



Bioinformatics: 6 (Applied Mycology and Biotechnology)

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The advances in genomic technologies, such as microarrays and high throughput sequencing, have expanded the realm of possibilities for capturing data and analyzing it using automated computer driven bioinformatics tools. With the completion of the sequencing of genomes of human and several model organisms, a quest for scientific discoveries being fueled by integrative and multidimensional techniques in mathematics and computational sciences. In this volume, leading researchers and experts have provided an overview of significant concepts from biological, mathematical, and computational perspectives. It provides a high level view of fungal genomic data integration and annotation, classification of proteins and identification of vaccine targets, identification of secretome or secreted proteins in fungal genomes, as well as tools for analyzing microarray expression profiles.

- * Provides a survey of theoretical underpinnings on the technological tools and applications
- * Discusses the tools utilized for the annotation of fungal genomes and addresses issues related to automated annotation generation in a high throughput biotechnology environment
- * describing the applications of the concepts and methodologies presented throughout the book

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