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Chapter 18: Regulation of Gene Expression**Overview**

The overview for Chapter 18 introduces the idea that while all cells of an organism have all genes in the genome, not all genes are expressed in every cell. What regulates gene expression? Gene expression in prokaryotic cells differs from that in eukaryotic cells. How do disruptions in gene regulation lead to cancer? This chapter gives you a look at how genes are expressed and modulated.

Concept 18.1 Bacteria often respond to environmental change by regulating transcription

1. All genes are not "on" all the time. Using the metabolic needs of *E. coli*, explain why not.

E. coli works by activating a metabolic pathway that makes tryptophan. This only happens when the human does not eat any meat. Once meat is eaten, *E. coli* stops to save its energy for when the meat is gone.

2. What are the two main ways of controlling metabolism in bacterial cells?

- Cells can adjust the activity of enzymes already present with chemical cues through feedback inhibition
- Cells can adjust the production level of the genes encoding the enzymes

3. *Feedback inhibition* is a recurring mechanism throughout biological systems. In the case of *E. coli* regulating tryptophan synthesis, is it *positive* or *negative inhibition*? Explain your choice.

It is *negative inhibition* because the accumulation of the product stops the enzyme from catalyzing it.

4. What is a *promoter*?

A *promoter* is a sequence of DNA nucleotides where RNA polymerase binds to in order to start transcription.

5. What is the *operator*? What does it do?

An *operator* is a sequence of nucleotides near the start of an operon where an active repressor can attach. It prevents RNA polymerase from attaching to the promoter and transcribing the genes of the operon.

6. What is an *operon*?

An *operon* is a unit of genetic function found in bacteria and phages. It includes a promoter, an operator, and the genes they control.

7. List the three components of an *operon*, and explain the role of each one.

- *Promoter*- starts transcription
- *Operator*- On/Off switch- synthesizes enzymes at once
- *Genes they control*- determines whether or not a certain amino acid will be coded for

8. How does a *repressor* protein work?

A *repressor* protein binds to the operator and blocks the attachment of RNA polymerase to the promoter, which prevents transcription of a gene.

9. What are *regulatory genes*?

Regulatory genes code for a protein that controls the transcription of another gene or group of genes.

10. Distinguish between *inducible* and *repressible operons*, and describe one example of each type.

- *Repressible operons* are usually on but can be inhibited when a specific small molecule binds allosterically to a regulatory protein. (ex: tryptophan operon)
- An *inducible operon* is usually off but can be stimulated when a specific small molecule interacts with regulatory proteins. (ex: lactose operon)

Gene Expression And Regulation Study Answers

D Keegan



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Study Guide and Solutions Manual Bruce A. Chase, Peter J. Russell, 2005-06 This student resource contains chapter outlines of text material solutions to all end of chapter problems key terms suggestions for analytical approaches problem solving strategies and a variety of additional questions for student practice Also featured are questions that relate to chapter specific animations and iActivities

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[From Functional Genomics to Biotechnology in Ornamental Plants](#) Swee-Suak Ko, Akira Kanno, Raquel Sánchez-Pérez, Hsin-Hung Yeh, Annette Hohe, Mariana Mondragón-Palomino, 2019-07-12 For centuries plants of a broad taxonomical background have been bred and commercialized because of the beauty of their flowers However until recently genomic analyses of ornamentals remained a challenge because of their large genome sizes and high ploidy levels In the last decade increasingly affordable sequencing technologies and powerful bioinformatic approaches resulted in the complete sequencing of several horticultural species genomes and the characterization of their transcriptomes These developments enabled research on many challenging topics This Research Topic gives you a primer into them by featuring a broad range of original research contributions on some of the most active areas of ornamental plant research the genetic basis of flower morphology scent and color the genetic regulation of physiology as well as the epigenetic factors affecting vegetative development In this context one of the most significant hurdles to functional genetic studies in ornamentals is achieving efficient genetic transformation Several articles in this Research Topic describe strategies to tackle this challenge and present insights into the way transgene activity renders novel flower phenotypes

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