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as a guide to its application.

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## **Computational and statistical methods for high-throughput ...**

Computational statistics, or statistical computing, is the interface between statistics and computer science. It is the area of computational science (or

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scientific computing) specific to the mathematical science of statistics. This area is also developing rapidly, leading to calls that a broader concept of computing should be taught as part of general statistical education. As in traditional statistics the goal is to transform raw data into knowledge, but the focus lies on computer intensive st

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## **Computational statistics - Wikipedia**

Statistics can be called that body of analytical and computational methods by which characteristics of a population are inferred through observations made in a representative sample from that population. Since scientists rarely observe entire populations, sampling

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and statistical inference are essential.

## **STATISTICAL METHODS - Swartz Center for Computational ...**

The use of statistical methods and tools from applied probability to address problems in computational molecular biology. Biological problems in sequence analysis, structure prediction, gene

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expression analysis, phy-logenetic trees, and statistical genetics will be addressed. The following

## **Statistical Methods for Computational Biology**

Computational and Statistical Methods for Protein Quantification by Mass Spectrometry: Introduces the use of

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mass spectrometry in protein quantification and how the bioinformatics challenges in this field can be solved using statistical methods and various software programs.

## **Computational and Statistical Methods for Protein ...**

In this review, we intend to provide an

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overview of computational and statistical methods that are commonly applied to analyze metabolomics data. The review is divided into five sections. The first two sections will introduce the background and the databases and resources available for metabolomics research. The third section will briefly ...

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The Computational and Statistical Genomics Branch focuses on the development and application of computationally intensive approaches to analyze large-scale genetic and genomic data, with a particular focus on identifying genetic contributions to

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human disease.

## **Computational and Statistical Genomics Branch**

Statistical methods for reproducible data analysis. Computational tools for reproducible data analysis and version control (Git/GitHub, Emacs/RStudio/Spyder), reproducible

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data (Data repositories/Dataverse) and  
reproducible dynamic report generation  
(Rmarkdown/R  
Notebook/Jupyter/Pandoc), and  
workflows.

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Statistical methods for reproducible data

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analysis Computational tools for  
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control (Git/GitHub,  
Emacs/RStudio/Spyder), reproducible  
data (Data repositories/Dataverse) and  
reproducible dynamic report generation  
(Rmarkdown/R  
Notebook/Jupyter/Pandoc), and  
workflows.

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uses the latest approaches in genomics,  
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develop and apply computationally  
efficient and statistically robust methods  
to sort through increasingly rich and  
massive genome wide data sets to  
identify complex genetic patterns, gene  
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