

BIOCHEMISTRY

THIRD EDITION

BIOCHEMISTRY

THIRD EDITION

LUBERT STRYER

Libert Sty-

STANFORD UNIVERSITY

Library of Congress Cataloging-in-Publication Data

Stryer, Lubert. Biochemistry.

Includes index.
1. Biochemistry. I. Title.
QP514.2.S66 1988 574.19'2 87-36486
ISBN 0-7167-1843-X
ISBN 0-7167-1920-7 (international student ed.)

Copyright © 1975, 1981, 1988 by Lubert Stryer

No part of this book may be reproduced by any mechanical, photographic, or electronic process, or in the form of a phonographic recording, nor may it be stored in a retrieval system, transmitted, or otherwise copied for public or private use, without written permission from the publisher.

Printed in the United States of America

234567890 RRD 6543210898

To my teachers

Paul F. Brandwein

Daniel L. Harris

Douglas E. Smith

Elkan R. Blout

Edward M. Purcell

PREFACE

to the Third Edition

Biochemistry has been profoundly transformed by recombinant DNA technology. The genome is now an open book—any passage can be read. The cloning and sequencing of millions of bases of DNA have greatly enriched our understanding of genes and proteins. Indeed, recombinant DNA technology has led to the integration of molecular genetics and protein chemistry. The intricate interplay of genotype and phenotype is now being unraveled at the molecular level. One of the fruits of this harvest is insight into how the genome is organized and its expression is controlled. The molecular circuitry of growth and development is coming into view. The reading of the genome is also providing a wealth of amino acid sequence information that illuminates the entire protein landscape. Scarce proteins can be produced in abundance by transfected cells. Moreover, precisely designed novel proteins can be generated by site-specific mutagenesis to elucidate how proteins fold, catalyze reactions, transduce signals, transport ions, and interconvert different forms of free energy.

Our understanding of molecular evolution also has been greatly enriched by the recombinant DNA revolution. Families and superfamilies of proteins have come into view. Theme and variation at the level of proteins are vivid expressions of the underlying processes of gene duplication and divergence. The genes of complex proteins display the coming together in evolution of exons encoding functional modules. The many recurring structural and mechanistic motifs seen throughout nature testify to the fundamental unity of all forms of life. The discovery of catalytic RNA enables us to envision an RNA world early in the evolution of life, prior to the appearance of DNA and protein. The ubiquity of ribonucleotides in metabolism and the central roles played

xxvi PREFACE by them are reflections of their ancient origins—of the early RNA world, when RNA served both as gene and enzyme.

These remarkable advances compel a major change in the way biochemistry is taught. I have altered the architecture of this book to provide a new framework for the exposition of fundamental themes and principles of biochemistry. The book begins with a new part, entitled Molecular Design of Life, that provides an overview of the central molecules of life—DNA, RNA, and proteins—and their interplay. Recombinant DNA technology and other experimental methods for exploring proteins and genes are also presented in this part. This introduction prepares the reader for the detailed consideration of protein structure and function that follows. The teaching of metabolism is likewise enriched by this new organization. The other major structural changes are the addition of a chapter on carbohydrates and one on protein targeting. Many sections of the book have been extensively revised and hundreds of new illustrations have been added. I have tried to preserve the unity of biochemistry as an intellectual discipline. My goal has been to make this powerful language comprehensible and to share its beautiful imagery.

I am grateful to Alexander Glazer, Daniel Koshland, Jr., and Alexander Rich for having encouraged me to write this edition. I would not have embarked on this endeavor had it not been for their warm support and good counsel.

The planning of this edition unexpectedly took place in terrain quite different from Yale, Stanford, and Aspen, where the first two editions took form. In December 1985, my family and I went to Nepal to trek in the Everest region. After two rewarding days in Katmandu, we were on the verge of boarding the plane to Lukla, only to be turned back with the disappointing news that the landing strip was closed because of snow. We then headed for the Annapurna region in four-wheel drive vehicles but had to return a day later because the road vanished in the heavy rain. The inaccessibility of the high Himalayas led to an abrupt change of itinerary. We flew to Bangkok and arrived in 95-degree heat, carrying our parkas and arctic sleeping bags. Instead of hiking at 12,000 feet, we found ourselves at a hotel pool at sea level. This dislocation led to a totally unforeseen benefit. I was able to unhurriedly reflect on the remarkable development of biochemistry since I wrote my last edition. I had the leisure to think and dream and plan this book. Best of all, I was able to share my thoughts with my son Daniel, who was then a senior majoring in human biology, and gain from his insights.

Alexander Glazer, Richard Gumport, Roger Koeppe, James Rawn, Carl Rhodes, and Peter Rubenstein read the entire manuscript. I have benefited greatly from their scholarly and perceptive criticism. Steve Block, Daniel Branton, Simone Brutlag, Carolyn Cohen, Jeffrey Critchfield, Peter Cullis, Russell Doolittle, Marilyn Farquhar, Robert Fletterick, Robert Fox, Michel Goldberg, Jack Griffith, James Hageman, Stephen Harrison, Brian Holl, Leroy Hood, Horace Jackson, Gunther Kohlhaw, Arthur Kornberg, Roger Kornberg, Stephen Kron, Michael Levitt, Bo Malmström, Lynne Mercer, Albert Mildvan, Jeremy Nathans, Christopher Newgard, Marion O'Leary, George Palade, Peter Parham, Frederic Richards, Ed Rock, Gottfried Schatz, Gray Scrimgeour, Paul Sigler, Jeffrey Sklar, James Spudich, Thomas Steitz, Nigel Unwin, Ronald Vale, William Wickner, and Robley Williams also gave valuable advice and help.

The contributors of many striking and informative illustrations are acknowledged in the figure legends. I am also indebted to crystallog-

xxvii PREFACE

raphers who have deposited the atomic coordinates of their solved structures in the Protein Data Bank, a valuable resource maintained by Brookhaven National Laboratory. Many new figures depicting molecular structure were generated on our departmental molecular graphics computer facility. David Austen and William Hurja helped me use this excellent system.

I was able to concentrate on the writing of this book because my office was in the capable hands of Joanne Tisch. She played a critical role in preparing the manuscript and reading the proofs. Her sensitivity, intelligence, and good spirits lightened my load. The Medline bibliographic retrieval system of the National Library of Medicine greatly facilitated my search of the literature. The staff of the Lane Medical Library and Falconer Biology Library of Stanford University were most helpful in locating books and references.

Andrew Kudlacik edited this manuscript with a fine sense of style and meaning. Mike Suh skillfully integrated word and picture in the design of each page. Susan Moran kept a watchful and discerning eye over many thousands of pages of manuscript, figures, and proofs. I also wish to thank Tom Cardamone and Shirley Baty for many outstanding drawings.

I am grateful to my family for their sustained support of this endeavor, which was more arduous than anticipated. My sons, Michael and Daniel, now embarked on their own careers, cheered me from afar. My wife, Andrea, provided criticism, advice, and encouragement in just the right proportions. I have been nurtured, too, by many who have reached out to express their warmth and interest in continuing this dialogue of biochemistry. I feel very fortunate and privileged to partake in this process at such a wonderful time.

Lubert Stryer
DECEMBER 1987

PREFACE

to the Second Edition

The pace of discovery in biochemistry has been exceptionally rapid during the past several years. This progress has greatly enriched our understanding of the molecular basis of life and has opened many new areas of inquiry. The sequencing of DNA, the construction and cloning of new combinations of genes, the elucidation of metabolic control mechanisms, and the unraveling of membrane transport and transduction processes are some of the highlights of recent research. One of my aims in this edition has been to weave new knowledge into the fabric of the text. I have sought to enhance the book's teaching effectiveness by centering the exposition of new material on common themes wherever feasible and by citing recurring motifs. I have also tried to convey a sense of the intellectual power and beauty of the discipline of biochemistry.

I am indebted to Thomas Emery, Henry Epstein, Alexander Glazer, Roger Kornberg, Robert Martin, and Jeffrey Sklar for their counsel, criticism, and encouragement in the preparation of this edition. Robert Baldwin, Charles Cantor, Richard Caprioli, David Eisenberg, Alan Fersht, Robert Fletterick, Herbert Friedmann, Horace Jackson, Richard Keynes, Sung-Hou Kim, Aaron Klug, Arthur Kornberg, Daniel Koshland, Jr., Samuel Latt, Vincent Marchesi, David Nelson, Garth Nicolson, Vernon Oi, Robert Renthal, Carl Rhodes, Frederic Richards, James Rothman, Peter Sargent, Howard Schachman, Joachim Seelig, Eric Shooter, Elizabeth Simons, James Spudich, Theodore Steck, Thomas Steitz, Judit C.-P. Stenn, Robert Trelstad, Christopher Walsh, Simon Whitney, and Bernhard Witkop also gave valuable advice.

Patricia Mittelstadt edited both editions of this text. I deeply appreciate her critical and sustained contributions. I am indebted to Donna

XXX PREFACE Salmon for her outstanding drawings. David Clayton, David Dressler, John Heuser, Lynne Mercer, Kenneth Miller, George Palade, Nigel Unwin, and Robley Williams generously provided many fine electron micrographs. Betty Hogan typed the manuscript and played an indispensable role in its preparation. Cary Leiden and Karen Marzotto carefully read the proofs. I also wish to thank Michael Graves for his excellent photographic work.

My wife, Andrea, and my sons, Michael and Daniel, have cheerfully allowed this text to become a member of the family. I am deeply grateful to them for their patience and buoyancy. Andrea provided much advice on style and design, as she did for the first edition.

I have been heartened by the many letters that I have received from readers of the first edition. Their comments and criticisms have enlightened, stimulated, and encouraged me. I look forward to a continuing dialogue with readers in the years ahead.

Lubert Stryer
AUGUST 1980

PREFACE

to the First Edition

This book is an outgrowth of my teaching of biochemistry to undergraduates, graduate students, and medical students at Yale and Stanford. My aim is to provide an introduction to the principles of biochemistry that gives the reader a command of its concepts and language. I also seek to give an appreciation of the process of discovery in biochemistry. My exposition of the principles of biochemistry is organized around several major themes:

- Conformation—exemplified by the relationship between the three-dimensional structure of proteins and their biological activity
- 2. Generation and storage of metabolic energy
- 3. Biosynthesis of macromolecular precursors
- 4. Information—storage, transmission, and expression of genetic information
- 5. Molecular physiology—interaction of information, conformation, and metabolism in physiological processes

The elucidation of the three-dimensional structure of proteins, nucleic acids, and other biomolecules has contributed much in recent years to our understanding of the molecular basis of life. I have emphasized this aspect of biochemistry by making extensive use of molecular models to give a vivid picture of architecture and dynamics at the molecular level. Another stimulating and heartening aspect of contemporary biochemistry is its increasing interaction with medicine. I have presented many examples of this interplay. Discussions of molecular diseases such as sickle-cell anemia and of the mechanism of action of drugs such as penicillin enrich the teaching of biochemistry. Finally, I have tried to define several challenging areas of inquiry in biochemistry today, such as the molecular basis of excitability.

xxxii PREFACE

In writing this book, I have benefitted greatly from the advice, criticism, and encouragement of many colleagues and students. Leroy Hood, Arthur Kornberg, Jeffrey Sklar, and William Wood gave me invaluable counsel on its overall structure. Richard Caprioli, David Cole, Alexander Glazer, Robert Lehman, and Peter Lengyel read much of the manuscript and made many very helpful suggestions. I am indebted to Frederic Richards for sharing his thoughts on macromolecular conformation and for extensive advice on how to depict threedimensional structures. Deric Bownds, Thomas Broker, Jack Griffith, Hugh Huxley, and George Palade made available to me many striking electron micrographs. I am also very thankful for the advice and criticism that were given at various times in the preparation of this book by Richard Dickerson, David Eisenberg, Moises Eisenberg, Henry Epstein, Joseph Fruton, Michel Goldberg, James Grisolia, Richard Henderson, Harvey Himel, David Hogness, Dale Kaiser, Samuel Latt, Susan Lowey, Vincent Marchesi, Peter Moore, Allan Oseroff, Jordan Pober, Russell Ross, Edward Reich, Mark Smith, James Spudich, Joan Steitz, Thomas Steitz, and Alan Waggoner.

I am grateful to the Commonwealth Fund for a grant that enabled me to initiate the writing of this book. The interest and support of Robert Glaser, Terrance Keenan, and Quigg Newton came at a critical time. One of my aims in writing this book has been to achieve a close integration of word and picture and to illustrate chemical transformations and three-dimensional structures vividly. I am especially grateful to Donna Salmon, John Foster, and Jean Foster for their work on the drawings, diagrams, and graphs. Many individuals at Yale helped to bring this project to fruition. I particularly wish to thank Margaret Banton and Sharen Westin for typing the manuscript, William Pollard for photographing space-filling models, and Martha Scarf for generating the computer drawings of molecular structures on which many of the illustrations in this book are based. John Harrison and his staff at the Kline Science Library helped in many ways.

Much of this book was written in Aspen. I wish to thank the Aspen Center of Physics and the Given Institute of Pathobiology for their kind hospitality during several summers. I have warm memories of many stimulating discussions about biochemistry and molecular aspects of medicine that took place in the lovely garden of the Given Institute and while hiking in the surrounding wilderness areas. The concerts in Aspen were another source of delight, especially after an intensive day of writing.

I am deeply grateful to my wife, Andrea, and to my children, Michael and Daniel, for their encouragement, patience, and good spirit during the writing of this book. They have truly shared in its gestation, which was much longer than expected. Andrea offered advice on style and design and also called my attention to the remark of the thirteenth-century Chinese scholar Tai T'ung (*The Six Scripts: Principles of Chinese Writing*): "Were I to await perfection, my book would never be finished."

I welcome comments and criticisms from readers.

Lubert Stryer
OCTOBER 1974

BIOCHEMISTRY

On the facing page: Proteins and nucleic acids are the central molecules of life. A complex of DNA bound to the protein that replicates it is shown here. DNA polymerase is shown in blue, one strand of the double-helical DNA template in green, and the other in red. [Courtesy of Dr. Thomas Steitz.]

Contents

List of Topics ix Preface to the Third Edition xxv Preface to the Second Edition xxix Preface to the First Edition xxxi

PART I MOLECULAR DESIGN OF LIFE 1

- CHAPTER 1. Prelude 3
 - 2. Protein Structure and Function 15
 - 3. Exploring Proteins 43
 - 4. DNA and RNA: Molecules of Heredity 71
 - 5. Flow of Genetic Information 91
 - 6. Exploring Genes: Analyzing, Constructing, and Cloning **DNA** 117

PART II PROTEIN CONFORMATION, DYNAMICS, AND **FUNCTION 141**

- CHAPTER 7. Oxygen-transporting Proteins: Myoglobin and Hemoglobin 143
 - 8. Introduction to Enzymes 177
 - 9. Mechanisms of Enzyme Action 201
 - 10. Control of Enzymatic Activity 233
 - 11. Connective-Tissue Proteins 261
 - 12. Introduction to Biological Membranes 283

CONTENTS

PART III GENERATION AND STORAGE OF METABOLIC ENERGY 313

- CHAPTER 13. Metabolism: Basic Concepts and Design 315
 - 14. Carbohydrates 331
 - 15. Glycolysis 349
 - 16. Citric Acid Cycle 373
 - 17. Oxidative Phosphorylation 397
 - 18. Pentose Phosphate Pathway and Gluconeogenesis 427
 - 19. Glycogen Metabolism 449
 - 20. Fatty Acid Metabolism 469
 - 21. Amino Acid Degradation and the Urea Cycle 495
 - 22. Photosynthesis 517

PART IV BIOSYNTHESIS OF MACROMOLECULAR PRECURSORS 545

- CHAPTER 23. Biosynthesis of Membrane Lipids and Steroid Hormones 547
 - 24. Biosynthesis of Amino Acids and Heme 575
 - 25. Biosynthesis of Nucleotides 601
 - 26. Integration of Metabolism 627

PART V GENETIC INFORMATION: storage, transmission, and expression 647

- CHAPTER 27. DNA Structure, Replication, and Repair 649
 - 28. Gene Rearrangements: Recombination and Transposition 687
 - 29. RNA Synthesis and Splicing 703
 - 30. Protein Synthesis 733
 - 31. Protein Targeting 767
 - 32. Control of Gene Expression in Procaryotes 799
 - 33. Eucaryotic Chromosomes and Gene Expression 823
 - 34. Viruses and Oncogenes 851

PART VI MOLECULAR PHYSIOLOGY: interaction of information, conformation, and metabolism in physiological processes 887

- CHAPTER 35. Molecular Immunology 889
 - 36. Muscle Contraction and Cell Motility 921
 - 37. Membrane Transport 949
 - 38. Hormone Action 975
 - 39. Excitable Membranes and Sensory Systems 1005 Appendixes 1044 Answers to Problems 1049 Index 1065

List of Topics

PART I MOLECULAR DESIGN OF LIFE 1



CHAPTER 1 Prelude 3

Molecular models depict three-dimensional structure 4 Space, time, and energy 5

Reversible interactions of biomolecules are mediated by three kinds of noncovalent bonds 7

The biologically important properties of water are its polarity and cohesiveness 9

Water solvates polar molecules and weakens ionic and hydrogen bonds 9

Hydrophobic interactions: nonpolar groups tend to associate in water 10

Design of this book 11

CHAPTER 2 Protein Structure and Function 15

Proteins are built from a repertoire of twenty amino acids 16

Amino acids are linked by peptide bonds to form polypeptide chains 22

Proteins have unique amino acid sequences that are specified by genes 23

Protein modification and cleavage confer new capabilities 24

The peptide unit is rigid and planar 25

Polypeptide chains can fold into regular structures: the α helix and β pleated sheets 25

Polypeptide chains can reverse direction by making β-turns 28

Proteins are rich in hydrogen-bonding potentiality 28 Water-soluble proteins fold into compact structures with nonpolar cores 29

Levels of structure in protein architecture 31 Amino acid sequence specifies three-dimensional structure 32

Proteins fold by the association of α -helical and β -strand segments 34

Prediction of conformation from amino acid sequence 35

Essence of protein action: specific binding and transmission of conformational changes 37 Appendix: Acid-base concepts 41

CHAPTER 3 Exploring Proteins 43

Proteins can be separated by gel electrophoresis and displayed 44

Proteins can be purified according to size, charge, and binding affinity 46

Ultracentrifugation is valuable for separating biomolecules and determining molecular weights 49

Amino acid sequences can be determined by automated Edman degradation 50

Proteins can be specifically cleaved into small peptides to facilitate analysis 55

Recombinant DNA technology has revolutionized protein sequencing 57

Amino acid sequences provide many kinds of insights

X-ray crystallography reveals three-dimensional structure in atomic detail 59

Proteins can be quantitated and localized by highly specific antibodies 62

Peptides can be synthesized by automated solid-phase methods 64

CHAPTER 4 DNA and RNA: Molecules of Heredity 71

DNA consists of four kinds of bases joined to a sugarphosphate backbone 72

Transformation of pneumococci by DNA revealed that genes are made of DNA 73

The Watson-Crick DNA double helix 76

The complementary chains act as templates for each other in DNA replication 78

DNA replication is semiconservative 79

The double helix can be reversibly melted 80

DNA molecules are very long 82

Some DNA molecules are circular and supercoiled 83 DNA is replicated by polymerases that take instructions from templates 84

Some viruses have single-stranded DNA during part of their life cycle 85

The genes of some viruses are made of RNA 86 RNA tumor viruses replicate through double-helical DNA intermediates 87

CHAPTER 5 Flow of Genetic Information 91

Several kinds of RNA play key roles in gene expression 92

Formulation of the concept of messenger RNA 93 Experimental evidence for messenger RNA, the informational intermediate in protein synthesis 94 Hybridization studies showed that messenger RNA is

complementary to its DNA template 95

Ribosomal RNA and transfer RNA are also synthesized on DNA templates 96

All cellular RNA is synthesized by RNA polymerases 96

RNA polymerase takes instructions from a DNA template 97

Transcription begins near promoter sites and ends at terminator sites 98

Transfer RNA is the adaptor molecule in protein synthesis 99

Amino acids are coded by groups of three bases starting from a fixed point 99

Deciphering the genetic code: synthetic RNA can serve as messenger 101

Trinucleotides promote the binding of specific transfer RNA molecules to ribosomes 103

Copolymers with a defined sequence were also instrumental in breaking the code 104 Major features of the genetic code 106 Start and stop signals for protein synthesis 107

The genetic code is nearly universal 108

The sequences of genes and their encoded proteins are colinear 109

Most eucaryotic genes are mosaics of introns and exons 110

Many exons encode protein domains 111 RNA probably came before DNA and proteins in evolution 113

CHAPTER 6 Exploring Genes: Analyzing, Constructing, and Cloning DNA 117

Restriction enzymes split DNA into specific fragments

Restriction fragments can be separated by gel electrophoresis and visualized 119

DNA can be sequenced by specific chemical cleavage (Maxam-Gilbert method) 120

DNA can be sequenced by controlled interruption of replication (Sanger dideoxy method) 121

DNA probes and genes can be synthesized by automated solid-phase methods 123

New genomes can be constructed, cloned, and expressed 124

Restriction enzymes and DNA ligase are key tools in forming recombinant DNA molecules 126

Plasmids and lambda phage are choice vectors for DNA cloning in bacteria 127

Specific genes can be cloned from a digest of genomic DNA 130

Complementary DNA (cDNA) prepared from mRNA can be expressed in host cells 132

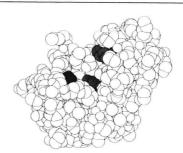
New genes inserted into eucaryotic cells can be efficiently expressed 133

Tumor-inducing (Ti) plasmids can be used to bring new genes into plant cells 135

Novel proteins can be engineered by site-specific mutagenesis 136

Recombinant DNA technology has opened new vistas 137

PART II PROTEIN CONFORMATION, DYNAMICS, AND FUNCTION 141



CHAPTER 7 Oxygen-transporting Proteins: Myoglobin and Hemoglobin 143

Oxygen binds to a heme prosthetic group 144 Myoglobin was the first protein to be seen at atomic resolution 144