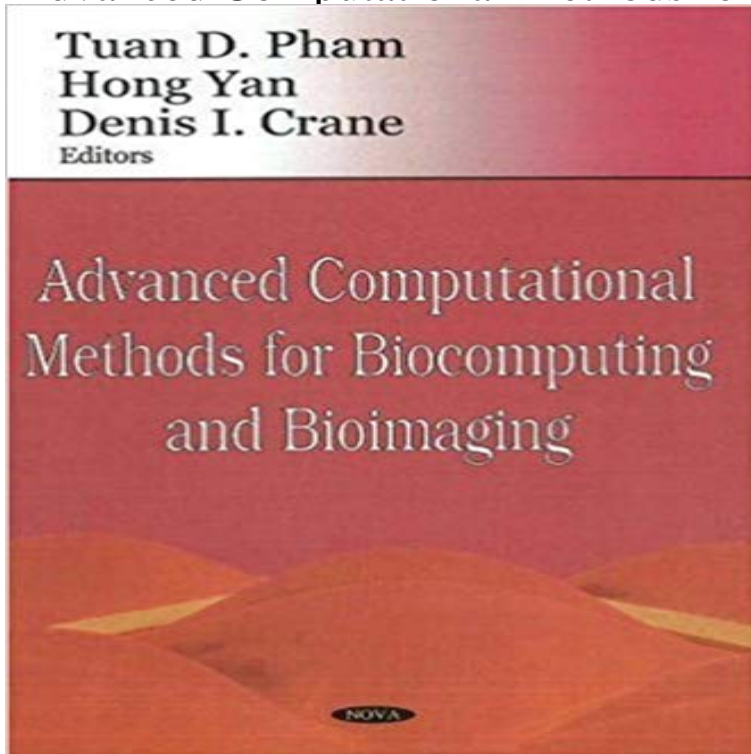


# Advanced Computational Methods for Biocomputing And Bioimaging



Computational models have been playing a significant role for the computer-based analysis of biological and biomedical data. Given the recent availability of genomic sequences and microarray gene expression data, there is an increasing demand for developing and applying advanced computational techniques for exploring these types of data, such as functional interpretation of gene expression data, deciphering of how genes and proteins work together in pathways and networks, extracting and analysing phenotypic features of mitotic cells for high throughput screening of novel anti-mitotic drugs. Successful applications of advanced computational algorithms to solving modern life-science problems will make significant impacts on several important and promising issues related to genomic medicine, molecular imaging, and the scientific knowledge of the genetic basis of diseases.

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