Preface

This book is intended as a guide to data analysis with the R system for statistical computing. R is an environment incorporating an implementation of the S programming language, which is powerful, flexible and has excellent graphical facilities (R Development Core Team, 2005). In the Handbook we aim to give relatively brief and straightforward descriptions of how to conduct a range of statistical analyses using R. Each chapter deals with the analysis appropriate for one or several data sets. A brief account of the relevant statistical background is included in each chapter along with appropriate references, but our prime focus is on how to use R and how to interpret results. We hope the book will provide students and researchers in many disciplines with a self-contained means of using R to analyse their data.

R is an open-source project developed by dozens of volunteers for more than ten years now and is available from the Internet under the General Public Licence. R has become the lingua franca of statistical computing. Increasingly, implementations of new statistical methodology first appear as R add-on packages. In some communities, such as in bioinformatics, R already is the primary workhorse for statistical analyses. Because the sources of the R system are open and available to everyone without restrictions and because of its powerful language and graphical capabilities, R has started to become the main computing engine for reproducible statistical research (Leisch, 2002a,b, 2003, Leisch and Rossini, 2003, Gentleman, 2005). For a reproducible piece of research, the original observations, all data preprocessing steps, the statistical analysis as well as the scientific report form a unity and all need to be available for inspection, reproduction and modification by the readers.

Reproducibility is a natural requirement for textbooks such as the ‘Handbook of Statistical Analyses Using R’ and therefore this book is fully reproducible using an R version greater or equal to 4.2.0. All analyses and results, including figures and tables, can be reproduced by the reader without having to retype a single line of R code. The data sets presented in this book are collected in a dedicated add-on package called HSAUR accompanying this book. The package can be installed from the Comprehensive R Archive Network (CRAN) via

R> install.packages("HSAUR")

and its functionality is attached by

R> library("HSAUR")

The relevant parts of each chapter are available as a vignette, basically a
document including both the R sources and the rendered output of every
analysis contained in the book. For example, the first chapter can be inspected by

R> vignette("Ch_introduction_to_R", package = "HSAUR")

and the R sources are available for reproducing our analyses by

R> edit(vignette("Ch_introduction_to_R", package = "HSAUR"))

An overview on all chapter vignettes included in the package can be obtained from

R> vignette(package = "HSAUR")

We welcome comments on the R package HSAUR, and where we think these
add to or improve our analysis of a data set we will incorporate them into the
package and, hopefully at a later stage, into a revised or second edition of the book.

Plots and tables of results obtained from R are all labelled as ‘Figures’ in
the text. For the graphical material, the corresponding figure also contains
the ‘essence’ of the R code used to produce the figure, although this code may
differ a little from that given in the HSAUR package, since the latter may include some features, for example thicker line widths, designed to make a
basic plot more suitable for publication.

We would like to thank the R Development Core Team for the R system, and
authors of contributed add-on packages, particularly Uwe Ligges and Vince
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course, the joint responsibility of the two authors.

Brian S. Everitt and Torsten Hothorn
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Bibliography


