**DESCRIPTION**

"I think this is an excellent book— I recommend it to anyone involved in molecular epidemiology... The 26 chapters are written by topic specialists, in an explanatory, east to read style." –BTS Newsletter, Summer 2009

“This text provides an accessible and useful handbook for the epidemiologist who wants to survey the field, to become better informed, to look at recent developments and get some background on these or simply to appreciate further the relatively rapid changes in informatic and analytical technologies which increasingly will serve and underpin future epidemiological studies. One of the strengths in this book is the extensive array of practical illustrative examples, and it would also in my opinion have useful potential as a teaching text." –American Journal of Human Biology, March 2009

With the sequencing of the human genome and the mapping of millions of single nucleotide polymorphisms, epidemiology has moved into the molecular domain. Scientists can now use molecular markers to track disease-associated genes in populations, enabling them to study complex chronic diseases that might result from the weak interactions of many genes with the environment. Use of these laboratory generated biomarker data and an understanding of disease mechanisms are increasingly important in elucidating disease aetiology.

*Molecular Epidemiology of Disease* crosses the disciplinary boundaries between laboratory scientists, epidemiologists, clinical researchers and biostatisticians and is accessible to all these relevant research communities in focusing on practical issues of application, rather than reviews of current areas of research.
• Covers categories of biomarkers of exposure, susceptibility and disease

• Includes chapters on novel technologies: genomics, transcriptomics, proteomics and metabonomics, which are increasingly finding application in population studies

• Emphasizes new statistical and bioinformatics approaches necessitated by the large data sets generated using these new methodologies

• Demonstrates the potential applications of laboratory techniques in tackling epidemiological problems while considering their limitations, including the sources of uncertainty and inaccuracy

• Discusses issues such as reliability (compared to traditional epidemiological methods) and the timing of exposure

• Explores practical elements of conducting population studies, including biological repositories and ethics

*Molecular Epidemiology of Disease* provides an easy-to-use, clearly presented handbook that allows epidemiologists to understand the specifics of research involving biomarkers, and laboratory scientists to understand the main issues of epidemiological study design and analysis. It also provides a useful tool for courses on molecular epidemiology, using many examples from population studies to illustrate key concepts and principles.

---

**ABOUT THE AUTHOR**

**Professor Chris Wild.** Molecular Epidemiology Unit, Head, Centre for Epidemiology and Biostatistics, Faculty of Medicine and Health, The LIGHT Laboratories, University of Leeds, UK.

**Professor Paolo Vineis.** Chair in Environmental Epidemiology, Div of Primary Care and Population Health Sciences, Medicine, Imperial College, London, UK.

**Professor Seymour Garte.** Professor of Environmental and Community Medicine, Robert Wood Johnson Medical School, University of Medicine and Dentistry of New Jersey; Scientific Director, Genetics Research Institute, Milan, Italy. Center for Environmental Oncology, University of Pittsburgh Cancer Institute, Pittsburgh, PA, USA.

For additional product details, please visit [https://www.wiley.com/en-us](https://www.wiley.com/en-us)