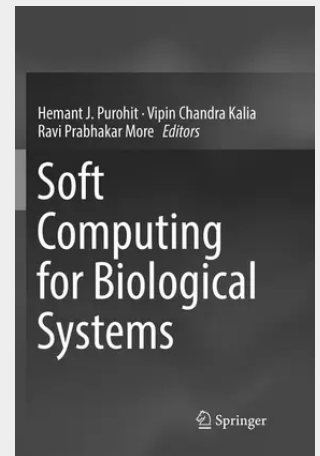


Soft Computing for Biological Systems

Includes well-illustrated chapters for easy understanding of methodologies
Provides the most recent research in the area of soft-computing analysis
Discusses how the insights from various computational studies can be collated and ultimately exploited practically

This book explains how the biological systems and their functions are driven by genetic information stored in the DNA, and their expression driven by different factors. The soft computing approach recognizes the different patterns in DNA sequence and try to assign the biological relevance with available information. The book also focuses on using the soft-computing approach to predict protein-protein interactions, gene expression and networks. The insights from these studies can be used in metagenomic data analysis and predicting artificial neural networks.



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