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THE MADER/WINDELSPECHT STORY... The twelfth edition of Biology is a traditional, comprehensive introductory biology textbook, with coverage from Cell Structure and Function to the Conservation of Biodiversity. The book, which centers on the evolution and diversity of organisms, is appropriate for any one- or two-semester biology course. Biology, 12th Edition is the epitome of Sylvia Mader's expertise. Its concise, precise writing-style employs lucid language to present the material as succinctly as possible, enabling students—even non-majors—to master the foundational concepts before coming to class. “Before You Begin”, “Following the Themes”, and “Thematic Feature Readings” piece together the three major themes of the text—evolution, nature of science, and biological systems. Students are consistently engaged in these themes, revealing the interconnectedness of the major topics in biology. Sylvia Mader typifies an icon of science education. Her dedication to her students, coupled with her clear, concise writing-style has benefited the education of thousands of students over the past three decades. The integration of the text and digital world has been achieved with the addition of Dr. Michael Windelspecht's facility for the development of digital learning assets. For over ten years, Michael served as the Introductory Biology Coordinator at Appalachian State University—a program that enrolls over 4,500 non-science majors annually. Michael

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is the lead architect in the design of McGraw-Hill's Connect Plus and LearnSmart media content for the Mader series. These assets allow instructors to easily design interactive tutorial materials, enhance presentations in both online and traditional environments, and assess the learning objectives and outcomes of the course.

Mader includes revised coverage of animal behaviour and ecology as well as a wealth of new focus boxes which highlight topics of high interest and relate biology to everyday life. This text is linked to a web site offering extended chapter outlines.

An immensely useful manual with many attractive features: comprehensive and lucid keys, precise diagrams, annotated checklists and up-to-date references. ... there is no doubt that it should be seen as an example of the type of manual which is so badly needed in the study of the fauna of many shores around the world.--Journal of Animal Ecology Congratulations to the editors, contributors, and publisher for a job well done. The third edition has been rewritten,

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corrected, and enlarged, so that while retaining the basic organization of the earlier ones, it is more useful, informative and up-to-date. The meticulous scholarship of Smith and Carlton is just what the revision needed.--Systematic Zoology This revision should serve for many years. It is therefore particularly commendable that the editing has been meticulous, perhaps flawless. ... thanks are due to the many contributors for a job well done.--The Quarterly Review of Biology As the Pacific Coast intertidal zone undergoes increasingly profound changes, knowing the sentinel invertebrates can foretell the future of the sea, and hence, of our species. Jim Carlton's hefty new update of The Light & Smith Manual, the comprehensive compendium of who's who between the tides, is the best and quickest way to do so.--Elliot A. Norse, President, Marine Conservation Biology Institute This much-anticipated modernization of Light's Manual is an astonishing accomplishment, blending state-of-the-art taxonomy with profusely illustrated and user-friendly keys to who's whom on marine shores from its stated boundaries of mid-California through Oregon, and clearly, much further north. It's also an informative, well referenced read. Marine biologists should not leave home without it.--Robert Paine, Professor Emeritus of Biology, University of Washington At this time of environmental change and loss of biodiversity, species identification has never been more important.

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The fourth edition of Light and Smith is more than just a field guide--it is a masterwork of research and description with a strong focus on morphological detail. No other book has such a broad scope, newly expanded to include even the most obscure taxa. The revised keys and beautiful anatomical illustrations make this classic guide more indispensable than ever. As taxonomists become extinct, there are fewer students to receive the vast body of knowledge accumulated by generations of careful study. I hope that the beauty and depth of this guide will inspire a generation of young scientists to continue this critical taxonomic work. It will have a place of honor in all marine labs.--Paul K. Dayton, Scripps Institution of Oceanography

This volume offers a comprehensive history of the Mount Desert Island Biological Laboratory (MDIBL), one of the major marine laboratories in the United States and a leader in using marine organisms to study fundamental physiological concepts. Beginning with its founding as the Harpswell Laboratory of Tufts University in 1898, David H. Evans follows its evolution from a teaching facility to a research center for distinguished renal and epithelial physiologists. He also describes how it became the site of major advances in cytokinesis, regeneration, cardiac and vascular physiology, hepatic physiology, endocrinology and toxicology, as well as studies of the comparative

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physiology of marine organisms. Fundamental physiological concepts in the context of the discoveries made at the MDIBL are explained and the social and administrative history of this renowned facility is described.

Textbook for Cell and Molecular Biology.

This handbook provides guidance on methods for use in fish ageing studies. Fish age, among other biological parameters, is one of the most relevant pieces of data to attain sustainable exploitation levels of fishery resources. Indeed, most analytical methods used in stock assessment require knowledge of demographic structure according to age of stocks, as well as to recruitment, growth, maturity, natural mortality, etc., which are strictly linked to information on age and age structure. The handbook was put together to deal with identified gaps - which affect both the precision and the accuracy of estimations - regarding ageing schemes, criteria and methodologies used in preparing calcified structures. It focuses on the general principles that underpin age analysis (assignment of birth date, preparation methods, ageing scheme reading and identification of true and false rings). Crucially, it aims to contribute to the establishment of common analysis methods, which can enable better calibration across

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the diverse institutes involved, thereby improving the quality and reliability of results.

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Sequence - Evolution - Function is an introduction to the computational approaches that play a critical role in the emerging new branch of biology known as functional genomics. The book provides the reader with an understanding of the principles and approaches of functional genomics and of the potential and limitations of computational and experimental approaches to genome analysis. Sequence - Evolution - Function should help bridge the "digital divide" between biologists and computer scientists, allowing biologists to better grasp the peculiarities of the emerging field of Genome Biology and to learn how to benefit from the enormous amount of sequence data available in the public databases. The book is non-technical with respect to the computer methods for genome analysis and discusses these methods from the user's viewpoint, without addressing mathematical and algorithmic details. Prior practical familiarity with the basic methods for sequence analysis is a major advantage, but a reader without such experience will be able to use the book as an

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introduction to these methods. This book is perfect for introductory level courses in computational methods for comparative and functional genomics.

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