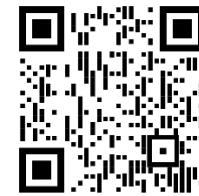
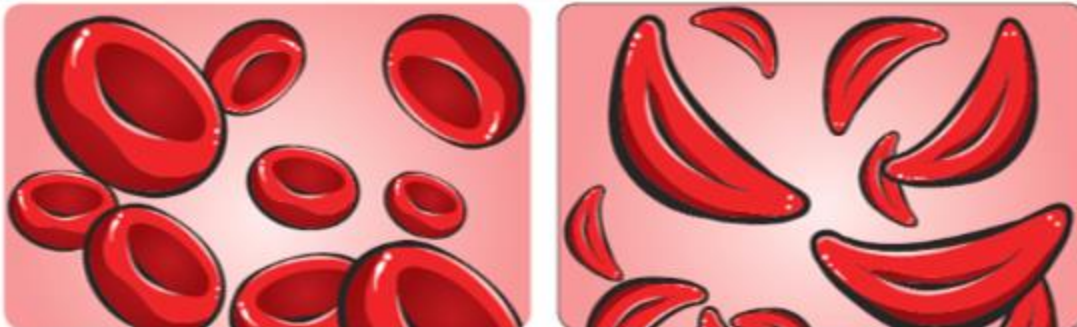


Carolina Biological Supply Company

Modeling DNA to Protein Go Hands-On with Protein Synthesis and Mutation



“How could Terra develop the genetic disease sickle-cell anemia if neither of her parents had it?”



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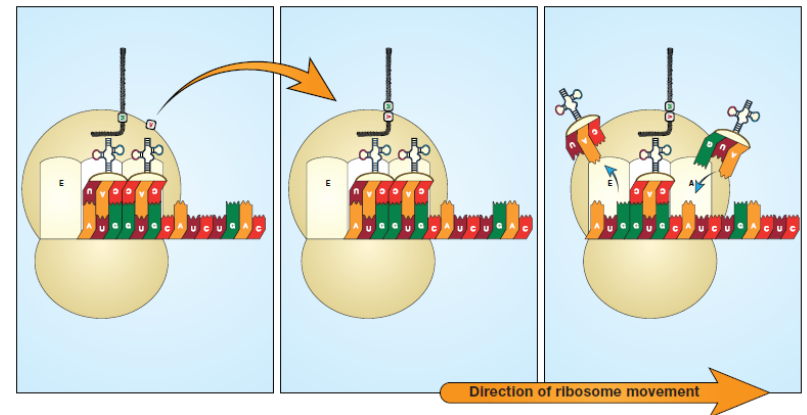
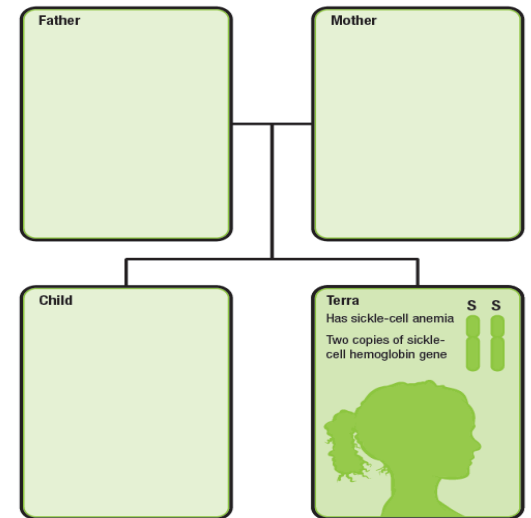
Workshop Overview

Phenomenon

“How could Terra develop the genetic disease sickle-cell anemia if neither of her parents had it?”

Objectives

- **Model** the processes of protein synthesis.
- **Construct and revise an explanation** for the mechanisms of sickle-cell disease **using models as evidence.**
- **Construct an explanation** for how DNA structure results in the production of different proteins and how a change can result in a genetic disease.



Building Toward 3-Dimensional Learning

Science and Engineering Practices	Disciplinary Core Ideas	Crosscutting Concepts
<ul style="list-style-type: none">• Developing and using models• Constructing explanations and designing solutions	LS1.A: Structure and function	<ul style="list-style-type: none">• Systems and system models• Structure and function¹


Materials

- Nucleotide Magnets
- tRNA Magnets
- Storage Cups with Lids
- Pipe Cleaner
- Red Clay
- Cornstarch or Flour
- Ribosome Transparency
- Magnetic Whiteboards
- Amino Acid 1-Letter Abbreviation Charms
- Funnel
- Ruler
- Whiteboard or Dry-Erase Markers

1. NGSS Lead States, *Next Generation Science Standards: For States, By States* (Washington, DC: The National Academies Press, 2013), retrieved from www.nextgenscience.org or ngss.nsta.org.

Time Required

Three 45-minute class periods

 Time Requirements	
Teacher Preparation (Total)	30 minutes
Previewing Videos and Interactive Lessons	10 minutes
General Preparation	10 minutes
Investigation Preparation	10 minutes
Introduction (Total)	30 minutes
Phenomenon	15 minutes
Introduction Questions	15 minutes
Investigation 1	30 minutes
Investigation 2 (Total)	50 minutes
Modeling	30 minutes
Analysis	20 minutes
Conclusion	25 minutes
Final Performance	25 minutes

To maximize the limited time of our workshop we will:

- **Split up the activities.** Work in 2 groups per table and share data between groups.
- **Discuss,** rather than complete, **some activities.**
- **Omit portions of the lab.**

Activate Background Knowledge and Reflect

1. What differences do you observe between normal and sickled red blood cells?
2. Use your prior knowledge and information from the illustrations to predict how the sickling of red blood cells could influence their functioning in Terra's body.
3. Neither Terra's older brother nor her parents suffer from sickle-cell disease, yet her medical team knows that Terra has inherited the disease from her parents. Record some questions you now have about sickle-cell anemia, the transmission of genetic diseases such as sickle-cell disease, Terra's family history, or the methods used to diagnose Terra.

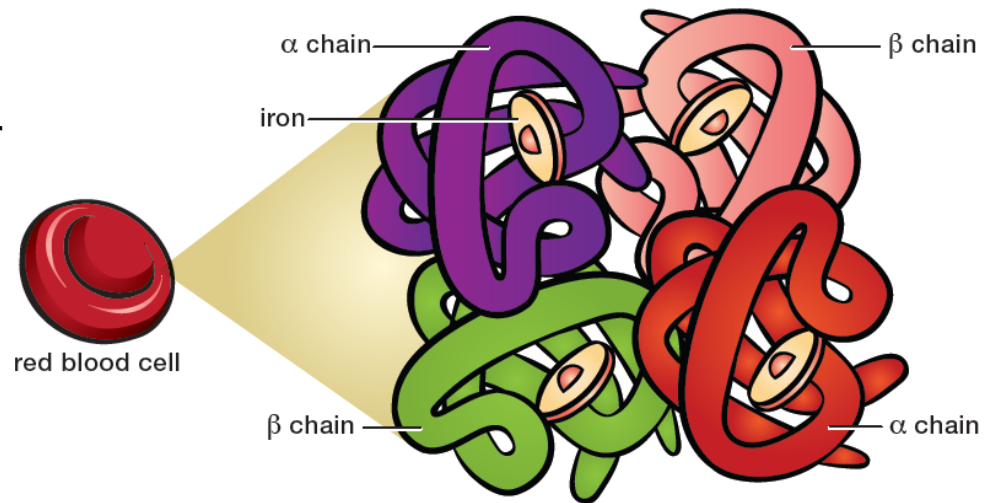
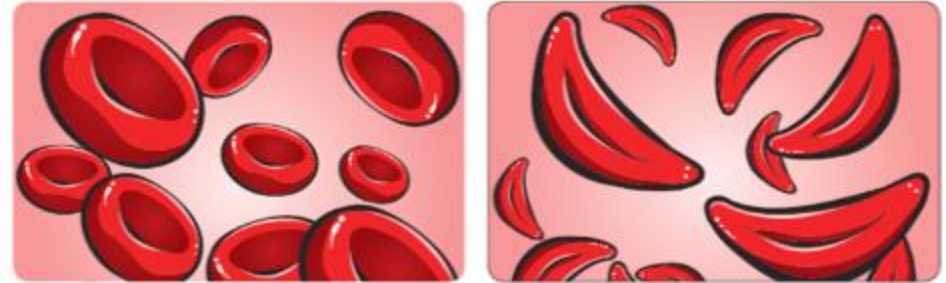


Figure 2. Normal hemoglobin protein

Procedure—Introduction

Objective: Model the process of **transcription** in the same region of the normal and the sickle-cell beta hemoglobin gene in humans to determine any differences in the resulting **mRNA sequences**.

Split each table into 2 groups of 4

Group 1: Normal Gene

TAC CAC GTA GAC TGA GGA CTC



1. Complete **steps 1-4** to transcribe the normal gene sequence.

DNA Base A – T
Pairing Rules G – C



Group 2: Mutant Gene

TAC CAC GTA GAC TGA GGA CAC



1. Complete **steps 5-8** to transcribe the normal gene sequence.

RNA Base A – U
Pairing Rules G – C

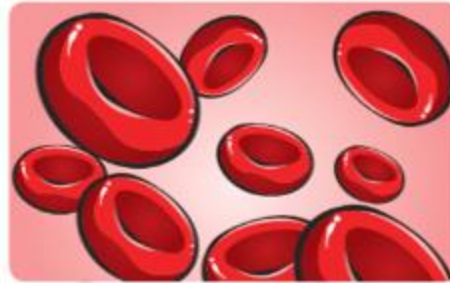


2. Share results between groups.
3. Complete introduction questions 1 and 2.

Procedure—Introduction

Objective: Model the process of **transcription** in the same region of the normal and the sickle-cell beta hemoglobin gene in humans to determine any differences in the resulting **mRNA sequences**.

1. Compare the DNA sequences in the normal and sickle-cell anemia hemoglobin genes, then make a claim about the structure of Terra's red blood cells.



2. After simulating transcription, identify any differences between the mRNA sequences in normal and sickle-cell hemoglobin. How did these differences arise?

Normal Gene

TAC CAC GTA GAC TGA GGA CTC



Mutant Gene

TAC CAC GTA GAC TGA GGA CAC



Procedure—Investigation 1

Objective: Model the process of **translation** in the same region of the normal and the sickle-cell beta hemoglobin gene in humans to determine any differences in the resulting **amino acid sequences**.

Table 1 (workshop handout page 6)

Type of Hemoglobin		1 st Codon	2 nd Codon	3 rd Codon	4 th Codon	5 th Codon	6 th Codon	7 th Codon
Normal (wild-type)	DNA	TAC	GAC	GTA	GAC	TGA	GGA	CTC
	mRNA	AUG	CUG	CAU	CUG	ACU	CCU	GAG
	Amino Acid							
Sickle-Cell (mutant)	DNA	TAC	CAC	GTA	GAC	TGA	GGA	CAC
	mRNA	AUG	GUG	CAU	CUG	ACU	CCU	GAG
	Amino Acid							

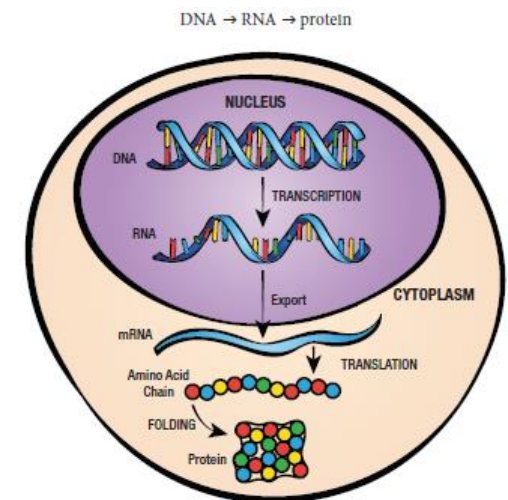


Figure 1. The central dogma of molecular biology: The flow of genetic information

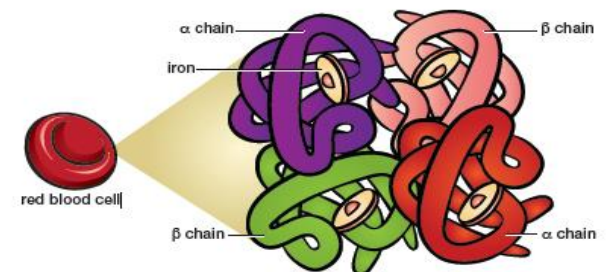
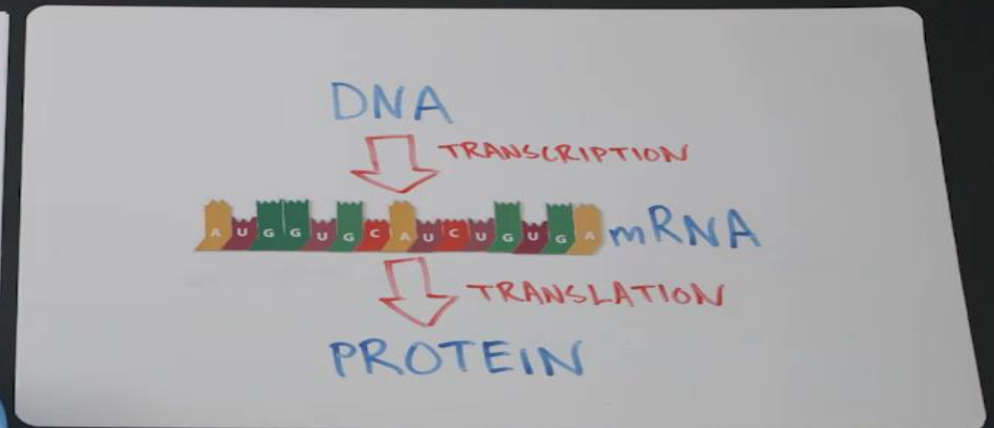
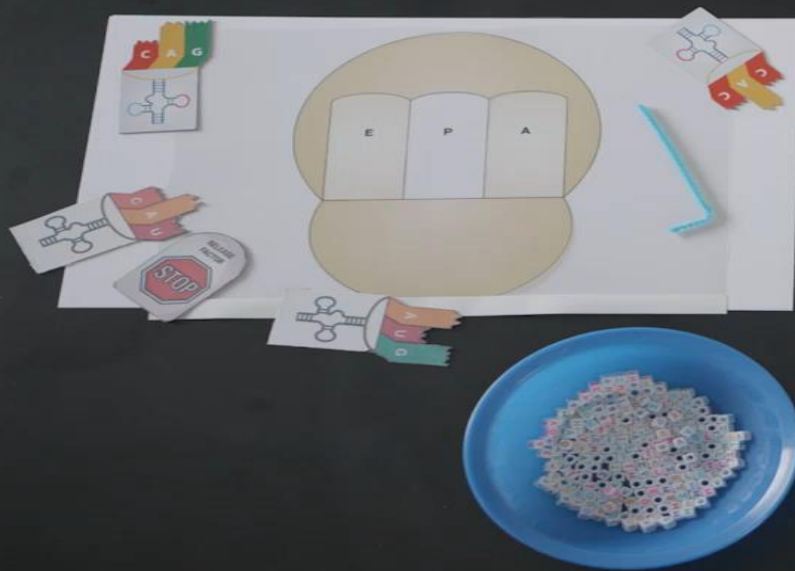


Figure 2. Normal hemoglobin protein

Procedure—Investigation 1

Objective: Model the process of **translation** in the same region of the normal and the sickle-cell beta hemoglobin gene in humans to determine any differences in the resulting **amino acid sequences**.



Procedure—Investigation 1

Objective: Model the process of **translation** in the same region of the normal and the sickle-cell beta hemoglobin gene in humans to determine any differences in the resulting **amino acid sequences**.

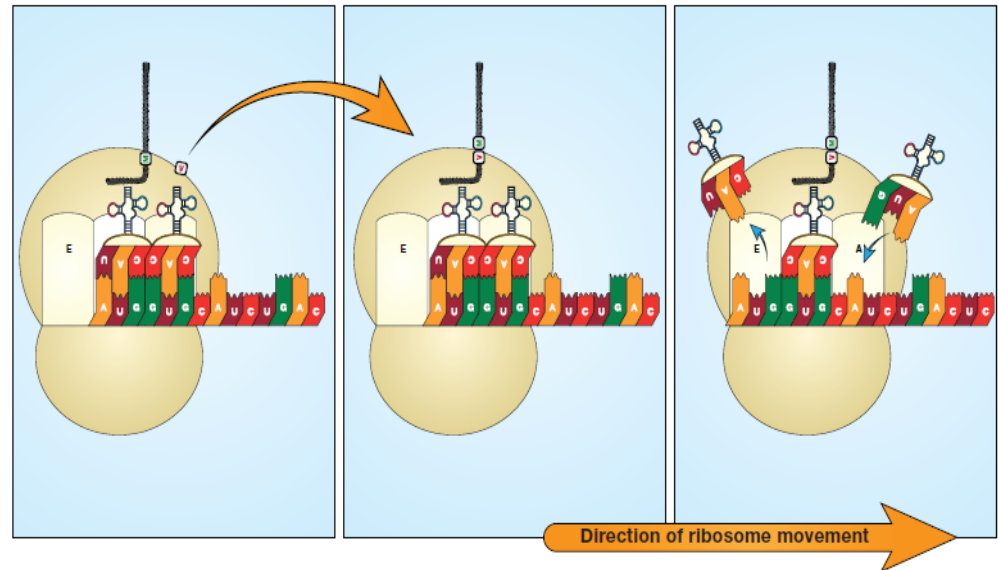
Stay in your groups

Group 1 and Group 2

1. Complete the 3 Steps of Translation (workshop handout pages 6-10, steps 2-13) using **your group's mRNA sequence**.
2. Complete steps 14-16 (workshop handout page 10) by sharing with the other group.

Helpful Tips:

- Workshop handout page 11 has a codon chart.
- Divide actions among group members, e.g.:
 - Member 1: uses codon chart to translate
 - Member 2: identifies and adds charms
 - Member 3: finds and moves tRNA
 - Member 4: moves mRNA for each step



Procedure—Investigation 2

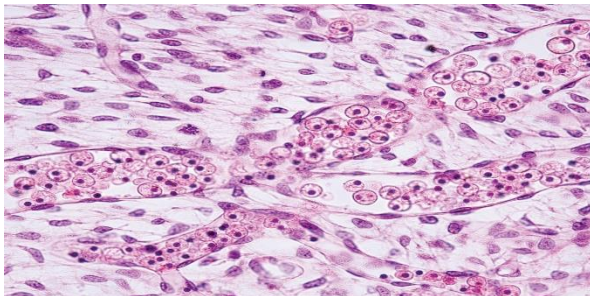
Objective: Model the effect of a change in the amino acid sequence of the hemoglobin gene **on red blood cell structure and function** when blood cells move from blood vessels into narrow capillaries.

Stay in your groups

Group 1: Normal Gene



1. Complete **steps 1-3** to model normal blood cells.



Group 2: Mutant Gene

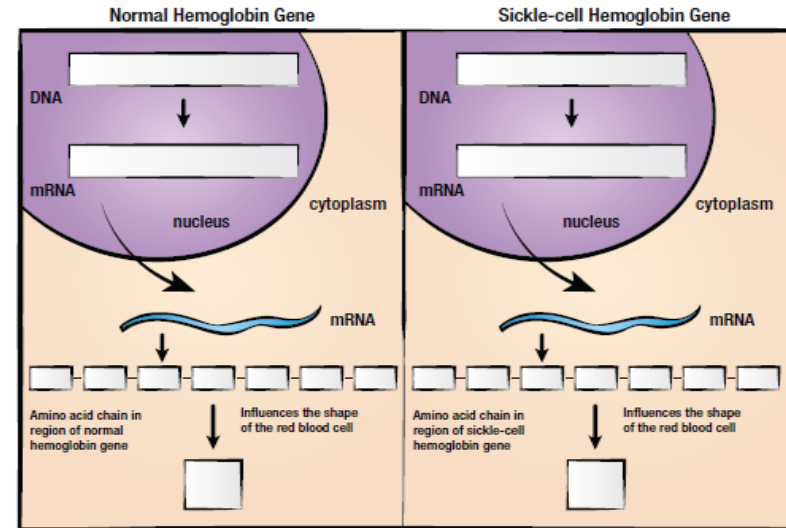


1. Complete **steps 4-6** to model sickle-cell blood cells.

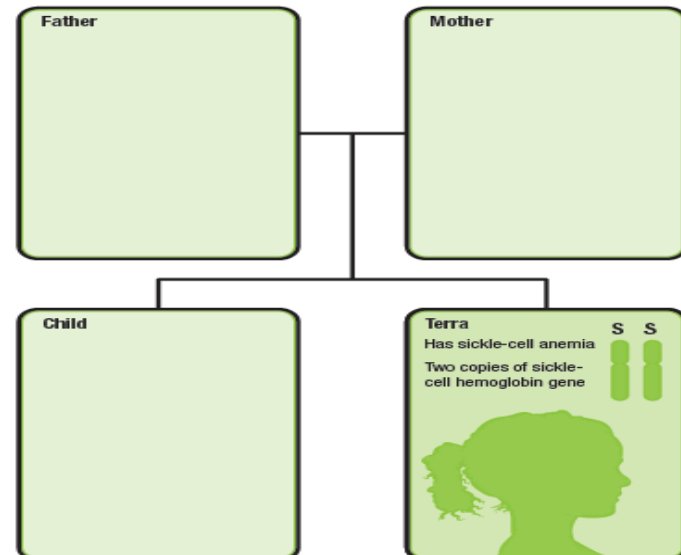
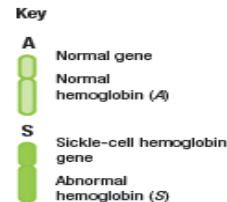
2. Share results between groups.

Analysis Examples

- This model represents the process of transcription and translation in a portion of normal and sickle-cell hemoglobin genes.
 - Fill in the blanks.
 - Label each process.
 - Revise model by drawing and labeling tRNAs, amino acids, and ribosomes.
 - Draw the shape of the resulting red blood cell represented by your clay model after each gene has been translated into normal or sickle-cell hemoglobin.



- Use the family tree template to construct a model that explains how Terra inherited sickle-cell disease, although her parents do not have it.
 - Label each family member's likely genotype (AA, AS, or SS), disease status (has sickle-cell anemia, is a carrier, or is normal), and number of each gene (normal or sickle) possessed.
 - List all likely possibilities for individuals when multiple genotypes are possible.



Resources from Carolina



Modeling DNA to Protein Kit

1-Station version: Item #211181

8-Station version: Item #211183

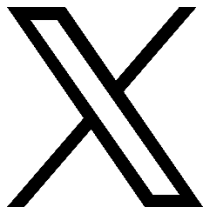
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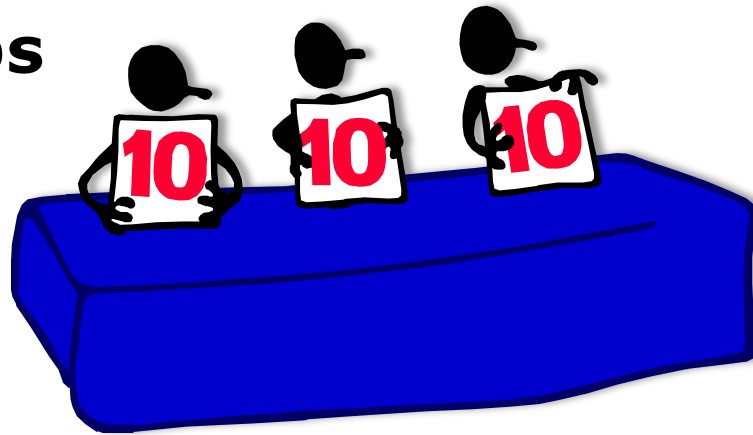
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Evaluations: Share Your Thoughts!

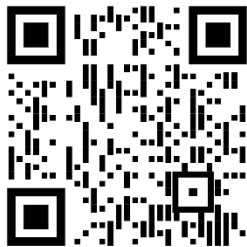
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