	c. Where and how do they function in cells?



Assume a scientist makes three artificial mRNA strands:

(x) 5' AAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 3'

(y) 5' AAACCCAAACCCAAACCCAAACCCAAA 3'

(z) 5' AUAUAUAUAUAUAUAUAUAUAUAUAU 3'

When he analyzes the polypeptides produced, he finds that:

x produces a polypeptide composed entirely of lysine.

y produces a polypeptide that is 50% phenylalanine and 50% proline.

z produces a polypeptide that is 50% isoleucine and 50% tyrosine.

c. Do these results support the three-bases-per-codon or the four-bases-per-codon hypothesis? Explain.

d. This type of experiment was used to discover the mRNA nucleotide codons for each of the 20 amino acids. If you were doing these experiments, what sequences would you try next? Explain your logic.

6. Now that the complete genetic code has been determined, you can use the strand of DNA shown here and the codon chart in Figure 17.4 (page 313) in *Biology*, 7th edition to answer the next questions.

Original template strand of DNA: 3' TAC GCA AGC AAT ACC GAC GAA 5'

a. If this DNA strand produces an mRNA, what is the sequence of the mRNA?

b. For what sequence of amino acids does this mRNA code? (Assume it does not contain introns.)

- c. The chart lists five point mutations that may occur in the original strand of DNA. What happens to the amino acid sequence or protein produced as a result of each mutation? (*Note:* Position 1 refers to the first base at the 3' end of the transcribed strand. The last base in the DNA strand, at the 5' end, is at position 21.)

Original template strand: 3' TAC GCA AGC AAT ACC GAC GAA 5'

Mutation	Effect on amino acid sequence
i. Substitution of T for G at position 8.	
ii. Addition of T between positions 8 and 9.	
iii. Deletion of C at position 15.	
iv. Substitution of T for C at position 18.	
v. Deletion of C at position 18.	
vi. Which of the mutations produces the greatest change in the amino acid sequence of the polypeptide coded for by this 21 base pair gene?	

7. Sickle-cell disease is caused by a single base substitution in the gene for the beta subunit of hemoglobin. This base substitution changes one of the amino acids in the hemoglobin molecule from glutamic acid to valine. Look up the structures of glutamic acid (glu) and valine (val) on page 79 of *Biology*, 7th edition. What kinds of changes in protein structure might result from this substitution? Explain.

8. Why do dentists and physicians cover patients with lead aprons when they take mouth or other X-rays?