

COMMITTEES

Steering Committee:

Dan Gusfield (UC Davis, USA)
Ion Mandoiu (UConn, USA)
Yi Pan (SIAT, China; GSU, USA)
Marie-France Sagot (INRIA, France)
Zhirong Sun (Tsinghua, China)
Ying Xu (UGA, USA)
Aidong Zhang (UVA, USA)

General Co-Chairs:

Sagi Snir (U. Haifa, Israel) Alexander Zelikovsky (GSU, USA)

Vice General Chair:

Tianwei Yu (CUHK-SZ, China)

Program Co-Chairs:

Zhipeng Cai (GSU, USA) Mukul Bansal (UConn, USA) Serghei Mangul (USC, USA)

Publicity Chairs:

Yanjie Wei (SIAT, China)

Publication Chairs:

TBA

Local Arrangement Chairs:

TBA

Workshop Chairs:

TBA

Web Chairs:

Jaden Moore (USC, USA) Filipp Rondel (GSU, USA)

Special Issues:

IEEEACM TRANSACTIONS ON COMPUTATIONAL BIOLOGY AND BIOINFORMATICS



18th International Symposium on Bioinformatics Research and Applications (ISBRA)

University of Haifa, Haifa, Israel November 14 - 17, 2022

https://mangul-lab-usc.github.io/ISBRA

SCOPE

The International Symposium on Bioinformatics Research and Applications (ISBRA) provides a forum for the exchange of ideas and results among researchers, developers, and practitioners working on all aspects of bioinformatics and computational biology and their applications. Submissions presenting original research are solicited in all areas of bioinformatics and computational biology, including the development of experimental or commercial systems. Topics of interest include but are not limited to:

Biomarker discovery High-performance bio-computing

Biomedical databases and data Metagenomics integration Molecular evolution

Biomedical text mining and Molecular modeling and simulation ontologies Next-generation sequencing data analysis Biomolecular imaging Pattern discovery and classification

Biomolecular imaging Pattern discovery and classification Comparative genomics Software tools and applications

Computational genetic epidemiology Structural biology Computational proteomics Systems biology

Data mining and visualization AI and machine learning methods in

Gene expression analysis biology and medicine

Genome analysis Big data analytics in biology and

Population genetics medicine

PAPER SUBMISSION

Please submit extended abstracts (up to 12 pages in Springer LNCS format) via the online submission system (https://easychair.org/conferences/?conf=isbra2022). Submission implies willingness of at least one author to register and present at the symposium. The authors should choose one of the following tracks:

Track 1: Submit up to 12 pages in LNCS format for (i) publication of the extended abstract in the Springer LNBI proceedings and (ii) invitation to submit a 30-50%-extended version to ISBRA'22 special issues of IEEE/ACM Transactions on Computational Biology and Bioinformatics, Journal of Computational Biology, Tsinghua Science and Technology or FGCS (Pending Approval).

Track 2: Submit up to 12 pages in LNCS format for (i) publication of an abridged two-page abstract in the Springer LNBI proceedings and (ii) invitation to submit a full-length paper to ISBRA'22 special issues of BMC Bioinformatics, BMC Genomics or Neurocomputing (Pending Approval).

IMPORTANT DATES

Submission Deadline: July 25, 2022
Notification: August 25, 2022
Final Version Due: September 7, 2022