14th International Conference on

Bioinformatics and Computational Biology (BICOB 2022)

PROGRAM

March 22, 2022

Virtual Conference Online

Sponsored by



International Society for Computers and Their Applications

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14th International Conference on Bioinformatics and Computational Biology (BICOB 2022)

SPONSOR

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All times are in US Central Daylight Time (CDT: UTC-05:00)

8:55 a.m. – 9:00 a.m. WELCOME By BICOB Organizers

Oliver Eulenstein, Iowa State University, USA Hisham Al-Mubaid, University of Houston-Clear Lake, USA Tamer Aldwairi, Temple University, USA

9:00 a.m. – 10:00 a.m. KEYNOTE SPEAKER

Dr. Fahad Saeed

Knight Foundation School of Computing and Information Sciences Florida International University

Challenges in Proteomics and Neuroscience with High-Performance Data Analytics and Machine-Learning

Session Chair: Dr. Oliver Eulenstein, Iowa State University, USA

10:00 a.m. - 10:05 a.m. 5 MINUTE BREAK



SESSION TITLE: Cancer Research

Tuesday March 22, 10:05 a.m. – 11:05 a.m. Session Chair: Oliver Eulenstein (Iowa State University, USA)

- 1. Transfer Learning Pre-training Dataset Effect Analysis for Breast Cancer Imaging Chanaka Bulathsinghalage and Lu Liu (North Dakota State University, USA)
- 2. Variation of miRNA target sites in the Human Genome Brad V. Bellomo, Helen Piontkivska and Arvind K. Bansal (Kent State University, USA)
- 3. Comparative Single-cell RNA-sequencing Cluster Analysis for Traumatic Brain Injury Marker Genes Detection Audra Addison and Tayo Obafemi-Ajayi (Missouri State University, USA)

11:05 a.m. – 11:10 a.m. 5 MINUTE BREAK

SESSION 2

SESSION TITLE: Machine Learning in Bioinformatics I

Tuesday March 22, 11:10 a.m. - 12:30 p.m. Session Chair: Hisham Al-Mubaid (University of Houston-Clear Lake, USA)

- 1. Learning Gene Regulatory Networks using Graph Granger Causality Pranita Patil and Maria Vaida (Harrisburg University of Science and Technology, USA)
- 2. Guiding Protein Conformation Sampling with Conformation Space Maps Ahmed Bin Zaman, Kenneth De Jong and Amarda Shehu (George Mason University, USA)
- 3. Simple evolutionary algorithm for quantifying how medical history factors predict disease outcomes James Camp and Hisham Al-Mubaid (University of Houston - Clear Lake, Houston, USA)
- 4. Analysis of AlphaFold2 for Modeling Structures of Wildtype and Variant Protein Sequences

Anowarul Kabir, Toki Tahmid Inan and Amarda Shehu (George Mason University, USA)

10 MINUTE BREAK 12:30 p.m. – 12:40 p.m.



SESSION TITLE: Bioinformatics

Tuesday March 22, 12:40 p.m. – 2:00 p.m. Session Chair: Tamer Aldwairi (Temple University, USA)

- 1. Identification and Analysis of Alternative Splicing in Soybean Plants Xiangjia Min, Theoni Kasamias, Mykaela Wagner, Atinuke Ogunbayi and Feng Yu (Youngstown State University, USA)
- 2. Signed Rearrangement Distances Considering Repeated Genes and Intergenic Regions Gabriel Sigueira, Alexsandro Oliveira Alexandrino, and Zanoni Dias (University of Campinas, Brazil)
- 3. A Novel Approach for Mapping Ambiguous Sequences of Transcriptomes Tamer Aldawiri (Temple University), Bindu Nanduri (Mississippi State University, USA), Mahalingam Ramkumar (Mississippi State University, USA) and Andy D. Perkins (Mississippi State University, USA)
- 4. Using Topological Data Analysis and RRT to Investigate Protein Conformational Spaces Ramin Dehghanpoor, Fatemeh Afrasiabi, and Nurit Haspel (University of Massachusetts, USA)

10 MINUTE BREAK 2:00 p.m. – 2:10 p.m.

SESSION 4

SESSION TITLE: Machine Learning in Bioinformatics II

Tuesday March 22, 2:10 p.m. – 3:50 p.m. Session Chair: Oliver Eulenstein (Iowa State University, USA)

1. SARS-CoV-2 variants classification and characterization

Sofia Borgato (Politecnico di Torino, Italy), Marco Bottino (Politecnico di Torino, Italy), Marta Lovino (Universita` degli Studi di Modena e Reggio Emilia, Italy), and Elisa Ficarra (Universita` degli Studi di Modena e Reggio Emilia, Italy)

2. Interpretable Image Classification Model Using Formal Concept Analysis Based Classifier

Minal Khatri (University of Nebraska-Lincoln, USA), Adam Voshall (University of Nebraska-Lincoln, USA), Surinder K. Batra (University of Nebraska Medical Center, USA), Sukhwinder Kaur (University of Nebraska Medical Center, USA), and Jitender S. Deogun (University of Nebraska-Lincoln, USA)

- 3. EMG-Based Feature Extraction and Classification for Prosthetic Hand Control Reza Bagherian Azhiri, Mohammad Esmaeili, and Mehrdad Nourani (University of Texas at Dallas, USA)
- 4. A Deep Dimensionality Reduction Method based on Variational Autoencoder for Antibody Complementarity Determining Region Sequences Analysis Saeed Khalilian, Mohammad N. Isfahani, Zahra Moti, Arian Baloochestani, Alireza Chavosh, and Zahra Hemmatian
- 5. Machine Learning Techniques in Structure-Property Optimization of Polymeric Scaffolds for Tissue Engineering Zigeng Wang, Xia Xiao, Syam P. Nukavarapu, Sangamesh G. Kumbar and Sanguthevar Rajasekaran (University of Connecticut, USA)

3:50 p.m. – 4:00 p.m. Best Paper Award Announcement

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