

Commenced Publication in 1973

Founding and Former Series Editors:

Gerhard Goos, Juris Hartmanis, and Jan van Leeuwen

Editorial Board

David Hutchison

Lancaster University, UK

Takeo Kanade

Carnegie Mellon University, Pittsburgh, PA, USA

Josef Kittler

University of Surrey, Guildford, UK

Jon M. Kleinberg

Cornell University, Ithaca, NY, USA

Friedemann Mattern

ETH Zurich, Switzerland

John C. Mitchell

Stanford University, CA, USA

Moni Naor

Weizmann Institute of Science, Rehovot, Israel

Oscar Nierstrasz

University of Bern, Switzerland

C. Pandu Rangan

Indian Institute of Technology, Madras, India

Bernhard Steffen

University of Dortmund, Germany

Madhu Sudan

Massachusetts Institute of Technology, MA, USA

Demetri Terzopoulos

New York University, NY, USA

Doug Tygar

University of California, Berkeley, CA, USA

Moshe Y. Vardi

Rice University, Houston, TX, USA

Gerhard Weikum

Max-Planck Institute of Computer Science, Saarbruecken, Germany

José M. Barreiro
Fernando Martin-Sanchez Víctor Maojo
Ferran Sanz (Eds.)

Biological and Medical Data Analysis

5th International Symposium, ISBMDA 2004
Barcelona, Spain, November 18-19, 2004
Proceedings

Volume Editors

José M. Barreiro
Polytechnical University of Madrid
School of Computer Science
Artificial Intelligence Department
Campus de Montegancedo s/n, 28660 Boadilla del Monte, Madrid, Spain
E-mail: jmbarreiro@fi.upm.es

Fernando Martin-Sanchez
Institute of Health Carlos III
Department of Medical Bioinformatics
Ctra. Majadahonda a Pozuelo, km. 2, 28220 Majadahonda, Madrid, Spain
E-mail: fmartin@isciis.es

Víctor Maojo
Polytechnical University of Madrid
School of Computer Science
Artificial Intelligence Lab
Boadilla del Monte, 28660 Madrid, Spain
E-mail: vmaoj@infomed.dia.fi.upm.es

Ferran Sanz
University Pompeu Fabra
Municipal Institute of Medical Research (IMIM)
Research Unit on Biomedical Informatics (GRIB)
Passeig Marítim de la Barceloneta 37-49
08003 Barcelona, Spain
E-mail: fsanz@imim.es

Library of Congress Control Number: 2004115460

CR Subject Classification (1998): H.2.8, I.2, H.3, G.3, I.5.1, I.4, J.3, F.1

ISSN 0302-9743

ISBN 3-540-23964-2 Springer Berlin Heidelberg New York

This work is subject to copyright. All rights are reserved, whether the whole or part of the material is concerned, specifically the rights of translation, reprinting, re-use of illustrations, recitation, broadcasting, reproduction on microfilms or in any other way, and storage in data banks. Duplication of this publication or parts thereof is permitted only under the provisions of the German Copyright Law of September 9, 1965, in its current version, and permission for use must always be obtained from Springer. Violations are liable to prosecution under the German Copyright Law.

Springer is a part of Springer Science+Business Media

springeronline.com

© Springer-Verlag Berlin Heidelberg 2004
Printed in Germany

Typesetting: Camera-ready by author, data conversion by Olgun Computergrafik
Printed on acid-free paper SPIN: 11362128 06/3142 5 4 3 2 1 0

Preface

This year, the 5th International Symposium on Medical Data Analysis has experimented an apparently slight modification. The word “biological” has been added to the title of the conferences. The motivation for this shift goes beyond the wish to attract a different kind of professional. It is linked to recent trends to produce a shift within various biomedical areas towards genomics-based research and practice. For instance, medical informatics and bioinformatics are being linked in a synergic area denominated biomedical informatics. Similarly, patient care is being improved, leading to concepts and areas such as molecular medicine, genomic medicine or personalized healthcare.

The results from different genome projects, the advances in systems biology and the integrative approaches to physiology would not be possible without new approaches in data and information processing. Within this scenario, novel methodologies and tools will be needed to link clinical and genomic information, for instance, for genetic clinical trials, integrated data mining of genetic clinical records and clinical databases, or gene expression studies, among others.

Genomic medicine presents a series of challenges that need to be addressed by researchers and practitioners. In this sense, this ISBMDA conference aimed to become a place where researchers involved in biomedical research could meet and discuss. For this conference, the classical contents of former ISMDA conferences were updated to incorporate various issues from the biological fields. Similarly to the incorporation of these new topics of the conference, data analysts will face, in this world of genomic medicine and related areas, significant challenges in research, education and practice.

For this conference, after a peer-review process, we selected the scientific papers that are now published in this issue. The editors would like to thank all the participants for their outstanding contributions and Springer for publishing the proceedings of this conference. Finally, we would like to acknowledge the collaboration of Prof. Rüdiger Brause, who was the first chair of the ISMDA conferences. Without his inspiration and support, this idea would not have been possible.

November 2004

Jose M. Barreiro
Victor Maojo
Fernando Martin-Sanchez
Ferran Sanz

Organization

Executive Committee

Chair	Fernando Martin-Sanchez, Institute of Health Carlos III, Spain
Scientific Committee Coordinators	Victor Maojo, Polytechnical Univ. of Madrid, Spain. Ferran Sanz, IMIM-Univ. Pompeu Fabra, Barcelona, Spain

Steering Committee

J.M. Barreiro (Polytechnical Univ. of Madrid, Spain)
R. Brause (J.W.G.-University, Frankfurt, Germany)
M. García-Rojo (Complejo Hospitalario Ciudad Real, Spain)
C. Kulikowski. (Rutgers University, USA)

Scientific Committee

J. Ares (Univ. of A Coruña, Spain)
H. Billhardt (Univ. Rey Juan Carlos, Spain)
V. Breton (CNRS, France)
J.M. Carazo (CNB, Spain)
A. Colosito (Univ. of Rome “La Sapienza”, Italy)
M. Dugas (Univ. of Munich, Germany)
A. Giuliani (Nat. Inst. of Health, Italy)
R. Guthke (Hans-Knoell Inst., Germany)
P. Larranaga (Univ. of the Basque Country, Spain)
N. Lavrac (J. Stefan Institute, Slovenia)
L. Ohno-Machado (Harvard Univ, USA)
E. Medico (IRCC, Italy)
X. Pastor (IDIBAPS, Spain)
A. Pazos (Univ. of A Coruña, Spain)
P. Perner (IBaI Leipzig, Germany)
G. Potamias (Institute of Computer Science, FORTH, Greece)
W. Sauerbrei (Univ. of Freiburg, Germany)
J. Sima (Academy of Sciences of the Czech Republic)
A. Silva (Polytechnical Univ. of Madrid, Spain)
A. Sousa (Univ. Aveiro, Portugal)
B. Zupan (Univ. of Ljubljana, Slovenia)
J. Zvávová (Charles Univ. and Academy of Sciences, Czech Republic)

Local Committee

Members of the Executive Board of SEIS, the Spanish Society of Health Informatics.

Table of Contents

Data Analysis for Image Processing

RF Inhomogeneity Correction Algorithm in Magnetic Resonance Imaging	1
<i>Juan A. Hernández, Martha L. Mora, Emanuele Schiavi, and Pablo Toharia</i>	
Fully 3D Wavelets MRI Compression	9
<i>Emanuele Schiavi, C. Hernández, and Juan A. Hernández</i>	
A New Approach to Automatic Segmentation of Bone in Medical Magnetic Resonance Imaging	21
<i>Gabriela Pérez, Raquel Montes Diez, Juan A. Hernández, and José San Martín</i>	
An Accurate and Parallelizable Geometric Projector/Backprojector for 3D PET Image Reconstruction	27
<i>Roberto de la Prieta</i>	

Data Visualization

EEG Data and Data Analysis Visualization	39
<i>Josef Rieger, Karel Kosar, Lenka Lhotska, and Vladimír Krajča</i>	
A Web Information System for Medical Image Management	49
<i>César J. Acuña, Esperanza Marcos, Valeria de Castro, and Juan A. Hernández</i>	
Reliable Space Leaping Using Distance Template	60
<i>Sukhyun Lim and Byeong-Seok Shin</i>	

Decision Support Systems

A Rule-Based Knowledge System for Diagnosis of Mental Retardation	67
<i>R. Sánchez-Morgado, Luis M. Laita, Eugenio Roanes-Lozano, Luis de Ledesma, and L. Laita</i>	
Case-Based Diagnosis of Dysmorphic Syndromes	79
<i>Tina Waligora and Rainer Schmidt</i>	
Bayesian Prediction of Down Syndrome Based on Maternal Age and Four Serum Markers	85
<i>Raquel Montes Diez, Juan M. Marin, and David Ríos Insua</i>	
SOC: A Distributed Decision Support Architecture for Clinical Diagnosis	96
<i>Javier Vicente, Juan M. García-Gómez, César Vidal, Luís Martí-Bonmati, Aurora del Arco, and Montserrat Robles</i>	

Decision Support Server Architecture for Mobile Medical Applications	105
<i>Marek Kurzynski and Jerzy Sas</i>	
Ordered Time-Independent CIG Learning	117
<i>David Riaño</i>	
SINCO: Intelligent System in Disease Prevention and Control. An Architectural Approach	129
<i>Carolina González, Juan C. Burguillo, Juan C. Vidal, and Martin Llamas</i>	
Could a Computer Based System for Evaluating Patients with Suspected Myocardial Infarction Improve Ambulance Allocation?	141
<i>Martin Gellerstedt, Angela Bång, and Johan Herlitz</i>	
On the Robustness of Feature Selection with Absent and Non-observed Features .	148
<i>Petra Geenen, Linda C. van der Gaag, Willie Loeffen, and Armin Elbers</i>	
Design of a Neural Network Model as a Decision Making Aid in Renal Transplant	160
<i>Rafael Magdalena, Antonio J. Serrano, Agustin Serrano, Jorge Muñoz, Joan Vila, and E. Soria</i>	
Learning the Dose Adjustment for the Oral Anticoagulation Treatment	171
<i>Giacomo Gamberoni, Evelina Lamma, Paola Mello, Piercamillo Pavesi, Sergio Storari, and Giuseppe Trocino</i>	

Information Retrieval

Thermal Medical Image Retrieval by Moment Invariants	182
<i>Shao Ying Zhu and Gerald Schaefer</i>	

Knowledge Discovery and Data Mining

Employing Maximum Mutual Information for Bayesian Classification	188
<i>Marcel van Gerven and Peter Lucas</i>	
Model Selection for Support Vector Classifiers via Genetic Algorithms. An Application to Medical Decision Support	200
<i>Gilles Cohen, Mélanie Hilario, and Antoine Geissbuhler</i>	
Selective Classifiers Can Be Too Restrictive: A Case-Study in Oesophageal Cancer	212
<i>Rosa Blanco, Linda C. van der Gaag, Iñaki Inza, and Pedro Larrañaga</i>	
A Performance Comparative Analysis Between Rule-Induction Algorithms and Clustering-Based Constructive Rule-Induction Algorithms. Application to Rheumatoid Arthritis	224
<i>J.A. Sanandrés-Ledesma, Victor Maojo, Jose Crespo, M. García-Remesal, and A. Gómez de la Cámara</i>	

Domain-Specific Particularities of Data Mining: Lessons Learned	235
<i>Victor Maojo</i>	

Statistical Methods and Tools for Biological and Medical Data Analysis

A Structural Hierarchical Approach to Longitudinal Modeling of Effects of Air Pollution on Health Outcomes	243
<i>Michael Friger, Arkady Bolotin, and Ulrich Ranft</i>	

Replacing Indicator Variables by Fuzzy Membership Functions in Statistical Regression Models: Examples of Epidemiological Studies	251
<i>Arkady Bolotin</i>	

PCA Representation of ECG Signal as a Useful Tool for Detection of Premature Ventricular Beats in 3-Channel Holter Recording by Neural Network and Support Vector Machine Classifier	259
<i>Stanisław Jankowski, Jacek J. Duszka, Mariusz Wierzbowski, and Artur Oręziak</i>	

Finding Relations in Medical Diagnoses and Procedures	269
<i>David Riaño and Ioannis Aslanidis</i>	

An Automatic Filtering Procedure for Processing Biomechanical Kinematic Signals	281
<i>Francisco Javier Alonso, José María Del Castillo, and Publio Pintado</i>	

Analysis of Cornea Transplant Tissue Rejection Delay in Mice Subjects	292
<i>Zdenek Valenta, P. Svozilkova, M. Filipec, J. Zvarova, and H. Farghali</i>	

Toward a Model of Clinical Trials	299
<i>Laura Collada Ali, Paola Fazi, Daniela Luzi, Fabrizio L. Ricci, Luca Dan Serbanati, and Marco Vignetti</i>	

Time Series Analysis

Predicting Missing Parts in Time Series Using Uncertainty Theory	313
<i>Sokratis Konias, Nicos Maglaveras, and Ioannis Vlahavas</i>	

Classification of Long-Term EEG Recordings	322
<i>Karel Kosar, Lenka Lhotska, and Vladimir Krajca</i>	

Application of Quantitative Methods of Signal Processing to Automatic Classification of Long-Term EEG Records	333
<i>Josef Rieger, Lenka Lhotska, Vladimir Krajca, and Milos Matousek</i>	

Semantic Reference Model in Medical Time Series	344
<i>Fernando Alonso, Loïc Martínez, César Montes, Aurora Pérez, Agustín Santamaría, and Juan Pedro Valente</i>	

Control of Artificial Hand via Recognition of EMG Signals	356
<i>Andrzej Wolczowski and Marek Kurzynski</i>	

Bioinformatics:

Data Management and Analysis in Bioinformatics

SEQPACKER: A Biologist-Friendly User Interface to Manipulate Nucleotide Sequences in Genomic Epidemiology	368
<i>Oscar Coltell, Miguel Arregui, Larry Parnell, Dolores Corella, Ricardo Chalmeta, and Jose M. Ordovas</i>	

Performing Ontology-Driven Gene Prediction Queries in a Multi-agent Environment	378
<i>Vassilis Koutkias, Andigoni Malousi, and Nicos Maglaveras</i>	

Protein Folding in 2-Dimensional Lattices with Estimation of Distribution Algorithms	388
<i>Roberto Santana, Pedro Larrañaga, and José A. Lozano</i>	

Bioinformatics: Integration of Biological and Medical Data

Quantitative Evaluation of Established Clustering Methods for Gene Expression Data	399
<i>Dörte Radke and Ulrich Möller</i>	

DiseaseCard: A Web-Based Tool for the Collaborative Integration of Genetic and Medical Information	409
<i>José Luís Oliveira, Gaspar Dias, Ilídio Oliveira, Patrícia Rocha, Isabel Hermosilla, Javier Vicente, Inmaculada Spiteri, Fernando Martin-Sánchez, and António Sousa Pereira</i>	

Biomedical Informatics: From Past Experiences to the Infobiomed Network of Excellence	418
<i>Victor Maojo, Fernando Martin-Sánchez, José María Barreiro, Carlos Diaz, and Ferran Sanz</i>	

Bioinformatics: Metabolic Data and Pathways

Network Analysis of the Kinetics of Amino Acid Metabolism in a Liver Cell Bioreactor	427
<i>Wolfgang Schmidt-Heck, Katrin Zeilinger, Michael Pfaff, Susanne Toepfer, Dominik Driesch, Gesine Pless, Peter Neuhaus, Joerg Gerlach, and Reinhard Guthke</i>	

Model Selection and Adaptation for Biochemical Pathways	439
<i>Rüdiger W. Brause</i>	

NeoScreen: A Software Application for MS/MS Newborn Screening Analysis . . .	450
<i>Miguel Pinheiro, José Luís Oliveira, Manuel A.S. Santos, Hugo Rocha, M. Luis Cardoso, and Laura Vilarinho</i>	

Bioinformatics: Microarray Data Analysis and Visualization

Technological Platform to Aid the Exchange of Information and Applications

Using Web Services 458
Antonio Estruch and José Antonio Heredia

Visualization of Biological Information with Circular Drawings 468
Alkiviadis Symeonidis and Ioannis G. Tollis

Gene Selection Using Genetic Algorithms 479
Bruno Feres de Souza and André C.P.L.F. de Carvalho

Knowledgeable Clustering of Microarray Data 491
George Potamias

Correlation of Expression Between Different IMAGE Clones
from the Same UniGene Cluster 498
*Giacomo Gamberoni, Evelina Lamma, Sergio Storari, Diego Arcelli,
Francesca Francioso, and Stefano Volinia*

Author Index 507