ORGANIZATION COMMITTEES Steering Committee:

Dan Gusfield (UC Davis, USA) Ion Mandoiu (UConn, USA) Yi Pan (SIAT,China),Chair Marie-France Sagot (INRIA,France) Zhirong Sun (Tsinghua, China)

Ying Xu (UGA, USA) Aidong Zhang (UVA, USA)

Zhipeng Cai (GSU, USA)

General Co-Chairs:

Yuyu Niu (KUST, China) Jianxin Wang (CSU, China) Alexander Zelikovsky (GSU, USA)

Program Co-Chairs:

Zhipeng Cai (GSU, USA) Pavel Skums(UConn, USA) Wei Peng (KUST, China)

Publicity Chairs:

Min Li (CSU, China) Yanjie Wei (SIAT, China) Xuefeng Cui(SDU, China)

Publication Chair:

Jin Liu (CSU, China)

Web Chair:

Hongdong Li (CSU, China) Wei Dai (KUST, China)

Special Issues:



Bioinformatics

Genomics

20th International Symposium on Bioinformatics Research and Applications (ISBRA)

Kunming University of Science and Technology, Kunming, China July 19-21, 2024

https://bio.csu.edu.cn/ISBRA2024/ISBRA2024 Home.html

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

Comparative genomics Software tools and applications

Computational genetic Structural biology epidemiology Systems biology

Computational proteomics AI and machine learning methods in

Data mining and visualization Bioinformatics and Medical Information

Gene expression analysis Big Data Analytics in Biology and

Genome analysis Medicine

Population genetics

PAPER SUBMISSION

Please submit extended abstracts (up to 12 pages in Springer LNCS format) via the online submission system (https://cmt3.research.microsoft.com/ISBRA2024). 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%-extend version to ISBRA'24 special issues. The pending approval journals are IEEE/ACM Transactions on Computational Biology and Bioinformatics, Journal of Computer Science and Technology, Tsinghua Science and Technology, Chinese Journal of Electronics, Big Data Mining and Analytics or IEEE Journal of Biomedical and Health Informatics.

Track 2: Submit up to 12 pages in LNCS format for (i) presentation at the conference and (ii) invitation to submit a full-length paper to ISBRA'24 special issues. The pending approval journals are Journal of Computational Biology, Methods, Interdisciplinary Sciences-Computational Life Sciences, BMC Bioinformatics or BMC Genomics.

IMPORTANT DATES

Submission Deadline: March 25, 2024
Notification: May 1, 2024
Final Version Due: May 12, 2024
Conference Date: July 19-21, 2024