

Chapman & Hall/CRC
**Handbooks of Modern
Statistical Methods**

Handbook of Discrete-Valued Time Series

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Handbook Of Discrete Valued Time Series Chapman Hallcrc Handbooks Of Modern Statistical Methods

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Ravishanker, 2016-01-06 Model a Wide Range of Count Time Series Handbook of Discrete Valued Time Series presents state of the art methods for modeling time series of counts and incorporates frequentist and Bayesian approaches for discrete valued spatio temporal data and multivariate data While the book focuses on time series of counts some of the techniques discussed ca Handbook of Statistical Methods and Analyses in Sports Jim Albert, Mark E. Glickman, Tim B. Swartz, Ruud

H. Koning, 2017-02-03 This handbook will provide both overviews of statistical methods in sports and in depth treatment of critical problems and challenges confronting statistical research in sports The material in the handbook will be organized by major sport baseball football hockey basketball and soccer followed by a section on other sports and general statistical design and analysis issues that are common to all sports This handbook has the potential to become the standard reference for obtaining the necessary background to conduct serious statistical analyses for sports applications and to appreciate scholarly work in this expanding area **Handbook of Statistical Methods for Case-Control Studies** Ørnulf

Borgan, Norman Breslow, Nilanjan Chatterjee, Mitchell H. Gail, Alastair Scott, Chris J. Wild, 2018-06-27 Handbook of Statistical Methods for Case Control Studies is written by leading researchers in the field It provides an in depth treatment of up to date and currently developing statistical methods for the design and analysis of case control studies as well as a review of classical principles and methods The handbook is designed to serve as a reference text for biostatisticians and quantitatively oriented epidemiologists who are working on the design and analysis of case control studies or on related statistical methods research Though not specifically intended as a textbook it may also be used as a backup reference text for graduate level courses Book Sections Classical designs and causal inference measurement error power and small sample inference Designs that use full cohort information Time to event data Genetic epidemiology About the Editors rnulf Borgan is Professor of Statistics University of Oslo His book with Andersen Gill and Keiding on counting processes in survival analysis is a world classic Norman E Breslow was at the time of his death Professor Emeritus in Biostatistics University of Washington For decades his book with Nick Day has been the authoritative text on case control methodology Nilanjan Chatterjee is Bloomberg Distinguished Professor Johns Hopkins University He leads a broad research program in statistical methods for modern large scale biomedical studies Mitchell H Gail is a Senior Investigator at the National Cancer Institute His research includes modeling absolute risk of disease intervention trials and statistical methods for epidemiology Alastair Scott was at the time of his death Professor Emeritus of Statistics University of Auckland He was a major contributor to using survey sampling methods for analyzing case control data Chris J Wild is Professor of Statistics University of Auckland His research includes nonlinear regression and methods for fitting models to response selective data **Handbook of Graphical**

Models Marloes Maathuis, Mathias Drton, Steffen Lauritzen, Martin Wainwright, 2018-11-12 A graphical model is a statistical

model that is represented by a graph The factorization properties underlying graphical models facilitate tractable computation with multivariate distributions making the models a valuable tool with a plethora of applications Furthermore directed graphical models allow intuitive causal interpretations and have become a cornerstone for causal inference While there exist a number of excellent books on graphical models the field has grown so much that individual authors can hardly cover its entire scope Moreover the field is interdisciplinary by nature Through chapters by leading researchers from different areas this handbook provides a broad and accessible overview of the state of the art Key features Contributions by leading researchers from a range of disciplines Structured in five parts covering foundations computational aspects statistical inference causal inference and applications Balanced coverage of concepts theory methods examples and applications Chapters can be read mostly independently while cross references highlight connections The handbook is targeted at a wide audience including graduate students applied researchers and experts in graphical models

Handbook of Big Data Peter Bühlmann, Petros Drineas, Michael Kane, Mark van der Laan, 2016-02-22 Handbook of Big Data provides a state of the art overview of the analysis of large scale datasets Featuring contributions from well known experts in statistics and computer science this handbook presents a carefully curated collection of techniques from both industry and academia Thus the text instills a working understanding of key statistical

Handbook of Forensic Statistics David L. Banks, Karen Kafadar, David H. Kaye, Maria Tackett, 2020-11-05 Handbook of Forensic Statistics is a collection of chapters by leading authorities in forensic statistics Written for statisticians scientists and legal professionals having a broad range of statistical expertise it summarizes and compares basic methods of statistical inference frequentist likelihoodist and Bayesian for trace and other evidence that links individuals to crimes the modern history and key controversies in the field and the psychological and legal aspects of such scientific evidence Specific topics include uncertainty in measurements and conclusions statistically valid statements of weight of evidence or source conclusions admissibility and presentation of statistical findings and the state of the art of methods including problems and pitfalls for collecting analyzing and interpreting data in such areas as forensic biology chemistry and pattern and impression evidence The particular types of evidence that are discussed include DNA latent fingerprints firearms and toolmarks glass handwriting shoeprints and voice exemplars

Handbook of Bayesian Variable Selection Mahlet G. Tadesse, Marina Vannucci, 2021-12-24 Bayesian variable selection has experienced substantial developments over the past 30 years with the proliferation of large data sets Identifying relevant variables to include in a model allows simpler interpretation avoids overfitting and multicollinearity and can provide insights into the mechanisms underlying an observed phenomenon Variable selection is especially important when the number of potential predictors is substantially larger than the sample size and sparsity can reasonably be assumed The Handbook of Bayesian Variable Selection provides a comprehensive review of theoretical methodological and computational aspects of Bayesian methods for variable selection The topics covered include spike and slab priors continuous

shrinkage priors Bayes factors Bayesian model averaging partitioning methods as well as variable selection in decision trees and edge selection in graphical models The handbook targets graduate students and established researchers who seek to understand the latest developments in the field It also provides a valuable reference for all interested in applying existing methods and or pursuing methodological extensions Features Provides a comprehensive review of methods and applications of Bayesian variable selection Divided into four parts Spike and Slab Priors Continuous Shrinkage Priors Extensions to various Modeling Other Approaches to Bayesian Variable Selection Covers theoretical and methodological aspects as well as worked out examples with R code provided in the online supplement Includes contributions by experts in the field Supported by a website with code data and other supplementary material

[Advances in Time Series Analysis and Forecasting](#) Ignacio Rojas,Héctor Pomares,Olga Valenzuela,2017-07-31 This volume of selected and peer reviewed contributions on the latest developments in time series analysis and forecasting updates the reader on topics such as analysis of irregularly sampled time series multi scale analysis of univariate and multivariate time series linear and non linear time series models advanced time series forecasting methods applications in time series analysis and forecasting advanced methods and online learning in time series and high dimensional and complex big data time series The contributions were originally presented at the International Work Conference on Time Series ITISE 2016 held in Granada Spain June 27 29 2016 The series of ITISE conferences provides a forum for scientists engineers educators and students to discuss the latest ideas and implementations in the foundations theory models and applications in the field of time series analysis and forecasting It focuses on interdisciplinary and multidisciplinary research encompassing the disciplines of computer science mathematics statistics and econometrics

Handbook of Spatial Epidemiology Andrew B. Lawson,Sudipto Banerjee,Robert P. Haining,Maria Dolores Ugarte,2016-04-06 Handbook of Spatial Epidemiology explains how to model epidemiological problems and improve inference about disease etiology from a geographical perspective Top epidemiologists geographers and statisticians share interdisciplinary viewpoints on analyzing spatial data and space time variations in disease incidences These analyses can provide imp

[Handbook of Approximate Bayesian Computation](#) Scott A. Sisson,Yanan Fan,Mark Beaumont,2018-09-03 As the world becomes increasingly complex so do the statistical models required to analyse the challenging problems ahead For the very first time in a single volume the Handbook of Approximate Bayesian Computation ABC presents an extensive overview of the theory practice and application of ABC methods These simple but powerful statistical techniques take Bayesian statistics beyond the need to specify overly simplified models to the setting where the model is defined only as a process that generates data This process can be arbitrarily complex to the point where standard Bayesian techniques based on working with tractable likelihood functions would not be viable ABC methods finesse the problem of model complexity within the Bayesian framework by exploiting modern computational power thereby permitting approximate Bayesian analyses of models that would otherwise be impossible to implement The Handbook of ABC provides illuminating insight into

the world of Bayesian modelling for intractable models for both experts and newcomers alike It is an essential reference book for anyone interested in learning about and implementing ABC techniques to analyse complex models in the modern world

Handbook of Methods for Designing, Monitoring, and Analyzing Dose-Finding Trials John O'Quigley,Alexia Iasonos,Björn Bornkamp,2017-04-27 Handbook of Methods for Designing Monitoring and Analyzing Dose Finding Trials gives a thorough presentation of state of the art methods for early phase clinical trials The methodology of clinical trials has advanced greatly over the last 20 years and arguably nowhere greater than that of early phase studies The need to accelerate drug development in a rapidly evolving context of targeted therapies immunotherapy combination treatments and complex group structures has provided the stimulus to these advances Typically we deal with very small samples sequential methods that need to be efficient while at the same time adhering to ethical principles due to the involvement of human subjects Statistical inference is difficult since the standard techniques of maximum likelihood do not usually apply as a result of model misspecification and parameter estimates lying on the boundary of the parameter space Bayesian methods play an important part in overcoming these difficulties but nonetheless require special consideration in this particular context The purpose of this handbook is to provide an expanded summary of the field as it stands and also through discussion provide insights into the thinking of leaders in the field as to the potential developments of the years ahead With this goal in mind we present An introduction to the field for graduate students and novices A basis for more established researchers from which to build A collection of material for an advanced course in early phase clinical trials A comprehensive guide to available methodology for practicing statisticians on the design and analysis of dose finding experiments An extensive guide for the multiple comparison and modeling MCP Mod dose finding approach adaptive two stage designs for dose finding as well as dose time response models and multiple testing in the context of confirmatory dose finding studies John O Quigley is a professor of mathematics and research director at the French National Institute for Health and Medical Research based at the Faculty of Mathematics University Pierre and Marie Curie in Paris France He is author of Proportional Hazards Regression and has published extensively in the field of dose finding Alexia Iasonos is an associate attending biostatistician at the Memorial Sloan Kettering Cancer Center in New York She has over one hundred publications in the leading statistical and clinical journals on the methodology and design of early phase clinical trials Dr Iasonos has wide experience in the actual implementation of model based early phase trials and has given courses in scientific meetings internationally Bj rn Bornkamp is a statistical methodologist at Novartis in Basel Switzerland researching and implementing dose finding designs in Phase II clinical trials He is one of the co developers of the MCP Mod methodology for dose finding and main author of the DoseFinding R package He has published numerous papers on dose finding nonlinear models and Bayesian statistics and in 2013 won the Royal Statistical Society award for statistical excellence in the pharmaceutical industry Handbook of Mixture Analysis Sylvia Fruhwirth-Schnatter,Gilles Celeux,Christian P. Robert,2019-01-04 Mixture models have been around

for over 150 years and they are found in many branches of statistical modelling as a versatile and multifaceted tool They can be applied to a wide range of data univariate or multivariate continuous or categorical cross sectional time series networks and much more Mixture analysis is a very active research topic in statistics and machine learning with new developments in methodology and applications taking place all the time The Handbook of Mixture Analysis is a very timely publication presenting a broad overview of the methods and applications of this important field of research It covers a wide array of topics including the EM algorithm Bayesian mixture models model based clustering high dimensional data hidden Markov models and applications in finance genomics and astronomy Features Provides a comprehensive overview of the methods and applications of mixture modelling and analysis Divided into three parts Foundations and Methods Mixture Modelling and Extensions and Selected Applications Contains many worked examples using real data together with computational implementation to illustrate the methods described Includes contributions from the leading researchers in the field The Handbook of Mixture Analysis is targeted at graduate students and young researchers new to the field It will also be an important reference for anyone working in this field whether they are developing new methodology or applying the models to real scientific problems Handbook of Cluster Analysis Christian Hennig, Marina Meila, Fionn Murtagh, Roberto Rocci, 2015-12-16 Handbook of Cluster Analysis provides a comprehensive and unified account of the main research developments in cluster analysis Written by active distinguished researchers in this area the book helps readers make informed choices of the most suitable clustering approach for their problem and make better use of existing cluster analysis tools The *Handbook of Infectious Disease Data Analysis* Leonhard Held, Niel Hens, Philip O'Neill, Jacco Wallinga, 2019-11-07 Recent years have seen an explosion in new kinds of data on infectious diseases including data on social contacts whole genome sequences of pathogens biomarkers for susceptibility to infection serological panel data and surveillance data The Handbook of Infectious Disease Data Analysis provides an overview of many key statistical methods that have been developed in response to such new data streams and the associated ability to address key scientific and epidemiological questions A unique feature of the Handbook is the wide range of topics covered Key features Contributors include many leading researchers in the field Divided into four main sections Basic concepts Analysis of Outbreak Data Analysis of Seroprevalence Data Analysis of Surveillance Data Numerous case studies and examples throughout Provides both introductory material and key reference material *Handbook of Quantile Regression* Roger Koenker, Victor Chernozhukov, Xuming He, Limin Peng, 2017-10-12 Quantile regression constitutes an ensemble of statistical techniques intended to estimate and draw inferences about conditional quantile functions Median regression as introduced in the 18th century by Boscovich and Laplace is a special case In contrast to conventional mean regression that minimizes sums of squared residuals median regression minimizes sums of absolute residuals quantile regression simply replaces symmetric absolute loss by asymmetric linear loss Since its introduction in the 1970 s by Koenker and Bassett quantile regression has

been gradually extended to a wide variety of data analytic settings including time series survival analysis and longitudinal data. By focusing attention on local slices of the conditional distribution of response variables it is capable of providing a more complete, more nuanced view of heterogeneous covariate effects. Applications of quantile regression can now be found throughout the sciences including astrophysics, chemistry, ecology, economics, finance, genomics, medicine, and meteorology. Software for quantile regression is now widely available in all the major statistical computing environments. The objective of this volume is to provide a comprehensive review of recent developments of quantile regression methodology illustrating its applicability in a wide range of scientific settings. The intended audience of the volume is researchers and graduate students across a diverse set of disciplines.

[Handbook of Neuroimaging Data Analysis](#) Hernando Ombao, Martin Lindquist, Wesley Thompson, John Aston, 2016-11-18. This book explores various state-of-the-art aspects behind the statistical analysis of neuroimaging data. It examines the development of novel statistical approaches to model brain data. Designed for researchers in statistics, biostatistics, computer science, cognitive science, computer engineering, biomedical engineering, applied mathematics, physics, and radiology, the book can also be used as a textbook for graduate-level courses in statistics and biostatistics or as a self-study reference for Ph.D. students in statistics, biostatistics, psychology, neuroscience, and computer science.

Economic Evaluation of Cancer Drugs Iftekhar Khan, Ralph Crott, Zahid Bashir, 2019-06-14. Cancer is a major healthcare burden across the world and impacts not only the people diagnosed with various cancers but also their families, carers, and healthcare systems. With advances in the diagnosis and treatment, more people are diagnosed early and receive treatments for a disease where few treatment options were previously available. As a result, the survival of patients with cancer has steadily improved, and in most cases, patients who are not cured may receive multiple lines of treatment, often with financial consequences for the patients, insurers, and healthcare systems. Although many books exist that address economic evaluation, *Economic Evaluation of Cancer Drugs using Clinical Trial and Real World Data* is the first unified text that specifically addresses the economic evaluation of cancer drugs. The authors discuss how to perform cost-effectiveness analyses while emphasising the strategic importance of designing cost-effectiveness into cancer trials and building robust economic evaluation models that have a higher chance of reimbursement if truly cost-effective. They cover the use of real-world data using cancer registries and discuss how such data can support or complement clinical trials with limited follow-up. Lessons learned from failed reimbursement attempts, factors predictive of successful reimbursement, and the different payer requirements across major countries including US, Australia, Canada, UK, Germany, France, and Italy are also discussed. The book includes many detailed practical examples, case studies, and thought-provoking exercises for use in classroom and seminar discussions. Iftekhar Khan is a medical statistician and health economist and a lead statistician at Oxford University's Center for Statistics in Medicine. Professor Khan is also a Senior Research Fellow in Health Economics at University of Warwick and is a Senior Statistical Assessor within the Licensing Division of the UK Medicine and Health Regulation Agency.

Ralph Crott is a former professor in Pharmacoeconomics at the University of Montreal in Quebec Canada and former head of the EORTC Health Economics Unit and former senior health economist at the Belgian HTA organization Zahid Bashir has over twelve years experience working in the pharmaceutical industry in medical affairs and oncology drug development where he is involved in the design and execution of oncology clinical trials and development of reimbursement dossiers for HTA submission Journal of the American Statistical Association ,2007 **Subject Guide to Books in Print** ,1991
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