Lecture Notes in Artificial Intelligence 5108

Edited by R. Goebel, J. Siekmann, and W. Wahlster

Subseries of Lecture Notes in Computer Science

Petra Perner Ovidio Salvetti (Eds.)

Advances in Mass Data Analysis of Images and Signals in Medicine, Biotechnology, Chemistry and Food Industry

Third International Conference, MDA 2008 Leipzig, Germany, July 14, 2008 Proceedings



Series Editors

Randy Goebel, University of Alberta, Edmonton, Canada Jörg Siekmann, University of Saarland, Saarbrücken, Germany Wolfgang Wahlster, DFKI and University of Saarland, Saarbrücken, Germany

Volume Editors

Petra Perner IBaI - Institute of Computer Vision and Applied Computer Sciences Arno-Nitzsche-Str. 43, 04277 Leipzig, Germany E-mail: pperner@ibai-institut.de

Ovidio Salvetti Istituto di Scienza e Tecnologie dell'Informazione (ISTI) Area della Ricerca CNR di Pisa Via G. Moruzzi 1, 56124 Pisa, Italy E-mail: ovidio.salvetti@isti.cnr.it

Library of Congress Control Number: 2008930273

CR Subject Classification (1998): H.2.8, I.4.6, J.3

LNCS Sublibrary: SL 7 – Artificial Intelligence

ISSN 0302-9743

ISBN-10 3-540-70714-X Springer Berlin Heidelberg New York ISBN-13 978-3-540-70714-1 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

springer.com

© Springer-Verlag Berlin Heidelberg 2008 Printed in Germany

Typesetting: Camera-ready by author, data conversion by Scientific Publishing Services, Chennai, India Printed on acid-free paper SPIN: 12441549 06/3180 543210

Preface

The automatic analysis of signals and images together with the characterization and elaboration of their representation features is still a challenging activity in many relevant scientific and hi-tech fields such as medicine, biotechnology, and chemistry. Multidimensional and multisource signal processing can generate a number of information patterns which can be useful to increase the knowledge of several domains for solving complex problems. Furthermore, advanced signal and image manipulation allows relating specific application problems into pattern recognition problems, often implying also the development of KDD and other computational intelligence procedures.

Nevertheless, the amount of data produced by sensors and equipments used in biomedicine, biotechnology and chemistry is usually quite huge and structured, thus strongly pushing the need of investigating advanced models and efficient computational algorithms for automating mass analysis procedures. Accordingly, signal and image understanding approaches able to generate automatically expected outputs become more and more essential, including novel conceptual approaches and system architectures.

The purpose of this third edition of the International Conference on Mass Data Analysis of Signals and Images in Medicine, Biotechnology, Chemistry and Food Industry (MDA 2008; www.mda-signals.de) was to present the broad and growing scientific evidence linking mass data analysis with challenging problems in medicine, biotechnology and chemistry. Scientific and engineering experts convened at the workshop to present the current understanding of image and signal processing and interpretation methods useful for facing various medical and biological problems and exploring the applicability and effectiveness of advanced techniques as solutions.

The primary goal of the conference was to disseminate this knowledge to a multidisciplinary community and encourage cooperative proactive collaboration in all the interested fields.

We were pleased to see that the idea of the conference was taken up by a growing number of researchers and that we could start to bundle the activities in this area.

We appreciate the help and understanding of the editorial staff at Springer, and in particular Alfred Hofmann, who supported the publication of these proceedings in the LNAI series.

Last, but not least, we wish to thank all the speakers and participants who contributed to the success of the conference.

The next International Conferences on Mass Data Analysis of Signals and Images (www.mda-signals.de) will be held in July 2009. We are looking forward to your submissions.

July 2008 Petra Perner
Ovidio Salvetti

International Conferences on Mass Data Analysis of Signals and Images in Medicine, Biotechnology, Chemistry, and Food Industry, MDA 2008

July 14, 2008, Leipzig, Germany

Organization

Institute of Computer Vision and applied Computer Sciences, IBaI, Germany

Chairs

Petra Perner IBaI, Germany Ovidio Salvetti CNR-ISTI, Italy

Committee

Walter Arnold Fraunhofer Institute of Non-destructive Testing, Germany

Ewert Bengsston University of Uppsala, Sweden Valentin Brimkov Buffalo State College, USA Hans du Buf University of Algarve, Portugal

Eugenio Fava Max Planck Institute of Molecular Cell Biology &

Genetics, Germany

Maria Frucci Istituto di Cibernetica, CNR, Italy Igor Gurevich Academy of Science, Russia Thomas Günther JenaBios GmbH, Germany

Giulio Iannello University Campus Bio-Medico of Rome, Italy

Xiaoyi Jiang University of Muenster, Germany

Montse Pardas Universitat Politècnica de Catalunya, Spain

Thang Viet Pham OncoProteomics Laboratory - VUmc, The Netherlands

Gabriella Saniti di Baja Istituto di Cibernetica, CNR, Italy

Arnold Smeulders University of Amsterdam, The Netherlands

Tuan Pham James Cook University, Australia Julie Wilson York Structural Biology Laboratory, UK

Aim of Conference

The automatic analysis of images and signals in medicine, biotechnology, and chemistry is a challenging and demanding field.

Signal-producing procedures by microscopes, spectrometers and other sensors have found their way into wide fields of medicine, biotechnology, economy and environmental analysis. With this arises the problem of the automatic mass analysis of

signal information. Signal-interpreting systems which generate automatically the desired target statements from the signals are therefore of compelling necessity. The continuation of mass analyses on the basis of the classical procedures leads to investments of proportions that are not feasible. New procedures and system architectures are therefore required.

Scope of Conference

The scope of the International Conference on Mass Data Analysis of Images and Signals in Medicine, Biotechnology, Chemistry and Food Industry (www.mda-signals.de) is to bring together researcher, practitioners and industry people who deal with mass analysis of images and signals to present and discuss recent research in these fields.

The goals of this workshop are to:

- Provide a forum for identifying important contributions and opportunities for research on mass data analysis on microscopic images
- Promote the systematic study of how to apply automatic image analysis and interpretation procedures to that field
- Show case applications of mass data analysis in biology, medicine, and chemistry

Topics

Topics of interest include (but are not limited to):

- Techniques and developments of signal and image producing procedures
- Object matching and object tracking in microscopic and video microscopic images
- 1D, 2D and 3D shape analysis and description
- 1D, 2D and 3D feature extraction of texture, structure and location
- Algorithms for 1D, 2D and 3D signal analysis and interpretation
- Image segmentation algorithms
- Parallelization of image analysis and interpretation algorithms
- Semantic tagging of images from life science applications
- Applications in medicine, biotechnology, chemistry and others
- Applications in crystallography
- Applications in proteomics
- Applications in 2D and 3D cell images analysis
- Image acquisition procedures for mass data analysis

Table of Contents

Calin Ciufudean, Otilia Ciufudean, Constantin Filote	1
User Assisted Substructure Extraction in Molecular Data Mining Burcu Yılmaz, Mehmet Göktürk, and Natalie Shvets	12
Fully Automatic Heart Beat Rate Determination in Digital Video Recordings of Rat Embryos	27
Biomedical Signal and Image Processing for Decision Support in Heart Failure	38
Automatic Data Acquisition and Signal Processing in the Field of Virology	52
Colorectal Polyps Detection Using Texture Features and Support Vector Machine	62
OplAnalyzer: A Toolbox for MALDI-TOF Mass Spectrometry Data Analysis	73
Classification of Mass Spectrometry Based Protein Markers by Kriging Error Matching	82
A Mathematical Operator for Automatic and Real Time Analysis of Sequences of Vascular Images	95
A Unified Mathematical Treatment of Regression Problems in Image Processing	108

X Table of Contents

Multi-scale Representation and Persistency for Shape Description Davide Moroni, Mario Salvetti, and Ovidio Salvetti	123
Novel Computerized Methods in System Biology – Flexible High-Content Image Analysis and Interpretation System for Cell Images	139
MDA 2006	
Automatic Segmentation of Unstained Living Cells in Bright-Field Microscope Images	158
Author Index	173