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About the Author. Robert R. Sokal is a Distinguished Professor Emeritus at Stony Brook University. A native of Vienna, Austria, he attended college in Shanghai, China, and obtained his Ph.D. from the University of Chicago. In both his research, which has ranged over a diverse group of topics, and his teaching, which spans almost a half-century, Dr. Sokal has always promoted the use of statistics in biology.

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Robert Sokal, James Rohlf, Biometry: The Principles and Practice of Statistics in Biological Research, MacMillan, 1995, ISBN13: 9780716724117, LC: QH323.5.S63. Robert Sokal, James Rohlf, Introduction to Biostatistics, Dover, 2009, ISBN13: 9780486469614, LC: QH323.5.S633. Datasets:

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Biometry by Robert R. Sokal

Robert Reuven Sokal was an Austrian-American biostatistician and entomologist. Distinguished Professor Emeritus at the Stony Brook University, Sokal was a member of the National Academy of Sciences and the American Academy of Arts and Sciences. He promoted the use of statistics in biology and co-founded the field of numerical taxonomy, together with Peter H. A. Sneath.

Robert R. Sokal - Wikipedia

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Offers students with little background in statistical analysis an introduction to a variety of statistical concepts and methods. In addition to the incorporation of computer calculation, this new edition expands on a number of important topics, including the revised Kolmogrov-Smirnov test.

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This separate compendium of tables used with Sokal/Rohlf, Biometry, Third Edition, eliminates the inconvenience of having to turn back and forth within the text to refer to data. It can also be used with other texts, or as an independent research resource.

Suitable for undergraduates with a minimal background in mathematics, this introduction ranges from descriptive statistics to fundamental distributions and the testing of hypotheses. Includes numerous worked-out problems and examples. 1987 edition.

A straightforward introduction to a wide range of statistical methods for field biologists, using thoroughly explained R code.

R — the statistical and graphical environment is rapidly emerging as an important set of teaching and research tools for biologists. This book draws upon the popularity and free availability of R to couple the theory and practice of biostatistics into a single treatment, so as to provide a textbook for biologists learning statistics, R, or both. An abridged description of biostatistical principles and analysis sequence keys are combined together with worked examples of the practical use of R into a complete practical guide to designing and analyzing real biological research. Topics covered include: simple hypothesis testing, graphing exploratory data analysis and graphical summaries regression (linear, multi and non-linear) simple and complex ANOVA and ANCOVA designs (including nested, factorial, blocking, spit-plot and repeated measures) frequency analysis and generalized linear models. Linear mixed effects modeling is also incorporated extensively throughout as an alternative to traditional modeling techniques. The book is accompanied by a companion website www.wiley.com/go/logan/r with an extensive set of resources comprising all R scripts and data sets used in the book, additional worked examples, the biology package, and other instructional materials and links.

The Biostatistics course is often found in the schools of public Health, medical schools, and, occasionally, in statistics and biology departments. The population of students in these courses is a diverse one, with varying preparedness. The book assumes the reader has at least two years of high school algebra, but no previous exposure to statistics is required. Written for individuals who might be fearful of mathematics, this book minimizes the technical difficulties and emphasizes the importance of statistics in scientific investigation. An understanding of underlying design and analysis is stressed. The limitations of the research, design and analytical techniques are discussed, allowing the reader to accurately interpret results. Real data, both processed and raw, are used extensively in examples and exercises. Statistical computing packages - MINITAB, SAS and Stata - are integrated. The use of the computer and software allows a sharper focus on the concepts, letting the computer do the necessary number-crunching. * Emphasizes underlying statistical concepts more than competing texts * Focuses on experimental design and analysis, at an elementary level * Includes an introduction to linear correlation and regression * Statistics are central: probability is downplayed * Presents life tables and survival analysis * Appendix with solutions to many exercises * Special instructor's manual with solution to all exercises

A thoroughly updated and expanded step-wise guide to the study of animal behaviour.

An essential textbook for any student or researcher in biology needing to design experiments, sample programs or analyse the resulting data. The text begins with a revision of estimation and hypothesis testing methods, covering both classical and Bayesian philosophies, before advancing to the analysis of linear and generalized linear models. Topics covered include linear and logistic regression, simple and complex ANOVA models (for factorial, nested, block, split-plot and repeated measures and covariance designs), and log-linear models. Multivariate techniques, including classification and ordination, are then introduced. Special emphasis is placed on checking assumptions, exploratory data analysis and presentation of results. The main analyses are illustrated with many examples from published papers and there is an extensive reference list to both the statistical and biological literature. The book is supported by a website that provides all data sets, questions for each chapter and links to software.

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