## Mathematics and Visualization

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# Visualization in Medicine and Life Sciences II

Progress and New Challenges

With 132 Figures, 116 in color



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### **Preface**

Medicine has for a long time been a major driver for the development of data processing and visualization techniques. Many exciting and challenging visualization problems are continually arising in medicine as a consequence of our ability to generate increasingly large and complicated data (image data, simulated data) that require us to devise effective tools for meaningful interpretation and utilization in medical practice. The first VMLS workshop, which led to the book entitled "Visualization in Medicine and Life Sciences (VMLS)," was driven by the fact that emerging technologies in the life sciences had produced significant data visualization challenges. One interesting question was: Can medical data visualization approaches be devised and/or improved to meet these challenges with the promise of ultimately being adopted by medical experts.

Life sciences are understood by us in a broad sense, including animal and human biology, biochemistry, bioinformatics, biomathematics, food sciences, environmental sciences, and pharmacology. Different data acquisition technologies lead to different types of data, including both spatial and non-spatial data. The aim of the second international VMLS workshop was to document and discuss the progress that had been made since the first workshop and to explore what novel solution approaches for data processing and visualization had been developed and what new challenges had come up.

Internationally leading experts from the visualization and driving medical application areas came together for this second workshop held in Bremerhaven, Germany, in July 2009. Research and survey papers were solicited and peerreviewed, ultimately leading to the collection of papers included in this book.

The research topics covered by the papers in this book deal with these themes:

- Feature Extraction
- Classification
- Volumes and Shapes
- Tensor Visualization
- Visualizing Genes, Proteins, and Molecules

vi Preface

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Bremen, Germany Kaiserslautern, Germany Davis, California, U.S.A. Berlin, Germany Lars Linsen Hans Hagen Bernd Hamann Hans-Christian Hege

#### **Contents**

Part I Feature Extraction

#### Discrete Distortion for 3D Data Analysis ..... 3 Leila De Floriani, Federico Iuricich, Paola Magillo, Mohammed Mostefa Mesmoudi, and Kenneth Weiss Interactive Visualization-A Key Prerequisite for Reconstruction and Analysis of Anatomically Realistic Neural Networks ..... 27 Vincent J. Dercksen, Marcel Oberlaender, Bert Sakmann, and Hans-Christian Hege MRI-Based Visualisation and Ouantification of Rheumatoid and Psoriatic Arthritis of the Knee..... 45 Ben Donlon, Douglas Veale, Patrick Brennan, Robert Gibney, Hamish Carr, Louise Rainford, ChinTeck Ng, Eliza Pontifex, Jonathan McNulty, Oliver FitzGerald, and John Ryan An Application for the Visualization and Quantification of HIV-Associated Lipodystrophy from Magnetic Resonance Imaging Datasets ..... 61 Tadhg O'Sullivan, Patrick Brennan, Peter Doran, Paddy Mallon, Stephen J. Eustace, Eoin Kavannagh, Allison Mcgee, Louise Rainford, and John Ryan Part II Classification Semi-Automatic Rough Classification of Multichannel Medical

Imaging Data .....

Ahmed Elmoasry, Mohamed Sadek Maswadah, and Lars Linsen

71

viii Contents

An Evaluation of Peak Finding for DVR Classification of Biological Data	91		
Aaron Knoll, Rolf Westerteiger, and Hans Hagen			
Part III Volumes and Shapes			
Vessel Visualization with Volume Rendering			
Efficient Selection of Representative Views and Navigation Paths for Volume Data Exploration Eva Monclús, Pere-Pau Vázquez, and Isabel Navazo	133		
Feature Preserving Smoothing of Shapes Using Saliency Skeletons Alexandru Telea	153		
Part IV Tensor Visualization			
Enhanced DTI Tracking with Adaptive Tensor Interpolation	173		
Image-Space Tensor Field Visualization Using a LIC-like Method Sebastian Eichelbaum, Mario Hlawitschka, Bernd Hamann, and Gerik Scheuermann	191		
Towards a High-quality Visualization of Higher-order Reynold's Glyphs for Diffusion Tensor Imaging  Mario Hlawitschka, Younis Hijazi, Aaron Knoll, and Bernd Hamann			
Part V Visualizing Genes, Proteins, and Molecules			
VENLO: Interactive Visual Exploration of Aligned Biological Networks and Their Evolution Steffen Brasch, Georg Fuellen, and Lars Linsen	229		
<b>Embedding Biomolecular Information in a Scene Graph System</b>	249		
Linking Advanced Visualization and MATLAB for the Analysis of 3D Gene Expression Data Oliver Rübel, Soile V.E. Keränen, Mark Biggin, David W. Knowles, Gunther H. Weber, Hans Hagen, Bernd Hamann, and E. Wes Bethel			
Index	285		