

GLOBAL
EDITION



Lisa A. Urry
Michael L. Cain
Steven A. Wasserman
Peter V. Minorsky
Jane B. Reece

CAMPBELL BIOLOGY IN FOCUS

SECOND EDITION

ALWAYS LEARNING

PEARSON

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Preface

Our goal in writing this text is to focus with high intensity on the core concepts that biology majors need to master in the introductory biology course.

The current explosion of biological information, while exhilarating in its scope, poses a significant challenge—how best to teach a subject that is constantly expanding its boundaries. In particular, instructors have become increasingly concerned that their students are overwhelmed by a growing volume of detail and are losing sight of the big ideas in biology. In response to this challenge, various groups of biologists have initiated efforts to refine and in some cases redesign the introductory biology course. In particular, the report *Vision and Change in Undergraduate Biology Education: A Call to Action** advocates focusing course material and instruction on key ideas while transforming the classroom through active learning and scientific inquiry. Many instructors have embraced such approaches and have changed how they teach. Cutting back on the amount of detail they present, they focus on core biological concepts, explore select examples, and engage in a rich variety of active learning exercises.

We were inspired by these ongoing changes in biology education to write the first edition of *CAMPBELL BIOLOGY IN FOCUS*, a new, shorter textbook that was received with widespread excitement by instructors. Guided by their feedback, we honed the Second Edition so that it does an even better job of helping students explore the key questions, approaches, and ideas of modern biology.

New to This Edition

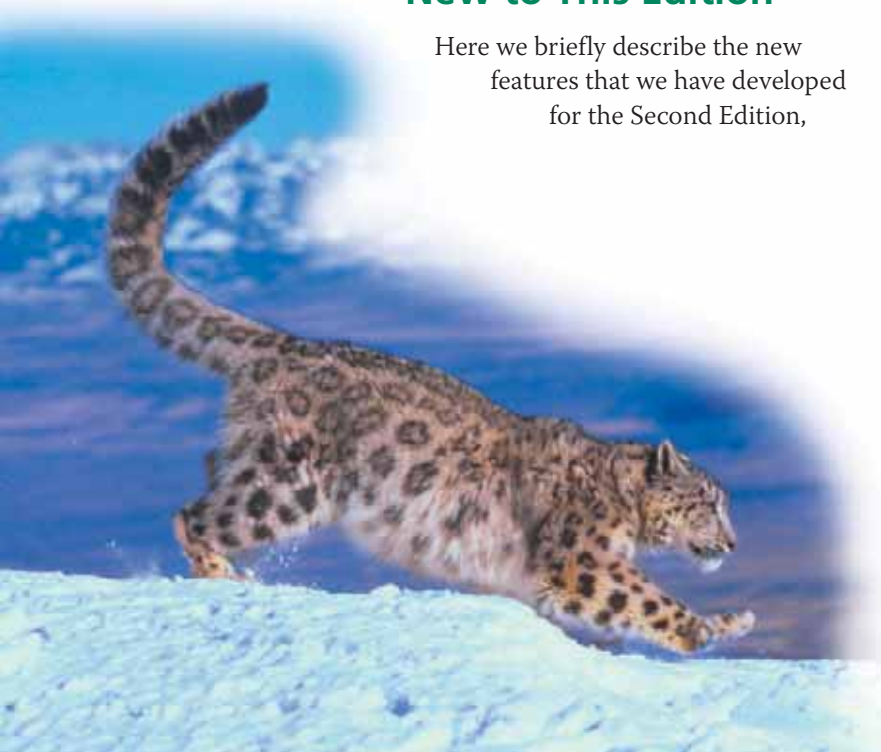
Here we briefly describe the new features that we have developed for the Second Edition,

but we invite you to explore pages xii–xxvi for more information and examples.

New in the Text

- The impact of **genomics** across biology is explored throughout the Second Edition with examples that reveal how our ability to rapidly sequence DNA and proteins on a massive scale is transforming all areas of biology, from molecular and cell biology to phylogenetics, physiology, and ecology. Illustrative examples are distributed throughout the text.
- The Second Edition provides increased coverage of the urgent issue of **global climate change**. Starting with a new figure (Figure 1.11) and discussion in Chapter 1 and concluding with significantly expanded material on causes and effects of climate change in Chapter 43, including a new Make Connections Figure (Figure 43.28), the text explores the impact of climate change at all levels of the biological hierarchy.
- Ten **Make Connections Figures** pull together content from different chapters to assemble a visual representation of “big-picture” relationships. By reinforcing fundamental conceptual connections throughout biology, these figures help overcome students’ tendencies to compartmentalize information.
- **Interpret the Data Questions** throughout the text engage students in scientific inquiry by asking them to analyze data presented in a graph, figure, or table. The Interpret the Data Questions can be assigned and automatically graded in MasteringBiology.®
- **Synthesize Your Knowledge Questions** at the end of each chapter ask students to synthesize the material in the chapter and demonstrate their big-picture understanding. A striking, thought-provoking photograph leads to a question that helps students realize that what they have learned in the chapter connects to their world and provides understanding and insight into natural phenomena.
- Scannable QR codes and URLs at the end of every chapter give students quick access to **Vocabulary Self-Quizzes** and **Practice Tests** that students can use on a smartphone, tablet, or computer.
- Detailed information about the organization of the text and new content in the Second Edition is provided on pages 6–9, following this Preface.

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New in MasteringBiology®

- New **MasteringBiology tutorials** extend the power of MasteringBiology:
 - **Interpret the Data Questions** ask students to analyze a graph, figure, or table.
 - **Solve It Tutorials** engage students in a multistep investigation of a “mystery” or open question in which they must analyze real data.
 - **HHMI Short Films**, documentary-quality movies from the Howard Hughes Medical Institute, engage students in topics from the discovery of the double helix to evolution, with assignable questions.
 - **Video Field Trips** allow students to study ecology by taking virtual field trips and answering follow-up questions.

Our Guiding Principles

Our key objective in creating *CAMPBELL BIOLOGY IN FOCUS* was to produce a shorter text by streamlining selected material, while emphasizing conceptual understanding and maintaining clarity, proper pacing, and rigor. Here, briefly, are the five guiding principles of our approach:

1. Focus on Core Concepts

We developed this text to help students master the fundamental content and scientific skills they need as college biology majors. In structuring the text, we were guided by discussions with biology professors across the country, analysis of hundreds of syllabi, study of the debates in the literature of scientific pedagogy, and our experience as instructors at a range of institutions. The result is a **briefer book for biology majors** that informs, engages, and inspires.

2. Establish Evolution as the Foundation of Biology

Evolution is the central theme of all biology, and it is the core theme of this text, as exemplified by the various ways that evolution is integrated into the text:

- Every chapter explicitly addresses the topic of evolution through an **Evolution section** that leads students to consider the material in the context of natural selection and adaptation.
- Each Chapter Review includes a **Focus on Evolution Question** that asks students to think critically about how an aspect of the chapter relates to evolution.
- Evolution is the unifying idea of **Chapter 1, Introduction: Evolution and the Foundations of Biology**, which devotes Concept 1.2 to the core theme of evolution, providing students with a foundation in evolution early in their study of biology.

- Following the in-depth coverage of **evolutionary mechanisms in Unit 3**, evolution also provides the storyline for the novel approach to presenting biological diversity in **Unit 4, The Evolutionary History of Life**. Focusing on landmark events in the history of life, Unit 4 highlights how key adaptations arose within groups of organisms and how evolutionary events led to the diversity of life on Earth today.

3. Engage Students in Scientific Thinking

Helping students learn to “think like a scientist” is a nearly universal goal of introductory biology courses. Students need to understand how to formulate and test hypotheses, design experiments, and interpret data. Scientific thinking and data interpretation skills top lists of learning outcomes and foundational skills desired for students entering higher-level courses. *CAMPBELL BIOLOGY IN FOCUS*, Second Edition, meets this need in several ways:

- **Scientific Skills Exercises** in every chapter use real data to build skills in graphing, interpreting data, designing experiments, and working with math—skills essential for students to succeed in biology. These exercises can also be assigned and automatically graded in MasteringBiology.
- New **Interpret the Data Questions** ask students to analyze a graph, figure, or table. These questions are also assignable in MasteringBiology.
- **Scientific Inquiry Questions** in the end-of-chapter material give students further practice in scientific thinking.
- **Inquiry Figures** and **Research Method Figures** reveal *how* we know *what* we know and model the process of scientific inquiry.

4. Use Outstanding Pedagogy to Help Students Learn

CAMPBELL BIOLOGY IN FOCUS, Second Edition, builds on our hallmarks of clear and engaging text and superior pedagogy to promote student learning:

- In each chapter, a framework of carefully selected **Key Concepts** helps students distinguish the “forest” from the “trees.”
- Questions throughout the text catalyze learning by encouraging students to **actively engage with and synthesize key material**. Active learning questions include Concept Check Questions, Make Connections Questions, What If? Questions, Figure Legend Questions, Draw It Exercises, Summary Questions, and the new Synthesize Your Knowledge and Interpret the Data Questions.

- **Test Your Understanding Questions** at the end of each chapter are organized into three levels based on **Bloom's Taxonomy**.

5. Create Art and Animations That Teach

Biology is a visual science, and students learn from the art as much as the text. Therefore, we have developed our art and animations to teach with clarity and focus. Here are some of the ways our art and animations serve as superior teaching tools:

- The ten new **Make Connections Figures** help students see connections between topics across the entire introductory biology course.
- Each unit in *CAMPBELL BIOLOGY IN FOCUS*, Second Edition, opens with a **visual preview** that tells the story of the chapters' contents, showing how the material in the unit fits into a larger context.
- **BioFlix® 3-D Animations** help students visualize biology with movie-quality animations that can be shown in class and reviewed by students in the Study Area. **BioFlix Tutorials** use the animations as a jumping-off point for MasteringBiology coaching assignments with feedback.
- By integrating text, art, and photos, **Exploring Figures** help students access information efficiently.
- **Guided Tour Figures** use descriptions in blue type to walk students through complex figures as an instructor would, pointing out key structures, functions, and steps of processes.
- Because text and illustrations are equally important for learning biology, the **page layouts** are carefully designed to place figures together with their discussions in the text.
- **PowerPoint®** slides are painstakingly developed for optimum presentation in lecture halls, with enlarged editable labels, art broken into steps, and links to animations and videos.
- Many **Tutorials** and **Activities** in MasteringBiology integrate art from the text, providing a unified learning experience.

MasteringBiology®

MasteringBiology is the most widely used online assessment and tutorial program for biology, providing an extensive library of homework assignments that are graded automatically. **Self-paced tutorials provide individualized coaching with specific hints and feedback** on the most difficult topics in the course. In addition to the new tutorials already mentioned, MasteringBiology includes hundreds of online exercises that can be assigned. For example:

- The **Scientific Skills Exercises** from the text can be assigned and automatically graded in MasteringBiology.
- **BioFlix® Tutorials** use 3-D animations to help students master tough topics.
- **Make Connections Tutorials** help students connect what they are learning in one chapter with material they have learned in another chapter.

- **BLAST Data Analysis Tutorials** teach students how to work with real data from the BLAST database.
- **Experimental Inquiry Tutorials** allow students to replicate a classic biology experiment and learn the conceptual aspects of experimental design.
- **Reading Quiz Questions** and approximately 3,000 **Test Bank Questions** are available for assignment.
- Optional **Adaptive Follow-up Assignments** are based on each student's performance on the original MasteringBiology assignment and provide additional coaching and practice as needed.

Every assignment is automatically graded and entered into a **gradebook**. Instructors can check the gradebook to see what topics students are struggling with and then address those topics in class.

The following resources are also available in MasteringBiology:

- The **Instructor Resources** area provides everything needed to teach the course.
- **Learning Catalytics™** allows students to use their smartphones, tablets, or laptops to respond to questions in class.
- **Dynamic Study Modules** provide students with multiple sets of questions with extensive feedback so that they can test, learn, and retest until they achieve mastery of the textbook material. Students can use these modules on their smartphones on their own or the modules can be assigned.
- Students can read the **eText** and use the self-study resources in the **Study Area**.

MasteringBiology and the text work together to provide an unparalleled learning experience. For more information about MasteringBiology, see pages 15–16 and 20–24.

* * *

Our overall goal in developing and revising this text was to assist instructors and students in their exploration of biology by emphasizing essential content and skills while maintaining rigor. Although this Second Edition is now completed, we recognize that *CAMPBELL BIOLOGY IN FOCUS*, like its subject, will evolve. As its authors, we are eager to hear your thoughts, questions, comments, and suggestions for improvement. We are counting on you—our teaching colleagues and all students using this book—to provide us with this feedback, and we encourage you to contact us directly by e-mail:

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Organization and New Content

CAMPBELL BIOLOGY IN FOCUS, Second Edition, is organized into an introductory chapter and seven units that cover core concepts of biology at a thoughtful pace. When we adapted *CAMPBELL BIOLOGY* to write the first edition of this text, we made informed choices about how to design each chapter of *CAMPBELL BIOLOGY IN FOCUS* to meet the needs of instructors and students. In some chapters, we retained most of the material; in other chapters, we pruned material; and in still others, we completely reconfigured the material. In creating the Second Edition, we solicited feedback from reviewers and used their thoughtful critiques to further fine-tune the content and pedagogy. We have also updated the content wherever appropriate, and in a few cases reintroduced material. Here, we present synopses of the seven units and highlight the major revisions made to the Second Edition of *CAMPBELL BIOLOGY IN FOCUS*.

CHAPTER 1 Introduction: Evolution and the Foundations of Biology

Chapter 1 introduces the **five biological themes** woven throughout the text: the core theme of **Evolution**, together with **Organization, Information, Energy and Matter**, and **Interactions**. Chapter 1 also explores the process of scientific inquiry through a case study describing experiments on the evolution of coat color in the beach mouse.

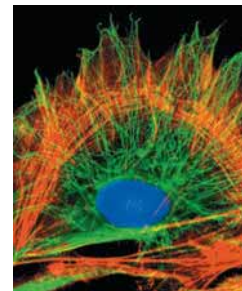


The chapter concludes with a discussion of the importance of diversity within the scientific community.

In the Second Edition, a new figure (Figure 1.8) on gene expression uses lens cells in the eye as an example of DNA → RNA → protein and introduces the terms transcription and translation. This new figure and text equip students from the outset with an understanding of how gene sequences determine an organism's characteristics. New text and a new photo (Figure 1.11) inform students about the effects of climate change in general, and global warming in particular, on species survival and diversity. Concept 1.3 has been thoroughly revised to more realistically reflect the process of science. A new section has been added on the Flexibility of the Scientific Process, accompanied by a new Figure 1.19 that depicts the more realistic and complex process of science. The text now discusses searching the scientific literature, and a new question in the Chapter Review asks students to use PubMed.

UNIT 1 Chemistry and Cells

A succinct, two-chapter treatment of basic chemistry (Chapters 2 and 3) provides the foundation for this unit focused on cell structure and function. The related topics of cell membranes and cell signaling are consolidated into one chapter (Chapter 5). Due to the importance of the fundamental concepts in Units 1 and 2, much of the material in the rest of these two units has been retained from *CAMPBELL BIOLOGY*.



For the Second Edition, a new table has been added to Chapter 2 detailing the elements in the human body, with an associated Interpret the Data question. Chapter 3 includes a new section on isomers, with an accompanying figure (Figure 3.5), and ends with a new Concept 3.7 that includes cutting-edge coverage of DNA sequencing and introduces genomics and proteomics, as well as bioinformatics. A new Make Connections Figure (Figure 3.30) entitled “Contributions of Genomics and Proteomics to Biology” provides an overview of areas in which genomics and proteomics have had significant impacts—including evolution, conservation biology, paleontology, medical science, and species interactions—with the aim of inspiring and motivating students. A striking photo of thermophilic cyanobacteria has been added to Figure 6.16 on environmental factors affecting enzyme activity. In Chapter 7, a computer model of ATP synthase has been added to Figure 7.13. The icon for this enzyme in Chapters 7 and 8 has been re-drawn to more closely represent its structure. A new Make Connections Figure (Figure 8.20, “The Working Cell”) integrates all the cellular activities covered in Chapters 3–8 in the context of a single working plant cell.

UNIT 2 Genetics

Topics in this unit include meiosis and classical genetics as well as the chromosomal and molecular basis for genetics and gene expression (Chapters 10–14). We also include a chapter on the regulation of gene expression (Chapter 15) and one on the role of gene regulation in development, stem cells, and cancer (Chapter 16). Methods in biotechnology



are integrated into appropriate chapters. The stand-alone chapter on viruses (Chapter 17) can be taught at any point in the course. The final chapter in the unit, on genome evolution (Chapter 18), provides both a capstone for the study of genetics and a bridge to the evolution unit.

Chapter 10 of the Second Edition includes a new section on “Crossing Over and Synapsis During Prophase I” that explains the events of prophase I in more detail, supported by new Figure 10.9, which clearly shows and describes these events. In Chapter 11, to incorporate more molecular biology into the discussion of Mendelian genetics, Figure 11.4 on alleles has been enhanced and a new Figure 11.16 on sickle-cell disease has been added. Chapter 13 includes new text and two new figures (Figures 13.29 and 13.30) covering advances in sequencing technology. Also in this chapter, a new section, including new Figure 13.31, describes gene editing using the CRISPR-Cas9 system. In Chapter 15, the section on noncoding RNAs has been updated, and Figure 15.14 on *in situ* hybridization has been expanded and enhanced to help students understand this important technique. Chapter 16 includes a new Inquiry Figure (Figure 16.16) on induced pluripotent stem cells (iPS cells). Material on embryonic stem cells and induced pluripotent stem cells has been significantly updated. A new Make Connections Figure (Figure 16.21), “Genomics, Cell Signaling, and Cancer,” illustrates recent research on subtypes of breast cancer, connecting content that students have learned in Chapters 5, 9, and 16. It also addresses treatment for one subtype of breast cancer as an example. In Chapter 17, the discussion of the importance of cell-surface proteins in determining host range has been enhanced. A new figure (Figure 17.9) presents the example of the receptor and co-receptor proteins for HIV. Coverage of the CRISPR system, as a bacterial “immune” system, has been added, supported by new Figure 17.6. Coverage of recent epidemics has been inserted (Ebola) or updated (H5N1). Chapter 18 has been significantly updated to reflect recent sequencing advances, including a discussion of the results of the ENCODE project, information on the bonobo genome, and use of high-throughput techniques to address the problem of cancer. Regarding protein structure, the discussion of BLAST searches has been enhanced, and computer models of lysozyme and α -lactalbumin have been added to support the discussion of the evolution of genes with novel functions.

UNIT 3 Evolution

This unit provides in-depth coverage of essential evolutionary topics, such as mechanisms of natural selection, population genetics, and speciation. Early in the unit, Chapter 20 introduces “tree thinking” to support students in interpreting phylogenetic trees and thinking about the big picture of evolution.



Chapter 23 focuses on mechanisms that have influenced long-term patterns of evolutionary change. Throughout the unit, new discoveries in fields ranging from paleontology to phylogenomics highlight the interdisciplinary nature of modern biology.

Revisions in the Second Edition aim to strengthen connections among fundamental evolutionary concepts. For example, Concept 20.5 includes new text on horizontal gene transfer among eukaryotes, reinforcing the overall discussion of how horizontal gene transfer has played an important role in the evolutionary history of life. Also in Concept 20.5, a new Scientific Skills Exercise walks students through the process of comparing and interpreting amino acid sequences to determine whether horizontal gene transfer may have occurred in certain organisms. Chapter 20 also includes more discussion of tree thinking, as well as a new figure (Figure 20.11) that distinguishes between paraphyletic and polyphyletic taxa. New material in Chapter 21 clarifies the interplay between mutation, genetic variation, and natural selection. A new Make Connections Figure (Figure 21.15, “The Sickle-Cell Allele”) integrates material from chapters across the book in exploring the sickle-cell allele and its impact from the molecular and cellular levels to the allele’s global distribution in the human population. Other changes in the unit include new examples and figures that reinforce evolutionary concepts. For example, a new introduction to Chapter 23 tells the story of the discovery of whale fossils from the Sahara Desert, striking evidence of how organisms in the past differed from organisms living today. In Chapter 22, a new figure (Figure 22.11) has been added to support the expanded text discussion of allopolyploid speciation in *Tragopogon* in the Pacific Northwest. Dates have also been revised in the text, Table 23.1 (The Geologic Record), and figures in Chapter 23 and throughout the Second Edition to reflect the International Commission on Stratigraphy 2013 revision of the Geologic Time Scale.

UNIT 4 The Evolutionary History of Life

This unit employs a novel approach to studying the evolutionary history of biodiversity. Each chapter focuses on one or more major steps in the history of life, such as the origin of cells or the colonization of land. Likewise, the coverage of natural history and biological diversity emphasizes the evolutionary process—how factors such as the origin of key adaptations have influenced the rise and fall of different groups of organisms over time.



In the Second Edition, we have expanded our coverage of genomic and other molecular studies. Examples include a new figure (Figure 24.25) and text on the potential use and significance of CRISPR-Cas systems, a new Scientific Skills Exercise in Chapter 26 on genomic analyses of mycorrhizal and nonmycorrhizal fungi, and a new figure (Figure 27.36) and text related to evidence of gene flow between Neanderthals and modern humans. In addition, many phylogenies have been revised to reflect recent miRNA and genomic data. The unit also includes more connections to other chapters. For instance, a new Make Connections Question in Figure 24.4 asks students to apply material from Chapter 3 to explain how a membrane-like bilayer can self-assemble and form a vesicle, and a new Make Connections Figure (Figure 26.14) explores the diverse structural solutions for maximizing surface area that have evolved in cells, organ systems, and whole organisms. Other changes enhance the evolutionary storyline of the unit. For example, in Chapter 26, the chapter title, Figure 26.2, Key Concept 26.2, and text in Concepts 26.1 and 26.2 have all been revised to emphasize and explain that fungi are not closely related to plants, although they likely played a role in facilitating the colonization of land by plants, and that fungi possess their own novel adaptations for terrestrial life. Likewise, in Chapter 27, the discussion of the evolutionary impact of animals has been expanded, and new text and four new figures (Figures 27.12, 27.13, 27.30, and 27.31) on molluscs, birds, and mammals have been added. The chapter also includes expanded coverage of human evolution, including three new figures (Figures 27.34, 27.35, and 27.36). Supporting the extensive revision of Chapter 27, the number of Key Concepts in this chapter has increased from five to seven.

UNIT 5 Plant Form and Function

The form and function of higher plants are often treated as separate topics, thereby making it difficult for students to make connections between the two. In Unit 5, plant anatomy (Chapter 28) and the acquisition and transport of resources (Chapter 29) are bridged by a discussion of how plant architecture influences resource acquisition.



Chapter 30 provides an introduction to plant reproduction and examines controversies surrounding the genetic engineering of crop plants. The final chapter (Chapter 31) explores how plants respond to environmental challenges and opportunities and how the integration of this diverse information by plant hormones influences plant growth and reproduction.

In the Second Edition, a new micrograph of parenchyma cells and new information relating to root hair density, length, and function have been added to Chapter 28. In Chapter 29, a new Make Connections Figure (Figure 29.10, “Mutualism Across Kingdoms and Domains”) enables students to integrate what they have learned about plant mutualisms with other examples across the natural realm. A new Inquiry Figure (Figure 29.11) examines the metagenomics of soil bacteria. A discussion on mycorrhizae and plant evolution has also been added in Chapter 29. In Chapter 30, the angiosperm life cycle figure and related text are more closely integrated, with all the numbered steps now identified in the text. Also, a discussion of coevolution of flowers and pollinators has been added. The in-depth discussion of the development from seed to flowering plant has been expanded to include the transition from vegetative growth to reproductive growth, making a connection to what students learned about development in Chapter 28. In addition, the depictions of the structure of maize root systems and raspberry fruit development have been improved. The information in Concept 31.4 concerning plant defenses against disease has been thoroughly revised and updated to reflect rapid advances in our understanding of plant immunity. Updated information relates to the two types of plant immunity: PAMP-triggered immunity and effector-triggered immunity. New Figure 31.23 highlights examples of physical, chemical, and behavioral defenses against herbivory.

UNIT 6 Animal Form and Function

In this unit, a focused exploration of animal physiology and anatomy applies a comparative approach to a limited set of examples to bring out fundamental principles and conserved mechanisms. Students are first introduced to the closely related topics of endocrine signaling and homeostasis in an integrative introductory chapter (Chapter 32). Additional melding of interconnected material is reflected in chapters that combine treatment of circulation and gas exchange, reproduction and development, neurons and nervous systems, and motor mechanisms and behavior.

In the Second Edition, we re-envisioned the introductory chapter of this unit (Chapter 32), as conveyed by its new title, “The Internal Environment of Animals: Organization and Regulation.” Endocrine signaling and the integration of nervous and endocrine system function now precede the introduction of homeostasis and the consideration of the two major examples: thermoregulation and osmoregulation. Figures on simple hormone and neurohormone pathways (Figures 32.6 and 32.7) and hormone cascades (Figure 32.8) have been substantially revised to provide clear and consistent presentation of hormone function and of the regulation of hormone secretion. The presentation of the mechanism for filtrate processing in the kidney has been substantially revised, with a single figure (Figure 32.22) in place of two and with the accompanying numbered text walking students through a carefully paced tour of the nephron. In this chapter and throughout the unit, figures illustrating homeostatic regulation have been revised to highlight the common principles and features of homeostatic mechanisms. The unit includes two new Make Connections Figures: Figure 32.3 illustrates shared and divergent solutions to fundamental challenges common to plants and animals, and Figure 37.8, on ion movements and gradients, explores the fundamental role of concentration gradients in life processes ranging from osmoregulation and gas exchange to locomotion. Also in Chapter 37, the treatments of synaptic signaling, summation, modulating signaling, and neurotransmitters have been revised to highlight key ideas, ensuring appropriate pacing and helping students focus on fundamental principles rather than memorization. Updates in Unit 6 informed by current research include new Figure 33.15 and text highlighting the explosion of interest in and understanding of the microbiome. Chapter 38 opens with a new photograph and introductory text that showcase the “brainbow” technique for labeling individual brain neurons.

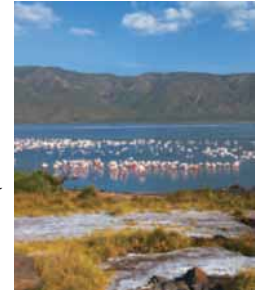


UNIT 7 Ecology

This unit applies the key themes of the text, including evolution, interactions, and energy and matter, to help students learn ecological principles. Chapter 40 integrates material on population growth and Earth’s environment, highlighting the importance of both biological and physical processes in determining where species are found.

Chapter 43 ends the book with a focus on global ecology and conservation biology. This chapter illustrates the threats to all species from increased human population growth and resource use. It begins with local factors that threaten individual species and ends with global factors that alter ecosystems, landscapes, and biomes.

The increased emphasis throughout the Second Edition on global climate change is capped by new discussions and figures in Unit 7. Chapter 43, for example, includes a new figure on the greenhouse effect (Figure 43.26) as well as new text examining aspects of climate change other than global warming. The chapter explores documented examples of the impacts to organisms in a new section on “Biological Effects of Climate Change” and a new Make Connections Figure (Figure 43.28, “Climate Change Has Effects at All Levels of Biological Organization”). Throughout the unit, the presentation of several other key topics has been revised. For example, in Chapter 40, the discussion of each of the following concepts or models was revised to standardize and clarify their meaning: life tables, per capita population growth, the per capita rate of increase (r), exponential population growth, and logistic population growth. The discussion of species interactions in Chapter 41 was modified to group species interactions according to whether they have positive (+) or negative (–) effects on survival and reproduction; as a result, there is a new section on “Exploitation” (which includes predation, herbivory, and parasitism) and another new section on “Positive Interactions” (which includes mutualism and commensalism). Material throughout Chapter 42 was revised to reinforce the fact that energy flows through ecosystems, whereas chemical elements cycle within ecosystems. New Figure Legend Questions give students practice in actively interpreting results; see, for example, the new questions with Figure 43.22 (biological magnification of PCBs) and Figure 43.31 (a new figure on per capita ecological footprints). The unit also includes a new Make Connections Figure (Figure 42.18, “The Working Ecosystem”) that ties together population, community, and ecosystem processes in the arctic tundra.



About the Authors

The author team's contributions reflect their biological expertise as researchers and their teaching sensibilities gained from years of experience as instructors at diverse institutions. They are also experienced textbook authors, having written *CAMPBELL BIOLOGY* in addition to *CAMPBELL BIOLOGY IN FOCUS*.

Lisa A. Urry



Lisa Urry (Chapter 1 and Units 1 and 2) is Professor of Biology and Chair of the Biology Department at Mills College in Oakland, California, and a Visiting Scholar at the University of California, Berkeley. After graduating from Tufts University with a double major in biology and French, Lisa completed her Ph.D. in molecular and developmental biology at Massachusetts Institute of Technology (MIT) in the MIT/Woods Hole Oceanographic Institution Joint Program. She has published a number of research papers, most of them focused on gene expression during embryonic and larval development in sea urchins. Lisa has taught a variety of courses, from introductory biology to developmental biology and senior seminar. As a part of her mission to increase understanding of evolution, Lisa also teaches a nonmajors course called Evolution for Future Presidents and is on the Teacher Advisory Board for the Understanding Evolution website developed by the University of California Museum of Paleontology. Lisa is also deeply committed to promoting opportunities for women and underrepresented minorities in science.

Michael L. Cain



Michael Cain (Chapter 1 and Units 3, 4, and 7) is an ecologist and evolutionary biologist who is now writing full-time. Michael earned a joint degree in biology and math at Bowdoin College, an M.Sc. from Brown University, and a Ph.D. in ecology and evolutionary biology from Cornell University. As a faculty member at New Mexico State University and Rose-Hulman Institute of Technology, he taught a wide range of courses, including introductory biology, ecology, evolution, botany, and conservation biology. Michael is the author of dozens of scientific papers on topics that include foraging behavior in insects and plants, long-distance seed dispersal, and speciation in crickets. In addition to his work on *CAMPBELL BIOLOGY IN FOCUS*, Michael is also the lead author of an ecology textbook.

Steven A. Wasserman



Steve Wasserman (Chapter 1 and Unit 6) is Professor of Biology at the University of California, San Diego (UCSD). He earned his A.B. in biology from Harvard University and his Ph.D. in biological sciences from MIT. Through his research on regulatory pathway mechanisms in the fruit fly *Drosophila*, Steve has contributed to the fields of developmental biology, reproduction, and immunity. As a faculty member at the University of Texas Southwestern Medical Center and UCSD, he has taught genetics, development, and physiology to undergraduate, graduate, and medical students. He currently focuses on teaching introductory biology. He has also served as the research mentor for more than a dozen doctoral students and more than 50 aspiring scientists at the undergraduate and high school levels. Steve has been the recipient of distinguished scholar awards from both the Markey Charitable Trust and the David and Lucille Packard Foundation. In 2007, he received UCSD's Distinguished Teaching Award for undergraduate teaching.

Peter V. Minorsky



Peter Minorsky (Chapter 1 and Unit 5) is Professor of Biology at Mercy College in New York, where he teaches introductory biology, evolution, ecology, and botany. He received his A.B. in biology from Vassar College and his Ph.D. in plant physiology from Cornell University. He is also the science writer for the journal *Plant Physiology*. After a postdoctoral fellowship at the University of Wisconsin at Madison, Peter taught at Kenyon College, Union College, Western Connecticut State University, and Vassar College. His research interests concern how plants sense environmental change. Peter received the 2008 Award for Teaching Excellence at Mercy College.

Jane B. Reece



The head of the author team for recent editions of *CAMPBELL BIOLOGY*, Jane Reece was Neil Campbell's longtime collaborator. Earlier, Jane taught biology at Middlesex County College and Queensborough Community College. She holds an A.B. in biology from Harvard University, an M.S. in microbiology from Rutgers University, and a Ph.D. in bacteriology from the University of California, Berkeley. Jane's research as a doctoral student and postdoctoral fellow focused on genetic recombination in bacteria. Besides her work on the Campbell textbooks for biology majors, she has been an author of *Campbell Biology: Concepts & Connections*, *Campbell Essential Biology*, and *The World of the Cell*.

Neil A. Campbell



Neil Campbell (1946–2004) combined the investigative nature of a research scientist with the soul of an experienced and caring teacher. He earned his M.A. in zoology from the University of California, Los Angeles, and his Ph.D. in plant biology from the University of California, Riverside, where he received the Distinguished Alumnus Award in 2001. Neil published numerous research articles on desert and coastal plants and how the sensitive plant (*Mimosa*) and other legumes move their leaves. His 30 years of teaching in diverse environments included introductory biology courses at Cornell University, Pomona College, and San Bernardino Valley College, where he received the college's first Outstanding Professor Award in 1986. He was a visiting scholar in the Department of Botany and Plant Sciences at the University of California, Riverside. Neil was the lead author of *Campbell Biology: Concepts & Connections*, *Campbell Essential Biology*, and *CAMPBELL BIOLOGY*, upon which this book is based.

Make Connections Visually

NEW! Ten **Make Connections Figures** integrate content from different chapters and provide a visual representation of “big picture” relationships.

Make Connections Figures include:

Figure 3.30 Contributions of Genomics and Proteomics to Biology, p. 112

Figure 8.20 The Working Cell, pp. 222–223

Figure 16.21 Genomics, Cell Signaling, and Cancer, pp. 382–383

Figure 21.15 The Sickle-Cell Allele, pp. 472–473

Figure 26.14 Maximizing Surface Area, p. 570

Figure 29.10 Mutualism Across Kingdoms and Domains, p. 647

Figure 32.3 Life Challenges and Solutions in Plants and Animals, shown at right and on pp. 710–711

Figure 37.8 Ion Movement and Gradients, p. 821

Figure 42.18 The Working Ecosystem, pp. 946–947

Figure 43.28 Climate Change Has Effects at All Levels of Biological Organization, pp. 968–969

▼ Figure 32.3

MAKE CONNECTIONS

Life Challenges and Solutions in Plants and Animals

Multicellular organisms face a common set of challenges. Comparing the solutions that have evolved in plants and animals reveals both unity (shared elements) and diversity (distinct features) across these two lineages.



Environmental Response

All forms of life must detect and respond appropriately to conditions in their environment. Specialized organs sense environmental signals. For example, the floral head of a sunflower (left) and an insect's eyes (right) both contain photoreceptors that detect light. Environmental signals activate specific receptor proteins, triggering signal transduction pathways that initiate cellular responses coordinated by chemical and electrical communication. (See Figure 31.12 and Figure 38.26.)



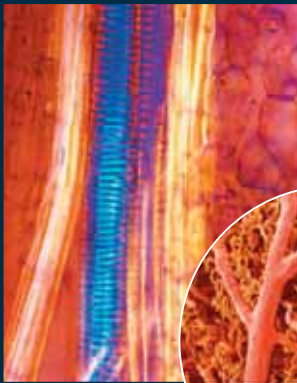
Nutritional Mode

All living things must obtain energy and carbon from the environment to grow, survive, and reproduce. Plants are autotrophs, obtaining their energy through photosynthesis and their carbon from inorganic sources, whereas animals are heterotrophs, obtaining their energy and carbon from food. Evolutionary adaptations in plants and animals support these different nutritional modes. The broad surface of many leaves (left) enhances light capture for photosynthesis. When hunting, a bobcat relies on stealth, speed, and sharp claws (right). (See Figure 29.2 and Figure 33.14.)

Growth and Regulation

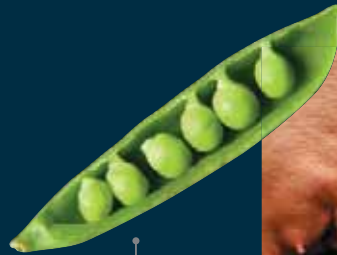
The growth and physiology of both plants and animals are regulated by hormones. In plants, hormones may act in a local area or be transported in the body. They control growth patterns, flowering, fruit development, and more (left). In animals, hormones circulate throughout the body and act in specific target tissues, controlling homeostatic processes and developmental events such as molting (below). (See Table 31.1 and Figure 33.19.)





Transport

All but the simplest multicellular organisms must transport nutrients and waste products between locations in the body. A system of tubelike vessels is the common evolutionary solution, while the mechanism of circulation varies. Plants harness solar energy to transport water, minerals, and sugars through specialized tubes (left). In animals, a pump (heart) moves circulatory fluid through vessels (right). (See Figure 28.9 and Figure 34.3.)



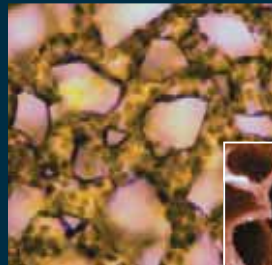
Reproduction

In sexual reproduction, specialized tissues and structures produce and exchange gametes. Offspring are generally supplied with nutritional stores that facilitate rapid growth and development. For example, seeds (left) have stored food reserves that supply energy to the young seedling, while milk provides sustenance for juvenile mammals (right). (See Figure 30.8 and Figure 32.7.)



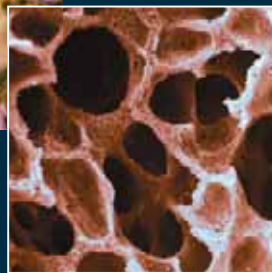
Absorption

Organisms need to absorb nutrients. The root hairs of plants (left) and the villi (projections) that line the intestines of vertebrates (right) increase the surface area available for absorption. (See Figure 28.4 and Figure 33.10.)



Gas Exchange

The exchange of certain gases with the environment is essential for life. Respiration by plants and animals requires taking up oxygen (O_2) and releasing carbon dioxide (CO_2). In photosynthesis, net exchange occurs in the opposite direction: CO_2 uptake and O_2 release. In both plants and animals, highly convoluted surfaces that increase the area available for gas exchange have evolved, such as the spongy mesophyll of leaves (left) and the alveoli of lungs (right). (See Figure 28.17 and Figure 34.20.)



MAKE CONNECTIONS Compare the adaptations that enable plants and animals to respond to the challenges of living in hot and cold environments. See Concepts 31.3 and 32.3.

ANIMATION



Visit the Study Area in **MasteringBiology** for the BioFlix® 3-D Animations on Water Transport in Plants (Chapter 29), Homeostasis: Regulating Blood Sugar (Chapter 33), and Gas Exchange (Chapter 34).

◀ **Make Connections Questions** ask students to relate content in the chapter to material presented earlier in the course.

Practice Scientific Skills

Scientific Skills Exercises in every chapter use real data to build key skills needed for biology, including **data analysis, graphing, experimental design, and math skills.**

Each Scientific Skills Exercise is based on an experiment related to the chapter content.

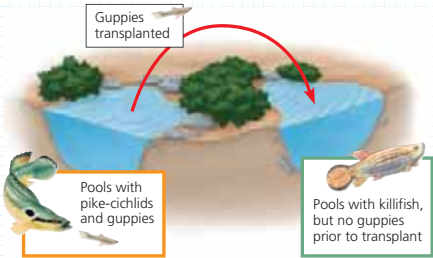
Scientific Skills Exercise

Making and Testing Predictions

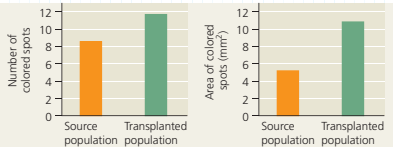
Can Predation Result in Natural Selection for Color Patterns in Guppies? What we know about evolution changes constantly as new observations lead to new hypotheses—and hence to new ways to test our understanding of evolutionary theory. Consider the wild guppies (*Poecilia reticulata*) that live in pools connected by streams on the Caribbean island of Trinidad. Male guppies have highly varied color patterns that are controlled by genes that are only expressed in adult males. Female guppies choose males with bright color patterns as mates more often than they choose males with drab coloring. But the bright colors that attract females also can make the males more conspicuous to predators. Researchers observed that in pools with few predator species, the benefits of bright colors appear to “win out,” and males are more brightly colored than in pools where predation is more intense.

One guppy predator, the killifish, preys on juvenile guppies that have not yet displayed their adult coloration. Researchers predicted that if adult guppies with drab colors were transferred to a pool with only killifish, eventually the descendants of these guppies would be more brightly colored (because of the female preference for brightly colored males).

How the Experiment Was Done Researchers transplanted 200 guppies from pools containing pike-cichlid fish, intense predators of adult guppies, to pools containing killifish, less active predators that prey mainly on juvenile guppies. They tracked the number of bright-colored spots and the total area of those spots on male guppies in each generation.



Data from the Experiment After 22 months (15 generations), researchers compared the color pattern data for guppies from the source and transplanted populations.




| Population | Number of colored spots | Area of colored spots (mm ²) |
|-------------------------|-------------------------|--|
| Source population | 9 | 5 |
| Transplanted population | 11 | 10 |

Data from J. A. Endler, Natural selection on color patterns in *Poecilia reticulata*, *Evolution* 34:76–91 (1980).

INTERPRET THE DATA

- Identify the following elements of hypothesis-based science in this example: (a) question, (b) hypothesis, (c) prediction, (d) control group, and (e) experimental group. (For additional information about hypothesis-based science, see Chapter 1 and the Scientific Skills Review in Appendix F and the Study Area of MasteringBiology.)
- Explain how the types of data the researchers chose to collect enabled them to test their prediction.
- What conclusion do you draw from the data presented above?
- Predict what would happen if, after 22 months, guppies from the transplanted population were returned to the source pool. Describe an experiment to test your prediction.

 A related version of this Scientific Skills Exercise can be assigned in MasteringBiology.

Most Scientific Skills Exercises use **data from published research**, cited in the exercise.

Questions build in **difficulty**, walking students through new skills step by step and providing opportunities for higher-level critical thinking.

Every chapter has a Scientific Skills Exercise:

- Interpreting a Pair of Bar Graphs, p. 62
- Interpreting a Scatter Plot with a Regression Line, p. 84
- Analyzing Polypeptide Sequence Data, p. 113
- Using a Scale Bar to Calculate Volume and Surface Area of a Cell, p. 124
- Interpreting a Scatter Plot with Two Sets of Data, p. 153
- Making a Line Graph and Calculating a Slope, p. 178
- Making a Bar Graph and Evaluating a Hypothesis, p. 199
- Making Scatter Plots with Regression Lines, p. 220
- Interpreting Histograms, p. 240
- Making a Line Graph and Converting Between Units of Data, p. 254
- Making a Histogram and Analyzing a Distribution Pattern, p. 271
- Using the Chi-Square (χ^2) Test, p. 290
- Working with Data in a Table, p. 301
- Interpreting a Sequence Logo, p. 338
- Analyzing DNA Deletion Experiments, p. 357
- Analyzing Quantitative and Spatial Gene Expression Data, p. 369
- Analyzing a Sequence-Based Phylogenetic Tree to Understand Viral Evolution, p. 397
- Reading an Amino Acid Sequence Identity Table, p. 414
- Making and Testing Predictions**, shown above and on p. 436
- NEW!** Using Protein Sequence Data to Test an Evolutionary Hypothesis, p. 454
- Using the Hardy-Weinberg Equation to Interpret Data and Make Predictions, p. 464
- Identifying Independent and Dependent Variables, Making a Scatter Plot, and Interpreting Data, p. 485
- Estimating Quantitative Data from a Graph and Developing Hypotheses, p. 503

Each Scientific Skills Exercise from the text also has an **assignable, interactive tutorial version in MasteringBiology** that is automatically graded and includes coaching feedback.

MasteringBiology®

34. Circulation and Gas Exchange Scientific Skills Exercise: Interpreting Data in Histograms

Item Type: Tutorial | Difficulty: 1 | Time: 5m | Learning Outcomes | Contact the Publisher

Manage this item: Standard View

Scientific Skills Exercise: Interpreting Data in Histograms

Individuals with an inactivating mutation in one copy of PCSK9 gene (study group)

Individuals with two functional copies of PCSK9 gene (control group)

Data from J.-C. Cohen et al., "Sequence variations in PCSK9, low LDL, and protection against coronary heart disease," *New England Journal of Medicine* 354:1264-1272 (2006).

Part A - Reading the histograms

The results are presented using a form of bar graph called a histogram. In a histogram, the variable on the x-axis is grouped into ranges. The height of each bar in this histogram represents the percentage of samples that fall into the range specified on the x-axis for that bar. For example, in the top histogram, about 4% of individuals studied had plasma LDL cholesterol levels in the 25–50 mg/dL (milligrams per deciliter) range.

What percentage of individuals in the study group had an LDL level below 100 mg/dL? (Hint: Add the percentages for the relevant bars.)

about 17%
 about 35%
 about 58%
 about 79%

Submit My Answers Give Up

Incorrect. Try Again

Make sure you add the values of the three bars representing plasma LDL levels below 100 mg/dL (the bars to the left of 100 mg/dL on the x-axis).

Part B

What percentage of individuals in the control group had an LDL level below 100 mg/dL?

about 5%
 about 17%
 about 43%
 about 58%

Submit My Answers Give Up

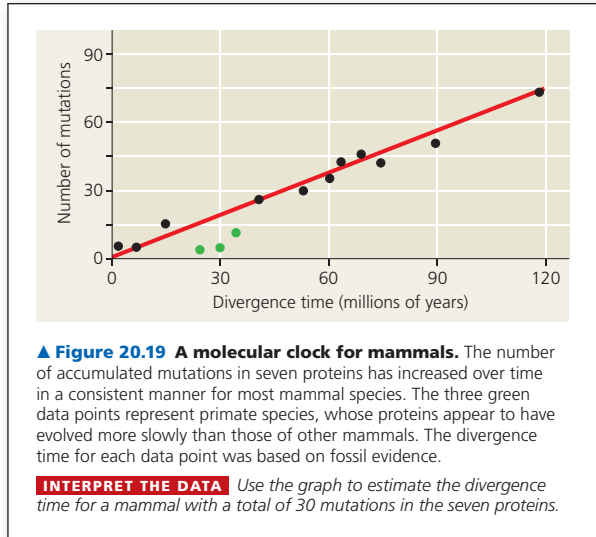
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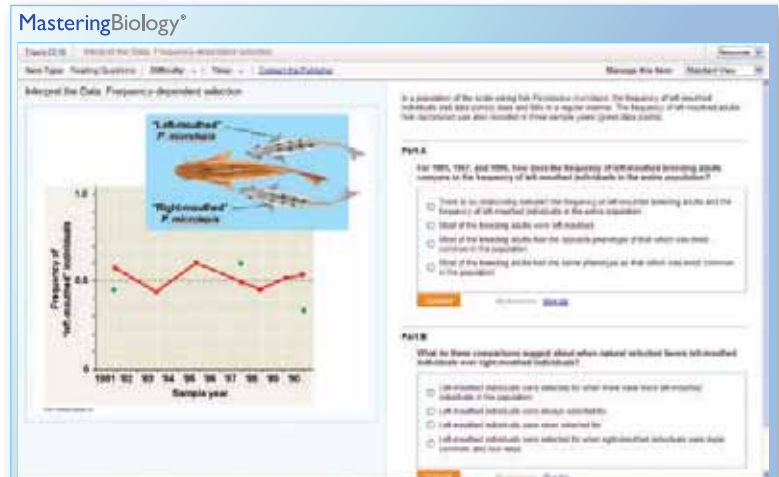
24. Making a Bar Graph and Interpreting Data, p. 537
25. Interpreting Comparisons of Genetic Sequences, p. 545
26. **NEW!** Interpreting Genomic Data and Generating Hypotheses, p. 573
27. Understanding Experimental Design and Interpreting Data, p. 614
28. Using Bar Graphs to Interpret Data, p. 626
29. Calculating and Interpreting Temperature Coefficients, p. 641
30. Using Positive and Negative Correlations to Interpret Data, p. 676
31. Interpreting Experimental Results from a Bar Graph, p. 700
32. Describing and Interpreting Quantitative Data, p. 723
33. Interpreting Data from an Experiment with Genetic Mutants, p. 748
34. **Interpreting Data in Histograms**, shown above and on p. 765
35. Comparing Two Variables on a Common x-Axis, p. 792
36. Making Inferences and Designing an Experiment, p. 805
37. Interpreting Data Values Expressed in Scientific Notation, p. 831
38. Designing an Experiment Using Genetic Mutants, p. 841
39. Interpreting a Graph with Log Scales, p. 869
40. Using the Logistic Equation to Model Population Growth, p. 904
41. Using Bar Graphs and Scatter Plots to Present and Interpret Data, p. 914
42. Interpreting Quantitative Data in a Table, p. 937
43. Graphing Cyclic Data, p. 966

Interpret Data

CAMPBELL BIOLOGY IN FOCUS, Second Edition, and MasteringBiology offer a wide variety of ways for students to move beyond memorization and to **think like a scientist**.



◀ **NEW! Interpret the Data Questions** throughout the text ask students to analyze a graph, figure, or table.



▲ **NEW! Every Interpret the Data** question from the text is **assignable in MasteringBiology**.

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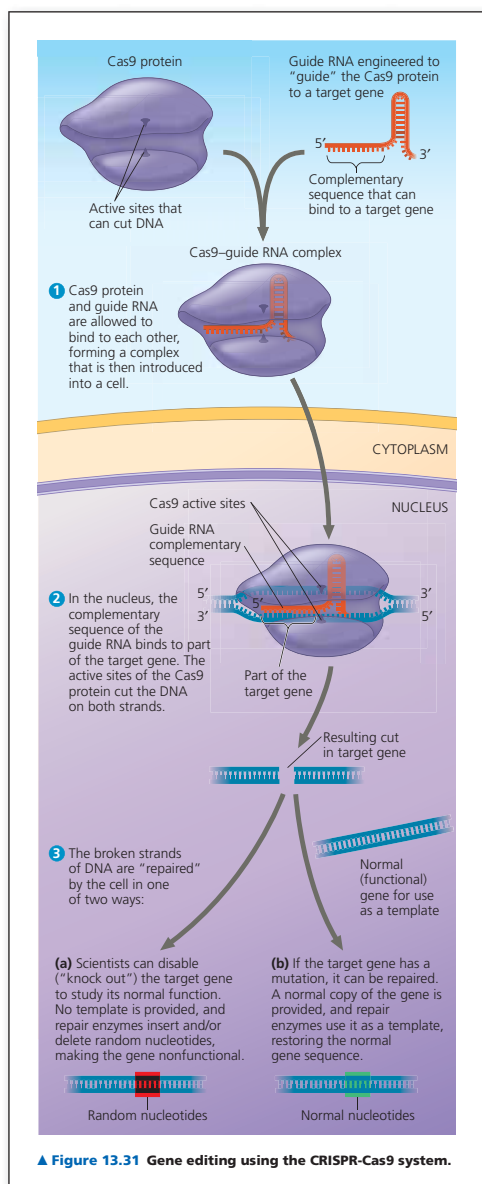
◀ **NEW! Solve It Tutorials** engage students in a multi-step investigation of a “mystery” or open question in which they must analyze real data. These are assignable in MasteringBiology. Topics include:

- Which Biofuel Has the Most Potential to Reduce Our Dependence on Fossil Fuels?
- Is It Possible to Treat Bacterial Infections Without Traditional Antibiotics?
- Which Insulin Mutations May Result in Disease?
- Are You Getting the Fish You Paid For?
- Why Are Honey Bees Vanishing?
- What Is Causing Episodes of Muscle Weakness in a Patient?
- How Can the Severity of Forest Fires Be Reduced?

Keep Current with New Scientific Advances

NEW! The Second Edition incorporates **up-to-date content** on genomics, gene editing, human evolution, microbiomes, climate change, and more.

NEW! The Second Edition shows students how our ability to **sequence DNA and proteins rapidly and inexpensively** is transforming every subfield of biology, from cell biology to physiology to ecology. For instance, the examples in this figure from Chapter 3 are explored in greater depth later in the text.



▲ **Figure 13.31** Gene editing using the CRISPR-Cas9 system.

◀ **NEW!** Chapter 13 describes **gene editing using the CRISPR-Cas9 system**, and Chapter 17 describes the basic biology of this system in bacteria.

NEW! Chapter 27 includes new material on human origins, including how sequencing DNA extracted from this fossil jawbone recently revealed **evidence of human-Neanderthal interbreeding**.

Figure 3.30

MAKE CONNECTIONS

Contributions of Genomics and Proteomics to Biology

Nucleotide sequencing and the analysis of large sets of genes and proteins can be done rapidly and inexpensively due to advances in technology and information processing. Taken together, genomics and proteomics have advanced our understanding of biology across many different fields.

Evolution


A major aim of evolutionary biology is to understand the relationships among species, both living and extinct. For example, genome sequence comparisons have identified the hippopotamus as the land mammal sharing the most recent common ancestor with whales. (See Figure 19.20.)



Hippopotamus Short-finned pilot whale


Conservation Biology

The tools of molecular genetics and genomics are increasingly used by forensic ecologists to identify which species of animals and plants are killed illegally. In one case, genomic sequences of DNA from illegal shipments of elephant tusks were used to track down poachers and pinpoint the territory where they were operating. (See Figure 43.8.)




Paleontology

New DNA sequencing techniques have allowed decoding of minute quantities of DNA found in ancient tissues from our extinct relatives, the Neanderthals (*Homo neanderthalensis*). Sequencing the Neanderthal genome has informed our understanding of their physical appearance as well as their relationship with modern humans. (See Figure 27.36.)




Medical Science

Identifying the genetic basis for human diseases like cancer helps researchers focus their search for potential future treatments. Currently, sequencing the sets of genes expressed in an individual's tumor can allow a more targeted approach to treating the cancer, a type of "personalized medicine." (See Concept 9.3 and Figure 16.21.)



Species Interactions

Most plant species exist in a mutually beneficial partnership with fungi (right) and bacteria associated with the plants' roots; these interactions improve plant growth. Genome sequencing and analysis of gene expression have allowed characterization of plant-associated communities. Such studies will help advance our understanding of such interactions and may improve agricultural practices. (See the Chapter 26 Scientific Skills Exercise and Figure 29.11.)



MAKE CONNECTIONS Considering the examples provided here, describe how the approaches of genomics and proteomics help us to address a variety of biological questions.

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▲ **Figure 27.36** Fossil evidence of human-Neanderthal interbreeding.

Focus on the Key Concepts

Each chapter is organized around a framework of **3 to 6 Key Concepts** that focus on the big picture and provide a context for the supporting details.

The **list of Key Concepts** ▶ introduces the big ideas covered in the chapter.

Every chapter opens with a visually dynamic photo accompanied by an **intriguing question** that invites students into the chapter.

CHAPTER

14

Gene Expression: From Gene to Protein

KEY CONCEPTS

- 14.1** Genes specify proteins via transcription and translation
- 14.2** Transcription is the DNA-directed synthesis of RNA: *a closer look*
- 14.3** Eukaryotic cells modify RNA after transcription
- 14.4** Translation is the RNA-directed synthesis of a polypeptide: *a closer look*
- 14.5** Mutations of one or a few nucleotides can affect protein structure and function



▲ **Figure 14.1** How does a single faulty gene result in the dramatic appearance of an albino donkey?

The Flow of Genetic Information

The island of Asinara lies off the coast of Sardinia, an Italian island. The name Asinara probably originated from the Latin word *sinuaria*, which means “sinus-shaped.” A second meaning of Asinara is “donkey-inhabited,” which is particularly appropriate because Asinara is home to a wild population of albino donkeys (**Figure 14.1**). The donkeys were brought to Asinara in the early 1800s and abandoned there in 1885 when the 500 residents were forced to leave the island so it could be used as a penal colony. What is responsible for the phenotype of the albino donkey, strikingly different from its pigmented relative?

Inherited traits are determined by genes, and the trait of albinism is caused by a recessive allele of a pigmentation gene (see Concept 11.4). The information content of genes is in the form of specific sequences of nucleotides along strands of DNA, the genetic material. But how does this information determine an organism’s traits? Put another way, what does a gene actually say? And how is its message

translated by cells into a specific trait, such as brown hair, type A blood, or, in the case of an albino donkey, a total lack of pigment? The albino donkey has a faulty version of a key protein, an enzyme required for pigment synthesis, and this protein is faulty because the gene that codes for it contains incorrect information.

This example illustrates the main point of this chapter: The DNA inherited by an organism leads to specific traits by dictating the synthesis of proteins and of RNA molecules involved in protein synthesis. In other words, proteins are the link between genotype and phenotype. **Gene expression** is the process by which DNA directs the synthesis of proteins (or, in some cases, just RNAs). The expression of genes that code for proteins includes two stages: transcription and translation. This chapter describes the flow of information from gene to protein and explains how genetic mutations affect organisms through their proteins. Understanding the processes of gene expression, which are similar in all three domains of life, will allow us to revisit the concept of the gene in more detail at the end of the chapter.

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After reading a Key Concept section, ▶ students can check their understanding using the **Concept Check questions**:

Make Connections questions ask students to relate content in the chapter to material presented earlier in the course.

What if? questions ask students to apply what they’ve learned.

Draw It Exercises ask students to put pencil to paper and draw a structure, annotate a figure, or graph experimental data.

CONCEPT CHECK 14.5

1. What happens when one nucleotide pair is lost from the middle of the coding sequence of a gene?
2. **MAKE CONNECTIONS** Individuals heterozygous for the sickle-cell allele show effects of the allele under some circumstances (see Concept 11.4). Explain in terms of gene expression.
3. **WHAT IF? DRAW IT** The template strand of a gene includes this sequence: 3'-TACTTGTCGATATC-5'. It is mutated to 3'-TACTTGTCGAATATC-5'. For both versions, draw the DNA, the mRNA, and the encoded amino acid sequence. What is the effect on the amino acid sequence?

For suggested answers, see Appendix A.

The Summary of Key Concepts refocuses students on the main points of the chapter.

14 Chapter Review

Go to MasteringBiology® for Assignments, the eText, and the Study Area with Animations, Activities, Vocab Self-Quiz, and Practice Tests.

SUMMARY OF KEY CONCEPTS

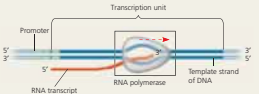
CONCEPT 14.1
Genes specify proteins via transcription and translation (pp. 323–328)

- Beadle and Tatum's studies of mutant strains of *Neurospora* led to the one gene–one polypeptide hypothesis. During **gene expression**, the information encoded in genes is used to make specific polypeptide chains (enzymes and other proteins) or RNA molecules.
- **Transcription** is the synthesis of RNA complementary to a **template strand of DNA**. **Translation** is the synthesis of a polypeptide whose amino acid sequence is specified by the nucleotide sequence in **mRNA**.
- Genetic information is encoded as a sequence of nonoverlapping nucleotide triplets, or **codons**. A codon in messenger RNA (mRNA) either is translated into an amino acid (61 of the 64 codons) or serves as a stop signal (3 codons). Codons must be read in the correct **reading frame**.

1 Describe the process of gene expression, by which a gene affects the phenotype of an organism.

CONCEPT 14.2
Transcription is the DNA-directed synthesis of RNA: a closer look (pp. 328–330)

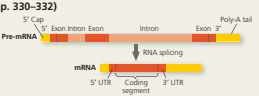
- RNA synthesis is catalyzed by **RNA polymerase**, which links together RNA nucleotides complementary to a DNA template strand. This process follows the same base-pairing rules as DNA replication, except that in RNA, uracil substitutes for thymine.



- The three stages of transcription are initiation, elongation, and termination. A **promoter**, often including a **TATA box** in eukaryotes, establishes where RNA synthesis is initiated. **Transcription factors** help eukaryotic RNA polymerase recognize promoter sequences, forming a **transcription initiation complex**. Termination differs in bacteria and eukaryotes.

2 What are the similarities and differences in the initiation of gene transcription in bacteria and eukaryotes?

CONCEPT 14.3
Eukaryotic cells modify RNA after transcription (pp. 330–332)



CONCEPT 14.4
Translation is the RNA-directed synthesis of a polypeptide: a closer look (pp. 332–342)

- A cell translates an mRNA message into protein using **transfer RNAs (tRNAs)**. After being bound to a specific amino acid by an **aminoacyl-tRNA synthetase**, a tRNA lines up via its **anticodon** at the complementary codon on mRNA. A ribosome, made up of **ribosomal RNAs (rRNAs)** and proteins, facilitates this coupling with binding sites for mRNA and tRNA. Ribosomes coordinate the three stages of translation: initiation, elongation, and termination. The formation of peptide bonds between amino acids is catalyzed by ribosomal RNAs as tRNAs move through the **A** and **P** sites and exit through the **E** site.
- After translation, modifications to proteins can affect their shape. Free ribosomes in the cytosol initiate synthesis of all proteins, but proteins with a **signal peptide** are synthesized on the ER.
- A gene can be transcribed by multiple RNA polymerases simultaneously. Also, a single mRNA molecule can be translated simultaneously by a number of ribosomes, forming a **polysome**. In bacteria, these processes are coupled, but in eukaryotes they are separated in time and space by the nuclear membrane.

3 What function do rRNAs serve in the process of translation?

CONCEPT 14.5
Mutations of one or a few nucleotides can affect protein structure and function (pp. 342–346)

- Small-scale mutations include **point mutations**, changes in one DNA nucleotide pair, which may lead to production of non-functional proteins. **Nucleotide-pair substitutions** can cause **missense** or **nonsense mutations**. Nucleotide-pair **insertions** or **deletions** may produce **frameshift mutations**.
- Spontaneous mutations can occur during DNA replication, recombination, or repair. Chemical and physical **mutagens** cause DNA damage that can alter genes.

CHAPTER 14. GENE EXPRESSION: FROM GENE TO PROTEIN 345

VOCAB SELF-QUIZ

NEW! QR Codes and URLs at the end of every chapter give students quick access to **Vocabulary Self-Quizzes** and **Practice Tests** on their smartphones, tablets, and computers.

PRACTICE TEST

TEST YOUR UNDERSTANDING

Level 1: Knowledge/Comprehension

- Which of the following is *not* true about mRNAs?
 - (A) Eukaryotic mRNAs lack introns.
 - (B) Prokaryotic mRNAs do not have poly-A tails.
 - (C) Prokaryotic cells do not cap their mRNAs on the 5'.
 - (D) Pre-mRNAs do not have 5' caps.
- Which of the following is *not* true of a codon?
 - (A) It may code for the same amino acid as another codon.
 - (B) It never codes for more than one amino acid.
 - (C) It extends from one end of a tRNA molecule.
 - (D) It is the basic unit of the genetic code.
- The anticodon of a particular tRNA molecule is
 - (A) complementary to the corresponding mRNA codon.
 - (B) complementary to the corresponding triplet in rRNA.
 - (C) the part of tRNA that bonds with a specific amino acid.
 - (D) catalytic, making the tRNA a ribozyme.
- Which of the following is *not* true of RNA processing?
 - (A) Exons are cut out before mRNA leaves the nucleus.
 - (B) Nucleotides may be added at both ends of the RNA.
 - (C) Ribozymes may function in RNA splicing.
 - (D) RNA splicing can be catalyzed by spliceosomes.
- Which of the following is *not* required for transcription?
 - (A) RNA polymerase
 - (B) Transcription factors
 - (C) Deoxyribonucleotides
 - (D) DNA promoter

Level 2: Application/Analysis

- Using Figure 14.6, identify a 5' → 3' sequence of nucleotides in the DNA template strand for an mRNA coding for the polypeptide sequence Phe-Pro-Lys.
 - (A) 5'-UUUCCCAAA-3'
 - (B) 5'-GAACCCCTT-3'
 - (C) 5'-CTTCGGGAA-3'
 - (D) 5'-AAACCCUUU-3'
- The following sequence of a DNA template strand: 5'-TAT GGT GCA-3' codes for a peptide. Determine the mutation that would have the most severe effect.
 - (A) 5'-TAT GGT ACA-3'
 - (B) 5'-TAT CGT GCA-3'
 - (C) 5'-GAT GGT GCA-3'
 - (D) 5'-TAT GTT GCA-3'
- Would the coupling of the processes shown in Figure 14.23 be found in a eukaryotic cell? Explain why or why not.

9. Fill in the following table:

| Type of RNA | Functions |
|----------------------|--|
| Messenger RNA (mRNA) | |
| Transfer RNA (tRNA) | |
| Primary transcript | Plays catalytic (ribozyme) roles and structural roles in ribosomes |


Level 3: Synthesis/Evaluation

10. SCIENTIFIC INQUIRY
A researcher wants to produce a eukaryotic protein of 45 kDa (around 350 amino acids). The researcher amplifies the gene from genomic DNA taking care to eliminate the eukaryotic promoter and starts the amplification at the start ATG codon and stops it right at the termination codon. Finally, the gene is cloned into a bacterial expression vector right after a bacterial promoter. When the researcher tries to produce the protein in bacteria no protein of the expected size can be detected. However, the researcher is able to identify some RNA for the gene that is around 10,000 base pairs. Explain what is happening in this experiment.

11. FOCUS ON EVOLUTION
The genetic code has a built-in redundancy as several codons can code for a single amino acid. This is the result of the availability of greater possible base combinations in codons (64 than availability of amino acids). Take a look at table 14.6 and you will realize that the third base of a codon or several amino acids can be changed without changing the translation. Discuss the possible implications this could have in mutation and evolution.

12. FOCUS ON INFORMATION
Evolution accounts for the unity and diversity of life, and the continuity of life is based on heritable information in the form of DNA. In a short essay (100–150 words), discuss how the fidelity with which DNA is inherited is related to the processes of evolution. (Review the discussion of proofreading and DNA repair in Concept 13.2.)

13. SYNTHESIZE YOUR KNOWLEDGE



Some mutations result in proteins that function well at one temperature but are nonfunctional at a different (usually higher) temperature. Siamese cats have such a "temperature-sensitive" mutation in a gene encoding an enzyme that makes dark pigment in the fur. The mutation results in the breed's distinctive point markings and lighter body color (see the photo). Using this information and what you learned in the chapter, explain the pattern of the cat's fur pigmentation.

For selected answers, see Appendix A.

- **Summary of Key Concepts questions** check students' understanding of a key idea from each concept.
- **Summary figures** recap key information visually.

Evolution, the fundamental theme of biology, is emphasized throughout. For example:

- Every Chapter Review includes a "Focus on Evolution" question (shown above right).
- Every chapter has a section explicitly relating the chapter content to evolution (shown at right).

Evolution of the Genetic Code

EVOLUTION The genetic code is nearly universal, shared by organisms from the simplest bacteria to the most complex plants and animals. The mRNA codon CCG, for instance, is translated as the amino acid proline in all organisms whose genetic code has been examined. In laboratory experiments, genes can be transcribed and translated after being transplanted from one species to another, sometimes with quite striking results, as shown in **Figure 14.7**. Bacteria can be programmed by the insertion of human genes to synthesize certain human proteins for medical use, such as insulin. Such applications have produced many exciting developments in the area of genetic engineering (see Concept 13.4).

Despite a small number of exceptions in which a few codons differ from the standard ones, the evolutionary significance of the code's *near* universality is clear. A language shared by all living things must have been operating very early in the

(a) Tobacco plant expressing a firefly gene. The yellow glow is produced by a chemical reaction catalyzed by the protein product of the firefly gene.

(b) Pig expressing a jellyfish gene. Researchers injected the gene for a fluorescent protein into fertilized pig eggs. One of the eggs developed into this fluorescent pig.

▲ Figure 14.7 Expression of genes from different species. Because diverse forms of life share a common genetic code, one species can be programmed to produce proteins characteristic of a second species by introducing DNA from the second species into the first.

FOCUS ON THE KEY CONCEPTS

19

Instructors can easily incorporate **active learning** into their courses using suggested activity ideas and questions for use with classroom response systems, including Learning Catalytics.

- ▼ **Learning Catalytics™** allows students to use their smartphone, tablet, or laptop to respond to questions in class. Visit www.learningcatalytics.com.



The following conditions were detected in a mutant cell:

- The cell is running out of ATP, while ADP is building up to very high levels.
- NADH is building up to very high levels, while the level of NAD^+ is becoming very low.
- The amount of protons in the intermembrane space and in the matrix is becoming more equal (the strength of the proton gradient is decreasing/less).

Use this information to predict which stage of cellular respiration is not functioning normally in this mutant cell.

The diagram illustrates the stages of cellular respiration. It shows glycolysis in the cytoplasm, the Krebs cycle in the mitochondrial matrix, and the electron transport chain in the intermembrane space. Key components labeled include: Glycolysis (Glucose to Pyruvate), Pyruvate to Acetyl-CoA, Krebs cycle (Acetyl-CoA to CO_2), and Electron transport chain (Electrons from NADH and FADH_2 to O_2). It also shows the production of ATP in the intermembrane space and the pumping of protons (H^+) across the membrane. A central red circle is labeled 'ATP synthase'.

More Instructor Resources

Instructor's Resource Materials

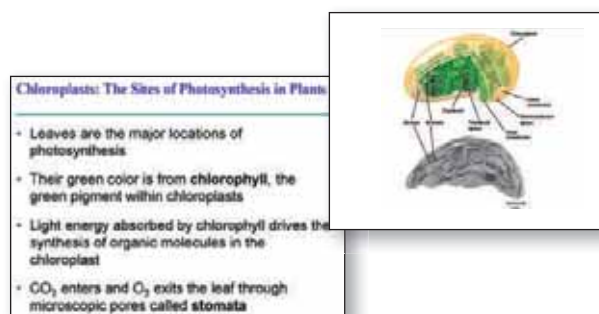
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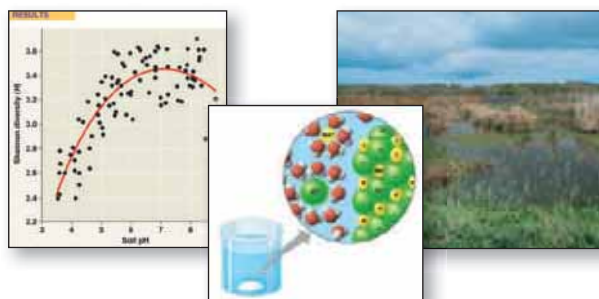
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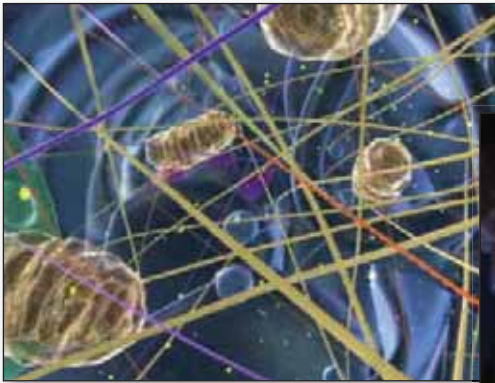
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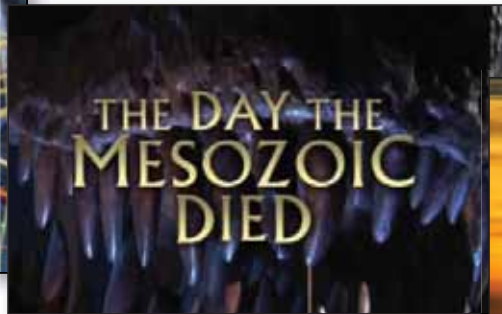
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| Assigned Points | | 25 | 10 | 10 | 10 | 10 | 10 | 10 | 10 | 10 | 10 | 10 | 130 |
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| Student 1 | | 70.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 2 | | 48.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 3 | | 54.0 | 81.0 | 81.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 4 | | 48.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 5 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 6 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 7 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 8 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 9 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 10 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 11 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 12 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 13 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 14 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 15 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 16 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 17 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 18 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 19 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |
| Student 20 | | 82.0 | 82.0 | 82.0 | 84.0 | 84.0 | 81.0 | 82.0 | 81.0 | 81.0 | 81.0 | 81.0 | 81.0 |

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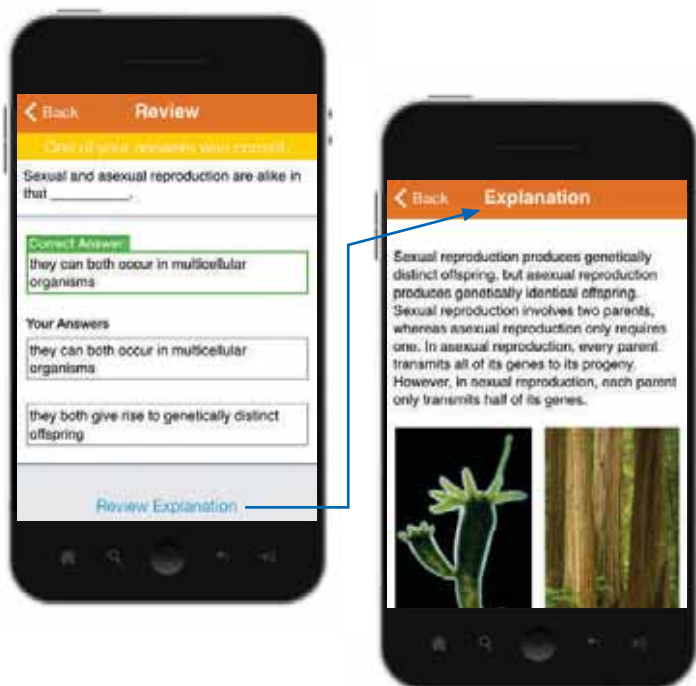
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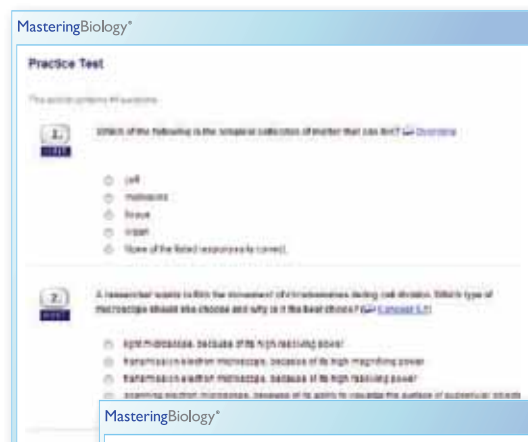
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^{*}The Inquiry Figure, original research paper, and a worksheet to guide you through the paper are provided in *Inquiry in Action: Interpreting Scientific Papers*.

[†]A related Experimental Inquiry Tutorial can be assigned in MasteringBiology.

Acknowledgments

The authors wish to express their gratitude to the global community of instructors, researchers, students, and publishing professionals who have contributed to the Second Edition of **CAMPBELL BIOLOGY IN FOCUS**.

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