



Genomes

By Terence A. Brown

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Genomes By Terence A. Brown

Genomes 2 covers modern molecular genetics from the genomics perspective, incorporating major advances made in the past three years, including the sequencing of the human genome, characterization of genome expression and replication processes, and transcriptomics and proteomics. The text is richly illustrated with clear, easy-to-follow, full-color diagrams, which are downloadable from the book's website (www.wiley.com/brown).

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Genomes By Terence A. Brown Bibliography

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Editorial Review

From Library Journal

This undergraduate textbook concentrates on the study of the genome (the total genetic makeup of an organism) as a whole rather than the traditional approach of studying specific genes. The writing is clear and straightforward, and the diagrams are excellent, making difficult topics like DNA sequencing, RNA replication, and genomic evolution, mutation, and recombination approachable and understandable.

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Review

"...this didactic masterpiece?model of a modern university textbook deserves to be read and studied by?students worldwide." (*Modern Pathology*, Vol. 15, No. 9, 2002)

"...a second edition to incorporate new developments in genome science and to make the book more 'user friendly'...the book substantially exceeds that modest objective; this is a text suitable for anyone who does not specialize in genome science." (*Clinical Chemistry*, Vol. 48, No. 12)

"...a marvelous text?plenty of definitions and excellent illustrations essential for an introductory textbook...the reader is...guided wonderfully through the text...this text is an essential cornerstone of information..." (*Pharmaceutical Research*, Vol. 19, No. 12, December 2002)

"...probably the most up-to-date textbook on molecular biology...a useful text...and reference..." (*Annals of Pharmacotherapy*, September 2003)

Review

"I have found Genomes 2 by Terence Brown to be quite up to date, insightful and written in an engaging fashion. It nicely covers areas on the cutting edge, such as proteomics." -- Richard D. Smith, Ph.D., Pacific Northwest National Laboratory, Richland, WA

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