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# Announcing New Clinical Genetics

By Andrew Read and Dian Donnai, *University of Manchester, UK*

## “Conceptually brilliant!”

David N. Cooper, *Professor of Human Molecular Genetics, Cardiff University, UK*

Written for students, genetic counselors and clinical geneticists, *New Clinical Genetics* provides the reader with a concise summary of post-genomic human genetics and guidance as to how our current understanding can be utilized in clinical practice.

The authors have taken an integrated case-based approach to the subject. Realistic case scenarios are used throughout the text so that readers can gain a practical understanding of clinical cases, their diagnoses and treatment options.

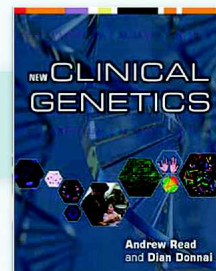
The book is based on the requirements of the ASHG Medical School Core Curriculum which states that “every physician who practices in the 21st century must have an in-depth knowledge of the principles of human genetics and their application to a wide variety of clinical problems.”

For details of how the book covers the core curriculum, please visit [www.scionpublishing.com/newclinicalgenetics](http://www.scionpublishing.com/newclinicalgenetics). With self-assessment questions, separate boxes containing diagnostic methods and approaches to common diseases, and over 300 clinical photograph and figures, the book is presented in a user-friendly manner to aid understanding.

2007, 428 pp., full-color throughout

Paperback \$54.95

ISBN: 9781904842316



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## About the authors

**Andrew Read** (PhD, FRCPath, FMedSci) is Professor of Human Genetics at the University of Manchester. He is co-author of Strachan and Read's *Human Molecular Genetics*, the leading advanced textbook on the subject. He established one of the first molecular genetic diagnostic services in the UK and has a long-standing interest in what genetic testing can and should achieve.

**Dian Donnai** (FRCP, FRCPCH, FMedSci) is Professor of Medical Genetics at the University of Manchester and Consultant Clinical Geneticist in the NW Regional Genetics Service. She is one of the world's leading authorities on clinical genetics, especially the study of dysmorphic syndromes, and is an advisor to many national and international bodies on the clinical application of genetics.

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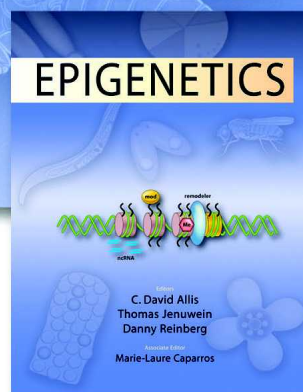
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# Epigenetics



Edited By C. David Allis, *The Rockefeller University, New York*; Thomas Jenuwein, *Research Institute of Molecular Pathology, Vienna*; Danny Reinberg, *Howard Hughes Medical Institute/University of Medicine and Dentistry of New Jersey–Robert Wood Johnson Medical School*; Marie-Laure Caparros, *Associate Editor*

The regulation of gene expression in many biological processes involves epigenetic mechanisms. In this new volume, 24 chapters written by experts in the field discuss epigenetic effects from many perspectives. There are chapters on the basic molecular mechanisms underpinning epigenetic regulation, discussion of cellular processes that rely on this kind of regulation, and surveys of organisms in which it has been most studied. Thus, there are chapters on histone and DNA methylation, siRNAs and gene silencing; X-chromosome inactivation, dosage compensation and imprinting; and discussion of epigenetics in microbes, plants, insects, and mammals. The last part of the book looks at how epigenetic mechanisms act in cell division and differentiation, and how errors in these pathways contribute to cancer and other human diseases. Also discussed are consequences of epigenetics in attempts to clone animals. This book is a major resource for those working in the field, as well as being a suitable text for advanced undergraduate and graduate courses on gene regulation.

2007, 502 pp., illus., appendices, index  
Hardcover \$158

ISBN 978-087969724-2

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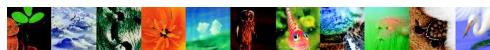
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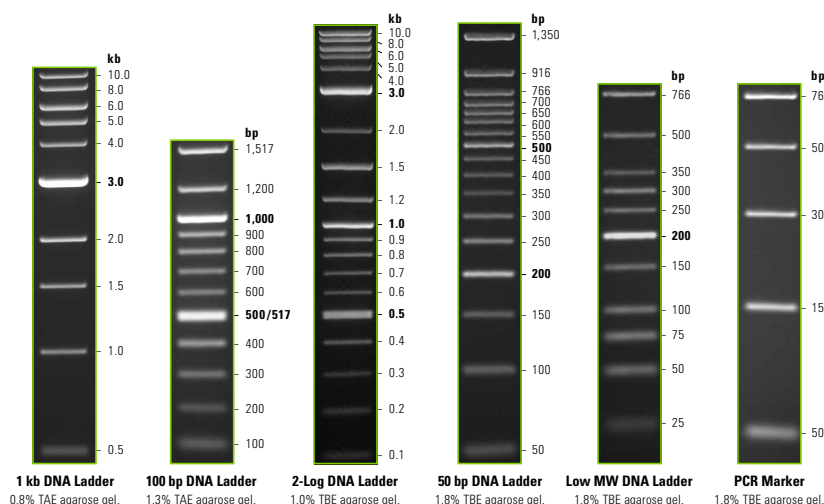
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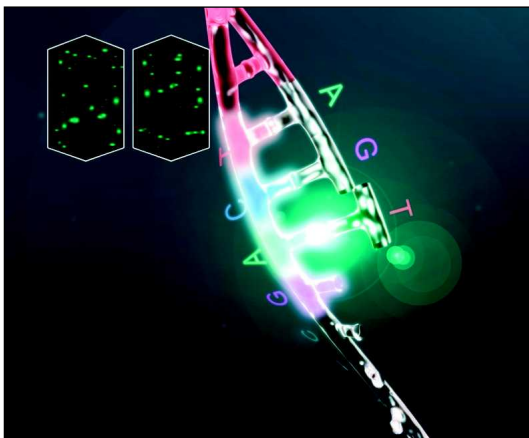
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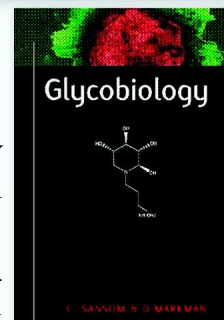
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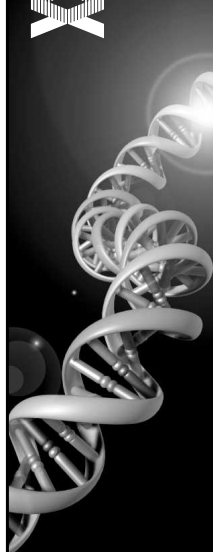
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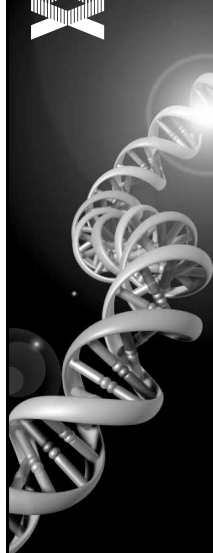
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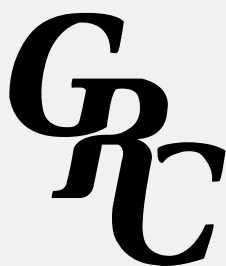
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The goal of this conference is to bring together representatives from this diversity of research areas to examine the most recent findings on the molecular bases of evolution. Molecular evolution underlies much of comparative genomics, informs models of computational biology, and provides a framework for understanding evolution of development and complex systems. The small size of the conference (restricted to 130 participants) and the location (on the beach in sunny Ventura, CA), coupled with the large blocks of time set aside for informal discussion are all critical features to the ongoing success of this conference. The following sessions and speakers have been confirmed:

### Exploring Adaptive Landscapes

Ben Kerr (University of Washington)  
Tony Dean (University of Minnesota)  
Dan Weinreich (Brown University)

### Measuring Evolutionary Timescales with classes of molecular markers

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### Molecular Basis of Heart Evolution

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Jose Xavier-Neto (Universidade de São Paulo)  
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### Positive and Negative Selection on Noncoding DNA

Peter Andolfatto (Univ. of California at San Diego)  
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### Evolvability

Günter Wagner (Yale University)  
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### Molecular Evolution of Body Axes

Mike Levine (University of California at Berkeley)  
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Mark Martindale (University of Hawaii)  
John Gerhart (University of California at Berkeley)

### Computational and Statistical Advances

Sudir Kumar (Arizona State University)  
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