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Biomedical Politics - Institute of Medicine 1991-02-01

The abortifacient RU-486 was born in the laboratory, but its history has been shaped by legislators, corporate marketing executives, and protesters on both sides of the abortion debate. This volume explores how society decides what to do when discoveries such as RU-486 raise complex and emotional policy issues. Six case studies with insightful commentary offer a revealing look at the interplay of scientists, interest groups, the U.S. Congress, federal agencies, and the public in determining biomedical public policy and suggest how decision making might become more reasoned and productive in the future. The studies are fascinating and highly readable accounts of the personal interactions behind the headlines. They cover dideoxyinosine (ddI), RU-486, Medicare coverage for victims of chronic kidney failure, the human genome project, fetal tissue transplantation, and the 1975 Asilomar conference on recombinant DNA.

Genomic Medicine - Dhavendra Kumar 2014-10-15

Preceded by Genomics and clinical medicine / edited by Dhavendra Kumar. [First edition]. 2008.

Molecular Biotechnology - Glick Bernard R 1998

The second edition explains the principles of recombinant DNA technology as well as other important techniques such as DNA sequencing, the polymerase chain reaction, and the production of monoclonal antibodies.

CRISPR-Cas Systems - Rodolphe Barrangou 2012-12-13

CRISPR/Cas is a recently described defense system that protects bacteria and archaea against invasion by mobile genetic elements such as viruses and plasmids. A wide spectrum of distinct CRISPR/Cas systems has been identified in at least half of the available prokaryotic genomes. On-going structural and functional analyses have resulted in a far greater insight into the functions and possible applications of these systems, although many secrets remain to be discovered. In this book, experts summarize the state of the art in this exciting field.

Basic Biotechnology - Colin Ratledge 2006-05-25

Biotechnology is one of the major technologies of the twenty-first century. Its wide-ranging, multi-disciplinary activities include recombinant DNA techniques, cloning and the application of microbiology to the production of goods from bread to antibiotics. In this new edition of the textbook Basic Biotechnology, biology and bioprocessing topics are uniquely combined to provide a complete overview of biotechnology. The fundamental principles that underpin all biotechnology are explained and a full range of examples are discussed to show how these principles are applied; from starting substrate to final product. A distinctive feature of this text are the discussions of the public perception of biotechnology and the business of biotechnology, which set the science in a broader context. This comprehensive textbook is essential reading for all students of biotechnology and applied microbiology, and for researchers in biotechnology industries.

Exploration and Analysis of DNA Microarray and Other High-Dimensional Data - Dhammika Amaratunga 2014-01-27

Praise for the First Edition "...extremely well written...a comprehensive and up-to-date overview of this important field." -Journal of Environmental Quality Exploration and Analysis of DNA Microarray and

Other High-Dimensional Data, Second Edition provides comprehensive coverage of recent advancements in microarray data analysis. A cutting-edge guide, the Second Edition demonstrates various methodologies for analyzing data in biomedical research and offers an overview of the modern techniques used in microarray technology to study patterns of gene activity. The new edition answers the need for an efficient outline of all phases of this revolutionary analytical technique, from preprocessing to the analysis stage. Utilizing research and experience from highly-qualified authors in fields of data analysis, Exploration and Analysis of DNA Microarray and Other High-Dimensional Data, Second Edition features: A new chapter on the interpretation of findings that includes a discussion of signatures and material on gene set analysis, including network analysis. New topics of coverage including ABC clustering, biclustering, partial least squares, penalized methods, ensemble methods, and enriched ensemble methods. Updated exercises to deepen knowledge of the presented material and provide readers with resources for further study. The book is an ideal reference for scientists in biomedical and genomics research fields who analyze DNA microarrays and protein array data, as well as statisticians and bioinformatics practitioners. Exploration and Analysis of DNA Microarray and Other High-Dimensional Data, Second Edition is also a useful text for graduate-level courses on statistics, computational biology, and bioinformatics.

Genes and Genomes - R.S. Verma 1998-06-03

The laws of inheritance were considered quite superficial until 1903, when the chromosome theory of heredity was established by Sutton and Boveri. The discovery of the double helix and the genetic code led to our understanding of gene structure and function. For the past quarter of a century, remarkable progress has been made in the characterization of the human genome in order to search for coherent views of genes. The unit of inheritance termed factor or gene, once upon a time thought to be a trivial and imaginary entity, is now perceived clearly as the precise unit of inheritance that has continually deluged us with amazement by its complex identity and behaviour, sometimes bypassing the universality of Mendel's law. The aim of the fifth volume, entitled Genes and Genomes, is to cover the topics ranging from the structure of DNA itself to the structure of the complete genome, along with everything in between, encompassing 12 chapters. These chapters relate much of the information accumulated on the role of DNA in the organization of genes and genomes per se. Several distinguished scientists, all pre-eminent authorities in each field to share their expertise. Obviously, since the historical report on the double helix configuration in 1953, voluminous reports on the meteoric advances in genetics have been accumulated, and to cover every account in a single volume format would be a Herculean task. Therefore, only a few topics are chosen, which are of great interest to molecular geneticists. This volume is intended for advanced graduate students who would wish to keep abreast with the most recent trends in genome biology.

Biotechnology - David P. Clark 2015-05-16

Biotechnology, Second Edition approaches modern biotechnology from a molecular basis, which has grown out of increasing biochemical understanding of genetics and physiology. Using straightforward, less-technical jargon, Clark and Pazdernik introduce each chapter with basic concepts that develop into more

specific and detailed applications. This up-to-date text covers a wide realm of topics including forensics, bioethics, and nanobiotechnology using colorful illustrations and concise applications. In addition, the book integrates recent, relevant primary research articles for each chapter, which are presented on an accompanying website. The articles demonstrate key concepts or applications of the concepts presented in the chapter, which allows the reader to see how the foundational knowledge in this textbook bridges into primary research. This book helps readers understand what molecular biotechnology actually is as a scientific discipline, how research in this area is conducted, and how this technology may impact the future. Up-to-date text focuses on modern biotechnology with a molecular foundation. Includes clear, color illustrations of key topics and concept. Features clearly written without overly technical jargon or complicated examples. Provides a comprehensive supplements package with an easy-to-use study guide, full primary research articles that demonstrate how research is conducted, and instructor-only resources.

Human Genetics and Genomics - Bruce R. Korf 2012-11-19

This fourth edition of the best-selling textbook, *Human Genetics and Genomics*, clearly explains the key principles needed by medical and health sciences students, from the basis of molecular genetics, to clinical applications used in the treatment of both rare and common conditions. A newly expanded Part 1, *Basic Principles of Human Genetics*, focuses on introducing the reader to key concepts such as Mendelian principles, DNA replication and gene expression. Part 2, *Genetics and Genomics in Medical Practice*, uses case scenarios to help you engage with current genetic practice. Now featuring full-color diagrams, *Human Genetics and Genomics* has been rigorously updated to reflect today's genetics teaching, and includes updated discussion of genetic risk assessment, "single gene" disorders and therapeutics. Key learning features include: Clinical snapshots to help relate science to practice 'Hot topics' boxes that focus on the latest developments in testing, assessment and treatment 'Ethical issues' boxes to prompt further thought and discussion on the implications of genetic developments 'Sources of information' boxes to assist with the practicalities of clinical research and information provision Self-assessment review questions in each chapter Accompanied by the Wiley E-Text digital edition (included in the price of the book), *Human Genetics and Genomics* is also fully supported by a suite of online resources at www.korfgenetics.com, including: Factsheets on 100 genetic disorders, ideal for study and exam preparation Interactive Multiple Choice Questions (MCQs) with feedback on all answers Links to online resources for further study Figures from the book available as PowerPoint slides, ideal for teaching purposes The perfect companion to the genetics component of both problem-based learning and integrated medical courses, *Human Genetics and Genomics* presents the ideal balance between the bio-molecular basis of genetics and clinical cases, and provides an invaluable overview for anyone wishing to engage with this fast-moving discipline.

Pan-genomics: Applications, Challenges, and Future Prospects - Debmalya Barh 2020-03-06

Pan-genomics: Applications, Challenges, and Future Prospects covers current approaches, challenges and future prospects of pan-genomics. The book discusses bioinformatics tools and their applications and focuses on bacterial comparative genomics in order to leverage the development of precise drugs and treatments for specific organisms. The book is divided into three sections: the first, an "overview of pan-genomics and common approaches, brings the main concepts and current approaches on pan-genomics research; the second, "case studies in pan-genomics, thoroughly discusses twelve case, and the last, "current approaches and future prospects in pan-multiomics, encompasses the developments on omics studies to be applied on bacteria related studies. This book is a valuable source for bioinformaticians, genomics researchers and several members of biomedical field interested in understanding further bacterial organisms and their relationship to human health. Covers the entire spectrum of pangenomics, highlighting the use of specific approaches, case studies and future perspectives Discusses current bioinformatics tools and strategies for exploiting pangenomics data Presents twelve case studies with different organisms in order to provide the audience with real examples of pangenomics applicability

The Gene - Siddhartha Mukherjee 2016-05-17

The #1 NEW YORK TIMES Bestseller The basis for the PBS Ken Burns Documentary *The Gene: An Intimate History* Now includes an excerpt from Siddhartha Mukherjee's new book *Song of the Cell!* From the Pulitzer Prize-winning author of *The Emperor of All Maladies*—a fascinating history of the gene and "a magisterial account of how human minds have laboriously, ingeniously picked apart what makes us tick"

(Elle). "Sid Mukherjee has the uncanny ability to bring together science, history, and the future in a way that is understandable and riveting, guiding us through both time and the mystery of life itself." —Ken Burns "Dr. Siddhartha Mukherjee dazzled readers with his Pulitzer Prize-winning *The Emperor of All Maladies* in 2010. That achievement was evidently just a warm-up for his virtuoso performance in *The Gene: An Intimate History*, in which he braids science, history, and memoir into an epic with all the range and biblical thunder of *Paradise Lost*" (The New York Times). In this biography Mukherjee brings to life the quest to understand human heredity and its surprising influence on our lives, personalities, identities, fates, and choices. "Mukherjee expresses abstract intellectual ideas through emotional stories...[and] swaddles his medical rigor with rhapsodic tenderness, surprising vulnerability, and occasional flashes of pure poetry" (The Washington Post). Throughout, the story of Mukherjee's own family—with its tragic and bewildering history of mental illness—reminds us of the questions that hang over our ability to translate the science of genetics from the laboratory to the real world. In riveting and dramatic prose, he describes the centuries of research and experimentation—from Aristotle and Pythagoras to Mendel and Darwin, from Boveri and Morgan to Crick, Watson and Franklin, all the way through the revolutionary twenty-first century innovators who mapped the human genome. "A fascinating and often sobering history of how humans came to understand the roles of genes in making us who we are—and what our manipulation of those genes might mean for our future" (Milwaukee Journal-Sentinel), *The Gene* is the revelatory and magisterial history of a scientific idea coming to life, the most crucial science of our time, intimately explained by a master. "The Gene is a book we all should read" (USA TODAY).

Gene Cloning and DNA Analysis - T. A. Brown 2013-04-25

Known world-wide as the standard introductory text to this important and exciting area, the sixth edition of *Gene Cloning and DNA Analysis* addresses new and growing areas of research whilst retaining the philosophy of the previous editions. Assuming the reader has little prior knowledge of the subject, its importance, the principles of the techniques used and their applications are all carefully laid out, with over 250 clearly presented four-colour illustrations. In addition to a number of informative changes to the text throughout the book, the final four chapters have been significantly updated and extended to reflect the striking advances made in recent years in the applications of gene cloning and DNA analysis in biotechnology. *Gene Cloning and DNA Analysis* remains an essential introductory text to a wide range of biological sciences students; including genetics and genomics, molecular biology, biochemistry, immunology and applied biology. It is also a perfect introductory text for any professional needing to learn the basics of the subject. All libraries in universities where medical, life and biological sciences are studied and taught should have copies available on their shelves. "... the book content is elegantly illustrated and well organized in clear-cut chapters and subsections... there is a Further Reading section after each chapter that contains several key references... What is extremely useful, almost every reference is furnished with the short but distinct author's remark." -Journal of Heredity, 2007 (on the previous edition)

Concepts and Techniques in Genomics and Proteomics - N Saraswathy 2011-07-01

Concepts and techniques in genomics and proteomics covers the important concepts of high-throughput modern techniques used in the genomics and proteomics field. Each technique is explained with its underlying concepts, and simple line diagrams and flow charts are included to aid understanding and memory. A summary of key points precedes each chapter within the book, followed by detailed description in the subsections. Each subsection concludes with suggested relevant original references. Provides definitions for key concepts Case studies are included to illustrate ideas Important points to remember are noted

Genomics in Drug Discovery and Development - Dimitri Semizarov 2008-11-03

Early characterization of toxicity and efficacy would significantly impact the overall productivity of pharmaceutical R&D and reduce drug candidate attrition and failure. By describing the available platforms and weighing their relative advantages and disadvantages, including microarray data analysis, *Genomics in Drug Discovery and Development* introduces readers to the biomarker, pharmacogenomic, and toxicogenomics toolbox. The authors provide a valuable resource for pharmaceutical discovery scientists, preclinical drug safety department personnel, regulatory personnel, discovery toxicologists, and safety scientists, drug development professionals, and pharmaceutical scientists.

Guidelines for Research Involving Recombinant DNA Molecules - 1980

DNA Technology in Forensic Science - National Research Council 1992-02-01

Matching DNA samples from crime scenes and suspects is rapidly becoming a key source of evidence for use in our justice system. DNA Technology in Forensic Science offers recommendations for resolving crucial questions that are emerging as DNA typing becomes more widespread. The volume addresses key issues: Quality and reliability in DNA typing, including the introduction of new technologies, problems of standardization, and approaches to certification. DNA typing in the courtroom, including issues of population genetics, levels of understanding among judges and juries, and admissibility. Societal issues, such as privacy of DNA data, storage of samples and data, and the rights of defendants to quality testing technology. Combining this original volume with the new update-The Evaluation of Forensic DNA Evidence- provides the complete, up-to-date picture of this highly important and visible topic. This volume offers important guidance to anyone working with this emerging law enforcement tool: policymakers, specialists in criminal law, forensic scientists, geneticists, researchers, faculty, and students.

Wilson and Walker's Principles and Techniques of Biochemistry and Molecular Biology - Andreas Hofmann 2018-04-19

Bringing this best-selling textbook right up to date, the new edition uniquely integrates the theories and methods that drive the fields of biology, biotechnology and medicine, comprehensively covering both the techniques students will encounter in lab classes and those that underpin current key advances and discoveries. The contents have been updated to include both traditional and cutting-edge techniques most commonly used in current life science research. Emphasis is placed on understanding the theory behind the techniques, as well as analysis of the resulting data. New chapters cover proteomics, genomics, metabolomics, bioinformatics, as well as data analysis and visualisation. Using accessible language to describe concepts and methods, and with a wealth of new in-text worked examples to challenge students' understanding, this textbook provides an essential guide to the key techniques used in current bioscience research.

Computational Genomics with R - Altuna Akalin 2020-12-16

Computational Genomics with R provides a starting point for beginners in genomic data analysis and also guides more advanced practitioners to sophisticated data analysis techniques in genomics. The book covers topics from R programming, to machine learning and statistics, to the latest genomic data analysis techniques. The text provides accessible information and explanations, always with the genomics context in the background. This also contains practical and well-documented examples in R so readers can analyze their data by simply reusing the code presented. As the field of computational genomics is interdisciplinary, it requires different starting points for people with different backgrounds. For example, a biologist might skip sections on basic genome biology and start with R programming, whereas a computer scientist might want to start with genome biology. After reading: You will have the basics of R and be able to dive right into specialized uses of R for computational genomics such as using Bioconductor packages. You will be familiar with statistics, supervised and unsupervised learning techniques that are important in data modeling, and exploratory analysis of high-dimensional data. You will understand genomic intervals and operations on them that are used for tasks such as aligned read counting and genomic feature annotation. You will know the basics of processing and quality checking high-throughput sequencing data. You will be able to do sequence analysis, such as calculating GC content for parts of a genome or finding transcription factor binding sites. You will know about visualization techniques used in genomics, such as heatmaps, meta-gene plots, and genomic track visualization. You will be familiar with analysis of different high-throughput sequencing data sets, such as RNA-seq, ChIP-seq, and BS-seq. You will know basic techniques for integrating and interpreting multi-omics datasets. Altuna Akalin is a group leader and head of the Bioinformatics and Omics Data Science Platform at the Berlin Institute of Medical Systems Biology, Max Delbrück Center, Berlin. He has been developing computational methods for analyzing and integrating large-scale genomics data sets since 2002. He has published an extensive body of work in this area. The framework for this book grew out of the yearly computational genomics courses he has been organizing and teaching since 2015.

Environmental Microbiology - Eugene L. Madsen 2015-09-28

New and expanded for its second edition, *Environmental Microbiology: From Genomes to Biogeochemistry*, Second Edition, is a timely update to a classic text filled with ideas, connections, and concepts that advance an in-depth understanding of this growing segment of microbiology. Core principles are highlighted with an emphasis on the logic of the science and new methods-driven discoveries. Numerous up-to-date examples and applications boxes provide tangible reinforcement of material covered. Study questions at the end of each chapter require students to utilize analytical and quantitative approaches, to define and defend arguments, and to apply microbiological paradigms to their personal interests. Essay assignments and related readings stimulate student inquiry and serve as focal points for teachers to launch classroom discussions. A companion website with downloadable artwork and answers to study questions is also available. *Environmental Microbiology: From Genomes to Biogeochemistry*, Second Edition, offers a coherent and comprehensive treatment of this dynamic, emerging field, building bridges between basic biology, evolution, genomics, ecology, biotechnology, climate change, and the environmental sciences.

DNA Recombination and Repair - Paul James Smith 1999

The processes of DNA recombination and repair are vital to cell integrity - an error can lead to disease such as cancer. It is therefore a large and exciting area of research and is also taught on postgraduate and undergraduate courses. This book is not a comprehensive view of the field, but a selection of the issues currently at the forefront of knowledge.

Mapping and Sequencing the Human Genome - National Research Council 1988-01-01

There is growing enthusiasm in the scientific community about the prospect of mapping and sequencing the human genome, a monumental project that will have far-reaching consequences for medicine, biology, technology, and other fields. But how will such an effort be organized and funded? How will we develop the new technologies that are needed? What new legal, social, and ethical questions will be raised? *Mapping and Sequencing the Human Genome* is a blueprint for this proposed project. The authors offer a highly readable explanation of the technical aspects of genetic mapping and sequencing, and they recommend specific interim and long-range research goals, organizational strategies, and funding levels. They also outline some of the legal and social questions that might arise and urge their early consideration by policymakers.

The Epigenetics Revolution - Nessa Carey 2012-03-06

Epigenetics can potentially revolutionize our understanding of the structure and behavior of biological life on Earth. It explains why mapping an organism's genetic code is not enough to determine how it develops or acts and shows how nurture combines with nature to engineer biological diversity. Surveying the twenty-year history of the field while also highlighting its latest findings and innovations, this volume provides a readily understandable introduction to the foundations of epigenetics. Nessa Carey, a leading epigenetics researcher, connects the field's arguments to such diverse phenomena as how ants and queen bees control their colonies; why tortoiseshell cats are always female; why some plants need cold weather before they can flower; and how our bodies age and develop disease. Reaching beyond biology, epigenetics now informs work on drug addiction, the long-term effects of famine, and the physical and psychological consequences of childhood trauma. Carey concludes with a discussion of the future directions for this research and its ability to improve human health and well-being.

Evolution of Translational Omics - Institute of Medicine 2012-09-13

Technologies collectively called omics enable simultaneous measurement of an enormous number of biomolecules; for example, genomics investigates thousands of DNA sequences, and proteomics examines large numbers of proteins. Scientists are using these technologies to develop innovative tests to detect disease and to predict a patient's likelihood of responding to specific drugs. Following a recent case involving premature use of omics-based tests in cancer clinical trials at Duke University, the NCI requested that the IOM establish a committee to recommend ways to strengthen omics-based test development and evaluation. This report identifies best practices to enhance development, evaluation, and translation of omics-based tests while simultaneously reinforcing steps to ensure that these tests are appropriately assessed for scientific validity before they are used to guide patient treatment in clinical trials.

Molecular Biology - David P. Clark 2012-03-20

Molecular Biology, Second Edition, examines the basic concepts of molecular biology while incorporating primary literature from today's leading researchers. This updated edition includes Focuses on Relevant Research sections that integrate primary literature from Cell Press and focus on helping the student learn how to read and understand research to prepare them for the scientific world. The new Academic Cell Study Guide features all the articles from the text with concurrent case studies to help students build foundations in the content while allowing them to make the appropriate connections to the text. Animations provided deal with topics such as protein purification, transcription, splicing reactions, cell division and DNA replication and SDS-PAGE. The text also includes updated chapters on Genomics and Systems Biology, Proteomics, Bacterial Genetics and Molecular Evolution and RNA. An updated ancillary package includes flashcards, online self quizzing, references with links to outside content and PowerPoint slides with images. This text is designed for undergraduate students taking a course in Molecular Biology and upper-level students studying Cell Biology, Microbiology, Genetics, Biology, Pharmacology, Biotechnology, Biochemistry, and Agriculture. NEW: "Focus On Relevant Research" sections integrate primary literature from Cell Press and focus on helping the student learn how to read and understand research to prepare them for the scientific world. NEW: Academic Cell Study Guide features all articles from the text with concurrent case studies to help students build foundations in the content while allowing them to make the appropriate connections to the text. NEW: Animations provided include topics in protein purification, transcription, splicing reactions, cell division and DNA replication and SDS-PAGE Updated chapters on Genomics and Systems Biology, Proteomics, Bacterial Genetics and Molecular Evolution and RNA Updated ancillary package includes flashcards, online self quizzing, references with links to outside content and PowerPoint slides with images. Fully revised art program

Biology for AP® Courses - Julianne Zedalis 2017-10-16

Biology for AP® courses covers the scope and sequence requirements of a typical two-semester Advanced Placement® biology course. The text provides comprehensive coverage of foundational research and core biology concepts through an evolutionary lens. Biology for AP® Courses was designed to meet and exceed the requirements of the College Board's AP® Biology framework while allowing significant flexibility for instructors. Each section of the book includes an introduction based on the AP® curriculum and includes rich features that engage students in scientific practice and AP® test preparation; it also highlights careers and research opportunities in biological sciences.

Biotechnology - David P. Clark 2015-06-22

Biotechnology, Second Edition approaches modern biotechnology from a molecular basis, which has grown out of increasing biochemical understanding of genetics and physiology. Using straightforward, less-technical jargon, Clark and Pazdernik introduce each chapter with basic concepts that develop into more specific and detailed applications. This up-to-date text covers a wide realm of topics including forensics, bioethics, and nanobiotechnology using colorful illustrations and concise applications. In addition, the book integrates recent, relevant primary research articles for each chapter, which are presented on an accompanying website. The articles demonstrate key concepts or applications of the concepts presented in the chapter, which allows the reader to see how the foundational knowledge in this textbook bridges into primary research. This book helps readers understand what molecular biotechnology actually is as a scientific discipline, how research in this area is conducted, and how this technology may impact the future. Up-to-date text focuses on modern biotechnology with a molecular foundation Includes clear, color illustrations of key topics and concept Features clearly written without overly technical jargon or complicated examples Provides a comprehensive supplements package with an easy-to-use study guide, full primary research articles that demonstrate how research is conducted, and instructor-only resources

Darwin in the Genome - Lynn Helena Caporale 2003

Smart genomes--an enthralling account of revolutionary discoveries at the cutting edge of genomics research Written by a molecular biologist at the forefront of genomics research, Darwin in the Genome is an exciting account of one of the hottest new theories in biology today: evolution by natural selection inevitably leads to strategic mutations. In the struggle for survival, from pathogens to flowers, birds to orangutans, baker's yeast to people, the fittest genomes are those that evolve effective molecular strategies that respond to, and in fact anticipate, challenges and opportunities in their environments. Writing in a

clear, accessible style, Lynn Caporale describes the emergence of genomic mutation strategies, which researchers are just beginning to uncover. She also spells out some of the more profound implications of these findings, including the importance of biodiversity, indeed human diversity, for survival, the possibility of bold new directions for medical research, and the inherent dangers of attempting to fix perceived "errors" in a human genome.

The Double Helix - James D. Watson 2011-08-16

The classic personal account of Watson and Crick's groundbreaking discovery of the structure of DNA, now with an introduction by Sylvia Nasar, author of A Beautiful Mind. By identifying the structure of DNA, the molecule of life, Francis Crick and James Watson revolutionized biochemistry and won themselves a Nobel Prize. At the time, Watson was only twenty-four, a young scientist hungry to make his mark. His uncompromisingly honest account of the heady days of their thrilling sprint against other world-class researchers to solve one of science's greatest mysteries gives a dazzlingly clear picture of a world of brilliant scientists with great gifts, very human ambitions, and bitter rivalries. With humility unspoiled by false modesty, Watson relates his and Crick's desperate efforts to beat Linus Pauling to the Holy Grail of life sciences, the identification of the basic building block of life. Never has a scientist been so truthful in capturing in words the flavor of his work.

The Absolute, Ultimate Guide to Lehninger Principles of Biochemistry - Marcy Osgood 2008-04-04

The Absolute, Ultimate Guide combines an innovative study guide with a reliable solutions manual in one convenient printed volume.

Gene Cloning and DNA Analysis - T. A. Brown 2001-10-15

The previous edition of Gene Cloning, has become known world-wide as the standard introductory text to this important and exciting subject. Now, the current importance of non-cloning approaches, notably PCR, in gene cloning has led to the creation of this fourth edition which explores DNA analysis. Retaining the philosophy of the previous editions, Gene Cloning and DNA Analysis assumes the reader has little prior knowledge of the subject and clearly explains its importance, the principles of the techniques used and their applications. It is carefully laid out, with over 250 two-colour illustrations. In addition to some organisational changes within the re-written text, the fourth edition has two new chapters. The first covers the methods used to sequence genomes and how to understand a sequence after it has been obtained. The second looks in detail at the applications of gene cloning and DNA analysis in forensic science; providing an excellent illustration of the applications of DNA analysis in the real world. In Gene Cloning and DNA Analysis Terry Brown has once again provided a resource of exceptional clarity, an essential introductory text to a wide range of biological sciences students, including genetics and genomics, molecular biology, biochemistry, immunology and applied biology. Also, as well as being required reading for many course modules, it is a perfect introductory text for any professional who needs to learn the basics of the subject. All libraries in universities where medical, life and biological sciences are studied or taught should have multiple copies of the book available on their shelves. Terry Brown is Professor of Biomolecular Archaeology at the Department of Biomolecular Sciences, University of Manchester Institute of Science and Technology, Manchester, UK. Visit the companion website for more details about the book: <http://www.blackwellpublishing.com/genecloning/> Fourth edition of internationally popular undergraduate text. Terry Brown is world famous for his work and writing in the area of genomes, genetics and gene cloning. Fully updated and revised. Two-colour text including approximately 250 two-colour line illustrations. New dedicated web pages for the book, including student questions, illustrations and supplementary material.

Concepts of Biology - Samantha Fowler 2018-01-07

Concepts of Biology is designed for the single-semester introduction to biology course for non-science majors, which for many students is their only college-level science course. As such, this course represents an important opportunity for students to develop the necessary knowledge, tools, and skills to make informed decisions as they continue with their lives. Rather than being mired down with facts and vocabulary, the typical non-science major student needs information presented in a way that is easy to read and understand. Even more importantly, the content should be meaningful. Students do much better when they understand why biology is relevant to their everyday lives. For these reasons, Concepts of Biology is

grounded on an evolutionary basis and includes exciting features that highlight careers in the biological sciences and everyday applications of the concepts at hand. We also strive to show the interconnectedness of topics within this extremely broad discipline. In order to meet the needs of today's instructors and students, we maintain the overall organization and coverage found in most syllabi for this course. A strength of Concepts of Biology is that instructors can customize the book, adapting it to the approach that works best in their classroom. Concepts of Biology also includes an innovative art program that incorporates critical thinking and clicker questions to help students understand--and apply--key concepts.

Biological Science - Jon (Emeritus Professor of Bioscience Education Scott, Emeritus Professor of Bioscience Education University of Leicester) 2022-06-24

Biological Science: Exploring the Science of Life responds to the key needs of lecturers and their students by placing a clear central narrative, carefully-structured active learning, and confidence with quantitative concepts and scientific enquiry central to its approach. Written by a team of dedicated and passionate academics, and shaped by feedback from over 55 institutions, its straightforward narrative, reinforced by key concept overview videos for every chapter, communicate key ideas clearly: the right information is provided at the right time, and at the right depth. Its pause and think features, self-check quizzes, and graded end of chapter questions, augmented by flashcards of key terms, directly support active learning. The combination of narrative text and learning features promote a rich, active learning experience: read, watch, and do. Its combination of Quantitative Toolkits, Scientific Process panels, and the Life and its Exploration chapters provide more insight and support than any other general biology text; they prepare students to engage with this quantitative and experimental discipline with confidence, and set them on a path for success throughout their future studies. With coverage that spans the full scale of biological science - from molecule to ecosystem - and with an approach that fully supports flexible, self-paced learning, Biological Science: Exploring the Science of Life will set you on a path towards a deeper understanding of the key concepts in biology, and a greater appreciation of biology as a dynamic experimental science. Digital formats and resources

Biological Science: Exploring the Science of Life is available for students and institutions to purchase in a variety of formats. The enhanced ebook is enriched with features that offer extra learning support: www.oxfordtextbooks.co.uk/ebooks- Key concepts videos support students from the start of every chapter and as they make their way through every Module.- Self-check questions at the end of each chapter section give students quick and formative feedback, building their confidence and comprehension as they study and revise.- Quantitative skills video screencasts help students to master the foundational skills required by this discipline.- Interactive figures give students the control they need to step through, and gain mastery over, key concepts.- Per-chapter flashcard glossaries help students to recall the key terms and concepts on which further study can be built.

Challenges in Delivery of Therapeutic Genomics and Proteomics - Ambikanandan Misra 2010-09-09

Delivery of therapeutic proteomics and genomics represent an important area of drug delivery research. Genomics and proteomics approaches could be used to direct drug development processes by unearthing pathways involved in disease pathogenesis where intervention may be most successful. This book describes the basics of genomics and proteomics and highlights the various chemical, physical and biological approaches to protein and gene delivery. Covers a diverse array of topics from basic sciences to therapeutic applications of proteomics and genomics delivery. Of interest to researchers in both academia and industry. Highlights what's currently known and where further research is needed.

DNA Technology - I. Edward Alcamo 2001

Gives the educated layperson a survey of DNA by presenting a brief history of genetics, an outline of techniques, and indications of breakthroughs in cloning and other DNA advances. This book helps students, business people, lawyers, and jurists gain confidence in their ability to understand and appreciate DNA technology and human genetics.

Biocode - Dawn Field 2015

The living world runs on genomic software - what Dawn Field and Neil Davies call the 'biocode' - the sum of all DNA on Earth. In Biocode, they tell the story of a new age of scientific discovery: the growing global effort to read and map the biocode, and what that might mean for the future. The structure of DNA was identified in 1953, and the whole human genome was mapped by 2003. Since then the new field of

genomics has mushroomed and is now operating on an industrial scale. Genomes can now be sequenced rapidly and increasingly cheaply. The genomes of large numbers of organisms from mammals to microbes, have been mapped. Getting your genome sequenced is becoming affordable for many. You too can check paternity, find out where your ancestors came from, or whether you are at risk of some diseases. Some check out the pedigree of their pets, while others turn genomes into art. A stray hair is enough to crudely reconstruct the face of the owner. From reading to constructing: the first steps to creating artificial life have already been taken. Some may find the rapidity of developments, and the potential for misuse, alarming. But they also open up unprecedented possibilities. The ability to read DNA has changed how we view ourselves and understand our place in nature. From the largest oceans, to the insides of our guts, we are able to explore the biosphere as never before, from the genome up. Sequencing technology has made the invisible world of microbes visible, and biodiversity genomics is revealing whole new worlds within us and without. The findings are transformational: we are all ecosystems now. Already the first efforts at 'barcoding' entire ecological communities and creating 'genomic observatories' have begun. The future, the authors argue, will involve biocoding the entire planet.

Fundamentals of Food Biotechnology - Byong H. Lee 2015-02-16

Fundamentals of Food Biotechnology Food biotechnology is the application of modern biotechnological techniques to the manufacture and processing of food; for example, through fermentation of food (which is the oldest biotechnological process) and food additives, as well as plant and animal cell cultures. New developments in fermentation and enzyme technological processes, molecular thermodynamics, genetic engineering, protein engineering, metabolic engineering, bioengineering, and processes involving monoclonal antibodies, nanobiotechnology and quorum sensing have introduced exciting new dimensions to food biotechnology, a burgeoning field that transcends many scientific disciplines. Fundamentals of Food Biotechnology, 2nd edition is based on the author's 25 years of experience in teaching on a food biotechnology course at McGill University in Canada. The book will appeal to professional food scientists as well as graduate and advanced undergraduate students by addressing the latest exciting food biotechnology research in areas such as genetically modified foods (GMOs), bioenergy, bioplastics, functional foods/ nutraceuticals, nanobiotechnology, quorum sensing and quenching. In addition, cloning techniques for bacterial and yeast enzymes are included in a "New Trends and Tools" section and selected references, questions, and answers appear at the end of each chapter. This new edition has been comprehensively rewritten and restructured to reflect the new technologies, products, and trends that have emerged since the original book. Many new aspects highlight the short- and longer-term commercial potential of food biotechnology. Food Biochemistry and Food Processing, 2nd Edition Edited by Benjamin K. Simpson, Leo M.L. Nollet, Fidel Toldra, et al. ISBN 978-0-8138-0874-1 Food Processing: Principles and Applications, 2nd Edition Edited by Stephanie Clark (Editor), Stephanie Jung, Buddhi Lamsal ISBN 978-0-470-67114-6

Molecular Biology of the Cell - Bruce Alberts 2004

Advances in Animal Genomics - Sukanta Mondal 2020-11-25

Advances in Animal Genomics provides an outstanding collection of integrated strategies involving traditional and modern - omics (structural, functional, comparative and epigenomics) approaches and genomics-assisted breeding methods which animal biotechnologists can utilize to dissect and decode the molecular and gene regulatory networks involved in the complex quantitative yield and stress tolerance traits in livestock. Written by international experts on animal genomics, this book explores the recent advances in high-throughput, next-generation whole genome and transcriptome sequencing, array-based genotyping, and modern bioinformatics approaches which have enabled to produce huge genomic and transcriptomic resources globally on a genome-wide scale. This book is an important resource for researchers, students, educators and professionals in agriculture, veterinary and biotechnology sciences that enables them to solve problems regarding sustainable development with the help of current innovative biotechnologies. Integrates basic and advanced concepts of animal biotechnology and presents future developments. Describes current high-throughput next-generation whole genome and transcriptome sequencing, array-based genotyping, and modern bioinformatics approaches for sustainable livestock

production. Illustrates integrated strategies to dissect and decode the molecular and gene regulatory networks involved in complex quantitative yield and stress tolerance traits in livestock. Ensures readers will gain a strong grasp of biotechnology for sustainable livestock production with its well-illustrated discussion.

Genetically Engineered Crops - National Academies of Sciences, Engineering, and Medicine 2017-01-28

Genetically engineered (GE) crops were first introduced commercially in the 1990s. After two decades of production, some groups and individuals remain critical of the technology based on their concerns about possible adverse effects on human health, the environment, and ethical considerations. At the same time, others are concerned that the technology is not reaching its potential to improve human health and the environment because of stringent regulations and reduced public funding to develop products offering more benefits to society. While the debate about these and other questions related to the genetic engineering techniques of the first 20 years goes on, emerging genetic-engineering technologies are adding new complexities to the conversation. *Genetically Engineered Crops* builds on previous related Academies reports published between 1987 and 2010 by undertaking a retrospective examination of the purported positive and adverse effects of GE crops and to anticipate what emerging genetic-engineering technologies

hold for the future. This report indicates where there are uncertainties about the economic, agronomic, health, safety, or other impacts of GE crops and food, and makes recommendations to fill gaps in safety assessments, increase regulatory clarity, and improve innovations in and access to GE technology.

Principles of Genome Analysis - Sandy B. Primrose 1998-03-30

Genome analysis and genomics are at the forefront of current research in the life sciences. Since the first edition of *Principles of Genome Analysis* was published, the sequencing of genomes has continued apace, with the major landmark of the human genome sequence being achieved in 2001. Now the emphasis of biological research is on genomics: the understanding of gene function and the interaction of gene products at the whole genome level. As before, this book provides a step-by-step outline of the techniques involved in genome mapping and sequencing. Additionally, the text has been greatly expanded to cover sub-disciplines of genomics, revisions of sections on genome sequencing and bioinformatics, and new chapters on comparative genomics, functional genomics and proteomics. The book concludes with an exciting new chapter describing a variety of ways to utilize genome analysis and sequencing in biology, medicine and agriculture. Aimed at advanced undergraduates, this text will follow the same format as the highly successful *Principles of Gene Manipulation* by Primrose, Twyman and Old, now in its sixth edition.