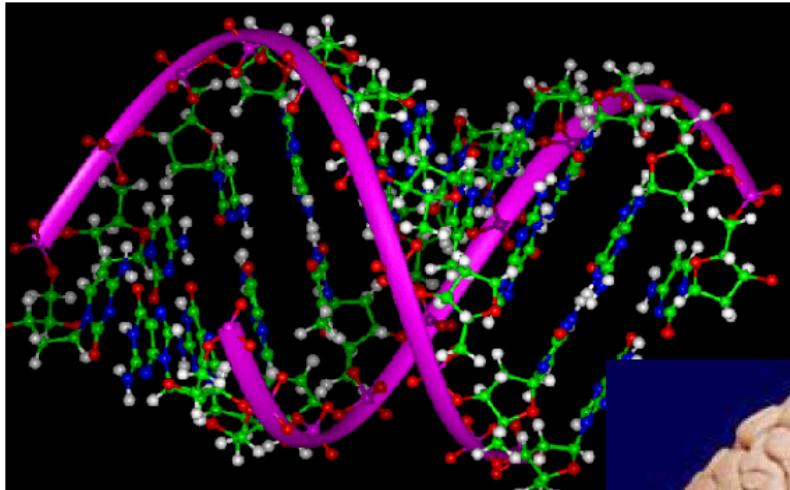
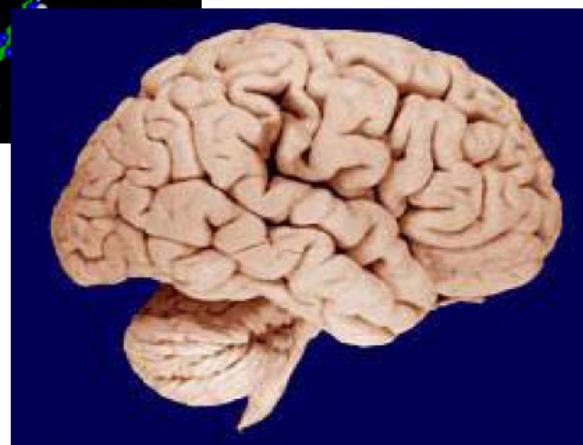


# Proceedings of the 7<sup>th</sup> IEEE International Conference on Bioinformatics and Bioengineering



***IEEE  
BIBE 2007***

*The Conference Center at  
Harvard Medical School  
Boston, Massachusetts USA  
October 14 - 17, 2007*



## Volume I

### Editors

**Jack Y. Yang, Mary Qu Yang, Michelle M. Zhu  
Yanqing Zhang, Hamid R. Arabnia, Youping Deng and Nikolaos G. Bourbakis**

Library of Congress 2007904763

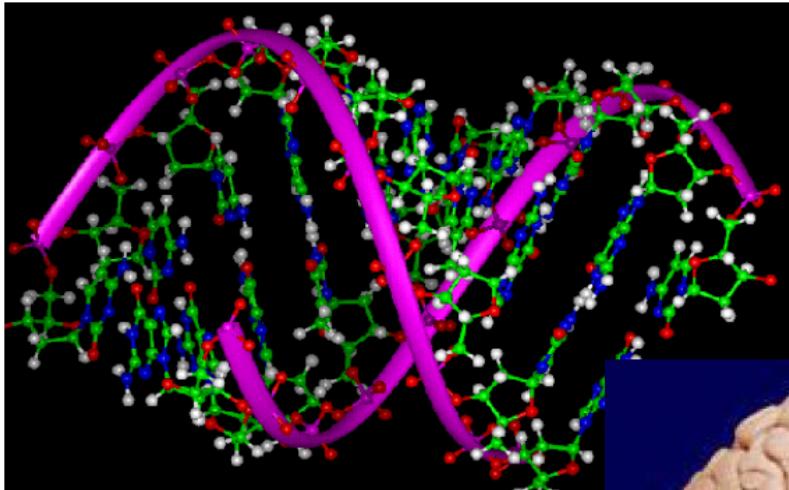
ISBN 1-4244-1509-8

IEEE Catalog Number 07EX1893

IEEE Press

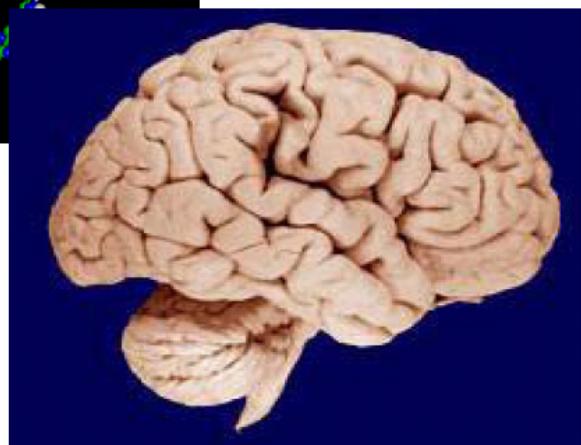


# Proceedings of the 7<sup>th</sup> IEEE International Conference on Bioinformatics and Bioengineering



***IEEE  
BIBE 2007***

*The Conference Center at  
Harvard Medical School  
Boston, Massachusetts USA  
October 14 - 17, 2007*



## Volume II

### Editors

**Jack Y. Yang, Mary Qu Yang, Michelle M. Zhu  
Yanqing Zhang, Hamid R. Arabnia, Youping Deng and Nikolaos G. Bourbakis**

Library of Congress 2007904763

ISBN 1-4244-1509-8

IEEE Catalog Number 07EX1893

IEEE Press



## Message from the General, Program and Steering Committee Chairs

On behalf of the committees of the 7<sup>th</sup> IEEE International Conference on Bioinformatics and Bioengineering (IEEE BIBE), we are delighted to welcome you to the conference center at Harvard Medical School, Boston, Massachusetts in October 14-17, 2007.

Bioinformatics and Bioengineering are complementary disciplines that hold great promise for the advancement of research and development in complex biomedical systems, agricultural, environmental, pharmaceutical and medical sciences as well as public health, drug designs, genomics and so on. The 7<sup>th</sup> IEEE BIBE provides a common platform for the cross fertilization of ideas, to help shape knowledge and scientific achievements by bridging these two very important and complementary disciplines into an interactive and attractive forum. The 7<sup>th</sup> IEEE BIBE is a large international conference with 7 years of tradition and leading reputation.

The synergy of bioinformatics and bioengineering has been well received by researchers. This year, the 7<sup>th</sup> IEEE BIBE has received more than 600 high-quality full research papers. Each paper was reviewed and ranked by at least 4 professors and scientists in the program and scientific review committee. As a result of the rigorous review process and stringent evaluation, the committee selected 66 papers as full regular research papers, 58 special interest research papers, 23 workshop research papers, and 77 regular research short papers. Lastly, there are 14 keynote and tutorial lecture papers and abstracts from world renowned scientists in the fields that are within the scope of the 7<sup>th</sup> IEEE BIBE. The acceptance rate for the full regular research papers is less than 12%. In addition to the 600+ submissions of full-length research papers, there are also large numbers of poster abstract submissions. The 272 professors and scientists in the program and scientific review committee have spent tremendous amount of time and efforts for this academic event in evaluating all submitted papers. The anticipated participation exceeds 500 people, most of them from academia, medical and research centers. All accepted papers (except for poster papers and abstracts) are published in the IEEE proceeding books indexed in EI, INSPEC, DBLP and Library of Congress. The conference also offers a supplementary journal issue of BMC Genomics (Impact Factor 4.09, PubMed/Medline and SCI indexed). The 7<sup>th</sup> IEEE BIBE is a large international gathering that is aimed to promote synergistic inter/multidisciplinary research and education to solve important but difficult biomedical problems that need both expertise in biomedical sciences and engineering.

Major organizers of the 7<sup>th</sup> IEEE BIBE who hold formal degrees in the fields of both engineering and biophysical/biomedical sciences with many years of research, teaching and engineering practice experiences realized the power and needs of synergistic engineering and biomedical research based on their professional experience. They and all other organizers are dedicated to promote such inter/multidisciplinary research and education. Therefore, the 7<sup>th</sup> IEEE BIBE has been designed to be responsive to the need for synergistic inter/multidisciplinary research and education. For that reason, it is the only meeting whose components are also defined dynamically in response to specific needs of largest number of keynote lectures, cutting-edge research tutorial lectures, special interest research workshops and special sessions with academic supports and contributions from leading scientists at NIH, national laboratories and research universities. Each proposal and nomination has been reviewed and voted by the 7<sup>th</sup> IEEE BIBE committee to ensure that participants will be benefited significantly from the academic event. We are very grateful for the IEEE BIBE Plenary Keynote Lectures given by:

Dr. A. Keith Dunker  
Dr. Mary Qu Yang

(Indiana University and Purdue University),  
(National Human Genome Research Institute, National Institutes  
of Health, U.S. Department of Health and Human Services and  
Oak Ridge, DOE),

Dr. John Quackenbush  
Dr. Andy Baxevanis

(Harvard School of Public Health and Dana-Farber Cancer Institute),  
(National Human Genome Research Institute, National Institutes of

Dr. Steven E. Seltzer  
Dr. Tony Xiaohua Hu  
Dr. Xudong Huang  
Dr. Yi Pan  
Dr. Jay Steve Loeffler  
Dr. Jun S. Liu  
Dr. Ferenc A. Jolesz  
Dr. Ruzena Bajcsy

Health, United States Department of Health and Human Services), (Harvard Medical School and Brigham & Women's Hospital), (Drexel University and the IEEE Computer Society), (Harvard Medical School and Brigham & Women's Hospital), (Georgia State University), (Harvard Medical School and Massachusetts General Hospital), (Harvard University and Massachusetts Institute of Technology), (Harvard Medical School and Brigham & Women's Hospital), (University of California at Berkeley and Member of United States National Academy of Engineering and, Member of United States Institute of Medicine of the National Academies).

And the Advanced Research Frontier and Education Tutorial Lectures given by:

Dr. Mark Borodovsky  
Dr. Vladimir N. Uversky  
Dr. Patrick S. P. Wang  
Dr. Arif Ghafoor  
Dr. David Lightfoot  
Dr. Igor B. Zhulin  
Dr. May D. Wang  
Dr. Yehuda Braiman  
Dr. Craig A. Stewart  
Dr. Tian Zheng  
Dr. Larry O. Hall

(Georgia Institute of Technology),  
(Indiana University School of Medicine),  
(Northeastern University and Massachusetts Institute of Technology)  
(Purdue University, West Lafayette),  
(Southern Illinois University, Carbondale),  
(Oak Ridge National Laboratory, DOE and University of Tennessee),  
(Georgia Institute of Technology and Emory University),  
(Oak Ridge National Laboratory, U.S. Department of Energy),  
(Indiana University, Bloomington),  
(Columbia University),  
(University of South Florida and the IEEE Systems, Man and Cybernetics Society).

The 7<sup>th</sup> IEEE BIBE brings together top researchers from the United States and around the world to exchange research results and address open issues in all aspects of bioinformatics and bioengineering. The IEEE conference hosts a number of cutting-edge research workshops in collaboration with the National Human Genome Research Institute (NHGRI) and National Cancer Institute (NCI), National Institutes of Health (NIH), U.S. Department of Health and Human Services, Oak Ridge National Laboratory, U.S. Department of Energy, Indiana University, Purdue University, Georgia Institute of Technology, Southern Illinois University, University of Illinois, University of Michigan, University of California, University of Massachusetts, Harvard Medical School, Harvard University and a number of other national laboratories and major research universities. We are especially grateful for the Cutting-Edge Research Workshops and the IEEE BIBE Plenary Keynote Lectures given by:

Dr. Laura L. Elnitski  
Dr. Brian D. Athey  
Dr. Linda Molnar  
Dr. Eric Jakobsson  
Dr. Chih-Ming Ho

(National Human Genome Research Institute, National Institutes of Health (NIH), U. S. Department of Health and Human Services),  
(University Michigan and the National Center for Integrative Biomedical Informatics),  
(National Cancer Institute, National Institutes of Health (NIH), United States Department of Health and Human Services),  
(University of Illinois at Urbana-Champaign and the National Center for Design of Biomimetic Nanoconductors  
(University of California at Los Angles and Member of United States National Academy of Engineering and Academician of Academia Sinica).

Obviously, the conference would not have achieved such a great success without the hard work and voluntary efforts by many contributors. Organizing such a major academic event in the fields is not possible without contributions from members of program and scientific review committee. Thanks must be given to them for their professionalisms. We must express our sincere gratitude to Mary Qu Yang, Vladimir N. Uversky, Yunlong Liu and the program and scientific review committee members for their high-quality timely evaluation of more than 600 full-length regular research papers, as well as an additional large set of poster submissions. We must extend our sincere thanks to all chairs, organizers and committee members (names and affiliations are listed below) for their dedications and professional services. In particular, Hamid R. Arabnia, Walker Land, Jr., Homayoun Valafar and the award committee members dedicated themselves to the unbiased evaluation and selection of the best papers, and the NSF student travel fellowships; Yanqing Zhang managed the paper submission system and handled various important organizing and academic affairs; Youping Deng handled not only registration and finance but also reviewed a large number of papers; Alex Zelikovsky, Yuehui Chen and Yufang Jin managed the overall publication matters; Jonathan Jesneck and Pengyu Hong helped local arrangements; Zejin Jason Ding, Qingzhong Liu and Bingxin Shen maintained the website. Walker Land, Guo-Zheng Li, Hamid R. Arabnia, Youping Deng, Michelle M. Zhu and Jack Y. Yang helped Scientific Review Committee Co-Chairs in the overall systematic evalutions of all reviewers' comments and ranks. Nikolaos G. Bourbakis, Michelle M. Zhu and Heng Huang helped in the overall organization of the conference and Michelle M. Zhu arranged the scientific presentations and program schedules; Jack Y. Yang initiated the organization of the 7<sup>th</sup> IEEE BIBE and guided the committees as well as managed the overall infrastructure of the IEEE conference. Mary Qu Yang and Jack Y. Yang initiated not only the journal issues but also the NSF proposals that provided funds to support students' travel; this is the first time that IEEE BIBE is now not only offering leading scientific journal issues, but also receiving grants and support from the National Science Foundation. Mary Qu Yang and Jack Y. Yang reformed the traditional IEEE BIBE with new components that are defined dynamically in response to specific needs of inter/multidisciplinary cutting-edge research and education, therefore, Mary Qu Yang and Jack Y. Yang initiated and arranged the organization of cutting-edge research workshops, tutorial lectures, special sessions and poster presentations in addition to the traditional keynote lectures. Mary Qu Yang and Jack Y. Yang took the initiatives and invested significant efforts to expand the size of IEEE BIBE from traditional 100-200 papers each year for past 6 years to more than 600 full-length regular research paper submissions this year.

The 7<sup>th</sup> IEEE BIBE committees would like to acknowledge our appreciation of the Institute of Electrical and Electronics Engineers (IEEE), IEEE Systems, Man and Cybernetics Society, IEEE Engineering in Medicine and Biology Society, the Biological and Artificial Intelligence Society (BAIS), the International Society of Intelligent Biological Medicine (ISIBM) and the National Science Foundation (NSF) for their academic supports and sponsorships, as well as the support and collaboration from the IEEE Computer Society. The 7<sup>th</sup> IEEE BIBE is 100% fully sponsored both financially and academically by the IEEE. Finally, we would like to express our sincere appreciation to the excellent professional services provided to the 7<sup>th</sup> IEEE BIBE by Elizabeth Aubrey, an IEEE officer. Once again, welcome to the 7<sup>th</sup> IEEE BIBE. We encourages participants to consider submitting proposals for workshops, special interest research sessions and tutorials in order to ensure that the conference continuously plays the leadership role in promoting inter/multidisciplinary research and education in the fields. We hope your experience at the conference center at Harvard Medical School, Boston, Massachusetts during October 14-17, 2007 would be pleasant, productive, and memorable.

Jack Y. Yang , General Chair

Michelle M. Zhu , Yanqing Zhang, Hamid R. Arabnia and Youping Deng, Program Chairs

Mary Qu Yang , Scientific Review, Advisory and Steering Committee Chair

Nikolas G. Bourbakos , Chair of Founding Steering Committee

## **The 7<sup>th</sup> IEEE BIBE Organization**

Jack Y. Yang

General Chair and Coordinator of Committees on the IEEE 7th BIBE  
Harvard University, Harvard Medical School, USA

Mary Qu Yang

Scientific Review, Advisory, and Steering Committee Chair  
National Human Genome Research Institute,  
National Institutes of Health (NIH), U.S. Department of Health and Human Services,  
and Oak Ridge, DOE, USA

Nikolaos G. Bourbakis

Chair of Founding Steering Committee  
Wright State University, USA

Michelle M. Zhu

Program Chair

Secretary-General of the International Society of Intelligent Biological Medicine  
Oak Ridge National Laboratory, DOE and Southern Illinois University, USA

Yanqing Zhang

Program Co-Chair and Bioinformatics Track Chair  
Georgia State University, USA

Hamid R. Arabnia

Program Co-Chair and Award Committee Chair  
Vice-President of the International Society of Intelligent Biological Medicine  
Head of Panel on Best Papers & Travel Support  
University of Georgia, USA

Youping Deng

Program Co-Chair and Registration/Budget Committee Chair  
Vice-President of the International Society of Intelligent Biological Medicine  
University of Southern Mississippi, USA

George T. Y. Chen

Program Co-Chair and Bioengineering Track Chair  
Harvard University, Harvard Medical School and Massachusetts General Hospital, USA

Lawrence O. Hall

President of IEEE/SMC  
University of South Florida, USA

Zhi-Pei Liang

Co-Chair of Advisory Committee  
IEEE/EMBS Vice-President for Conferences  
University of Illinois at Urbana Champaign, USA

A. Keith Dunker  
Indiana University and Purdue University, USA  
Ying Xu  
University of Georgia and Oak Ridge National Laboratory, DOE, USA  
Co-Chairs of Workshop Proposal Committee

Organizers of Workshop on Bio-Nano-Info Integration for Personalized Medicine  
May D. Wang, Georgia Institute of Technology and Emory University School of Medicine.  
Linda Molnar, National Cancer Institute, NIH, U.S. Dept. of Health and Human Services, USA.  
Eric Jakobsson, National Center for Design of Biomimetic Nanoconductors, and National Center  
for Supercomputing Applications, University of Illinois at Urbana-Champaign, USA

Organizer of Workshop on Progress toward Petascale Applications in Bioinformatics and  
Computational Biology  
Craig A. Stewart and Malinda Lingwall, Indiana University, Bloomington, USA  
David Bader, Georgia Institute of Technology

Organizers of Workshop on Joint Research in the Southern Illinois University, University of Illinois,  
and Oak Ridge National Laboratory, U.S. Department of Energy, U.S.A.  
Michelle M. Zhu and William P. Osborne, Southern Illinois University, USA  
Laura L. Elnitski, National Human Genome Research Institute, NIH, U.S. Department of Health  
Zhi-Pei Liang, IEEE/EMBS Vice-President for Conference, University of Illinois, USA  
Igor B. Zhulin, University of Tennessee and Oak Ridge National Laboratory, DOE, USA  
Yehuda Braiman, Oak Ridge National Lab, DOE, USA

Organizers of 7<sup>th</sup> IEEE BIBE Special Workshop - The 4th Annual University of Massachusetts  
Bioinformatics Conference (folded into main conference)  
Georges Grinstein and Ken Marx, University of Massachusetts Lowell, USA

Organizers of Pacific Bioinformatics and Biomedical Engineering Research Workshop  
(folded into main conference)  
Ron Kikinis, Harvard University, Harvard Medical School and BWH, USA  
Yuehui Chen, University of Jinan, China  
J. Paul Robinson, Purdue University, West Lafayette, USA  
Andrew H. Sung, New Mexico Tech., USA  
Guo-Zheng Li, Shanghai University, China

Organizer of Special Session on Sequence Alignment and Phylogenetic Analysis.  
Mark Clement, Brigham Young University, USA

Organizers of Special Session on DNA Microarray Data Analysis.  
Chris Ding, Lawrence Berkeley National Laboratory and University of California, USA  
Heng Huang, University of Texas at Arlington, USA

Organizers of Special Session on Research in Bioinformatics, Neuroinformatics, and Systems Biology  
in East Asia.  
Hyunsoo Kim, Georgia Institute of Technology, USA  
Sun Kim, Indiana University, Bloomington, USA  
Doheon Lee, Korea Advanced Institute of Science and Technology, Korea

Organizers of Special Session on Bio-Medical Soft Computing  
Qingzhong Liu, Srinivas Mukkamala and Andrew H. Sung, New Mexico Tech., USA

Organizers of Special Session: Developing Algorithms for Solving Problems in Molecular Biology and Bio-Medicine.

Guo-Zheng Li, Shanghai University, China  
Saurabh Sinha, University of Illinois at Urbana-Champaign, USA

Organizer of Special Session on High-throughput Data Analysis for Genomics and Proteomics.  
Jean Gao, University of Texas at Arlington, USA.

Organizers of Special Session on Computational Intelligence in Medical Informatics.  
Jianlin Cheng, University of Missouri, USA  
Jijun Tang, University of South Carolina, USA

Organizers of Special Session on Bio-Complexity.  
Matthias Dehmer, Max F. Perutz Laboratories, Vienna Bio Center, Austria  
Frank Emmert-Streib, Washington University, USA

Organizers of Special Session on Evolutionary Systems Biology.  
Xun Gu, Iowa State University, USA  
Yufeng Wang, University of Texas at San Antonio, USA

Organizer of Special Session on Pattern Recognition and Gene Discovery in Molecular Genetics.  
Sridhar Ramachandran, Indiana University, Southeast, USA

Organizer of Special Session on Machine Learning Methods in Structural/Functional Genomics.  
Rui Kuang, University of Minnesota, USA

Organizers of Special Session on Biomedical Engineering for Handicapped Individuals  
Nikolaos G. Bourbakis, Wright State University, USA  
Jack Y. Yang, Harvard University, Harvard Medical School, USA

Inter/Multidisciplinary Chair  
Mary Qu Yang, National Human Genome Research Institute, NIH, and Oak Ridge, DOE, USA

Co-Chairs of Publication Committee  
Yuehui Chen, University of Jinan, China  
Yufang Jin, University of Texas at San Antonio, USA  
Alex Zelikovsky, Georgia State University, USA

Co-Chairs of Award Committee  
Hamid R. Arabnia, University of Georgia, USA  
Walker Land, Jr., Binghamton University, USA  
Homayoun Valafar, University of South Carolina, Columbia, USA

Head of Panel on Best Papers and NSF Travel Fellowships  
Hamid R. Arabnia, University of Georgia, USA

**Co-Chairs of Tutorial Proposal Committee**  
Joydeep Ghosh, University of Texas at Austin, USA  
Vasile Palade, Oxford University, United Kingdom  
My Tra Thai, University of Florida, USA

**Poster Chairs**  
Chung-Kuan Cheng, University of California at San Diego, USA  
Jonathan Jesneck, Broad Institute of Massachusetts Institute of Technology and Harvard, USA  
Michael L. Raymer, Wright State University, USA

**Publicity Chairs**  
Okan K. Ersoy, Purdue University, West Lafayette, USA  
Lihua Li, Hangzhou Dianzi University, China  
Yingshu Li, Georgia State University, USA  
Liqiang Zhang, Indiana University, South Bend USA

**Web Chairs**  
Zejin Jason Ding, Georgia State University, USA  
Qingzhong Liu, New Mexico Tech, USA  
Binxin Shen, Brookhaven National Lab and Stony Brook University, USA

**Industry Chairs**  
Kevin Daimi, University of Detroit Mercy, USA  
Deanne Taylor, Harvard School of Public Health, Harvard University, USA

**International Chair (Visa Request Chair)**  
Anu Bourgeois, Georgia State University, USA

**Local Chairs**  
Gil Alterovitz, Harvard Medical School and Massachusetts Institute of Technology, USA  
Marek Ancukiewicz, Harvard Medical School and Massachusetts General Hospital, USA  
Jonathan Jesneck, Harvard Medical School and Massachusetts Institute of Technology, USA  
Pengyu Hong, Brandeis University and National Center of Behavioral Genomics, USA  
Marco Ramoni, Harvard Medical School and Massachusetts Institute of Technology, USA

**Organizing Chair**  
Heng Huang, University of Texas at Arlington, USA

**Program Vice-Chairs**  
Yuehui Chen, University of Jinan, USA  
Jun Ni, University of Iowa USA  
Vladimir N. Uversky, Indiana University School of Medicine, USA  
Patrick S. Wang, Northeastern University, USA

**Co-Chairs of Scientific Review Committee**  
Yunlong Liu, Indiana University Purdue University, USA  
Vladimir Uversky, Indiana University School of Medicine, USA  
Mary Qu Yang, National Human Genome Research Institute, NIH, & Oak Ridge, DOE, USA

**Co-Chairs of Advisory Committee**

Zhi-Pei Liang, IEEE/EMBS Vice-President for Conferences, University of Illinois, USA.

Mary Qu Yang, National Human Genome Research Institute, NIH, and Oak Ridge, DOE, USA

**Members of Program and Scientific Review Committee**

1. Purang Abolmaesumi, Queen's University, Canada
2. Ajith Abraham, Chung-Ang University, South Korea
3. Osman Abul, TOBB University of Economics and Technology, Turkey
4. R. Acharya, Pennsylvania State University, USA
5. Reda Alhajj, University of Calgary, Canada
6. Gil Alterovitz, Massachusetts Institute of Technology and Harvard University, USA
7. Sergio A. Alvarez, Boston College, USA
8. Marek Ancukiewicz, Harvard University, Harvard Medical School and MGH, USA
9. Neculai Archip, Harvard University, USA
10. Babak Akhgar, Sheffield Hallam University, U.K.
11. Hisham Al-Mubaid, University of Houston- Clear Lake, USA
12. Hamid R. Arabnia, University of Georgia, USA
13. Brian D. Athey, University of Michigan, USA.
14. Fred Azar, University of Pennsylvania, USA
15. David Bader, Georgia Tech, USA
16. Ruzena Bajcsy, University of California-Berkeley, USA
17. Chengpeng Bi, University of Missouri Kansas City & Children's Mercy Hospitals, USA
18. Emad Boctor, Johns Hopkins University, USA
19. Nazeih Botros, Southern Illinois University, Carbondale, USA
20. Nikolaos G. Bourbakis, Wright State University, USA
21. Anu Bourgeois, Georgia State University, USA
22. Yehuda Braiman, Oak Ridge National Lab, DOE, USA
23. Hong Cai, University of Texas at San Antonio, USA
24. Liming Cai, University of Georgia, USA
25. Rui Camacho, LIACC/FEUP University of Porto, Portugal
26. Ada Chen, Southern Illinois University, USA
27. Bin Chen, Duke University, USA
28. George T. Y. Chen, Harvard University, Harvard Medical School and MGH, USA
29. Jianer Chen, Texas A&M University, USA
30. Luonan Chen, Osaka Sangyo University, Japan
31. Yuehui Chen, University of Jinan, USA.
32. Chao Cheng, University of Southern California, USA
33. Jianlin Cheng, University of Missouri, USA
34. Chung-Kuan, Cheng, University of California (UCSD), USA
35. Qiang Cheng, Wayne State University and Southern Illinois University, USA
36. Sheng Cheng, Shanghai Jiaotong University, China
37. Annie Chiang, Stanford University, USA
38. Sung-Bae Cho, Yonsei University, Korea
39. Albert C. S. Chung, Hong Kong University of Science and Technology, Hong Kong
40. Mark Clement, Brigham Young University, USA
41. Craig W. Codrington, Purdue University, USA
42. Jason Corso, University at Buffalo, USA
43. Jessica Crouch, Old Dominion University, USA

44. Kevin Daimi, University of Detroit Mercy, USA
45. Phuongan Dam, University of Georgia, USA
46. Karen Daniels, University of Massachusetts Lowell, USA
47. Matthias Dehmer, Vienna Bio Center, Austria
48. Youping Deng, University of Southern Mississippi, USA
49. Chris Ding, Lawrence Berkeley National Lab. – University of California, USA
50. Zejin Ding, Georgia State University, USA
51. Karin Dorman, Iowa State University, USA
52. Cheng Du, Chinese Academy of Science, China
53. Ye Duan, University of Missouri, USA
54. Zhong-Hui Duan, University of Akron, USA
55. Werner Dubitzky, University of Ulster, Coleraine, U.K.
56. A. Keith Dunker, Indiana University Purdue University, U.S.A.
57. Mark Ebbert, Brigham Young University, USA
58. Laura L. Elnitski, National Human Genome Research Institute, NIH, USA
59. Frank Emmert-Streib, University of Washington, USA
60. Okan K. Ersoy, Purdue University, West Lafayette, U.S.A.
61. Michael Folkert, Massachusetts Institute of Technology and Harvard University, USA
62. Jean Gao, University of Texas at Arlington, USA
63. Jane Geisler-Lee, Southern Illinois University, USA
64. Arif Ghafoor, Purdue University, West Lafayette, USA
65. Joydeep Ghosh, University of Texas at Austin, USA
66. Vicente Grau, University of Oxford, U.K.
67. Georges Grinstein, University of Massachusetts Lowell, USA
68. Jianying Gu, City University of New York, USA
69. Juntao Guo, University of Georgia, USA
70. Aili Han, Shandong University at Weihai, China
71. Cliff S. Han, Los Alamos National Laboratory, DOE. USA
72. Robert Harrison, Georgia State University, USA
73. Aristotelis Hatzioannou, the National Science Foundation of Greece, Greece
74. Haibo He, Stevens Institute of Technology, USA
75. Rattikorn Hewett, Texas Tech University, USA
76. Vasant Honavar, Iowa State University, USA
77. Pengyu Hong, Brandeis University and National Center for Behavior Genomics, USA
78. David Hsu, National University of Singapore, Singapore
79. Wen-Lian Hsu, Academia Sinica, Taiwan
80. Tony Xiaohua Hu, Drexel University, USA
81. Yuh-Jyh Hu, National Chiao Tung University, Taiwan
82. Heng Huang, University of Texas at Arlington, USA
83. Xudong Huang, Harvard University and Massachusetts Institute of Technology, USA
84. Yalou Huang, Naikai University, China
85. Eric Jakobsson, University of Illinois at Urbana-Champaign, USA
86. Jonathan Jesneck, Harvard University, Harvard Medical School and DFCI, USA
87. Tao Jiang, University of California-Riverside, USA
88. Tianzi Jiang, Chinese Academy of Science, China
89. Yuan Jiang, Nanjing University, China
90. Bo Jin, Medical University of South Carolina, USA
91. Yufang Jin, University of Texas at San Antonio, USA
92. Mehmed Kantardzic, University of Louisville, USA

93. George Karypis, University of Minnesota, USA
94. Paul Keall, Stanford University, USA
95. Arpad Kelemen, Niagara University, USA
96. Daisuke Kihara, Purdue University, West Lafayette, USA
97. Ron Kikinis, Harvard University, Harvard Medical School and BWH, USA
98. Hyunsoo Kim, Georgia Institute of Technology, USA
99. Sun Kim, Indiana University, USA
100. Meena Kishore Sakharkar, Nanyang Technological University, Singapore
101. Thomas Knudsen, University of Louisville, USA
102. Shawn Konecni, University of Massachusetts Lowell, USA
103. Stefan C. Kremer, University of Guelph, Canada
104. Rui Kuang, University of Minnesota, USA
105. Vipin Kumar, University of Minnesota, USA
106. Michael Kwok-Po Ng, Hong Kong Baptist University, Hong Kong
107. Walter Land, Jr., Binghamton University, USA
108. Chang-Shing Lee, National University of Taiwan, Taiwan
109. Doheon Lee, KIST, Korea
110. Haim Levkowitz, University of Massachusetts Lowell, USA
111. Christina M. Li, Duke University, USA
112. Guo-Zheng Li, Shanghai University, China
113. Guoliang Li, National University of Singapore, Singapore
114. Jinyan Li, Institute for Infocomm Research, Singapore
115. Lihua Li, Hangzhou Dianzi University, China
116. Yingshu Li, Georgia State University, USA
117. Hongen Liao, University of Tokyo, Japan
118. Li Liao, University of Delaware, USA
119. Lily Liang, University of the District of Columbia, USA
120. Yanchun Liang, Jilin University, China
121. Yulan Liang, University at Buffalo, USA
122. Zhi-Pei Liang, University of Illinois at Urbana-Champaign, USA
123. David Lightfoot, Southern Illinois University, USA
124. Timothy G. Lilburn, ATCC, USA
125. Guohui Lin, University of Alberta, Canada
126. Malinda Lingwall, Indiana University, Bloomington, USA
127. Qingzhong Liu, New Mexico Tech, USA
128. Tianming Liu, Harvard University, USA
129. Wei Liu, Hangzhou Dianzi University, China
130. Ying Liu, University of Texas at Dallas, USA
131. Yunlong Liu, Indiana University School of Medicine, USA
132. Xiaole Shirley Liu, Harvard University, Harvard School of Public Health, USA
133. Gary Livingston, University of Massachusetts Lowell, USA
134. Marco Loog, University of Copenhagen, Denmark
135. Shiyong Lu, Wayne State University, USA
136. Xiaochun Lu, Duke University, USA
137. Xinghua Lu, Medical University of South Carolina, USA
138. Yi Lu, University of Maryland East Shore, USA
139. Dijun Luo, University of Texas at Arlington, USA
140. Tiejian Luo, Chinese Academy of Science, China
141. Zuojie Luo, Guangxi Medical University, China

- 142. Carol Lushbough, University of South Dakota, USA
- 143. Yan Ma, Guangxi Medical University, China
- 144. Anant Madabhushi, Rutgers University, USA
- 145. Man-Wai Mak, Hong Kong Polytechnic University, Hong Kong
- 146. Ion Mandoiu, University of Connecticut, USA
- 147. Ilias Maglogiannis, University of Aegean, Greece
- 148. Kezhi Mao, Nanyang Technological University, Singapore
- 149. Marcos Martin-Fernandez, Harvard University, USA
- 150. Osamu Maruyama, Kyushu University, Japan
- 151. Kenneth Marx, University of Massachusetts Lowell, USA
- 152. Vasilis Megalooikonomou, Temple University, USA
- 153. Linda Molnar, National Cancer Institute, NIH, US Dept. of Health, USA
- 154. Jason H. Moore, Dartmouth College, Hitchcock Medical Center, USA
- 155. Shahriar Movafaghi, University of Southern New Hampshire, USA
- 156. Palaniswami Mprof, Melbourne University, Australia
- 157. Srinivas Mukkamala, New Mexico Tech, USA
- 158. Kayvan Najarian, University of North Carolina at Charlotte, USA
- 159. Giri Narasimhan, Florida International University, USA
- 160. Jennifer Neary, University of Texas at San Antonio, USA
- 161. Kathleen Neumann, Western Illinois University, USA
- 162. See-Kiong Ng, Institute for Infocomm Research, Singapore
- 163. Jun Ni, University of Iowa, USA
- 164. Juan J. Nieto, Universidad de Santiago de Compostela, Spain
- 165. Zoran Obradovic, Temple University, USA
- 166. Christopher Oldfield, Indiana University - Purdue University, USA
- 167. William Osborne, Southern Illinois University Carbondale, USA
- 168. Vasile Palade, Oxford University, U.K.
- 169. Haesun Park, Georgia Institute of Technology, USA
- 170. Minseo Park, University of Massachusetts Lowell, USA
- 171. Peter Park, Harvard University, USA
- 172. Srinivasan Parthasarathy, Ohio State University, USA
- 173. William Perrizo, North Dakota State University, USA
- 174. Vinhthuy Phan, Univerisity of Memphis, USA
- 175. Aleksandar Poleksic, University of Northern Iowa, USA
- 176. David Pollack, University of Colorado Health Science Center, USA
- 177. Jun Qin, Duke University, USA
- 178. Yifeng Qin, Guangxi Medical University, China
- 179. Sridhar Ramachandran, Indiana University Southeast, USA
- 180. Marco Ramoni, Massachusetts Institute of Technology and Harvard University, USA
- 181. Michael L. Raymer, Wright State University, USA
- 182. Arun Rawat, University of Southern Mississippi, USA
- 183. Chandan Reddy, Wayne State University, USA
- 184. J. Paul Robinson, Purdue University, USA
- 185. Usman W. Roshan, New Jersey Institute of Technology, USA
- 186. Hagit Shatkay, Queen's University, Canada
- 187. John Sharko, University of Massachusetts Lowell, USA
- 188. Bingxin Shen, Brookhaven National Lab. - Stony Brook University (SUNY), USA
- 189. Hong-Bin Shen, Harvard University, USA
- 190. Li Shen, Indiana University, USA

191. Prawal Sinha, Indian Institute of Technology Kanpur, India
192. Saurabh Sinha, University of Illinois at Urbana-Champaign, USA
193. Scott Smith, Boise State University, USA
194. Craig A. Stewart, Indiana University, Bloomington, USA
195. Yijun Sun, University of Florida, USA
196. Andrew Sung, New Mexico Tech, USA
197. Wing-Kin Sung, National University of Singapore, USA
198. Jack Szostak, Harvard University, USA
199. David Taniar, Monash University, Australia
200. Jijun Tang, University of South Carolina, USA
201. Yuchun Tang, Georgia State University, USA
202. Rahman Tashakkori, Appalachian State University, USA
203. Jeff Thai, University of Illinois, USA
204. My T. Thai, University of Florida, USA
205. Homayoun Valafar, University of South Carolina (Columbia), USA
206. Thanos Vasilakos, University of Western Macedonia, Greece
207. Vladimir N. Uversky, Indiana University School of Medicine, USA
208. Jason T.L. Wang, New Jersey Institute of Technology, USA
209. Jianzhong Wang, Northeast Normal University, China
210. Li-San Wang, University of Pennsylvania, USA
211. Liangjiang Wang, Clemson University and Greenwood Genetic Center, USA
212. Lipo Wang, Nanyang Technological University, Singapore
213. May D. Wang, Georgia Institute of Technology, USA
214. Partrick Wang, Northeastern University, USA
215. Wei Wang, University of North Carolina at Chapel Hill, USA
216. Xiangyun Wang, Amgen Inc., USA
217. Yufeng Wang, University of Texas at San Antonio, USA
218. Yuhang Wang, Southern Methodist University, USA
219. Zhengyuan Wang, Washington University, USA
220. Lonnie R. Welch, Ohio University, USA
221. Limsoon Wong, National University of Singapore, Singapore
222. Stephen Wong, Harvard University, Harvard Medical School and BWH, USA
223. Hongwei Wu, University of Georgia, USA
224. Qishi Wu, University of Memphis, USA
225. Eric P. Xing, Carnegie Mellon University, USA
226. Fei Xiong, Dartmouth College, USA
227. Dong Xu, University of Missouri-Columbia, USA
228. Jinbo Xu, Toyota Technological Institute at Chicago, USA
229. Min Xu, University of Southern California, USA
230. Ying Xu, University of Georgia and Oak Ridge National Lab., USA
231. Changhui Yan, Utah State University, USA
232. Hong Yan, City University of Hong Kong, Hong Kong
233. Xiting Yan, University of Southern California, USA
234. Jie Yang, Shanghai Jiaotong University, China
235. Laurence T. Yang, St. Francis Xavier University, Canada
236. Luge Yang, University of California at Los Angeles, USA
237. Jack Y. Yang, Harvard University, Harvard Medical School, USA
238. Mary Qu Yang, National Human Genome Research Inst., NIH; & Oak Ridge, DOE, USA
239. Mohammed Yeasin, University of Memphis, USA

- 240. Tulay Yildirim, Yildiz Technical University, Turkey
- 241. Jingkai Yu, Wayne State University, USA
- 242. Jiazheng Yuan, Southern Illinois University, USA
- 243. P. Yuan, IBM, USA
- 244. Feng Yue, University of South Carolina, USA
- 245. Mehdi Zagerham, Southern Illinois University, USA
- 246. Mohammed Zaki, Rensselaer Polytechnic Institute, USA
- 247. Leonid Zaslavsky, NCBI, NLM, National Institutes of Health, USA
- 248. Alex Zelikovsky, Georgia State University, USA
- 249. Jianyang Zeng, Duke University, USA
- 250. Aidong Zhang, University at Buffalo (SUNY), USA
- 251. Byoung-Tak Zhang, Seoul National University, South Korea
- 252. Kangyu Zhang, University of Southern California, USA
- 253. Liqiang Zhang, Indiana University South Bend, USA
- 254. Meng Zhang, Jilin University, China
- 255. Michael Q. Zhang, Cold Spring Harbor Laboratory, USA
- 256. Qing Zhang, Pharmaceutical Industry, USA
- 257. Yanqing Zhang, Georgia State University, USA
- 258. Weixiong Zhang, Washington University (St. Louis), USA
- 259. Xiang-Sun Zhang, Chinese Academy of Sciences, China
- 260. Mingyuan Zhao, University of Electronic Science and Technology of China, China
- 261. Ying Zhao, Tsinghua University, China
- 262. Jim Zheng, Medical University of South Carolina, USA
- 263. Tian Zheng, Columbia University, USA
- 264. Wei Zhong, University of South Carolina Upstate, USA
- 265. Jianping Zhou, University of Massachusetts Lowell, USA
- 266. Jie Zhou, Northern Illinois University, USA
- 267. Joe Zhou, Monash University, Australia
- 268. Joe Jizhong Zhou, Oak Ridge National Lab. DOE and University of Oklahoma, USA
- 269. Zhi-Hua Zhou, Nanjing University, China
- 270. Michelle M. Zhu, Oak Ridge National Lab. and Southern Illinois University, USA
- 271. Hanqi Zhuang, Florida Atlantic University, USA
- 272. Igor B. Zhulin, Oak Ridge National Lab, DOE and University of Tennessee, USA

## **Members of Steering and Advisory Committee**

1. Hamid R. Arabnia, University of Georgia, USA
2. Brian D. Athey, University of Michigan and NCIBI, USA.
3. Ruzena Bajcsy, University of California, Berkeley, USA.
4. Mark Borodovsky, Georgia Institute of Technology, USA
5. Nikolaos G. Bourbakis, Wright State University, USA.
6. Philip E. Bourne, University of California at San Diego, U.S.A.
7. George T. Y. Chen, Harvard University and Massachusetts General Hospital, USA.
8. Yuehui Chen, University of Jinan, China
9. Keith A. Crandall, Brigham Young University, USA.
10. Youping Deng, University of Southern Mississippi, USA.
11. Laura L. Elnitski, National Human Genome Research Institute, NIH, USA.
12. Okan K. Ersoy, Purdue University, West Lafayette, USA
13. Joydeep Ghosh, University of Texas, Austin, USA.
14. Georges Grinstein, University of Massachusetts, Lowell, USA
15. Tony Xiaohua Hu, Drexel University, USA
16. Xudong Huang, Harvard University, Harvard Medical School and BWH, USA.
17. Eric Jakobsson, University of Illinois at Urbana-Champaign, USA
18. George J. Klir, Binghamton University, USA
19. Thomas Knudsen, University of Louisville, USA
20. Ming Li, University of Waterloo, Canada
21. Zhi-Pei Liang, IEEE/EMBS Vice-President for Conferences, University of Illinois, USA
22. Jun Ni, University of Iowa, USA.
23. Andrzej Niemierko, Harvard University and Massachusetts General Hospital, USA.
24. Saranummi Niilo, Technical Research Center, Finland
25. Yi Pan, Georgia State University, USA
26. Corrado Priami, University of Trento, Italy.
27. John Quackenbush, Harvard School of Public Health and Danna-Faber Cancer Inst., USA
28. Carmelina Ruggiero, University of Genova, Italy
29. Patrick S. Wang, Northeastern University, USA
30. Limsoon Wong, National University of Singapore, Singapore.
31. Jonathan Wren, University of Oklahoma, USA.
32. Jack Y. Yang, Harvard University, Harvard Medical School, USA
33. Mary Qu Yang, National Human Genome Research Inst., NIH; & Oak Ridge, DOE, USA
34. Xin Yao, University of Birmingham, U.K.
35. Lotfi A. Zadeh, University of California, Berkeley, USA.
36. Yanqing Zhang, Georgia State University, USA.
37. Michelle M. Zhu, Oak Ridge National Lab, DOE and Southern Illinois University, USA.

Library of Congress 2007904763

ISBN 1-4244-1509-8

IEEE Catalog Number 07EX1893

IEEE Press

## TABLE OF CONTENTS

### VOLUME I

#### IEEE BIBE Plenary Keynote Lecture Notes (Part I)

<b>Intrinsically Disordered Proteins: Predictions and Applications.....</b>	<b>1</b>
A. Keith Dunker, ( <i>Indiana University School of Medicine and School of Informatics, USA</i> ); Jack Y. Yang, ( <i>Harvard Medical School, Harvard University, USA</i> ); Christopher S. Oldfield ( <i>Indiana University School of Informatics, USA</i> ); Zoran Obradovic, ( <i>Temple University, USA</i> ); Jingwei Meng, Pedro Romero ( <i>Indiana University School of Informatics, USA</i> ), Vladimir N. Uversky ( <i>Indiana University School of Medicine, USA</i> )	
<b>Stochasticity and Networks in Genomic Data.....</b>	<b>2</b>
John Quackenbush ( <i>Harvard School of Public Health, Harvard University, USA</i> )	
<b>Decoding Novel Genomes: From Microbiomes to the Eukaryota.....</b>	<b>3</b>
Mark Borodovsky ( <i>Georgia Institute of Technology, USA</i> )	
<b>Promoter Studies in the Human Genome: One Perspective on an Unfinished Story.....</b>	<b>4</b>
Mary Qu Yang ( <i>National Human Genome Research Institute, National Institutes of Health (NIH), U.S. Department of Health and Human Services and Oak Ridge, DOE, USA</i> ); Laura L. Elnitski ( <i>National Human Genome Research Institute, National Institutes of Health (NIH), U.S. Department of Health and Human Services, USA</i> )	
<b>Protein Structure Prediction and Its Understanding Based on Machine Learning Methods.....</b>	<b>7</b>
Yi Pan ( <i>Georgia State University, USA</i> )	
<b>Statistical Analysis of Nucleosome Occupancy and Histone Modification Data.....</b>	<b>8</b>
Guocheng Yuan, Jun S. Liu ( <i>Harvard University, USA</i> )	
<b>An Investigation into the Feasibility of Detecting Microscopic Disease using Machine Learning.....</b>	<b>9</b>
Mary Qu Yang ( <i>National Human Genome Research Institute, National Institutes of Health (NIH) and Oak Ridge, DOE, USA</i> ); Jack Y. Yang, ( <i>Harvard Medical School, Harvard University, USA</i> )	
<b>Nanobioinformatics: The Enabling Technology of Personalized Medicine .....</b>	<b>11</b>
Linda K. Molnar ( <i>National Cancer Institute, National Institutes of Health (NIH), U.S. Department of Health and Human Services, USA</i> )	
<b>Intrinsically Disordered Proteins in Human Diseases.....</b>	<b>12</b>
Vladimir N. Uversky ( <i>Indiana University School of Medicine, USA</i> ); Christopher S. Oldfield ( <i>Indiana University School of Informatics, USA</i> ); A. Keith Dunker ( <i>Indiana University, USA</i> )	
<b>Metallobiochemistry of Alzheimer's Disease and Its Therapeutic Agent Development.....</b>	<b>13</b>
Xudong Huang ( <i>Harvard Medical School and Brigham &amp; Women's Hospital, USA</i> )	

#### Regular Research Papers (Part I).

##### Session 1: Protein Structure and Function

<b>Identification of Conserved Domain Combinations in <i>S.cerevisiae</i> Proteins .....</b>	<b>14</b>
Suk Hoon Jung, Hee-Young Hur, Desok Kim, Dong-Soo Han ( <i>Information and Communications University, Korea</i> )	
<b>Multiclass Fuzzy Clustering Support Vector Machines for Protein Local Structure Prediction.....</b>	<b>21</b>
Wei Zhong ( <i>University of South Carolina Upstate, USA</i> ); Jieyue He ( <i>Southeast University, China</i> ); Yi Pan ( <i>Georgia State University, USA</i> )	
<b>A New Alignment-Independent Algorithm for Clustering Protein Sequences .....</b>	<b>27</b>
Abdellali Kelil, Shengrui Wang, Ryszard Brzezinski ( <i>Sherbrooke University, Canada</i> )	

<b>Metropolis-Hastings Algorithm and Continuous Regression for Finding Next-State Models of Protein Modification Using Information Scores .....</b>	<b>35</b>
David J. John, Jacquelyn S. Fetrow, James L. Norris ( <i>Wake Forest University, USA</i> )	
<b>Protein Function Prediction from Interaction Networks Using a Random Walk Ranking Algorithm .....</b>	<b>42</b>
Valerio Freschi ( <i>University of Urbino, Italy</i> )	
<b>Intrinsically Disordered Proteins: An Update .....</b>	<b>49</b>
A. Keith Dunker, ( <i>Indiana University School of Medicine and School of Informatics, USA</i> ); Jack Y. Yang, ( <i>Harvard Medical School, Harvard University, USA</i> ); Christopher S. Oldfield, ( <i>Indiana University School of Informatics, USA</i> ); Zoran Obradovic, ( <i>Temple University, USA</i> ); Jingwei Meng, Pedro Romero ( <i>Indiana University School of Informatics, USA</i> ); Vladimir N. Uversky ( <i>Indiana University School of Medicine, USA</i> );	
<b>Linking Protein Mass with Function via Organismal Massome Networks.....</b>	<b>59</b>
Gil Alterovitz ( <i>Harvard Medical School and Massachusetts Institute of Technology, USA</i> ); Eugenia Lyashenko ( <i>Columbia University, USA</i> ); Michael Xiang ( <i>Harvard Medical School, USA</i> ); Marco F. Ramoni ( <i>Harvard University and Massachusetts Institute of Technology, USA</i> )	
<b>Ensemble of Probabilistic Neural Networks for Protein Fold Recognition.....</b>	<b>66</b>
Yuehui Chen, Xueqin Zhang ( <i>University of Jinan, China</i> ); Mary Qu Yang ( <i>National Human Genome Research Institute, National Institutes of Health (NIH) and Oak Ridge, DOE, USA</i> ); Jack Y. Yang, ( <i>Harvard Medical School, Harvard University, USA</i> )	
 <b>Session 2: Microarray Data and Applications (Part A)</b>	
<b>HICCUP: Hierarchical Clustering Based Value Imputation Using Heterogeneous Gene Expression Microarray Datasets .....</b>	<b>71</b>
Qiankun Zhao ( <i>AOL Labs, China</i> ); Prasenjit Mitra, Dongwon Lee ( <i>Penn State University, USA</i> ); Jaewoo Kang ( <i>Korea University, Korea</i> )	
<b>On the Effectiveness of Constraints Sets in Clustering Genes.....</b>	<b>79</b>
Erliang Zeng, Chengyong Yang, Tao Li, Giri Narasimhan ( <i>Florida International University, USA</i> )	
<b>Integrate Qualitative Biological Knowledge to Build Gene Networks by Parallel Dynamic Bayesian Network Structure Learning.....</b>	<b>87</b>
Song Li ( <i>Iowa State University, USA</i> )	
<b>Finding Clusters of Positive and Negative Coregulated Genes in Gene Expression Data .....</b>	<b>93</b>
Kerstin Koch ( <i>Hasselt University, Belgium</i> ); Stefan Schönauer ( <i>University of Helsinki, Finland</i> ); Ivy Jansen, Jan van den Bussche, Tomasz Burzykowski ( <i>Hasselt University, Belgium</i> )	
<b>Pattern Cores and Connectedness in Cancer Gene Expression .....</b>	<b>100</b>
Noha A. Yousi, Mohamed S. Kamel ( <i>University of Waterloo, Canada</i> ); Mohamed A. Ismail ( <i>University of Alexandria, Egypt</i> )	
<b>PANP – A New Method of Gene Detection on Oligonucleotide Expression Arrays.....</b>	<b>108</b>
Peter Warren ( <i>Wyeth Research, USA</i> ); Deanne Taylor ( <i>Harvard School of Public Health, USA</i> ); Paolo G.V. Martini, Jennifer Jackson ( <i>EMD Serono, USA</i> ); Jadwiga Bienkowska ( <i>Biogen-Idec</i> )	
 <b>Session 3: Microarray Data and Applications (Part B)</b>	
<b>Cooperative Partitional-Divisive Clustering and Its Application in Gene Expression Analysis .....</b>	<b>116</b>
Rasha Kashef, Mohamed S. Kamel ( <i>University of Waterloo, Canada</i> )	
<b>On the Design of Oligos for Gene Synthesis.....</b>	<b>123</b>
Chris Thachuk ( <i>Simon Fraser University, Canada</i> ); Anne Condon ( <i>University of British Columbia, Canada</i> )	
<b>Evidence for Proximal to Distal Appendage Amputation Site Effects from Global Gene Expression Correlations Found in Newt Microarrays .....</b>	<b>131</b>
Kenneth A. Marx, John Sharko, Georges G. Grinstein ( <i>University of Massachusetts Lowell, USA</i> ); Shannon Odelberg ( <i>University of Utah School of Medicine, USA</i> ); Hans-Georg Simon ( <i>Northwestern University, USA</i> )	

<b>Denoising of Array-Based DNA Copy Number Data Using the Dual-tree Complex Wavelet Transform .....</b>	<b>137</b>
Nha Nguyen, Heng Huang, Soontorn Oraintara ( <i>University of Texas at Arlington, USA</i> ); Yuhang Wang ( <i>Southern Methodist University, USA</i> )	
<b>Quality Assessment of Affymetrix GeneChip Data Using the EM Algorithm and a Naïve Bayes Classifier .....</b>	<b>145</b>
Brian E. Howard, Imara Perera, Yang Ju Im, Heike Winter-Sederoff ( <i>North Carolina State University, USA</i> ); Beate Sick ( <i>Zurich University of Applied Science Winterthur, Switzerland</i> ); Steffen Heber ( <i>North Carolina State University, USA</i> )	
<b>Comparing Cancer and Normal Gene Regulatory Networks Based on Microarray Data and Transcription Factor Analysis.....</b>	<b>151</b>
Yu-Chun Lin, Hsiang-Yuan Yeh, Shih-Wu Cheng ( <i>National Tsing Hua University, Taiwan</i> ); Von-Wun Soo ( <i>National University of Kaohsiung, Taiwan</i> )	

## Session 4: Microarray Data and Applications (Part C)

<b>Finding Cancer-Related Gene Combinations Using a Molecular Evolutionary Algorithm .....</b>	<b>158</b>
Chan-Hoon Park, Soo-Jin Kim, Sun Kim, Dong-Yeon Cho, Byoung-Tak Zhang ( <i>Seoul National University, Korea</i> )	
<b>A Two-Stage Gene Selection Algorithm by Combining ReliefF and mRMR .....</b>	<b>164</b>
Yi Zhang ( <i>Florida International University, USA</i> ); Chris Ding ( <i>University of Texas, USA</i> ); Tao Li ( <i>Florida International University, USA</i> )	
<b>Estimating Classification Error to Identify Biomarkers in Time Series Expression Data .....</b>	<b>172</b>
John H. Phan, May D. Wang ( <i>Georgia Institute of Technology, USA</i> )	
<b>Detection and Prediction of Alternative Splicing within Acceptor/Donor Sites in Pre-mRNA of Arabidopsis Thaliana.....</b>	<b>180</b>
Minseo Park, Deane L. Falcone, Kil-Young Yun, ( <i>University of Massachusetts, USA</i> ); Kil-Young Yun ( <i>Universitiy of Maine, USA</i> ); Karen M. Daniels ( <i>University of Massachusetts, USA</i> )	
<b>An Effective Interwoven Loop Design Application for Two-Channel Microarray Experiments .....</b>	<b>187</b>
Mehdi Pirooznia, Arun Rawat ( <i>The University of Southern Mississippi, USA</i> ); Ping Gong ( <i>SpecPro Inc., USA</i> ); Jack Y. Yang ( <i>Harvard Medical School, Harvard University, USA</i> ); Edward J. Perkins ( <i>US Army Engineer Research and Development Center, USA</i> ); Mary Qu Yang ( <i>National Human Genome Research Institute, National Institutes of Health (NIH), USA</i> ); Youping Deng ( <i>The University of Southern Mississippi, USA</i> )	
<b>SNPMiner: A Domain-Specific Deep Web Mining Tool.....</b>	<b>192</b>
Fan Wang, Gagan Agrawal ( <i>Ohio State University, USA</i> ); Ruoming Jin, Helen Piontkivska ( <i>Kent State University, USA</i> )	

## Session 5: Biomedical Engineering

<b>Fast and Robust Detection of Epilepsy in Noisy EEG Signals Using Permutation Entropy.....</b>	<b>200</b>
Iman Veisi, Naser Pariz, Ali Karimpour ( <i>Ferdowsi University, Iran</i> )	
<b>A Robotic System for Real-Time Tumor Manipulation During Image Guided Breast Biopsy .....</b>	<b>204</b>
Vishnu Mallapragada, Nilanjan Sarkar, ( <i>Vanderbilt University, USA</i> ); Tarun K. Podder ( <i>Thomas Jefferson University Hospital, USA</i> )	
<b>Morphometric Analysis of Hippocampal Shape in Mild Cognitive Impairment: An Imaging Genetics Study.....</b>	<b>211</b>
Li Shen, Andrew J. Saykin ( <i>Indiana University School of Medicine, USA</i> ); Moo K. Chung ( <i>University of Wisconsin Madison, USA</i> ); Heng Huang ( <i>University of Texas at Arlington, USA</i> )	
<b>A Compartmentalized Approach to the Assembly of Physical Maps .....</b>	<b>218</b>
Serdar Bozdag, Timothy J. Close, Stefano Lonardi ( <i>University of California, Riverside, USA</i> )	
<b>Graph Theory Application in Cell Nucleus Segmentation, Tracking and Identification .....</b>	<b>226</b>
Lelin Zhang, Hongkai Xiong, Kai Zhang ( <i>Shanghai Jiao Tong University, P.R. China</i> ); Xiaobo Zhou ( <i>Harvard Medical School, USA</i> )	
<b>An End-to-End Process for Cancer Identification from Images of Lung Tissue.....</b>	<b>233</b>

Walker H. Land, Jr. (*Binghamton University, USA*); Dan McKee (*Mansfield University, USA*); Tatyana Zhukov, Dansheng Song, Wei Qian (*University of South Florida, USA*)

**Rat Mammary Fat Pad Segmentation and Growth Rate Evaluation in T1 Weighted MR Images..... 240**  
Bin Wang, Jianhua Xuan (*Virginia Tech, USA*); Matthew T. Freedman, Peter G. Shields (*Georgetown University, USA*);  
Yue Wang (*Virginia Tech, USA*)

**A Simulation-Based 3D Axon Axis Extraction in Confocal Fluorescence Microscopy Images ..... 246**  
Kai Zhang, Hongkai Xiong (*Shanghai Jiao Tong University, China*); Xiaobo Zhou (*The Methodist Hospital Research Institute and Cornell University, USA*)

**Advancements in Automated Diagnostic Mammography ..... 251**  
Walker H. Land, Jr., Alda Mizaku, Thomas Raway (*Binghamton University, USA*); John Heine, Claudia Berman, Nataliya Kovalchuk (*H. Lee Moffitt Cancer Center and Research Institute and University of South Florida, USA*)

**Calcium De-blooming in Coronary CT Images ..... 257**  
Zhuangli Liang, W. Clem Karl (*Boston University, USA*); Synho Do, Udo Hoffmann, Thomas Brady, Homer Pien (*Massachusetts General Hospital and Harvard Medical School, USA*)

## Session 6: Computational Methods in Bioinformatics

**Distance Preserving Dimension Reduction Using the QR Factorization or the Cholesky Factorization ..... 263**  
Hyunsoo Kim, Haesun Park, Hongyuan Zha (*Georgia Institute of Technology, USA*)

**Discrete Methods for Association Search and Status Prediction in Genotype Case-Control Studies..... 270**  
Dumitru Brinza (*University of California at San Diego, USA*); Alexander Zelikovsky (*Georgia State University, USA*)

**A Multi-Objective Genetic Algorithm that Employs a Hybrid Approach for Isolating Codon Usage Bias Indicative of Translational Efficiency ..... 278**  
Douglas Raiford, Dan E. Krane, Travis E. Doom, Michael L. Raymer (*Wright State University, USA*)

**Rapid Detection and Classification of Bacterial Contamination Using Grid Computing..... 286**  
Wamiq M. Ahmed, Bulent Bayraktar, Arun K. Bhunia, E. Dan Hirleman, J. Paul Robinson, Bartek Rajwa (*Purdue University, USA*)

**Sensitivity Analysis of Biomolecular Simulations Using Symbolic Models ..... 294**  
Sadaf R. Alam, Nikhil Bhatia, Jeffrey S. Vetter (*Oak Ridge National Laboratory, DOE, USA*)

## Session 7: Sequence Analysis

**Detecting and Assessing Conserved Stems for Accurate Structural Alignment of RNA Sequences..... 301**  
Xiaoyong Fang, Zhigang Luo (*National University of Defense Technology, China*); Bo Yuan, Zhenghua Wang (*Ohio State University, USA*)

**The RNA String Kernel for siRNA Efficacy Prediction ..... 307**  
Shibin Qiu (*Pathwork Diagnostics, Inc., USA*); Terran Lane (*University of New Mexico, USA*)

**A Machine Learning Approach for Prediction of Lipid-Interacting Residues in Amino Acid Sequences..... 315**  
Stephanie Jiménez Irausquin (*University of South Carolina, USA*); Liangjiang Wang (*Clemson University, USA*)

**Assessing the Performance of Macromolecular Sequence Classifiers ..... 320**  
Cornelia Caragea, Jivko Sinapov, Vasant Honavar, Drena Dobbs (*Iowa State University, USA*)

**Shortest Path Approaches for the Longest Common Subsequence of a Set of Strings ..... 327**  
Marina Barsky, Ulrike Stege, Alex Thomo, Chris Upton (*University of Victoria, Canada*)

**Constrained RNA Structural Alignment: Algorithms and Application to Motif Detection in the Untranslated Regions of *Trypanosoma Brucei* mRNAs ..... 334**  
Mugdha Khaladkar (*New Jersey Institute of Technology, USA*); Vivian Bellofatto (*University of Medicine and Dentistry of New Jersey*); Jason T.L. Wang, Vandananben Patel, Marvin K. Nakayama (*New Jersey Institute of Technology, USA*)

## Session 8: Systems Biology

<b>Optimizing Flow-based Modularization by Iterative Centroid Search in Protein Interaction Networks .....</b>	<b>342</b>
Young-Rae Cho, Woochang Hwang, Aidong Zhang ( <i>State University of New York at Buffalo, USA</i> )	
<b>Homomorphisms of Multisource Trees into Networks with Applications to Metabolic Pathways.....</b>	<b>350</b>
Qiong Cheng, Robert Harrison, Alexander Zelikovsky ( <i>Georgia State University, USA</i> )	
<b>Review of Systems Biology Simulation Tools for Translational Research .....</b>	<b>358</b>
Melissa Freedenberg, Chanchala Kaddi, Chang F. Quo, May D. Wang ( <i>Georgia Institute of Technology, USA</i> )	
<b>Combined Expression Data with Missing Values and Gene Interaction Network Analysis:</b>	
<b>A Markovian Integrated Approach.....</b>	<b>366</b>
Juliette Blanchet ( <i>INRIA Rhône-Alpes, France</i> ); Matthieu Vignes ( <i>BioSS at Scottish Crop Research Institute, Scotland</i> )	
<b>A Systematic Approach to Quantifying Evolutionary Functional Trends Across the Universal Tree of Life.....</b>	<b>374</b>
Gil Alterovitz, Taro Muso ( <i>Harvard Medical School, USA</i> ); Paresh Malalur ( <i>Massachusetts Institute of Technology, USA</i> ); Marco F. Ramoni ( <i>Harvard Medical School, USA</i> )	

## Session 9: Sequence and Structure

<b>Super Granular Shrink-SVM Feature Elimination (Super GS-SVM-FE) Model for Protein Sequence Motif Information Extraction.....</b>	<b>379</b>
Bernard Chen, Stephen Pellicer, Phang C. Tai, Robert Harrison, Yi Pan ( <i>Georgia State University, USA</i> )	
<b>False Discovery Rates in Identifying Functional DNA Motifs.....</b>	<b>387</b>
Osman Abul ( <i>TOBB University of Economics and Technology, Turkey</i> ); Geir Sandve, Finn Drablos ( <i>Norwegian University of Science and Technology, Norway</i> )	
<b>A Geometrical Model for the SNP Motif Identification Problem .....</b>	<b>395</b>
Gaofeng Huang, Peter Jeavons ( <i>Oxford University, UK</i> )	
<b>A Hierarchical Grow-and-Match Algorithm for Backbone Resonance Assignments Given 3D Structure.....</b>	<b>403</b>
Fei Xiong, Chris Bailey-Kellogg ( <i>Dartmouth College, USA</i> )	
<b>A Flexible Stem-Based Local Search Algorithm for Predicting RNA Secondary Structures Including Pseudoknots.....</b>	<b>411</b>
Xiang Chen, Si-Min He, Dong-Bo Bu, Run-Sheng Chen, Wen Gao ( <i>Chinese Academy of Sciences, China</i> )	
<b>Mining Conserved Structures of Enzymes from Functional Hierarchical Classification.....</b>	<b>418</b>
Yu-Feng Huang, Yu-Shin Lin, Tian-Wei Hsu, Chien-Kang Huang ( <i>National Taiwan University, Taiwan</i> )	
<b>Database Approaches and Data Representation in Structural Bioinformatics .....</b>	<b>425</b>
Kreshna Gopal, James C. Sacchettini, Thomas R. Ioerger ( <i>Texas A&amp;M University, USA</i> )	

## Session 10: Bioinformatics Applications

<b>Identifying Genomic Regulators of Set-Wise Co-Expression .....</b>	<b>433</b>
Jung Hoon Woo ( <i>Macrogen Inc., China</i> ); Tian Zheng ( <i>Columbia University, USA</i> ); Ju Han Kim ( <i>Seoul National University, China</i> )	
<b>Tracing Lineage in Multi-Version Scientific Databases.....</b>	<b>440</b>
Mingwu Zhang, Daisuke Kihara, Sunil Prabhakar ( <i>Purdue University, USA</i> )	
<b>Resolving Scientific Service Interoperability with Schema Mapping .....</b>	<b>448</b>
Nadia Yacoubi Ayadi, Zoe Lacroix ( <i>Arizona State University, USA</i> )	
<b>A Machine Learning Approach to Pharmacological Profiling of the Quinone Scaffold in the NCI Database:</b>	
<b>A Compound Class Enriched in Those Effective Against Melanoma and Leukemia Cell Lines .....</b>	<b>456</b>
M.L. Ujwal ( <i>Eli Lilly &amp; Co. USA</i> ); Patrick Hoffman ( <i>Inforsense, USA</i> ); Kenneth A. Marx ( <i>University of Massachusetts Lowell, USA</i> )	
<b>Biomarker Selection for Predicting Alzheimer Disease using High-Resolution MALDI-TOF Data .....</b>	<b>464</b>

Jung Hun Oh, Young Bun Kim (*The University of Texas, Arlington, USA*); Prem Gurnani, Kevin P. Rosenblatt (*University of Texas at Southwestern Medical Center, USA*); Jean Gao (*University of Texas at Arlington, USA*)

**Multivariate Analysis of Imaging Mass Spectrometry Data ..... 472**

Eric R. Muir, Ibrahima J. Ndiour, Nolwenn A. Le Goasduff, Richard A. Moffitt, Ying Liu, M.C. Sullards, A.H. Merrill,Jr., Yanfeng Chen, May D. Wang (*Georgia Institute of Technology, USA*)

**Phagocyte Transmigration Modeling Using System Dynamic Controls..... 480**

Jiaxing Xue, Jean Gao, Liping Tang (*University of Texas at Arlington, USA*)

## WORKSHOP PAPERS

### WORKSHOP on Bio-Nano-Info Integration for Personalized Medicine

**Toward a Nanobioinformatics Infrastructure for Nanotechnology-Based Prostate Cancer**

**Therapeutic Response Tracking ..... 486**

David Paik (*Stanford University, USA*)

**A Novel Approach for Signal Transduction Networks Simulation at a Mesoscopic Level..... 487**

Chenxi Shao, Hongli Deng (*University of Science and Technology of China, China*)

**Microtubule Dynamics Classification using a Statistical Model of the Movement of Outer Tips..... 495**

Christopher Alberti, Jean-Phillipe Villareal, Delano Billingsley, Koon Yin Kong (*Georgia Institute of*

*Technology, USA*); Adam I. Marcus (*Emory University, USA*); Paraskevi Giannakakou (*Cornell University, USA*);

May D. Wang (*Georgia Institute of Technology, USA*)

**Computer Aided Histopathological Classification of Cancer Subtypes ..... 503**

Sohaib Waheed, Richard A. Moffitt, Qaiser Chaudry (*Georgia Institute of Technology, USA*); Andrew N. Young (*Emory University, USA*); May D. Wang (*Georgia Institute of Technology, USA*)

**Evolving Biological Behavior in Gene-Based Cellular Simulations ..... 509**

John H. Phan, Richard A. Moffitt, Todd H. Stokes, May D. Wang (*Georgia Institute of Technology, USA*)

**Engineering Multifunctional Biologically-Amenable Nanomaterials for Interfacial Therapeutic Delivery and  
Substrate-Based Cellular Interrogation ..... 517**

Mark Chen, Brian Huang, Eric Shin, Erik Robinson, Erik Pierstorff, Houjin Huang, Dean Ho (*Northwestern University, USA*)

**Non-Monotonic Radio-Sensitivity over Tumor Volumes on Adjuvant Radio Therapy -**

***A New Insight on Cell Killing and Modeling ..... 524***

Jack Y. Yang, (*Harvard Medical School, Massachusetts General Hospital and Brigham & Women's Hospital, USA*);

Mary Qu Yang (*National Human Genome Research Institute, National Institutes of Health (NIH)*,

*and Oak Ridge, DOE, USA*); Xudong Huang (*Harvard Medical School and Brigham & Women's Hospital, USA*);

Mengxia Michelle Zhu (*Sothern Illinois University, and Oak Ridge, DOE, USA*); Andrzej Niemierko (*Harvard Medical School and Massachusetts General Hospital, USA*); Youping Deng (*University of Southern Mississippi, USA*)

### WORKSHOP on Joint Research in the Southern Illinois University, University of Illinois, and Oak Ridge National Labortaroy, U.S. Department of Energy.

**Novel Materials for the Direct Removal of Water and Ions from the Body for Patients with Dialysis Needs..... 532**

Lisa Furby, Ravinder Gupta, Ajay Mahajan, Jarlen Don, Tsuchin Chu, Bakul Dave (*Southern Illinois University at Carbondale, USA*); Brad Schwartz (*Southern Illinois University School of Medicine at Springfield, USA*)

**A Robust Method for Generating Discriminative Gene Clusters ..... 538**

Min Xu (*University of Southern California, USA*); Louxin Zhang (*National University of Singapore, Singapore*);  
Pei Li Zhou (*Monash Univeristy, Australia*)

**Biological Mechanism on a Chip: Modeling and Realization of Growth Hormone Secretion Mechanism..... 546**

John R. Shell, Yonglian Wang, Nazeih M. Botros (*Southern Illinois University at Carbondale, USA*)

<b>Retrieval Analysis of a Cementless Modular Total Hip Arthroplasty Prosthesis.....</b>	<b>553</b>
Manish Paliwal ( <i>The College of New Jersey, USA</i> ); D. Gordon Allan ( <i>Southern Illinois University School of Medicine at Springfield, USA</i> ); Peter Filip ( <i>Sothern Illinois University at Carbondale, USA</i> )	
<b>Trabecular Metal Patella Implanted into Soft-Tissue in a Post-Patelloctomized Knee: A Case Report .....</b>	<b>559</b>
D. Gordon Allan ( <i>Sothern Illinois University School of Medicine at Springfield, USA</i> ); Manish Paliwal ( <i>The College of New Jersey, USA</i> ); Peter Filip ( <i>Southern Illinois University at Carbondale, USA</i> )	
<b>A Comparison Between Traditional Shift-and-Add (SAA) and Point-By-Point Back Projection (BP) - Relevance to Morphology of Microcalcifications for Isocentric Motion in Digital Breast Tomosynthesis (DBT) .....</b>	<b>563</b>
Ying Chen ( <i>Southern Illinois University Carbondale, USA</i> ); Joseph Y. Lo, James T. Dobbins, III ( <i>Duke University and Duke University Medical Center, USA</i> )	
<b>Methodology for Evaluating DNA Pattern Searching Algorithms on Multiprocessor .....</b>	<b>570</b>
Benfano Soewito, Ning Weng ( <i>Southern Illinois University at Carbondale, USA</i> )	
<b>An Efficient Compression Method for Multiplannar Reformulated Biomedical Images .....</b>	<b>578</b>
Qiang Cheng, Mehdi Zargham ( <i>Southern Illinois University at Carbondale, USA</i> )	
<b>A Computational Approach to Understand <i>Arabidopsis thaliana</i> and Soybean Resistance to <i>Fusarium solani</i> (Fsg).....</b>	<b>585</b>
Jiazheng Yuan, Mengxia Michelle Zhu, M. Javed Iqbal ( <i>Southern Illinois University at Carbondale, USA</i> ); Jack Y. Yang ( <i>Harvard Medical School, Harvard University, USA</i> ); David A. Lightfoot ( <i>Southern Illinois University at Carbondale, USA</i> )	
<b>Biostatistical Considerations of the Use of Genomic DNA Reference in Microarrays.....</b>	<b>593</b>
Yunfeng Yang ( <i>Oak Ridge National Laboratory, USA</i> ); Mengxia Michelle Zhu ( <i>Southern Illinois University at Carbondale, USA</i> ); Liyou Wu, Jizhong Zhou ( <i>University of Oklahoma, USA</i> )	
<b>Dynamic Load Balancing for Mining of Molecular Substructures using Genetic Algorithm .....</b>	<b>601</b>
Salahuddin Masum, Mohammed Yeasin ( <i>The University of Memphis, USA</i> )	

## **WORKSHOP on Progress Toward Petascale Applications in Bioinformatics And Computational Biology**

<b>I/O Induced Scalability Limits of Bioinformatics Applications .....</b>	<b>609</b>
Robert Henschel, Matthias Mueller ( <i>Technische Universität Dresden, Germany</i> )	
<b>Large-Scale QM/MM Calculations of Electronic Excitations in Yellow Protein: Toward Petascale Level of Protein Calculations .....</b>	<b>614</b>
Marek Freindorf, Matthew D. Jones ( <i>State University at Buffalo, USA</i> ); Yihan Shao, Jing Kong ( <i>State University at Buffalo, and Q-Chem Inc. USA</i> ); Thomas R. Furlani ( <i>State University at Buffalo, USA</i> )	
<b>MotifNetwork: A Grid-Enabled Workflow for High-Throughput Domain Analysis of Biological Sequences: Implications for Annotation and Study of Phylogeny, Protein Interactions, and Intraspecies Variation.....</b>	<b>620</b>
Jeffrey Tilson ( <i>Renaissance Computing Institute, USA</i> ); Gloria Rendon ( <i>National Center for Supercomputing Applications, USA</i> ); Mao-Feng Ger, Eric Jakobsson ( <i>University of Illinois at Urbana-Champaign, USA</i> )	
<b>Toward Petascale Simulation of Cellular Microphysiology .....</b>	<b>628</b>
Scott B. Baden ( <i>University of California, San Diego, USA</i> ); Terrence J. Sejnowski, Thomas M. Bartol ( <i>The Salk Institute, USA</i> ); Joel Stiles ( <i>Pittsburgh Supercomputing Center and Carnegie Mellon University, USA</i> )	

## **VOLUME II**

### **Special Sessions Papers**

#### **Session 11: Special Session on Computational Intelligence in Medical Informatics**

Silent Killing: An Object-Oriented View of Hypertension and Kidney Failures - Part I .....	635
Jacqueline Signorini, Patrick Greussay ( <i>University Paris 8, France</i> )	
Medical Image Retrieval Based on Bidimensional Empirical Mode Decomposition .....	641
Wei Liu, Weidong Xu, Lihua Li ( <i>Hangzhou Dianzi University, China</i> )	
A Modified Fuzzy Kohonen's Competitive Learning Algorithms Incorporating Local Information for MR Image Segmentation .....	647
Jun Kong, Wenjing Lu, Jianzhong Wang, Na Che, Yinghua Lu ( <i>Northeast Normal University, China</i> )	
An Efficient Modeling and Simulation System on the Kidney Matching, Distribution and Exchange Problems .....	654
Siyuan Liu, Chao Liu ( <i>Chinese Academy of Sciences, China</i> ); Yu Liu ( <i>Shandong University, China</i> ); Chunzhe Zhao ( <i>Chinese Academy of Sciences, China</i> )	
SiteSeeker – A Motif Discovery Tool .....	662
Klaus Ecker, Lonnie Welch, Dazhang Gu ( <i>Ohio University, USA</i> )	

#### **Session 12: Special Session on Evolutionary Systems Biology**

Modeling Protein Interaction Network and Mechanisms in Exocytosis.....	665
Wen Zhou, Tian Xia, Jiansong Tong, Julie Dickerson, Bo Su, Xun Gu ( <i>Iowa State University, USA</i> )	
Toward a Realistic Model for Gene Network Evolution.....	673
Wenhai Chen ( <i>Chinese Academy of Science, China</i> ); Tian Xia ( <i>Iowa State University, USA</i> ); Tianzi Jiang ( <i>Chinese Academy of Science, China</i> ); Xun Gu ( <i>Iowa State University, USA</i> )	
Application of Wavelet Transform to the MS-Based Proteomics Data Preprocessing .....	680
Pan Du ( <i>Northwestern University, USA</i> ); Haihui Wang ( <i>Beihang University, China</i> ); Simon M. Lin, Warren A. Kibbe ( <i>Northwestern University, USA</i> )	
Pathway Complements of Four <i>Yersinia</i> .....	687
Jennifer L. Neary, Maribel Sanchez ( <i>University of Texas at San Antonio, USA</i> ); Timothy G. Lilburn ( <i>American Type Culture Collection, USA</i> ); Yufeng Wang ( <i>University of Texas at San Antonio, USA</i> )	
An Improved Ant Colony Optimization Algorithm Based on Route Optimization and Its Applications in Travelling Salesman Problem .....	693
Yi Zhang, Zhi-li Pei, Jin-hui Yang, Yan-chun Liang ( <i>Jilin University, China</i> )	

Characterization of <i>MYC</i> Expression in Gefitinib Versus Acute Myeloid Leukemia Reveals Novel Therapeutic Targets .....	699
James F. Courage, Saurabh Gupta, Yufeng Wang ( <i>University of Texas at San Antonio, USA</i> )	

Hot and Cold: Spatial Fluctuation in HIV-1 Recombination Rates.....	707
Misha L. Rajaram ( <i>Iowa State University, USA</i> ); Vladimir N. Minin ( <i>University of Washington, USA</i> ); Marc A. Suchard ( <i>University of California, Los Angeles, USA</i> ); Karin S. Dorman ( <i>Iowa State University, USA</i> )	

#### **Special Session 13: Special Session on Research in Bioinformatics, Neuroinformatics, and Systems Biology in East Asia**

Computational Genome-Wide Discovery of Aberrant Splice Variations with Exon Expression Profiles.....	715
--	-----

Ryo Yoshida (*Institute of Statistical Mathematics, Japan*); Kazuyuki Numata, Seiya Imoto, Masao Nagasaki (*University of Tokyo, Japan*); Atsushi Doi (*Gene Network International, Japan*); Kazuko Ueno, Satoru Miyano (*University of Tokyo, Japan*)

**Mining Frequent Contiguous Sequence Patterns in Biological Sequences.....** 723  
Tae Ho Kang, Jae Soo Yoo, Hak Yong Kim (*Chungbuk National University, Korea*)

**Efficient Methods for Biomedical Named Entity Recognition .....** 729  
Shing-Kit Chan, Wai Lam (*The Chinese University of Hong Kong, China*)

**Stochastic Simulation Model for Patterned Neural Multi-Electrode Arrays.....** 736  
Dong-Soo Kahng, Yoonkey Nam, Doheon Lee (*Korea Advanced Institute of Science Technology, Korea*)

**Classification of Enzyme Function from Protein Sequence based on Feature Representation .....** 741  
Bum Ju Lee, Heon Gyu Lee, Jong Yun Lee, Keun Ho Ryu (*Chungbuk National University, Korea*)

**Inferring Behavioral-Level Circuits of *Caenorhabditis elegans* from the Topology of Its Wiring Diagram .....** 748  
Yunkyu Sohn (*Korea Advanced Institute of Science Technology, Korea*); Jaeseung Jeong (*Korea Advanced Institute of Science Technology, Korea and Columbia University, USA*)

**HAMMER Algorithm: Hashing with Arithmetic Modulo-4 for Motif Extraction of Regulatory Elements.....** 753  
Huitao Sheng, Kishan Mehrotra, Chilukuri Mohan, Ramesh Raina (*Syracuse University, USA*)

**A Tale of Intronless Genes in Eukaryotic Genomes.....** 759  
Meena Sakarkar, Kishore R. Sakarkar (*Nanyang Technological University, Singapore*)

**To Fuse or Not to Fuse .....** 766  
Kishore R. Sakarkar, Meena Sakarkar (*Nanyang Technological University, Singapore*)

## Session 14: Special Session on Sequence Alignment and Phylogenetic Analysis

**Using Parsimony to Guide Maximum Likelihood Searches .....** 774  
Kenneth Sundberg, Timothy O'Connor, Hyrum Carroll, Mark Clement, Quinn Snell (*Brigham Young University, USA*)

**QOMA2: Optimizing the Alignment of Many Sequences.....** 780  
Xu Zhang, Tamer Kahveci (*University of Florida, USA*)

**A Reliable Metric for Quantifying Multiple Sequence Alignment.....** 788  
Ken D. Nguyen, Yi Pan (*Georgia State University, USA*)

**Amino Acid Substitution Matrices Based on 4-Body Delaunay Contact Profiles.....** 796  
Ahmet Sacan, I. Hakkı Toroslu (*Middle East Technical University, Turkey*)

**A Heuristic for Phylogenetic Reconstruction Using Transposition .....** 802  
Feng Yue (*University of South Carolina, USA*); Meng Zhang (*Jilin University, China*); Jijun Tang (*University of South Carolina, USA*)

**Phylogeny By Top Down Clustering Using a Given Multiple Alignment.....** 809  
Abdullah N. Arslan, Peyman Bizargity (*University of Vermont, USA*)

**Reconstructing Mutational Pathways from Serial Evolutionary Trees.....** 815  
Patricia Buendia (*University of Miami, USA*)

## Session 15: Special Session on Development of Algorithms for Solving Problems in Molecular Biology

**Towards Construction of Optimal Strip-Exchanging Moves .....** 821  
Swapnoneel Roy, Ashok KumarThakur (*IBM India Pvt. Ltd., India*)

**Rough Overlapping Biclustering of Gene Expression Data.....** 828  
Ruizhi Wang, Duoqian Miao, Gang Li, Hongyun Zhang (*Tongji University, P.R. China*)

**Using Domain-Based Structural Ensemble to Improve Structure Modeling.....** 835

Fa Zhang, Zhaoyun Ma, Zhiyong Liu (*Chinese Academy of Sciences, China*); Bo Yuan (*Shanghai Jiao Tong University, China*)

- Support Vector Regression with Feature Selection for the Multivariate Calibration of Spectrofluorimetric Determination of Aromatic Amino Acids .....** 842  
Guo-Zheng Li, Hao-Hua Meng (*Nanjing University, China*); Mary Qu Yang (*National Human Genome Research Institute, National Institutes of Health (NIH), USA*); Jack Y. Yang (*Harvard Medical School, Harvard University, USA*)

## Session 16: Special Session on High-Throughput Data Analysis for Genomics and Proteomics

- Bottom-Up Multiple Row Addition Algorithms for the Bioclustering-Problem.....** 849  
Hyung-Won Koh (*ISAS - Institute for Analytical Sciences, Germany*); Lars Hildebrand (*University of Dortmund, Germany*)
- Fast Computation of Human Genetic Linkage.....** 857  
Hongling Wang (*Columbus Children's Research Institute, USA*); Alberto Maria Segre (*University of Iowa, USA*); Yungui Huang (*Columbus Children's Research Institute, USA*); Jeffrey R. O'Connell (*University of Maryland, USA*); Veronica J. Vieland (*The Ohio State University, USA*)
- 3D Spectrum Analysis of DNA Sequence: Application to *Caenorhabditis elegans* Genome .....** 864  
Afef Elloumi Oueslati, Zied Lachiri, Noureddine Ellouze (*ENIT, Tunis*)
- MotifNetwork: Genome-Wide Domain Analysis Using Grid-Enabled Workflows .....** 872  
Jeffrey L. Tilson, Alan Blatecky (*Renaissance Computing Institute, USA*); Gloria Rendon (*National Center for Supercomputing Applications, USA*); Eric Jakobsson, Mao-Feng Ger (*University of Illinois at Urbana-Champaign, USA*)
- Profilins, Formins and Katanins as Flagellar Proteins of *Leishmania spp.*: A Genome-Based, Multi-StepBioinformatics-Driven Description .....** 880  
Elton J.R. Vasconcelos, Ana C.L. Pacheco, João J.S. Gouveia, Fabiana F. Araújo, Michely C. Diniz, Michel T. Kamimura, Marcília P. Costa, Rodrigo Maggioni, Raimundo Araujo-Filho, Raimundo B. Costa, and Diana Magalhães de Oliveira (*Universidade Estadual do Ceará - UECE, Brazil*)

## Session 17: Special Session on Bio-Medical Soft Computing

- FPGA Acceleration of Phylogeny Reconstruction for Whole Genome Data .....** 888  
Jason D. Bakos, Panormitis E. Elenis, Jijun Tang (*University of South Carolina, USA*)
- A Preliminary Study of Correlation between Depth and Path Length of GO Nodes with Gene Sequence Similarity.....** 896  
Elham Khabiri (*University of Houston-Clear Lake, USA*)
- Mathematical Models and Optimization Discussions on EA System on AIDS/HIV Spread Estimating and Countermeasures Evaluating .....** 900  
Siyuan Liu, Chao Liu, Chunzhe Zhao (*Chinese Academy of Sciences, China*); Yu Liu (*Shandong University, China*)

## Session 18: Special Session on Bio-Complexity

- Fractal Dimension of Mie Scattering Spectra for the Appraisal of Infected HeLa Cells in Cultures.....** 908  
Radu Dobrescu (*Politehnica University of Bucharest, Romania*); Loretta Ichim (*Politehnica University of Bucharest, Romania and Romanian Academy, Romania*)
- Complexity-Theoretic Modeling of Biological Cyanide Poisoning as Security Attack in Self-Organizing Networks .....** 914  
Jiejun Kong (*University of Florida, USA*); Xiaoyan Hong (*University of Alabama, USA*); Dapeng Wu (*University of Florida, USA*); Mario Gerla (*Univeristy of California, Los Angeles, USA*)
- Quantitative and Probabilistic Modeling in Pathway Logic.....** 922  
Alessandro Abate (*University of California, Berkeley, USA*); Yu Bai (*Stanford University, USA*); Nathalie Sznajder, *Ecole Normale Supérieure, France*; Carolyn Talcott, Ashish Tiwari (*SRI International, USA*)

<b>A 2D Vibration Array as an Assistive Device for Visually Impaired.....</b>	<b>930</b>
Dimitrios Dakopoulos, Sanjay K. Boddhu, Nikolaos Bourbakis ( <i>Wright State University, USA</i> )	

## Session 19: Special Session on Machine Learning Methods in Structural and Functional Genomics

<b>Characterizing and Predicting Catalytic Residues in Enzyme Active Sites Based on Local Properties: A Machine Learning Approach .....</b>	<b>938</b>
Leonardo Bobadilla, Ferno Niño, Edilberto Cepeda ( <i>National University of Colombia, Colombia</i> ); Manuel A. Patarroyo ( <i>Fundacion Instituto de Inmunologia de Colombia, Colombia</i> )	
<b>Predicting Protein-Protein Interaction Based on Fisher Scores Extracted from Domain Profiles .....</b>	<b>946</b>
Tapan Patel, Li Liao ( <i>University of Delaware, USA</i> )	
<b>A Novel Sequence-Structure Approach for Accurate Prediction of Resistance to HIV-1 Protease Inhibitors.....</b>	<b>952</b>
Majid Masso, Iosif I. Vaisman ( <i>George Mason University, USA</i> )	
<b>Prediction of Histone Modifications in DNA Sequences .....</b>	<b>959</b>
Tho Hoan Pham ( <i>Hanoi National University of Education, Vietnam</i> ); Tu Bao Ho, Dang Hung Tran, Kenji Satou ( <i>Japan Advanced Institute of Science and Technology, Japan</i> )	
<b>Feature Selection and Partial Least Squares Based Dimension Reduction for Tumor Classification.....</b>	<b>967</b>
Hua-Long Bu, Guo-Zheng Li ( <i>Shanghai University, China</i> ); Jack Y. Yang ( <i>Harvard Medical School, USA</i> ); Mary Qu Yang ( <i>National Institutes Health (NIH), USA</i> )	

## Session 20: Special Session on Pattern Recognition and Gene Discovery in Molecular Genetics

<b>aCORR: A Novel Algorithm for Clustering Gene Expression Data.....</b>	<b>974</b>
Hossam Sharara, Mohamed A. Ismail ( <i>Alexandria University, Egypt</i> )	
<b>An Intelligent System for Searching Genomic Sequences .....</b>	<b>982</b>
Vana Gummuluru ( <i>University of Florida, USA</i> ); Su-Shing Chen ( <i>University of Florida, USA; and Shaghai Institute of Biological Sciences, China</i> )	
<b>XML Encoding of Features Describing Rule-Based Modeling of Reaction Networks with Multi-Component Molecular Complexes .....</b>	<b>987</b>
Michael L. Blinov, Ion Moraru ( <i>University of Connecticut Health Center, USA</i> )	

## Session 21: Special Session on DNA Microarray Data Analysis

<b>BALBOA: Extending Bicluster Analysis to Classify ORFs using Expression Data .....</b>	<b>995</b>
Kenneth Bryan, Pádraig Cunningham ( <i>University College Dublin, Ireland</i> )	
<b>A Comprehensive Fuzzy-Based Framework for Cancer Microarray Data Gene Expression Analysis .....</b>	<b>1003</b>
Zhenyu Wang, Vasile Palade ( <i>Oxford University, UK</i> )	
<b>Ensemble of Kernel based Classifiers to Improve the Human Cancer Prediction using DNA Microarrays.....</b>	<b>1011</b>
Ángela Blanco, Manuel Martín-Merino ( <i>Universidad Pontificia de Salamanca, Spain</i> ); Javier De Las Rivas ( <i>Cancer Research Center (CIC-IBMCC, CSIC/USAL), Spain</i> )	
<b>Mining Order Preserving Patterns in Microarray Data by Finding Frequent Orders.....</b>	<b>1019</b>
Li Teng, Laiwan Chan ( <i>The Chinese University of Hong Kong</i> )	
<b>A New Smoothing Model for Analyzing Array CGH Data .....</b>	<b>1027</b>
Nha Nguyen, Heng Huang, Soontorn Oraintara, An Vo ( <i>University of Texas at Arlington, USA</i> )	
<b>Inference of Gene Regulatory Networks using Time Sliding Comparison and Transcriptional Lagging Time from Time Series Gene Expression Profiles .....</b>	<b>1035</b>
Sheehyun Kim, Dongsup Kim ( <i>Korea Advanced Institute of Science and Technology, Korea</i> )	

## Regular Research Papers (Part II)

### Session 22: Microarray Gene Expression Analysis

<b>Multivariate Feature Selection using Random Subspace Classifiers for Gene Expression Data .....</b>	<b>1041</b>
Vidya Kamath ( <i>University of South Florida and H. Lee Moffitt Cancer Center &amp; Research Institute, USA</i> ); Lawrence Hall ( <i>University of South Florida, USA</i> ); Timothy J. Yeatman, Steven A. Eschrich ( <i>H. Lee Moffitt Cancer Center &amp; Research Institute, USA</i> )	
<b>Gene Selection via Matrix Factorization .....</b>	<b>1046</b>
Fei Wang ( <i>Tsinghua University, China</i> ); Tao Li ( <i>Florida International University, USA</i> )	
<b>Mutual Information based Minimum Spanning Trees Model for Selecting Discriminative Genes .....</b>	<b>1051</b>
Fang Zhou, Jieyue He ( <i>Southeast University, China</i> ); Wei Zhong ( <i>University of South Carolina Upstate, USA</i> )	
<b>Gene-Markers Representation for Microarray Data Integration.....</b>	<b>1056</b>
Elena Baralis, Elisa Ficarra, Alessandro Fiori, Enrico Macii ( <i>Politecnico di Torino, Italy</i> )	
<b>Partial Mixture Model for Tight Clustering in Exploratory Gene Expression Analysis .....</b>	<b>1061</b>
Yinyin Yuan, Chang-Tsun Li ( <i>University of Warwick, UK</i> )	
<b>rSWTi: A Robust Stationary Wavelet Denoising Method for Array CGH Data .....</b>	<b>1066</b>
Yuhang Wang, Siling Wang ( <i>Southern Methodist University, USA</i> ); Andrew R. Zinn ( <i>UT Southwestern Medical Center at Dallas, USA</i> )	
<b>An Empirical CDF Approach to Estimate the Significance of Gene Ranking for Finding Differentially Expressed Genes .....</b>	<b>1071</b>
Jahangheer Shaik, E.O. George, Mohammed Yeasin ( <i>University of Memphis, USA</i> )	
<b>Gene Expression Profiling and Machine Learning to Understand and Predict Primary Graft Dysfunction .....</b>	<b>1076</b>
Monika Ray, Sekhar Dharmarajan ( <i>Washington University, USA</i> ); Johannes Freudenberg ( <i>Cincinnati Children's Hospital, USA</i> ); G. Alexander Patterson, Weixiong Zhang ( <i>Washington University, USA</i> )	
<b>Sensitivity and Consistency of Affymetrix GeneChip Normalization Methods .....</b>	<b>1081</b>
Jared Fox, Nik Brown ( <i>UCLA, USA</i> )	

### Session 23: Bioengineering (Part A) -- Deformable and 3D Modeling

<b>Statistical Shape Analysis of the Corpus Callosum in Subtypes of Autism.....</b>	<b>1087</b>
Qing He, Ye Duan, Judith Miles, Nicole Takahashi ( <i>University of Missouri-Columbia, USA</i> )	
<b>Simulation and Visualization of Menisci-Femur Contact using Patient-Specific Deformable Models.....</b>	<b>1092</b>
Ying Zhu ( <i>Georgia State University, USA</i> )	
<b>Improved Edge Map of Geometrical Active Contour Model based on Coupling to Anisotropic Diffusion Filtering .....</b>	<b>1097</b>
Foued Derraz, Abdelmalik Taleb-Ahmed ( <i>Valenciennes and Hainaut Cambresis University, France</i> ); Azzeddine Chikh, Fethi Bereksi-Reguig ( <i>Abou Bekr Belkaid University, Algeria</i> )	
<b>Analysis of Cardiac Wall Motion Estimation Methods.....</b>	<b>1102</b>
Soroor Behbahani, Keivan Magholi ( <i>Islamic Azad University, Iran</i> )	
<b>Partial Transmission High-Speed Continuous Tracking Multi-Leaf Collimator for 4D Adaptive Radiation Therapy .....</b>	<b>1108</b>
Tarun Podder, Ivan Buzurovic, Yida Hu, James M. Galvin, Yan Yu ( <i>Thomas Jefferson University, USA</i> )	
<b>Quantitative Analysis of Inter-Object Spatial Relationships in Biological Images .....</b>	<b>1113</b>
Wamiq Ahmed, Magdalena Jonczyk, Ali Shamsaie, Arif Ghafoor, J. Paul Robinson ( <i>Purdue University, USA</i> )	

### Session 24: Bioengineering (Part B) – Biomedical Image Processing I

<b>Histogram Transformation for Inter-Modality Image Registration.....</b>	<b>1118</b>
Joerg Meyer ( <i>University of California at Irvine, USA</i> )	
<b>The GPU on Biomedical Image Processing for Color and Phenotype Analysis.....</b>	<b>1124</b>
Antonio Ruiz, Manuel Ujaldón, José Antonio Andrades, José Becerra ( <i>University of Malaga, Spain</i> ); Kun Huang, Tony Pan, Joel Saltz ( <i>Ohio State University, USA</i> )	
<b>Tissue Identification in Ultrasound Images using Rayleigh Local Parameter Estimation .....</b>	<b>1129</b>
Santiago Aja-Fernández, Marcos Martín-Fernández, Carlos Alberola-López ( <i>Universidad de Valladolid, Spain</i> )	
<b>Exploring Cardioneural Signal from Noninvasive ECG Measurement .....</b>	<b>1134</b>
Amirali Shayan Arani, Yi Zhu, Yi-Ning Cheng ( <i>University of California San Diego, USA</i> ); Shien-Fong Lin, Peng-Sheng Chen ( <i>Indiana University, USA</i> ); Chung-Kuan Cheng ( <i>University of California San Diego, USA</i> )	
<b>MS Lesions Detection in MRI using Grouping Artificial Immune Networks.....</b>	<b>1139</b>
Akmal Younis, Ahmed T. Soliman, Mansur R. Kabuka, Nigel M. John ( <i>University of Miami, USA</i> )	
<b>Non-Negative Tensor Factorization Based on Alternating Large-Scale</b>	
<b>Non-Negativity-Constrained Least Squares.....</b>	<b>1147</b>
Hyunsoo Kim, Haesun Park ( <i>Georgia Institute of Technology, USA</i> ); Lars Eldén ( <i>Linkoping University, Sweden</i> )	

## Session 25: Bioengineering (Part C) – Biomedical Image Processing II

<b>A Computer Aided Tool for the Assessment of Human Sperm Morphology.....</b>	<b>1152</b>
Henry Carrillo Lindado, Jorge Villarreal, Miguel Sotaquirá, Álvaro Goelkel, Ricardo Gutiérrez ( <i>Universidad del Norte, Colombia</i> )	
<b>Feature Estimation for Vocal Fold Edema Detection Using Short-Term Cepstral Analysis.....</b>	<b>1158</b>
Benedito G. Aguiar Neto, Joseana M. Fechine, Silvana Cunha Costa ( <i>Federal University of Campina Grande, Brazil</i> ); Menaka Muppa ( <i>University of Washington, USA</i> )	
<b>A New Hybrid Technique for Dermatological Image Registration.....</b>	<b>1163</b>
Heng Huang ( <i>University of Texas at Arlington, USA</i> ); Paul Bergstresser ( <i>University of Texas Southwestern Medical Center, USA</i> )	
<b>PID Control based on BP Neural Network for the Regulation of Blood Glucose Level in Diabetes.....</b>	<b>1168</b>
Chengwei Li ( <i>Harbin Institute of Technology, China</i> ); Ruiqiang Hu ( <i>Yan Shan University, China</i> )	
<b>Spatiotemporal Analysis of an Agent-Based Model of a Colony of Keratinocytes:</b>	
<b>A First Approach for the Development of Validation Methods .....</b>	<b>1173</b>
Cesar Pichardo-Almarza, Rod Smallwood, S.A. Billings ( <i>University of Sheffield, UK</i> )	

## Session 26: Sequence Analysis and Sequence Alignment

<b>A Suffix Tree Construction Algorithm for DNA Sequences.....</b>	<b>1178</b>
Hongwei Huo ( <i>Xidian University, China</i> ); Vojislav Stojkovic ( <i>Morgan State University, USA</i> )	
<b>Identifying Fungal Regulatory Motif Patterns Using SCOPE, an Ensemble Learning Method Motif Finder ....</b>	<b>1183</b>
Viktor Martyanov, Larkin Elderon, Amy Gladfelter, Dhanalakshmi R. Nair, Robert H. Gross ( <i>Dartmouth College, USA</i> )	
<b>Computational Identification of Cis-regulatory Elements Associated with Pungency of Chili Peppers .....</b>	<b>1188</b>
Tieming Ji, Desh Ranjan, Jeanne Curry, Mary O'Connell ( <i>New Mexico State University, USA</i> )	
<b>Speeding Up Pairwise Sequence Alignments: A Scoring Scheme Reweighting Based Approach .....</b>	<b>1194</b>
Yong Gao, Michael Henderson ( <i>University of British Columbia Okanagan, Canada</i> )	
<b>Iterative Refinement of Repeat Sequence Specification Using Constrained Pattern Matching.....</b>	<b>1199</b>
Dan He, Abdullah N. Arslan, Yu He, Xindong Wu ( <i>The University of Vermont, USA</i> )	
<b>Differential Scoring for Systolic Sequence Alignment .....</b>	<b>1204</b>
Antonio de la Serna ( <i>IEEE</i> )	

<b>Determining Domain Similarity and Domain-Protein Similarity Using Functional Similarity Measurements of Gene Ontology Terms.....</b>	<b>1209</b>
Jennifer Leopold, Anne Maglia, Lisa Guntly ( <i>University of Missouri - Rolla, USA</i> )	
<b>Signal Representation and Processing of Nucleotide Sequences .....</b>	<b>1214</b>
Paul Cristea, Rodica Tuduce, Iulian Nastac ( <i>University "Politehnica" of Bucharest, Romania</i> ); Jan Cornelis, Rudi Deklerck ( <i>Vrije Universiteit Brussel, Belgium</i> ); Marius Andrei ( <i>Google Inc., USA</i> )	
<b>Cluster Analysis of Regulatory Sequences with a Log Likelihood Ratio Statistics-Based Similar Measure .....</b>	<b>1220</b>
Huiru Zheng, Haiying Wang ( <i>University of Ulster, UK</i> ); Jinglu Hu ( <i>Waseda UniversityJapan</i> )	

## Session 27: Bioengineering (Part D) – Biomedical Applications

<b>A PQRM-Based PACS System for Advanced Medical Services Under Grid Environment.....</b>	<b>1225</b>
Yong-Jie Ni, Chan-Hyun Youn, Byoung-Jin Kim, Young-Joo Han, Peng Liu ( <i>Information and Communications University, Korea</i> )	
<b>Middleware Based Inpatient Healthcare Information System.....</b>	<b>1230</b>
S.H. Hsieh, S.L. Hsieh, Y.C. Weng, T.H. Yang, Feipei Lai, P.H. Cheng , X.O. Ping, M.Y. Jan, J.C. Lin, C.H. Peng, K.H. Huang, L.F. Ko, C.H. Chen , K.P. Hsu ( <i>National Taiwan University, Taiwan</i> )	
<b>Generating Social Networks of Intimate Contacts for the Study of Public Health Intervention Strategies.....</b>	<b>1235</b>
Courtney R. Corley, Lindsey Brown, Armin R. Mikler ( <i>University of North Texas, USA</i> ); Diane J. Cook ( <i>Washington State University, USA</i> ); Karan Singh ( <i>University of North Texas Health Science Center, USA</i> )	
<b>YinYang Mental Squares – An Equilibrium-Based System For Bipolar Neurobiological Pattern Classification and Analysis .....</b>	<b>1240</b>
Wen-Ran Zhang, Karl Peace ( <i>Georgia Southern University, USA</i> )	
<b>Automatically Extracting Acronyms from Biomedical Text .....</b>	<b>1245</b>
Jared Fox, Nik Brown ( <i>UCLA, USA</i> )	
<b>Feature Extraction and Parameters Selection of Classification Model on Brain-Computer Interface .....</b>	<b>1249</b>
Mingyuan Zhao, Mingtian Zhou, Qingxin Zhu, Ping Yang ( <i>University of Electronic Science Technology of China</i> )	

## Session 28: Biological Networks

<b>Consistent Modeling, Integration and Simulation of Molecular Interaction Networks in Space-Time Dimension.....</b>	<b>1254</b>
Rui Chang ( <i>Technical University of Munich, Germany</i> )	
<b>Feature Reduction for Gene Regulatory Network Control .....</b>	<b>1260</b>
Mehmet Tan ( <i>University of Calgary, Canada</i> ); Faruk Polat ( <i>Middle East Technical University, Turkey</i> ); Reda Alhajj ( <i>University of Calgary, Canada</i> )	
<b>Graph Mining of Networks from Genome Biology .....</b>	<b>1265</b>
George Chin, Jr., Grant C. Nakamura, Daniel G. Chavarria, Heidi J. Sofia ( <i>Pacific Northwest National Laboratory, USA</i> )	
<b>Structural Prediction of Protein-Protein Interactions in <i>Saccharomyces Cerevisiae</i> .....</b>	<b>1270</b>
Martin S.R. Paradesi, Doina Caragea, William H. Hsu ( <i>Kansas State University, USA</i> )	
<b>Complete Backtranslation of Oligopeptides for Metabolic Pathways Exploration of Complex Environments Using Functional Microarrays .....</b>	<b>1275</b>
Mohieddine Missaoui, David R.C. Hill ( <i>Laboratoire d'informatique et de Modélisation et Optimisation des Systèmes (LIMOS), France</i> ); Cécile Militon, Pierre Peyret ( <i>Laboratoire de Biologie des Protistes (LBP), France</i> )	
<b>A Structure Learning Algorithm for Inference of Gene Networks from Microarray Gene Expression Data Using Bayesian Networks.....</b>	<b>1280</b>
Kazuyuki Numata, Seiya Imoto, Satoru Miyano ( <i>University of Tokyo, Japan</i> )	

## Session 29: Algorithms in Bioinformatics

<b>An Algorithm for Mining Fuzzy Association Rules Based on Immune Principles .....</b>	<b>1285</b>
Zhang Lei, Li Ren-hou ( <i>Xi'an Jiaotong University, China</i> ); Zhang Lei ( <i>Henan University of Science and Technology, China</i> )	
<b>Toward The Recognition Code Of Protein-DNA Recognition .....</b>	<b>1290</b>
Juan Shan, Yuxuan Wang, Changhui Yan ( <i>Utah State University, USA</i> )	
<b>Self-Optimizing Parallel Algorithms for Haplotype Reconstruction and Their Evaluation on the JPT and CHB Genotype Data.....</b>	<b>1294</b>
Dragoș Trincă, Sanguthevar Rajasekaran ( <i>University of Connecticut, USA</i> )	
<b>Reducing Folding Scenario Candidates in Pseudoknots Detection Using PLMM_DPSS Algorithm Integrated with Energy Filters .....</b>	<b>1299</b>
Xiaolu Huang, Hesham Ali ( <i>University of Nebraska at Omaha, USA</i> )	
<b>Exploring Topological Properties of NMR Graphs.....</b>	<b>1304</b>
Paulius Micikevicius ( <i>Armstrong Atlantic State University, USA</i> ); Narsingh Deo ( <i>University of Central Florida, USA</i> )	
<b>Probability Analysis on Associations of Adverse Drug Events with Drug-Drug Interactions.....</b>	<b>1308</b>
Yu-Ting Huang, Shih-Fang Lin, Chung-Cheng Chiu, Hsiang-Yuan Yeh, Von-Wun Soo ( <i>National Tsing Hua University, Taiwan</i> )	

## Session 30: Bioinformatics Tools

<b>Combining Semantics, Context, and Statistical Evidence in Genomics Literature Search .....</b>	<b>1313</b>
Jay Urbain, Nazli Goharian ( <i>Illinois Institute of Technology, USA</i> ); Ophir Frieder ( <i>Georgetown University, USA</i> )	
<b>Graph and Topological Structure Mining on Scientific Articles.....</b>	<b>1318</b>
Fan Wang ( <i>The Ohio State University, USA</i> ); Ruoming Jin ( <i>Kent State University, USA</i> ); Gagan Agrawal ( <i>Ohio State University, USA</i> ); Helen Piontkivska ( <i>Kent State University, USA</i> )	
<b>Supervised HITS Algorithm for MEDLINE Citation Ranking .....</b>	<b>1323</b>
Ying Liu, Yongjing Lin ( <i>University of Texas at Dallas, USA</i> )	
<b>Storing Efficiently Bioinformatics Workflows.....</b>	<b>1328</b>
Michel Kinsky, Zoe Lacroix ( <i>Arizona State University, USA</i> )	
<b>Intelligent Interfaces for Mining Large-Scale RNAi-HCS Image Databases.....</b>	<b>1333</b>
Chen Lin, Wayne Mak, Pengyu Hong ( <i>Brandeis University, USA</i> ); Katharine Sepp ( <i>Harvard Medical School, USA</i> ); Norbert Perrimon ( <i>Harvard Medical School and Howard Hughes Medical Institute, USA</i> )	
<b>GPX: A Tool for the Exploration and Visualization of Genome Evolution .....</b>	<b>1338</b>
Neha Nahar, Lutz Hamel ( <i>University of Rhode Island, USA</i> ); Maria S. Poptsova, J. Peter Gogarten ( <i>University of Connecticut, USA</i> )	
<b>SBLAST: Structural Basic Local Alignment Searching Tools using Geometric Hashing .....</b>	<b>1343</b>
Tom Milledge ( <i>Florida International University, USA</i> ); Gaolin Zheng ( <i>North Carolina Central University, USA</i> ); Tim Mullins ( <i>IBM Systems and Technology Group, USA</i> ); Giri Narasimhan ( <i>Florida International University, USA</i> )	

## Session 31: Protein Structural Prediction and Functional Analysis

<b>G-Protein Coupled Receptor Subfamilies Prediction Based on Nearest Neighbor Approach .....</b>	<b>1348</b>
Mudassir Fayyaz, Adnan Mujahid ( <i>Ghulam Ishaq Khan Institute of Engineering Science &amp; Technology, Pakistan</i> ); Asifullah Khan, Tae-Sun Choi ( <i>Gwangju Institute of Science Technology, Korea</i> ); Nadeem Iqbal ( <i>Korea Advanced Institute of Science and Technology KAIST, Korea</i> )	
<b>Protein Secondary Structure Prediction Using Genetic Neural Support Vector Machines .....</b>	<b>1355</b>
Anjum Reyaz-Ahmed, Yan-Qing Zhang ( <i>Georgia State University, USA</i> )	
<b>An HV-SVM Classifier to Infer TF-TF Interactions Using Protein Domains and GO Annotations .....</b>	<b>1360</b>

Xiao-Li Li (*Institute for Infocomm Research, Singapore*); Jun-Xiang Lee, Bharadwaj Veeravalli (*National University of Singapore, Singapore*); See-Kiong Ng (*Institute for Infocomm Research, Singapore*)

**Supervised Statistical and Machine Learning Approaches to Inferring Pairwise and Module-Based Protein Interaction Networks .....** **1365**  
Fiona Browne, Haiying Wang, Huiru Zheng, Francisco Azuaje (*University of Ulster at Jordanstown, UK*)

**Predicting Protein Subcellular Localizations Using Weighted Euclidian Distance.....** **1370**  
Jing Hu, Changhui Yan (*Utah State University, USA*)

**An Efficient Data Structure for Applying Multiple Seeds in Homology Search.....** **1374**  
Alireza Hadj Khodabakhshi (*Simon Fraser University, Canada*); Mehdi Mirzazadeh (*University of Waterloo, Canada*); Arvind Gupta (*Simon Fraser University, Canada*)

**A Mixture of Experts Method for Predicting Domain Boundaries in Proteins .....** **1379**  
Ian MacDonald (*College of Saint Rose, USA*); George Berg (*University at Albany, SUNY, USA*)

**SIGN: Reliable Protein Interaction Identification by Integrating the Similarity in GO and the Similarity in Protein Interaction Networks.....** **1384**  
Woochang Hwang, Taehyong Kim, Young-rae Cho, Aidong Zhang, Murali Ramanathan (*State University of New York at Buffalo, USA*)

## Session 32: Bioinformatics Applications

**Identification of Differential Flow Cytometry Expression .....** **1389**  
Elizabeth Rossin, Saumyadipta Pyne (*Broad Institute of MIT and Harvard, USA*); Florian Hahne (*German Cancer Research Center, Germany*); Philip L. De Jager (*Brigham & Women's Hospital, USA*)

**Performance Characterization of BLAST for the Grid .....** **1394**  
Enis Afgan, Purushotham Bangalore (*University of Alabama at Birmingham, USA*)

**Large-Scale Discovery of Regulatory Motifs Involved in Alternative Splicing .....** **1399**  
Sihui Zhao, Jihye Kim, Steffen Heber (*North Carolina State University, USA*)

**Semiparametric RMA Background-Correction for Oligonucleotide Arrays .....** **1404**  
Ionut Bebu, Françoise Seillier-Moiseiwitsch, Hongfang Liu (*Georgetown University, USA*)

**Alternative Splicing: Associating Frequency with Isoforms.....** **1409**  
Anuradha Roy, Jennifer L. Leopold, Anne M. Maglia (*University of Missouri - Rolla, USA*)

**A Proposed Statistical Protocol for the Analysis of Metabolic Toxicological Data  
Derived from NMR Spectroscopy .....** **1414**  
Benjamin Kelly, Paul Anderson, Nicholas Reo (*Wright State University, USA*); Nicholas DelRaso (*Wright-Patterson AFB, USA*); Travis Doom, Michael Raymer (*Wright State University, USA*)

**A Framework for Mass Spectral Quality Assessment without Prior Information.....** **1419**  
Fang-Xiang Wu, Jiarui Ding (*University of Saskatchewan, Canada*); Guy G. Poirier (*Laval University Medical Research Center (CHUL), Canada*)

**Using Long Fragments to Reconstruct RNA Backbones .....** **1424**  
Tapani Utriainen, Graham Kemp (*Chalmers University of Technology, Sweden*)

**A Modified Uniformization Method for the Chemical Master Equation .....** **1429**  
Jingwei Zhang, Layne Watson (*Virginia Polytechnic Institute and State University, USA*)

**Decision Fusion of Circulating Markers for Breast Cancer Detection in Premenopausal Women .....** **1434**  
Jonathan L. Jesneck, Sayan Mukherjee, Loren W. Nolte (*Duke University, USA*); Anna E. Lokshin (*University of Pittsburgh, USA*); Jeffrey R. Marks, Joseph Lo (*Duke University, USA*)

## IEEE BIBE Plenary Keynote Lecture Notes (Part II)

<b>Partial Least Squares Based Dimension Reduction with Gene Selection for Tumor Classification .....</b>	<b>1439</b>
Guo-Zheng Li, Xue-Qiang Zeng ( <i>Shanghai University, China</i> ); Jack Y. Yang ( <i>Harvard Medical School, Harvard University, USA</i> ); Mary Qu Yang ( <i>National Human Genome Research Institute, National Institutes of Health (NIH), USA</i> )	
<b>Functional and Molecular Imaging: Key Components of Personalized Healthcare .....</b>	<b>1445</b>
Steven E. Seltzer ( <i>Harvard Medical School and Brigham &amp; Women's Hospital, USA</i> )	
<b>Biomedical Literature Mining.....</b>	<b>1446</b>
Xiaohua Hu ( <i>Drexel University, USA</i> )	
<b>An Overview of Bidirectional Promoters .....</b>	<b>1447</b>
Mary Qu Yang ( <i>National Human Genome Research Institute, National Institutes of Health (NIH), U.S. Department of Health and Human Services and Oak Ridge, DOE, USA</i> ); Laura L. Elnitski ( <i>National Human Genome Research Institute, National Institutes of Health (NIH), U.S. Department of Health and Human Services, USA</i> )	
<b>Distributed wireless sensors on the Human Body .....</b>	<b>1448</b>
Ruzena Bajcsy ( <i>University of California at Berkeley, USA</i> )	
<b>Transforming Medicine: Genomics, Bioinformatics, and Human Health .....</b>	<b>1449</b>
Andreas D. Baxevanis ( <i>National Human Genome Research Institute, National Institutes of Health (NIH), U.S. Department of Health and Human Services, USA</i> )	
<b>Integrative Biomedical Informatics.....</b>	<b>1450</b>
Brain D. Athey ( <i>University of Michigan Medical School and National Center for Integrative Biomedical Informatics (NCIBI), USA</i> )	
<b>High-Performance Supercomputing for Drug Design .....</b>	<b>1451</b>
Jack Y. Yang ( <i>Harvard Medical School, Harvard University, USA</i> ); Mary Qu Yang ( <i>National Human Genome Research Institute, National Institutes of Health (NIH), and Oak Ridge, DOE, USA</i> ); Xudong Huang ( <i>Harvard Medical School and Brigham &amp; Women's Hospital, USA</i> ); Mengxia Michelle Zhu ( <i>Southern Illinois University, and Oak Ridge, DOE, USA</i> ), Lawrence O. Hall ( <i>University of South Florida and IEEE</i> ); Hamid R. Arabnia ( <i>University of Georgia, USA</i> )	
<b>Bio-Nano-Info Integration for Personalized Medicine .....</b>	<b>1452</b>
Eric Jakobsson ( <i>National Center for Biomimetic Nanoconductors and University of Illinois at Urbana-Champaign, USA</i> ); May D. Wang ( <i>Georgia Tech and Emory University</i> ); Linda Molnar ( <i>National Cancer Institute, National Institutes of Health, U.S. Department of Health and Human Services, USA</i> )	
<b>Tutorial Lecture Notes</b>	
<b>Using the Soybean Genome Database (SoyGD) To Display and Analyze a 1 Gbp Genome Sequence.....</b>	<b>1453</b>
David Lightfoot, Mengxia Michelle Zhu ( <i>Southern Illinois University, Carbondale, USA</i> )	
<b>Biometrics Intelligence Information Systems and Applications .....</b>	<b>1454</b>
Patrick S. P. Wang ( <i>Northeastern University, USA</i> )	
<b>Knowledge Modeling for High Content Screening of Multimedia Biological Data .....</b>	<b>1455</b>
Arif Ghafoor, J. Paul Robinson ( <i>Purdue University, West Lafayette, USA</i> )	
<b>Reconstructing signal transduction from raw genomic data .....</b>	<b>1456</b>
Igor B. Zhulin ( <i>Oak Ridge National Laboratory, DOE, USA</i> )	
<b>Studying co-regulation and inter-regulation of genes via eQTL mapping.....</b>	<b>1457</b>
Tian Zheng ( <i>Columbia University, USA</i> )	
<b>Lecture on Progress Toward Petascale Applications in Bioinformatics and Computational Biology .....</b>	<b>1458</b>
Craig A. Stewart, Malinda Lingwall ( <i>Indiana University, USA</i> ); David Bade ( <i>Georgia Institute of Technology, USA</i> )	
<b>Performance Evaluation of a Scalable Molecular Dynamics Simulation Framework on a Massively-Parallel System .....</b>	<b>1459</b>
Sadaf R. Alam, Pratul K. Agarwal and Jeffery A. Kuehn ( <i>Oak Ridge National Laboratory, DOE, USA</i> )	

<i>Author Index for Volume I and II</i> .....	<i>Follows page 634</i>
<i>Author Index for Volume I and II</i> .....	<i>Follows page 1466</i>

Library of Congress 2007904763  
 ISBN 1-4244-1509-8  
 IEEE catalog number 07EX1893  
 IEEE Press

