PACIFIC SYMPOSIUM ON BIOCOMPUTING 2021

2021 marks the 26th Pacific Symposium on Biocomputing (PSB). Unfortunately, circumstances surrounding the Covid-19 pandemic prevent us from gathering together on the Big Island to engage in the traditional scientific sessions, invited lectures, workshops and collegiality-by-the-beach (and pool) that has become the hallmark of PSB. Fortunately, we still received very high-quality submissions and these will be presented virtually and allow continued (socially distanced) interaction and scientific exchange. We are also arranging virtual workshops and a virtual poster session. We have shortened the total length of the meeting, to help participants avoid "zoom fatigue" but many of the presentations will be recorded and available, so the reach and impact of the meeting may in some ways be extended.

Covid-19 has changed life on earth in 2020 in ways that few of us could have imagined. Many of the most basic life and work practices and assumptions have been challenged. The biomedical informatics, bioinformatics and computational biology communities have played a special role in contributing to the fight against the virus. While many of our experimental colleagues had to shut down their labs for weeks to months during the pandemic, many computational labs were able to continue their work with access to their computing facilities and a move to virtual meetings. There are many examples of early contributions to understanding the biology of the SARS-Cov-2 virus based on computational analysis of its genome, epitope targets for vaccine development, and its proteome and connections to the host. In addition, the tsunami of publications (particularly preprints) has created challenges in automatic understanding and integration (and triage) of scientific findings, in order to help scientists track the rapidly evolving landscape of our understanding of the virus. Finally, there have been opportunities in tracking population health and the delivery of clinical care—in many cases, informatics technologies have helped decision makers understand the details of the pandemic, and how best to deploy resources. At the same time, the pandemic has exposed weaknesses in our health information infrastructure-for both communicating health data, storing it, integrating it and analyzing it. For example, computational researchers who may have never sent or received a fax (!) are now trying to help figure out how to automate their analysis (many Covid-19 case reports are still faxed to local health departments) and ensure that they are soon replaced as first-line communication methods. There are also great challenges in modeling the pandemic and its impacts on health and the economy. Never before have analytic and computational capabilities in biology and medicine been so important. So, we persevere under difficult circumstances to continue the mission of PSB to bring scientists together as they approach some of the most challenging and important problems.

In addition to being published by World Scientific and indexed in PubMED, the proceedings from all PSB meetings are available online at <u>http://psb.stanford.edu/psb-online/</u>. PSB has 1227 papers listed in PubMED (as of today). These papers are routinely cited in archival journal articles and often represent important early contributions in new subfields—many times before there is an established literature in more traditional journals; for this reason, many papers have garnered hundreds of citations.

The Twitter handle PSB 2021 is @PacSymBiocomp and the hashtag this year will be #psb21.

The efforts of a dedicated group of session organizers have produced an outstanding program. The sessions of PSB 2021 and their hard-working organizers are as follows:

Advanced Methods for Big Data Analytics in Women's Health

Organizers: Graciela Gonzalez-Hernandez, Karin Verspoor, Maricel G. Kann, Su Golder, Lisa Levine, Mary Regina Boland, Natalia Villanueva-Rosales, Karen O'Connor

Achieving Trustworthy Biomedical Data

Organizers: Dennis Wall, Nicholas Tatonetti, Jan Liphardt, Bethany Percha, Serena Yeung, Peter Washington

Pattern Recognition in Biomedical Data for Discovery

Organizers: Brett Beaulieu-Jones, Christian Darabos, Dokyoon Kim, Shilpa Kobren, Anurag Verma

What about the environment? Leveraging multi-omic datasets to characterize the environment's role in human <u>health</u>

Organizers: Kristin Passero, Shefali Setia Verma, Kimberly McAllister, Arjun Manrai, Chirag Patel, Molly A. Hall

Biocomputing and AI for infectious disease modelling and therapeutics Organizers: Gil Alterovitz, Wei-Lun Alterovitz, Gail H. Cassell, Lixin Zhang, A. Keith Dunker

Computational Challenges and Artificial Intelligence in Precision Medicine

Organizers: Olga Afanasiev, Joanne Berghout, Steven Brenner, Martha L. Bulyk, Dana Crawford, Jonathan H. Chen, Roxana Daneshjou, Łukasz Kidziński

We are also pleased to present four 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:

Bioinformatics of Corals Organizers: Lenore J. Cowen, Judith Klein-Seetharaman, Hollie Putnam

Translational Bioinformatics

Organizers: Jason Moore, Ju Han Kim, Dokyoon Kim

Making Tools that People Will Use: User-Centered Design in Computational Biology Research Organizers: Mary Goldman, Nils Gehlenborg

<u>Raising the Stakeholders: Improving Patient Outcomes Through Interprofessional Collaborations in AI for</u> <u>Healthcare</u>

Organizers: Carly A. Bobak, Kristine A. Giffin, Marek Svoboda, Jason Moore, Dennis P. Wall

Establishing the Reliability of Algorithms in Biomedical Informatics

Organizers: Lara Mangravite, Sean Mooney, Iddo Freidberg, Justin Guinney

Tiffany Murray has managed the peer review process and assembly of the proceedings since 2001 and plays a key role in many aspects of the meeting. We are grateful for the support of the National Institutes of Health¹. The Research Parasite Awards benefit from support from GigaScience, Jeff Stibel, Mr. and Mrs. Stephen Canon, and Drs. Casey and Anna Greene. The Research Symbiont Awards benefit from support from the Wellcome Trust, Springer-Nature and the DragonMaster Foundation.

We are particularly grateful to the PSB staff Cynthia Paulazzo and Ryan Whaley 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 virtual meeting and very much hope to see you all again on the Big Island in 2022. Aloha!

Pacific Symposium on Biocomputing Co-Chairs, October 8, 2020

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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 to 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.

Abubakar Abid Rocky Aikens Tatsuya Akutsu Wei-Lun Alterovitz Raquel Aoki Brittany Baur Amber Beitelshees Mary Regina Boland Nicolas Buchler Sophie Burkhardt Laura Cao Nyasha Chambwe Ed Chan Tianchi Chen Xiangyin Chen Yu Wai Chen Abu Chowdhury Evan Cofer Jessica Cooke Bailey Dana Crawford Sara Cromer Gargi Datta Devendra Singh Dhami Alexander D. Diehl Jessilyn Dunn Jason Ernst Nicole Ferraro Noah R. Flynn Juan Fuxman Bass Ethan Garner Mario Giacobini Benjamin Glicksberg

Graciela Gonzalez-Hernandez Molly Hall Jie Hao Bryan He Ting Hu Tai-huang Huang Hyungsoon Im Alokkumar Jha Mingon Kang Lukasz Kidzinski Benjamin Kompa Dhireesha Kudithipudi Chirag Lakhani Avantika Lal Jerry Lee Tom Lenaerts Lisa Levine Joshua Levy Li Li Pinghua Liu David Magnus Bill Majoros Dmitry Maslov Magdalena Matusiak Jason Miller Riccardo Miotto Elias Chaibub Neto Karen O'Connor Chris Oldfield Andrew Patterson Vivek Philip Vasiliki Nataly Rahimzadeh

Maya Ramchandran David Rhoinev Pedro Romero Daniel Rotroff Max Salfinger Indra Neil Sarka Jacob Schreiber Alejandro Schuler Lulu Shang Li Shen Masaki Shimomura Kyung-Ah Sohn Yosuke Tanigawa Ling Teng Elizabeth Torrres Anna Tyler Martie VanderWalt Karin Verspoor Kailas Vodrahalli Jennifer Wagner Haohan Wang Yixin Wang Jennifer Williams Kevin Wu Jeff Xia Eva Zhang Lixin Zhang Ren Zhiyun Xinxin (Katie) Zhu Marinka Zitnik