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This book features complete and original labs for the integrated laboratory. All materials, protocols, and equipment are spelled out. Each lab is customizable for your department. The book introduces and explains a wide range of lab techniques, and is geared to various ability levels. This volume is intended for chemistry instructors seeking to provide engaging and challenging labs that combine all the features and benefits of the integrated laboratory. Written by educators from around the country, each chapter of the

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book contains a fully detailed and explained experiment, with guidance for student questions and possible customization. The book offers students and instructors a wealth of learning opportunities in experiment preparation, measurement, recording and analysis from disciplines extending from biology and microbiology to geology, nanotechnology, and microelectronics. All experiments have been classroom tested, with safety and monitoring issues given precedence. Many of the experiments contain modules that permit the instructor to make the lab more challenging as time and student ability dictate.

DNA Methods in Clinical Microbiology describes the novel DNA-based technology now used in the diagnosis and management of infectious diseases. It is a concise, yet readable, overview written

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primarily for clinicians, clinical microbiologists, medical students and undergraduates in medical and veterinary microbiology. The book has two primary aims. First, to explain the principles of these methods at the 'molecular' level. Second, to provide a clinical perspective by reporting results from actual DNA-based investigations on a range of specimens. Those approaching DNA methods for the first time are assisted by a brief résumé of the relevant features of nucleic acids (Chapter 2): this information is essential for an understanding of later chapters. Subsequent text covers detection, characterization and quantification of pathogens by a variety of methods - e.g., target amplification (PCR, LCR, NASBA, TMA and SDA), signal amplification (bDNA) and probe-based techniques; the chapter on typing describes nearly twenty named molecular methods, including spoligotyping and MLST. All

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chapters include an adequate range of current reference from which, if required, detailed protocols can be obtained. The diagrams are clear, and readers are assisted by a detailed index.

This synthesis of the latest knowledge on homework presents unique findings by researchers from various countries and diverse professional backgrounds. It approaches the topic of homework from several perspectives, including its political and cultural contexts aspects of parental involvement and parent-child relationships school contexts and practices observable impacts It highlights homework-specific concerns and considers two principal solutions. Firstly, support initiatives from schools and communities. Secondly, improved homework design, aimed at attracting greater student interest and promoting communication within families.

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Recommendations for practice and future research are also discussed. A political analysis shows that current confusion about homework might stem from a tension between two prevailing ideologies, one stressing students' improved achievement in competing economies, and the other privileging collaboration between the school, family and community. This book considers how both teachers and parents can ensure a balance between the child's school life and his or her overall development. Certain community resources are available to parents and students, but these in no way absolve parents of the responsibility to maintain an interest in their child's school activities. *International Perspectives on Student Outcomes and Homework* proposes ideas and actions of relevance to everyone interested in the issue: school administrations, teachers in training and in practice, parents, and

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researchers eager to contribute to the advancement of knowledge in the field. It is a perfect companion to *International Perspectives on Contexts, Communities and Evaluated Innovative Practices*, also edited by Rollande Deslandes, and published simultaneously by Routledge.

PCR in situ hybridization allows the detection of specific nucleic acid sequences and their distribution in the cell. It combines two powerful techniques in situ hybridization (ISH), which allows cellular localization of DNA and RNA sequences in cells and tissues, and the polymerase chain reaction (PCR), which allows reproducible amplification of rare nucleic acid sequences. The combined technique and its variants greatly enhance the sensitivity of in situ hybridization and add morphological localization to the

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sensitivity of PCR. Such techniques have enormous potential for research and diagnosis but problems with reproducibility and reliability are often encountered. This book overcomes these problems by describing the key procedures in step-by-step detail and by providing the essential advice needed for success. Topics include: DNA in situ PCR and DNA PCR in situ hybridization (PCR ISH) for the detection of DNA targets in cells; reverse transcriptase in situ PCR (RT-PCR) and RT-PCR ISH for the detection of RNA targets; and PRINS (primed in situ synthesis) for chromosomal analysis in interphase nuclei and metaphase chromosome spreads. There are further chapters on fixation of tissues for PCR, selective ultraviolet radiation fractionation (SURF), application of in situ PCR to human tissues, applications and modifications of PCR-ISH, and automation of in situ

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amplification. PCR In Situ Hybridization is a unique and timely collection of well-tested protocols for the amplification of DNA and RNA in cells and tissues, drawing on the accumulated knowledge and experience of leading exponents of these techniques. For each topic covered, the authors provide detailed guidance on the key steps in the protocols, numerous hints and tips for success, and advice on trouble-shooting. PCR In Situ Hybridization will be invaluable to molecular biologists, pathologists, geneticists, and all those seeking to perform in situ analyses of nucleic acid molecules.

The book provides an updated overview of molecular analysis of human tissues, and the impact this analysis has on diagnosis and

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prognosis of human diseases. Special emphasis is placed on human cancer and the future directions of the field. Methods of handling clinical tissue samples, including the impact of handling on subsequent molecular analysis, are also discussed. In addition, detailed protocols for molecular analysis of DNA, RNA and protein, with special emphasis on molecular analysis of highly complex human tissue samples containing mixtures of cell populations, are provided.

Over the last decade there has been a rapid development of molecular techniques, with an increasing range of instrumentation now available. The development of accompanying reference literature has not kept pace with technological advances and this poses significant challenges to the analyst. Essentials of Nucleic

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Acid Analysis sets out to guide the analyst through the steps needed to obtain good quality results in DNA analysis. The underlying principles for achieving this goal were formulated by LGC (formerly the Laboratory of the Government Chemist) as the six principles for ensuring valid analytical measurement, which are detailed in the introduction. The reader is also provided with guidelines for method validation and quality control of established and emerging DNA measurement techniques. The authors of each chapter are practitioners of the art of DNA analysis in areas where the quality of the result is critical. Technical details and examples of application of key techniques in nucleic acid analysis are provided while highlighting best practice, available standards and practical advice on improving measurement quality. This book provides an indispensable handbook and premier reference for those working in

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the widely varying areas and specifically in the fields of food analysis and forensic applications.

Preceded by Avian influenza / edited by David E. Swayne. 2008.

Advanced Methods in Molecular Biology and Biotechnology: A Practical Lab Manual is a concise reference on common protocols and techniques for advanced molecular biology and biotechnology experimentation. Each chapter focuses on a different method, providing an overview before delving deeper into the procedure in a step-by-step approach. Techniques covered include genomic DNA extraction using cetyl trimethylammonium bromide (CTAB) and chloroform extraction, chromatographic techniques, ELISA, hybridization, gel electrophoresis, dot blot analysis and methods for

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studying polymerase chain reactions. Laboratory protocols and standard operating procedures for key equipment are also discussed, providing an instructive overview for lab work. This practical guide focuses on the latest advances and innovations in methods for molecular biology and biotechnology investigation, helping researchers and practitioners enhance and advance their own methodologies and take their work to the next level. Explores a wide range of advanced methods that can be applied by researchers in molecular biology and biotechnology Features clear, step-by-step instruction for applying the techniques covered Offers an introduction to laboratory protocols and recommendations for best practice when conducting experimental work, including standard operating procedures for key equipment

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Advanced Topics in Forensic DNA Typing: Interpretation builds upon the previous two editions of John Butler's internationally acclaimed Forensic DNA Typing textbook with forensic DNA analysts as its primary audience. Intended as a third-edition companion to the Fundamentals of Forensic DNA Typing volume published in 2010 and Advanced Topics in Forensic DNA Typing: Methodology published in 2012, this book contains 16 chapters with 4 appendices providing up-to-date coverage of essential topics in this important field. Over 80 % of the content of this book is new compared to previous editions. Provides forensic DNA analysts coverage of the crucial topic of DNA mixture interpretation and statistical analysis of DNA evidence Worked mixture examples illustrate the impact of different statistical approaches for reporting results Includes allele frequencies for 24 commonly used autosomal

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