

1. Record Nr.	UNINA9910830637803321
Autore	McLachlan Geoffrey J. <1946->
Titolo	Analyzing microarray gene expression data [[electronic resource] ] / Geoffrey J. McLachlan, Kim-Anh Do, Christopher Ambroise
Pubbl/distr/stampa	Hoboken, N.J., : Wiley-Interscience, c2004
ISBN	1-280-25332-0 9786610253326 0-470-35030-X 0-471-72612-5 0-471-72842-X
Descrizione fisica	1 online resource (366 p.)
Collana	Wiley series in probability and statistics
Altri autori (Persone)	DoKim-Anh <1960-> AmbroiseChristophe <1969->
Disciplina	572.8636 572.865
Soggetti	DNA microarrays - Statistical methods Gene expression - Statistical methods
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Description based upon print version of record.
Nota di bibliografia	Includes bibliographical references and index.
Nota di contenuto	Analyzing Microarray Gene Expression Data; Contents; Preface; 1 Microarrays in Gene Expression Studies; 1.1 Introduction; 1.2 Background Biology; 1.2.1 Genome, Genotype, and Gene Expression; 1.2.2 Of Wild-Types and Other Alleles; 1.2.3 Aspects of Underlying Biology and Physiochemistry; 1.3 Polymerase Chain Reaction; 1.4 cDNA; 1.4.1 Expressed Sequence Tag; 1.5 Microarray Technology and Application; 1.5.1 History of Microarray Development; 1.5.2 Tools of Microarray Technology; 1.5.3 Limitations of Microarray Technology; 1.5.4 Oligonucleotides versus cDNA Arrays 1.5.5 SAGE: Another Method for Detecting and Measuring Gene Expression Levels1.5.6 Emerging Technologies; 1.6 Sampling of Relevant Research Entities and Public Resources; 2 Cleaning and Normalization; 2.1 Introduction; 2.2 Cleaning Procedures; 2.2.1 Image Processing to Extract Information; 2.2.2 Missing Value Estimation; 2.2.3 Sources of Nonlinearity; 2.3 Normalization and Plotting Procedures for Oligonucleotide Arrays; 2.3.1 Global Approaches for Oligonucleotide

Array Data; 2.3.2 Spiked Standard Approaches; 2.3.3 Geometric Mean and Linear Regression Normalization for Multiple Arrays  
2.3.4 Nonlinear Normalization for Multiple Arrays Using Smooth Curves  
2.4 Normalization Methods for cDNA Microarray Data; 2.4.1 Single-Array Normalization; 2.4.2 Multiple Slides Normalization; 2.4.3 ANOVA and Related Methods for Normalization; 2.4.4 Mixed-Model Method for Normalization; 2.4.5 SNOMAD; 2.5 Transformations and Replication; 2.5.1 Importance of Replication; 2.5.2 Transformations; 2.6 Analysis of the Alon Data Set; 2.7 Comparison of Normalization Strategies and Discussion; 3 Some Cluster Analysis Methods; 3.1 Introduction; 3.2 Reduction in the Dimension of the Feature Space  
3.3 Cluster Analysis  
3.4 Some Hierarchical Agglomerative Techniques; 3.5 k-Means Clustering; 3.6 Cluster Analysis with No A Priori Metric; 3.7 Clustering via Finite Mixture Models; 3.7.1 Definition; 3.7.2 Advantages of Model-Based Clustering; 3.8 Fitting Mixture Models Via the EM Algorithm; 3.8.1 E-Step; 3.8.2 M-Step; 3.8.3 Choice of Starting Values for the EM Algorithm; 3.9 Clustering Via Normal Mixtures; 3.9.1 Heteroscedastic Components; 3.9.2 Homoscedastic Components; 3.9.3 Spherical Components; 3.9.4 Choice of Root; 3.9.5 Available Software; 3.10 Mixtures of t Distributions  
3.11 Mixtures of Factor Analyzers  
3.12 Choice of Clustering Solution; 3.13 Classification ML Approach; 3.14 Mixture Models for Clinical and Microarray Data; 3.14.1 Unconditional Approach; 3.14.2 Conditional Approach; 3.15 Choice of the Number of Components in a Mixture Model; 3.15.1 Order of a Mixture Model; 3.15.2 Approaches for Assessing Mixture Order; 3.15.3 Bayesian Information Criterion; 3.15.4 Integrated Classification Likelihood Criterion; 3.16 Resampling Approach; 3.17 Other Resampling Approaches for Number of Clusters; 3.17.1 The Gap Statistic  
3.17.2 The Cleft Method for the Number of Clusters

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## Sommario/riassunto

A multi-discipline, hands-on guide to microarray analysis of biological processes Analyzing Microarray Gene Expression Data provides a comprehensive review of available methodologies for the analysis of data derived from the latest DNA microarray technologies. Designed for biostatisticians entering the field of microarray analysis as well as biologists seeking to more effectively analyze their own experimental data, the text features a unique interdisciplinary approach and a combined academic and practical perspective that offers readers the most complete and applied coverage of the subject

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2. Record Nr.	UNISA996218266003316
Autore	Lewalski Barbara Kiefer <1931->
Titolo	The life of John Milton [[electronic resource] ] : a critical biography // Barbara K. Lewalski
Pubbl/distr/stampa	Oxford ; ; Malden, MA, : Blackwell Publishers, 2000
ISBN	1-281-31076-X 9786611310769 0-470-77567-X 0-470-77684-6
Edizione	[1st ed.]
Descrizione fisica	1 online resource (814 p.)
Collana	Blackwell critical biographies
Disciplina	821/.4 B
Soggetti	Poets, English - Early modern, 1500-1700
Lingua di pubblicazione	Inglese
Formato	Materiale a stampa
Livello bibliografico	Monografia
Note generali	Description based upon print version of record.
Nota di bibliografia	Includes bibliographical references (p. [705]-753) and index.
Nota di contenuto	The Life of John Milton: A Critical Biography Revised Edition; Contents; List of Plates; Preface; Acknowledgments; List of Abbreviations; 1 "The Childhood Shews the Man" 1608-1625; 2 "To Cambridge . . . for Seven Years" 1625-1632; 3 "Studious Retirement": Hammersmith and Horton 1632-1638; 4 "I Became Desirous . . . of Seeing Foreign Parts, Especially Italy" 1638-1639; 5 "All Mouths Were Opened Against . . . the Bishops" 1639-1642; 6 "Domestic or Personal Liberty" 1642-1645; 7 "Service . . . Between Private Walls" 1645-1649 8 "The So-called Council of State . . . Desired to Employ My Services" 1649-1652 9 "Tireless . . . for the Sake of Liberty" 1652-1654; 10 "I . . . Still Bear Up and Steer Right Onward" 1654-1658; 11 "The Last Words of Our Expiring Libertie" 1658-1660; 12 "In Darknes, and with Dangers Compast Round" 1660-1665; 13 "Higher Argument": Completing and Publishing Paradise Lost 1665-1669; 14 "To Try, and Teach the Erring Soul" 1669-1674; Epilogue: "Something . . . Written to Aftertimes"; Notes; Select Bibliography; Index
Sommario/riassunto	Providing a close examination of Milton's wide-ranging prose and poetry at each stage of his life, Barbara Lewalski reveals a rather different Milton from that in earlier accounts. Provides a close analysis

of each of Milton's prose and poetry works. Reveals how Milton was the first writer to self consciously construct himself as an 'author'. Focuses on the development of Milton's ideas and his art.

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