

Sa3	ASVKAVGRVCYSALPSQAHDTLDLIDEGGPFPY	33
Sa	-DVS--GTVCLSALPPEATDTLNLIASDGPFPY	30
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Sa3	SQDGVVVFQNREGLLPAHSTGYYYHEYTVITPGSP	66
Sa	SQDGVVVFQNRESVLPTQSYGYYYHEYTVITPGAR	63
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Sa3	TRGARRIITGQQWQEDYYTADHYASFRRVDFAC	99
Sa	TRGTRRIITGEATQEDYYTGDHYATFSLIDQTC	96
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Microbial Ribonucleases With 5 Figures

E. Heinz



Microbial Ribonucleases With 5 Figures:

Microbial Ribonucleases Fujio Egami, K. Nakamura, 2012-12-06 **The Bacteriocins** Peter Reeves, 2012-12-06

In recent years bacteriocins especially colicins have become widely known to molecular biologists as proteins with peculiar ways of killing bacteria. These same bacteriocins have been known for a long time to bacteriology for their unusual activity spectra and enormous variety. In this monograph I have attempted to bring together our detailed knowledge of those few bacteriocins which have already received attention from molecular biologists and our less detailed but extensive knowledge of the variety of bacteriocins which exist. The field has been reviewed in whole or in part by several authors: FREDERICQ 1957, 1964, 1965; IVANOVICS 1962; HAMON 1965; REEVES 1965. These reviews have been very useful to the author and readers will find further references in them and sometimes alternative viewpoints. We have already referred to bacteriocins as proteins and in doing so have excluded many more complex antibacterial agents which resemble bacteriophages or their tails. In the author's view these phage-like particles are probably not bacteriocins but many authors include them within the definition. The more restrictive definition used here has meant omitting discussion of some excellent studies on what the present author would term defective bacteriophages. In the first chapter we look at the discovery of bacteriocins and an outline of their classification. With this background we can discuss in Chapters 2 to 6 the chemistry, genetics and mode of action of the more intensively studied bacteriocins.

Protein Sequence Determination Saul B. Needleman, 2013-06-29

All the King's horses and all the King's men couldn't put Humpty Dumpty together again. It is entirely possible that the difficulty facing all the King's men was principally the lack of a sufficient guide to the techniques of reassembling from a series of small components the original entity. It is the sincere hope of the editor of the present work and of each of the contributing authors that the modern researcher will not face a similar predicament in his endeavours to reconstruct the complete primary sequence of a protein from the array of component amino acids. Rather it is the intent that with this volume he may proceed untimorously if not with outright confidence toward achieving his purpose. To the newcomer in protein sequencing, compelled by necessity or fascination to investigate the exact order of amino acids in proteins, the question of where to begin or how to do it is urgent. To those more skilled, a ready source of additional techniques should nevertheless be of value. This volume attempts to present in a single source a discussion of the methods and techniques useful to the determination of the primary structures of proteins and peptides. Hopefully this book will tell the reader how to do it.

The Ribosome Aleksandr S. Spirin, Lidija P. Gavrilova, 2012-12-06

This monograph is neither a historical outline of the development of the concepts of protein biosynthesis and the structure and functions of the ribosomes nor an exhaustive survey of the literature on these questions. The monograph is based upon an analysis of the modern trends in this field. The purpose of the monograph was to formulate more or less generalized representations of the structure and function of the ribosome as we envision it at the present day. It may be that this attempt is premature for a number of reasons and the concepts outlined here will very soon

be revised Nonetheless despite this risk we believe it to be advisable to undertake this attempt for the following reasons firstly the undertaken analysis could aid in the comprehension of the substantial mass of extremely scattered experimental data on the ribosomes presently available secondly in any event even if most of the concepts outlined rapidly become obsolete they can still serve as a stimulus for a whole series of experiments and thirdly we hope that some of the concepts outlined will still remain essentially correct and relatively stable In view of the aforementioned we should make the following reservations First of all we made no attempt to cite all the literature on the problems discussed but considered it sufficient to illustrate the various premises with one or several sample references Protamines Toshio Ando,M. Yamasaki,K.

Suzuki,2012-12-06 A century has already passed since FRIEDRICH MIESCHER working at Strasbourg and Basel began his study of protamine one of the basic nuclear proteins of cells It was first established by KOSSEL that protamine represents the simplest known protein In the conviction that research into the nature of protamine would shed light on that of other typical proteins a group of researchers in Germany followed MIESCHER and laid the foundations of protein chemistry A general view of prot amines was thus built up by KOSSEL working at Strasbourg Berlin Marburg an der Lahn and Heidelberg FELIX at Heidelberg Munich and Frankfurt am Main and WALDSCHMIDT LEITZ at Prague and Munich Concepts and techniques established by these studies have been widely utilized for research on other typical proteins The revolutionary advances in chemical and physical techniques after W orld War II extended the sphere of research to Tokyo in the Far East Prof FELIX visit in 1955 greatly encouraged our research group in Tokyo His death in August 1960 constituted a sad loss to protein chemistry and stimulated our group to assume responsibility for carrying on the studies In the following decade we in Tokyo have been able to add a new development to the results on the chemical structure of protamines accumulated by the Eurqpean researchers over a period of about fifty years *Bacteriocins, Microcins and Lantibiotics* Richard James,Claude Lazdunski,Franc Pattus,2013-06-29 Presented here are recent findings on bacteriocins plasmid encoded toxins produced by bacteria which differ from traditional antibiotics in killing only bacteria that are closely related to the producing strain Included are introductory chapters on bacteriocins microcins low molecular weight bacteriocins and lantibiotics peptide antibiotics containing lanthionine further contributions on pore forming bacteriocins the mechanisms of immunity to bacteriocins uptake and secretion as well as evolution of bacteriocins It is of particular interest that the lantibiotic nisin is approved for use as food preservative and another lantibiotic epidermin has potential as a therapeutic drug against acne

Micromethods in Molecular Biology Volker Neuhoff,2012-12-06 This book is based on practical experience and is therefore written as a prac tical manual The fore runners of the book were the manuals of the first and second EMBO Courses on Micromethods in Molecular Biology which were held in G6ttingen in the spring of 1970 and the autumn of 1971 This book may serve as a manual not only for the participants of the third EMBO Course to be held in G6ttingen in autumn 1973 but also for all experimenters who are interested in using micromethods It must be emphasized from the outset that

this book is conceived as a cook book and not as a monograph which attempts to review the literature on micromethods critically. The methods described here in detail are performed routinely in the authors' laboratories and include all the practical details necessary for the successful application of the micromethods. There are many other sensitive and excellent micro methods which are not included in this book because the authors feel that in a cook book only methods for which they have personal experience and proficiency should be described. Some readers may feel that the title promises more than the present contents of this book; however, if sufficient interest is shown in this volume, it may be possible to remedy such deficiencies in future editions.

Quantitative Aspects of Allosteric Mechanisms A. Levitzki, 2013-11-11. The aim of this monograph is to summarize the essential features which characterize the behavior of regulatory systems. Firstly, we discuss the laws which govern ligand binding in thermodynamic terms. The basic cooperative and allosteric phenomena are described in thermodynamic terms without assuming any particular model. Then the molecular models developed by Monod, Wyman, and Changeux and by Koshland, Nemethy, and Filmer are presented in detail. Special emphasis has been given to the analysis of the Hill coefficient and its meaning both in thermodynamic terms and in terms of the two allosteric models: the concerted model of Monod, Wyman, and Changeux and the sequential model of Koshland, Nemethy, and Filmer. Special types of cooperativities are discussed in some detail, namely cooperativity stemming from ligand-coupled protein association or dissociation, negative cooperativity, and half of the sites reactivity. A slightly extended space was devoted to the discussion of negative cooperativity and half of the sites reactivity since the existence of these phenomena and their possible biological importance is less of a common knowledge than positive cooperativity. This monograph does not attempt to be a review of specific examples analyzed according to one model or another. Rather, an attempt is made to provide the reader with the quantitative tools to analyze any specific regulatory system. Last but not least, I would like to thank Prof. F. W. Dahlquist from the Institute of Molecular Biology, the University of Oregon, Eugene, and Prof. D. E. Koshland, Jr.

Advanced Methods in Protein Sequence Determination Saul B. Needleman, 2012-12-06. Confusion now hath made his masterpiece. Macbeth II.iii.72. Whence and what are those execrable shapes? Paradise Lost. Ib. 1.681. Confusion worse confounded. Paradise Lost. Ib. 1.995. When the manuscript for the first part of this book was proposed, it was anticipated that the discussion of the entire field of protein sequencing could be covered in a single volume, from purification and characterization of the protein through fragmentation by chemical or enzymic means and finally to reassembly of the identified individual peptides into the reconstructed total sequence. It soon became evident that this would not be possible. While the intent was to restrict the exposure of procedures only to that information concerned with hands-on wet chemistry, it became apparent that a thorough presentation would require, in addition, a discussion of certain instrumental and more theoretical approaches not included in the first volume. Furthermore, the entire understanding of the field of protein sequencing has advanced appreciably since the inception of this book. The purpose of the first volume was to provide practical information in sufficient detail to permit the

researcher to undertake the actual sequencing procedures in his own laboratory *Lampbrush Chromosomes* Harold G. Callan, 2012-12-06 In 1965 I was asked by Dr Konrad Springer whether I would consider writing a monograph on Lampbrush chromosomes and their physiological meaning and although I accepted in principle I refused to write there and then or to meet a deadline I wanted to continue with my own research and I had other responsibilities that left me with little free time but a much more important consideration was that in the 1960s the subject was beset by a host of unresolved questions I felt that to write a review then would be premature full of speculations many of which would be refuted and indeed were refuted within the next decade or two Had I written at that time the only real advantage over the present would have been that few biologists were studying lampbrush chromosomes and the published literature was therefore scanty I am glad that I insisted on delay and am grateful for Springer Verlag's patient acceptance of my decision The first chapter of this monograph describes the history of research on lampbrush chromosomes from their discovery towards the end of the 19th century until the early 1960s By then several facts concerning their structure and chemistry had been firmly established including the evidence that a lampbrush chromatid is uninear it contains a single uninterrupted DNA duplex This exposed a major problem the C value paradox grades of complexity of organization in eukaryotes are unrelated to their genome sizes

Electrical Potentials in Biological Membrane Transport E. Heinz, 2013-03-08 The material of this volume was originally planned to be incorporated in the preceding monograph *Mechanics and Energetics of Biological Transport* A separate and coherent treatment of the variety of bioelectrical phenomena was considered preferable mainly for didactic reasons Usually the biologist has to gather the principles of bioelectricity he needs from different sources and on different levels The present book intends to provide these principles in a more uniform context and in a form adjusted to the problems of a biologist rather than of a physicist or electrical engineer The main emphasis is put on the molecular aspect by relating the bioelectrical phenomena such as the membrane diffusion potentials pump potentials or redox potentials to the properties of the membrane concerned and as far as possible to specific steps of transport and metabolism of ions and nonelectrolytes Little space is devoted to the familiar and widely used representation of bioelectrical phenomena in terms of electrical networks of equivalent circuits with batteries resistances capacities etc In order to elucidate the basic principles the formal treatment is kept as simple as possible using highly simplified models based on biological systems The corresponding equations are derived in two ways kinetically i.e. in terms of the Law of Mass Action as well as energetically i.e. in terms of Nonequilibrium Thermodynamics **Connective Tissue** M.B. Mathews, 2012-12-06 This book deals primarily with the

principal extracellular macromolecules of animal connective tissues It attempts to answer some general questions about the biological organization of the tissues What is the nature of this organization at various dimensional levels What functions does the organization serve How has it evolved I have given major emphasis to the structures and properties of the macromolecular components of extracellular matrices from a wide range of invertebrates and vertebrates In doing so

however I have treated cursorily many important aspects of connective tissue biology that appeared to be only indirectly relevant to the principal questions asked. On the other hand I have not hesitated to search for broad biological principles outside the prevailing conceptual boundaries of animal connective tissues and the lower molecular dimensional levels. The numerous speculations presented will I hope stimulate the reader to further thought and investigation.

Acknowledgements

My previously unpublished data that have been included in this volume could not have been obtained without the generous gifts of specimens and other assistance by a number of individuals and institutions. These are Dr G BERENSON of Louisiana State University, Drs W P BRAKER and D ZUMWALD of the Shedd Aquarium, Dr S H CHUANG of the University of Singapore, Dr L G CLARK and Mr U M VARELA DIAZ of the University of Pennsylvania, the Government of Nicaragua, Dr E CLARK of Cape Haze Marine Laboratories, Mr C E Sensory Reception Y.A. Vinnikov, 2012-12-06. This book presents a distillation of many years of investigation by the author and his associates on the problem of sensory reception. Both our own data and data from the scientific literature on the electron microscopy, cytochemistry, biochemistry and electrophysiology of the organs of vision, taste, smell, hearing and gravitation are presented to show that the evolution of the sense organs of all animals on our planet is based on a receptor cell equipped with a motile antenna, a biological recorder of information concerning certain types of energy reaching the animal from the environment. The conversion or encoding of this energy into information is effected with the aid of special protein molecules positioned in the plasma membrane of the antennae. The action of the unit of energy of a stimulus on such a specific protein molecule causes a change of shape and this is the basis of the trigger mechanism of reception leading to the stimulation of the receptor cell and the transmission of the information encoded in this cell in the form of nerve impulses to the central nervous system. The present monograph summarizes over 30 years of working experience by the author and his associates in the field of evolution of sense organs. Material is used here from his earlier monographs: *The Retina of the Eye* (Vertebrates) 1947, *The Morphology of the Organ of Smell* 1957, *The Organ of Corti: Its Histophysiology and Histochemistry* 1964, written jointly with L K Chemical Relaxation in Molecular Biology I. Pecht, R. Rigler, 2012-12-06. The development of an area of scientific research is a dynamic process with its own kinetic equations and its own physical mechanism. The study of fast chemical interactions and transformations is such an area and while it is tempting to draw analogies or to speculate about the simplest model system, the lack of adequately averaged observables is an annoying obstacle to such an undertaking. Sciences suffering from such conditions usually avoid quantitative models, be they primitive or complex. Instead they prove their point by case histories. Chemical relaxation kinetics started as an offspring of research in acoustics. In some aqueous ionic solutions, anomalous acoustic absorption had been observed. A systematic study traced the cause of this absorption, showing that the covered frequency range and the intensity of the absorption were related in a predictable manner to the rate at which ions can interact and form structures differing in volume from the non-interacting species. The step from this experimental observation and its correct non-trivial explanation

to the discovery that all fast chemical processes must reveal themselves quantitatively in the relaxation rate of a perturbed equilibrium state and that perturbation parameters other than sound waves can be used for its exploitation was made by MANFRED EIGEN in 1954 The foresightedness of K F *Inhibitors of Nucleic Acid Synthesis* H. Kersten, W.

Kersten, 2013-11-11 During the last decade physical and chemical methods have improved rapidly a fact which allowed the mode of action of antibiotics to be studied and many biochemically oriented scientists have devoted their research to the following questions 1 What is the metabolic pathway that is inhibited selectively and what are the target molecules within a sensitive cell 2 What are the relationships between the chemical structure of an antibiotic and the physicochemical properties of the sensitive molecules 3 Why and how far is the action selective 4 Is it possible to correlate the interaction with the target molecules with the particular biological activities observed This monograph deals with those antibiotics which interfere with the biosynthesis of nucleic acids The idea was to provide an insight into how to investigate the preceding questions experimentally and to solve as yet unresolved problems rather than to give a review of the current state of knowledge Although the biochemistry of nucleic acid synthesis is known in general the precise molecular mechanisms by which deoxyribonucleic acid is replicated or transcribed has still to be clarified For this reason it is not yet possible to describe the molecular mechanisms by which the inhibitors of nucleic acid and protein synthesis exhibit their effects The fact that the inhibitors of nucleic acid and protein synthesis themselves served as useful tools to obtain an insight into the mechanisms of replication transcription and translation was one of the most exciting discoveries in this field **Mechanics**

and Energetics of Biological Transport E. Heinz, 2012-12-06 This book deals with energetics of transport processes largely expressed in terms of the thermodynamics of irreversible processes Since at the present time too little is known about the molecular mechanism of transport the present treatment is based largely on hypothetical models Care has been taken however to define the crucial features of these models as generally as possible so that the equations do not depend too much on hypothetical details Accordingly most equations though developed on the basis of a mobile carrier ferryboat model should apply equally to a conformational model with an appropriate reinterpretation of the symbols To better elucidate the essentials the models are greatly simplified by special assumptions Maximally only two flows are assumed to be present in each model at one time e.g. two solute flows the flow of solvent and of one solute the flow of solvent and of heat The simplifying assumptions may often be unreal Hence the equations should not be applied uncritically to actual mechanisms They may at best serve as a basis on which the more appropriate equations may be developed The book is not designed to give a complete kinetic analysis of the transport processes described The kinetic equations are kept to the minimum required to describe the model concerned and to relate it to the corresponding thermodynamic equations The intention is to stress the close relationship between biosmotic transport and biochemical processes in metabolism **Regulation of**

Transcription and Translation in Eukaryotes Ekkehard K.F. Bautz, P. Karlson, H. Kersten, 2012-12-06 This volume

represents the proceedings of the 24th Mosbach Colloquium on Regulation of Transcription and Translation in Eukaryotes which was held April 26-28 1973 in Mosbach Germany under the auspices of the Gesellschaft für Biologische Chemie To the three of us H KERSTEN P KARLSON and myself who were commissioned with the invitation of speakers it was a difficult decision as to whether we should attempt to cover with some twenty contributions as many aspects of this broad topic as possible or to sacrifice the intellectually perhaps more pleasing but more speculative concepts and to concentrate on a few aspects of gene expression in reasonable detail We unanimously decided on the latter course leaving such important and timely topics as for example hormone action cyclic AMP and reverse transcription to the proceedings of other symposia and concentrating on the four questions which are most basic to an understanding of the mechanisms of transcription and translation and for which fragmentary but nonetheless reliable experimental results have become available within the last few years These are the structure of chromatin the synthesis of messenger RNA the structure of the active ribosome and the role of initiation factors in protein synthesis

Transition Metals in Biochemistry A. S. Brill, 2012-12-06 Transition metal ions in biological systems are of interest in biology biochemistry chemistry medicine and physics Scientists with rather different viewpoints employing many methods have contributed to this area A concise review of the current state of the field will to some extent reflect the special knowledge of the person writing it in this case application of physical methods to the investigation of metal coordination X-ray diffraction is one of the most important of these methods but a useful treatment of X-ray structure analysis would be comparable in size with and beyond the scope of the monograph Many results of X-ray diffraction studies are of course presented Electron paramagnetic resonance spectroscopy has played a major part in the rapid advance in knowledge of the electronic structures of transition metal ions in biological systems More generally measurements involving light microwaves and magnetic fields are capable of producing much new information and the required instrumentation is available at most research institutions Therefore light absorption and paramagnetic resonance are treated in depth The principles described in the latter discussions are broadly applicable for example to the promising techniques of X-ray spectroscopy utilizing synchrotron radiation and lanthanide perturbed very high resolution nuclear magnetic resonance spectroscopy

Molecular Biology, Biochemistry, and Biophysics, 1975 **Ribonucleases, Part A: Functional Roles and Mechanisms of Action**, 2001-09-26 This first of two volumes provides up to date methods related information on ribonuclease functions assays and applications Chapter topics include the identification of characterization of and assays for secreted ribonucleases viral ribonucleases artificial and engineered ribonucleases and ribozymes 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

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