



DNA and RNA Structure

IST-1.K.1

DNA, and in some cases RNA, is the primary source of heritable information.

IST-1.K.2

Genetic information is transmitted from one generation to the next through DNA or RNA—

- a. Genetic information is stored in and passed to subsequent generations through DNA molecules and, in some cases, RNA molecules.**
- b. Prokaryotic organisms typically have circular chromosomes, while eukaryotic organisms typically have multiple linear chromosomes.**



DNA and RNA Structure

IST-1.K.3

Prokaryotes and eukaryotes can contain plasmids, which are small extra chromosomal, double-stranded, circular DNA molecules.

IST-1.L.1

DNA, and sometimes RNA, exhibits specific nucleotide base pairing that is conserved

through evolution: adenine pairs with thymine or uracil (A-T or A-U) and cytosine pairs with guanine (C-G)–

- a. Purines (G and A) have a double ring structure.
- b. Pyrimidines (C, T, and U) have a single ring structure.



What is the primary source of genetic information?

- A. Carbohydrates**
- B. DNA/RNA**
- C. Lipids**
- D. Proteins**

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**What is the primary
source of genetic
information?**

B. DNA/RNA



**DNA and RNA are nucleic acids.
Nucleic acids are the source of
genetic information for all
organisms.**



Prokaryotic DNA is...

- A. Double stranded and circular**
- B. Double stranded and linear**
- C. Single stranded and circular**
- D. Single stranded and linear**

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Prokaryotic DNA is...

A. Double stranded and circular



Prokaryotes have a single circular DNA strand while eukaryotes have multiple linear DNA strands.

DNA is traditionally double stranded.



Eukaryotic DNA is...

- A. Double stranded and circular**
- B. Double stranded and linear**
- C. Single stranded and circular**
- D. Single stranded and linear**

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Eukaryotic DNA is...

**B. Double stranded and
linear**



Eukaryotes have multiple linear DNA strands while prokaryotes have a single circular DNA strand.

DNA is traditionally double stranded.

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How do the number of origins of replication differ and why?

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How do the number of origins of replication differ and why?

Prokaryotic cells have ONE origin of replication, and Eukaryotic cells have MULTIPLE origins of replication. Prokaryotic cells have single circular DNA strands, so it would be more efficient to start at one location and move around the entire circle to make two DNA strands. Eukaryotic cells have multiple linear DNA strands, so they have multiple origins of replication to increase efficiency and allow for DNA to replicate quickly.

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What is a plasmid?

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6.1

What is a plasmid?



**Small, extra chromosomal,
double stranded, circular DNA
molecule.**

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Both prokaryotic and eukaryotic cells have plasmids?

A. True

B. False

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Both prokaryotic and eukaryotic cells have plasmids?

A. True



Although traditionally discussed in prokaryotes, plasmids have been found in eukaryotes.

Plasmids carry a minimum of one gene that is beneficial to the host but do not carry genetic information about the organism.

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Plasmids can provide genetic variation to prokaryotes?

- A. True**
- B. False**

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6.1

Plasmids can provide genetic variation to prokaryotes?

A. True



Plasmids are small, extra chromosomal, double stranded, circular DNA molecule. This is additional genetic information that can provide additional genetic information thus genetic variation.

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In RNA, adenine pairs with...

- A. Cytosine**
- B. Guanine**
- C. Thymine**
- D. Uracil**

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TOPIC

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**In RNA, adenine pairs
with...**

D. Uracil



**RNA has four nucleotides:
adenine, cytosine, guanine, and
uracil. Base pairing is Adenine
with Uracil and Cytosine with
Guanine.**

**Apples Under the Tree (A-U)
Cars in the Garage (C-G)**

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In DNA, adenine pairs with...

- A. Cytosine**
- B. Guanine**
- C. Thymine**
- D. Uracil**

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TOPIC

6.1

**In DNA, adenine pairs
with...**

C. Thymine



**DNA has four nucleotides:
adenine, cytosine, guanine, and
thymine. Base pairing is Adenine
with Thymine and Cytosine with
Guanine.**

Apples in the Tree (A-T)

Cars in the Garage (C-G)

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**In RNA or DNA, cytosine pairs
with...**

- A. Adenine**
- B. Guanine**
- C. Thymine**
- D. Uracil**

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6.1

**In RNA or DNA, cytosine
pairs with...**

B. Guanine



**There are five nucleotides: adenine,
cytosine, guanine, thymine, and
uracil. Base pairing is Adenine with
Thymine or Uracil and Cytosine with
Guanine.**

**Apples in the Tree (A-T)
Cars in the Garage (C-G)**

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What is a purine?

- A. Carbohydrate with a single ring**
- B. Carbohydrate with a double ring**
- C. Nitrogenous base with a single ring**
- D. Nitrogenous base with a double ring**

What is a purine?

**D. Nitrogenous base with
a double ring**



Nucleotides are made up of pentose sugar, nitrogenous base, and phosphate group. There are two types of nitrogenous bases: purine (double-ring) and pyrimidine (single-ring). Adenine & Guanine are Purines and Cytosine, Uracil, and Thymine are Pyrimidines.



Which bases are purines?

- A. Adenine and Cytosine**
- B. Adenine and Guanine**
- C. Cytosine and Guanine**
- D. Cytosine and Thymine**

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6.1

Which bases are purines?

B. Adenine and Guanine



Nucleotides are made up of pentose sugar, nitrogenous base, and phosphate group. There are two types of nitrogenous bases: purine (double-ring) and pyrimidine (single-ring). Adenine & Guanine are Purines and Cytosine, Uracil, and Thymine are Pyrimidines.

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What are pyrimidines?

- A. Carbohydrate with a single ring**
- B. Carbohydrate with a double ring**
- C. Nitrogenous base with a single ring**
- D. Nitrogenous base with a double ring**

What are pyrimidines?

**C. Nitrogenous base with
a single ring**



Nucleotides are made up of pentose sugar, nitrogenous base, and phosphate group. There are two types of nitrogenous bases: purine (double-ring) and pyrimidine (single-ring). Adenine & Guanine are Purines and Cytosine, Uracil, and Thymine are Pyrimidines.



Which bases are pyrimidines?

- A. Adenine and Cytosine**
- B. Adenine and Guanine**
- C. Cytosine and Guanine**
- D. Cytosine and Thymine**

**Which bases are
pyrimidines?**

D. Cytosine and Thymine



Nucleotides are made up of pentose sugar, nitrogenous base, and phosphate group. There are two types of nitrogenous bases: purine (double-ring) and pyrimidine (single-ring). Adenine & Guanine are Purines and Cytosine, Uracil, and Thymine are Pyrimidines.



Contrast shape of DNA between prokaryotes and eukaryotes.

A. Eukaryotes have circular and prokaryotes have linear DNA

B. Prokaryotes have circular and eukaryotes have linear DNA

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6.1

Contrast shape of DNA between prokaryotes and eukaryotes.

B. Prokaryotes have circular and eukaryotes have linear DNA



The DNA structure differs between prokaryotes and eukaryotes. Prokaryotes have a single circular DNA strand while eukaryotes have multiple linear DNA strands. Prokaryotes store theirs in the nucleoid while eukaryotes store theirs in the nucleus.

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Contrast amount of DNA between prokaryotes and eukaryotes.

A. Eukaryotes have multiple DNA strands while prokaryotes have one DNA strand

B. Prokaryotes have multiple DNA strands while eukaryotes have one DNA strand

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6.1

Contrast amount of DNA between prokaryotes and eukaryotes.

A. Eukaryotes have multiple DNA strands while prokaryotes have one DNA strand



The DNA structure differs between prokaryotes and eukaryotes. Prokaryotes have a single circular DNA strand while eukaryotes have multiple linear DNA strands. Prokaryotes store theirs in the nucleoid while eukaryotes store theirs in the nucleus.



Which organism has plasmids?

- A. Bacteria (prokaryotes)**
- B. Yeast (eukaryotes)**
- C. Both bacteria and yeast**

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6.1

Which organism has plasmids?

C. Both bacteria and yeast



Both prokaryotes and eukaryotes have plasmids (small, extra chromosomal, double stranded, circular DNA molecule). It has been found in fungi, yeast, and some higher animals.

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DNA is more stable than RNA

- A. True**
- B. False**

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6.1

**DNA is more stable than
RNA**

A. True



DNA is double stranded while RNA is single stranded. The double stranded component provides structural stability as it is harder to break and there is another strand to compare when an error occurs in replication.



Replication

IST-1.M.1

DNA replication ensures continuity of hereditary information—

- a. DNA is synthesized in the 5' to 3' direction.**
- b. Replication is a semiconservative process—that is, one strand of DNA serves as the template for a new strand of complementary DNA.**
- c. Helicase unwinds the DNA strands.**
- d. Topoisomerase relaxes supercoiling in front of the replication fork.**



Replication

IST-1.M.1

DNA replication ensures continuity of hereditary information—

- e. DNA polymerase requires RNA primers to initiate DNA synthesis.**
- f. DNA polymerase synthesizes new strands of DNA continuously on the leading strand and discontinuously on the lagging strand.**
- g. Ligase joins the fragments on the lagging strand.**



Which direction is DNA read for DNA replication?

A. 3' → 5'

B. 5' → 3'

C. C terminus to N terminus

D. N terminus to C terminus

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TOPIC

6.2



Which direction is DNA read for DNA replication?

A. 3' → 5'

DNA is antiparallel, so it is read 3' → 5' and synthesized 5' → 3'. Recall the 5' end is the phosphate group and the 3' end is the hydroxyl of the deoxyribose.



Which direction is DNA synthesized in DNA replication?

A. 3' → 5'

B. 5' → 3'

C. C terminus to N terminus

D. N terminus to C terminus

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6.2

Which direction is DNA synthesized in DNA replication?

B. $5' \rightarrow 3'$



DNA is antiparallel, so it is synthesized

$5' \rightarrow 3'$ and read $3' \rightarrow 5'$.

Recall the $5'$ end is the phosphate group and the $3'$ end is the hydroxyl of the deoxyribose.

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**What is found at the 3' end
and 5' ends of DNA?**

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6.2



What is found at the **3'** end and **5'** ends of DNA?

5' end: phosphate

3' end: hydroxyl

Students a couple years ago would say the DNA is synthesized in the order of "P-OH".



How is DNA replicated?

- A. Conservative**
- B. Dispersive**
- C. Lagging**
- D. Semiconservative**

How is DNA replicated?

D. Semiconservative



The parent strand of DNA separates to provide a template strand for base pairing. The two separate strands base pair to synthesize the daughter strand. Each daughter strand has one parent strand (old) and one daughter strand (new) so the original sequence is conserved.

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6.2



**What does it mean to say the
DNA replicates
semiconservative?**

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What does it mean to say the DNA replicates semiconservative?

The parent strand of DNA separates to provide a template strand for base pairing. The two separate strands base pair to synthesize the daughter strand. Each daughter strand has one parent strand (old) and one daughter strand (new) so the original sequence is conserved.

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6.2



**Describe the experiment from
Meselson & Stahl that proved
semiconservative**

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**Describe the experiment
from Meselson & Stahl
that proved
semiconservative**



They grew bacteria in the presence of nitrogen 15 (heavy nitrogen). Then put the bacteria into presence of nitrogen 14 (light nitrogen). After one round of replication, the DNA was a hybrid of heavy and light nitrogen. After two rounds of replication, there was a hybrid of heavy & light and a light strand.



**Enzymes responsible for sealing
phosphodiester linkage**

- A. Helicase**
- B. Ligase**
- C. Polymerase**
- D. Topoisomerase**

Enzymes responsible for sealing phosphodiester link

B. Ligase



DNA ligase is used to form the phosphodiester linkage between two fragments of DNA on the sugar-phosphate backbone. This will seal the Okazaki fragments together (and the fragments from different origins of replication)



**Enzymes responsible for relieve
strand of double helix**

- A. Helicase**
- B. Ligase**
- C. Polymerase**
- D. Topoisomerase**

**Enzymes responsible for
relieve strain of double
helix**

D. Topoisomerase



As helicase pulls apart the two strands of DNA (breaks the H bonds between the two strands), the strand gets supercoiled upstream. The topoisomerase will break the H bonds relieve the strain and then re-form the bonds.



**Enzymes responsible for breaking
H bonds between strands**

- A. Helicase**
- B. Ligase**
- C. Polymerase**
- D. Topoisomerase**

Enzymes responsible for breaking H bonds between strands

A. Helicase



The helicase will break the H bonds between the two strands to allow access to the nucleotides for DNA replication. Recall, enzymes tell you what they do... helicase will break the bonds in the helix.



**Enzymes responsible for
synthesis of new DNA**

- A. Helicase**
- B. Ligase**
- C. Polymerase**
- D. Topoisomerase**

Enzymes responsible for
synthesis of new DNA

C. Polymerase



DNA polymerase synthesizes the new DNA strand. It will bind to an open 3' end to add the next nucleotide to the polynucleotide strand. Recall, enzymes tell you what they do... DNA polymerase makes a DNA polymer.

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6.2



DNA polymerase can start replication independently

- A. True**
- B. False**

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DNA polymerase can start replication independently

B. False

DNA polymerase requires an open 3' end for synthesis.

Primase will synthesize an RNA primer of 5-10 RNA nucleotides before the DNA strand.

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**What does DNA Polymerase
require to initiate replication?**

What does DNA Polymerase require to initiate replication?



An open 3' end. To initiate replication, primase will make an RNA primer. This primer has the open 3' end that is required for DNA polymerase for replication. The RNA will be replaced later with DNA by another type of DNA polymerase.

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TOPIC

6.2



DNA is...

A. Antiparallel

B. Parallel

DNA is...

A. Antiparallel



The strands of DNA are antiparallel. The strands are equidistant due to a purine and pyrimidine bonding. The strands run in opposite directions. Notice in the figure, there's a 5' across from a 3'.

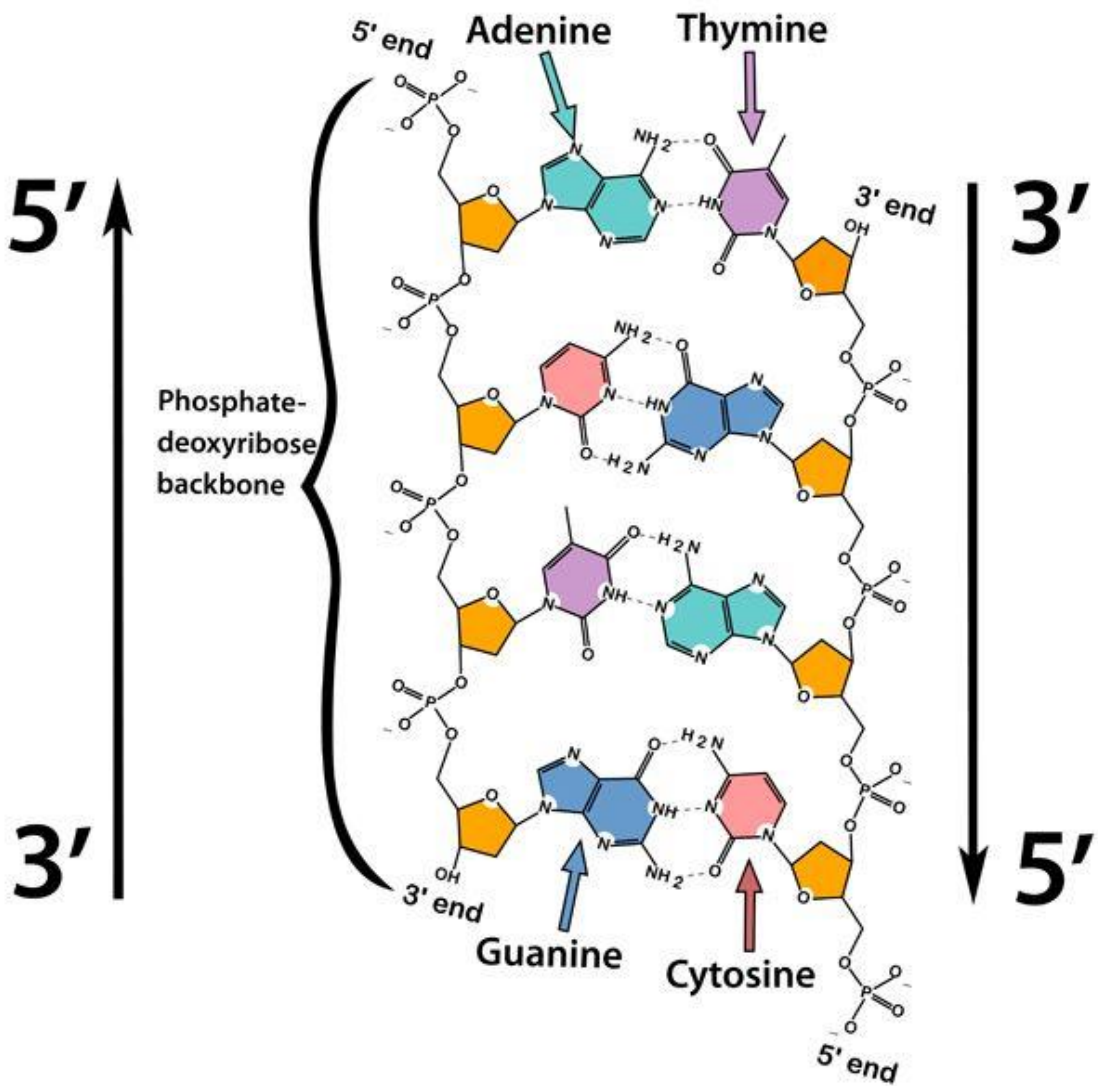
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DNA is...

A. Antiparallel



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TOPIC

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What is meant by leading and lagging strand?

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What is meant by leading and lagging strand?

Leading is synthesized continuous with replication fork with the replication fork while lagging strand is synthesized discontinuously. This occurs due to the antiparallel nature of DNA. The two strands when pulled apart only have one strand that can be read 3' to 5' (leading strand's template). Due to this, the lagging strand needs to move away from the replication fork so it can read the template strand in 3' to 5' direction.



Which enzyme connects the Okazaki fragments (lagging)

- A. Helicase**
- B. Ligase**
- C. Polymerase**
- D. Topoisomerase**

**Which enzyme connects
the Okazaki fragments
(lagging)**

B. Ligase



**DNA ligase will form the
phosphodiester linkage between
DNA fragments to form a
continuous sugar-phosphate
backbone.**

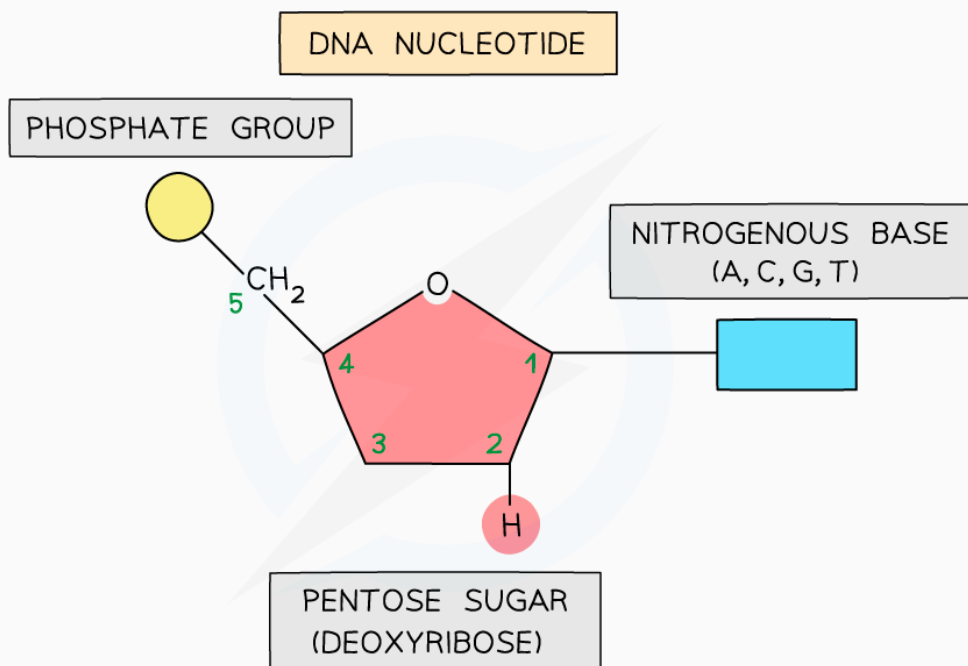


What is at the **5'** end of DNA?

- A. Hydrogen
- B. Hydroxyl
- C. Nitrogenous Base
- D. Phosphate

What is at the **5'** end of DNA?

D. Phosphate



The **5'** end has the phosphate group.
The **1'** end has the nitrogenous base.
The **3'** end has the hydroxyl group of the pentose sugar.

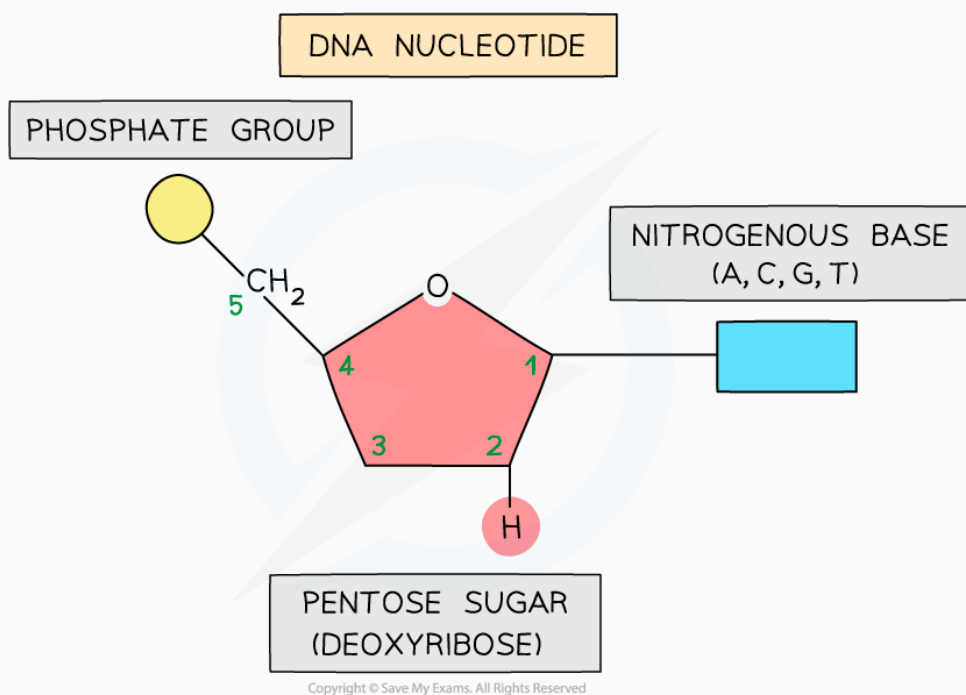


What is at the 3' end of DNA?

- A. Hydrogen**
- B. Hydroxyl**
- C. Nitrogenous Base**
- D. Phosphate**

What is at the **3'** end of DNA?

B. Hydroxyl



The **3'** end has the hydroxyl group of the pentose sugar.

The **5'** end has the phosphate group.

The **1'** end has the nitrogenous base.



**Describe the directionality of
DNA**

- A. Antiparallel**
- B. Parallel**
- C. Perpendicular**
- D. Single Stranded 5' to 3'**

**Describe the directionality
of DNA**

A. Antiparallel



The strands of DNA are antiparallel. The strands are equidistant due to a purine and pyrimidine bonding. The strands run in opposite directions.

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6.2



**What does it mean that the
DNA is antiparallel?**

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What does it mean that the DNA is antiparallel?



DNA is equidistant (recall it's a purine and pyrimidine so it has the same width). The two strands run in opposite directions.

5' → 3'

3' ← 5'



What is the enzyme responsible for breaking hydrogen bonds between nitrogenous bases?

- A. DNA polymerase**
- B. Helicase**
- C. Ligase**
- D. Primase**

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TOPIC

6.2

What is the enzyme responsible for breaking hydrogen bonds between nitrogenous bases?

B. Helicase



Helicase will break the hydrogen bond between the nitrogenous bases. Recall, the enzymes tell you what they do so helicase will separate the helix.



What enzyme is responsible for synthesizing RNA start?

- A. DNA polymerase**
- B. Helicase**
- C. Ligase**
- D. Primase**

What enzyme is responsible for synthesizing RNA start?

D. Primase



DNA polymerase requires an RNA primer to provide the open 3' end to attach the DNA nucleotides. Recall, enzymes tell you what they do so primase will make a primer.



What enzyme is responsible for making the new DNA strand?

- A. DNA polymerase**
- B. Helicase**
- C. Ligase**
- D. Primase**

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6.2



What enzyme is responsible for making the new DNA strand?

A. DNA polymerase

DNA polymerase will synthesize the new DNA strand. Recall, enzymes tell you what they do so DNA polymerase makes a DNA polymer.

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6.2



Why does DNA require an RNA start (primer)?

Why does DNA require an RNA start (primer)?



DNA polymerase requires an open 3' end to build upon for replication

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TOPIC

6.2



Which direction is DNA read?

- A. 3' to 5'**
- B. 5' to 3'**
- C. Right to Left**
- D. Left to Right**

Which direction is DNA read?

A. 3' to 5'



DNA is antiparallel, so it is read 3' \rightarrow 5' and synthesized 5' \rightarrow 3'. Recall the 5' end is the phosphate group and the 3' end is the hydroxyl of the deoxyribose.



Which direction is DNA synthesized?

- A. 3' to 5'**
- B. 5' to 3'**
- C. Right to Left**
- D. Left to Right**

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TOPIC

6.2

Which direction is DNA synthesized?

B. **5' to 3'**



DNA is antiparallel, so it is synthesized **5' -> 3'** and read **3' -> 5'**. Recall the **5'** end is the phosphate group and the **3'** end is the hydroxyl of the deoxyribose.



Which strand is formed continuously into the replication fork?

- A. Lagging strand**
- B. Leading strand**

Which strand is formed continuously into the replication fork?

B. Leading strand



The leading strand is synthesized continuously. This strand is synthesized towards the replication fork. DNA is antiparallel so moving in one direction, one strand will be in the correct orientation of 3' to 5' while the other strand will be "backwards" in the 5' to 3' orientation.



Which strand is formed discontinuously away from the replication fork?

- A. Lagging strand**
- B. Leading strand**



Which strand is formed discontinuously away from the replication fork?

A. Lagging strand

The lagging strand is synthesized discontinuously. This strand is synthesized away from the replication fork. DNA is antiparallel so moving in one direction, one strand will be in the correct orientation of 3' to 5' while the other strand will be "backwards" in the 5' to 3' orientation.

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**Lagging strand is made 3' to 5'
and leading strand is made 5'
to 3'.**

A. True

B. False

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TOPIC

6.2

Lagging strand is made 3' to 5' and leading strand is made 5' to 3'.

B. False



DNA is ALWAYS synthesized 5' to 3'. The leading strand is synthesized continuously towards the replication fork while the lagging strand is synthesized discontinuously away from the replication fork.



Transcription and RNA Processing

IST-1.N.1

The sequence of the RNA bases, together with the structure of the RNA molecule, determines RNA function—

- a. mRNA molecules carry information from DNA to the ribosome.



Transcription and RNA Processing

IST-1.N.1

- b. Distinct tRNA molecules bind specific amino acids and have anti-codon sequences that base pair with the mRNA. tRNA is recruited to the ribosome during translation to generate the primary peptide sequence based on the mRNA sequence.**
- c. rRNA molecules are functional building blocks of ribosomes.**



Transcription and RNA Processing

IST-1.N.2

Genetic information flows from a sequence of nucleotides in DNA to a sequence of bases in an mRNA molecule to a sequence of amino acids in a protein.

IST-1.N.3

RNA polymerases use a single template strand of DNA to direct the inclusion of bases in the newly formed RNA molecule. This process is known as transcription.



Transcription and RNA Processing

IST-1.N.4

The DNA strand acting as the template strand is also referred to as the noncoding strand, minus strand, or antisense strand. Selection of which DNA strand serves as the template strand depends on the gene being transcribed.

IST-1.N.5

The enzyme RNA polymerase synthesizes mRNA molecules in the 5' to 3' direction by reading the template DNA strand in the 3' to 5' direction.



Transcription and RNA Processing

IST-1.N.6

In eukaryotic cells the mRNA transcript undergoes a series of enzyme-regulated modifications—

- a. Addition of a poly-A tail.
- b. Addition of a GTP cap.
- c. Excision of introns & splicing & retention of exons.
- d. Excision of introns and splicing and retention of exons can generate different versions of the resulting mRNA molecule; this is known as alternative splicing.



Which RNA functions as site of protein synthesis?

- A. mRNA**
- B. rRNA**
- C. tRNA**

Which RNA functions as site of protein synthesis?

B. rRNA



rRNA is the ribosomal RNA. Ribosomes, the site of protein synthesis, is composed of protein and rRNA.



Which RNA functions to bring amino acids to the ribosome?

- A. mRNA**
- B. rRNA**
- C. tRNA**

Which RNA functions to bring amino acids to the ribosome?

C. tRNA



tRNA is the transfer RNA. There are two important binding sites on the tRNAs. At one end, the tRNA is bound to an amino acid and the other end has an anticodon which pairs with the codon on the mRNA.



**Which RNA is the transcript
from DNA template?**

- A. mRNA**
- B. rRNA**
- C. tRNA**

Which RNA is the transcript from DNA template?

A. mRNA



mRNA is messenger RNA. It functions as a transcript of the DNA to carry the genetic code to the ribosome for protein synthesis.



Which RNA has an anticodon that pairs with codon?

- A. mRNA**
- B. rRNA**
- C. tRNA**

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6.3

Which RNA has an anticodon that pairs with codon?

C. tRNA



tRNA is the transfer RNA. The main function of the tRNA is to bring amino acids to the ribosome. The tRNA has an anticodon that pairs with the mRNA codon to ensure the correct amino acid is added.



**Which RNA has the codons
sequence for translation?**

- A. mRNA**
- B. rRNA**
- C. tRNA**

Which RNA has the codons sequence for translation?

A. mRNA



mRNA is the messenger RNA. It is a transcript of the DNA to carry to message to the ribosome for protein synthesis. The triplet code of nucleotides on the mRNA is called a codon.

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6.3



Big picture, how do the three different RNA molecules work together?

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6.3

Big picture, how do the three different RNA molecules work together?



mRNA is the product of transcription in the nucleus. It brings the "message" from the nucleus about the order for protein synthesis.

mRNA binds to the small subunit of the ribosome.

tRNA brings the amino acids The tRNA has an anticodon at one end that base pairs with the codons on the mRNA and an amino acid on the other end. It binds to the large subunit of the ribosome.

rRNA is the site of protein synthesis. The rRNA binds with protein to form the ribosome.

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How does the location of transcription differ in prokaryotes vs. Eukaryotes?

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How does the location of transcription differ in prokaryotes vs. eukaryotes?



Due to the absence of a nuclear envelope, transcription occurs in the cytosol (nucleoid) of the prokaryote.

Due to the presence of a nuclear envelope, transcription occurs in the nucleus of an eukaryote.



**Which enzyme is responsible
for transcription?**

- A. Ligase**
- B. Primase**
- C. RNA Polymerase**
- D. Transcriptase**

**Which enzyme
is responsible
for transcription?**

C. RNA Polymerase



**RNA Polymerase is an enzyme
that binds to the DNA and
synthesizes a RNA strand.
Remember: enzymes tell you
what they do. RNA Polymerase
makes an RNA polymer.**

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6.3



**What is the name of the strand
read for transcription?**

What is the name of the strand read for transcription?



- > **Template strand**
- > **Noncoding strand**
- > **Minus strand**
- > **Antisense strand**

(Note: you should know all 4 of these terms as they are directly in the CED and can be used in the prompt.)



**In transcription, which direction
is DNA read?**

- A. 3' to 5'**
- B. 5' to 3'**
- C. C terminus to N terminus**
- D. N terminus to C terminus**

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TOPIC

6.3

In transcription, which direction is DNA read?

A. 3' to 5'



DNA and RNA are antiparallel. The DNA is read 3' to 5' while the RNA is synthesized 5' to 3'. Remember the 3' is the hydroxyl group of the pentose sugar and the 5' is the phosphate.



**In transcription, which direction
is RNA made?**

- A. 3' to 5'**
- B. 5' to 3'**
- C. C terminus to N terminus**
- D. N terminus to C terminus**

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TOPIC

6.3

In transcription, which direction is RNA made?

B. 5' to 3'



DNA and RNA are antiparallel. The RNA is synthesized 5' to 3' while the DNA is read 3' to 5'.

Remember the 3' is the hydroxyl group of the pentose sugar and the 5' is the phosphate.

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TOPIC

6.3



Describe the three changes made during post-transcription

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TOPIC

6.3



**Describe the three changes
made during post-
transcription**

- > Guanine cap added to 5' end**
- > Poly A tail added to 3' end**
- > Introns are removed by
splicing**



Function of 5' Cap...

- A. Add phosphates to the RNA**
- B. Provide ATP for transcription**
- C, Protect from hydrolytic enzymes**
- D. Site of ribosome binding**

Function of 5' Cap...

D. Site of ribosome binding



The 5' Cap has two functions. It binds with the ribosome for translation (which reads the RNA 5' to 3') at the 5' end. It facilitates the mRNA leaving the nucleus.



Function of Poly A tail...

- A. Add phosphates to the RNA**
- B. Provide ATP for transcription**
- C. Protect from hydrolytic enzymes**
- D. Site of ribosome binding**

Function of Poly A tail...

C. Protect from hydrolytic enzymes



The poly-A tail is made up of multiple adenines at the end of the mRNA. As the hydrolytic enzymes in the cytosol hydrolyze the mRNA, it prolongs its "life" in the cytosol to maximize protein synthesized.

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TOPIC

6.3



**What process allows the
synthesis of different proteins
from same DNA**

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TOPIC

6.3

What process allows the synthesis of different proteins from same DNA



Alternative RNA splicing



What is the process of transcription?

- A. Synthesizing DNA under the direction of DNA**
- B. Synthesizing DNA under the direction of RNA**
- C. Synthesizing RNA under the direction of DNA**
- D. Synthesizing RNA under the direction of RNA**

What is the process of transcription?

C. Synthesizing RNA under the direction of DNA



Transcription is the process of synthesizing a RNA molecule from an DNA template. The DNA must be protected as its the genetic code for the cell, so the RNA is a disposable copy of the information for cellular usage.

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TOPIC

6.3



What is the order in the central dogma?

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TOPIC

6.3

What is the order in the central dogma?



DNA → RNA → polypeptide



What type of virus will violate the central dogma?

- A. Bacteriophage**
- B. DNA virus**
- C. Retrovirus**
- D. All of the above**

What type of virus will violate the central dogma?

C. Retrovirus



Retroviruses have a RNA genome. They have reverse transcriptase, enzyme that synthesizes DNA from RNA template. The central dogma states DNA \rightarrow RNA, but the retrovirus goes RNA \rightarrow DNA.

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TOPIC

6.3



What is the function of mRNA, rRNA, and tRNA?

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TOPIC

6.3



What is the function of mRNA, rRNA, and tRNA?

mRNA is the messenger RNA, which brings the information about the order of amino acids to the ribosomes and base pairs the codon with the anticodon (on the tRNA)

tRNA is the transfer RNA, which brings the amino acids to the ribosome and base pairs the anticodon with the codon (on the mRNA)

rRNA is the ribosomal RNA, which makes up the ribosome in addition to proteins.



What enzyme is responsible for transcription?

- A. DNA polymerase**
- B. Helicase**
- C. Primase**
- D. RNA polymerase**

What enzyme is responsible for transcription?

D. RNA polymerase



RNA polymerase is the enzyme that functions in transcription.

It will synthesize a RNA molecule from the DNA template.



Which strand is read for transcription?

- A. 3' to 5' coding strand**
- B. 5' to 3' coding strand**
- C. 3' to 5' noncoding strand**
- D. 5' to 3' noncoding strand**

Which strand is read for transcription?

C. 3' to 5' noncoding strand



DNA and RNA are antiparallel, so the DNA is read 3' to 5' and the RNA is synthesized 5' to 3'. Note: the DNA template strand can be called 4 different names: noncoding strand, minus strand, or antisense strand.



Which direction is RNA synthesized?

- A. 3' to 5'**
- B. 5' to 3'**

AP BIO INSTA-REVIEW

TOPIC

6.3

Which direction is RNA synthesized?

B. 5' to 3'



DNA and RNA are antiparallel, so the RNA is synthesized 5' to 3' and DNA is read 3' to 5'. Recall: the 5' end is the site of the phosphate and the 3' end is the hydroxyl group of the pentose sugar.

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TOPIC

6.3



What three things happen to the mRNA before it leaves the nucleus?

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TOPIC

6.3



What three things happen to the mRNA before it leaves the nucleus?

- > **Add a 5' guanine cap**
- > **Add a poly A tail**
- > **Cut out introns**



What is the function of the poly A tail?

- A. Binds to the promoter region**
- B. Decrease degradation by hydrolytic enzymes**
- C. Remove noncoding information**
- D. Site of ribosome binding**

What is the function of the poly A tail?

B. Decrease degradation by hydrolytic enzymes



The poly-A tail is made up of multiple adenines at the end of the mRNA. As the hydrolytic enzymes in the cytosol hydrolyze the mRNA, it prolongs its "life" in the cytosol to maximize protein synthesized.



What is the function of the 5' cap?

- A. Binds to the promoter region**
- B. Decrease degradation by hydrolytic enzymes**
- C. Remove noncoding information**
- D. Site of ribosome binding**

What is the function of the 5' cap?

D. Site of ribosome binding



The 5' Cap has two functions. It binds with the ribosome for translation (which reads the RNA 5' to 3') at the 5' end. It facilitates the mRNA leaving the nucleus.

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TOPIC

6.3



How can you have the same DNA, but result in multiple different proteins?

AP BIO INSTA-REVIEW

TOPIC

6.3

How can you have the same DNA, but result in multiple different proteins?



Alternative gene splicing



Translation

IST-1.0.1

Translation of the mRNA to generate a polypeptide occurs on ribosomes that are present in the cytoplasm of both prokaryotic and eukaryotic cells and on the rough endoplasmic reticulum of eukaryotic cells.



Translation

IST-1.0.2

In prokaryotic organisms, translation of the mRNA molecule occurs while it is being transcribed.

IST-1.0.3

Translation involves energy and many sequential steps, including initiation, elongation, and termination.



Translation

IST-1.0.4

The salient features of translation include—

- a. Translation is initiated when the rRNA in the ribosome interacts with the mRNA at the start codon.
- b. The sequence of nucleotides on the mRNA is read in triplets called codons.



Translation

IST-1.0.4

The salient features of translation include—

c. Each codon encodes a specific amino acid, which can be deduced by using a genetic code chart. Many amino acids are encoded by more than one codon.

d. Nearly all living organisms use the same genetic code, which is evidence for the common ancestry of all living organisms.



Translation

IST-1.0.4

The salient features of translation include—

- e. tRNA brings the correct amino acid to the correct place specified by the codon on the mRNA.
- f. The amino acid is transferred to the growing polypeptide chain.
- g. The process continues along the mRNA until a stop codon is reached.
- h. The process terminates by release of the newly synthesized polypeptide/protein.



Translation

IST-1.0.5

Genetic information in retroviruses is a special case and has an alternate flow of information: from RNA to DNA, made possible by reverse transcriptase, an enzyme that copies the viral RNA genome into DNA. This DNA integrates into the host genome and becomes transcribed and translated for the assembly of new viral progeny.



What is the function of the ribosomes?

- A. ATP Synthesis**
- B. Digestion**
- C. Protein Synthesis**
- D. Storage**

What is the function of the ribosomes?

C. Protein Synthesis



Ribosomes are the site of protein synthesis. The mRNA is the transcript that brings the message to the ribosome. The tRNA transfers the amino acids so the ribosome can assemble the protein. The rRNA makes up the ribosome.

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TOPIC

6.4



**Identify two locations ribosomes
are found in eukaryotic cells**

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TOPIC

6.4

**Identify two locations
ribosomes are found in
eukaryotic cells**



- > Cytosol**
- > Endoplasmic Reticulum
(rough)**

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TOPIC

6.4



Transcription & Translation are simultaneously completed in a prokaryote

- A. True**
- B. False**

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TOPIC

6.4

**Transcription &
Translation are
simultaneously completed
in a prokaryote**

A. True



Prokaryotes do not have a nuclear membrane. This means that the ribosomes are in the same area as the DNA, so as soon as the mRNA is being synthesized (transcription) the ribosomes can attach to start translation.

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TOPIC

6.4



Why does transcription and translation take place simultaneously in prokaryotic cell?

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TOPIC

6.4



Why does transcription and translation take place simultaneously in prokaryotic cell?

The absence of the nuclear membrane allows the ribosomes to gain access to the mRNA as it is synthesized.



Which describes initiation of translation?

- A. mRNA binds to tRNA with start codon**
- B. mRNA binds to small subunit**
- C. tRNA binds to rRNA with start codon**
- D. tRNA binds to Methionine**

Which describes initiation of translation?

B. mRNA binds to small subunit



Initiation is the start. To start translation, the mRNA binds to the small subunit of the ribosome. The ribosome searches for the start codon (AUG) before the large subunit binds with the tRNA containing methionine.

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TOPIC

6.4



What is the start codon?

A. AUG

B. GUA

C. TAC

D. UAC

What is the start codon?

A. AUG



The start codon is AUG, which will bring the amino acid methionine into the ribosome. This methionine is removed during post-translational processing.

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TOPIC

6.4



**Describe steps in elongation
phase of translation**

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TOPIC

6.4

**Describe steps in
elongation phase of
translation**



**Polypeptide is attached to tRNA at the P site
New tRNA enters bringing the next amino acid
(the anticodon pairs with the codon) at the A
site**

**A new peptide bond forms between the growing
polypeptide and the new amino acid**

**Translocation to move the empty tRNA from the
P site to the E site (and to exit), the tRNA with
the polypeptide to the P site, and the A site
ready to accept a new amino acid**

THEN REPEAT

@APBIOPENGUINS

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TOPIC

6.4



One codon codes for multiple amino acids...

- A. True**
- B. False**

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TOPIC

6.4

One codon codes for multiple amino acids...

B. False



Each codon will only code for ONE amino acid, but there can be multiple codons that code for the SAME amino acid.

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TOPIC

6.4



One amino acid is coded by multiple codons...

- A. True**
- B. False**

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TOPIC

6.4

One amino acid is coded by multiple codons...

A. True



There can be multiple codons that code for the SAME amino acid, but each codon will only code for ONE amino acid.

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TOPIC

6.4



		Second Base in Codon					
		U	C	A	G		
First Base in Codon	U	UUU } Phe UUC } UUA } Leu UUG }	UCU } UCC } Ser UCA } UCG }	UAU } Tyr UAC } UAA Stop UAG Stop	UGU } Cys UGC } UGA Stop UGG Trp	U C A G	
	C	CUU } Leu CUC } CUA } CUG }	CCU } CCC } Pro CCA } CCG }	CAU } His CAC } CAA } Gln CAG }	CGU } CGC } Arg CGA } CGG }	U C A G	
	A	AUU } Ile AUC } AUA } AUG Met or Start	ACU } ACC } Thr ACA } ACG }	AAU } Asn AAC } AAA } Lys AAG }	AGU } Ser AGC } AGA } Arg AGG }	U C A G	
	G	GUU } GUC } Val GUA } GUG }	GCU } GCC } Ala GCA } GCG }	GAU } Asp GAC } GAA } Glu GAG }	GGU } GGC } Gly GGA } GGG }	U C A G	
		Third Base in Codon					

What does AGU code for?

- A. Arg**
- B. Leu**
- C. Phe**
- D. Ser**

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TOPIC

6.4



What does **AGU** code for?

D. Ser

Look in the codon chart for **AGU**:
Left column for the **A** – third row
Top row for the **G** – fourth column
Check the square for the **AGU** – Ser

		Second Base in Codon				
		U	C	A	G	
U	UUU	UCU	UAU	UGU	U C A G	
	UUC	UCC	UAC	UGC		
	UUA	UCA	UAA	UGA		Stop
	UUG	UCG	UAG	UGG		Trp
C	CUU	CCU	CAU	CGU	U C A G	
	CUC	CCC	CAC	CGC		
	CUA	CCA	CAA	CGA		Arg
	CUG	CCG	CAG	CGG		
A	AUU	ACU	AAU	AGU	U C A G	
	AUC	ACC	AAC	AGC		Ser
	AUA	ACA	AAA	AGA		Arg
	AUG	ACG	AAG	AGG		Arg
G	GUU	GCU	GAU	GGU	U C A G	
	GUC	GCC	GAC	GGC		Gly
	GUA	GCA	GAA	GGA		
	GUG	GCG	GAG	GGG		

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TOPIC

6.4



**Prokaryotes can synthesize
human insulin...**

- A. True**
- B. False**

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TOPIC

6.4

**Prokaryotes can
synthesize human
insulin...**

A. True



All organisms have the same genetic code. The same codons code for the same amino acids. A gene from one organism can be inserted into another organism and the same protein can be synthesized.

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TOPIC

6.4



What does it tell us that prokaryotes are able to synthesize human insulin?

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TOPIC

6.4



What does it tell us that prokaryotes are able to synthesize human insulin?

Common ancestry of all living things because of common genetic code among all living things.

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TOPIC

6.4



Which is not a stop codon?

- A. UAA**
- B. UAG**
- C. UGA**
- D. UGG**

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TOPIC

6.4

Which is not a stop codon?

D. UGG



The three stop codons are UGA, UAA, and UAG. The code of UGG is not a stop codon, but it codes for Trp (tryptophan).

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TOPIC

6.4



What is added when the stop codon is reached

- A. Amino Acid**
- B. ATP**
- C. Stop Codon**
- D. Water**

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TOPIC

6.4

What is added when the stop codon is reached

D. Water



The stop codon causes a release factor to bind which will allow water to be used to break the bond. Recall: hydrolysis is the breaking of a water molecule to break a bond.

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TOPIC

6.4



HIV is a retrovirus. What does that mean?

HIV is a retrovirus. What does that mean?



Retroviruses have an RNA genome. They use the enzyme reverse transcriptase to catalyze the synthesis of DNA from their RNA template for insertion in host DNA.



What is translation?

- A. Synthesize DNA under direction of DNA**
- B. Synthesize RNA under direction of DNA**
- C. Synthesize polypeptides under direction of DNA**
- D. Synthesize polypeptides under direction of RNA**

What is translation?

**D. Synthesize polypeptides
under direction of RNA**



Translation is the using a mRNA transcript to synthesize a polypeptide. The mRNA is a disposable copy of the genetic information that is used to assemble the amino acids to form the polypeptide.



Where does translation take place?

- A. Cytosol**
- B. Golgi Bodies**
- C. Nucleus**
- D. Ribosome**

**Where does translation
take place?**

D. Ribosome



Translation takes place in the ribosome. The ribosome will bind to the mRNA (for the message) and the tRNA (for the amino acids) to synthesize the polypeptide.



What is the function of the A site on the ribosome?

- A. Adds the new amino acids**
- B. Attached the new amino acids to the polypeptide**
- C. Location where adenine pairs with uracil**
- D. Location where the empty tRNA leaves the ribosome**

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TOPIC

6.4

What is the function of the A site on the ribosome?

A. Adds the new amino acids



The A site is where the tRNA that is bringing in the next amino acid binds. The A site is where we ADD the next AMINO ACID.



What is the function of the P site on the ribosome?

- A. Holds the growing polypeptide**
- B. Holds phosphorylated polypeptide**
- C. Site that binds to the primary structure of a polypeptide**
- D. Site where the phosphate group binds**

What is the function of the P site on the ribosome?

A. Holds the growing polypeptide



A peptide bond forms between the growing polypeptide and the new amino acid. Then translocation will move the tRNA with the polypeptide to the P site. The P site has the growing **POLYPEPTIDE chain.**

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TOPIC

6.4



**How many nucleotides make up
a codon?**

- A. 1**
- B. 2**
- C. 3**
- D. 4**

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TOPIC

6.4

How many nucleotides
make up a codon?

c. 3



There are three nucleotides in each codon.

If there was only **1** nucleotide per codon, there are only **4** different combinations which isn't enough for the **20** amino acids. If there was only **2** nucleotides per codon, there are only **16** different combinations which isn't enough for the **20** amino acids. If there were **3** nucleotides per codon, it would be able to code for **64** different combinations which is enough for the **20** amino acids.

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TOPIC

6.4

		Second Base in Codon					
		U	C	A	G		
U	First Base in Codon	UUU } Phe	UCU } Ser	UAU } Tyr	UGU } Cys	U	Third Base in Codon
		UUC } Phe	UCC } Ser	UAC } Tyr	UGC } Cys	C	
		UUA } Leu	UCA } Ser	UAA Stop	UGA Stop	A	
		UUG } Leu	UCG } Ser	UAG Stop	UGG Trp	G	
C	First Base in Codon	CUU } Leu	CCU } Pro	CAU } His	CGU } Arg	U	Third Base in Codon
		CUC } Leu	CCC } Pro	CAC } His	CGC } Arg	C	
		CUA } Leu	CCA } Pro	CAA } Gln	CGA } Arg	A	
		CUG } Leu	CCG } Pro	CAG } Gln	CGG } Arg	G	
A	First Base in Codon	AUU } Ile	ACU } Thr	AAU } Asn	AGU } Ser	U	Third Base in Codon
		AUC } Ile	ACC } Thr	AAC } Asn	AGC } Ser	C	
		AUA } Met or Start	ACA } Thr	AAA } Lys	AGA } Arg	A	
		AUG } Met or Start	ACG } Thr	AAG } Lys	AGG } Arg	G	
G	First Base in Codon	GUU } Val	GCU } Ala	GAU } Asp	GGU } Gly	U	Third Base in Codon
		GUC } Val	GCC } Ala	GAC } Asp	GGC } Gly	C	
		GUA } Val	GCA } Ala	GAA } Glu	GGA } Gly	A	
		GUG } Val	GCG } Ala	GAG } Glu	GGG } Gly	G	



What amino acid has a codon of GAC?

- A. Asn**
- B. Asp**
- C. Glu**
- D. Gly**

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TOPIC

6.4

What amino acid has a codon of GAC?

B. Asp



Look in the codon chart for GAC:
Left column for the G – fourth row
Top row for the A – third column
Check the square for the GAC – Asp

		Second Base in Codon					
		U	C	A	G		
First Base in Codon	U	UUU } Phe UUC } UUA } Leu UUG }	UCU } UCC } Ser UCA } UCG }	UAU } Tyr UAC } UAA } Stop UAG } Stop	UGU } Cys UGC } UGA } Stop UGG } Trp	U C A G	
	C	CUU } CUC } Leu CUA } CUG }	CCU } CCC } Pro CCA } CCG }	CAU } His CAC } CAA } Gln CAG }	CGU } CGC } Arg CGA } CGG }	U C A G	
	A	AUU } AUC } Ile AUA } AUG } Met or Start	ACU } ACC } Thr ACA } ACG }	AAU } Asn AAC } AAA } Lys AAG }	AGU } Ser AGC } AGA } Arg AGG }	U C A G	
	G	GUU } GUC } Val GUA } GUG }	GCU } GCC } Ala GCA } GCG }	GAA } Asp GAC GAA } Glu GAG }	GGU } GCC } Gly GGA } GGG }	U C A G	
		Third Base in Codon					

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TOPIC

6.4



**How do retroviruses violate the
Central Dogma?**

**How do retroviruses
violate the Central
Dogma?**



**Retroviruses have an RNA
genome. They use their RNA to
make a DNA template then
insert into the host DNA to
remain dormant until
environmental cue**

AP BIO INSTA-REVIEW

TOPIC

6.4



**Describe the difference between
the protein made by free vs
bound ribosomes**

**How do retroviruses
violate the Central
Dogma?**



**Free ribosomes are freely
floating in the cytosol –
responsible for cytosolic proteins**

**Bound ribosomes are bound to
the rough ER – responsible for
membrane proteins or proteins
for secretion**

AP BIO INSTA-REVIEW

TOPIC

6.4



Free ribosomes and bound ribosomes are the same.

- A. True**
- B. False**

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TOPIC

6.4

Free ribosomes and bound ribosomes are the same.

A. True



All ribosomes begin as free ribosomes. There is a signal peptide which will bring to a SRP (signal recognition particle) to move the ribosome to the rough ER membrane to finish translation as a bound ribosome.



Regulation of Gene Expression

IST-2.A.1

Regulatory sequences are stretches of DNA that interact with regulatory proteins to control transcription.

IST-2.A.2

Epigenetic changes can affect gene expression through reversible modifications of DNA or histones.



Regulation of Gene Expression

IST-2.A.3

The phenotype of a cell or organism is determined by the combination of genes that are expressed and the levels at which they are expressed—

- a. Observable cell differentiation results from the expression of genes for tissue specific proteins.
- b. Induction of transcription factors during development results in sequential gene expression.



Regulation of Gene Expression

IST-2.B.1

Both prokaryotes and eukaryotes have groups of genes that are coordinately regulated—

- a. In prokaryotes, groups of genes called operons are transcribed in a single mRNA molecule. The lac operon is an example of an inducible system.
- b. In eukaryotes, groups of genes may be influenced by the same transcription factors to coordinately regulate expression.

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TOPIC

6.5



**Describe the effect of DNA
methylation**



Describe the effect of DNA methylation

DNA methylation involves binding methyl groups to the DNA. This causes the strand to condense which will inhibit transcription (so decreases gene expression)

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TOPIC

6.5



Describe the effect of histone acetylation

Describe the effect of histone acetylation



Firstly, histones are the proteins that DNA wraps around to condense.

The acetylation decreases the binding between DNA and histone which causes the DNA to loosen.

This increases transcription or stimulates gene expression.



What are the transcription factors?

- A. Molecules that provide ATP to transcription unit**
- B. Molecules that synthesize RNA during transcription**
- C. Molecules that bring amino acids to the transcription unit**
- D. Molecules that bind to enhance/inhibit transcription.**

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TOPIC

6.5

**What are the
transcription factors?**

**D. Molecules that bind to
enhance/inhibit
transcription.**



**Transcription factors bind to the
enhancer regions to assist RNA
polymerase to bind to the
promoter to initiate transcription**

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TOPIC

6.5



All somatic (body) cells have the same genetic information

- A. True**
- B. False**

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TOPIC

6.5

**All somatic (body) cells
have the same genetic
information**

A. True



**All of your cells resulted from a
single cell (zygote) that
underwent mitosis so you could
grow. All of your cells have the
SAME DNA (except for your
gametes due to meiosis)**

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TOPIC

6.5



**If all the cells have the same
DNA how are your cells so
different?**

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TOPIC

6.5

If all the cells have the same DNA how are your cells so different?



Cell differentiation

The cells have different transcription factors/activators which will enhance transcription of certain genes.



What types of cells have operons?

- A. Eukaryotes**
- B. Prokaryotes**
- C. Neither eukaryotes nor prokaryotes**
- D. Both eukaryotes and prokaryotes**

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TOPIC

6.5

What types of cells have operons?

B. Prokaryotes



Operons are only found in prokaryotes. Operons are made up of a promoter, operator, and the genes that they control. It is a simple way for the prokaryotes to organize for gene expression.

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TOPIC

6.5



What are three components of an operon?

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TOPIC

6.5



**What are three
components of an operon?**

- > Promoter**
- > Operator**
- > Genes it regulates**



What binds to the promoter?

- A. DNA polymerase**
- B. Operator**
- C. Repressor**
- D. RNA polymerase**

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TOPIC

6.5

What binds to the promoter?

D. RNA polymerase



The promoter is where the RNA polymerase to initiate transcription.

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TOPIC

6.5



What makes up the TATA box?

- A. Adenine & Cytosine**
- B. Cytosine & Guanine**
- C. Guanine & Thymine**
- D. Thymine & Adenine**

What makes up the TATA box?

D. Thymine & Adenine



The TATA box is made up of T nucleotides (thymine) and A nucleotides (adenine). This is a site on the DNA where transcription factors will bind to aid in RNA polymerase binding.



Where is the TATA box located?

- A. Near the introns**
- B. Near the exons**
- C. Near the poly adenylation sequence**
- D. Near the promoter**

Where is the TATA box located?

D. Near the promoter



The TATA box is near the promoter. The TATA box indicates the DNA strand to be read. Transcription factors bind to this region to assist the RNA polymerase to bind to the promoter.



Describe the effect of adding methyl group to DNA.

- A. Decreases condensation which decreases transcription**
- B. Decreases condensation which increases transcription**
- C. Increases condensation which decreases transcription**
- D. Increases condensation which increases transcription**

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TOPIC

6.5

Describe the effect of adding methyl group to DNA.

C. Increases condensation which decreases transcription



Adding methyl groups to the DNA will cause the DNA to condense which decreases transcription as the RNA polymerase is unable to access the DNA to transcribe.

Recall: XX individuals will undergo methylation to their extra X chromosome to form Barr Bodies.



**Describe effect of acetylating
the histone tails**

- A. Decreases condensation which
decreases transcription**
- B. Decreases condensation which
increases transcription**
- C. Increases condensation which
decreases transcription**
- D. Increases condensation which
increases transcription**

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TOPIC

6.5

**Describe effect of
acetylating the histone
tails**

**B. Decreases condensation
which increases
transcription**



When you acetylate the histone tails, it will cause the histone to be more negatively charged. The DNA and the histone will repel causing the DNA to decrease condensation and increase transcription as the RNA polymerase is able to access the DNA for transcription.

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TOPIC

6.5



All somatic cells within one organism have the same DNA.

- A. True**
- B. False**

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TOPIC

6.5

All somatic cells within one organism have the same DNA.

A. True



All cells originated from the zygote. The cells undergo mitosis forming identical cells that make up your body (somatic cells). The only cells in your body that are different are gametes due to the process of meiosis.

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TOPIC

6.5



What is the function of transcription factors?

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TOPIC

6.5

What is the function of transcription factors?



Binds to initiate transcription or inhibit transcription

Will turn on and off the genes of interest



Where does the repressor in an operon bind?

- A. Operator**
- B. Promoter**
- C. Regulatory sequence**
- D. TATA box**

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TOPIC

6.5

**Where does the repressor
in an operon bind?**

A. Operator



The operator is located in the promoter region. This is the site of binding for the repressor.

If the repressor is bound to the operator, the operon will be OFF. If the repressor is NOT bound to the operator, the operon will be ON.



If the repressor is bound to the operator...

- A. RNA polymerase is able to bind, the operon is off**
- B. RNA polymerase is able to bind, the operon is on**
- C. RNA polymerase is unable to bind, the operon is off**
- D. RNA polymerase is unable to bind, the operon is on**

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TOPIC

6.5

If the repressor is bound to the operator...

C. RNA polymerase is unable to bind, the operon is off



Due to the repressor binding to the operator located in the promoter region, the repressor will block the binding of the RNA polymerase. If RNA polymerase is unable to bind, transcription will be inhibited.



The lac operon is...

- A. Inducible operon so starts off and is activated by lactose**
- B. Inducible operon so starts on and is deactivated by lactose**
- C. Repressible operon so starts off and is activated by lactose**
- D. Repressible operon so starts on and is deactivated by lactose**

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TOPIC

6.5

The lac operon is...

A. Inducible operon so starts off and is activated by lactose



The lac operon synthesizes the enzymes to break down lactose. This operon will start as OFF and be stimulated by the presence of lactose. The lactose acts as an inducer to deactivate the repressor and turn the operon ON. This is an inducible operon.

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TOPIC

6.5



The trp operon is a(n)...

- A. Inducible operon – activated by trp**
- B. Inducible operon – deactivated by trp**
- C. Repressible operon – activated by trp**
- D. Repressible operon – deactivated by trp**

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TOPIC

6.5

The trp operon is a(n)...

**D. Repressible operon –
deactivated by trp**



The trp operon synthesizes the enzymes needed to synthesize tryptophan. This operon will start ON. When trp is present, it will bind to the repressor activating it so it binds to the operator to inhibit transcription and turn the operon OFF. This operon is repressible by the presence of trp.

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TOPIC

6.5



Note:

Histones are proteins that the DNA wraps around to organize it. I think of it like spaghetti (DNA) wrapped around a fork (histone) will allow you to access specific segments of the noodles versus the plate of unorganized pasta.

The histone is positively charged so attracted to DNA which is negatively charged (the phosphate groups each have 2 minus charge)

@APBIOPENGUINS



Gene Expression and Cell Specialization

IST-2.C.1

Promoters are DNA sequences upstream of the transcription start site where RNA polymerase and transcription factors bind to initiate transcription.

IST-2.C.2

Negative regulatory molecules inhibit gene expression by binding to DNA and blocking transcription.



Gene Expression and Cell Specialization

IST-2.D.1

Gene regulation results in differential gene expression and influences cell products and function.

IST-2.D.2

Certain small RNA molecules have roles in regulating gene expression.

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TOPIC

6.6



What is the function of the promoter?

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TOPIC

6.6

What is the function of the promoter?



Location for RNA polymerase to bind.



What is the function of RNA Polymerase?

- A. Synthesize RNA primer**
- B. Synthesize RNA transcript**
- C. Use RNA template to synthesize DNA**
- D. Use RNA template to synthesize protein**

**What is the function of
RNA Polymerase?**

**B. Synthesize RNA
transcript**



RNA polymerase is an enzyme that uses a DNA template to synthesize a RNA transcript. Recall enzymes tell you what they do, RNA Polymerase makes an RNA polymer.

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TOPIC

6.6



**How do transcription factors
affect transcription?**

How do transcription factors affect transcription?



They regulate transcription/gene expression. They can bind to enhance the binding of RNA polymerase.

I explain it like a baseball/softball glove. When you catch a ball, you would rather catch it in your glove with TWO hands instead of bare handling it. Why? It's more stable. The transcription factors create a "secure" fit for the RNA polymerase to get started



What are siRNA?

- A. Small interfering RNA**
- B. Short introns of RNA**
- C. Serine in RNA**

What are siRNA?

A. Small interfering RNA



siRNA are small interfering RNA. These are single stranded RNA that bind to mRNA to inhibit translation or degrade the mRNA.

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TOPIC

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**Describe what is the function of
siRNA**

Describe what is the function of siRNA



Small interfering RNAs (siRNA)
> short segments of RNA (21-28 bases) bind to mRNA, create sections of double-stranded mRNA, "death" tag for mRNA (triggers degradation of mRNA)
> cause gene "silencing" by post-transcriptional control, turns off gene so no protein produced

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TOPIC

6.6



**Describe the mechanism involved
with miRNA**



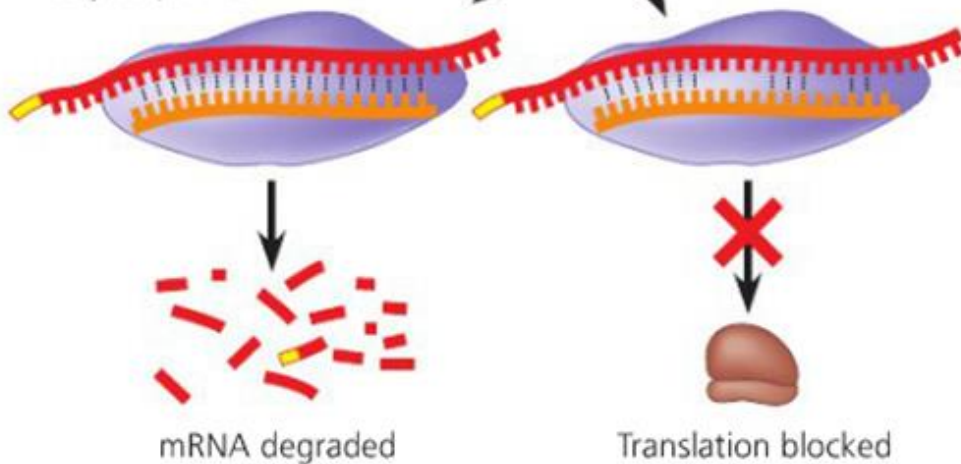
Describe the mechanism involved with miRNA

1 An enzyme cuts each hairpin from the primary miRNA transcript.

2 A second enzyme, called Dicer, trims the loop and the single-stranded ends from the hairpin, cutting at the arrows.

3 One strand of the double-stranded RNA is degraded; the other strand (miRNA) then forms a complex with one or more proteins.

4 The miRNA in the complex can bind to any target mRNA that contains at least 7 bases of complementary sequence.



5 If miRNA and mRNA bases are complementary all along their length, the mRNA is degraded (left); if the match is less complete, translation is blocked (right).



What is the function of the promoter?

- A. Activators bind to promote transcription**
- B. Motor proteins binds with actin for transcription**
- C. Repressors bind to initiate transcription**
- D. Site of RNA polymerase binding for transcription**

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TOPIC

6.6

What is the function of the promoter?

D. Site of RNA polymerase binding for transcription



The promoter is the region of DNA where the RNA polymerase binds to initiate transcription.

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TOPIC

6.6



What is the function of transcription factors?

What is the function of transcription factors?



Molecules bind to the enhancer region to promote transcription

The binding of transcription factors enhances the binding of RNA polymerase to promoter region to facilitate transcription

They are component of the initiation transcription complex



What is the function of RNA polymerase?

- A. Binds to DNA to synthesize an RNA primer**
- B. Binds to DNA to synthesize an RNA transcript**
- C. Binds to RNA to base pair with other RNA strand**
- D. Binds to RNA to seal phosphodiester linkages**

What is the function of RNA polymerase?

B. Binds to DNA to synthesize an RNA transcript



RNA polymerase is an enzyme that initiates transcription. The RNA polymerase binds to the DNA at the promoter. It uses the DNA as a template to synthesize a RNA transcript.

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TOPIC

6.6



What is differential gene expression?

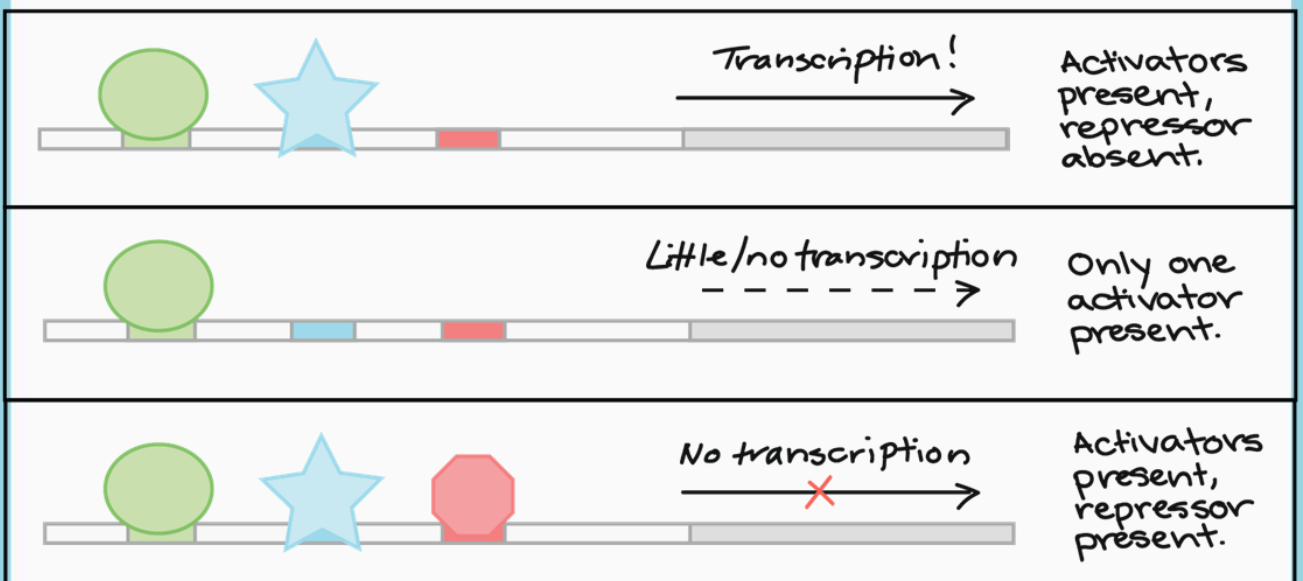
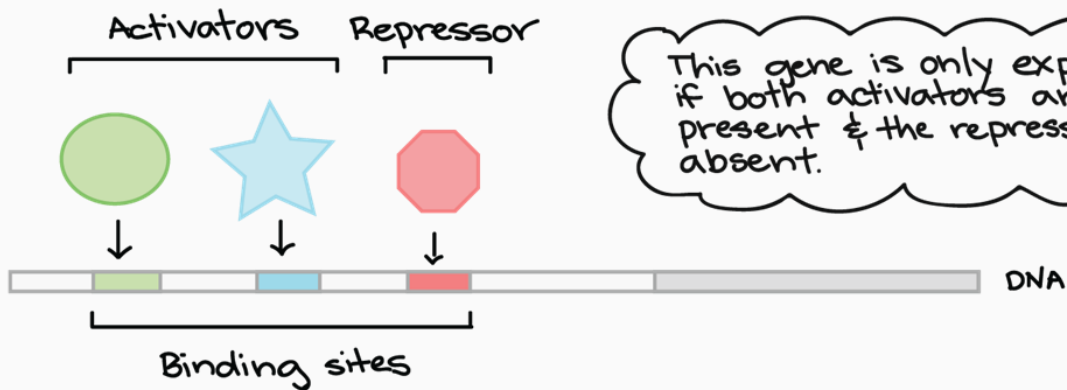
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What is differential gene expression?



Credit: Khan Academy

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TOPIC

6.6



What is the function of transcription factors?

What is the function of transcription factors?



Different genes are expressed by different cells

So even though all of your cells have the same DNA, a liver cell has the components and functionality of a liver cell vs your lens (eye) cell has the components and functionality of a lens cell.



What is siRNA?

- A. Small RNA strand that binds to mRNA to inhibit translation
- B. Small RNA strand that binds to DNA to inhibit transcription
- C. Small RNA strand that removes introns from mRNA
- D. Small RNA strand that removes exons from mRNA

What is siRNA?

A. Small RNA strand that binds to mRNA to inhibit translation



siRNA is a small interfering RNA. This single stranded RNA will bind to mRNA to inhibit translation or degrade the mRNA.



Mutations

IST-2.E.1

Changes in genotype can result in changes in phenotype—

- a. The function and amount of gene products determine the phenotype of organisms.
 - i. The normal function of the genes and gene products collectively comprises the normal function of organisms.
 - ii. Disruptions in genes and gene products cause new phenotypes.



Mutations

IST-2.E.2

Alterations in a DNA sequence can lead to changes in the type or amount of the protein produced and the consequent phenotype. DNA mutations can be positive, negative, or neutral based on the effect or the lack of effect they have on the resulting nucleic acid or protein and the phenotypes that are conferred by the protein.



Mutations

IST-4.A.1

Errors in DNA replication or DNA repair mechanisms, and external factors, including radiation and reactive chemicals, can cause random mutations in the DNA—

- a. Whether a mutation is detrimental, beneficial, or neutral depends on the environmental context.
- b. Mutations are the primary source of genetic variation.



Mutations

IST-4.A.2

Errors in mitosis or meiosis can result in changes in phenotype—

- a. Changes in chromosome number often result in new phenotypes, including sterility caused by triploidy, and increased vigor of other polyploids.
- b. Changes in chromosome number often result in human disorders with developmental limitations, including Down syndrome/ Trisomy 21 and Turner syndrome.



Mutations

IST-4.B.1

Changes in genotype may affect phenotypes that are subject to natural selection. Genetic changes that enhance survival and reproduction can be selected for by environmental conditions—

- a. The horizontal acquisitions of genetic information primarily in prokaryotes via transformation (uptake of naked DNA), transduction (viral transmission of genetic information), conjugation (cell-to-cell transfer of DNA), and transposition (movement of DNA segments within and between DNA molecules) increase variation.



Mutations

IST-4.B.1

Changes in genotype may affect phenotypes that are subject to natural selection. Genetic changes that enhance survival and reproduction can be selected for by environmental conditions—

- b. Related viruses can combine/recombine genetic information if they infect the same host cell.**
- c. Reproduction processes that increase genetic variation are evolutionarily conserved and are shared by various organisms.**

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TOPIC

6.7



What are mutations?

What are mutations?



**Changes in the DNA sequence or
the DNA quantity in a cell.**



Which mutation results in a change in the amino acid?

- A. Frameshift**
- B. Missense**
- C. Nonsense**
- D. Silent**

Which mutation results in a change in the amino acid?

B. Missense



A missense mutation is a substitution point mutation. One nucleotide is substituted with another. This substitution causes **ONE amino acid to be substituted for **ANOTHER** amino acid.**

This is similar to a word being substituted in a sentence.

I took my dog for a walk.

I took my fish for a walk.



Which mutation results in no change in the amino acid?

- A. Frameshift**
- B. Missense**
- C. Nonsense**
- D. Silent**

Which mutation results in no change in the amino acid?

D. Silent



A silent mutation is a substitution point mutation. One nucleotide is substituted with another. This substitution causes ONE amino acid to be substituted for the SAME amino acid.

Since you have the same amino acid, it is not obvious that there has been a mutation so it is SILENT.



**Which mutation results from
+/- one nucleotide base?**

- A. Frameshift**
- B. Missense**
- C. Nonsense**
- D. Silent**

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TOPIC

6.7

Which mutation results from +/- one nucleotide base?

A. Frameshift



If ONE nucleotide has been inserted or deleted, this will shift the reading frame. Recall, the reading frame is the group of triplets read as codons.

If there is a shift by 1 or 2, the ribosome will be pairing the wrong codons. This will cause a change to all amino acids downstream from the insertion or deletion.



Frameshift mutation in what area causes most damage?

- A. Intron**
- B. Start of exon**
- C. Middle of exon**
- D. End of exon**

Frameshift mutation in what area causes most damage?

B. Start of exon



If the frameshift mutation occurs at the start of the expressed exons, all of the codons downstream will be affected. This will cause the most damage of the options.

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TOPIC

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How does a heterozygote and homozygous dominant produce same phenotype?

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TOPIC

6.7



How does a heterozygote and homozygous dominant produce same phenotype?

The single dominant allele produces enough enzyme/protein products to produce the desired phenotypic result.



What is the primary source of genetic variation?

- A. Crossing over**
- B. Independent assortment**
- C. Mutations**
- D. Random fertilization**

What is the primary source of genetic variation?

C. Mutations



Mutations will add NEW variation to the population. This will increase genetic variation.



Which of the following is the cause of Trisomy 21?

- A. Duplication**
- B. Frameshift mutation**
- C. Nondisjunction**
- D. Point mutation**



Which of the following is the cause of Trisomy 21?

C. Nondisjunction

Individuals with Trisomy 21 has an additional chromosome 21. This results from nondisjunction. Nondisjunction results from homologous chromosomes or sister chromatids not separating during anaphase I or II.

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TOPIC

6.7



What phase of meiosis is cause of nondisjunction? Describe nondisjunction.

**What phase of meiosis is cause of nondisjunction?
Describe nondisjunction.**



Anaphase I or II

Nondisjunction involves the chromatids or homologous chromosomes not separating during anaphase. This results in a gamete with an additional chromosome or missing a chromosome.

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TOPIC

6.7



**Triploid organisms are sterile
(unable to reproduce)**

- A. True**
- B. False**

Triploid organisms are sterile (unable to reproduce)

A. True



Triploid organisms lack homologous chromosomes. They have THREE of each chromosome thus cannot pair during prophase I, line up during metaphase I, nor segregate evenly during anaphase I.

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TOPIC

6.7



**Why are triploid organisms
sterile?**

Why are triploid organisms sterile?



Unable for homologous chromosomes to pair up during meiosis.

(Triploid is three sets, how can you get paired off with three)



Which enzyme has proofreading capabilities?

- A. DNA Polymerase**
- B. Helicase**
- C. Ligase**
- D. Topoisomerase**

Which enzyme has proofreading capabilities?

A. DNA Polymerase



DNA polymerase is the enzyme responsible for DNA replication. A $3' \rightarrow 5'$ proofreading exonuclease domain is intrinsic to most DNA polymerases. It allows the enzyme to check each nucleotide during DNA synthesis and excise mismatched nucleotides in the $3'$ to $5'$ direction.



Which is responsible for halting DNA replication because of error?

- A. DNA Polymerase**
- B. Helicase**
- C. p53**
- D. Ras**

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TOPIC

6.7

Which is responsible for halting DNA replication because of error?

C. **p53**



p53 is a tumor suppressor gene/protein. **p53** is a transcription factor that activates many genes involved in essential maintenance of genetic stability. This gene will inhibit the cell cycle in the presence of DNA damage.



In disease with too many copies of genes (Huntington's)...

- A. Deletion**
- B. Duplication**
- C. Inversion**
- D. Translocation**

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TOPIC

6.7

In disease with too many
copies of genes
(Huntington's)...

B. Duplication



Duplication results when a gene is duplicated resulting in multiple copies of the gene.

Huntington's Disease is an autosomal dominant trait that results from duplication.

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6.7



**Involved in cri-du-chat where
chromosome is shorter**

- A. Deletion**
- B. Duplication**
- C. Inversion**
- D. Translocation**

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TOPIC

6.7

Involved in cri-du-chat
where chromosome is
shorter

A. Deletion



Deletion results in a region of
the chromosome being removed.

Cri-du-chat ("cry of the cat")
is a genetic disorder resulting
from a piece of chromosome 5 is
missing.



**Involved in down syndrome
without nondisjunction**

- A. Deletion**
- B. Duplication**
- C. Inversion**
- D. Translocation**

**Involved in down
syndrome without
nondisjunction**

D. Translocation



**Translocation results from
chromosome segments changing
positions.**

**This sometimes takes place in
Down Syndrome and Patau
syndrome resulting an additional
chromosome in the offspring.**

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TOPIC

6.7



What is horizontal transmission of traits?

**What is horizontal
transmission of traits?**



**Transmission of traits NOT from
parent-progeny**



Cell-to-cell transmission?

- A. Conjugation**
- B. Transduction**
- C. Transformation**
- D. Transposition**



Cell-to-cell transmission?

A. Conjugation

Prokaryotes are able to undergo cell-to-cell transmission by passing genetic information through sex pili in a process called conjugation.



Movement of DNA between or within segments?

- A. Conjugation**
- B. Transduction**
- C. Transformation**
- D. Transposition**

**Movement of DNA
between or within
segments**

D. Transposition



Transposition involves the DNA moving to a new location so it transfers position when it undergoes transposition.



Viral transmission of traits

- A. Conjugation**
- B. Transduction**
- C. Transformation**
- D. Transposition**

**Viral transmission of
traits**

B. Transduction



As seen during the Hershey and Chase experiment, bacteriophages traditionally will transduce their genetic information into the host nucleus.



Uptake naked DNA

- A. Conjugation**
- B. Transduction**
- C. Transformation**
- D. Transposition**

Uptake naked DNA

C. Transformation



During the transformation experiment, bacteria will take up genetic information (plasmids) from the environment to become antibiotic resistant bacteria/green glowing bacteria.

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6.7



What type of virus is able to insert its genome into host genome?



What type of virus is able to insert its genome into host genome?

Retrovirus

Viruses like HIV have an RNA genome. They can use reverse transcriptase to make a complement DNA strand then inserts in genome to become a pro virus.



What is the genotype?

- A. Genes in a genetic library for transformation**
- B. Physical characteristics due to gene expression**
- C. Specific genes an organism has (ex: heterozygous)**
- D. The type of genetic material the organism has**

What is the genotype?

C. Specific genes an organism has (ex: heterozygous)



Genotype is the genetic information within the DNA. This is traditionally discussed as homozygous dominant, heterozygous, or homozygous recessive.

For example, Pp.



What is the phenotype?

- A. Genes in an organism causing physical traits**
- B. Good genes that increase reproductive fitness**
- C. Physical characteristics of an organism**
- D. Showing types of genes (ex. homozygous dominant)**

What is the phenotype?

**C. Physical characteristics
of an organism**



**Phenotype is the physical
characteristics that result from
the genotype.**

For example, purple flowers.



Mutation resulting in a different amino acid

- A. Chromosomal**
- B. Missense**
- C. Nonsense**
- D. Silent**

**Mutation resulting in a
different amino acid**

B. Missense



**Missense mutation results in a
DIFFERENT amino acid. This is
due to a single nucleotide
substitution.**

**Example: Sickle Cell,
Cystic Fibrosis**



Mutation resulting in the same amino acid

- A. Chromosomal**
- B. Missense**
- C. Nonsense**
- D. Silent**

**Mutation resulting in
the same amino acid**

D. Silent



**Silent mutations have no
observable difference in the
amino acid sequence. The codon
before and after will code for
the same amino acid.**



Mutation resulting in a stop codon (premature stop)

- A. Chromosomal**
- B. Missense**
- C. Nonsense**
- D. Silent**

Mutation resulting in a stop codon (premature stop)



C. Nonsense

A nonsense mutation results in a stop codon for a substitution of codon that codes for an amino acid which leads to a shorter protein product.



If a nucleotide base pair was deleted in an intron, what would be the result?

- A. Frameshift mutation**
- B. Longer protein product**
- C. No change in synthesized protein**
- D. Shorter protein product**

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TOPIC

6.7

If a nucleotide base pair was deleted in an intron, what would be the result?

C. No change in synthesized protein



Recall, the introns are removed during post-transcriptional modification. If a mutation takes place in the intron, then it will not be observed as the introns are removed.



What happens if a nucleotide base pair inserted into an exon?

- A. Frameshift mutation**
- B. Longer protein product**
- C. No change in synthesized protein**
- D. Shorter protein product**

What happens if a nucleotide base pair inserted into an exon?

A. Frameshift mutation



If a single nucleotide base pair is inserted, this will shift the reading frame resulting in a frameshift mutation.



What is the result of a mutation?

- A. Gain new trait**
- B. Loss of trait**
- C. No change in trait**
- D. All of the above**

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TOPIC

6.7

What is the result of a mutation?

**D. All of the above
(Gain, Loss, or No change
in trait)**



The mutation could cause a different ligand to bind which could result in a new function.

The mutation could inhibit binding of a ligand which could result in the absence of a function.

The mutation could have no effect on the binding site or on the folding of the protein resulting in no change in function.



Which results in new genetic variability?

- A. Crossing over**
- B. Independent assortment**
- C. Mutations**
- D. Random fertilization**

Which results in new genetic variability?

C. Mutations



**Mutations results in new traits.
If there are more traits available, the genetic variability has increased.**



What does it mean to be triploid?

- A. Three units of chromosomes**
- B. Three sets of a chromosome**
- C. Three sets of organisms born**
- D. Three sets of population**

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TOPIC

6.7

What does it mean to be triploid?

A. Three units of chromosomes



The prefix of "tri" means **THREE**.

The suffix of "ploid" means having **CHROMOSOME SETS**

Triploid has **THREE SETS** of **CHROMOSOMES**

AP BIO INSTA-REVIEW

TOPIC

6.7



**Why are triploid organisms
sterile?**

Why are triploid organisms sterile?



They are unable to pair up the homologous chromosomes during meiosis to synthesize the gametes

AP BIO INSTA-REVIEW

TOPIC

6.7



**Natural selection acts on
genotype.**

- A. True**
- B. False**

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TOPIC

6.7

Natural selection acts on genotype.

B. False



Natural selection cannot act on what it cannot observe. Natural selection acts on phenotype, but it does affect the genotype resulting is an increase in allele frequency.

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TOPIC

6.7



**Natural selection affects
genotype.**

- A. True**
- B. False**

Natural selection affects genotype.

A. True



Natural selection will select individuals with favorable traits that align with the environment. As these traits are selected, the allele frequency will change which affects the genotype of the organisms.



Taking up DNA from the environment

- A. Conjugation**
- B. Transduction**
- C. Transformation**
- D. Transposition**

Taking up DNA from the environment

C. Transformation



Transformation was observed in the Griffith experiment. The living R cells are transformed into living S cells as they took up DNA from the heat-killed S cells.



**Insertion of DNA by
bacteriophage**

- A. Conjugation**
- B. Transduction**
- C. Transformation**
- D. Transposition**

**Insertion of DNA by
bacteriophage**

B. Transduction



Transduction is the process of a bacteriophage injecting the generic information which transduces the host cell.



DNA moves to new position in genome

- A. Conjugation**
- B. Transduction**
- C. Transformation**
- D. Transposition**

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TOPIC

6.7

**DNA moves to new
position in genome**

D. Transposition



**Genetic information is able to be
transferred to another position
through transposition.**



**Genetic exchange between
bacteria**

- A. Conjugation**
- B. Transduction**
- C. Transformation**
- D. Transposition**

**Genetic exchange between
bacteria**

A. Conjugation



**Conjugation is the attachment
of two prokaryotes then
transferring genetic information
between the two which increases
the genetic variability.**



Biotechnology

IST-1.P.1

Genetic engineering techniques can be used to analyze and manipulate DNA and RNA—

- a. Electrophoresis separates molecules according to size and charge.
- b. During polymerase chain reaction (PCR), DNA fragments are amplified.
- c. Bacterial transformation introduces DNA into bacterial cells.
- d. DNA sequencing determines the order of nucleotides in a DNA molecule



Which DNA tech separates fragments by size/charge?

A. Gel electrophoresis

B. PCR

C. Sequencing

D. Transformation

Which DNA tech separates fragments by size/charge?

A. Gel electrophoresis



Gel electrophoresis involves using an electrical current to draw DNA, RNA, or protein through a gel. The gel has pores which allows the small fragments to travel farther.



Why does the DNA move towards positive end?

- A. Deoxyribose is negatively charged**
- B. Deoxyribose is positively charged**
- C. Phosphate is negatively charged**
- D. Phosphate is positively charged**

Why does the DNA move towards positive end?

C. Phosphate is negatively charged



Recall, DNA is made up of a nitrogenous base, deoxyribose, and a phosphate group.

The phosphate group is negatively charged with PO_4^{2-} so it will be attracted to the positive end.

AP BIO INSTA-REVIEW

TOPIC

6.8



Which fragments travel farther?

- A. Small**
- B. Large**

Which fragments travel farther?

A. Small



Smaller fragments are able to pass through the pores quicker.

Another teacher describes it visually as comparing an adult to a kid in the ballpit of the playground. The child will be able to get out of the ballpit easier than the adult.



What cuts the DNA prior to running gel electrophoresis?

- A. DNA polymerase**
- B. Helicase**
- C. Ligase**
- D. Restriction enzyme**

AP BIO INSTA-REVIEW

TOPIC

6.8

What cuts the DNA prior to running gel electrophoresis?

D. Restriction enzyme



Restriction enzymes will cut the DNA at specific restriction sites. This will form fragments that are run on a gel electrophoresis. When comparing two strands of DNA, researchers will use the same restriction enzymes.

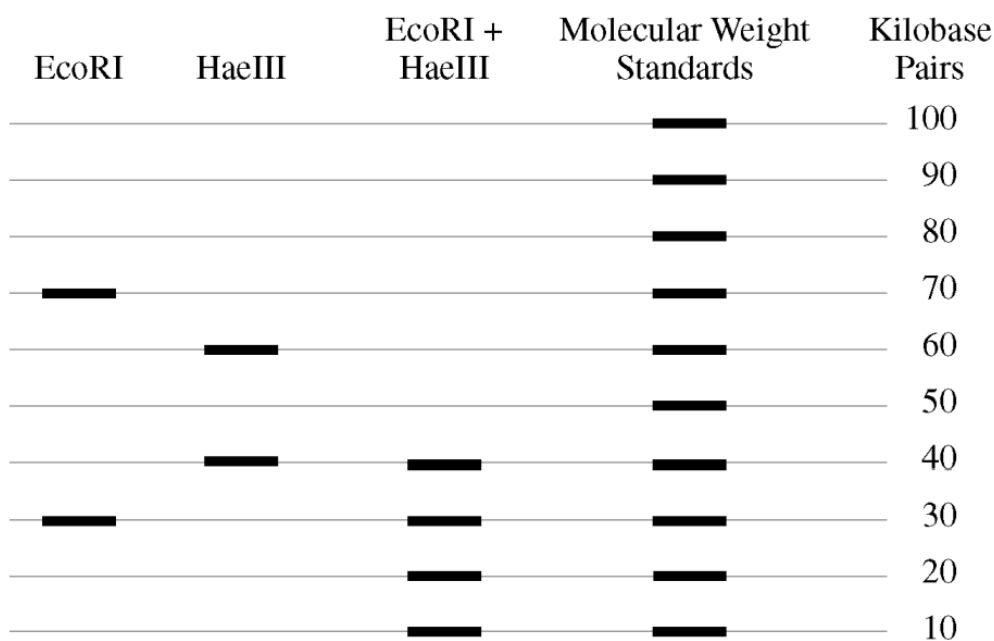
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RESULTS OF GEL ELECTROPHORESIS



Minimum restriction sites for ECO on plasmid to get this?

- A. 1**
- B. 2**
- C. 3**
- D. 4**

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Minimum restriction sites for ECO on plasmid to get this?

B. 2



Plasmids are circular DNA. If you look at the gel electrophoresis, you would observe that ECO has 2 fragments which means it was cut 2 times. If it was cut 2 times, there must be 2 restriction enzymes

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Evidence	A	B	C	D
—	—	—	—	—
==	—	==	—	—
—	—	—	—	—
—	—	—	—	—
—	—	—	—	—
—	—	—	—	—
—	—	—	—	—



Which suspect was at crime scene?

A. A

B. B

C. C

D. D

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**Which suspect was at
crime scene?**

B. B



**When comparing the banding
pattern between the crime scene
and the suspects, it is
observable that section B has
the same bands as the
restaurant.**



Which DNA tech is responsible for multiple copies of a gene?

- A. Gel electrophoresis**
- B. PCR**
- C. Sequencing**
- D. Transformation**

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Which DNA tech is responsible for multiple copies of a gene?

B. PCR



PCR stands for polymerase chain reaction. This procedure will create multiple copies of a certain DNA to compare and run testing on.

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What does PCR stand for?

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**What does PCR stand
for?**



Polymerase Chain Reaction

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PCR uses a specific type of polymerase, why and where does it come from?

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PCR uses a specific type of polymerase, why and where does it come from?

PCR involves a heating up then cooling cycle. Proteins (polymerase) denature in high temperatures so heat resistant polymerase from thermophile prokaryotes is needed to allow the enzymes to remain functional after heating phase.

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What is the function of the heating process in PCR?

What is the function of the heating process in PCR?



- > Break the hydrogen bonds between the two strands**
- > Allows the strands to be single stranded to allow for the replication to take place in future steps**



Which DNA tech involves adding plasmid to prokaryote?

- A. Gel electrophoresis**
- B. PCR**
- C. Sequencing**
- D. Transformation**

Which DNA tech involves adding plasmid to prokaryote?

D. Transformation



Transformation is a process by which foreign DNA is introduced into a cell. During this lab, we added a plasmid to a solution with E. coli then heat shocked to introduce the plasmid into the bacteria. It is then transformed into an antibiotic resistant bacteria.

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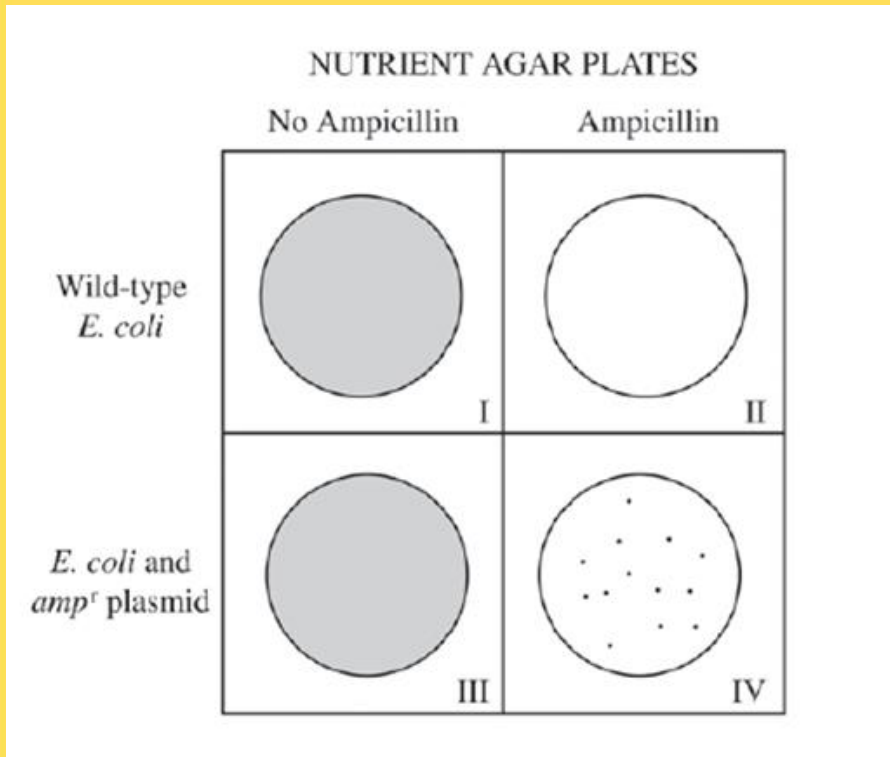


Plate has non transformed bacteria & no antibiotics

- A. I**
- B. II**
- C. III**
- D. IV**

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Plate has non transformed bacteria & no antibiotics

A. I

Plate I has the wild-type E. coli (non-transformed bacteria) and no ampicillin (no antibiotics)

It has a lawn of growth because there is no antibiotic to inhibit the bacterial growth.

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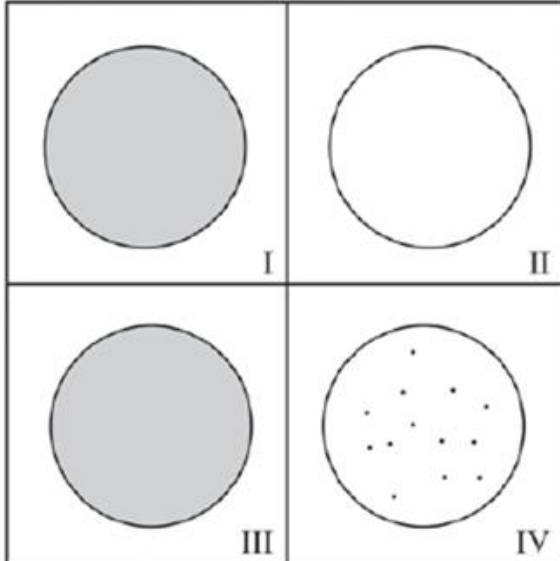


NUTRIENT AGAR PLATES

No Ampicillin

Ampicillin

Wild-type
E. coli



I

II

III

IV

E. coli and
amp^r plasmid

**Why is there no growth
on plate II?**

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**Why is there no growth
on plate II?**

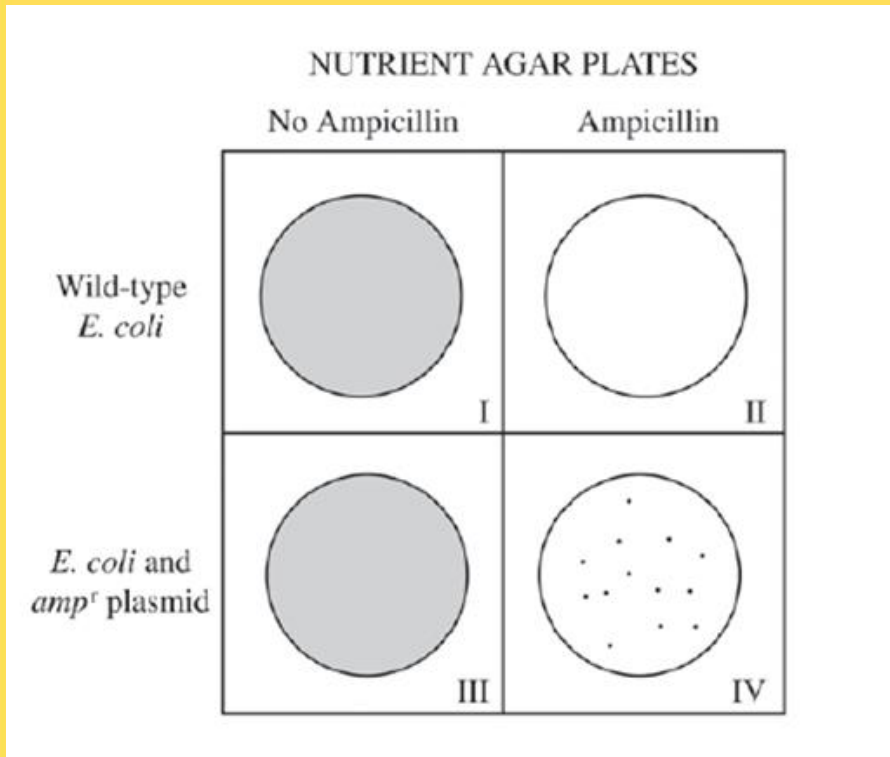


**Bacteria is not transformed and
there is antibiotic in the agar
which restricts growth.**

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What is function of plates I and III?



What is function of plates I and III?

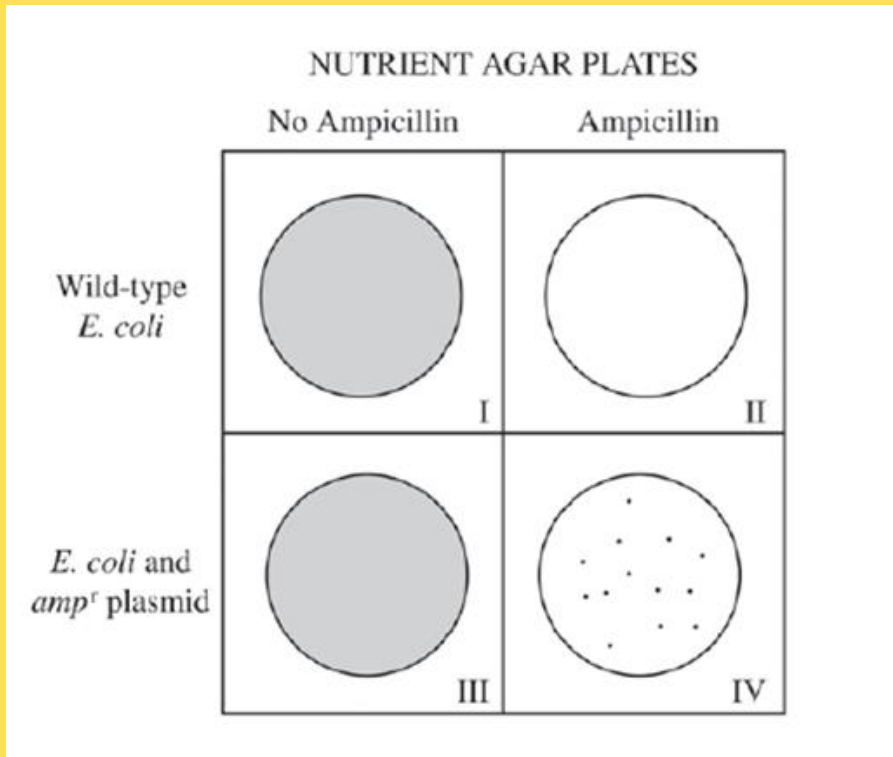
Control:

Positive control- you are verifying that the bacteria is viable. You need to be sure that the bacteria would grow in the absence of the antibiotic to know that the antibiotic killed non-resistant bacteria (explains plates II and IV)

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If plasmid also had the gene to synthesize insulin...

Which plate has the highest % of insulin producing bacteria?

- A. I
- B. II
- C. III
- D. IV

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If plasmid also had the gene to synthesize insulin...

Which plate has the highest % of insulin producing bacteria?

D. IV



Plate IV has *E. coli/amp* plasmid and ampicillin. The ampicillin will restrict the growth of the non-transformed bacteria. The only bacteria that will grow on this plate is the bacteria with the plasmid containing the gene to synthesize insulin, so **100%** of the bacteria are insulin producing bacteria.

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**DNA is ___ because
of the ____.**

- A. Negatively charged; hydroxyl group**
- B. Negatively charged; phosphate group**
- C. Positively charged; hydroxyl group**
- D. Positively charged; phosphate group**

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**DNA is ___ because
of the ____.**

**B. Negatively charged;
phosphate group**



**DNA is negatively charged
because of the phosphate group.
Phosphate is PO_4^{2-} and DNA
has an alternating sugar and
phosphate backbone.**



Gel electrophoresis separates molecules based on

- A. Charge and Polarity**
- B. Polarity and Weight**
- C. Weight and Size**
- D. Size and Charge**

**Gel electrophoresis
separates molecules based
on**

D. Size and Charge



Gel electrophoresis involves a gel with pores that allow for materials to be separated by size. Electrophoresis involves using an electrical current so the charge of the material will allow for movement through the gel.



In a gel electrophoresis...

- A. More numerous sizes are near the wells**
- B. Less numerous sizes are near the wells**
- C. Larger molecules are near the wells**
- D. Shorter molecules are near the wells**

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In a gel electrophoresis...

**C. Larger molecules
are near the wells**



The large molecules will be unable to pass through the pores as quickly and will be located near the wells (where the DNA was added to the gel). Smaller molecules will travel faster and thus farther.



What does a thicker band represent in a gel?

- A. There's more DNA material present in that band**
- B. There's more glucose available in that band**
- C. There's more mutation in the DNA fragment present**
- D. There's more radioactive material present in that band**

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What does a thicker band represent in a gel?

A. There's more DNA material present in that band



The DNA that is the same length will travel the same distance. If multiple fragments travel the same distance, it will cause the band in the gel to be thicker.



What is used to cut the DNA into fragments for the gel?

- A. Ligase from eukaryotes**
- B. Ligase from prokaryotes**
- C. Restriction enzymes from eukaryotes**
- D. Restriction enzymes from prokaryotes**

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What is used to cut the DNA into fragments for the gel?

D. Restriction enzymes from prokaryotes



Restriction enzymes will cut the DNA at restriction sites. The prokaryotes use restriction enzymes to cleave foreign DNA.



What does PCR stand for?

- A. Place for cellular respiration**
- B. Primase cytosine reactants**
- C. Polymerase chain reaction**
- D. Preferred chemical reaction**

What does PCR stand for?

C. Polymerase chain reaction



PCR stands for Polymerase Chain Reaction. This is a process that will amplify a segment of DNA using a thermocycler, DNA polymerase, a primer and DNA nucleotides.

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**What are the steps
involved in PCR?**

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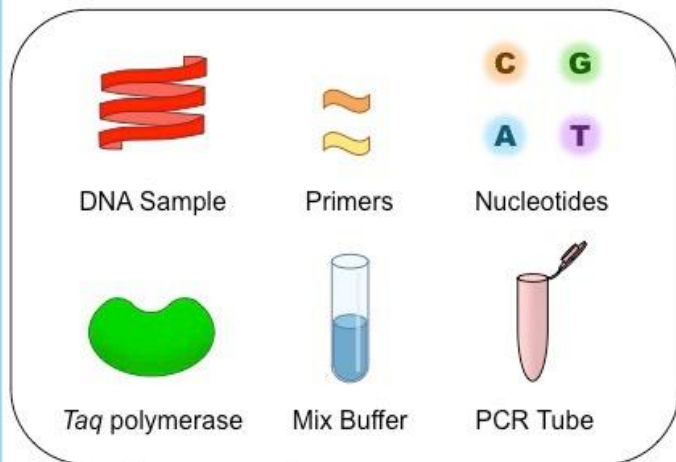
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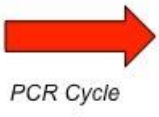


What are the steps involved in PCR?

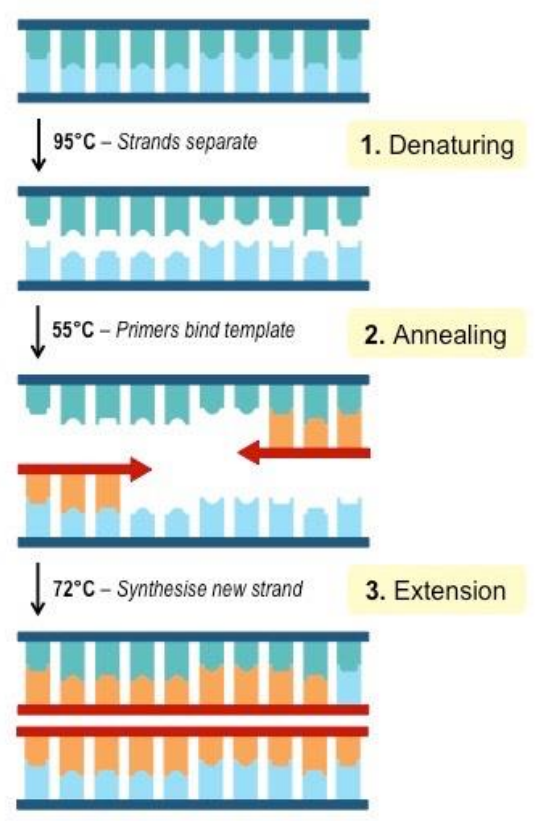
PCR Components



Thermal Cycler



PCR Process (ONE Cycle)





Where does the heat resistant polymerase come from?

- A. Fire-resistant plants**
- B. Monkeys**
- C. Nucleus**
- D. Thermophiles**

Where does the heat resistant polymerase come from?

D. Thermophiles



DNA polymerase is an enzyme (and a protein). Recall, heat will denature proteins. The process of PCR has repeated heating and cooling which would denature the enzyme. The thermophiles are bacteria that thrive at extremely high temperatures.



Plasmid makes bacteria resistant to ampicillin and able to synthesize the green glow protein. Which plate will have highest percent of growth?

- A. Plate with nutrients and no ampicillin present**
- B. Plate with ampicillin and nutrient present**

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Plasmid makes bacteria resistant to ampicillin and able to synthesize the green glow protein. Which plate will have highest percent of growth?

A. Plate with nutrients and no ampicillin present

Not all bacteria will be transformed, so the plate with nutrient only would have a lawn of growth while the plate with ampicillin would have colonies. Read the question carefully, it said highest percent of growth.



Plasmid makes bacteria resistant to ampicillin and able to synthesize the green glow protein. Which plate will have higher amount of glowing bacteria?

- A. Plate with nutrients and no ampicillin present**
- B. Plate with ampicillin and nutrient present**

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Plasmid makes bacteria resistant to ampicillin and able to synthesize the green glow protein. Which plate will have higher amount of glowing bacteria?

B. Plate with ampicillin and nutrient present



The plate with the ampicillin will restrict the growth to the transformed bacteria only. All of the bacteria on this plate will be transformed, thus will glow.

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What happens during DNA sequencing?

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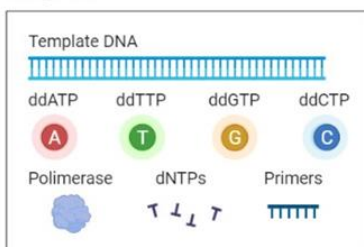
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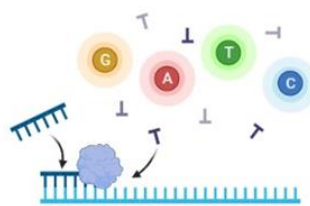


What happens during DNA sequencing?

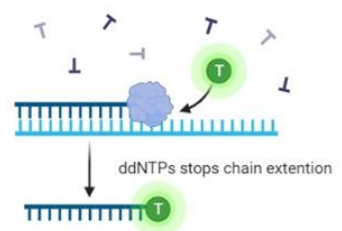
Reagents



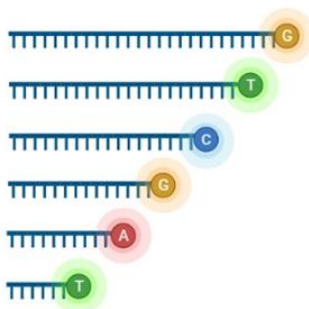
① Primer annealing and chain extension



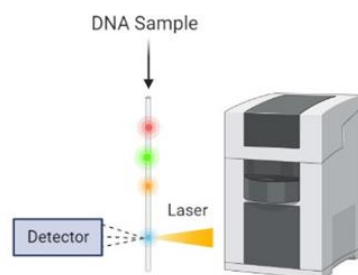
② ddNTP binding and chain termination



③ Fluorescently labelled DNA sample



④ Capillary gel electrophoresis and fluorescence detection



⑤ Sequence analysis and reconstruction

