

A Cell Biologist's Guide to Modeling and Bioinformatics

By Raquell M. Holmes



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A step-by-step guide to using computational tools to solve problems in cell biology

Combining expert discussion with examples that can be reproduced by the reader, *A Cell Biologist's Guide to Modeling and Bioinformatics* introduces an array of informatics tools that are available for analyzing biological data and modeling cellular processes. You learn to fully leverage public databases and create your own computational models. All that you need is a working knowledge of algebra and cellular biology; the author provides all the other tools you need to understand the necessary statistical and mathematical methods.

Coverage is divided into two main categories:

- Molecular sequence database chapters are dedicated to gaining an
 understanding of tools and strategies—including queries, alignment methods,
 and statistical significance measures—needed to improve searches for sequence
 similarity, protein families, and putative functional domains. Discussions of
 sequence alignments and biological database searching focus on publicly
 available resources used for background research and the characterization of
 novel gene products.
- Modeling chapters take you through all the steps involved in creating a
 computational model for such basic research areas as cell cycle, calcium
 dynamics, and glycolysis. Each chapter introduces a new simulation tooland is
 based on published research. The combination creates a rich context for
 ongoing skill and knowledge development in modeling biological research
 systems.

Students and professional cell biologists can develop the basic skills needed to learn computational cell biology. This unique text, with its step-by-step instruction, enables you to test and develop your new bioinformatics and modeling skills. References are provided to help you take advantage of more advanced techniques, technologies, and training.

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Editorial Review

Review

"A Cell Biologist's Guide to Modeling and Bioinformatics is well written, well illustrated...this volume is a must-read for any biologist who wants an introduction to quantitative methods. It will also be a useful textbook for students." (*The Quarterly Review of Biology*, September 2008)

From the Back Cover

A step-by-step guide to using computational tools to solve problems in cell biology

Combining expert discussion with examples that can be reproduced by the reader, this text introduces an array of informatics tools that are available for analyzing biological data and modeling cellular processes. You learn to fully leverage public databases and create your own computational models. All that you need is a working knowledge of algebra and cellular biology; the author provides all the other tools you need to understand the necessary statistical and mathematical methods.

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About the Author

Dr. Holmes is a cell biologist who has worked to develop the Bioinformatics Graduate Program at Boston University and works with computational scientists in the Education, Outreach, and Training Partnership for Advanced Computational Infrastructure. She is presently a Program Manager and researcher at the Center for Computational Science at Boston University.

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