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Statistical Genetics: Analyses with R

Here comes a dedication

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Preface

Hi there, this is my great book.

Why read this book

It is very important...

Structure of the book

Chapters ?? introduces a new topic, and ...

Software information and conventions

I used the **knitr** package (?) and the **bookdown** package (Xie, 2016) to compile my book. My R session information is shown below:

```
sessionInfo()
```

```
## R version 3.3.3 (2017-03-06)
## Platform: x86_64-apple-darwin13.4.0 (64-bit)
## Running under: macOS Sierra 10.12.3
```

```
##
## locale:
## [1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8
##
## attached base packages:
## [1] stats      graphics  grDevices  utils      datasets
## [6] methods    base
##
## loaded via a namespace (and not attached):
## [1] backports_1.0.5 bookdown_0.3   magrittr_1.5
## [4] rprojroot_1.2   tools_3.3.3   htmltools_0.3.5
## [7] rstudioapi_0.6  yaml_2.1.14   Rcpp_0.12.10
## [10] stringi_1.1.5   rmarkdown_1.4  knitr_1.15.1
## [13] stringr_1.2.0   digest_0.6.12  evaluate_0.10
```

Package names are in bold text (e.g., **rmarkdown**), and inline code and filenames are formatted in a typewriter font (e.g., `knitr::knit('foo.Rmd')`). Function names are followed by parentheses (e.g., `bookdown::render_book()`).

Acknowledgments

A lot of people helped me when I was writing the book.

Frida Gomam
on the Mars

About the Author





Introduction to Statistical Genetics



○

Evolutionary Genetics



○

Population Genetics



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Quantitative Genetics



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Genetics of complex diseases



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Genetic Epidemiology



generate a BibTeX database automatically for some R packages



Bibliography

Xie, Y. (2016). *bookdown: Authoring Books and Technical Documents with R Markdown*. R package version 0.3.



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