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Volume Editor

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Preface

The RECOMB conference series – with the full name of the Annual International Conference on Research in Computational Molecular Biology – was started in 1997 by Sorin Istrail, Pavel Pevzner, and Michael Waterman. The 18th RECOMB conference, RECOMB 2014, was held in Pittsburgh, PA, USA during April 2–5, 2014. It was hosted by Carnegie Mellon University and the University of Pittsburgh.

This volume contains the 35 extended abstracts selected for oral presentation at RECOMB 2014 by the Program Committee (PC) out of 154 submissions. Each submission was assigned to at least three PC members and reviewed with the help of many external reviewers. Following the initial reviews, there was an extensive discussion of the submissions among the members of the PC, leading to the final decisions.

This year RECOMB teamed up with *PLoS Computational Biology* to allow parallel submissions to the proceedings and the journal in a coordinated manner. I would like to thank Thomas Lengauer, deputy editor of *PLoS Computational Biology*, for the countless hours he spent on coordinating the review process and ensuring the success of this partnership. In total, 55 papers were submitted for consideration by RECOMB and *PLoS Computational Biology*. Eighteen of those papers passed an initial pre-screening for the journal and underwent a full review process. Five papers were subsequently accepted to both venues and appear as two-page abstracts in this volume; a few additional papers were accepted to only one of the venues. The five papers were published in full in a special RECOMB 2014 section of *PLoS Computational Biology*. All other papers that were accepted to RECOMB 2014 were invited for submission of an edited journal version to a special issue of the *Journal of Computational Biology*.

In addition to the paper presentations, RECOMB 2014 featured six invited keynote talks by leading scientists world-wide. The keynote speakers were Ian T. Baldwin (Max Planck Institute for Chemical Ecology), Atul Butte (Stanford University), James J. Collins (Harvard University), Trey Ideker (University of California at San Diego), Tom Mitchell (Carnegie Mellon University), and Sarah A. Tishkoff (University of Pennsylvania). Following the tradition started at RE-COMB 2010, RECOMB 2014 also featured a special highlights track containing computational biology papers that were published in journals during the last 18 months. There were 48 submissions, eight of which were selected for oral presentation.

The success of RECOMB depends on the effort, dedication, and devotion of many colleagues who contributed to the organization of the conference. We thank the PC members and the external reviewers for the timely review of the assigned papers despite their busy schedules; Teresa Przytycka for chairing the highlights track; Carl Kingsford for chairing the posters track; the Steering Committee and

its chair, Martin Vingron, for many excellent suggestions on the organization of the conference; the local Organizing Committee members, especially Thom Gulish and the Co-chairs Russell Schwartz and Panayiotis (Takis) Benos, for hosting the conference and providing the administrative, logistic, and financial support; and the authors of the papers, highlights, and posters and all the attendees for their enthusiastic participation in the conference. We also thank our generous sponsors, including the International Society of Computational Biology (ISCB), the US National Science Foundation, Biomed Central GigaScience, Carnegie Mellon's Lane Center for Computational biology, and University of Pittsburgh's Department of Computational and Systems Biology. Finally, I would like to thank Yehuda Afek (Tel Aviv University), Ron Shamir, and Benny Chor (Program Chairs of RECOMB 2000 and 2012, respectively) for their support and advice.

February 2014 Roded Sharan

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