

Name:

Date:

Section:

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# DNA Replication Simulator Activity: Modeling DNA Synthesis

## Understanding DNA Replication at the Molecular Level

### Phase 1: ENGAGE (5 minutes)

#### Getting Started:

Open [peebedu.com](http://peebedu.com) and navigate to DNA Replication Simulator

Read the introduction popup about DNA replication.

#### Essential Question:

How do cells accurately copy their entire genome before division? \_\_\_\_\_

#### Pre-Activity Review:

1. DNA polymerase can only add nucleotides in the \_\_\_\_' to \_\_\_\_' direction

2. Base pairing rules: A pairs with \_\_\_\_, G pairs with \_\_\_\_

#### Initial Hypothesis:

Why might replication be different on the two strands?

### Phase 2: EXPLORE (20 minutes)

#### Interactive DNA Replication Process

## Part A: Initiation

### 1. Step 1 - Topoisomerase:

- Click Topoisomerase and apply to DNA

- Why is this necessary? \_\_\_\_\_

### 1. Step 2 - Helicase:

- Apply Helicase to the relaxed DNA

- What forms at this location? \_\_\_\_\_

## Part B: Primer Addition

### 1. Step 3 - Primase:

- Apply Primase (RNA Polymerase)
- Count RNA primers added:

- Lagging strand: \_\_\_\_\_ primer(s)

## Part C: DNA Synthesis

### 1. Step 4 - DNA Polymerase:

- Apply to BOTH strands separately
- Observe synthesis direction:

- Lagging strand moves \_\_\_\_\_ from the fork

### 1. Interactive Synthesis:

- Drag correct nucleotides to match template
- Record any errors and corrections:

## Part D: Completion

### 1. Step 5 - DNA Ligase:

- Apply Ligase

- What happens to RNA primers? \_\_\_\_\_

## Phase 3: EXPLAIN (10 minutes)

### Analysis of Replication Mechanisms

#### 1. Key Patterns (Identify 3):

- Pattern 2: Leading strand is \_\_\_\_\_, lagging is \_\_\_\_\_

### 1. Cause-Effect Relationships:

Complete the chains:

- Antiparallel strands → Different synthesis patterns → \_\_\_\_\_ fragments

- DNA Pol can't start synthesis → Primase required → \_\_\_\_\_

### 1. Enzyme Function Summary:

- -----

Ligase

## Phase 4: ELABORATE (10 minutes)

### Applying Concepts

### Scenario Analysis:

#### 1. Mutation in Helicase Gene:

Effect on cell division: \_\_\_\_\_

#### 1. Telomere Problem:

The lagging strand can't replicate the very end of linear chromosomes.

- Why not? \_\_\_\_\_

### 1. Replication Speed:

E. coli replicates at ~1000 nucleotides/second

Humans replicate at ~50 nucleotides/second

- Why the difference? \_\_\_\_\_

### 1. Drug Target Design:

Many antibiotics target bacterial DNA replication.

Design a drug that would:

- Target: \_\_\_\_\_ (which enzyme)

- Why selective for bacteria? \_\_\_\_\_

## Phase 5: EVALUATE (5 minutes)

### Assessment Questions

1. **Process Understanding:** Explain why DNA replication is called "semiconservative" using evidence from the simulation. Include the fate of original strands. (3 pts)
1. **Pattern Application:** A new polymerase mutant can synthesize in both 5'→3' AND 3'→5' directions. How would this change replication? Would Okazaki fragments still form? (3 pts)

1. **Systems Thinking:** Connect DNA replication to:

- Mutations and variation (Unit 7): \_\_\_\_\_

(4 pts)

**Model Evaluation:**

- What aspects of replication are simplified? \_\_\_\_\_

**Research Topic:**

Investigate one DNA replication defect disease:

- Bloom syndrome
- Werner syndrome
- Cockayne syndrome

Explain which enzyme is affected and consequences: \_\_\_\_\_