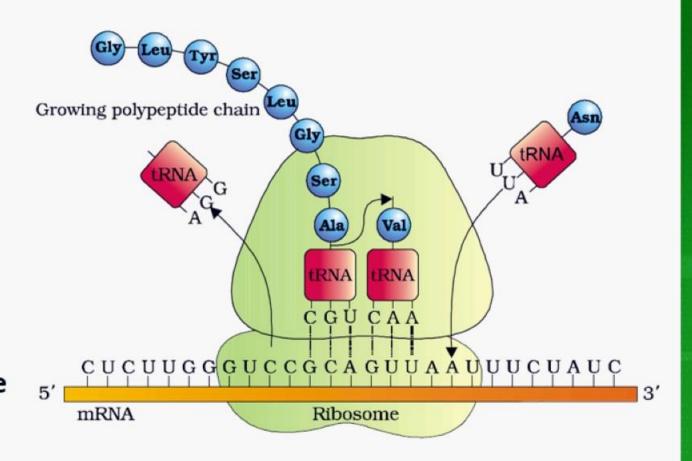


TRANSLATION (PROTEIN SYNTHESIS)

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- It is the process of polymerisation of amino acids to form a polypeptide based on the sequence of codons in mRNA.
- It takes place in ribosomes.
- Ribosome consists of structural RNAs and about 80 types of proteins.
- Ribosome also acts as a catalyst (235 rRNA in bacteria is the enzymeribozyme) for the formation of peptide bond (peptidyl transferase enzyme in large subunit of ribosome).



TRANSLATION (PROTEIN SYNTHESIS)

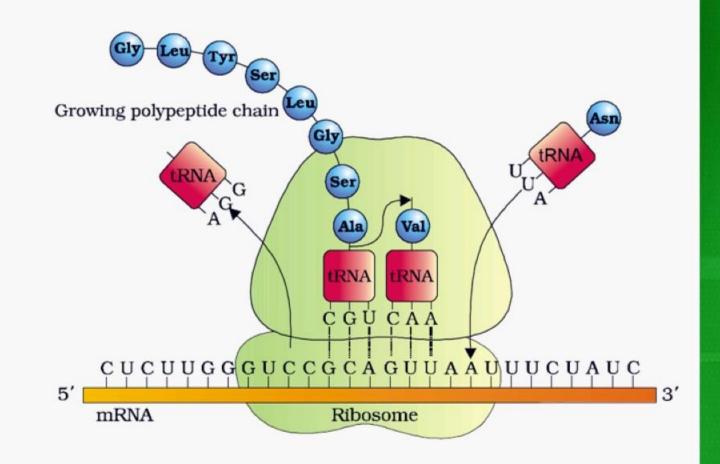
Steps of Translation

CHARGING OF IRNA

INITIATION

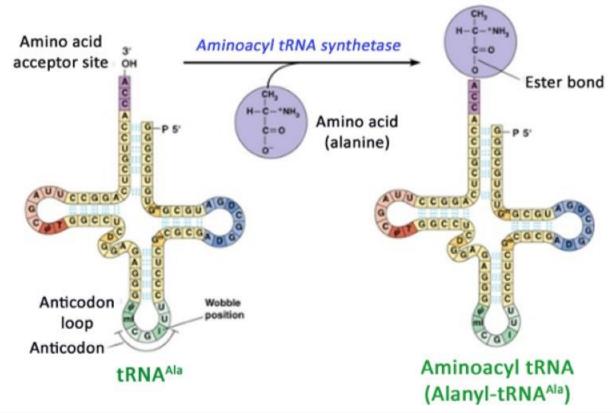
ELONGATION

TERMINATION



1. CHARGING (AMINOACYLATION) OF tRNA

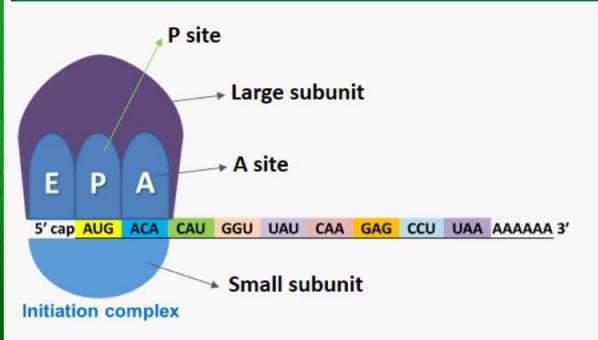


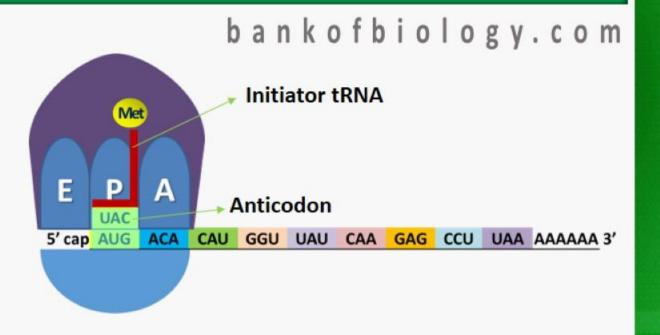


- Formation of peptide bond needs energy obtained from ATP.
- For this, amino acids are activated (amino acid + ATP) and linked to their cognate tRNA in presence of aminoacyl tRNA synthetase. Thus, the tRNA becomes charged.

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2. INITIATION





- In this, small subunit of ribosome binds to mRNA at the start codon (AUG).
- Now large subunit binds to small subunit to form initiation complex.
- Large subunit consists of aminoacyl tRNA binding site (A site) and peptidyl site (P site).

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 The initiator tRNA (which carries methionine) binds on P site. Its anticodon (UAC) recognizes start codon AUG.

3. ELONGATION

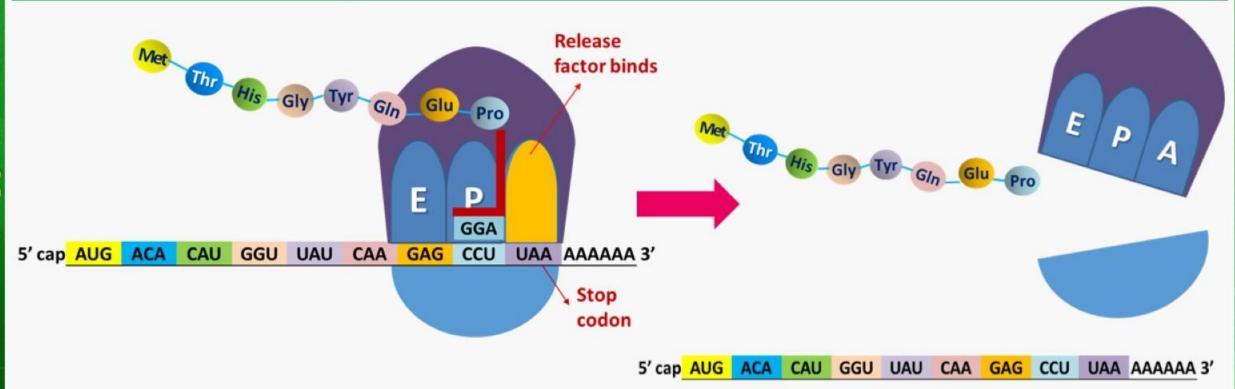


- Second aminoacyl tRNA binds to the A site of ribosome. Its anticodon binds to the second codon on the mRNA and a peptide bond is formed between first and second amino acids in presence of peptidyl transferase.
- First amino acid and its tRNA are broken. This tRNA is removed from P site and second tRNA from A site is pulled to P site along with mRNA. This is called translocation.

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- These processes are repeated for other codons in mRNA.
- During translation, ribosome moves from codon to codon.

4. TERMINATION

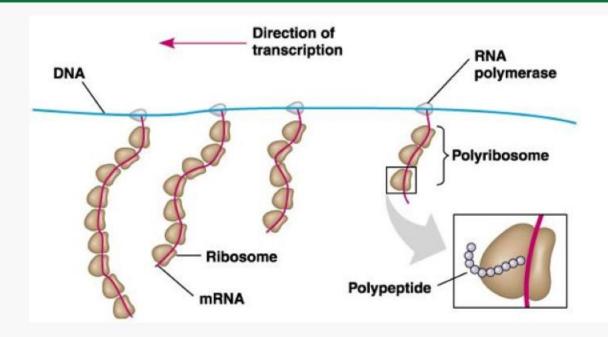


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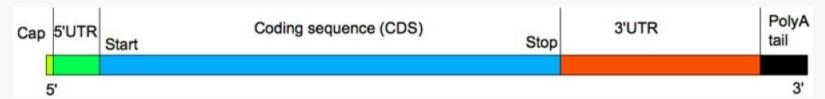
- When a release factor binds to the stop codon, the translation terminates.
- The polypeptide and tRNA are released from the ribosomes.
- The ribosome dissociates into large and small subunits.

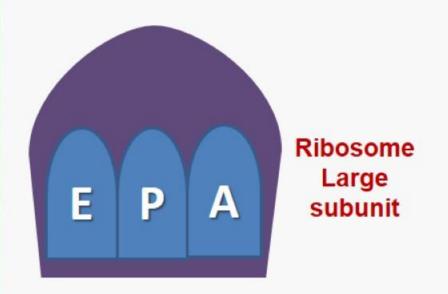
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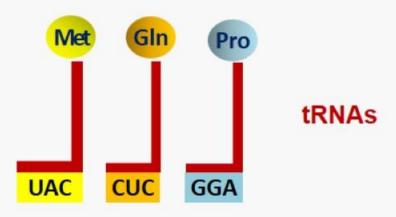
 A group of ribosomes associated with a single mRNA for translation is called a polyribosome (polysomes).



 An mRNA has additional sequences that are not translated (untranslated regions or UTR). UTRs are present at both 5'-end (before start codon) and 3'-end (after stop codon). They are required for efficient translation process.







5' cap AUG ACA CAU GGU UAU CAA GAG CCU UAA AAAAAA 3' mRNA





5' cap AUG ACA CAU GGU UAU CAA GAG CCU UAA AAAAAA 3'
MRNA

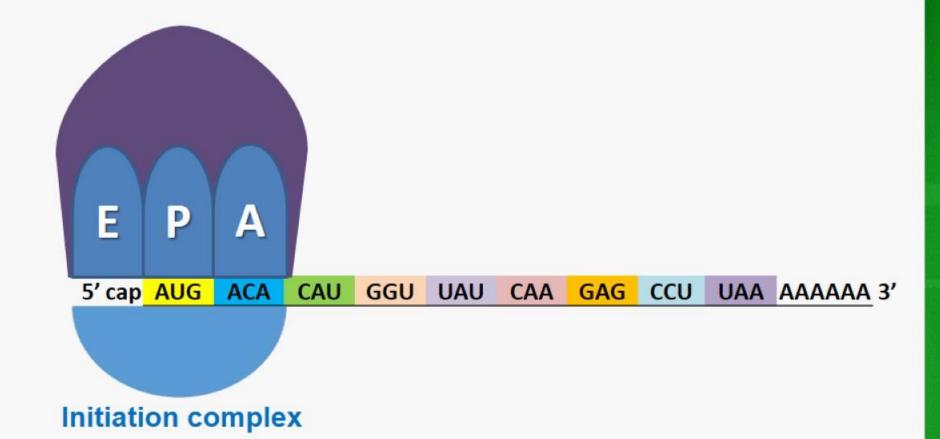
1.Initiation





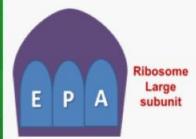




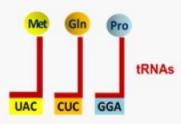


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5' cap AUG ACA CAU GGU UAU CAA GAG CCU UAA AAAAAA 3' mRNA



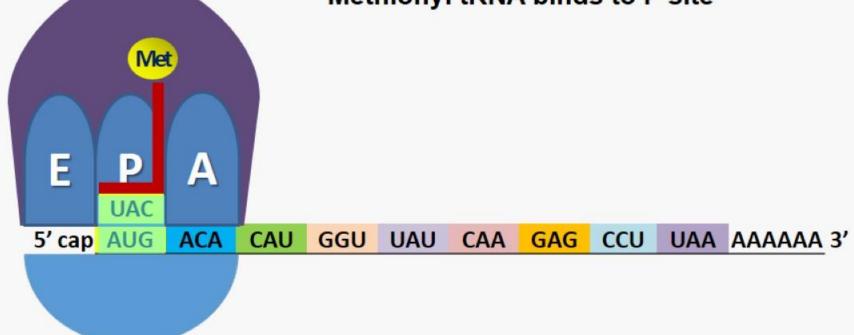






1.Initiation

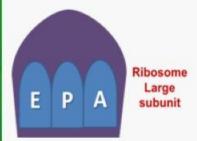
Methionyl tRNA binds to P site



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5' cap AUG ACA CAU GGU UAU CAA GAG CCU UAA AAAAAA 3' mRNA

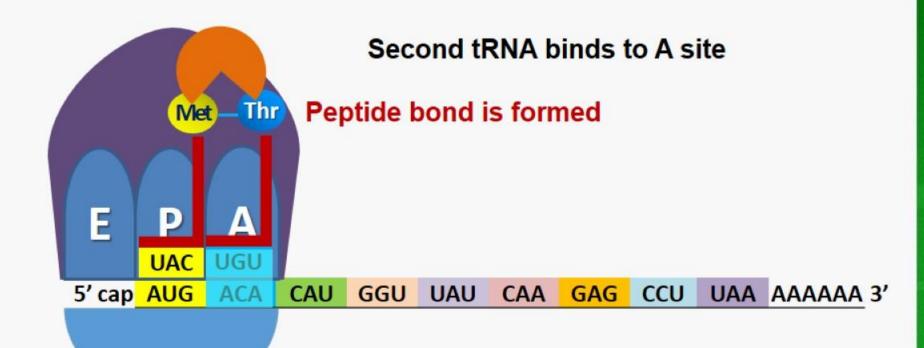








2. Elongation



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5' cap AUG ACA CAU GGU UAU CAA GAG CCU UAA AAAAAA 3' mRNA



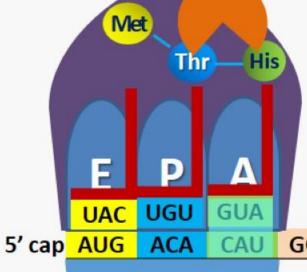






2. Elongation

Translocation from A site to P site



GGU UAU CAA GAG CCU UAA AAAAAA 3'



ACA

CAU

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GGU

UAU

GAG

CAA

CCU

5' cap AUG

UAA AAAAAA 3'



tRNAs



CAU

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GGU

UAU

CAA

GAG

5' cap AUG

GGA

CCU

UAA

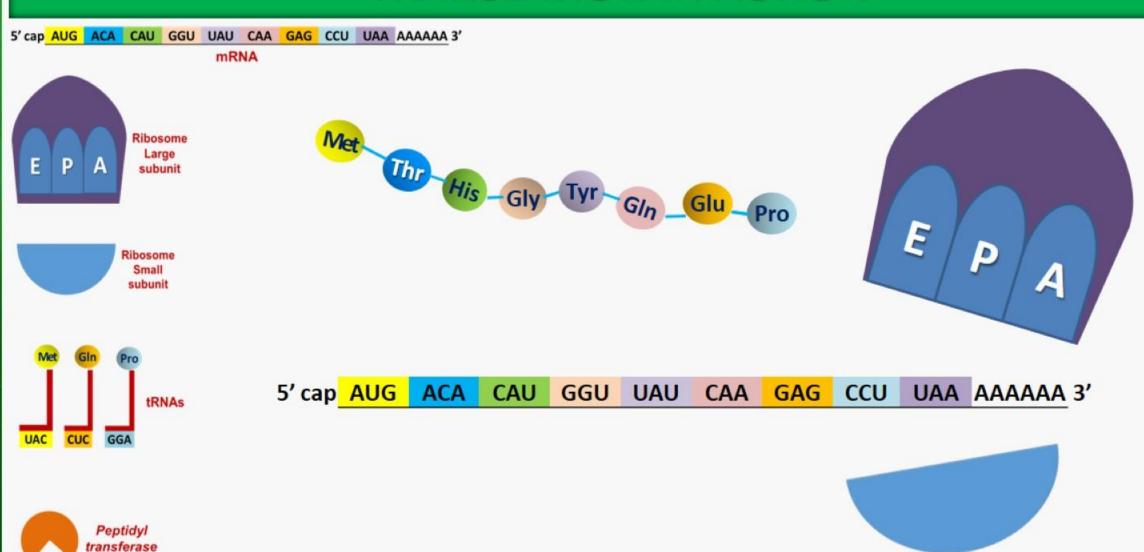
AAAAAA 3'

Stop

codon



tRNAs

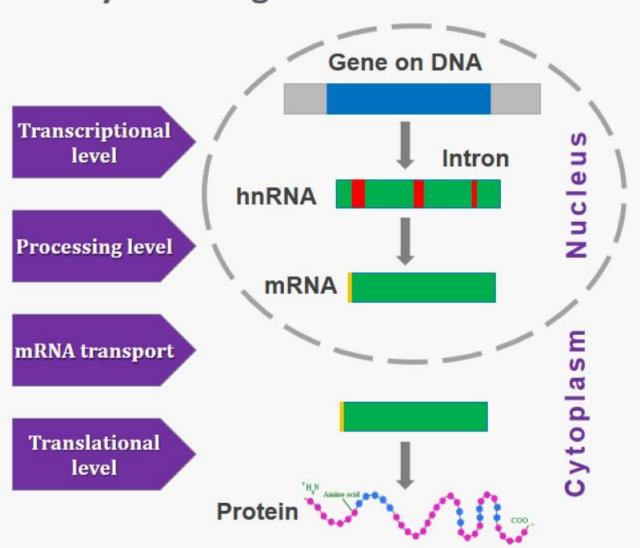






In eukaryotes, gene expression occurs by following levels:

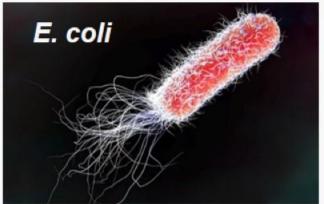
- Transcriptional level (formation of primary transcript).
- 2. Processing level (splicing, capping etc.).
- Transport of mRNA from nucleus to the cytoplasm.
- 4. Translational level (formation of a polypeptide).



The metabolic, physiological and environmental conditions regulate the gene expression. E.g.

- ✓ In E. coli, the beta-galactosidase enzyme hydrolyses lactose into galactose and glucose. In the absence of lactose, the synthesis of beta-galactosidase stops.
- ✓ The development and differentiation of embryo into adult are a result of the expression of several set of genes.







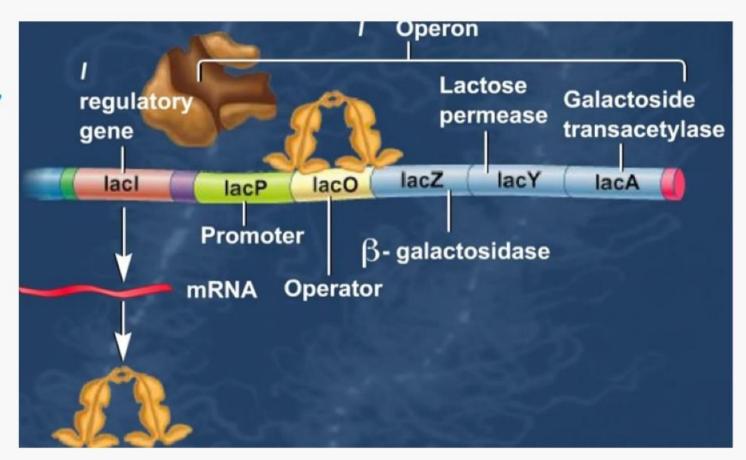
- If a substrate is added to growth medium of bacteria, a set of genes is switched on to metabolize it. It is called induction.
- When a metabolite (product) is added, the genes to produce it are turned off. This is called repression.



E. Coli growth medium

OPERON CONCEPT

- "Each metabolic reaction is controlled by a set of genes."
- All the genes regulating a metabolic reaction constitute an Operon.
- E.g. lac operon, trp operon, ara operon, his operon, val operon etc.

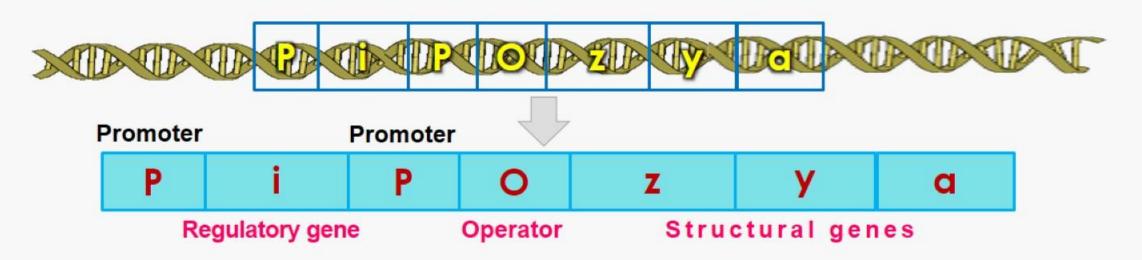


LAC OPERON

- Lac operon is the operon controlling lactose metabolism.
- It is proposed by Francois
 Jacob & Jacque Monod.



LAC OPERON

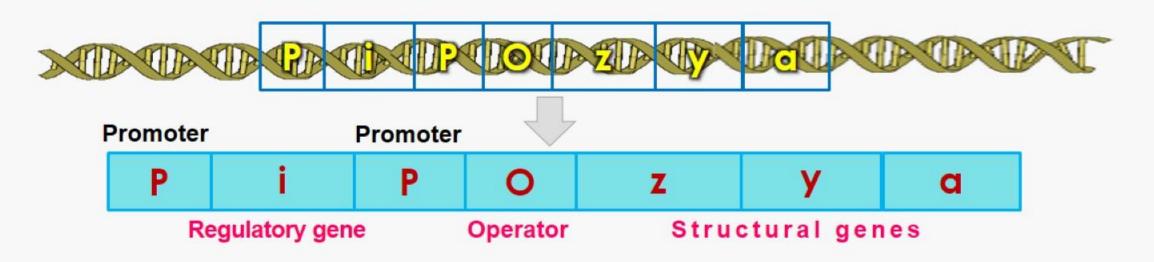


Lac operon consists of

- a. A regulatory or inhibitor (i) gene: Codes for repressor protein.
- b. 3 structural genes: z gene, y gene and a gene.
- c. An Operator region.

Genes in the operon function together in the same or related metabolic pathway.

LAC OPERON

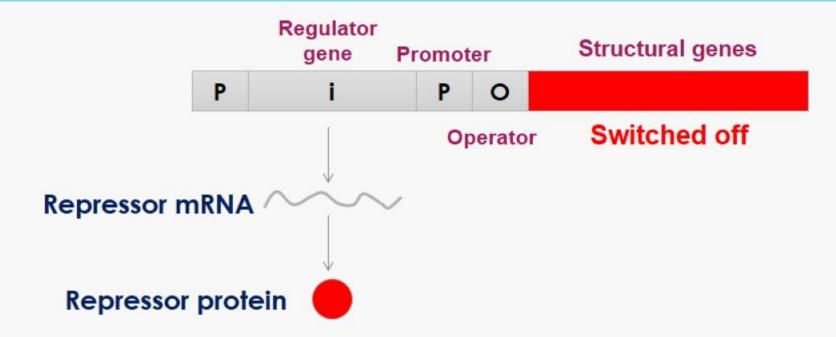


Role of structural genes:

- i. z gene: Codes for β galactosidase. It hydrolyzes lactose to galactose & glucose.
- ii. y gene: Codes for permease. It increases permeability of the cell to β -galactosides (lactose).
- iii. a gene: Codes for a transacetylase.

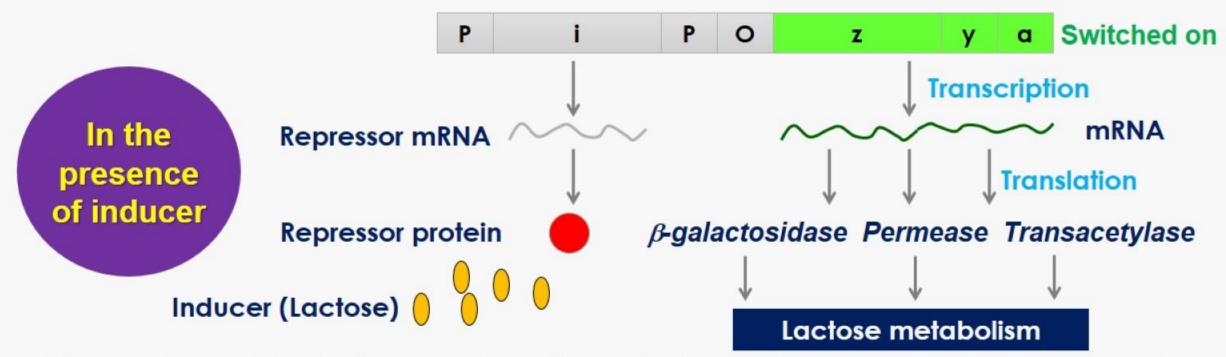
LAC OPERON

In the absence of inducer



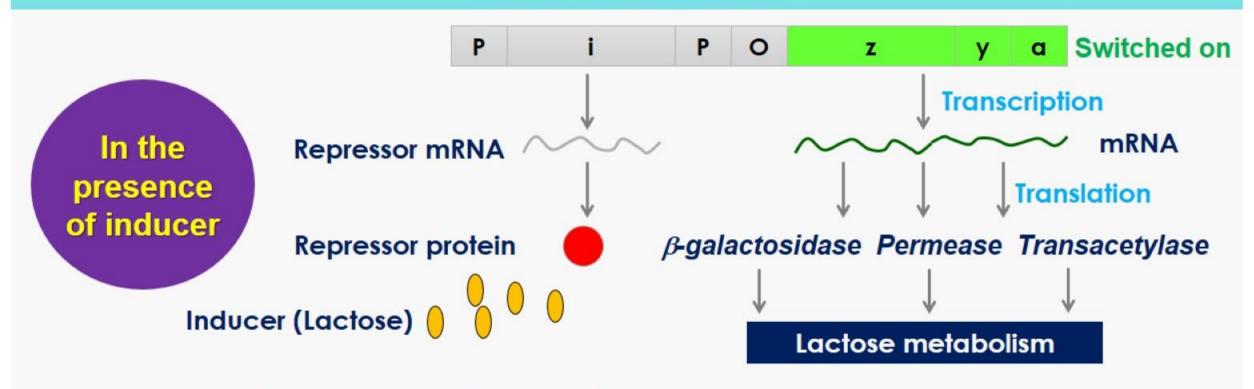
- If there is no lactose (inducer), lac operon remains switched off.
- The regulator gene synthesizes mRNA to produce the repressor protein. This
 protein binds to the operator region and blocks RNA polymerase movement. So
 the structural genes are not expressed.

LAC OPERON



• If lactose is provided in the growth medium, the lactose is transported into the E. coli cells by the action of permease. Lactose (inducer) binds with repressor protein. So repressor protein cannot bind to operator gene. The operator region becomes free and induces the RNA polymerase to bind with promoter gene. Then transcription starts.

LAC OPERON



- Glucose or galactose cannot act as inducers for lac operon.
- Regulation of lac operon by repressor is called negative regulation.
- Lac operon is under control of positive regulation as well.