



# Kansas Corn: Decoding DNA - Modeling Protein Synthesis



[qrco.de/decodedna](https://qrco.de/decodedna)

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[kscorn.com](https://kscorn.com)

# Kansas Corn: Decoding DNA-Modeling Protein Synthesis

## Grade Level: Middle School / High School

### Overview

When we look at a living organism, including ourselves, we are seeing proteins or the results of proteins at work. DNA is the instructions in the cell to make these proteins, and as a result it determines nearly everything about us including our eye color, hair color, and hair texture. It determines how the cell functions by ordering the production of enzymes, receptor proteins, and other important proteins. By adding or changing a DNA sequence, we can directly influence the traits of the organism because this new DNA will be used to produce a protein that was not previously produced by the organism. Because DNA is such an important molecule, it is essential to understand how it is used to carry information and produce proteins.

In this activity, students will work in groups of three or four to model protein synthesis. Genes, sections of DNA that code for a protein, will be located in a central location in the room, which will be considered the “nucleus”. The tRNA cards will be located at the group’s workspace, which will serve as the “ribosome”. One group member will go to the nucleus and select a “gene” to transcribe. When finished, the student will take the mRNA back to the ribosome where the group will translate the mRNA strand.

### Kansas College and Career Ready Standards

#### *Disciplinary Core Ideas*

- **MS-LS1-2.** Develop and use a model to describe the function of a cell as a whole and ways parts of cells contribute to the function.
- **HS-LS1-1.** Construct an explanation based on evidence for how the structure of DNA determines the structure of proteins, which carry out the essential functions of life through systems of specialized cells.

#### *Science and Engineering Practices*

- **Developing and Using Models.** A practice of both science and engineering is to use and construct models as helpful tools for representing ideas and explanations. These tools include diagrams, drawings, physical replicas, mathematical representations, analogies, and computer simulations.

### Learning Objectives

- Students will model that DNA is the storage molecule for genetic information in the cell.
- Students will perform transcription of DNA molecules into mRNA.
- Students will translate the mRNA molecules into sentences that represent proteins.
- Students will reverse the process to code a DNA molecule that will produce a given sentence.

# Kansas Corn: Decoding DNA-Modeling Protein Synthesis

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### Materials

- Decoding DNA PowerPoint (available at [www.kscorn.com](http://www.kscorn.com))
- DNA Sentence Strips – High School (pg. S1-2, or available online at [www.kscorn.com](http://www.kscorn.com))
- DNA Sentence Strips – Middle School (pg. S3-4, or available online at [www.kscorn.com](http://www.kscorn.com))
- tRNA cards; 1 set for each group (pg. S6-9, or available online at [www.kscorn.com](http://www.kscorn.com))
- Blank strips to transcribe mRNA; 1 for each sentence to be transcribed (pg. S5, or available online at [www.kscorn.com](http://www.kscorn.com))
- Decoding DNA Student Handout (pg. S10-12 or available online at [www.kscorn.com](http://www.kscorn.com))
- Decoding DNA PowerPoint (available online at [www.kscorn.com](http://www.kscorn.com))

### Procedures for Instruction

#### *Length of Time for Classroom Teaching:*

Activity 1: Transcription – 30 minutes

Activity 2: Designer DNA – Genetic Engineering -10 minutes

#### *Preparation Procedure*

Print out the DNA Sentence Strips, tRNA cards; 1 set for each group, have blank strips to transcribe mRNA; 1 for each sentence to be transcribed. Print enough copies of the Decoding DNA Student Handout and have the Decoding DNA PowerPoint ready.

#### *Instructions*

This lab has two activities; Transcription and Designer DNA – Genetic Engineering. Hand out the Decoding DNA Student Handout (pg. S10-12, or available online at [www.kscorn.com](http://www.kscorn.com)). The handout includes the instructions for the transcription activity. In addition, the handout provides places for students to write down their sentences for both activities, and also includes reflection and conclusion, as well as assessment questions.

### Background Information

Proteins are polymers of amino acids, produced in the cell by structures called “ribosomes”. There are 20 different amino acids that are analogous to letters in the alphabet. We have only 26 letters in our alphabet, but they can be ordered and rearranged to produce thousands of words and a near-infinite number of sentences. Amino acids work similarly in making proteins, where the order and arrangement of these amino acids determines the function of the protein.

# Kansas Corn: Decoding DNA-Modeling Protein Synthesis

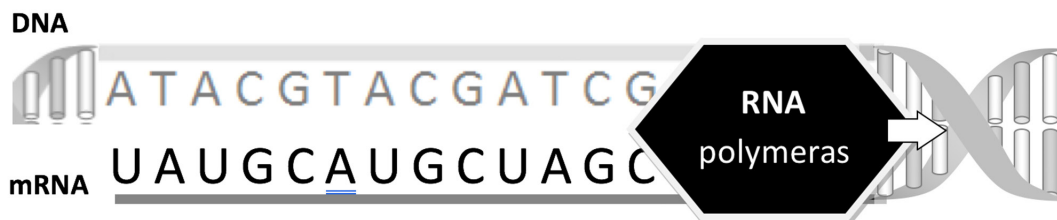
## Grade Level: Middle School / High School

DNA is a polymer of four different nucleotides: adenine (A), thymine (T), guanine (G), and cytosine (C). It is arranged in long chains that form a double helix. On each side of the helix is a complementary, matching, nucleotide. Adenine (A) always pairs with thymine (T), while cytosine (C) always pairs with guanine (G). This enables DNA to replicate, or copy itself, by “unzipping” and adding complementary nucleotides to each single strand. This allows new cells to be produced with identical copies of the DNA found in the original cell.



### *Transcription: DNA to mRNA*

In plant and animal cells, DNA stays in the nucleus. The instructions in DNA need to be copied into another molecule, RNA, to be carried to the ribosomes where they can be read to make proteins. RNA is another polymer of four nucleotides very similar to DNA. The only nucleotide that is different from those in DNA is uracil (U), which replaces thymine and complements adenine. mRNA is only a single strand that can leave the nucleus and be used by ribosomes as a recipe for making the protein. mRNA is produced in the nucleus in a process called “transcription”. The DNA strand is separated and the enzyme RNA polymerase adds complementary nucleotides to build a matching mRNA strand.



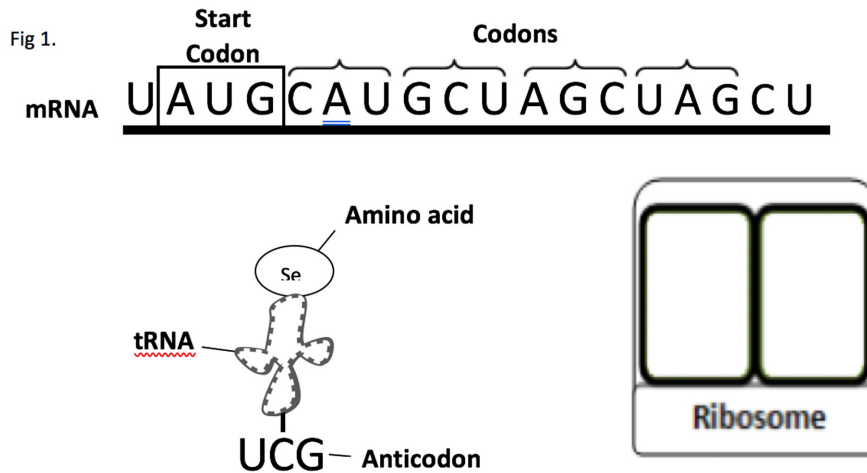
After the mRNA molecule is transcribed, it can leave the nucleus and move out into the cell where the ribosomes can translate the RNA into a protein.

# Kansas Corn: Decoding DNA-Modeling Protein Synthesis

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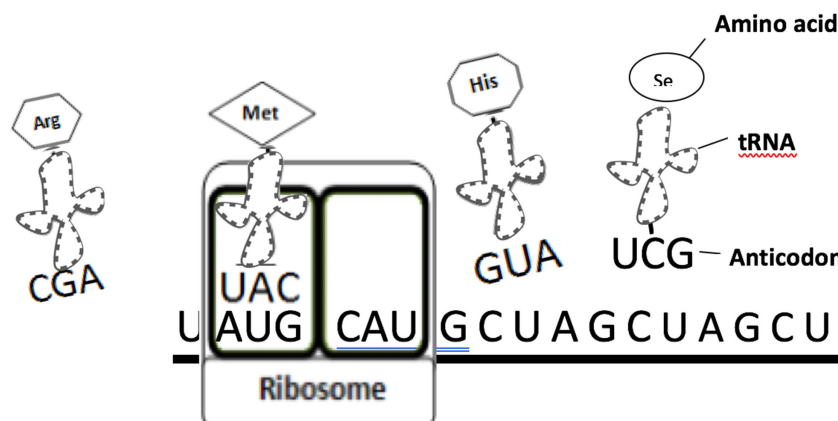
### Translation

Translation is the production of a protein from an mRNA strand by a ribosome. During translation, a ribosome uses tRNA molecules to determine the order of amino acids. The tRNA reads sections of mRNA three nucleotides at a time. These three nucleotide sections are called “codons”, and they are complemented by an anticodon on the tRNA molecule. AUG is the “Start” codon. Translation is divided into three parts: initiation, elongation, and termination.



### Initiation

A tRNA molecule attaches to the “Start” codon, AUG. This allows a ribosome to attach to the mRNA.

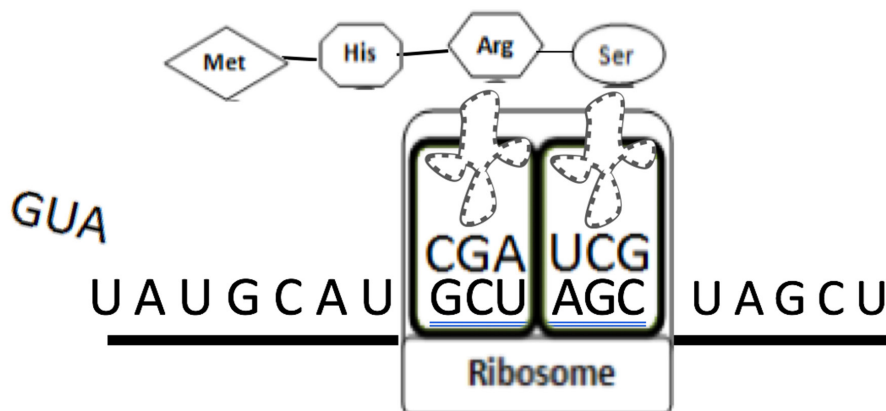
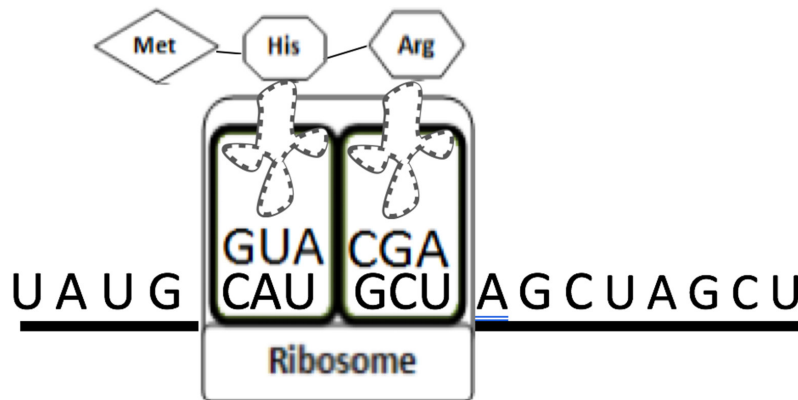
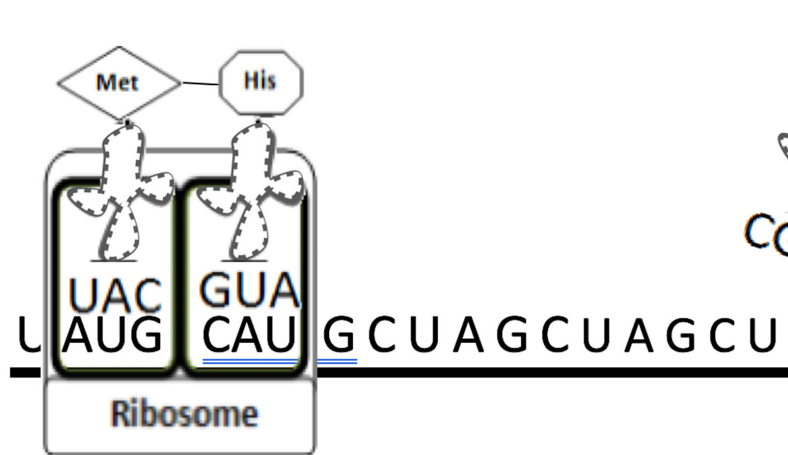


# Kansas Corn: Decoding DNA-Modeling Protein Synthesis

Grade Level: Middle School / High School

## Elongation

The ribosome continues to match anticodons to codons and add amino acids to the protein.

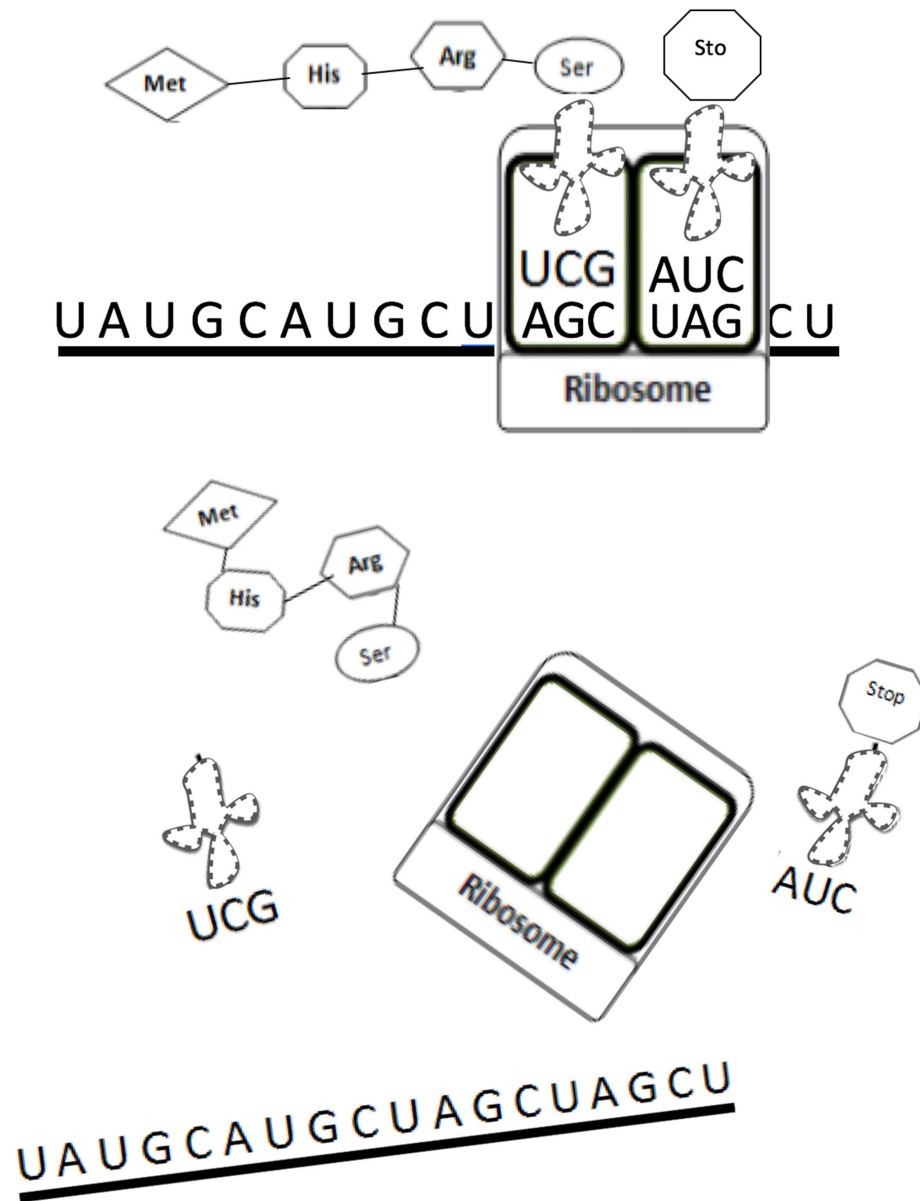


# Kansas Corn: Decoding DNA-Modeling Protein Synthesis

## Grade Level: Middle School / High School

### Termination

Occurs when a “Stop” codon is reached. The mRNA, ribosome, tRNA, and protein are all released. The protein folds into its shape and starts to work in the cell. The other components can be reused to make the protein again.



# Kansas Corn: Decoding DNA-Modeling Protein Synthesis

## Grade Level: Middle School / High School

### Classroom Discussion

Introduce the topic and assess students for prior understanding.

- Where does transcription occur? (Nucleus)
- Why do the DNA strips have to stay at the original location? (DNA can't leave the nucleus)
- What is the universal start codon? (AUG)
- Where does translation occur? (Ribosome)
- Which base pairs with A? (U)
- Which base pairs with C? (G)

### Activity 1: Transcription- 30 Minutes


#### Instructions

Hand out the Decoding DNA Student Handout (pg. S10-12, or available online at kscorn.com). The handout includes the instructions below and places to write down their sentences.

1. Number group members so that there will be an order to transcribe DNA into mRNA.
2. Group member #1 will go to the nucleus with a blank strip of mRNA strand, and choose a gene to transcribe that the group hasn't synthesized yet.
3. Write the number of the gene on the mRNA strand.
4. Transcribe the DNA strand into the complementary base pairs. For example:

C→G G→C A→U T→A

Gene #1	A T A C G T A C G A T C G A T C G A
mRNA # <u>1</u>	U A U G C A U G



5. Leave the DNA strip in the nucleus and return to the group with mRNA for translation.



# Kansas Corn: Decoding DNA-Modeling Protein Synthesis

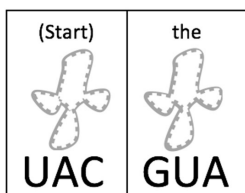
## Grade Level: Middle School / High School

### Translation Instructions

1. Once the mRNA strand has reached the group (ribosome), scan for the first “Start” codon, AUG, and highlight it. This is where you will begin translating the protein sentence.

mRNA # 1 **UAUGCAUGCUAGCUAGCU**

2. Match the complimentary anticodon from the tRNA cards to the codons and record the word on the other end of the card. This represents an amino acid in the protein that is being built. AUG (Start) is not a word to be included in the sentence, but does indicate the next word should be capitalized.



mRNA # 1 **UAUGCAUGCUAGCUAGCU**

Sentence # 1 **The** \_\_\_\_\_.



- Continue until the sentence is complete with punctuation.
- Group member #2 will move to the nucleus to transcribe another gene.
- Translate and repeat until all of the group members have transcribed a gene or completed four sentences.

Sentence # \_\_\_\_\_

Sentence # \_\_\_\_\_

Sentence # \_\_\_\_\_

Sentence # \_\_\_\_\_

# Kansas Corn: Decoding DNA-Modeling Protein Synthesis

## Grade Level: Middle School / High School

### Activity 2: Designer DNA – Genetic Engineering (5-10 minutes)

#### Overview

With the development of new technologies, such as CRISPR, DNA can be edited more easily inside living cells. This means scientists can design a protein and construct a DNA sequence that will code for that protein. These proteins are made up of the same building blocks as all other proteins, but they can introduce new traits to an organism.

#### Instructions

1. Each group will be given a sentence to code into a DNA strand.
2. Using tRNA cards, determine and record the order of anticodons that would correlate with the correct sentence. Don't forget a "start" and "stop" codon.
3. Use complimentary pairs to code for the mRNA strand that would complement the tRNA anticodons, as shown below.

tRNA → mRNA

A → U      U → A      C → G      G → C

4. Using the following base pairing, perform reverse transcription to produce a DNA strand from the mRNA strand, as shown below.

mRNA → DNA

A → T      U → A      C → G      G → C

5. When completed, insert the new gene into the nucleus.

Sentence for students to engineer: Biotechnology can improve our quality of life.

#### Teacher Resources

1. Set out the genes (DNA strands) in a central location; this will represent the nucleus.
2. Place students in groups of four; each group will synthesize four protein sentences.
3. Give each group a complete set of cards and four blank mRNA strands.

#### Reflection and Conclusion

Discuss with the students the questions on their handout. Students should understand the processes involved in producing proteins as well as how inserting engineered DNA can enable organisms to produce proteins that are new to that organism.

# Kansas Corn: Decoding DNA-Modeling Protein Synthesis

## Grade Level: Middle School / High School

### Science and Agriculture Careers

To learn more about agriculture careers visit [agexplorer.com](http://agexplorer.com). You can also find career profiles at [www.kscorn.com](http://www.kscorn.com).

### Sources

Ohio Corn and Wheat Curriculum <http://ohiocorneducation.org/>

Any educator electing to perform demonstrations is expected to follow *NSTA Minimum Safety Practices and Regulations for Demonstrations, Experiments, and Workshops*, which are available at <http://static.nsta.org/pdfs/MinimumSafetyPracticesAndRegulations.pdf>, as well as all school policies and rules and all state and federal laws, regulations, codes and professional standards. Educators are under a duty of care to make laboratories and demonstrations in and out of the classroom as safe as possible. If in doubt, do not perform the demonstrations.

# Kansas Corn: Decoding DNA-Modeling Protein Synthesis

## Grade Level: Middle School / High School

*Key: Sentences with DNA codes - High School*

1. Ethanol decreases CO<sub>2</sub> production.

TAC AGT CCG TAG TGA ATT

2. Ethanol is clean energy from corn.

TAC AGT TCC GAC ATC ATG AGG ATT

3. GMOs improve nutrition.

TAC AAA CCC AGC ATT

4. DNA is the molecule of life.

TAC TTA TCC TCG TGG TTT TAA ATT

5. Biotechnology increases yield.

TAC CGG CCC CGT ATT

6. Biotechnology decreases pesticide use.

TAC CGG CCG AAC AGA ATT

7. GMOs will meet the increasing demand for food.

TAC AAA GCG CGA TCG CCC ATA TAT CAA ATT

8. GMOs decrease poverty and increase sustainability.

TAC AAA CCG TGC GAG CCC AAG ATT

9. GMOs can grow on marginal land

TAC AAA ACG TGT TTT GCA CCT ATT

10. Some corn is drought tolerant.

TAC ACT AGG TCC CAC TTC AAT

11. One bushel of corn used for ethanol makes 17.5 pounds of distillers dried grains.

TAC CTT GAT TTT AGG AGA TAT AGT CAG TTG ACC TTT CCA ATT

12. Corn ethanol meets the demand for fuel and food.

TAC AGG AGT CGA TCG ATA TAT ATC GAG CAA ATT

13. One bushel of corn equals 56 pounds.

TAC CTT GAT TTT AGG CAG GGC ACC ATT

14. One bushel of corn makes 2.8 gallons of ethanol.

TAC CTT GAT TTT AGG CAG GGA ACC TTT AGT ATT

15. GMOs grow more food on less land.

TAC AAA TGT CCC CAA TTT CCG CCT ATT

16. Biotechnology decreases energy use.

TAC CGG CCG ATC AGA ATT

17. Biotechnology improves water and soil quality.

TAC CGG CCG TAG TGA ATT

18. Biotechnology decreases CO<sub>2</sub> production.

TAC CGG CCC GTT GAG AAT GCT ATT

19. 40% of corn is used to produce ethanol.

TAC GGT ACC TTT AGG TCC AGA CAT TGA AGT ATT

20. Dent corn produces most corn products.

TAC GAA AGG TGA ACA AGG TGA ATT

*Key for engineered DNA strand: Biotechnology can improve our quality of life.*

TAC CGG ACG CCC GTA GCT TTT TAA ATT

## DNA Sequencing Strips - High School

1 AGAGCTAGGATACAGTCCGTAGTGAATTCGCGATCGACCGG

2 GTATAGAACTACAGTTCGACATCATGAGGATTCCGAGCAG

3 GATTAGAGTTATTGTACAAACCCAGCATTGTAGCCGTAGAA

4 GTAGGTATACTTATCCTCGTGGTTTTAAATTGTAGTGACG

5 GCGAGCTTAGTACCGGCCCGTATTGCTAGCGGATCAGCTT

6 AGAGGCTGACTACCGGCCGAACAGAATTGATCGAGGTCAG

7 GAGCGTAGGTACAAACCGTGCGAGCCCAAGATTGAGCTAGG

8 GACGGATACAAAGCGCGATCGCCCATATATCAAATTGAGCGT

9 GACCAGCTTACAAAACGTGTTTTGCACCTATTGAGCGAGCTA

10 GGACGATCGTACACTAGGTCCCCTTCATTGGACGAGCTGA

11 ATACCTTGATTTTAGGAGATATAGTCAGTTGACCTTTCCAATT

12 GAGCATAACAGGAGTCGATCGATATATATCGAGCAAATTGAC

13 GACTACCTTGATTTTAGGCAGGGCACCTTTAGTATTCCT

14 GAGGACCAGGTACCTTGATTTTAGGCAGGGAACCTTTAGTATTG

15 GGACTAGGTACAAATGTCCCAATTTCCGCCTATTGAGG

16 CCCATGACGATTACCGGCCGATCAGAATTCCATCACGGA

17 GACAGCTGCGACGACTACCGGCCGTAGTGAATTGACGTTC

18 TTCATGCATACCGGCCCGTTGAGAATGCTATTTTCATAGTA

19 AGCACTACGGTACCTTTAGGTCCAGACATTGAAGTATTTCA

20 AACATAACAATACGAAAGGTGAACAAGGTGAATTGCTAT

## DNA Sequencing Strips - Middle School

1 TAC AGT CCG TAG TGA ATT

2 TAC AGT TCC GAC ATC ATG AGG ATT

3 TAC TTA TCC TCG TGG TTT TAA ATT

4 TAC AAA CCC AGC ATT

5 TAC CGG CCC CGT ATT

6 TAC CGG CCG AAC AGA ATT

7 TAC AAA CCG TGC GAG CCC AAG ATT

8 TAC AAA GCG CGA TCG CCC ATA TAT CAA ATT

9 TAC AAA ACG TGT TTT GCA CCT ATT

10 TAC ACT AGG TCC CAC TTC ATT

11 TAC CTT GAT TTT AGG AGA TAT AGT CAG TTG ACC TTT CCA ATT

12 TAC AGG AGT CGA TCG ATA TAT ATC GAG CAA ATT

13 TAC CTT GAT TTT AGG CAG GGC ACC TTT AGT ATT

14 TAC CTT GAT TTT AGG CAG GGA ACC TTT AGT ATT

15 TAC AAA TGT CCC CAA TTT CCG CCT ATT

16 TAC CGG CCG ATC AGA ATT

17 TAC CGG CCG TAG TGA ATT

18 TAC CGG CCC GTT GAG AAT GCT ATT

19 TAC GGT ACC TTT AGG TCC AGA CAT TGA AGT ATT

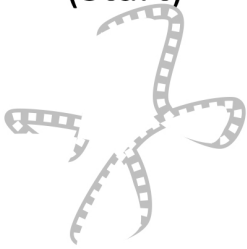
20 TAC GAA AGG TGA ACA AGG TGA ATT



## Blank Strips

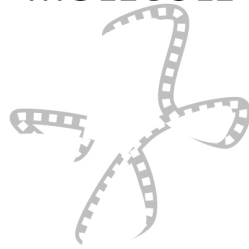
Blank writing lines for student use.

(Start)



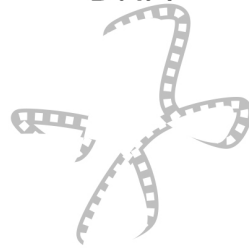
UAC

MOLECULE



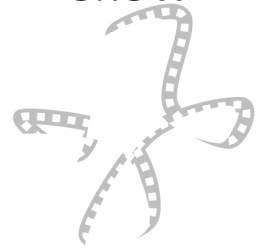
UGG

DNA



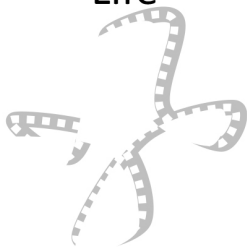
UUA

GROW



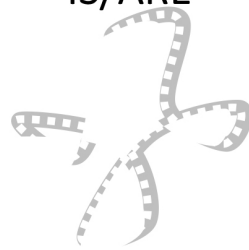
UGU

Life



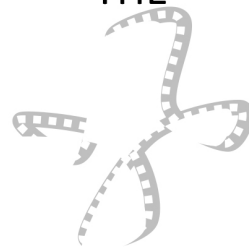
UAA

IS/ARE



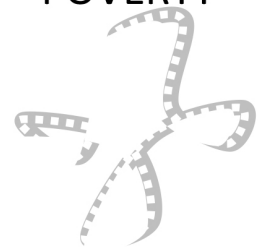
UCC

THE



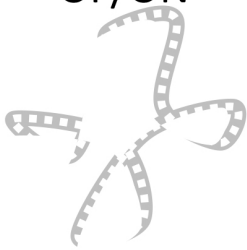
UCG

POVERTY



UGC

OF/ON



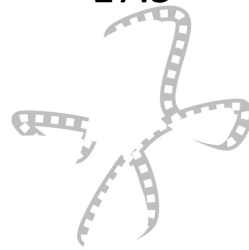
UUU

TOLERANT



UUC

17.5



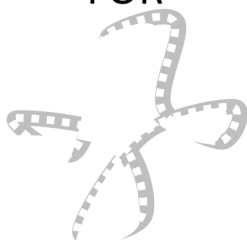
UUG

Produces/Production



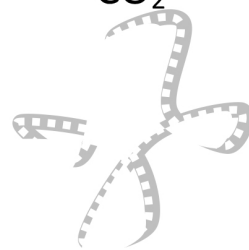
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FOR



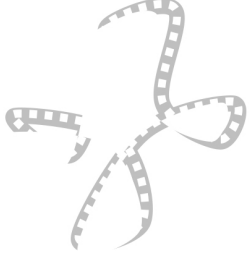
UAU

CO<sub>2</sub>



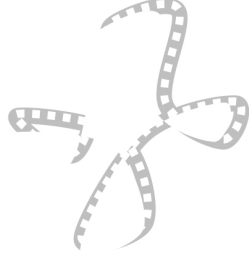
UAG

WATER



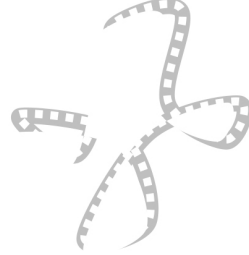
**GUU**

OUR



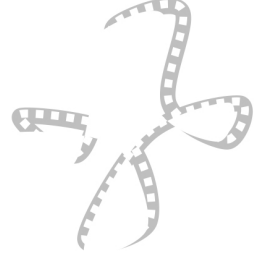
**GUA**

QUALITY



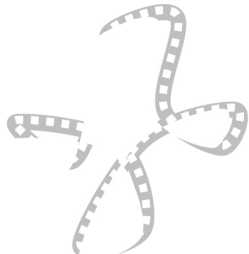
**GCU**

WILL



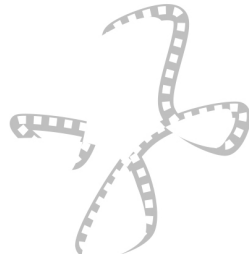
**GCG**

MARGINAL



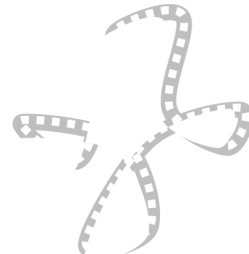
**GCA**

AND



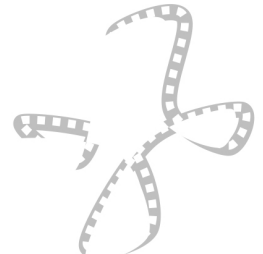
**GAG**

BUSHEL



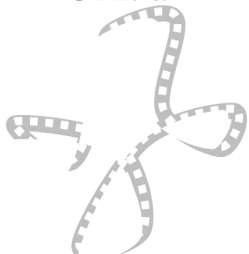
**GAU**

DENT



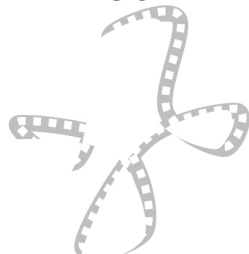
**GAA**

CLEAN



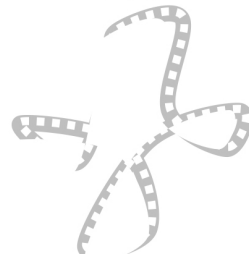
**GAC**

56



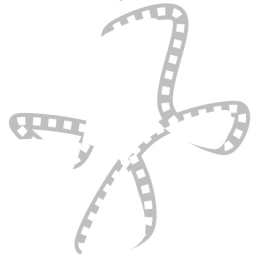
**GGC**

2.8



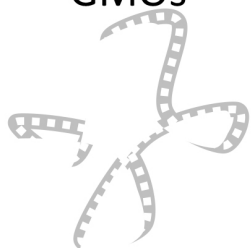
**GGA**

40

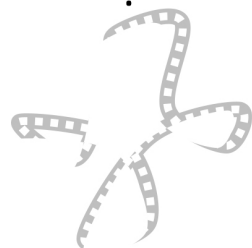


**GGU**

GMOs



AAA



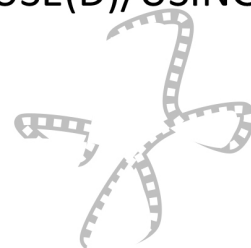
AUU

NUTRITION



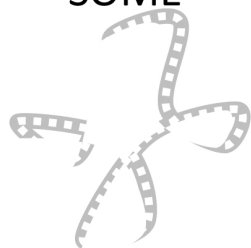
AGC

USE(D)/USING



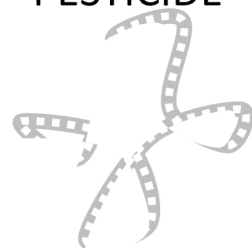
AGA

SOME



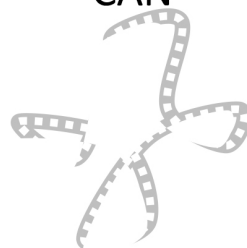
ACU

PESTICIDE



AAC

CAN



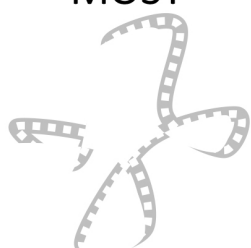
ACG

POUNDS/GALLONS  
/PERCENT



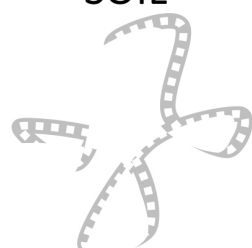
ACC

MOST



ACA

SOIL



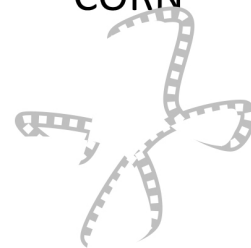
AAU

SUSTAINABILITY



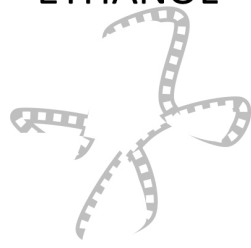
AAG

CORN



AGG

ETHANOL



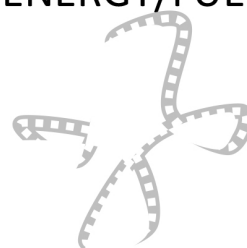
AGU

DEMAND



AUA

ENERGY/FUEL



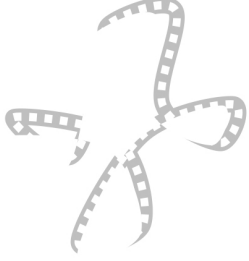
AUC

FROM



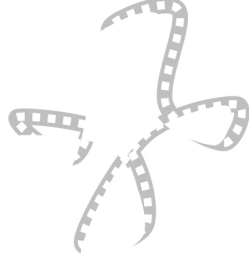
AUG

FOOD



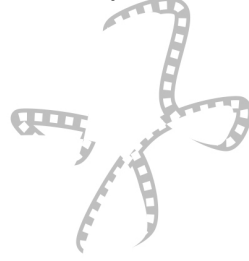
CAA

DROUGHT



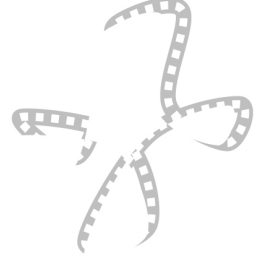
CAC

MAKES/EQUALS



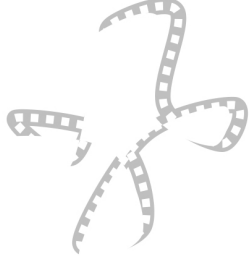
CAG

ONE



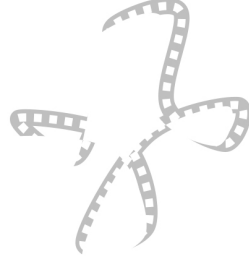
CUU

YIELD



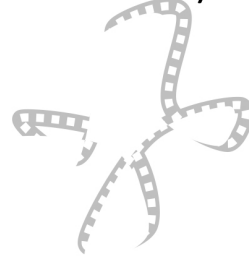
CGU

BIOTECHNOLOGY



CGG

MEET(S)



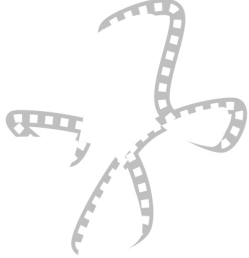
CGA

LESS/DECREASES/  
DECREASING



CCG

LAND



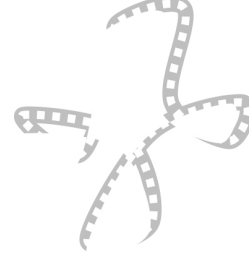
CCU

DISTILLERS  
DRIED GRAIN



CCA

TO



CAU

MORE/  
INCREASE(ING)/  
IMPROVE(ING)



CCC

## Decoding DNA: Student Handout

Name: \_\_\_\_\_


Lab Group: \_\_\_\_\_

### Transcription Instructions

1. Number group members so that there will be an order to transcribe DNA into mRNA.
2. Group member #1 will go to the nucleus with a blank strip of mRNA strand, and choose a gene to transcribe that the group hasn't synthesized yet.
3. Write the number of the gene on the mRNA strand.
4. Transcribe the DNA strand into the complementary base pairs. For example:

C→G G→C A→U T→A

Gene #1	ATACGTACGATCGATCGA
mRNA # <u>1</u>	UAUGCAUG



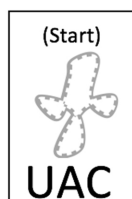
5. Leave the DNA strip in the nucleus and return to the group with mRNA for translation.

### Translation Instructions

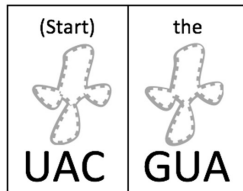
1. Once the mRNA strand has reached the group (ribosome), scan for the first start codon, AUG, and highlight it. This is where you will begin translating the protein sentence.

mRNA # <u>1</u>	UAUGCAUGCUAGCUAGCU
-----------------	--------------------

2. Match the complimentary anticodon from the tRNA cards to the codons and record the word on the other end of the card. This represents an amino acid in the protein that is being built. AUG (Start) is not a word to be included in the sentence, but does indicate the next word should be capitalized.




mRNA # <u>1</u>	UAUGCAUGCUAGCUAGCU
-----------------	--------------------



mRNA # 1 **UAUGCAUGCUAGCUAGCU**

Sentence # 1 *The* \_\_\_\_\_.



- Continue until the sentence is complete with punctuation.
- Group member #2 will move to the nucleus to transcribe another gene.
- Translate and repeat until all of the group members have transcribed a gene or completed four sentences.

Sentence # \_\_\_\_\_

Sentence # \_\_\_\_\_

Sentence # \_\_\_\_\_

Sentence # \_\_\_\_\_

Engineered DNA Strand

\_\_\_\_\_

## Reflection and Conclusion

1. How are chromosomes, DNA, genes, and proteins related?
2. What area of the cell does the table holding DNA represent in this modeling activity?
3. Why can the DNA strand not be brought back to your group?
4. What area of the cell does your table represent?
5. What do the words represent? The completed sentences?
6. What do you think the consequences might be if an error occurred in the cell as it goes through the process of protein synthesis?

## Assessment

1. Transcribe the following DNA strand. **GCTACGGACG**  
\_\_\_\_\_
2. Circle the first start codon in the mRNA strand you transcribed.
3. Write the sequence of the first codon following the start codon. \_\_\_\_\_
4. What would the anticodon be that would complement the codon in #3? \_\_\_\_\_