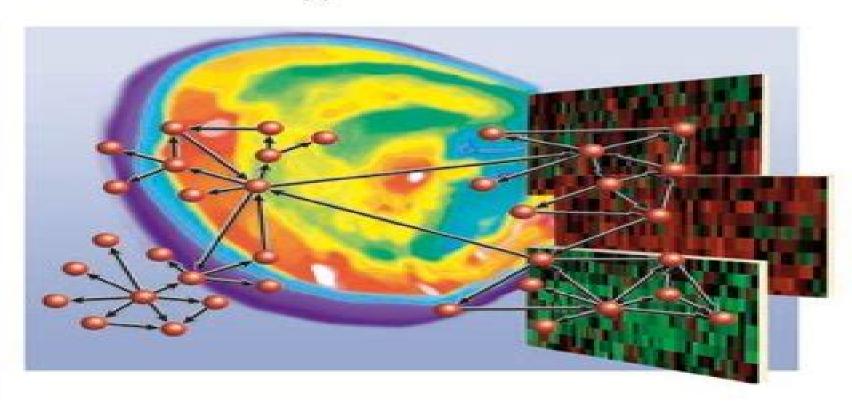


# Analysis of Microarray Data

A Network-Based Approach



# **Analysis Of Microarray Data A Network Based Approach**

**Matthias Dehmer, Frank Emmert-Streib** 

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on experimental data the book uniquely sets itself apart from the current literature by exploring the application of machine learning techniques to various types of complex networks Comprised of chapters written by internationally renowned researchers in the field of interdisciplinary network theory the book presents current and classical methods to analyze networks statistically Methods from machine learning data mining and information theory are strongly emphasized throughout Real data sets are used to showcase the discussed methods and topics which include A survey of computational approaches to reconstruct and partition biological networks An introduction to complex networks measures statistical properties and models Modeling for evolving biological networks The structure of an evolving random bipartite graph Density based enumeration in structured data Hyponym extraction employing a weighted graph kernel Statistical and Machine Learning Approaches for Network Analysis is an excellent supplemental text for graduate level cross disciplinary courses in applied discrete mathematics bioinformatics pattern recognition and computer science The book is also a valuable reference for researchers and practitioners in the fields of applied discrete mathematics machine learning data mining and biostatistics

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Analyzing the Large Number of Variables in Biomedical and Satellite Imagery Phillip I. Good, 2011-05-18 This book grew out of an online interactive offered through statcourse com and it soon became apparent to the author that the course was too limited in terms of time and length in light of the broad backgrounds of the enrolled students The statisticians who took the course needed to be brought up to speed both on the biological context as well as on the specialized statistical methods needed to handle large arrays Biologists and physicians even though fully knowledgeable concerning the procedures used to

generate microaarrays EEGs or MRIs needed a full introduction to the resampling methods the bootstrap decision trees and permutation tests before the specialized methods applicable to large arrays could be introduced As the intended audience for this book consists both of statisticians and of medical and biological research workers as well as all those research workers who make use of satellite imagery including agronomists and meteorologists the book provides a step by step approach to not only the specialized methods needed to analyze the data from microarrays and images but also to the resampling methods step down multi comparison procedures multivariate analysis as well as data collection and pre processing While many alternate techniques for analysis have been introduced in the past decade the author has selected only those techniques for which software is available along with a list of the available links from which the software may be purchased or downloaded without charge Topical coverage includes very large arrays permutation tests applying permutation tests gathering and preparing data for analysis multiple tests bootstrap applying the bootstrap classification methods decision trees and applying decision trees Computational Intelligence in Cancer Diagnosis Janmenjoy Nayak, Danilo Pelusi, Bighnaraj Naik, Manohar Mishra, Khan Muhammad, David Al-Dabass, 2023-04-12 Computational Intelligence in Cancer Diagnosis Progress and Challenges provides insights into the current strength and weaknesses of different applications and research findings on computational intelligence in cancer research The book improves the exchange of ideas and coherence among various computational intelligence methods and enhances the relevance and exploitation of application areas for both experienced and novice end users Topics discussed include neural networks fuzzy logic connectionist systems genetic algorithms evolutionary computation cellular automata self organizing systems soft computing fuzzy systems and hybrid intelligent systems The book s chapters are written by international experts from both cancer research oncology and computational sides to cover different aspects and make it comprehensible for readers with no background on informatics Contains updated information about advanced computational intelligence spanning the areas of neural networks fuzzy logic connectionist systems genetic algorithms evolutionary computation cellular automata self organizing systems soft computing fuzzy systems and hybrid intelligent systems in diagnosing cancer diseases Discusses several cancer types including their detection treatment and prevention Presents case studies that illustrate the applications of intelligent computing in data analysis to help readers to analyze and advance their research in cancer **Computational Intelligence and** Bioinformatics De-Shuang Huang, George William Irwin, 2006-08-03 This book constitutes the refereed proceedings of the International Conference on Intelligent Computing ICIC 2006 held in Kunming China in August 2006 The book presents 165 revised full papers carefully reviewed Topics covered include ant colony optimization particle swarm optimization swarm intelligence autonomy oriented computing quantum and molecular computations biological and DNA computing intelligent computing in bioinformatics intelligent computing in computational biology and drug design computational genomics and proteomics and more

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