PACIFIC SYMPOSIUM ON BIOCOMPUTING 2014

2014 marks the 19th Pacific Symposium on Biocomputing. The past year has been a magnificent one for computational biology. Dr. Michael Levitt won the Nobel Prize in Chemistry in October 2013 for his work modeling and simulating biological macromolecular structure. The demand for bioinformatics and computational biology skills in the marketplace has exploded, and there is a robust industry emerging providing "Big Data" services in support of genomics, pharmaceuticals and health care. The NIH has announced a \$100M program in "Big Data to Knowledge" or BD2K, to support a broad portfolio of work across all biomedical disciplines.

In addition to being published by World Scientific and indexed in PubMED, the proceedings from all previous meetings are available online at http://psb.stanford.edu/psb-online. PSB provides sessions focusing on emerging areas in biomedical computation. These sessions are typically conceived at the previous PSB meeting as people discuss the opportunities for new sessions. Once again, we have a very exciting set of areas that build on previous sessions and introduce new topics. There are revolutions occurring in cancer, drug repurposing, personalized medicine, the characterization of complex phenotypes, and our ability to extract knowledge from text. Each of these is associated with a session at this year's meeting. In addition, we see increasing interest in non-coding RNA, quantitative training in the biosciences, and applications in autism, which form the basis for our workshops. The efforts of a dedicated group of leaders has produced an outstanding set of sessions, with associated introductory tutorials. These organizers provide the scientific core of PSB and their sessions are as follows:

Cancer Panomics: Computational Methods and Infrastructure for Integrative Analysis of Cancer High-Throughput "Omics" Data

Søren Brunak, Francisco M. De La Vega, Gunnar Rätsch, Joshua M. Stuart

Computational Approaches to Drug Repurposing and Pharmacology

Zhiyong Lu, Pankaj Agarwal, Atul Butte, S. Joshua Swamidass

Detecting and Characterizing Pleiotropy: New Methods for Uncovering the Connection Between the Complexity of Genomic Architecture and Multiple Phenotypes

Sarah A. Pendergrass, Dana C. Crawford, Anna L. Tyler

Personalized Medicine: From Genotypes and Molecular Phenotypes Towards Therapy

Oliver Stegle, Jennifer Listgarten, Steven Brenner, Leopold Parts, Quaid Morris

Text and Data Mining for Biomedical Discovery

Graciela H. González, Kevin Bretonnel Cohen, Maricel G. Kann, Casey Greene, Robert Leaman, Nigam Shah, Jieping Ye

We are also pleased to present three workshops in which investigators with a common interest come together to exchange results and new ideas in a format that is more informal than the peer-reviewed sessions. For this year, the workshops and their organizers are:

Applications of Bioinformatics to Non-Coding RNAs in the Era of Next-Generation Sequencing

Chao Cheng, Jason Moore, Casey Greene

Building the Next Generation of Quantitative Biologists

Kristine A. Pattin, Anna C. Greene, James A. Foster, Jason Moore

Uncovering the Etiology of Autism Spectrum Disorders: Genomics, Bioinformatics, Environment, Data Collection and Exploration, and Future Possibilities

Sarah A. Pendergrass & Scott Selleck

We thank our keynote speakers Atul Butte (Science keynote) and Douglas Fridsma (Ethical, Legal and Social Implications keynote).

Tiffany Murray manages the peer review process and assembly of the proceedings, and also plays a key role in many other aspects of the meeting. We also thank the National Institutes of Health and the International Society for Computational Biology (ISCB) for travel grant support. We are particularly grateful to the onsite PSB staff Al Conde, Brant Hansen, Georgia Hansen, BJ McKay-Morrison, Jackson Miller, Kasey Miller, and Paul Murray for their assistance. We also acknowledge the many busy researchers who reviewed the submitted manuscripts on a very tight schedule. The partial list following this preface does not include many who wished to remain anonymous, and of course we apologize to any who may have been left out by mistake.

We look forward to a great meeting once again.

Aloha!

Pacific Symposium on Biocomputing Co-Chairs, October 14, 2013

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Thanks to the reviewers...

Finally, we wish to thank the scores of reviewers. PSB aims for every paper in this volume be reviewed by three independent referees. Since there is a large volume of submitted papers, paper reviews require a great deal of work from many people. We are grateful to all of you listed below and to anyone whose name we may have accidentally omitted or who wished to remain anonymous.

Zaky Adam Pankag Agarwal Uri David Akavia Gary Bader Joel Bader Amol Bhalla Ziv Bar-Joseph Alexis Battle Gurkan Bebek Gill Bejerano Riccardo Bellazzi Stephen Benz Andreas Beyer Sourav Bhowmick Inanc Birol Josh Bittker Sébastien Boisvert Karsten Borgwardt Andrew Brown Søren Brunak Atul Butte Colin Campbell Elias Chaibub Neto Sreenivas Chavali Keira Cheetham Bin Chen Su-Shing Chen Jie Cheng John Cleary Greg Cooper Eivind Coward Francisco De La Vega Jeroen De Ridder Emek Demir Josh Denny Jen Doherty Frank Dudbridge Sean Ekins Kyle Ellrott Barbara Engelhardt Elana Fertig Iddo Freidberg Johannes Freundenberg Brooke Fridely Nicolo Fusi

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Robert Leaman

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Christina Leslie

Havan Lee

Haiquan Li

Wenyuan Li

Hua Li

Jiao Li

Jason Li

Yen Yi Lin Christoph Lippert Chunyu Liu Haiguang Liu Yi Liu Xinghua Lou Zhiyong Lu Yves Lussier Florian Markowetz Brett McKinney Michael Menden Jill Mesirov Sara Mostafavi Sayan Mukherjee Chad Myers Dvir Netanely William Stafford Noble Chris O'Donnell Hatice Osmanbeyoglu Princy Parsana Leopold Parts Chirag Patel Bethany Percha Fernando Perez-Cruz Jason Piper Feng Oi Francis Ouetier Barbara Rakitsch Huzefa Rangwala Ben Raphael Gunnar Ratsch Ben Readhead Mireille Regnier Juri Reimand Susanna Repo Anna Ritz Luis Rocha Sushmita Roy Julio Saez-Rodriguez S. Cenk Sahinalp Matthew Scotch Jun Sese

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