

# Protein Multifunctionality

## Molecular multifunctionality: Aro1

### Molecular function

3-Dehydroquinate dehydratase  
3-Dehydroquinate synthase  
3-Phosphoshikimate 1-cv-transferase  
Shikimate 3-dehydrogenase  
Shikimate kinase

### Biological function

Chorismate biosynthesis

## Biological multifunctionality: Sod1

### Molecular function

Superoxide dismutase

### Biological function

Superoxide metabolic process  
Regulation of cellular respiration  
Ion homeostasis  
Protein stabilization

## Molecular & biological multifunctionality: Aco1

### Molecular function

Aconitase hydratase  
Double-stranded DNA binding

### Biological function

Tricarboxylic-acid cycle  
Mitochondrial genome maintenance

# Multifunctional Proteins

**D. Grahame Hardie, John R. Coggins**

## **Multifunctional Proteins:**

**Multifunctional Proteins** James F. Kane, 2019-06-04 First Published in 1983 this book offers a full comprehensive guide into proteins and their functions Carefully compiled and filled with a vast repertoire of notes diagrams and references this book serves as a useful reference for Students of Alternative Medicine and other practitioners in their respective fields

**Multifunctional Proteins** Kasper Kirschner, Hans Bisswanger, 1976      **Multifunctional Proteins** Hans Bisswanger, Eva Schmincke-Ott, 1980      **Multifunctional Proteins** Hans Bisswanger, Eva Schmincke-Ott, 1980      *Cryptic Enzymes and Moonlighting Proteins* Helen Irving, Chris Gehring, Aloysius Wong, 2025-04-24 Cryptic Enzymes and Moonlighting Proteins a new volume in the Foundations and Frontiers in Enzymology series offers a thorough overview of cryptic enzymes and moonlighting proteins in signaling cascades In early chapters leading international contributors discuss evolutionary considerations for moonlighting proteins moonlighting interactions in the extracellular matrix eukaryotic moonlighting proteins modulating moonlighting kinases moonlighting proteins in neurobiology signaling metabolic enzymes moonlighting as RNA binding and regulatory proteins Later methods driven chapters discuss practical aspects of identifying hidden moonlighting domains in proteins computational approaches and bioinformatic tools for the identification of cryptic enzymes establishing cryptic enzyme interactomes and assessing contributions of moonlighting proteins to signal cascades The book also explores recent advances in research and brings together an array of information across different fields to enable better targeting of these exciting proteins and their interactomes With a clear focus on the role of moonlighting and cryptic enzymes in signal transduction users will find examples of cryptic enzymes across species as well as those in human healthy biology and pathogenesis Covers recent advances in our understanding of cryptic moonlighting proteins in signal cascades highlighting and examining key themes across disciplines Empowers researchers to better target cryptic enzymes and moonlighting proteins and their interactomes Features chapter contributions from international leaders in the field

**Protein Structure and Function** Gregory A. Petsko, Dagmar Ringe, 2004 Each title in the Primers in Biology series is constructed on a modular principle that is intended to make them easy to teach from to learn from and to use for reference

**Informational Biopolymers of Genes and Gene Expression** R.D. Blake, 2005-03-23 This new text examines the biophysics and biochemistry of nucleic acids and proteins carving out the dynamic interface between chemistry and molecular biology and providing a detailed picture of nucleic acids and proteins their structures biological properties and origins and evolution Full of novel insights informed by years of research and teaching R.D. Blake has written a new text that examines the biophysics and biochemistry of nucleic acids and proteins This book carves out the dynamic interface between chemistry and molecular biology and provides a detailed picture of nucleic acids and proteins their structures biological properties and origins and evolution While pitched at a level that is accessible to upper level undergraduate students of biology the book will likewise be of keen interest to researchers and graduate students in biology chemistry and the physical

sciences A broad range of topics accompanied with extensive references footnotes boxed asides chapter ending problems and detailed illustrations throughout combine to make this an ideal choice for courses on advanced biochemistry physical biochemistry nucleic acids and proteins biophysics and biochemistry molecular evolution and protein nucleic acid interactions *Multidomain Proteins* D. Grahame Hardie, John R. Coggins, 1986 **Microbial Biochemistry** G.N. Cohen, 2013-03-19

Microbial physiology biochemistry and genetics allowed the formulation of concepts that turned out to be important in the study of higher organisms. In the first section the principles of bacterial growth are given as well as the description of the different layers that enclose the bacterial cytoplasm and their role in obtaining nutrients from the outside media through different permeability mechanism described in detail. A chapter is devoted to allosteric and is indispensable for the comprehension of many regulatory mechanisms described throughout the book. Another section analyses the mechanisms by which cells obtain the energy necessary for their growth glycolysis the pentose phosphate pathway the tricarboxylic and the anaplerotic cycles. Two chapters are devoted to classes of microorganisms rarely dealt with in textbooks namely the Archaea mainly the methanogenic bacteria and the methylotrophs. Eight chapters describe the principles of the regulations at the transcriptional level with the necessary knowledge of the machineries of transcription and translation. The next fifteen chapters deal with the biosynthesis of the cell building blocks amino acids purine and pyrimidine nucleotides and deoxynucleotides water soluble vitamins and coenzymes isoprene and tetrapyrrole derivatives and vitamin B12. The two last chapters are devoted to the study of protein-DNA interactions and to the evolution of biosynthetic pathways. The considerable advances made in the last thirty years in the field by the introduction of gene cloning and sequencing and by the exponential development of physical methods such as X-ray crystallography or nuclear magnetic resonance have helped presenting metabolism under a multidisciplinary attractive angle. The level of readership presupposes some knowledge of chemistry and genetics at the undergraduate level. The target group is graduate students researchers in Academia and industry.

*Structural and Functional Characterization of Staphylococcus aureus Enolase* Vijay Hemmadi, 2023-01-09 Doctoral Thesis Dissertation from the year 2022 in the subject Biology Micro and Molecular Biology grade 9.5 course PhD language English abstract The presence of metabolic enzymes like enolase on the surface of *S. aureus* has sparked considerable scientific debate. Enolase rSaeno is found in *S. aureus* as a fragile octamer and a robust dimer. The catalytic octamer resides in the cytoplasm and engages in energy metabolism metal interchangeability and averting the host's massive metal exploitation. In contrast, the membrane-bound dimer possesses a high affinity for host plasminogen and is involved in binding and immobilizing the host plasminogen on the *S. aureus*. The interaction of host plasminogen and rSaeno on the bacterial surface mediates the adherence of *S. aureus* to the host and subsequently arbitrates plasminogen activator-mediated activation of plasminogen. The binding of dimeric rSaeno to the kringle domain of plasminogen prevents 2 antiplasmin-mediated inhibition of plasmin. Enolase is a moonlighting protein that serves two unrelated functions in two separate cell compartments using

two oligomeric conformations Moonlighting proteins for pathogenicity saves a great deal of energy for the pathogen and allows the bacterial invasion system to remain incognito to the host The pathogen develops an endless supply of host protease and disguises the host immune system by targeting copious host plasminogen Although enolase is conserved across species prominent structural variations can be utilized as potential competitive drug targets As *S aureus* employs numerous multifunctional proteins for pathological invasion targeting enolase and other moonlighting proteins engaged in various steps of pathological invasion may serve as a potential means by which the pathological invasion of antibiotic resistant *S aureus* can be prevented

**Multifunctional Proteins** George R. Stark, Neil B. Madsen, 1977 Proteins in Biology and Medicine

Ralph A. Bradshaw, 1982 Proteins in Biology and Medicine contains the proceedings of the 1981 U S China Conference on Proteins in Biology and Medicine held in Shanghai China The papers explore the structure function relationships of proteins including their regulatory properties Topics range from the regulation of biological processes to the structure function relationships of enzymes and blood proteins along with protein protein interactions Organized into four sections encompassing 23 chapters this book begins with an overview of structure function relationships in phospholipase A2 including the enzyme for *Gene Expression, Translation and the Behavior of Proteins* Lester Goldstein, 1980

Multidomain Proteins László Patthy, Peter Friedrich, 1986 Biosyntheses Georges N. Cohen, 1998-08-31 This book emphasizes the importance of microbial physiology biochemistry genetics and evolution in the study of biosynthesis It takes into account the considerable advances made in the field in the last twenty years by the introduction of gene cloning and sequencing and by the exponential development of physical methods such as X ray crystallography of proteins

**Annual Review of Biochemistry** James Murray Luck, 1987 Annual Review of Biochemistry E. E. Snell, 1976 The Product of the His4 Gene Cluster in *Saccharomyces Cerevisiae* Joseph Kauffman Keesey, 1978

**Biocatalyst Design for Stability and Specificity** Michael E. Himmel, 1993 For graduate students and professionals in biochemistry and related fields these 25 papers highlight the diversity of approaches used in understanding the function of proteins and how this understanding is employed in improving protein design for biotechnological applications Papers are grouped into five sections on protein stability protein folding multifunctional proteins design of cellulases by recombinant methods and improving natural enzymes by chemical cross linking Annotation copyright by Book News Inc Portland OR

**Metabolism of Aromatic Amino Acids and Amines** Seymour Kaufman, 1987-05-28 The critically acclaimed laboratory standard 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 The series contains much material still relevant today truly an essential publication for researchers in all fields of life sciences

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