

Fundamental Molecular Biology

基础分子生物学

影印版

Lizabeth A. Allison



高等教育出版社 Higher Education Press



教育部高等教育司推荐 国外优秀生命科学教学用书

The right of Lizabeth A. Allison to be identified as the Author of this Work has been asserted in accordance with the UK Copy-

First published 2007 by Blackwell Publishing Ltd

Fundamental rubished 2007 by Blackwell Publishing Ltd This edition is published by arrangement with Blackwell Publishin a Molecular Biology

甚础分子生物

影印版

Lizabeth A. Allison

苏工业学院图书馆

010 - 58581000蓝色畅想图书发行有限公司 刷。北京信成印刷有限公司

本书加有缺页。阅页。脱页等质量问题,请到所购图书销售部门联系漏秧。



物料号 25057-90

图字: 01-2008-2867号

reprinted from

Lizabeth A. Allison



Fundamental Molecular Biology, first edition

© 2007 Lizabeth A. Allison

The right of Lizabeth A. Allison to be identified as the Author of this Work has been asserted in accordance with the UK Copyright, Designs, and Patents Act 1988.

First published 2007 by Blackwell Publishing Ltd

This edition is published by arrangement with Blackwell Publishing Ltd, Oxford

图书在版编目(CIP)数据

基础分子生物学 = Fundamental Molecular Biology:

英文/(美)阿利森(Allison, L. A.)著.一影印本.

北京: 高等教育出版社, 2008.6

ISBN 978 -7 -04 -025057 -2

I. 基… Ⅱ. 阿… Ⅲ. 分子生物学 - 英文 Ⅳ. Q7

中国版本图书馆 CIP 数据核字(2008)第 091178 号

策划编辑 赵晓媛 责任编辑 赵晓媛 封面设计 责任印制 尤

高等教育出版社 购书热线 010-58581118 出版发行 北京市西城区德外大街 4号 800 - 810 - 0598 免费咨询 100120 http://www.hep.edu.cn 邮政编码 http://www.hep.com.cn 机 010-58581000 网上订购 http://www.landraco.com http://www.landraco.com.cn 蓝色畅想图书发行有限公司 经 http://www.widedu.com 北京铭成印刷有限公司 畅想教育 印 次 2008年6月第1版 889 × 1194 1/16 版 46.75 即 2008年6月第1次印刷 印 张 69.00元(含光盘) 950 000 定

本书如有缺页、倒页、脱页等质量问题, 请到所购图书销售部门联系调换。

版权所有 侵权必究

物料号 25057-00

Preface

Chapter Summary
Analytical questions
Suggestions for juriner reading

The fast pace of modern molecular biology research is driven by intellectual curiosity and major challenges in medicine, agriculture, and industry. No discipline in biology has ever experienced the explosion in growth and popularity that molecular biology is now undergoing. There is intense public interest in the Human Genome Project and genetic engineering, due in part to fascination with how our own genes influence our lives. With this fast pace of discovery, it has been difficult to find a suitable, up-to-date textbook for a course in molecular biology. Other textbooks in the field fall into two categories: they are either too advanced, comprehensive, and overwhelmingly detailed, with enough material to fill an entire year or more of lectures, or they are too basic, superficial, and less experimental in their approach. It is possible to piece together literature for a molecular biology course by assigning readings from a variety of sources. However, some students are poorly prepared to learn material strictly from lectures and selected readings in texts and the primary literature that do not match exactly the content of the course. At the other end, instructors may find it difficult to decide what topics are the most important to include in a course and what to exclude when presented with an extensive array of choices. This textbook aims to fill this perceived gap in the market. The intent is to keep the text to a manageable size while covering the essentials of molecular biology. Selection of topics to include or omit reflects my view of molecular biology and it is possible that some particular favorite topic may not be covered to the desired extent. Students often complain when an instructor teaches "straight from the textbook," so adding favorite examples is encouraged to allow instructors to enrich their course by bringing to it their own enthusiasm and insight.

Approach

A central theme of the textbook is the continuum of biological understanding, starting with basic properties of genes and genomes, RNA and protein structure and function, and extending to the complex, hierarchical interactions fundamental to living organisms. A comprehensive picture of the many ways molecular biology is being applied to the analysis of complex systems is developed, including advances that reveal fundamental features of gene regulation during cell growth and differentiation, and in response to a changing nvironment, as well as developments that are more related to commercial and medical applications. Recent advances in technology, the process and thrill of discovery, and ethical considerations in molecular biology research are emphasized.

The text highlights the process of discovery – the observations, the questions, the experimental designs to test models, the results and conclusions – not just presenting the "facts." At the same time the language of molecular biology is emphasized, and a foundation is built that is based in fact. It is not feasible to examine every brick in the foundation and still have time to view the entire structure. However, as often as possible real examples of data are shown, e.g. actual results of an EMSA, Western blot, or RNA splicing assay. Experiments are selected either because they are classics in the field or because they illustrate a particular approach frequently used by molecular biologists to answer a diversity of questions.

Organization

The textbook is designed for a one-term course on molecular biology (or molecular genetics) for undergraduate students who are primarily majoring in biology or chemistry, with a large percentage of premedical students. First-year graduate students with a minimal background in molecular genetics/biology would also benefit from this course. Students would be expected to have completed, at a minimum, a two-term introductory biology course and to have completed at least 1 year of chemistry. Each chapter opens with a conceptual statement and historical perspective, followed by explanation and elaboration. Chapters end with a list of key references from the primary literature, and a series of analytical questions.

The book begins with a five chapter sequence that should be a review for most students, but with more detail than they would have encountered in an introductory biology or genetics course. Students of molecular biology need to have a solid grasp of these concepts so they may need to refresh their understanding of them. Depending on the curriculum at a particular institution, more or less time may need to be spent on these introductory chapters. Chapter 1 is a brief history of genetics and the beginnings of molecular biology. Chapter 2 discusses the structure and chemical properties of DNA. Chapter 3 discusses the organization of genomes and eukaryotic chromatin. Chapter 4 deals with the versatility of RNA structure and function, and Chapter 5 provides an overview of the flow of genetic information from DNA to RNA to protein and covers basic protein structure and function. The genetic code for amino acids is presented along with protein structure and function. This order reflects, the view that to understand protein structure and function it is essential to first understand the flow of genetic information and where the primary sequence of amino acids derives from, and the consequences of alterations in the genetic code.

Chapters 6 and 7 cover DNA replication, telomere maintenance, and DNA repair and recombination. Although some instructors prefer to cover DNA replication later in a course, my view is that the information is essential early on, particularly to be able to understand many of the experimental strategies used in studying genes and their activities at the molecular level.

There is always debate on where to place methods and techniques – scattered throughout, in an appendix, or as specific chapters. I have mainly taken the latter approach, with the intent that this textbook will be a useful resource for students well beyond the course. Many undergraduate programs now include a research component and having a compilation of the standard techniques in molecular biology along with theoretical background and how they arose from basic research provides an essential aid. My approach in teaching is to cover some of the very basic methods in a series of "recombinant DNA technology" lectures, but to introduce others as needed to understand experiments discussed throughout the course. For example, Chapters 8 and 9, which cover recombinant DNA technology, molecular cloning, and tools for analyzing gene expression would certainly not be covered from start to finish. Covering method after method would become tedious. In addition, appreciation of the concepts behind techniques is much greater after students have acquired more experience in molecular biology.

Eukaryotic molecular biology is emphasized, although where details are better understood from bacteria, these are included. When fundamental processes such as DNA replication, repair, and recombination are discussed, the focus is on eukaryotes because the basic process is similar to that in prokaryotes, although the components of the machinery and the specific names of the players may differ. Prokaryotic transcription is given a separate chapter (Chapter 10), however, since some aspects of transcriptional regulation are fundamentally different than in eukaryotes, e.g. the concept of the operon, attenuation, and riboswitches. The basic transcription apparatus is introduced in Chapter 10 and how transcripts are initiated, elongated, and terminated is covered.

Chapter 11 covers the control of transcription in eukaryotes, introducing the regulatory elements, part the general transcription factors, the interaction of DNA-binding proteins and DNA targets, the role of coactivators and corepressors, and regulated nuclear import of transcription factors. Chapter 12 covers the emerging field of epigenetics and monoallelic gene expression.

Chapter 13 introduces RNA processing and post-transcriptional gene regulation in eukaryotes, while 140 can Chapter 14 covers the mechanism of translation, with a focus on eukaryotic translation.

Chapters 15–17 cover some of the many applications of molecular biology. Chapter 15 introduces genetically modified organisms and their use in basic and applied research. Chapter 16 covers genome analysis, including DNA typing, genomics, and proteomics. Chapter 17 covers aspects of medical molecular biology including the molecular biology of cancer, gene therapy, and human behavior.

The course length is easily adjustable. The book is designed so that more or less time can be spent on particular topics according to an instructor's preference. The material in boxes can be treated as supplementary material if the course is too long for the needs of a particular class. On the other hand, there will be additional readings in these sections for students who want to go beyond the material in the main text to gain a deeper understanding of a particular topic.

Special features

Unique aspects of the book include a cohesive discussion of epigenetics and medical molecular biology, and the use of boxes to highlight molecular tools (Tool boxes), and to provide a more detailed treatment of material that will be of interest to the very keen student (Focus boxes). In addition, the textbook has a strong emphasis on biomedical research, which will appeal to the many premedical students who are likely to take the course prior to taking the MCATs. "Disease boxes" use diseases resulting from defects in a key gene to illustrate many principles of molecular biology. These examples place complex regulatory pathways, such as nucleotide excision repair, in a relevant context, making them more memorable for students.

• Book features:

- Tool boxes explore key experimental methods and techniques in molecular biology
- Focus boxes offer more detailed treatment of topics, delve into experimental strategies, and suggest areas for further exploration
- Disease boxes illustrate key principles of molecular biology by examining diseases that result from key gene defects
- Chapter-opening quotes, outlines, and introductions and the solutions tiend bine solutions
- End-of-chapter analytical questions
- End-of-book glossary.
- Interactive website features:
 - Interactive animations (based on art from the book and identified in the book with a special icon)
 - Interactive student tutorials and pdb files
 - Interactive student exercises
 - Answers to end-of-chapter analytical questions
 - Additional student and instructor resources cloudes DNA technology and the student and instructor resources
 - Downloadable artwork from the text. The most part to be a solution of the property of the
- CD-ROM features:
 - Downloadable artwork from the text
 - Sample interactive animation and tutorial who demonstrated although the Sample interactive animation and tutorial who demonstrated and tutorial who demonstrated animation and the demonstrated an
 - Sample syllabus
 - Link to website.

Acknowledgments

I am forever indebted to my undergraduate mentor L. Gerard Swartz, my master's thesis advisor, Gerald Shields, my PhD thesis advisor, Aimee Bakken, and my faculty mentors, Frank Sin and Larry Wiseman for their inspiration and belief in me throughout my education and career. My husband, Michael Levine and my son Andrew (born May 2003) deserve special thanks for their patience and encouragment ("good job Mommy!"). I thank my parents for nurturing my creativity and teaching me to follow my dreams. This book is dedicated to my mother Marjorie Allison (1929–1999) who remembered I was a molecular biologist as opposed to a microbiologist by thinking of the moles in her flower garden, and to my father Jack Allison (1918–2004) who sparked my interest in science while allowing me to wash beakers and flasks in his high school chemistry lab during the summer. At Blackwell Publishing, Nancy Whilton was my visionary cheerleader and Elizabeth Frank dealt very efficiently with all the nuts and bolts of the process. Rosie Hayden capably managed all the behind-the-scenes editorial work, Sarah Edwards heroically orchestrated the design and production aspects, and Jane Andrew skillfully handled the copy editing with an excellent eye for detail. In addition, Kieran Thomas designed a creative and easy-to-use website and Matt Payne spearheaded the marketing and publicity efforts. I also thank the members of the Allison lab who have put up with me sequestering myself in my office for many months at a time.

discussed, the focus is on entaryotes because the basic process is

Finally, I acknowledge the contributions of my outside reviewers: Brian Ashburner (University of Toledo), Alice Cheung (University of Massachusetts, Amherst), Robert S. Dotson (Tulane University), Jutta Heller

(Loyola University), Daniel Herman (University of Wisconsin–Eau Claire), Jerry Honts (Drake University), Jason Kahn (University of Maryland), Chentao Lin (University of California, Los Angeles), Alison Liu (Rutgers, The State University of New Jersey), Hao Nguyen (California State University, Sacramento), Rekha C. Patel (University of South Carolina), Ravinder Singh (University of Colorado), and Scott A. Strobel (Yale University).

I appreciate greatly the time spent by these reviewers and thank them for their insightful and exceptionally helpful comments, most of which I hope I have addressed. Any remaining errors are mine and I welcome comments and suggestions for improvement.

Lizabeth A. Allison Williamsburg, VA 2006

Contents

Preface, xviii

1	The	beginnings	of mole	cular	biology, 1
---	-----	------------	---------	-------	------------

- 1.1 Introduction
- 1.2 Historical perspective

Insights into heredity from round and wrinkled peas: Mendelian genetics Insights into the nature of hereditary material: the transforming principle is DNA Creativity in approach leads to the one gene-one enzyme hypothesis The importance of technological advances: the Hershey-Chase experiment

A model for the structure of DNA: the DNA double helix

Chapter summary Analytical questions

Suggestions for further reading

2 The structure of DNA, 13

- 2.1 Introduction
- 2.2 Primary structure: the components of nucleic acids

Five-carbon sugars

Nitrogenous bases

The phosphate functional group Nucleosides and nucleotides

- 2.3 Significance of 5' and 3'
- 2.4 Nomenclature of nucleotides
- 2.5 The length of RNA and DNA
- 2.6 Secondary structure of DNA

Hydrogen bonds form between the bases

Base stacking provides chemical stability to the DNA double helix

Structure of the Watson-Crick DNA double helix

Distinguishing between features of alternative double-helical structures

DNA can undergo reversible strand separation

2.7 Unusual DNA secondary structures

Slipped structures

Cruciform structures

Triple helix DNA

Disease box 2.1 Friedreich's ataxia and triple helix DNA 4.5. RNA is involved in a wide range of cellular processes

2.8 Tertiary structure of DNA

Supercoiling of DNA

Topoisomerases relax supercoiled DNA

What is the significance of supercoiling in vivo?

Disease box 2.2 Topoisomerase-targeted anticancer drugs

Chapter summary

Analytical questions

Suggestions for further reading

3 Genome organization: from nucleotides to chromatin, 37

- 3.1 Introduction
- 3.2 Eukaryotic genome

4.7 Ribozymes catalyze a variety of chemical reactions

Chromatin structure: historical perspective Histones Nucleosomes Beads-on-a-string: the 10 nm fiber The 30 nm fiber Loop domains Metaphase chromosomes Alternative chromatin structures 3.3 Bacterial genome 3.4 Plasmids 3.5 Bacteriophages and mammalian DNA viruses Bacteriophages Mammalian DNA viruses 3.6 Organelle genomes: chloroplasts and mitochondria Chloroplast DNA (cpDNA) Mitochondrial DNA (mtDNA) Disease box 3.1 Mitochondrial DNA and disease 3.7 RNA-based genomes Eukaryotic RNA viruses Retroviruses Viroids Other subviral pathogens Disease box 3.2 Avian flu Chapter summary **Analytical questions** Suggestions for further reading 4 The versatility of RNA, 54 4.1 Introduction 4.2 Secondary structure of RNA Secondary structure motifs in RNA particulative double-health and principle of the secondary structure motifs in RNA particular and principle of the secondary structure motifs in RNA particular and principle of the secondary structure motifs in RNA particular and particular a Base-paired RNA adopts an A-type double helix RNA helices often contain noncanonical base pairs 4.3 Tertiary structure of RNA tRNA structure: important insights into RNA structural motifs Common tertiary structure motifs in RNA 4.4 Kinetics of RNA folding 4.5 RNA is involved in a wide range of cellular processes 4.6 Historical perspective: the discovery of RNA catalysis Tetrahymena group I intron ribozyme RNase P ribozyme Focus box 4.1: The RNA world

Small ribozymes
Chapter summary
Analytical questions
Suggestions for further reading

Mode of ribozyme action

Large ribozymes

4.7 Ribozymes catalyze a variety of chemical reactions

5.1 Introduction 5.2 The central dogma 5.3 The genetic code Translating the genetic sode and general and gener
5.2 The central dogma 5.3 The genetic code Translating the genetic code The 21st and 22nd genetically encoded amino acids lown amage to primare 15 as the avoid modified nucleotides in decoding the primary and properties and acids lown amage to primare 15 as the avoid modified nucleotides in decoding the primary structure and properties are properties. 5.4 Protein structure Primary structure Secondary structure Guaternary structure Quaternary structure Guaternary structure Prediction of protein and protein activity by post-translational modifications are protein activity by post-translational modifications are protein assemblages 5.5 Protein function Enzymes are biological catalysts Regulation of protein activity by post-translational modifications are processed as a protein activity of protein activity model to acid activity of protein activity model to acid activity model to acid activity of protein activity model to acid activity model to acid activity of protein activity model to acid activity model to acid activity model to acid activity model to acid activity of protein activity model to acid activity model acid activity model to acid activity model to acid activity model to acid activity model acid acid activity model acid acid acid acid acid acid acid acid
Translating the genetic code Translating the genetic code The 21st and 22nd genetically encoded amino acids form senge to general activity and accessed implications of codon bias for molecular biologists 5.4 Protein structure Primary structure Secondary structure Secondary structure Guaternary structure Guaternary structure Guaternary structure Size and complexity of proteins Proteins contain multiple functional domains Prediction of protein structure Secondary structure From the complexity of proteins Proteins contain multiple functional domains Prediction of protein structure 5.5 Protein function Enzymes are biological catalysts Regulation of protein activity by post-translational modifications Regulation of protein activity by post-translational modifications Allosteric regulation of protein activity Cyclin-dependent kinase activation Macromolecular assemblages 5.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading Cyclin-dependent and telomere maintenance, 108 6 DNA replication and telomere maintenance, 108 engages and complexity of protein activity Analytical questions Suggestions for further reading Cyclin-dependent activity Cyclin-dependent complexity of protein degradation Cyclin-dependent c
Translating the genetic code The 21st and 22nd genetically encoded amino acids power among to primare and 5 arous according to the 21st and 22nd genetically encoded amino acids power among to primare and 5 arous according to the 21st and 22nd according to the 21st and
The 21st and 22nd genetically encoded amino acids for earling to prime and 2 and according to the prime and 2 and
Implications of codon bias for molecular biotogists AND alternation and codon bias for molecular biotogists AND alternation and codon bias for molecular biotogists Protein structure Primary structure Secondary structure Tertiary structure Quaternary structure Quaternary structure Size and complexity of proteins Proteins contain multiple functional domains Prediction of protein structure Enzymes are biological catalysts Regulation of protein activity by post-translational modifications Allosteric regulation of protein activity Cyclin-dependent kinase activation Macromolecular assemblages 5.6 Protein folding and misfolding Molecular-chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading Cyclin-dependent the reading industrial protein and protein and protein activation and protein degradation Suggestions for further reading industrial protein and protein and protein degradation Suggestions and telomere maintenance, 108 DNA replication and telomere maintenance, 108 AND protein folding the protein degradation and telomere maintenance, 108 AND protein folding the protein degradation and telomere maintenance, 108 AND protein folding the protein degradation and telomere maintenance, 108 AND protein folding the protein degradation and telomere maintenance, 108 AND protein folding the protein folding the protein folding the protein folding the prot
Implications of codon bias for molecular biologists 7.4 Protein structure Primary structure Secondary structure Countering structure Quaternary structure Quaternary structure Size and complexity of proteins Proteins contain multiple functional domains Prediction of protein structure Enzymes are biological catalysts Regulation of protein activity by post-translational modifications Allosteric regulation of protein activity Cyclin-dependent kinase activation Macromolecular assemblages 7.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading Cyclin-dependent and telomere maintenance, 108 AND Protein folding and telomere maintenance, 108 AND Protein folding and telomere maintenance, 108 AND Protein folding and telomere maintenance, 108 AND Protein architecture AND Protein folding and telomere maintenance, 108 AND Protein folding and telomere maintenance, 108
Protein structure Primary structure Secondary structure Secondary structure Secondary structure Secondary structure Guaternary structure Guaternary structure Guaternary structure Size and complexity of proteins Proteins contain multiple functional domains Prediction of protein structure Maintenance in telephone in the protein structure Protein function Enzymes are biological catalysts Regulation of protein activity by post-translational modifications Allosteric regulation of protein activity Cyclin-dependent kinase activation Macromolecular assemblages Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading middle family and the protein structure and telement and the protein summary and and telemere maintenance, 108 5.5 DNA replication and telemere maintenance, 108
Primary structure Secondary structure Tertiary structure Quaternary structure Quaternary structure Size and complexity of proteins Proteins contain multiple functional domains Prediction of protein structure Enzymes are biological catalysts Regulation of protein activity by post-translational modifications Allosteric regulation of protein activity Macromolecular assemblages 5.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading middlesses and summary appears to the summary and
Secondary structure Tertiary structure Quaternary structure Guaternary structure Size and complexity of proteins Proteins contain multiple functional domains Prediction of protein structure Enzymes are biological catalysts Regulation of protein activity by post-translational modifications Allosteric regulation of protein activity Cyclin-dependent kinase activation Macromolecular assemblages 5.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading
Tertiary structure Quaternary structure Size and complexity of proteins Proteins contain multiple functional domains Prediction of protein structure 5.5 Protein function Enzymes are biological catalysts Regulation of protein activity by post-translational modifications Allosteric regulation of protein activity Cyclin-dependent kinase activation Macromolecular assemblages 5.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading Cyclin-dependent experiments and telomere maintenance, 108 DNA replication and telomere maintenance, 108 Posterior dependent experiments and the protein degradation and telomere maintenance, 108 Posterior folding and misfolding and misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading diseases and protein degradation and telomere maintenance, 108
Quaternary structure Size and complexity of proteins Proteins contain multiple functional domains Prediction of protein structure 5.5 Protein function Enzymes are biological catalysts Regulation of protein activity by post-translational modifications Allosteric regulation of protein activity Cyclin-dependent kinase activation Macromolecular assemblages 5.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading Molecular chapero and telomere maintenance, 108 Analytical complexity of proteins and telomere maintenance, 108 mediator and telomere maintenance, 108 mediator and telomere maintenance, 108
Size and complexity of proteins Proteins contain multiple functional domains Prediction of protein structure 5.5 Protein function Enzymes are biological catalysts Regulation of protein activity by post-translational modifications, and a parameter and a
Proteins contain multiple functional domains Prediction of protein structure 5.5 Protein function Enzymes are biological catalysts Regulation of protein activity by post-translational modifications, and the protein activity of protein activity o
Prediction of protein structure 5.5 Protein function Enzymes are biological catalysts Regulation of protein activity by post-translational modifications Allosteric regulation of protein activity Cyclin-dependent kinase activation Macromolecular assemblages 5.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading DNA replication and telomere maintenance, 108 Enzymes are biological catalysts and telomere maintenance, 108 Protein function activity by post-translational modifications and telomere maintenance, 108 Allosteric regulation of protein activity by post-translational modifications and telomere maintenance, 108
Enzymes are biological catalysts Regulation of protein activity by post-translational modifications because as a proper activity of protein activi
Enzymes are biological catalysts Regulation of protein activity by post-translational modifications has been proposed as a proposed activity. Allosteric regulation of protein activity. Cyclin-dependent kinase activation Macromolecular assemblages 5.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading understanding and protein and telomere maintenance, 108 DNA replication and telomere maintenance, 108
Regulation of protein activity by post-translational modifications Allosteric regulation of protein activity Cyclin-dependent kinase activation Macromolecular assemblages 5.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading Chapter summary Chapter summary Analytical questions Suggestions for further reading Chapter summary Chapter summary Chapter summary Analytical questions Suggestions for further reading Chapter summary C
Allosteric regulation of protein activity Cyclin-dependent kinase activation Macromolecular assemblages 5.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading Suggestion and telomere maintenance, 108 Allosteric regulation of protein activity Analytical question of protein activity Cyclin-dependent kinase activation yrammus related anoiseural paralytical protein activity Analytical questions Suggestions for further reading Suggestion and telomere maintenance, 108 Analytication and telomere maintenance, 108
Macromolecular assemblages 5.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading industrial industrial and industrial
Macromolecular assemblages 5.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading mints am a protein and protein a
5.6 Protein folding and misfolding Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading indistance in the summary and th
Molecular chaperones Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading industrial industr
Ubiquitin-mediated protein degradation Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading plats and plats and plats are plated to be all plated to be all plats are plated to be all plated to be al
Protein misfolding diseases Disease box 5.1 Prions Chapter summary Analytical questions Suggestions for further reading Chapter reading Analytical questions Suggestions for further reading Chapter reading Chapter summary Analytical questions Suggestions for further reading Chapter reading Chapter summary Chapter s
Disease box 5.1 Prions Chapter summary Transitions and transversions can lead to silent, missense, or nonsense and transversions can lead to silent, missense, or nonsense and transversions can cause frameshiftemutations Suggestions for further reading applications to genetic instability and transverses of UNA replication and telomere maintenance, 108 DNA replication and telomere maintenance, 108
Chapter summary and transversions can lead to site in the sense of parameters and transversions of the sense frame of the sense of the sense frame of the sense o
Analytical questions Suggestions for further reading photosic particular and several several several photosic particular and p
Suggestions for further reading physical papers of abselvation and telegrated physical physic
6 DNA replication and telomere maintenance, 108
6 DNA replication and telomere maintenance, 108
0 1 1111 000001011
6.2 Historical perspective
Insight into the mode of DNA replication: the Meselson-Stahl experiment reports and the Meselson-Stahl experiment
Insight into the mode of DNA replication: visualization of replicating bacterial DNA values and A
6.3 DNA synthesis occurs from $5' \rightarrow 3'$
6.4 DNA polymerases are the enzymes that catalyze DNA synthesis and past elipnia to his gas?
Focus box 6.1 Bacterial DNA polymerases
6.5 Semidiscontinuous DNA replication
Leading strand synthesis is continuous risgan noise example.
Lagging strand synthesis is discontinuous appendix applications with the strand synthesis is discontinuous.
Lagging strand synthesis is discontinuous spendoo also plant of the send of th
6.6 Nuclear DNA replication in eukaryotic cells, Mg to Javomen yet nisqen Asend branza-elduod. 7
6.6 Nuclear DNA replication in eukaryotic cells. In the Javones and Javones Replication factories
6.6 Nuclear DNA replication in eukaryotic cells. In the Javomen video is a send for a se
6.6 Nuclear DNA replication in eukaryotic cells Mg to Javomen vd disgent send brante-educid. The Replication factories Histone removal at the origins of replication Prereplication complex formation at the origins of replication as a major of the second
6.6 Nuclear DNA replication in eukaryotic cells. In the Javomen yet insert branches are placed in the Proposition of the Proposition in Eukaryotic cells. In the Propo

Polymerase switching

Elongation of leading strands and lagging strands

Proofreading

Maturation of nascent DNA strands

Termination

Histone deposition

Focus box 6.2 The naming of genes involved in DNA replication the standard back breats and

Disease box 6.1 Systemic lupus erythematosus and PCNA strong asphrosinum

6.7 Replication of organelle DNA

Models for mtDNA replication

Replication of cpDNA

Disease box 6.2 RNase MRP and cartilage-hair hypoplasia

6.8 Rolling circle replication

6.9 Telomere maintenance: the role of telomerase in DNA replication, aging, and cancer

Telomeres

Solution to the end replication problem

Maintenance of telomeres by telomerase

Other modes of telomere maintenance

Regulation of telomerase activity

Telomerase, aging, and cancer

Disease box 6.3 Dyskeratosis congenita: loss of telomerase function to noissuper analaolia

Chapter summary

Analytical questions

Suggestions for further reading

7 DNA repair and recombination, 152

7.1 Introduction

7.2 Types of mutations and their phenotypic consequences

Transitions and transversions can lead to silent, missense, or nonsense mutations Insertions or deletions can cause frameshift mutations

Expansion of trinucleotide repeats leads to genetic instability

7.3 General classes of DNA damage

Single base changes

Structural distortion

DNA backbone damage

Insight into the mode of DNA replication: the Meselson-Stagement AND and the mode of DNA replication.

7.4 Lesion bypass limit the mode of BNA replication; visualization of replicating bacterial seasons

7.5 Direct reversal of DNA damage

7.6 Repair of single base changes and structural distortions by removal of DNA damage AMU AM

Base excision repair

Mismatch repair

Nucleotide excision repair

Disease box 7.1 Hereditary nonpolyposis colorectal cancer: a defect in mismatch repair

7.7 Double-strand break repair by removal of DNA damage (15) us ni noits iller AMD real publication and account of DNA damage (15) us ni noits iller AMD real publication and account of DNA damage (15) us ni noits iller amb (15) us ni noi

Homologous recombination

Nonhomologous end-joining

Disease box 7.2 Xeroderma pigmentosum and related disorders: defects in nucleotide and excision repair

Disease box 7.3 Hereditary breast cancer syndromes: mutations in *BRCA1* published salgud and *BRCA2*

Chapter summary
Analytical questions
Suggestions for further reading

8 Recombinant DNA technology and molecular cloning, 180

8.1 Introduction

8.2 Historical perspective

Insights from bacteriophage lambda (λ) cohesive sites senses allocations of languages and salvages are proposed and salv

8.3 Cutting and joining DNA

Major classes of restriction endonucleases

Restriction endonuclease nomenclature adjulum and Jacobian sequences for type III restriction endonucleases.

Recognition sequences for type II restriction endonucleases and to level entries in a local mention.

DNA ligase

Focus box 8.1 Fear of recombinant DNA molecules

8.4 Molecular cloning

Vector DNA

Choice of vector is dependent on insert size and application length to level entitle aleval entitle and application length to level entitle aleval entitle and application length to level entitle and application length to length to level entitle and application length to length to level entitle and application length and application length to le

Plasmid DNA as a vector

Bacteriophage lambda (λ) as a vector

Artificial chromosome vectors

Sources of DNA for cloning

Focus box 8.2 EcoRI: kinking and cutting DNA

Tool box 8.1 Liquid chromatography

8.5 Constructing DNA libraries

Genomic library cDNA library

8.6 Probes

Heterologous probes

Homologous probes

Tool box 8.2 Complementary DNA (cDNA) synthesis

Tool box 8.3 Polymerase chain reaction (PCR) and sensitive missing of property and the sensitive missing of the sensitive missing missing of the sensitive missing missing

Tool box 8.4 Radioactive and nonradioactive labeling methods

Tool box 8.5 Nucleic acid labeling

8.7 Library screening

Transfer of colonies to a DNA-binding membrane intensity grane spanness enough Colony hybridization

Detection of positive colonies

Wide publishers veries

8.8 Expression libraries

8.9 Restriction mapping

8.10 Restriction fragment length polymorphism (RFLP)

RFLPs can serve as markers of genetic diseases

Tool box 8.6 Electrophoresis non ecommoned possessided has esta

Tool box 8.7 Southern blot

Disease box 8.1 PCR-RFLP assay for maple syrup urine disease association and avid

8.11 DNA sequencing

Manual DNA sequencing by the Sanger "dideoxy" DNA method and sequencing Automated DNA sequencing

7	Tool	s for analyzing gene expression, 232					
	9.2	Transient and stable transfection assays					
	9.3	Reporter genes	2 Historical perspective				
		Commonly used reporter genes					
		Analysis of gene regulation amateva node					
		Purification and detection tags: fusion protei	. The first cloning experiments an				
		Tool box 9.1 Production of recombinant prote	3 Cutting and joining DNA				
	9.4	In vitro mutagenesis					
		Tool box 9.2 Fluorescence, confocal, and mu	Restriction endonucless exposoroim motorial				
	9.5	Analysis at the level of gene transcription:	RNA expression and localization				
		Northern blot					
		In situ hybridization					
		RNase protection assay (RPA)					
		Reverse transcription-PCR (RT-PCR)					
	9.6	Analysis at the level of translation: protein	expression and localization 1010 8V 10, 9010110				
		Western blot					
		<i>In situ</i> analysis					
		Enzyme-linked immunosorbent assay (ELISA					
		Tool box 9.3 Protein gel electrophoresis					
		Tool box 9.4 Antibody production					
	9.7	Antisense technology					
		Antisense oligonucleotides	Constructing DNA libraries				
		RNA interference (RNAi)					
	9.8	Analysis of DNA-protein interactions					
		Electrophoretic mobility shift assay (EMSA)					
		DNase I footprinting					
		Chromatin immunoprecipitation (ChIP) assa	y and an application of the second supplication				
			Tool box 8.2 Complementary DNA [cDNA] synth				
	9.9		Tool box 8.3 Polymerase chain reaction IPCRI.				
		Yeast two-hybrid assay	Tool box 8.5 Nucleic acid labeling				
		Coimmunoprecipitation assay	Library screening				
			Transfer of colonies to a DNA-binding mer(TBR				
	9.10	Structural analysis of proteins					
		X-ray crystallography					
		Nuclear magnetic resonance (NMR) spectro					
		Cryoelectron microscopy	Restriction mapping				

Atomic force microscopy (AFM) 9.11 **Model organisms**

Yeast: Saccharomyces cerevisiae and Schizosaccharomyces pombe anadomosia & Rod IcoT

Worm: Caenorhabditis elegans

Fly: Drosophila melanogaster seed annu quays elgem not yesse 9199-939 1.8 xod esseed

Fish: Danio rerio

Manual DNA sequencing by the Sanger "dideoxy" DNA method and size objects and size of the sequencing by the Sanger "dideoxy" DNA method

Mouse: Mus musculus

		Suggestions for further reading		
10		scription in prokaryotes, 278		
			Focus box 11.4 Homeoboxes and homeodomains	
			Disease box 11.2 Greig cephalopolarinatad ni ba	
	10.3	Mechanism of transcription dust a season	Disease box 11,3 Defective historie acetyltransfe	
		Bacterial promoter structure	Transcriptional coactivators and corepressors	
		Structure of bacterial RNA polymerase		
		Stages of transcription		
		Proofreading		
		Direction of transcription around the E. c	Focus box 11.5 is there a histon emozomorno ilo	
		Focus box 10.1 Which moves - the RNA	polymerase or the DNA? xelgmos notiginarisiT	
	10.4	Historical perspective: the Jacob-Mono	d operon model of gene regulation and to help 0	
		The operon model led to the discovery of	Enhanceosome model ANRm	
		Characterization of the Lac repressor		
		Lactose (lac) operon regulation		
		Lac operon induction	Mechanism of RNA polymerase II transcription	
		Basal transcription of the <i>lac</i> operon		
		Regulation of the <i>lac</i> operon by Rho		
			ne are widely used in molecular biology research	
			Transcription elongation through the nuclecarot	
			Disease box 11.4 Defects in Etongator and famili	
		Allosteric modifications and DNA binding		
		DNA looping	Karyophenns	
	10.7	Control of gene expression by RNA	Nuclear localization sequences (NLSs)	
	10.7		al attenuation of the tryptophan operon	
9.		Riboswitches	Nuclear import pathway	
		Riboswitch ribozymes		
		Chapter summary		
			Regulated nuclear import and signal transducti	
		Suggestions for further reading they had	Regulated nuclear import and signal transduction	
11	Tran	scription in eukarvotes. 312		
	11.1	Introduction	Chapter summary	
		Overview of transcriptional regulation		
	11.3	Protein-coding gene regulatory eleme		
	11.0	Structure and function of promoter eler		
			egulatory elements (99 oil oil sonom bas eoil and	
		Focus box 11.1 Position effect and long		
			and DNase I hypersensitive sites 25 m and 25 m a	
			Cytosine DNA methylation marks genes for siles?x	
			s and transcription factories example maintains.	
	11 /			
	11.4		Disease box 12.1 Cancer and epigenetics	
		Components of the general transcription		
		Structure of RNA polymerase II	Establishing and maintaining the imprint	
			itiation complex formation la	
		Mediator: a molecular bridge	Genomic imprinting is essential for normal develo	

Frog: Xenopus laevis and Xenopus tropicalis

Chapter summary notifectors tendification to the summary notifectors are the summary notifectors.

Analytical questions

12 Epigenetics and monoallelic gene expression, 392 mast-paol to modernal bas enulsural

Suggestions for further reading

12.1 Introduction

12.2 Epigenetic markers a will see and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNase I inverse said by a Historical trades and DNASE I inverse said by a Historical tr

Analytical questions

Cytosine DNA methylation marks genes for silencing management at 2011 xod auto-3
Stable maintenance of histone modifications are applicable for silencing management for si

12.3 Genomic imprinting

Structure of RNA polymerasa II

General transcription factors and preintiation composition factors and preintiation commission for more development.

Mediators a molecular bridge

Mediators a molecular bridge

Origins of genomic imprinting

Disease box 12.3 Genomic imprinting and neurodevelopmental disorders

12.4 X chromosome inactivation

Random X chromosome inactivation in mammals

Molecular mechanisms for stable maintenance of X chromosome inactivation and AMRI Is there monoallelic expression of all X-linked genes?

12.5 Phenotypic consequences of transposable elements

Historical perspective: Barbara McClintock's discovery of mobile genetic elements in maize

DNA transposons have a wide host range

DNA transposons move by a "cut and paste" mechanism of particular and paste and paste

Retrotransposons move by a "copy and paste" mechanism and behing rolls all born see 8 9 8 1

Some LTR retrotransposons are active in the mammalian genome a langitude and the state of the st

Non-LTR retrotransposons include LINEs and SINEs viewozaib ent evitagearea Isaliotzi H

Tool box 12.1 Transposon tagging

Disease box 12.4 Jumping genes and human disease bot degradations and human disease box 12.4 Jumping genes and human di

12.6 Epigenetic control of transposable elements as godyo bas augloun ent ni revonaut AMR 41.87

Methylation of transposable elements

Heterochromatin formation mediated by RNAi and RNA-directed DNA methylation willoud

12.7 Allelic exclusion

Yeast mating-type switching and silencing

Antigen switching in trypanosomes

V(D)J recombination and the adaptive immune response public resulting and the adaptive immune response respon

Disease box 12.5 Trypanosomiasis: human "sleeping sickness"

Focus box 12.1 Did the V(D)J system evolve from a transposon?

Chapter summary

Analytical questions

Suggestions for further reading

13 RNA processing and post-transcriptional gene regulation, 452

- 13.1 Introduction
- 13.2 RNA splicing: historical perspective and overview
- 13.3 Group I and group II self-splicing introns

Group I introns require an external G cofactor for splicing sonims to vividas pribas floorig

Group II introns require an internal bulged A for splicing

Ternary complex formation and loading onto the 40s ribosomerorini II bns I quorp

Focus box 13.1 Intron-encoded small nucleolar RNA and "inside-out" genes and publications

13.4 Archael and nuclear transfer RNA introns

Archael introns are spliced by an endoribonuclease Jamesodn 200 bns 204 and 16 gniniol. Some nuclear tRNA genes contain an intron eyeses polynoged neighbors I LAT xod loo T

13.5 Cotranscriptional processing of nuclear pre-mRNA outside additional processing outside additional pro

Addition of the 5'-7-methylguanosine cap

Termination and polyadenylation

Splicing

Disease box 13.1 Oculopharyngeal muscular dystrophy: trinucleotide repeat expansion in a poly(A)-binding protein gene

Disease box 13.2 Spinal muscular atrophy: defects in snRNP biogenesis

Disease box 13.3 Prp8 gene mutations cause retinitis pigmentosa Jeog bos Jenoitalans T VAL

13.6 Alternative splicing

Effects of alternative splicing on gene expression of your amount of the splicing on gene expression of the splicing on gene expression of the splicing of the splicing on gene expression of the splicing of