## SKCOND EDITION

# Modern Genetics

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## SECOND EDITION

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Francisco J. Ayala John A. Kiger, Jr.

University of California, Davis



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# Modern Genetics



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# Preface

Genetics is a rapidly advancing science. With these words we opened the preface to the first edition of *Modern Genetics*. The continuing truth of these words, along with the valuable comments of many instructors of genetics who have used the first edition, has necessitated this major revision only four years after the appearance of the first edition.

Modern Genetics is a basic genetics text intended for a one-semester course. Prior college courses in general biology and chemistry are assumed, although some essential concepts (for example, mitosis and meiosis) that the student should have learned in such courses are reviewed here. We have intentionally included more material than can be covered in most one-semester courses. However, the book is written so as to enable instructors to choose topics according to their preferences.

The molecular basis of heredity remains a dominant theme of modern genetics and of this revised text. Recombinant DNA technology has made great advances in the last four years and opened wide the door to a molecular understanding of the genetics of humans and other higher organisms that was only anticipated previously. But in-depth coverage of population genetics and evolutionary genetics remains a distinctive feature of *Modern Genetics*.

As with the first edition, we have exploited our complementary areas of expertise in writing *Modern Genetics*. F.J.A. wrote the first draft of most or all of Chapters 1–3, 10, 18, 19 and 21–26 and Appendix 1; J.A.K. wrote the draft of Chapters 4–9 and 11–17. Chapter 20 was a joint effort. The flip of a coin determined the order of authorship.

## Organization

The basic organization of this edition is unchanged from that of the first in spite of the rearrangement of some topics and addition of entirely new chapters on recombinant DNA technology, eukaryotic gene regulation, and somatic cell genetics. *Modern Genetics* is composed of three parts: Part I, Organization and Transmission of the Genetic Materials; Part II, Expression of the Genetic Materials; and Part III, Evolution of the Genetic Materials. We remain convinced that this format is both logical and of practical value in the classroom. It should aid the student in understanding why the various topics are considered and how they contribute to a full comprehension of the processes of heredity. Yet the text is designed to give the instructor maximum flexibility in devising a syllabus. The order in which chapters are taught can be substantially altered without creating serious gaps in knowledge. We believe, however, that each chapter forms a logical unit and may best be taught as such.

The major changes in organization and approach from the first edition begin in Part I. The section on chromosome mapping in eukaryotes (Chapter 5) has been rewritten and expanded based upon comments from instructors and our own students. Complementation analysis and fine-structure mapping in both prokaryotes and eukaryotes form a new Chapter 6. The previous Chapter 9 (DNA Replication, Repair, and Recombination) has become Chapters 13 and 14, which have been moved to Part II, because of the emphasis on the genetic functions that are involved in the processes of DNA replication and recombination. A new Chapter 9 (DNA Manipulation) concludes Part I since the tools of recombinant DNA technology and DNA sequence analysis are properly a part of a section on "Organization and Transmission of the Genetic Materials." The previous Chapters 6 and 7 have been updated and become Chapters 7 and 8. Much of the content of the previous Chapter 8 can now be found in Chapters 6 and 14.

In Part II, an entirely new Chapter 16, Regulation of Gene Expression in Eukaryotes, appears. It is based largely on the fruits of recombinant DNA research. The material in the previous chapter by this name has also been updated and is found in the new Chapter 17, Genetic Analysis of Development. A new Chapter 18, Genetics with Somatic Cells, where we have included the recent spectacular advances in mapping the human genome, has been added. Chapters 11 and 12 have been updated to incorporate our deepened understanding of an evolving genetic code and information flow in cells; the incorporation of Chapters 13 and 14 has been mentioned previously.

In Part III, we have enlarged in Chapter 21 the section on duplications because of the increased understanding of this topic provided by the availability of DNA sequences, and we have added a section on the evolution of the human chromosomes. In Chapter 22 the discussion of gene variation incorporates the results from the studies of cryptic protein variation and of nucleotide heterogeneity. The topic of linkage disequilibrium reviewed in Chapter 25 receives new light from examination of long DNA sequences.

\* Chapter 26 has new sections on the reconstruction of phylogeny from DNA sequences, on the evolution of the genome by gene elongation, adjoining, and duplication, and on the intriguing question of horizontal gene transfer between species.

## Special Features

 This edition, like the previous one, is generously illustrated to enhance the clarity of the text.

Special topics of various kinds are set apart as "boxes," most of which can
be skipped without loss of comprehension of the text. These boxes provide
an additional element of choice in selecting material for study.

 Important terms and concepts are now listed at the end of each chapter as an aid for review. A glossary is provided at the end of the book for easy

reference.

• The problems at the ends of the chapters are an integral part of the book; some contain new information that we consider subsidiary or too detailed to be included in the text proper. At the request of many instructors and students, we have included simple problems within each chapter in addition to more difficult ones, and have added the answers to the odd-numbered problems in Appendix 2 at the back of the book. (The answers to the even-numbered problems are available to instructors in a separate supplement.)

 Students who have not taken a course in statistics will find in Appendix 1 the concepts and methods needed to understand the text and solve the

problems.

 The bibliography at the end of each chapter lists many basic references that document the material covered in the text; the credit lines for figures and tables cite additional sources of information.

## Acknowledgments

Our most fundamental debt is to all the scientists whose brilliant efforts have made genetics such an exciting science. To many of them we are particularly thankful for their generosity in providing photographs, permission to reproduce illustrations, and other needed items. The manuscript of the first edition was extensively reviewed by both specialists and generalists to ensure a sound and balanced coverage of the subject. The new material of this second edition has also been extensively reviewed. Although we alone bear the responsibility for any shortcomings, they are all the fewer because of these helpful reviews.

We thank Crystal DiModica and Candy Miller, who typed the manuscript, and Lorraine Barr who helped with proofreading. Valuable assistance in preparing this edition was also provided by Dr. Helen K. Salz, Boni Gregory, and Elizabeth Harper. Lieselotte Hofmann has been a skilled editor of the manuscript; we acknowledge the lasting contribution of Fred Raab as

production editor of the first edition. The book has greatly benefited from the skill of Georg Klatt who did most of the drawings. We gratefully acknowledge the continued dedication of the Benjamin/Cummings staff, in particular Jim Behnke, Jane Gillen, and Pat Waldo. Working with a publisher is always a challenging experience; in this case, it was also a pleasure.

Davis, California January, 1984 Francisco J. Ayala John A. Kiger, Jr.

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