

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

Date: \_\_\_\_\_

Section: \_\_\_\_\_

## Epigenetics Activity

### Exploring the Central Dogma Through Interactive Modeling

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#### Phase 1: ENGAGE (5 minutes)

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**Getting Started:** Open [peebedu.com](http://peebedu.com) and navigate to Epic Genetics

Click 'Tutorial to review the interface, then Generate DNA' to begin.

**Essential Questions:** How does genetic information flow from DNA to proteins? \_\_\_\_\_ What molecular mechanisms ensure accurate gene expression? \_\_\_\_\_ How do mutations and RNA processing create protein diversity? \_\_\_\_\_

**Initial Analysis:** Examine your generated DNA sequence:

- Number of exons: \_\_\_\_\_

- Start codon (TAC in DNA → AUG in RNA) present? Yes / No

## Phase 2: EXPLORE (20 minutes)

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### Part A: Transcription Process

#### Promoter Addition:

- Select 'Promoter' and execute

- Why is this necessary? -----

#### RNA Synthesis:

- Select 'RNA Polymerase' alone and execute

- Now select both 'RNA Polymerase + Transcription Factor'

- Effect of transcription factor: -----

#### Transcript Analysis: Record the RNA sequence for one transcript:

- \_\_\_\_\_

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#### Regulatory Predictions:

If a mutation occurred in the promoter region:

- Effect on transcription: -----

- Clinical significance: -----

## Phase 5: EVALUATE (5 minutes)

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### Assessment Questions

**Process Understanding:** Explain why both 5' cap and poly-A tail are required for translation in eukaryotes. Include their molecular functions. (3 pts)

**Data Analysis:** You observe a protein that's shorter than expected. List three possible molecular explanations based on your simulation experience. (3 pts)

**Systems Integration:** Design an experiment using the simulation tools to test whether a specific exon is essential for protein function. Include:

- Tools needed: \_\_\_\_\_

- Expected results: \_\_\_\_\_

(4 pts)

### Model Evaluation:

- Most accurate representation: \_\_\_\_\_

- Missing element: \_\_\_\_\_

**Research Challenge:** Investigate one disease caused by:

- Splicing defects (e.g., -thalassemia)
- Nonsense mutations (e.g., DMD)
- Promoter mutations (e.g., some cancers)

Explain how the simulation helps understand the molecular basis: \_\_\_\_\_