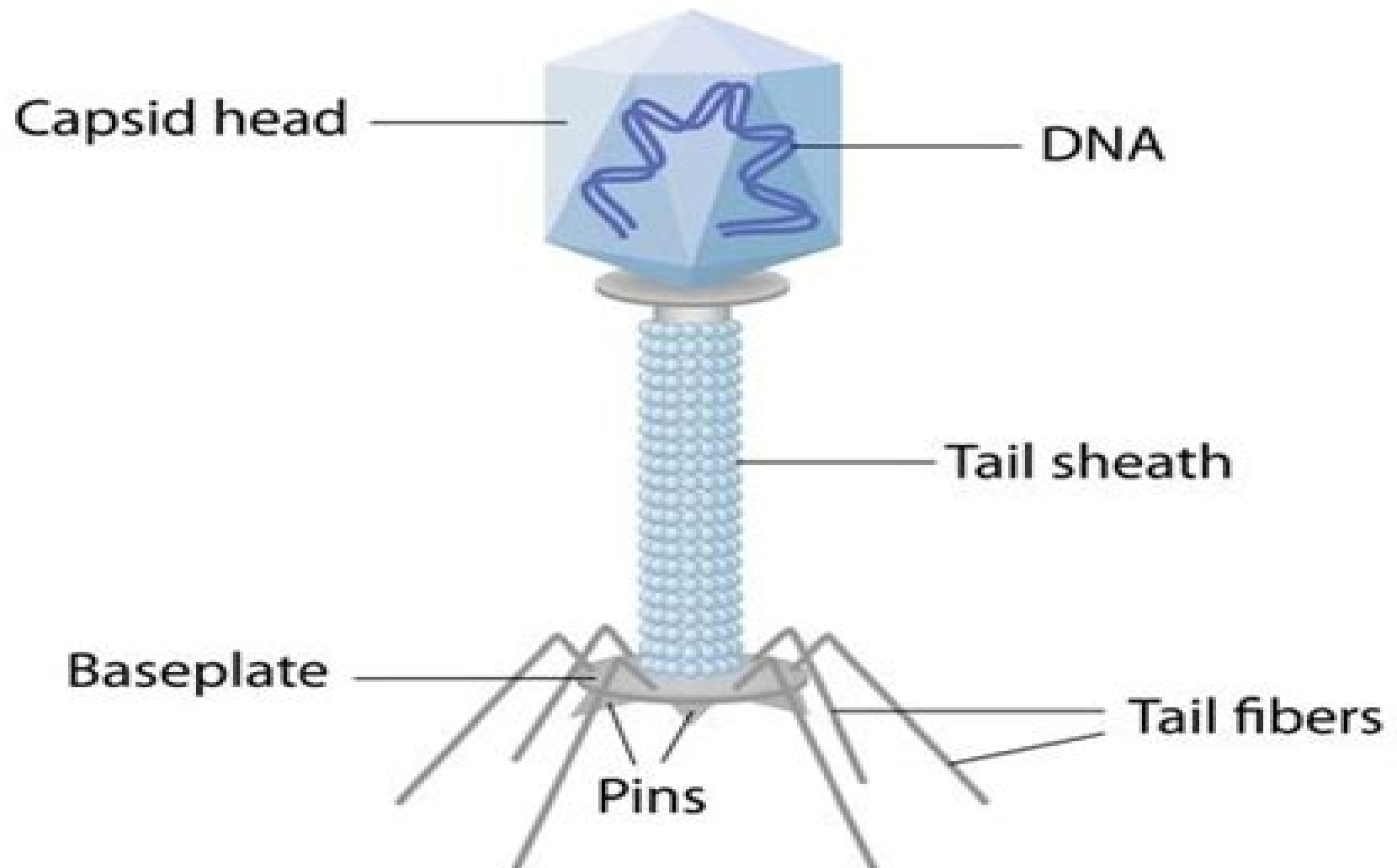


Structure of a Bacteriophage



Bacteriophage Biochemistry

Stephen T. Abedon



Bacteriophage Biochemistry:

Bacteriophage Biochemistry Christopher K. Mathews, 1971 *Bacteriophage Ecology* Stephen T. Abedon, 2008-05-01

Bacteriophages or phages are viruses that infect bacteria and are believed to be the most abundant and genetically diverse organisms on Earth. As such, their ecology is vast both in quantitative and qualitative terms. Their abundance makes an understanding of phage ecology increasingly relevant to bacterial ecosystem ecology, bacterial genomics, and bacterial pathology. Abedon provides the first text on phage ecology for almost 20 years. Written by leading experts synthesizing the three key approaches to studying phage ecology—namely studying them in natural environments *in situ*, experimentally in the lab, or theoretically using mathematical or computer models—With strong emphasis on microbial population biology and distilling cutting edge research into basic principles, this book will complement other currently available volumes. It will therefore serve as an essential resource for graduate students and researchers, particularly those with an interest in phage ecology and evolutionary biology. Comprehensive Virology H. Fraenkel-Conrat, 2012-12-06 The time seems ripe for a critical compendium of that segment of the biological universe we call viruses. Virology as a science having passed only recently through its descriptive phase of naming and numbering has probably reached that stage at which relatively few new truly new viruses will be discovered. Triggered by the intellectual probes and techniques of molecular biology, genetics, biochemical cytology, and high resolution microscopy and spectroscopy, the field has experienced a genuine information explosion. Few serious attempts have been made to chronicle these events. This comprehensive series, which will comprise some 6000 pages in a total of about 22 volumes, represents a commitment by a large group of active investigators to analyze, digest, and expostulate on the great mass of data relating to viruses, much of which is now amorphous and disjointed and scattered throughout a wide literature. In this way, we hope to place the entire field in perspective and to develop an invaluable reference and sourcebook for researchers and students at all levels. This series is designed as a continuum that can be entered anywhere but which also provides a logical progression of developing facts and integrated concepts.

Bacteriophages John Douglas, 2013-11-11 The contribution of bacteriophages to the development of modern biology cannot be overestimated; yet sixty years after their discovery, they are as remote and mysterious to many scientists as they are to most laymen. This book endeavours to remedy the situation: an attempt has been made to provide in readily comprehensible form a nucleus of information essential to anyone embarking on the study of bacteriophages or using them in their work for the first time. It shows the range of bacteriophage structure and behaviour; it illustrates the role of bacteriophage in molecular biology; it surveys the current state of the art; it presents the medical and industrial aspects. Some simple experimental procedures are given in sufficient detail for the beginner to attempt them successfully. Other more sophisticated procedures are presented so as to impart a feeling of intimate reality without dazzling the reader with technical complexity. I hope that young readers will forgive me for assuming that they have some knowledge of bacteria.

nucleic acids antibodies and isotopes Likewise I would ask mature workers to excuse the omission of cherished specialties To have included all these valuable though they are might have put this book beyond the reach of the phage novices for whom it is intended Specific references save a few of particular interest have been omitted Other books on bacteriophage v

vi BACTERIOPHAGES provide them in abundance The Bacteriophages Richard Calendar, 2006 This authoritative timely and comprehensively referenced compendium on the bacteriophages explores current views of how viruses infect bacteria In combination with classical phage molecular genetics new structural genomic and single molecule technologies have rendered an explosion in our knowledge of phages Bacteriophages the most abundant and genetically diverse type of organism in the biosphere were discovered at the beginning of the 20th century and enjoyed decades of use as anti bacterial agents before being eclipsed by the antibiotic era Since 1988 phages have come back into the spotlight as major factors in pathogenesis bacterial evolution and ecology This book reveals their compelling elegance of function and their almost inconceivable diversity Much of the founding work in molecular biology and structural biology was done on bacteriophages These are widely used in molecular biology research and in biotechnology as probes and markers and in the popular method of assessing gene expression

Ultrastructure of Bacterial Viruses Anna S. Tikhonenko, 2012-12-06 After the discovery of the tobacco mosaic virus by D I Ivanovskii in 1892 the new science of virology was born and began to develop rapidly The number of viruses now known is enormous and they can infect nearly all animal and plant organisms Microorganisms themselves are no exception to this rule Despite intensive study of viruses their origin and nature are still a subject for speculation and hypothesis The general concept of viruses embraces a wide group of biologically active structures occupying an intermediate position between living and nonliving matter The dual character of viruses is determined by the fact that while they do not possess an independent system of metabolism which is a characteristic feature of every living being they nevertheless carry within themselves all the necessary information for autoreproduction A striking feature of the virus is that it consists essentially of two components a protein envelope and the nucleic acid contained within it In contrast to the elementary structural unit of the living organism the cell which contains two types of nucleic acid DNA and RNA the virus particle contains only one type of nucleic acid either DNA or RNA It is perhaps this which is responsible for the imperfection of the virus as a living organism

Bacteriophages David R. Harper, Stephen T. Abedon, Benjamin H. Burrowes, Malcolm L. McConville, 2021-01-30 This first major reference work dedicated to the manifold industrial and medical applications of bacteriophages provides both theoretical and practical insights into the emerging field of bacteriophage biotechnology The book introduces to bacteriophage biology ecology and history and reviews the latest technologies and tools in bacteriophage detection strain optimization and nanotechnology Usage of bacteriophages in food safety agriculture and different therapeutic areas is discussed in detail This book serves as essential guide for researchers in applied microbiology biotechnology and medicine coming from both academia and industry

Biochemistry and Cell Biology, 2003 The

Enzymes of Biological Membranes Anthony Martonosi, 2012-12-06 A 1 Enzymes of Membrane Phospholipid Metabolism in Animals I Introduction II Type 1 Reactions A Acylation of Glycerol 3 phosphate B Esterification of Saturated Fatty Acids to Phospholipids G Hydrolysis of the 1 Acyl Ester in Phospholipids D Other Lysophospholipase Activities III Type 2 Reactions A Formation of the 2 Acyl Ester of Phosphatidic Acid B Esterification of Unsaturated Fatty Acids to Phospholipids C Hydrolysis of the 2 Acyl Ester IV Type 3 Reactions A Diacylglycerol Kinase B Choline and Ethanolamine Phosphotransferase G Hydrolysis of Phosphatidylcholine

Progress in Molecular and Subcellular Biology, 2012-12-06 **Cumulated Index Medicus**, 1994 **Origin and Evolution of Viruses** Esteban Domingo, Colin R. Parrish, John J. Holland, 2008-06-23 New viral diseases are emerging continuously Viruses adapt to new environments at astounding rates Genetic variability of viruses jeopardizes vaccine efficacy For many viruses mutants resistant to antiviral agents or host immune responses arise readily for example with HIV and influenza These variations are all of utmost importance for human and animal health as they have prevented us from controlling these epidemic pathogens This book focuses on the mechanisms that viruses use to evolve survive and cause disease in their hosts Covering human animal plant and bacterial viruses it provides both the basic foundations for the evolutionary dynamics of viruses and specific examples of emerging diseases NEW methods to establish relationships among viruses and the mechanisms that affect virus evolution UNIQUE combines theoretical concepts in evolution with detailed analyses of the evolution of important virus groups SPECIFIC Bacterial plant animal and human viruses are compared regarding their interaction with their hosts **Research Awards Index**, 1979 Handbook of Genetics Robert C. King, 2013-11-11 The purpose of this and future volumes of the Handbook of Genetics is to bring together a collection of relatively short authoritative essays or annotated compilations of data on topics of significance to geneticists Many of the essays will deal with various aspects of the biology of certain species selected because they are favorite subjects for genetic investigation in nature or the laboratory Often there will be an encyclopedic amount of information available on such a species with new papers appearing daily Most of these will be written for specialists in a jargon that is bewildering to a novice and sometimes even to a veteran geneticist working with evolutionarily distant organisms For such readers what is needed is a written introduction to the morphology life cycle reproductive behavior and culture methods for the species in question What are its particular advantages and disadvantages for genetic study and what have we learned from it Where are the classic papers the key bibliographies and how does one get stocks of wild type or mutant strains The chapters devoted to different species will contain information of this sort Only a few hundreds of the millions of species available to biologists have been subjected to detailed genetic study However those that have make up a very heterogeneous sample of the living world **Research Grants Index** National Institutes of Health (U.S.). Division of Research Grants, 1972 *The Genetic Mechanism and the Origin of Life* Lawrence Dillan, 2012-12-06 As shown in the text there can be little doubt that the genetic mechanism is for all practical purposes equivalent to life itself Consequently it is unrealistic to seek knowledge of the

origin of life and its subsequent evolution without simultaneously searching for an understanding of how this apparatus arose and evolved. Fortunately, the annual publication over the recent years of thousands of papers dealing with the genetic processes has brought the state of knowledge to a level where a synthesis of their major details in relation to life's history is feasible. Because of the voluminous body of literature, no single book can possibly treat all the ramifications of this fundamental subject. Subdivision into multiple volumes is necessary. This volume, the first of a trilogy, explores the molecular aspects of the problem in connection with the precellular aspects up to the point of the origin of the cell. The second, currently in progress, is concerned with the subsequent evolution of the cell as revealed by the energy-related organelles and their genetic apparatuses and by ultrastructural details of other cellular parts. The third volume, as presently planned, deals with developmental, immunological, and other complexities at the organismic level and in so doing throws additional light on basic properties of the genetic processes themselves. Thus, the genetic apparatus provides the warp and evolution the woof of the intricate fabric that emerges.

Chemical Recognition in Biology F. Chapeville, A. L. Haenni, 2012-12-06. Studies of chemical recognition in biology were initiated about half a century ago with the first kinetic data obtained on enzyme catalysis and inhibition. They led to a rather static representation of the recognition process illustrated by the lock and key model that still continues to influence our overall image of recognition and its specificity. In several cases, crystallographic studies of enzyme-substrate complexes have supported this model. Indeed, in a crystallized ligand-enzyme complex, a close fitting is observed between the active center of the enzyme and the functional groups of the ligand. However, this does not necessarily result from a direct recognition process between rigid structures but may result from a progressive adaptation during which the initial structures of the enzyme and the ligand are modified. Induced fit mechanism. Recently, a great deal of work has been devoted to the study of recognition in more complex systems such as the replication or the translation machinery. Clearly, the extraordinary precision of such systems cannot be explained solely in terms of physical matching between enzymes and their substrates. This has led to a noticeable change of perspective in these areas. As a result of the new kinetic viewpoint, one rather focuses on the time course of the processes, on the kinetic balance between steps of the reaction, on the energy-accuracy relationships, and on the strategies which permit the achievement of high precision using relatively error-prone components in an appropriate dynamic interplay.

NMR of Proteins Clore, Marius Clore, A. M. Gronenborn, 1993. Determination of structures of larger proteins in solution by three and four dimensional heteronuclear magnetic resonance spectroscopy. Methodological advances in protein NMR. Determination of high resolution NMR structures of proteins. Multidimensional NMR studies of immunosuppressant immunophilin complexes. NMR studies of the structure and role of modules involved in protein-protein interactions. NMR structural studies of membrane proteins. Heteronuclear NMR studies of the molecular dynamics of staphylococcal nuclease. Study of protein dynamics by NMR. The folding stability and dynamics of T4 lysozyme: a perspective using nuclear magnetic resonance.

Bioremediation of Chlorinated and Polycyclic Aromatic

Hydrocarbon Compounds Battelle Memorial In, 1994-02-28 This timely reference presents the state of the art of the emerging and rapidly changing field of bioremediation of chlorinated solvents PCBs and other chlorinated compounds as well as PAHs both in situ and on site This landmark publication reports significant advances in bioremediation with an emphasis on practical applications and state of the art developments Laboratory and field oriented reviews are presented with the objective of tying treatability studies and recent laboratory developments to field applications No other reference source gives you access to the most current techniques and methods for the bioremediation of chlorinated and polycyclic aromatic hydrocarbon compounds This book represents the work of leading experts in the fields of in situ and on site bioremediation from North America Europe and Asia The chapters include current field applications and laboratory studies undertaken in some cases in countries with regulatory standards more stringent than those of the United States

Circular Dichroism and the Conformational Analysis of Biomolecules G.D. Fasman, 2013-11-11 Excellent and very timely It will undoubtedly become a standard reference for the application of circular dichroism CD to biomolecules Quarterly Review of Biology March 1997 T estament to the book s utility is the fact that during the course of my review I had to rescue it from the desks of graduate students on an almost daily basis In summary this is a great book American Scientist Well documented chapters provide a very good insight into the problems surrounding the conformation of biomacromolecules An indispensable source of information Nahrung 42 2 1998 Renowned experts present the first state of the art description of circular dichroism spectroscopy CD Chapters present in depth discussions of the history of the field the theory of CD for application to globular proteins membrane proteins peptides nucleic acids and their interactions carbohydrates and instrumentation Discussions also feature new techniques using synchrotron radiation vibrational Raman optical activity and vibrational CD More than 250 illustrations supplement the text

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