

Statistics for Biology and Health

David G. Kleinbaum
Mitchel Klein

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Third Edition

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David G. Kleinbaum, Mitchel Klein



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Survival Analysis David G. Kleinbaum, Mitchel Klein, 2014-01-15 Survival Analysis David G. Kleinbaum, 2008-08-01 **Biometrics - Volume I** Susan R. Wilson, Conrad Burden, 2009-02-18 Biometrics is a component of Encyclopedia of Mathematical Sciences in the global Encyclopedia of Life Support Systems EOLSS which is an integrated compendium of twenty one Encyclopedias Biometry is a broad discipline covering all applications of statistics and mathematics to biology The Theme Biometrics is divided into areas of expertise essential for a proper application of statistical and mathematical methods to contemporary biological problems These volumes cover four main topics Data Collection and Analysis Statistical Methodology Computation Biostatistical Methods and Research Design and Selected Topics These volumes are aimed at the following five major target audiences University and College students Educators Professional practitioners Research personnel and Policy analysts managers and decision makers and NGOs

Statistical Methods in Molecular Evolution Rasmus Nielsen, 2005-04-21 In the field of molecular evolution

inferences about past evolutionary events are made using molecular data from currently living species. With the availability of genomic data from multiple related species, molecular evolution has become one of the most active and fastest growing fields of study in genomics and bioinformatics. Most studies in molecular evolution rely heavily on statistical procedures based on stochastic process modelling and advanced computational methods including high dimensional numerical optimization and Markov Chain Monte Carlo. This book provides an overview of the statistical theory and methods used in studies of molecular evolution. It includes an introductory section suitable for readers that are new to the field, a section discussing practical methods for data analysis, and more specialized sections discussing specific models and addressing statistical issues relating to estimation and model choice. The chapters are written by the leaders of the field and they will take the reader from basic introductory material to the state of the art statistical methods. This book is suitable for statisticians seeking to learn more about applications in molecular evolution and molecular evolutionary biologists with an interest in learning more about the theory behind the statistical methods applied in the field. The chapters of the book assume no advanced mathematical skills beyond basic calculus, although familiarity with basic probability theory will help the reader. Most relevant statistical concepts are introduced in the book in the context of their application in molecular evolution, and the book should be accessible for most biology graduate students with an interest in quantitative methods and theory.

Rasmus Nielsen received his Ph.D. from the University of California at Berkeley in 1998 and after a postdoc at Harvard University he assumed a faculty position in Statistical Genomics at Cornell University. He is currently an Ole R. Merz Fellow at the University of Copenhagen and holds a Sloan Research Fellowship. He is an associate editor of the *Journal of Molecular Evolution* and has published more than fifty original papers in peer-reviewed journals on the topic of this book.

From the reviews: Overall, this is a very useful book in an area of increasing importance. *Journal of the Royal Statistical Society*. I find *Statistical Methods in Molecular Evolution* very interesting and useful. It delves into problems that were considered very difficult just several years ago; the book is likely to stimulate the interest of statisticians that are unaware of this exciting field of applications. It is my hope that it will also help the wet lab molecular evolutionist to better understand mathematical and statistical methods. Marek Kimmel for the *Journal of the American Statistical Association*, September 2006.

Who should read this book? We suggest that anyone who deals with molecular data, who does not, and anyone who asks evolutionary questions, who should not, ought to consult the relevant chapters in this book.

Dan Graur and Dror Berel for *Biometrics*, September 2006.

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