



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

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



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

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Transcription & Translation are simultaneously completed in a prokaryote

A. True

B. False

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

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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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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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**Describe steps in elongation
phase of translation**

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**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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One codon codes for multiple amino acids...

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

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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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One amino acid is coded by multiple codons...

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

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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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| | | 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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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 | 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 } | GAU } Asp GAC } GAA } Glu GAG } | GGU } GGC } Gly GGA } GGG } | U C A G |

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

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

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

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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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Which is not a stop codon?

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

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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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What is added when the stop codon is reached

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

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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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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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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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How many nucleotides make up a codon?

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

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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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| | | 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 } Ile | 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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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 | Third Base in Codon | 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 |

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**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**

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**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**

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Free ribosomes and bound ribosomes are the same.

A. True

B. False

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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.