CALL FOR PAPERS

6TH WORKSHOP ON ALGORITHMS IN BIOINFORMATICS



WABI 2006



Zürich, Switzerland, September 11–13, 2006

WABI is sponsored by *EATCS*, the European Association for Theoretical Computer Science, and *ISCB*, the International Society for Computational Biology.

Important Dates:

 Submission
 May 15, 2006

 Notification
 June 9, 2006

 Final Version
 June 25, 2006

 Workshop
 September 11–13, 2006

Keynote Speaker:

RON SHAMIR, TEL AVIV U.

Program Committee:

Philipp Bücher, ISREC, Switzerland (co-chair) Rita Casadio, U. Bologna, Italy Nadia El-Mabrouk, U. Montréal, Canada Olivier Gascuel, LIRMM, U. Montpellier, France Raffaele Giancarlo, U. Palermo, Italy David Gilbert, U. Glasgow, Scotland Roderic Guigo, U. Pompeu Fabra, Spain Vasant Honavar, Iowa State U., USA Daniel Huson, U. of Tübingen, Germany Jens Lagergren, KTH, Sweden Randall Linder, U. Texas Austin, USA Joao Meidanis, Unicamp, Brazil Satoru Miyano, Tokyo U., Japan Bernard Moret, U. New Mexico, USA (co-chair) Gene Myers, HHMI Janelia Farm, USA Luay Nakhleh, Rice U., USA Cedric Notredame, CNRS Marseille, France Sven Rahmann, U. Bielefeld, Germany Knut Reinert, Freie U. Berlin, Germany Marie-France Sagot, U. Claude Bernard, France David Sankoff, U. of Ottawa, Canada Adam Siepel, Cornell U., USA Jijun Tang, U. South Carolina, USA Olga Troyanskaya, Princeton U., USA Alfonso Valencia, CNB-CSIC, Spain Jaak Vilo, Egeen Inc., Estonia Tandy Warnow, U. Texas Austin, USA Lusheng Wang, City U. Hong Kong, China Tiffani Williams, Texas A&M U., USA Louxin Zhang, National U. Singapore

Organizing Committee:

Michael Hoffman, ETHZ (chair) Angelika Steger, ETHZ Emo Welzl, ETHZ Peter Widmayer, ETHZ

Web site:

http://algo06.inf.ethz.ch/wabi/

Scope:

The workshop covers research in all aspects of algorithmic work in bioinformatics. The emphasis is on discrete algorithms and machine-learning methods that address important problems in molecular biology, that are founded on sound models, that are computationally efficient, and that have been implemented and tested in simulations and on real datasets. The goal is to present recent research results, including significant work-in-progress, and to identify and explore directions of future research.

Topics:

Original research papers (including significant work-in-progress) or state-of-the-art surveys are solicited in all aspects of algorithms in bioinformatics, including, but not limited to:

- Exact, approximate, and machine-learning algorithms for sequence analysis, gene and signal recognition, alignment and assembly, molecular evolution, structure determination or prediction, gene expression, gene networks, proteomics, functional genomics, and drug design.
- Methods, software, and dataset repositories for development and testing of such algorithms and their underlying models.
- High-performance computing approaches to computationally hard learning and optimization problems in bioinformatics.

Submissions:

Authors are invited to submit papers no exceeding 12 pages when formatted in the Springer LNCS style (www.springer.de/comp/lncs/authors.html). A paper should contain a succinct statement of the issues and of their motivation, a summary of the main results, and a brief explanation of their significance, all accessible to non-specialist readers. A more technical exposition should follow and an optional short appendix may contain proofs or additional data to be consulted at the discretion of the program committee.

Electronic submission is required; procedures for submission can be found on the WABI'06 web site. In case of problems, authors may, by prior arrangement, email a PDF file to the chairs, Philipp Bücher (Philipp.Bucher@isrec.unil.ch) or Bernard Moret (moret@cs.unm.edu).

Simultaneous submission to another conference with published proceedings is not permitted, but simultaneous submission to a journal is allowed, provided that the authors notify the program chairs; if published in a journal, such a contribution will be published as a short abstract in the WABI proceedings.

Proceedings:

Accepted papers will be published in the Springer series Lecture Notes in Computer Science, under the series heading of Lecture Notes in Bioinformatics and will be allotted 12 pages. Previous proceedings of WABI appeared in LNCS/LNBI volumes 2149 (Aarhus 2001), 2452 (Rome 2002), 2812 (Budapest 2003), 3240 (Bergen 2004), and 3692 (Mallorca 2005). By arrangement with the Editor-in-Chief, selected papers from WABI will be invited for an extended publication in a special issue of the IEEE/ACM Transactions on Computational Biology and Bioinformatics.