

## Statistical Bioinformatics With R

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Statistical Bioinformatics provides a balanced treatment of statistical theory in the context of bioinformatics applications. Designed for a one or two semester senior undergraduate or graduate bioinformatics course, the text takes a broad view of the subject – not just gene expression and sequence analysis, but a careful balance of statistical theory in the context of bioinformatics ...

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Statistical Bioinformatics With R This course provides an introduction to some statistical techniques through the use of the R language. Topics covered include: Chi2 and Fisher tests, descriptive statistics, t-test, analysis of variance and regression. Students will run analyses using statistical and graphical skills taught during the session.

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### *Statistical Bioinformatics With R*

R is a rapidly growing language making basic as well as advanced statistical programming easy. From an educational point of view, R provides the possibility to combine the learning of statistical concepts by mathematics, programming, and visualization. The plots and tables produced by R can

### *Wim P. Krijnen November 10, 2009 - The Comprehensive R ...*

extensible, R can unify most (if not all) bioinformatics data analysis tasks in one program with add-on packages. Rather than learn multiple tools, students and researchers can use one consistent environment for many tasks. It is because of the price of R, extensibility, and the growing use of R in bioinformatics that R

### *Statistics Using R with Biological Examples*

Designed for a one or two semester senior undergraduate or graduate bioinformatics course, the text takes a broad view of the subject – not just gene expression and sequence analysis, but a careful balance of statistical theory in the context of bioinformatics applications. The inclusion of R&SAS code as well as the development of advanced methodology such as Bayesian and Markov models provides students with the important foundation needed to conduct bioinformatics.

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This course teaches the R programming language in the context of statistical data and statistical analysis in the life sciences. We will learn the basics of statistical inference in order to understand and compute p-values and confidence intervals, all while analyzing data with R code.

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### *Introduction to Bioinformatics with R: A Practical Guide ...*

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