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A Doody's Core Title for 2015. Molecular Biology, 5/e by Robert Weaver, is designed for an introductory course in molecular biology. Molecular Biology 5/e focuses on the fundamental concepts of molecular biology emphasizing experimentation. In particular author, Rob Weaver, focuses on the study of genes and their activities at the molecular level. Through the combination of excellent illustrations and clear, succinct writing students are presented fundamental molecular biology concepts.

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As the pharmaceutical industry continues to advance, new techniques in drug design are emerging. In order to deliver optimum care to patients, the development of innovative pharmacological techniques has become a widely studied topic. Applied Case Studies and Solutions in Molecular Docking-Based Drug Design is a pivotal reference source for the latest scholarly research on the progress of pharmaceutical design and computational approaches in the field of molecular docking. Highlighting innovative research perspectives and real-world applications, this book is ideally designed for professionals, researchers, practitioners, and medical chemists actively involved in computational chemistry and pharmaceutical sciences.

This book is the first of its kind to provide a large collection of bioinformatics problems with accompanying solutions. Notably, the problem set includes all of the problems offered in Biological Sequence Analysis (BSA), by Durbin et al., widely adopted as a required text for bioinformatics courses at leading universities worldwide. Although many of the problems included in BSA as exercises for its readers have been repeatedly used for homework and tests, no detailed solutions for the problems were available. Bioinformatics instructors had therefore frequently expressed a need for fully worked solutions and a larger set of problems for use on courses. This book provides just that: following the same structure as BSA and significantly extending the set of workable problems, it will facilitate a better understanding of the contents of the chapters in BSA and will help its readers develop problem-solving skills that are vitally important for conducting successful research in the growing field of bioinformatics. All of the material has been class-tested by the authors at Georgia Tech, where the first ever M.Sc. degree program in Bioinformatics was held.

Based on formerly untapped archival sources as well as on interviews of participants, and building upon prior historical literature, Shaping Biology covers new ground and raises significant issues for further research on postwar biology and on federal funding of science in general.

Small-angle scattering of X-rays or neutrons is a technique that allows one to study the structures and interactions of disordered materials like polymers in the solid state, melt or solution or metal clusters in alloys. It is also the method of choice to characterize biological macromolecules in solution, in particular when they cannot be crystallized. A further advantage of the technique is that it can easily be combined with standard perturbation methods such as temperature and pressure jumps and stopped flow mixing thus offering useful information complementary to spectroscopic methods. The book describes all aspects of the technique: instrumentation, sample requirements, data interpretation and modelling methods in a comprehensive way and gives examples of applications in various fields of biophysics and biochemistry. Appendices describe the mathematical background and additional resources relevant to the method.

A leading microbiologist provides thought-provoking insights into the question of "What is Life?" as he examines the relationship of living things to the inorganic realms of physics and chemistry, explains how lifeless chemicals come together to form living beings, and details the true complexity of seemingly simple microorganisms such as E. coli.

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Developed to provide a comprehensive guide, the Handbook of Medicinal Chemistry has been revised and brought up to date to cover the past, present and future of the entire drug development process.

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