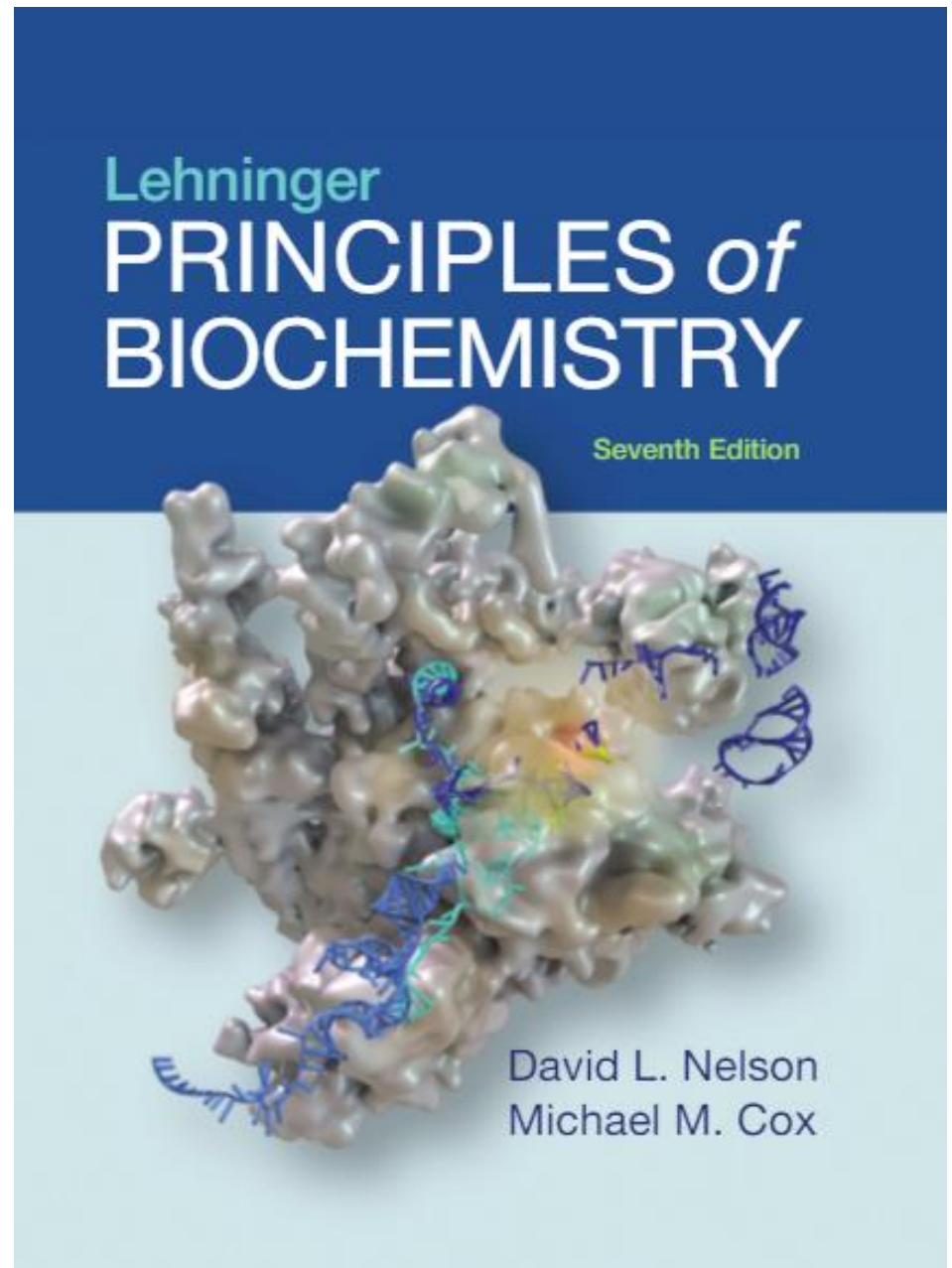


## 28 | Regulation of Gene Expression

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# Ways to Regulate Protein Concentration in a Cell

- Synthesis of primary RNA transcript
- How to process this RNA into mRNA
- Posttranscriptional modifications of mRNA
- Degradation of mRNA
- Protein synthesis
- Posttranslational modification of protein
- Targeting and transport of the protein
- Degradation of the protein
- **Seven Processes That Affect the Steady-State Concentration of a Protein**

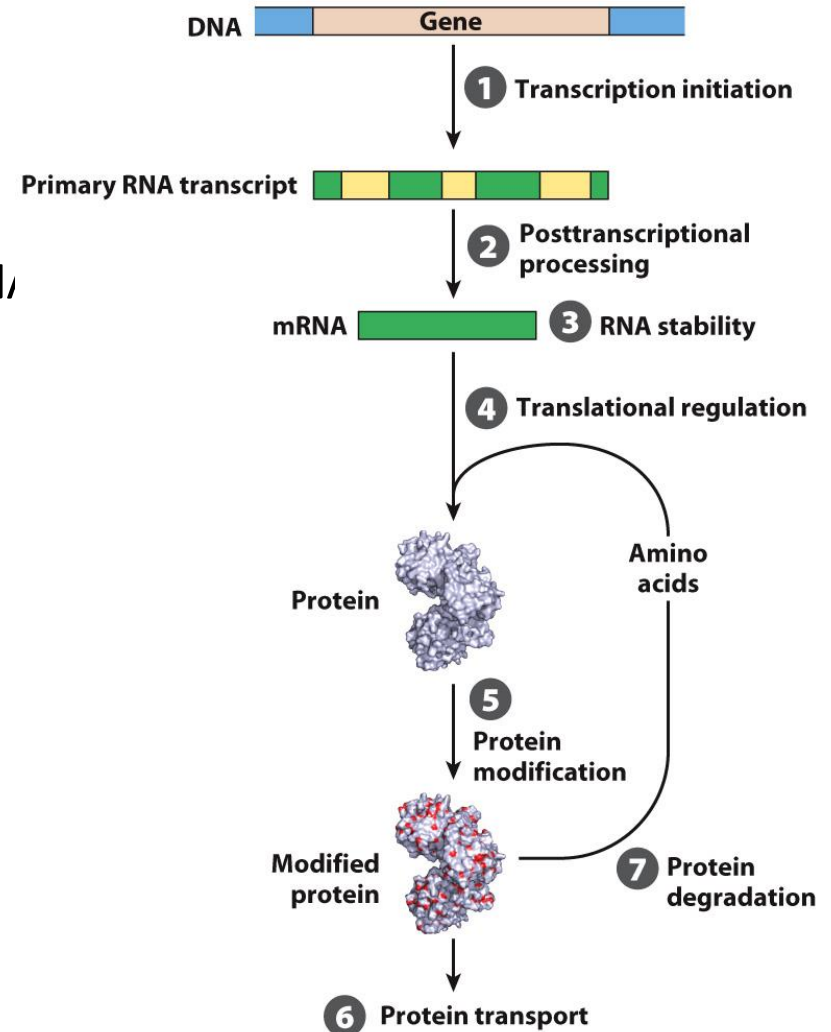


Figure 28-1

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# The Vocabulary of Gene Regulation

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- **Housekeeping gene**
  - under **constitutive expression**
  - constantly expressed in approximately all cells
- **Regulated gene**
  - Levels of the gene product rise and fall with the needs of the organism.
  - Such genes are **inducible**.
    - able to be turned on
  - Such genes are also **repressible**.
    - able to be turned off

# Small-Molecule Effectors Can Regulate Activators and Repressors

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- **Repressors** reduce RNA Pol-promoter interactions or block the polymerase.
  - bind to operator sequences on DNA
    - usually near a promoter in bacteria but further away in many eukaryotes
- **Effectors** can bind to repressor and induce a conformational change.
  - change may increase or decrease repressor's affinity for the operator and thus may increase or decrease transcription

# Activators Improve Contacts Between RNA Polymerase and the Promoter

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- Binding sites in DNA for activators are called **enhancers**.
- In bacteria, enhancers are usually adjacent to the promoter.
  - often adjacent to promoters that are “weak” (bind RNA polymerase weakly), so the activator is necessary
- In eukaryotes, enhancers may be very distant from the promoter.

# Negative Regulation

- **Negative regulation involves repressors.**
  - Example: Repressor binds to DNA and shuts down transcription
  - Alternative: Signal causes repressor to dissociate from DNA; transcription induced

*Despite opposite effects on transcription, both are negative regulation*

**Negative regulation**  
Molecular signal causes dissociation of repressor from DNA, inducing transcription.

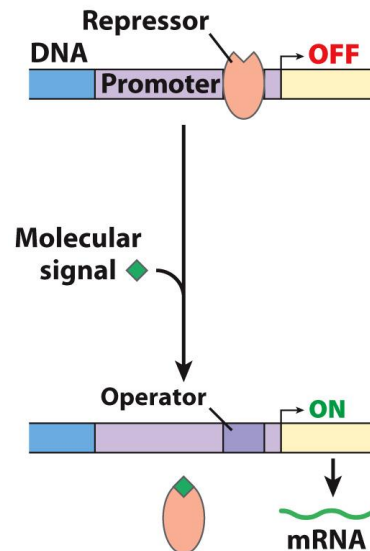


Figure 28-4a  
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**Negative regulation**  
Molecular signal causes binding of repressor to DNA, inhibiting transcription.

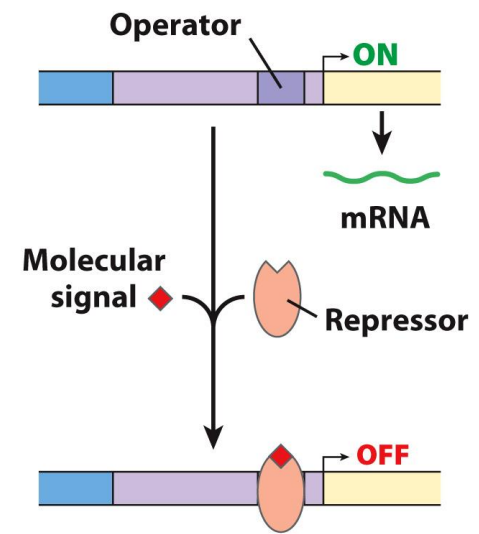


Figure 28-4b  
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# Positive Regulation

- Positive regulation involves **activators**.
- Enhance activity of RNA polymerase
  - Activator-binding sites are near promoters that weakly bind RNA Pol or do not bind at all.
  - It may remain bound until a molecule signals dissociation.
  - Alternatively, the activator may only bind when signaled.

**Positive regulation**  
Molecular signal causes dissociation of activator from DNA, inhibiting transcription.

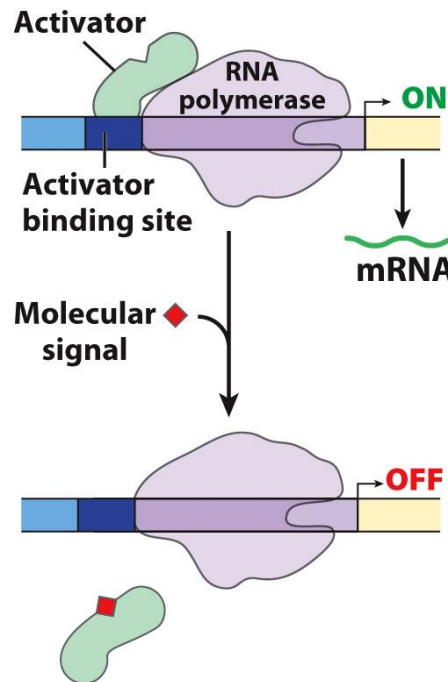


Figure 28-4c  
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**Positive regulation**  
Molecular signal causes binding of activator to DNA, inducing transcription.

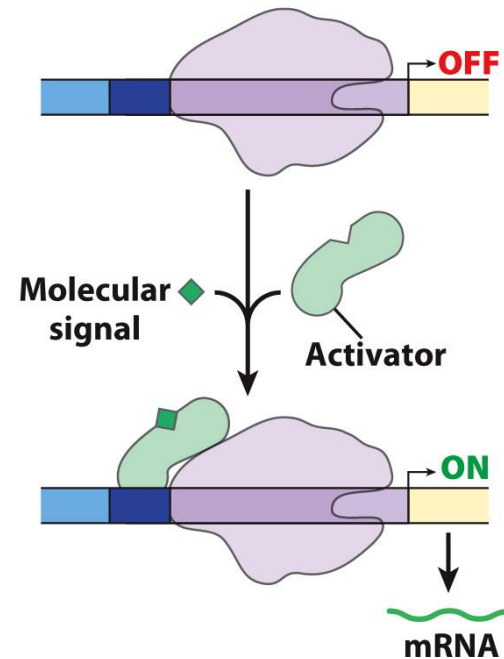
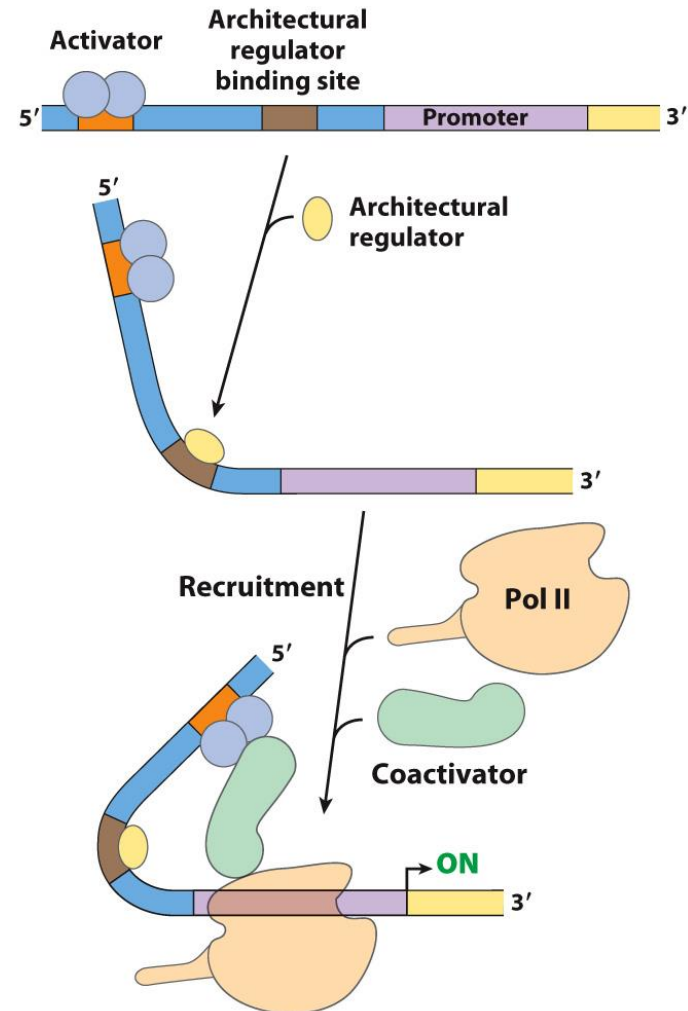


Figure 28-4d  
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# DNA Looping Allows Eukaryotic Enhancers to Be Far from Promoters

- Activators can influence transcription at promoters thousands of bp away.
- How? Via **formation of DNA loops**
- Looping can be facilitated by **architectural regulator proteins**.
- **Co-activators** may mediate binding by binding to both activator and RNA polymerase.



**Figure 28-5**  
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# Many Bacterial Genes Are Transcribed And Regulated Together in an Operon

- An **operon** is a cluster of genes sharing a promoter and regulatory sequences.
  - Genes are transcribed together, so mRNAs are several genes represented on one mRNA (polycistronic).
- First example: the *lac* operon

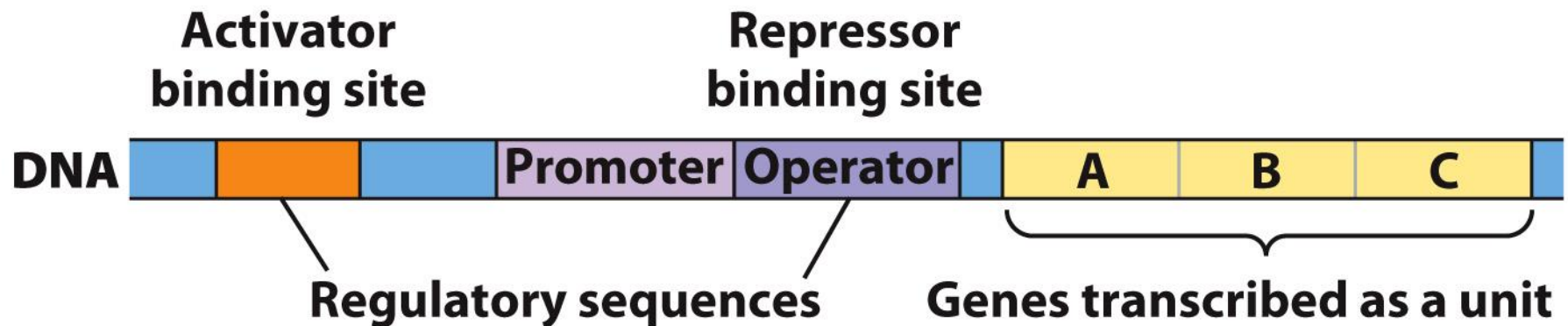


Figure 28-6

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# The *lac* Operon Reveals Many Principles of Gene Regulation

- Work of Jacob and Monod – 1960
- Shows how three genes for metabolism of lactose are regulated together as an operon:
  - $\beta$ -galactosidase (*lacZ*)
    - cleaves lactose to yield glucose and galactose
  - lactose permease (galactoside permease; *lacY*)
    - transports lactose into cell
  - thiogalactoside transacetylase (*lacA*)
- Rely on negative regulation via a repressor.

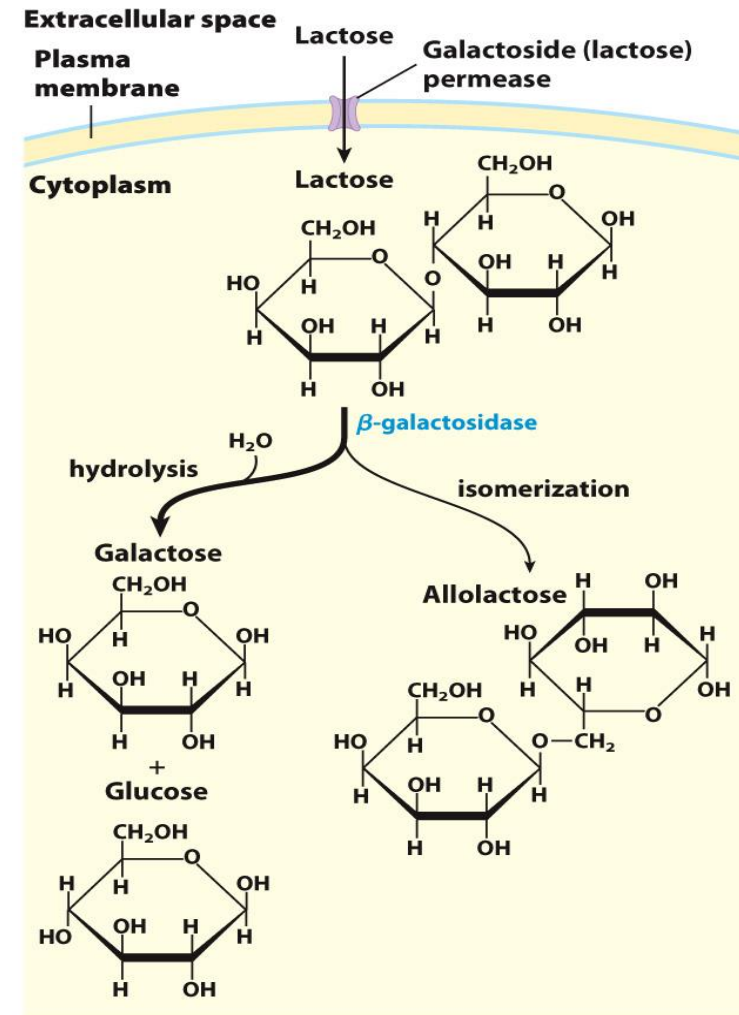


Figure 28-7

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# Lactose Metabolism in *E. Coli*

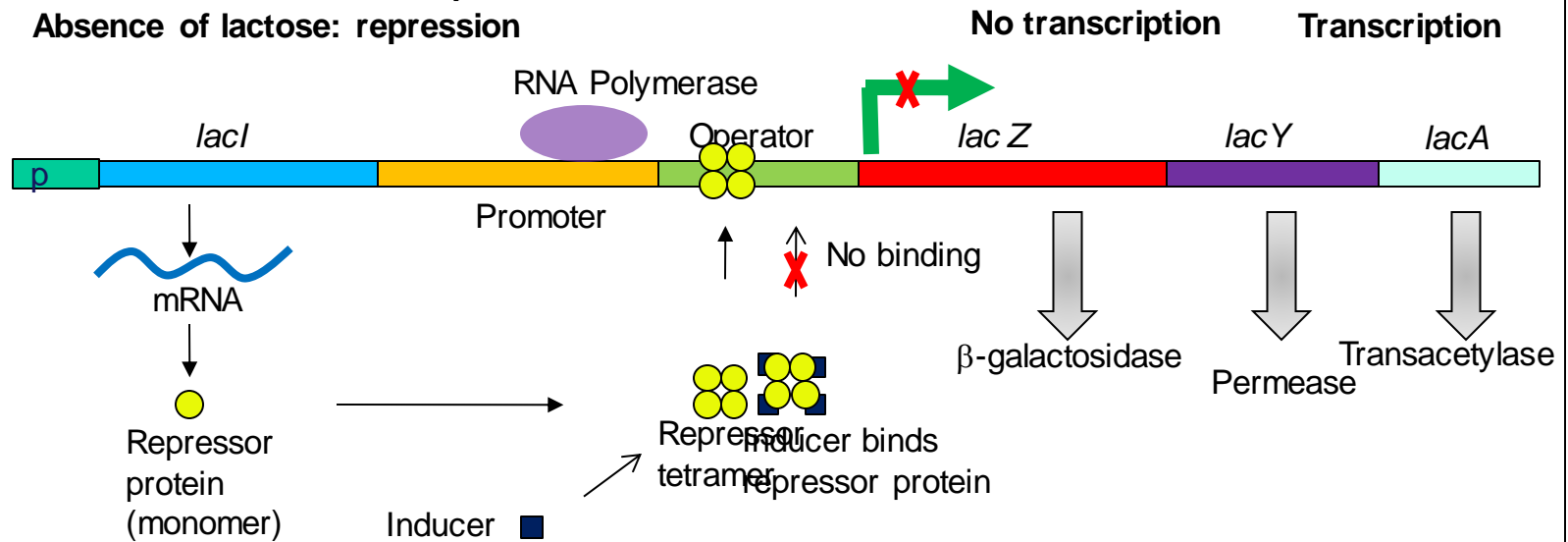
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- When glucose is abundant and lactose is lacking, cells make only very low levels *of enzymes for lactose metabolism*.
  - Transcription is **repressed**.
- If glucose is scarce and cells are **fed lactose**, the cells can use it as their energy source.
- The cells suddenly **express the genes for the enzymes for lactose metabolism**.
  - ***Transcription is no longer repressed.***

# The *lac* operon: negative control by *lacI* repressor

Presence of lactose: de-repression

Absence of lactose: repression



# Inhibiting the Transcription of the *lac* Operon via a Repressor Protein

---

- A gene called *lacI* encodes a repressor called the **Lac repressor**.
  - It has its own promoter  $P_I$ .
    - Transcription of the repressor is independent of transcription of the enzymes the repressor regulates.
  - The repressor can bind to three operator sites ( $O_1$ – $O_3$ ).
- The Lac repressor **binds primarily to the operator  $O_1$** .
  - $O_1$  is adjacent to the promoter.
  - Binding of the repressor helps prevent RNA polymerase from binding to the promoter.
- The repressor also binds to one of two secondary operators, with the **DNA looped between this secondary operator and  $O_1$**  (see Fig. 28-8b).

It reduces transcription, but ***transcription occurs at a low, basal rate***, even with the repressor bound.

# Structure of the *lac* Operon

(a)

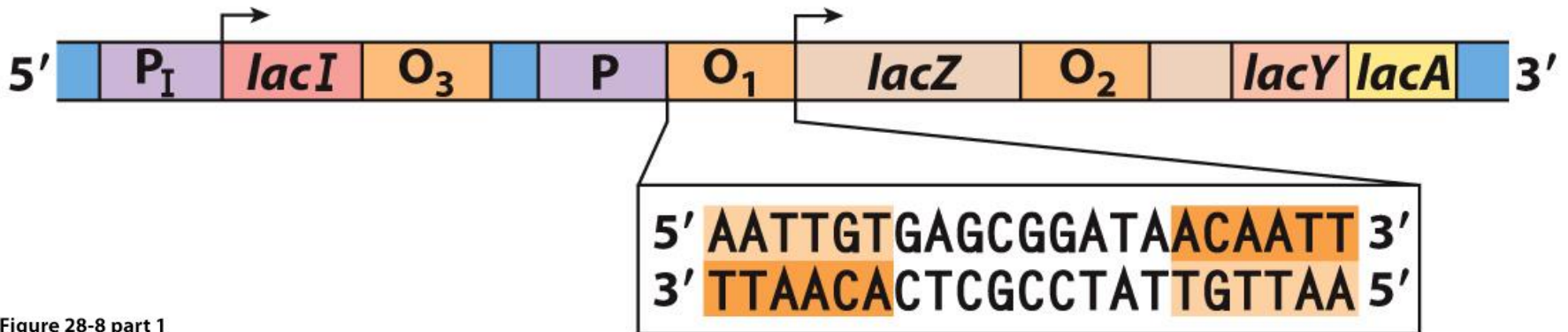


Figure 28-8 part 1

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## Lac Repressor Bound to O<sub>1</sub> and O<sub>3</sub> with DNA Looped Between

(b)

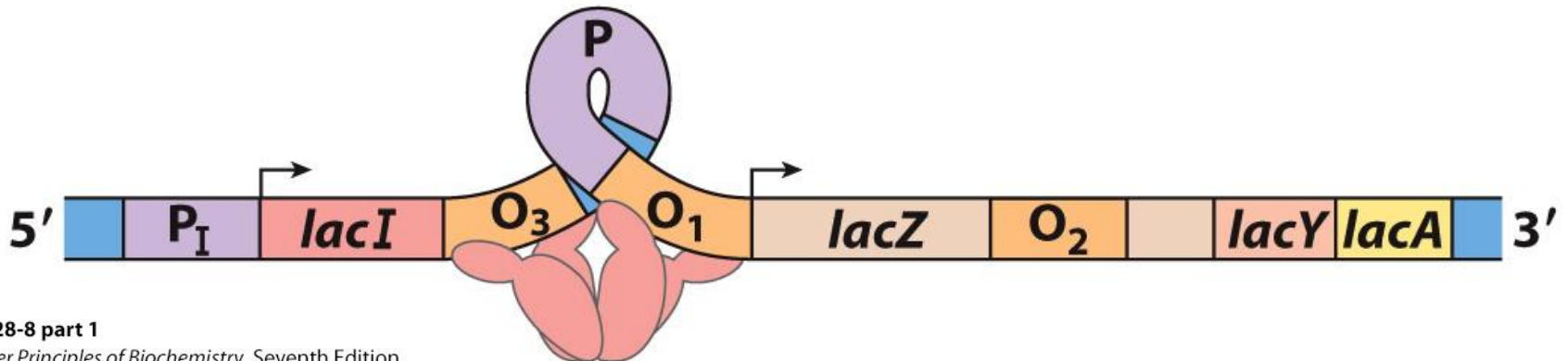


Figure 28-8 part 1

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# How Lac Repressor Binds to DNA

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- Lac repressor is a **tetramer**.
  - **dimer of dimers**
  - Each dimer binds to the palindromic operator sequence.
- The  $O_1$  sequence reflects the symmetry of the repressor.
- There are approximately 20 repressors per cell.

# The *lac* Operon Is Governed by More Than Repressor Binding

---

- The availability of glucose governs expression of lactose-digesting genes via **catabolite repression**.
  - When glucose is present, lactose genes are turned off.
  - It is mediated by cAMP and cAMP receptor protein (CRP or CAP for catabolite activator protein).



# When Glucose Is Absent, *lac* Operon Transcription Is Stimulated by CRP-cAMP

- cAMP binds near the promoter.
  - stimulates transcription 50-fold
  - bends DNA
  - open complex doesn't form readily without CRP-cAMP
- CRP-cAMP only has this effect when the Lac repressor has dissociated.
- cAMP is made when [glucose] is low.

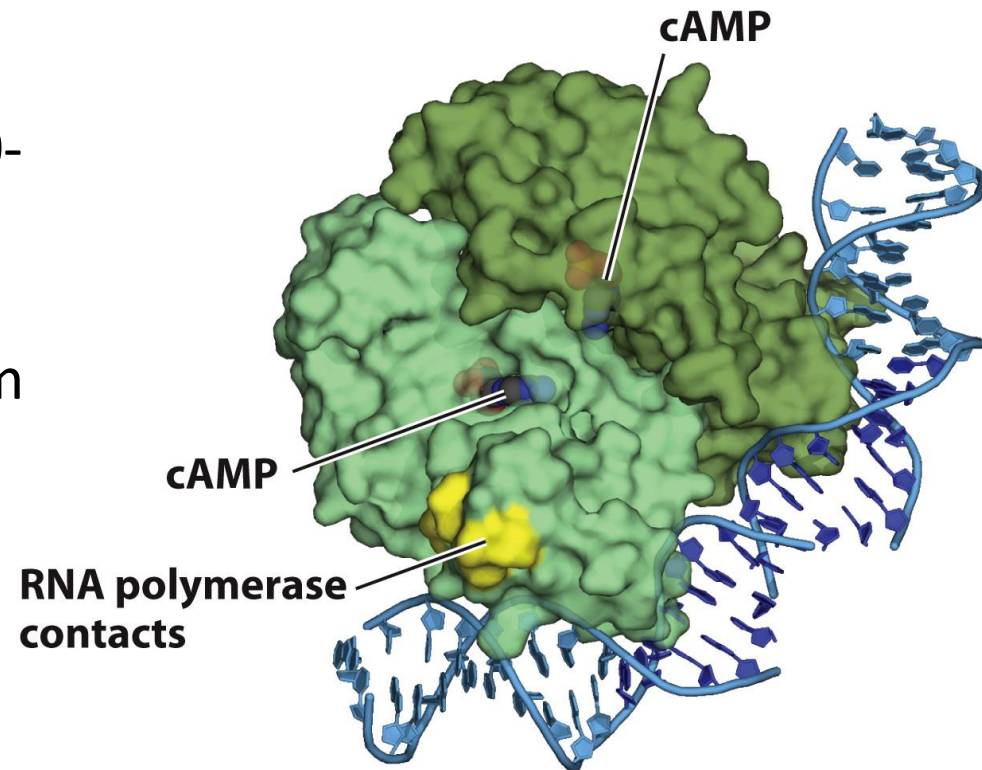


Figure 28-17  
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# When Lactose Is Absent Little to No Transcription Occurs

Whether [glucose] is high or low, if lactose is absent  
→ repressor stays bound....→ no transcription even  
when CRP-cAMP bind.

**Glucose high, cAMP low, lactose absent**

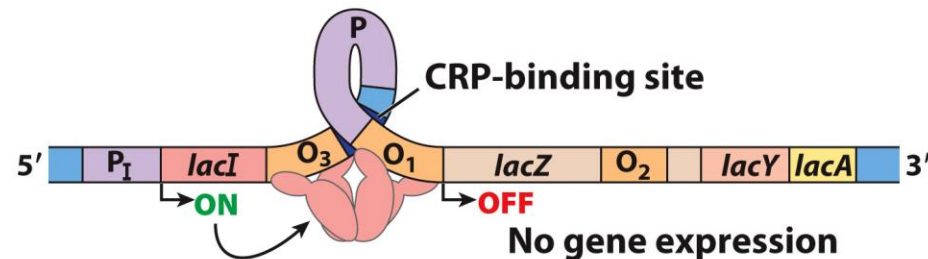


Figure 28-18a  
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**Glucose low, cAMP high, lactose absent**

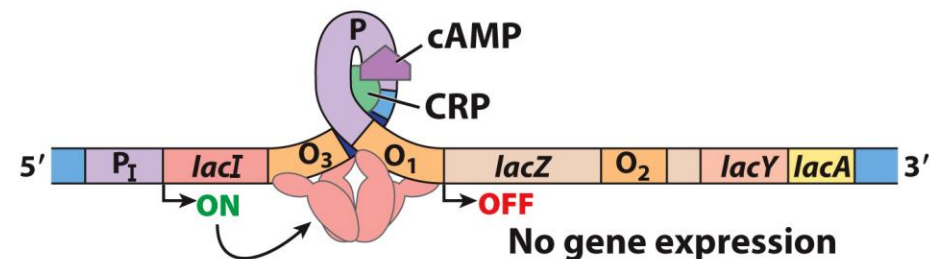


Figure 28-18b  
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# When Lactose Is Present, Transcription Depends On Glucose Level

- Repressor dissociates, but transcription is only stimulated significantly if cAMP rises.

**Glucose high, cAMP low, lactose present**

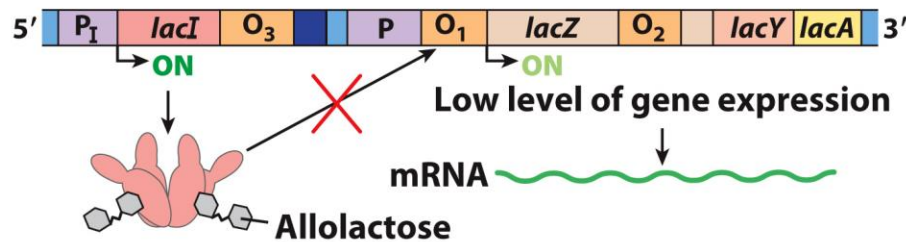


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**Glucose low, cAMP high, lactose present**

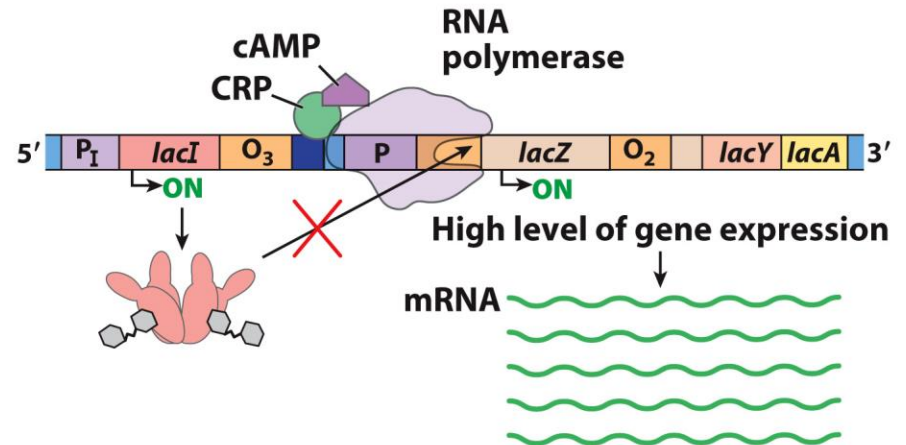


Figure 28-18d  
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# Two Requirements for Strongest Induction of the *lac* Operon

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1. Lactose must be present to form allolactose to bind to the repressor and cause it to dissociate from the operator.
  - reducing repression
2. [Glucose] must be low so that cAMP can increase, bind to CRP, and the complex can bind near the promoter
  - causing activation

# Combined Effects of Glucose and Lactose on the *lac* Operon

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- When **lactose is low**, repressor is bound:  
→ **inhibition**
- When **lactose is high**, repressor dissociates  
→ **permitting transcription**
- When **glucose is high**, CRP is not bound and  
→ **transcription is dampened**
- When **glucose is low**, cAMP is high and CRP is bound  
→ **activation**

# The *trp* Operon

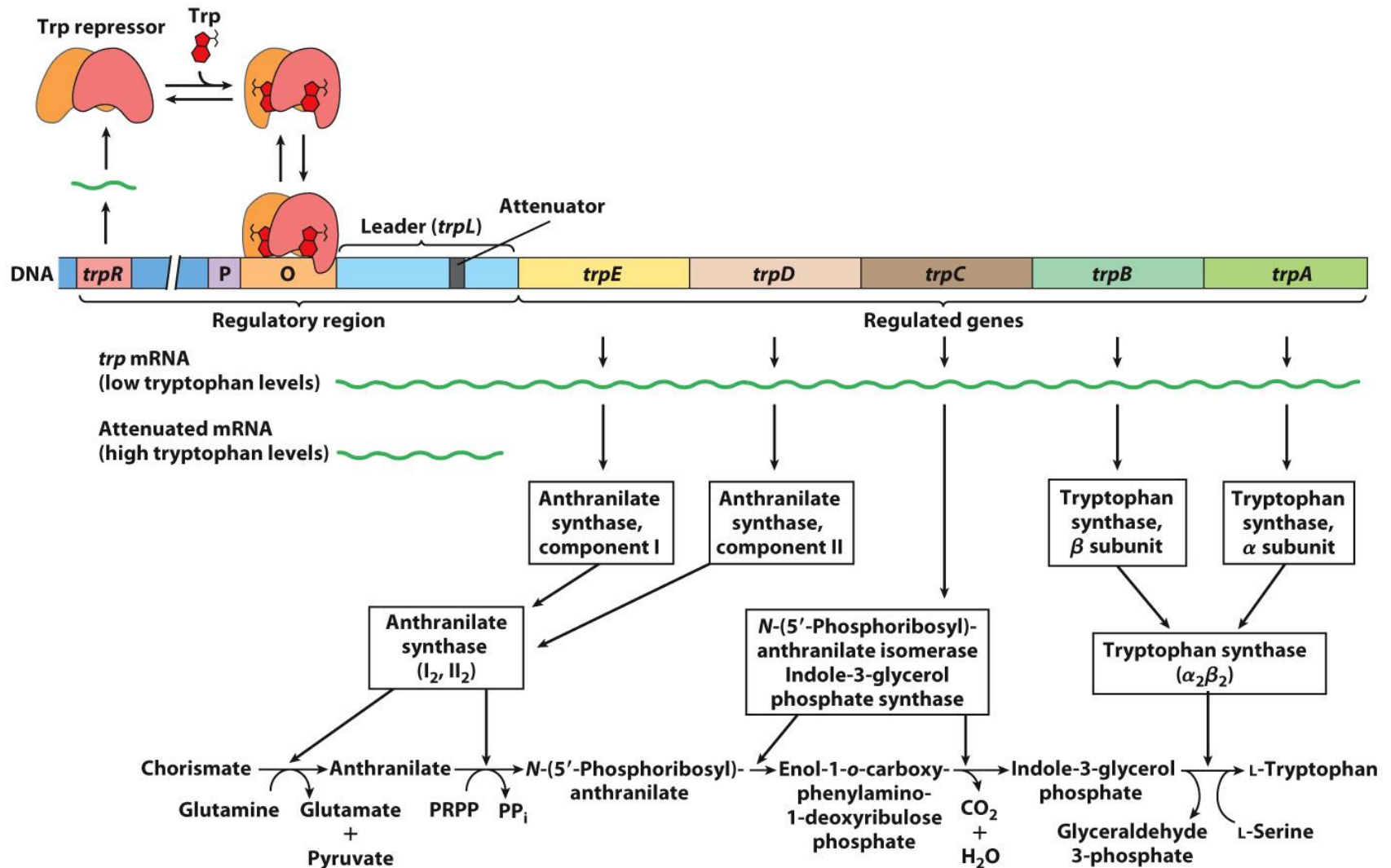


Figure 28-19

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