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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

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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

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–Journal of Heredity, 2007 (on the previous edition)

Genetic programming (GP) is a systematic, domain-independent method for getting computers to solve problems automatically starting from a high-level statement of what needs to be done. Using ideas from

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natural evolution, GP starts from an ooze of random computer programs, and progressively refines them through processes of mutation and sexual recombination, until high-fitness solutions emerge. All this without the user having to know or specify the form or structure of solutions in advance. GP has generated a plethora of human-competitive results and applications, including novel scientific discoveries and patentable inventions. This unique overview of this exciting technique is written by three of the most active scientists in GP. See www.gp-field-guide.org.uk for more information on the book.

Biology of *Drosophila* was first published by John

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Wiley and Sons in 1950. Until its appearance, no central, synthesized source of biological data on *Drosophila melanogaster* was available, despite the fly's importance to science for three decades. Ten years in the making, it was an immediate success and remained in print for two decades. However, original copies are now very hard to find. This facsimile edition makes available to the fly community once again its most enduring work of reference.

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The genomics revolution has expanded from its origins in molecular biology to impact upon every discipline in the life sciences, including ecology.

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Several lines of ecological research can now be profitably addressed using genomics technology, including issues of nutrient cycling, population structure, life-history variation, trophic interaction, stress responses, and adaptation to environmental change. This new edition addresses a series of fundamental ecological questions: the relationship between community structure and ecological function in ecosystems; how variation in life-history patterns among species can be explained from interaction between the genome and the environment; the molecular responses to changing and toxic environmental conditions; adaptive phenotypes and their relationship to genetic variation. Each of these

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questions is evaluated in the light of recent advances in genomics research, paying particular attention to data obtained from sequencing and screening of environmental genomes (metagenomics), microarray-based transcription profiling, gene expression directed by signal-transduction pathways, and the analysis of genomic polymorphisms. The chapters covering these key areas are preceded by discussions of genomics methodology (including an overview of next-generation sequencing technologies) and comparative genomics, and the book concludes with a chapter on integrative approaches such as ecological control analysis. The authors also provide a comparative survey of the properties of genomes (genome size,

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gene families, synteny, and polymorphism) for prokaryotes as well as the main eukaryotic models. An Introduction to Ecological Genomics incorporates a balance of plant, animal, and microbial examples, and continues to define the new and exciting field of ecological genomics.

This publication is a comprehensive assessment of leading risks to global health. It provides detailed global and regional estimates of premature mortality, disability and loss of health attributable to 24 global risk factors.--Publisher's description.

Scores of talented and dedicated people serve the

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forensic science community, performing vitally important work. However, they are often constrained by lack of adequate resources, sound policies, and national support. It is clear that change and advancements, both systematic and scientific, are needed in a number of forensic science disciplines to ensure the reliability of work, establish enforceable standards, and promote best practices with consistent application. Strengthening Forensic Science in the United States: A Path Forward provides a detailed plan for addressing these needs and suggests the creation of a new government entity, the National Institute of Forensic Science, to establish and enforce standards within the forensic science community. The

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benefits of improving and regulating the forensic science disciplines are clear: assisting law enforcement officials, enhancing homeland security, and reducing the risk of wrongful conviction and exoneration. Strengthening Forensic Science in the United States gives a full account of what is needed to advance the forensic science disciplines, including upgrading of systems and organizational structures, better training, widespread adoption of uniform and enforceable best practices, and mandatory certification and accreditation programs. While this book provides an essential call-to-action for congress and policy makers, it also serves as a vital tool for law enforcement agencies, criminal prosecutors and

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attorneys, and forensic science educators.

This book provides a practical and self-contained overview of the Gene Ontology (GO), the leading project to organize biological knowledge on genes and their products across genomic resources. Written for biologists and bioinformaticians, it covers the state-of-the-art of how GO annotations are made, how they are evaluated, and what sort of analyses can and cannot be done with the GO. In the spirit of the Methods in Molecular Biology book series, there is an emphasis throughout the chapters on providing practical guidance and troubleshooting advice. Authoritative and accessible, The Gene Ontology

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