



Equilibria and Kinetics of Biological Macromolecules

Jan Hermans Barry Lentz



WILEY

Copyright © 2014 by John Wiley & Sons, Inc. All rights reserved.

Published by John Wiley & Sons, Inc., Hoboken, New Jersey. Published simultaneously in Canada.

No part of this publication may be reproduced, stored in a retrieval system, or transmitted in any form or by any means, electronic, mechanical, photocopying, recording, scanning, or otherwise, except as permitted under Section 107 or 108 of the 1976 United States Copyright Act, without either the prior written permission of the Publisher, or authorization through payment of the appropriate per-copy fee to the Copyright Clearance Center, Inc., 222 Rosewood Drive, Danvers, MA 01923, (978) 750-8400, fax (978) 750-4470, or on the web at www.copyright.com. Requests to the Publisher for permission should be addressed to the Permissions Department, John Wiley & Sons, Inc., 111 River Street, Hoboken, NJ 07030, (201) 748-6011, fax (201) 748-6008, or online at http://www.wiley.com/go/permission.

Limit of Liability/Disclaimer of Warranty: While the publisher and author have used their best efforts in preparing this book, they make no representations or warranties with respect to the accuracy or completeness of the contents of this book and specifically disclaim any implied warranties of merchantability or fitness for a particular purpose. No warranty may be created or extended by sales representatives or written sales materials. The advice and strategies contained herein may not be suitable for your situation. You should consult with a professional where appropriate. Neither the publisher nor author shall be liable for any loss of profit or any other commercial damages, including but not limited to special, incidental, consequential, or other damages.

For general information on our other products and services or for technical support, please contact our Customer Care Department within the United States at (800) 762-2974, outside the United States at (317) 572-3993 or fax (317) 572-4002.

Wiley also publishes its books in a variety of electronic formats. Some content that appears in print may not be available in electronic formats. For more information about Wiley products, visit our web site at www.wiley.com.

Library of Congress Cataloging-in-Publication Data:

Hermans, Jan.

Equilibria and kinetics of biological macromolecules / by Jan Hermans and Barry Lentz.

p.; cm.

Includes index.

ISBN 978-1-118-47970-4 (cloth)

I. Lentz, Barry. II. Title.

[DNLM: 1. Biocompatible Materials-pharmacokinetics. 2. Biophysical Processes. 3.

Macromolecular Substances-pharmacokinetics. 4. Molecular Conformation. QT 37] RM301.5

615.7-dc23

2013013996

Printed in the United States of America

Equilibria and Kinetics of Biological Macromolecules

For Harold Scheraga

... Bin' i

此为试读,需要完整PDF请访问: www.ertongbook.com

Preface

It is only by attempting to explain our science to each other that we find out what we really know.

—J. M. Ziman, Nature 252: 318-324 (1969)

This book has grown out of circa 12 years of collaborative teaching of a 6-credit biophysics course that forms the core of the didactic teaching for the Molecular and Cellular Biophysics Program at UNC CH. Thus the book is directed at an audience of first year graduate students. However, the book has grown well beyond the content of those courses, also thanks to input and suggestions from colleagues who have shared our teaching the course (see Acknowledgments), and it is our hope that it will prove useful to working biochemists who seek a deeper understanding of modern biophysics.

The book is not meant to be a complete text in biophysics, as it focuses on the input of physics and physical chemistry to experimental studies and theoretical models of equilibria and kinetics of biological macromolecules (largely, proteins). A chapter is devoted to methods of molecular simulations; applications of molecular dynamics are included in several chapters. On the other hand, we limited the size of this book by devoting no space to spectroscopy and structure determination.

The book assumes some knowledge of physics and/or physical chemistry, but in Part 1, the chapters on thermodynamics, simple quantum mechanics and molecular structure and intra- and intermolecular forces shore up what may be shaky backgrounds of some students, and provide references for later chapters. Part 1 concludes with a chapter on water and the hydrophobic effect.

Two chapters in Part 2 introduce various ensembles of statistical mechanics, and these are followed by the aforementioned chapter on molecular simulations.

Next, in Part 3, we discuss equilibria of binding of "ligands" to macromolecules from different standpoints: chemical equilibrium theory, thermodynamics, and statistical mechanics. These are followed by a discussion of linked equilibria, and a chapter that focuses on hemoglobin as an example of allosteric control of function.

Part 3 concludes with a chapter on charge-charge interactions of macromolecules in solution.

In Part 4, we deal with folding equilibria. A brief overview of the physics of polymer solutions is followed by a chapter on the theory of helix-coil transitions of polypeptides and its many applications, and it ends with a section on helix-coil equilibria of double-stranded nucleic acids. This is followed by a long chapter on equilibria of protein folding. Part 4 concludes with a chapter on elasticity with elastin and tenascin as examples of two different mechanisms.

The final part of the book is devoted to kinetics. The first chapter describes kinetic measurement methods and a variety of kinetic models, ranging from simple rate equations to transition state theory. This is followed by a chapter on experiments and theory of kinetics of protein folding. Part 5 concludes with a chapter on stochastic processes and theories from the Langevin equation to Kramers' theory of reaction rates.

Finally, in a series of Appendices we have covered technical (mostly mathematical) details which we had skipped earlier to make the main content of this book easier to follow.

The authors will maintain a web page devoted to corrections and discussion of this book. Please consult the authors' personal web pages at the University of North Carolina.

Acknowledgments

This book's inception was in the form of lecture notes for the introductory class in molecular biophysics given at UNC each fall semester. An enormous help has been the feedback we received from students taking the class.

We have received help from many colleagues. We are grateful to professors Papoian (now at the University of Maryland) and Dokholyan, who have each taught part of the course, for letting us base important sections of the book on new presentations given by them in their lectures. Individual chapters have had input from Gary Ackers at Washington University, from Gary Pielak in the UNC Chemistry Department, from Gary Felsenfeld at the NIH, from Andy McCammon at UCSD, from Weitao Yang at Duke, from Austen Riggs at the University of Texas, from Robert Baldwin at Stanford and from Hao Hu at the University of Hong Kong.

We thank Dr. M. Hanrath, University of Cologne for the computer drawings of hydrogen atom wave functions shown in Chapter 2, and Dr. Chad Petit for microcalorimeter results discussed in Chapter 8. Some figures of molecular structures were prepared with the vmd graphics program.* We acknowledge many answers to questions involving basic Physics, found by consulting Wikipedia.

JH and BRL

September 2012

^{*}Humphrey, W., Dalke, A., Schulten, K. VMD - visual molecular dynamics. *J. Mol. Graphics Modell.* **14**: 33–38 (1996).

CONSTANTS

Elementary charge	$e = 1.602189 \times 10^{-19} \text{ C}$
Atomic mass unit	amu = $1.660566 \times 10^{-27} \text{ kg}$
Proton rest mass	$m_{\rm p} = 1.672648 \times 10^{-27} \text{ kg} = 1.0073 \text{ amu}$
Electron rest mass	$m_{\rm e} = 9.109534 \times 10^{-31} \text{ kg}$
Boltzmann's constant	$k_{\rm B} = 1.380662 \times 10^{-23} \mathrm{J K^{-1}}$
	$= 8.31441 \text{ J K}^{-1} \text{ mol}^{-1} = 1.987 \text{ cal K}^{-1} \text{ mol}^{-1}$
Planck's constant	$h = 6.626176 \times 10^{-34} \text{ J s}$
Avogadro's number	$N_{\rm A} = 6.022 \times 10^{23} \rm mol^{-1}$
$k_{\rm B}T$ at 300 K	$k_{\rm B}T = 4.142 \text{ zJ} = 2.494 \text{ kJ mol}^{-1} = 0.596 \text{ kcal mol}^{-1}$
Freezing point of water	273.15 K
Viscosity of water at 20 °C	$1.003 \times 10^{-3} \text{ Pa s}$
Gravity	$g = 9.8066 \text{ m s}^{-2}$
Permittivity of vacuum	$\varepsilon_0 = 8.854 \times 10^{-12} \text{ C}^2 \text{ N}^{-1} \text{ m}^{-2}$

PREFIXES OF THE METRIC SYSTEM

Name	Prefix	Value	Name	Prefix	Value
Name			Name	FIEIIX	
yotta	Y	10^{24}	yocto	y	10^{-24}
zotta	Z	10^{21}	zepto	Z	10^{-21}
exa	E	10^{18}	atto	a	10^{-18}
peta	P	10^{15}	femto	f	10^{-15}
tera	T	10^{12}	pico	p	10^{-12}
giga	G	10^{9}	nano	n	10^{-9}
mega	M	10^{6}	micro	μ	10^{-6}
kilo	k	10^{3}	milli	m	10^{-3}
		10^{0}			10^{0}

Also deka (da, 10^1), hekto (h, 10^2), deci (d, 10^{-1}) and centi (c, 10^{-2}).

GREEK ALPHABET

α	A	alpha	η	Н	eta	ν	N	nu	τ	T	tau
β	В	beta	θ , ϑ	Θ	theta	ξ	Ξ	xi	υ	Y,Υ	upsilon
γ	Γ	gamma	ι	I	iota	0	O	omicron	ϕ	Φ	phi
δ	Δ	delta	κ	K	kappa	π	П	pi	χ	X	chi
ε	E	epsilon	λ	Λ	lambda	ρ	P	rho	ψ	Ψ	psi
5	Z	zeta	μ	M	mu	σ, ς	Σ	sigma	ω	Ω	omega

Suggested pronunciation:

like "pie": π (pie), χ (kie);

like "bee": ξ (ksee), φ (fee), ψ (psee)

Contents

Pre	reface			
Acl	cknowledgments			
PA]	RT 1	BASIC PRINCIPLES		
1.	Ther	modynamics		
	1.1	Introduction / 3		
	1.2	The fundamental postulates or Laws of thermodynamics / 4		
	1.3	Other useful quantities and concepts / 14		
	1.4	Thermodynamics of the ideal gas / 19		
	1.5	Thermodynamics of solutions / 20		
	1.6	Phase equilibria / 25		
	1.7	Chemical equilibria / 29		
	1.8	Temperature dependence of chemical equilibria: The van't Hoff equation $/$ 31		
	1.9	Microcalorimetry / 31		
		Notes / 33		

2.		Four Basic Quantum Mechanical Models of Nuclear and Electronic Motion: A Synopsis		
	2.1	Introduction / 35		
	2.2	Fundamental hypotheses of quantum theory / 36		
	2.3	Three simple models of nuclear motion / 38		
	2.4	Hydrogen atomic orbitals: A simple model of electronic motion in atoms / 44		
	2.5	Many electron atoms / 47		
		Notes / 49		
		Suggested reading / 49		
3.	Mole	cular Structure and Interactions	51	
	3.1	Introduction / 51		
	3.2	Chemical bonding: Electronic structure of molecules / 51		
	3.3	Empirical classical energy expressions / 58		
	3.4	Noncovalent forces between atoms and molecules / 62		
	3.5	Molecular mechanics / 70		
		Notes / 75		
		Suggested reading / 76		
4.	Wate	r and the Hydrophobic Effect	77	
	4.1	Introduction / 77		
	4.2	Structure of liquid water / 78		
	4.3	The hydrophobic effect / 84		
		Notes / 89		
		Suggested reading / 89		
PA]	RT 2	STATISTICAL MECHANICS: THE MOLECULAR BASIS OF THERMODYNAMICS	91	
5.	The N	Molecular Partition Function	93	
	5.1	Introduction / 93		
	5.2	The Maxwell-Boltzmann distribution / 93		

5.3	The molecular partition function and thermodynamic functions / 99	
5.4	Application to macromolecules / 101	
	Notes / 108	
	Suggested reading / 110	
Syste	m Ensembles and Partition Functions	111
6.1	Introduction / 111	
6.2	Closed systems: The canonical ensemble / 112	
6.3	The canonical partition function of systems with continuous energy distributions: The phase-space integral / 119	
6.4	Application: Relation between binding and molecular interaction energy $/\ 123$	
6.5	Application: Binding of ligand to a macromolecule / 125	
6.6	Open systems: The grand canonical ensemble or grand ensemble / 127	
6.7	Fluctuations / 131	
6.8	Application: Light scattering as a measure of fluctuations of concentration / 134	
	Notes / 135	
	Suggested reading / 136	
Samp	oling Molecular Systems with Simulations	137
7.1	Introduction / 137	
7.2	Background / 138	
7.3	Molecular dynamics / 139	
7.4	Metropolis Monte Carlo / 142	
7.5	Simulation of a condensed system / 143	
7.6	Connecting microscopic and macroscopic system properties / 144	ļ
7.7	An example: Dynamics of Ace-Ala-Nme in solution / 146	
7.8	Forced transitions / 149	
7.9	Potential of mean force for changes of chemistry: "Computer Alchemy" / 152	

6.

7.

		Notes / 158	
		Suggested reading / 159	
PAI	RT 3	BINDING TO MACROMOLECULES	161
8.	Binding Equilibria		163
	8.1	Introduction / 163	
	8.2	Single-site model / 163	
	8.3	Measuring ligand activity and saturation / 166	
	8.4	Multiple sites for a single ligand / 173	
	8.5	A few practical recommendations / 182	
		Notes / 183	
		Suggested reading / 184	
9.	Ther	modynamics of Molecular Interactions	185
	9.1	Introduction / 185	
	9.2	Relation between binding and chemical potential: Unified formulation of binding and "exclusion" / 186	
	9.3	Free energy of binding / 187	
	9.4	Mutual response / 188	
	9.5	Volume exclusion / 189	
	9.6	Accounting for interactions of macromolecule and solvent components / 193	
		Notes / 196	
		Suggested reading / 196	
10.		ents of Statistical Mechanics of Liquids Solutions	197
	10.1	Introduction / 197	
	10.2	Partition function of ideal solution from thermodynamics / 198	
	10.3	Statistical mechanics of the ideal solution / 200	

7.10 The potential of mean force and the association equilibrium

constant of methane / 157

	10.5	A purely statistical mechanical formulation of molecular binding interactions $\ / \ 204$	
	10.6	Statistical mechanical models of nonideal solutions and liquids / 208	
		Notes / 211	
		Suggested reading / 211	
11.	Analy Funct	ysis of Binding Equilibria in Terms of Partition tions	213
	11.1	Alternate equivalent representations of the partition function / 21	13
	11.2	General implications / 215	
	11.3	Site-specific binding: General formulation / 216	
	11.4	Use of single-site binding constants / 218	
	11.5	Partition function for site binding: One type of ligand, independent multiple sites / 220	
	11.6	Site binding to interdependent or coupled sites / 221	
		Suggested reading / 222	
12.	Coup	led Equilibria	223
	12.1	Introduction / 223	
	12.2	Simple case: Coupling of binding (one site) and conformation change / 224	
	12.3	Coupling of binding to multiple sites and conformation change / 225	
	12.4	Free energy of binding can "drive" conformation change / 230	
	12.5	Formation of oligomers and polymers / 232	
	12.6	Coupled polymerization and ligand binding / 237	
		Notes / 238	
		Suggested reading / 238	
13.	Allost	eric Function	239
	13.1	Introduction / 239	

10.4 Formulation of molecular binding interactions in terms of a partition function: Empirical approach based on

thermodynamics / 202

14.

13.2 Background on hemoglobin / 240

13.3	The allosteric or induced-fit model of hemoglobin / 241
13.4	Simplified allosteric models: Concerted and sequential / 242
13.5	Numeric example / 244
13.6	Comparison of oxygen binding curves / 245
13.7	Separating oxygen binding and conformation change of hemoglobin / 246
13.8	Experiments with hybrid hemoglobins / 248
13.9	Two-site proteins, half-the-sites reactivity, and negative cooperativity / 248
13.10	Allosteric effects in protein function / 249
13.11	Sickle cell hemoglobin / 250
13.12	Hill plot / 250
	Notes / 252
	Suggested reading / 253
_	ged Groups: Binding of Hydrogen Ions, Solvation, Charge-Charge Interactions 255
14.1	Introduction / 255
14.2	Ionizable groups in peptides / 256
14.3	pH titration of a protein: Ribonuclease—normal and abnormal ionizable groups / 257
14.4	Local interactions cause pK_a s to be abnormal / 260
14.5	Internal charge-charge interactions: Ion pairs or salt bridges / 260
14.6	Measuring stability of salt bridges from double mutant cycles / 261
14.7	Salt bridges stabilize proteins from thermophilic organisms / 262
14.8	Charged side chains in enzyme catalysis and protein solubility / 263
14.9	Accounting for charge-charge and charge-solvent interactions / 263
14.10	
	The continuum dielectric model / 264
14.11	The continuum dielectric model / 264 Application to a charged spherical particle / 266

	14.13	Numerical treatment via finite differences / 268	
	14.14	Strengths and limitations of the continuum dielectric model / 269	
	14.15	Applications of the continuum dielectric model to macromolecules / 270	
		Notes / 273	
		Suggested reading / 275	
PAI	RT 4	CONFORMATIONAL STABILITY AND CONFORMATION CHANGE	277
15.	Some	Elements of Polymer Physics	279
	15.1	Introduction / 279	
	15.2	Conformational variation in small molecules / 280	
	15.3	Conformational variation in chain molecules / 280	
	15.4	The ideal random coil and the characteristic ratio / 281	
	15.5	The persistence length as a measure of chain flexibility / 282	
	15.6	Conformation of self-avoiding chains / 283	
	15.7	Dependence of chain conformation on solvent conditions; "Theta" conditions / 284	
	15.8	Relating chain statistics to molecular structure / 286	
	15.9	Polyelectrolytes / 287	
		Notes / 288	
		Suggested reading / 289	
16.	Helix-	Coil Equilibria	291
	16.1	Introduction: Multistate transitions of helical polymers / 291	
	16.2	Single-stranded poly (A): A completely non-cooperative transition / 291	
	16.3	Synthetic polypeptides / 292	
	16.4	Zimm-Bragg, Gibbs-DiMarzio, and Lifson-Roig analyses / 295 $$	
	16.5	Solution of the partition function / 297	
	16.6	Experiments on synthetic homo-polypeptides and protein fragments / 299	

17.

18.

16.7	Experimental determination of helix propensities in synthetic peptides / 299
16.8	Helix stabilization by salt bridges in oligomers containing Glu and Lys / 301
16.9	Helix stabilization by charged groups interacting with the helix dipole / 303
16.10	Helix-coil equilibria of nucleic acids / 303
16.11	Melting transition of DNA / 306
	Notes / 309
Protei	in Unfolding Equilibria 311
17.1	Introduction / 311
17.2	The two-state approximation / 312
17.3	Working with the two-state model / 314
17.4	Calorimetric measurements of the thermodynamics of protein unfolding / 316
17.5	Unfolding thermodynamics of ribonuclease / 318
17.6	Cold denaturation / 322
17.7	Solvent-induced unfolding: Guanidine hydrochloride and urea / 322
17.8	Mixed solvents: Denaturants and stabilizers / 324
17.9	Unfolding is not two-state under native conditions: Hydrogen exchange / 328
17.10	Nature of the two states / 332
17.11	A third state: The molten globule / 336
17.12	Range of stability / 338
17.13	Decomposition of the thermodynamic parameters for unfolding / 340
	Notes / 342
	Suggested reading / 345
Elasti	city of Biological Materials 347
18.1	Background / 347
18.2	Rubber-like elasticity of polymer networks / 348