

Methods in ENZYMولوجY

Volume 188
Hydrocarbons and Methylotrophy

edited by
Mary E. Lidstrom

Methods In Enzymology Volume 188

H Kauffman



Methods In Enzymology Volume 188:

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

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

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DNA Microarrays, Part A: Array Platforms and Wet-Bench Protocols, 2011-08-19 Modern DNA microarray technologies have evolved over the past 25 years to the point where it is now possible to take many million measurements from a single experiment These two volumes Parts A B in the Methods in Enzymology series provide methods that will shepherd any molecular biologist through the process of planning performing and publishing microarray results Part A starts with an overview of a number of microarray platforms both commercial and academically produced and includes wet bench protocols for performing traditional expression analysis and derivative techniques such as detection of transcription factor occupancy and chromatin status Wet bench protocols and troubleshooting techniques continue into Part B These techniques are well rooted in traditional molecular biology and while they require traditional care a researcher that can reproducibly generate beautiful Northern or Southern blots should have no difficulty generating beautiful array hybridizations Data management is a more recent problem for most biologists The bulk of Part B provides a range of techniques for data handling This includes critical issues from normalization within and between arrays to uploading your results to the public repositories for array data and how to integrate data from multiple sources There are chapters in Part B for both the debutant and the expert

bioinformatician Provides an overview of platforms Includes experimental design and wet bench protocols Presents statistical and data analysis methods array databases data visualization and meta analysis

Ubiquitin and Protein Degradation Raymond Joseph Deshaies, 2005 Ubiquitin and Protein Degradation Part B will cover chemical biology ubiquitin derivatives

and ubiquitin like proteins deubiquitinating enzymes proteomics as well as techniques to monitor protein degradation The chapters are highly methodological and focus on application of techniques Second part of the Ubiquitin and Protein Degradation series Topics include E1 Enzymes E2 Enzymes E3 Enzymes Proteasomes and Isopeptidases **Applications of Chimeric Genes and Hybrid Proteins, Part C: Protein-Protein Interactions and Genomics** ,2000-10-28 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 *Regulators of G Protein Signaling, Part B* ,2004-11-16 Regulators of G Protein Signaling Part B continues the in depth treatment of the topic and covers the RSG protein superfamily including RZ R4 R7 R12 RhoGEF and GRK as well as other heterotrimeric G protein signaling regulators Table of Contents RZ Subfamily R4 Subfamily R7 Subfamily R12 Subfamily RhoGEF Subfamily GRK Subfamily Other RGS proteins Activators Inhibitors Other Modulators **Advanced Bacterial Genetics: Use of Transposons and Phage for Genomic Engineering** ,2007-02-27 The critically acclaimed laboratory standard for more than fifty 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 over 400 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 This new volume presents methods related to the use of bacterial genetics for genomic engineering The book includes sections on strain collections and genetic nomenclature transposons and phage

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