

Methods in ENZYMولوجY

Volume 188
Hydrocarbons and Methylotrophy

edited by
Mary E. Lidstrom

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**Amy Claire Rosenzweig, Stephen W.
Ragsdale**



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The Enzyme Reference Daniel L. Purich, R. Donald Allison, 2003-01-04 The aim of this work is to provide a fuller spectrum of information in a single source on enzyme catalyzed reactions than is currently available in any published reference work or as part of any Internet database The Enzyme Reference A Comprehensive Guidebook to Enzyme Nomenclature Reactions and Methods includes 20 000 review articles and seminal research papers Additionally it provides a novel treatment of so called ATPase and GTPase reactions to account for the noncovalent substratelike and productlike states of molecular motors elongation factors transporters DNA helicases G regulatory proteins and other energases Includes a compendium of over 6 000 enzyme reactions including enzyme commission numbers alternative names substrates products alternative substrates and properties Covers over 900 chemical structures of key metabolites and cofactors Index directs readers to the exact pages for over 9 500 enzyme names

Methods in Methane Metabolism, Part A Amy Rosenzweig, Stephen W. Ragsdale, 2011-05-11 Produced by microbes on a large scale methane is an important alternative fuel as well as a potent greenhouse gas This volume focuses on microbial methane metabolism which is central to the global carbon cycle Both methanotrophy and methanogenesis are covered in detail Topics include isolation and classification of microorganisms metagenomics approaches biochemistry of key metabolic enzymes gene regulation and genetic systems and field measurements The state of the art techniques described here will both guide researchers in specific pursuits and educate the wider scientific community about this exciting and rapidly developing field Topics include isolation and classification of microorganisms metagenomics approaches biochemistry of key metabolic enzymes gene regulation and genetic systems and field measurements The state of the art techniques described here will both guide researchers in specific pursuits and educate the wider scientific community about this exciting and rapidly developing field

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Selected Works in Bioinformatics Xuhua Xia, 2011-10-21 This book consists of nine chapters covering a variety of bioinformatics subjects ranging from database resources for protein allergens unravelling genetic determinants

of complex disorders characterization and prediction of regulatory motifs computational methods for identifying the best classifiers and key disease genes in large scale transcriptomic and proteomic experiments functional characterization of inherently unfolded proteins regions protein interaction networks and flexible protein protein docking The computational algorithms are in general presented in a way that is accessible to advanced undergraduate students graduate students and researchers in molecular biology and genetics The book should also serve as stepping stones for mathematicians biostatisticians and computational scientists to cross their academic boundaries into the dynamic and ever expanding field of bioinformatics

RNA Turnover in Eukaryotes: Nucleases, Pathways and Analysis of mRNA Decay Lynne E. Maquat, Megerditch Kiledjian, 2009-01-30 Specific complexes of protein and RNA carry out many essential biological functions including RNA processing RNA turnover RNA folding as well as the translation of genetic information from mRNA into protein sequences Messenger RNA mRNA decay is now emerging as an important control point and a major contributor to gene expression Continuing identification of the protein factors and cofactors and mRNA instability elements responsible for mRNA decay allow researchers to build a comprehensive picture of the highly orchestrated processes involved in mRNA decay and its regulation Covers the nonsense mediated mRNA decay NMD or mRNA surveillance pathway Expert researchers introduce the most advanced technologies and techniques to identify mRNA processing transport localization and turnover which are central to the process of gene expression Offers step by step lab instructions including necessary equipment and reagents

Chromatin and Chromatin Remodeling Enzymes, Part B Carl Wu, C. David Allis, 2004-03-07 DNA in the nucleus of plant and animal cells is stored in the form of chromatin Chromatin and the Chromatin remodelling enzymes play an important role in gene transcription

RNA Polymerase and Associated Factors, Part C Sankar Adhya, Susan Garges, 2003-12-18 RNA polymerase is molecule important to gene transcription Along with associated factors RNA polymerase is part of the process in which RNA is transcribed to produce a protein Construction and purification of RNA polymerases DNA microarrays and bacterial gene expression Functional analysis of transcription factors

RNA Polymerase and Associated Factors, Part D Sankar Adhya, Susan Garges, 2003-12-18 RNA polymerase is molecule important to gene transcription Along with associated factors RNA polymerase is part of the process in which RNA is transcribed to produce a protein Models and methods for studying polymerase translocation Assay for movements of RNA polymerase along DNA Engineering of elongation complexes of bacterial and yeast RNA polymerases

Quinones and Quinone Enzymes, Part A Helmut Sies, Lester Packer, 2004-05-01 Quinones are members of a class of aromatic compounds with two oxygen atoms bonded to the ring as carbonyl groups This volume covers the role of quinines enzymes in cellular signalling and modulation of gene expression Coenzyme Q Detection and Quinone Reductases Plasma Membrane Quinone Reductases Quinones Cellular Signaling and Modulation of Gene Expression

Liposomes, Part B Nejat Duzgunes, 2003-11-19 Liposomes are cellular structures made up of lipid molecules Important as a cellular model in the study of basic biology liposomes are also

used in clinical applications such as drug delivery and virus studies Liposomes in Biochemistry Liposomes in Molecular Cell Biology Liposomes in Molecular Virology *Macromolecular Crystallography, Part C* Charles W. Carter Jr.,2003-12-15

Accurate molecular structures are vital for rational drug design and for structure based functional studies directed toward the development of effective therapeutic agents and drugs Crystallography can reliably predict structure both in terms of folding and atomic details of bonding Methodological methods in crystals Methodological methods data analysis

Combinatorial Chemistry, Part B ,2004-01-26 Combinatorial Chemistry encompasses both the design of compounds for specific pharmacological use and the screening of molecules in high throughput automated tests to find active agents with specific functions Analytical techniques Direct sorting split and pool combinatorial synthesis Linkers and their applications Microwave assisted synthesis Oligosaccharide chemistry Peptide Synthesis and Screening Polymer assisted approaches Small molecule and heterocycle synthesis Macromolecular Crystallography, Part D Charles W. Carter Jr.,Charles W.

Carter,Robert M. Sweet,2003-12-10 Accurate molecular structures is vital for rational drug design and for structure based functional studies directed toward the development of effective therapeutic agents and drugs Crystallography can reliably predict structure both in terms of folding and atomic details of bonding Phases Map interpretation and refinement Analysis and software **Chromatin and Chromatin Remodeling Enzymes Part C** Carl Wu,C. David Allis,2004-04-07 DNA in the

nucleus of plant and animal cells is stored in the form of chromatin Chromatin and the chromatin remodelling enzymes play an important role in gene transcription Genetic assays of chromatin modification and remodeling Histone modifying enzymes ATP dependent chromatin remodeling enzymes Enzyme Kinetics and Mechanisms, Part E, Energetics of Enzyme Catalysis

,1999-09-06 This volume supplements Volumes 63 64 87 and 249 of Methods in Enzymology These volumes provide a basic source for the quantitative interpretation of enzyme rate data and the analysis of enzyme catalysis Among the major topics covered are Energetic Coupling in Enzymatic Reactions Intermediates and Complexes in Catalysis Detection and Properties of Low Barrier Hydrogen Bonds Transition State Determination and Inhibitors The critically acclaimed laboratory standard for more than forty years Methods in Enzymology is one of the most highly respected publications in the field of biochemistry Since 1955 each volume has been eagerly awaited frequently consulted and praised by researchers and reviewers alike Now with more than 300 volumes all of them still in print the series contains much material still relevant today truly an essential publication for researchers in all fields of life sciences *Handbook of Human Toxicology* Edward J. Massaro,1997-07-09

Covering some of the most important topics in modern toxicology the Handbook of Human Toxicology is a unique and valuable addition to the current literature It addresses issues answers questions and provides data related to Within each of these five major sections are several carefully selected topics that reflect the current state of human to *Sphingolipid Metabolism and Cell Signaling, Part B* ,2000-10-25 This volume contains information on analyzing sphingolipids sphingolipid transport and trafficking and sphingolipid protein interactions and cellular targets Its companion Volume 311 presents

methods used in studying enzymes of sphingolipid biosynthesis and turnover including inhibitors of some of these enzymes genetic approaches and organic and enzymatic syntheses of sphingolipids and analogs The critically acclaimed laboratory standard for more than forty years *Methods in Enzymology* is one of the most highly respected publications in the field of biochemistry Since 1955 each volume has been eagerly awaited frequently consulted and praised by researchers and reviewers alike Now with more than 300 volumes all of them still in print the series contains much material still relevant today truly an essential publication for researchers in all fields of life sciences *Energetics of Biological Macromolecules, Part E* ,2004-04-02 *Energetics of Biological Macromolecules Part E* focuses on methods related to allosteric enzymes and receptors including fluorescent probes spectroscopic methods and quantitative analysis as well as on cooperativity in protein folding NMR and mass spectrometry methods are discussed *Allosteric Enzymes and Receptors Cooperativity in Protein Folding and Assembly Measuring Biological Responses with Automated Microscopy* ,2006-11-02 The critically acclaimed laboratory standard for more than forty years *Methods in Enzymology* is one of the most highly respected publications in the field of biochemistry Since 1955 each volume has been eagerly awaited frequently consulted and praised by researchers and reviewers alike Now with more than 300 volumes all of them still in print the series contains much material still relevant today truly an essential publication for researchers in all fields of life sciences *Adult Stem Cells* ,2006-12-11 This is the second of three planned volumes in the *Methods in Enzymology* series on the topic of stem cells This volume is a unique anthology of stem cell techniques focusing on adult stem cells and written by experts from the top laboratories in the world The contributors not only have hands on experience in the field but often have developed the original approaches that they share with great attention to detail The chapters provide a brief review of each field followed by a cookbook and handy illustrations The collection of protocols includes the isolation and maintenance of stem cells from various species using conventional and novel methods such as derivation of ES cells from single blastomeres differentiation of stem cells into specific tissue types isolation and maintenance of somatic stem cells stem cell specific techniques and approaches to tissue engineering using stem cell derivatives The reader will find that some of the topics are covered by more than one group of authors and complement each other Comprehensive step by step protocols and informative illustrations can be easily followed by even the least experienced researchers in the field and allow the setup and troubleshooting of these state of the art technologies in other laboratories Provides complete coverage spanning from derivation isolation of stem cells and including differentiation protocols characterization and maintenance of derivatives and tissue engineering Presents the latest most innovative technologies Addresses therapeutic relevance including FDA compliance and tissue engineering

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