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Tom Strachan is Emeritus Professor of Human Molecular Genetics at Newcastle University, Newcastle, UK, and is a Fellow of the Royal Society of Edinburgh and a Fellow of the Academy of Medical Sciences. He was the founding Head of Institute at Newcastle University's Institute of Human Genetics (now the Institute of Genetic Medicine) and its Scientific Director from 2001 to 2009.

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Tom Strachan is Scientific Director of the Institute of Human Genetics and Professor of Human Molecular Genetics at Newcastle University, UK, and is a Fellow of the Academy of Medical Sciences and a Fellow of the Royal Society of Edinburgh. Tom's early research interests were in multigene family evolution and interlocus sequence exchange, notably in the HLA and 21-hydroxylase gene clusters.

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Rev. ed. of: Human molecular genetics 3 / Tom Strachan and Andrew Read. 3rd ed. c2004.

Human Molecular Genetics is an established and class-proven textbook for upper-level undergraduates and graduate students which provides an authoritative and integrated approach to the molecular aspects of human genetics. While maintaining the hallmark features of previous editions, the Fourth Edition has been completely updated. It includes new Key Concepts at the beginning of each chapter and annotated further reading at the conclusion of each chapter, to help readers navigate the wealth of information in this subject. The text has been restructured so genomic technologies are integrated throughout, and next generation sequencing is included. Genetic testing, screening, approaches to therapy, personalized medicine, and disease models have been brought together in one section. Coverage of cell biology including stem cells and cell therapy, studying gene function and structure, comparative genomics, model organisms, noncoding

RNAs and their functions, and epigenetics have all been expanded.

Professors Tom Strachan & Andrew Read awarded the Education Award 2007 of the ESHG for their outstanding contribution to the dispersal of knowledge of modern human molecular genetics among students and professionals. Following the completion of the Human Genome Project the content and organization of the third edition of Human Molecular Genetics has been thoroughly revised. \* Part One (Chapters 1-7) covers basic material on DNA structure and function, chromosomes, cells and development, pedigree analysis and the basic techniques used in the laboratory. \* Part Two (Chapters 8-12) discusses the various genome sequencing projects and the insights they provide into the organisation, expression, variation and evolution of our genome. \* Part Three (Chapters 13-18) focuses on mapping, identifying and diagnosing the genetic causes of mendelian and complex diseases and cancer. \* Part Four (Chapters 19-21) looks at the wider horizons of functional genomics, proteomics, bioinformatics, animal models and therapy. There are new chapters on cells and development and on functional genomics. The sections on complex diseases have been completely rewritten and reorganized, as has the chapter on Genome Projects. Other changes include a new section on molecular phylogenetics (Chapter 12) and the introduction of 'Ethics Boxes' to discuss some of the implications of the new knowledge. Virtually every page has been revised and updated to take account of the stunning developments of the past four years since the publication of the last edition of Human Molecular Genetics. Features: \* Integration of Human Genome Project data throughout the book \* Two new chapters 'Cells and Development' (Chapter 3) and 'Beyond the Genome Project: Functional Genomics, Proteomics and Bioinformatics' (Chapter 19) \* Completely rewritten and reorganised coverage of complex disease genetics \* Increased emphasis on gene function and on applications of genetic knowledge, including ethical issues \* More prominence given to novel approaches to treating disease, such as cell-based therapies, pharmacogenomics, and personalised medicine \* Special topic boxes that include detailed coverage of ethical, legal and social issues, including eugenics, genetic testing and discrimination, germ-line gene therapy and genetic enhancement, and human cloning \* Contains two indices: a general index and one that contains names of diseases and disorders Supplements: Art of HMG3 (CD-ROM) 0-8153-4183-0: £34.00

Human Molecular Genetics has been carefully crafted over successive editions to provide an authoritative introduction to the molecular aspects of human genetics, genomics and cell biology. Maintaining the features that have made previous editions so popular, this fifth edition has been completely updated in line with the latest developments in the field. Older technologies such as cloning and hybridization have been merged and summarized, coverage of newer DNA sequencing technologies has been expanded, and powerful new gene editing and single-cell genomics technologies have been added. The coverage of GWAS, functional genomics, stem cells, and disease modeling has been expanded. Greater focus is given to inheritance and variation in the context of populations and on the role of epigenetics in gene regulation. Key features: Fully integrated approach to the molecular aspects of human genetics, genomics, and cell biology Accessible text is supported and enhanced throughout by superb artwork illustrating the key concepts and mechanisms Summary boxes at the end of each chapter provide clear learning points Annotated further reading helps readers navigate the wealth of additional information in this complex subject and provides direction for further study Reorganized into five sections for improved access to related topics Also new to this edition □ brand new chapter on evolution and anthropology from the authors of the highly acclaimed Human Evolutionary Genetics A proven and popular textbook for upper-level undergraduates and graduate students, the new edition of Human Molecular Genetics remains the □go-to□ book for those

studying human molecular genetics or genomics courses around the world.

Genetics and Genomics in Medicine is a new textbook written for undergraduate students, graduate students, and medical researchers that explains the science behind the uses of genetics and genomics in medicine today. Rather than focusing narrowly on rare inherited and chromosomal disorders, it is a comprehensive and integrated account of how geneti

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Recent developments within molecular biology and genetic engineering have led to huge advances and changes within the biological sciences especially within the field of human genetics. Diagnostic Techniques in Genetics offers an important overview of how DNA or RNA technology may be applied to a large set of genetic diagnoses. The first part of the book focuses on DNA/RNA applications and includes many of the latest developments in the field combined with routine procedures of genetic diagnoses, for example cloning and sequencing DNA. The DNA applications presented in the first chapter are then each applied to a specific kind of genetic diagnosis and the text concludes with a chapter devoted to population genetics. First published in French by Dunod in 2002, this book is an excellent reference for students taking courses in molecular biology, medicine and medical genetics. It is also a useful introduction for postgraduate students and researchers in the field who require a general overview of genetic diagnoses.

This book provides an introduction to human cytogenetics. It is also suitable for use as a text in a general cytogenetics course, since the basic features of chromosome structure and behavior are shared by all eukaryotes. Because my own background includes plant and animal cytogenetics, many of the examples are taken from organisms other than man. Since the book is written from a cytogeneticist's point of view, human syndromes are described only as illustrations of the effects of abnormal chromosome constitutions on the phenotype. The selection of the phenomena to be discussed and of the photographs to illustrate them is, in many cases, subjective and arbitrary and is naturally influenced by my interests and the work done in our laboratory. The approach to citations is the exact opposite of that usually used in scientific papers. Whenever possible, the latest and/or most comprehensive review has been cited, instead of the original publication. Thus the reader is encouraged to delve deeper into any question of interest to him or her. I am greatly indebted to many colleagues for suggestions and criticism. However, my special thanks are due to Dr. JAMES F. CROW, Dr. TRAUTE M. SCHROEDER, and Dr. CARTER DENNISTON for their courage in reading the entire manuscript. I wish to express my gratitude also to the cytogeneticists and editors who have generously permitted the use of published and unpublished photographs.

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