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<u>Analysis Of Microarray Gene Expression Data Trends In</u> <u>Logic</u>

Y Pai

Analysis of Microarray Gene Expression Data Mei-Ling Ting Lee, 2004-04-30 After genomic sequencing microarray technology has emerged as a widely used platform for genomic studies in the life sciences Microarray technology provides a systematic way to survey DNA and RNA variation With the abundance of data produced from microarray studies however the ultimate impact of the studies on biology will depend heavily on data mining and statistical analysis The contribution of this book is to provide readers with an integrated presentation of various topics on analyzing microarray data in Intelligent Computing Research and Development Tripathy, B.K., Acharjya, D. P., 2013-12-31 As the amount of accumulated data across a variety of fields becomes harder to maintain it is essential for a new generation of computational theories and tools to assist humans in extracting knowledge from this rapidly growing digital data Global Trends in Intelligent Computing Research and Development brings together recent advances and in depth knowledge in the fields of knowledge representation and computational intelligence Highlighting the theoretical advances and their applications to real life problems this book is an essential tool for researchers lecturers professors students and developers who have seek insight into knowledge representation and real life applications An Introduction to Toxicogenomics Michael E. Burczynski, 2003-03-26 Since the advent of cDNA microarrays oligonucleotide array technology and gene chip analysis genomics has revolutionized the entire field of biomedical research A byproduct of this revolution toxicogenomics is a fast rising star within toxicological analysis Gathering together leading authors and scientists at the forefront of the field An Introduction to Toxicogenomics provides a comprehensive overview of this new discipline With a focus on toxicology it introduces the basic principles of microarray oligonucleotide array based genomic analysis and explains how it fits into the field of biomedical research These discussions provide an overview to the actual mechanics of the analyses themselves and offer insights on handling and quality control Then the book features an important section on the basics of data analysis and clustering methods such as genetic algorithms Finally it covers the application of expression profiling in the field of toxicology and addresses the two fundamental types of analysis in detail with sections dedicated to both mechanistic and predictive studies Although toxicogenomics promises fast efficient techniques and information rich data much of its potential remains untapped An Introduction to Toxicogenomics consolidates the concepts underlying the field to provide a solid foundation from which to begin your research endeavors Scalable Pattern Recognition Algorithms Pradipta Maji, Sushmita Paul, 2014-03-19 This book addresses the need for a unified framework describing how soft computing and machine learning techniques can be judiciously formulated and used in building efficient pattern recognition models. The text reviews both established and cutting edge research providing a careful balance of theory algorithms and applications with a particular emphasis given to applications in computational biology and bioinformatics Features integrates different soft computing and machine learning methodologies with pattern recognition tasks discusses in detail the integration of different

techniques for handling uncertainties in decision making and efficiently mining large biological datasets presents a particular emphasis on real life applications such as microarray expression datasets and magnetic resonance images includes numerous examples and experimental results to support the theoretical concepts described concludes each chapter with directions for future research and a comprehensive bibliography *Computational Intelligence in Bioinformatics* Arpad Kelemen, Ajith Abraham, 2008-01-03 Bioinformatics involve the creation and advancement of algorithms using techniques including computational intelligence applied mathematics and statistics informatics and biochemistry to solve biological problems usually on the molecular level This book deals with the application of computational intelligence in bioinformatics Addressing the various issues of bioinformatics using different computational intelligence approaches is the novelty of this edited volume

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deals with methods and protocols for the preparation of microarrays The second volume details applications and data analysis which is important in analyzing the enormous data coming out of microarray experiments Among the topics discussed in Volume 1 Synthesis Methods are matrices in the synthesis of microarrays array optimization processes array based comparative genomic hybridization 60 mer oligonucleotide probes bifunctional reagents NTMTA and NTPAC and high density arrays using digital microarray synthesis platforms Other topics include multiplex ligation dependent probe amplification MLPA hybridization conditions in situ synthesized oligo arrays peptide arrays high density replication tools HDRT protocols for the quantification of oligo hybridization glyco bead arrays and an investigation into the emerging nano technology Microarray Technology Volumes 1 and 2 provide ample information to all levels of scientists from novice to those intimately familiar with array technology

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