The 6th International Workshop on Computational Network Biology: Modeling, Analysis, and Control (CNB-MAC 2019)

September 7, 2019 | Niagara Falls, NY

CNB-MAC 2019 Workshop Chairs:

Byung-Jun Yoon (Texas A&M) Xiaoning Qian (Texas A&M)
Tamer Kahveci (University of Florida) Ranadip Pal (Texas Tech University)

Workshop Program:

09:00-09:15 Opening Remarks – CNB-MAC 2019 Workshop Chairs

09:15-10:30 Session 1 – Session chair: Dr. Young (Yang) Cao

- Cell Cycle Control and Environmental Response by Second Messenger Networks in Caulobacter Crescentus – by Chunrui Xu, Bronson Weston and Yang Cao
- Analysis of two mechanisms of telomere maintenance based on the theory of G-Networks and Stochastic Automata Networks – by <u>Kyung Hyun Lee</u> and Marek Kimmel
- Network-based Method for Loci with Statistically Frequent Inter-Chromosomal Interactions at Single-cell Resolution – by Chanaka Bulathsinghalage and <u>Lu Liu</u>

10:30-10:45 Coffee Break

10:45-12:00 Session 2 – Session chair: Dr. Dougu Nam

- Leveraging TCGA gene expression data to build predictive models for cancer drug response
 by Evan Clayton, Toyya Pujol, John McDonald and Peng Qiu
- Robust edge-based biomarker discovery improves prediction of breast cancer metastasis by Nahim Adnan, Chengwei Lei and Jianhua Ruan
- A Meta-Analysis of Genomic and Transcriptomic Variations in Lung Adenocarcinoma by Talip Zengin and <u>Tuğba Önal-Süzek</u>

12:00-13:30 Lunch Break

13:30-14:45 Session 3 – Session chair: Dr. Peng Qiu

- (Highlight Presentation) Biclustering analysis of transcriptome big data identifies condition-specific microRNA targets by Sora Yoon, Hai Nguyen and <u>Dougu Nam</u>
- Bayesian Gamma-Negative Binomial Modeling of Single-Cell RNA Sequencing Data by
 Siamak Zamani Dadaneh, Paul de Figueiredo, Sing-Hoi Sze, Mingyuan Zhou and Xiaoning
 Qian
- ComHapDet: A Spatial Community Detection Algorithm for Haplotype Assembly by <u>Abishek Sankararaman</u>, Haris Vikalo and François Baccelli

14:45-15:00 Coffee Break

15:00-16:20 Session 4 – *Session chair: Dr. Anna Ritz*

- DeeplyEssential: A Deep Neural Network for Predicting Essential Genes in Microbes by Md Abid Hasan and Stefano Lonardi
- Grow Lineages along Minimum Spanning Tree for B Cell Receptor Sequencing Data by Xingyu Yang, Christopher Tipton, Matthew Woodruff, Enlu Zhou, Frances Eun-Hyung Lee, Ignacio Sanz and Peng Qiu
- (Lightning Talk) A Bayesian Rank-Based Pathway Classifier by <u>Emre Arslan</u> and Ulisses M. Braga-Neto.
- (Lightning Talk) A Meta-analysis of Gene Expression Data in Neurodegenerative Diseases by Abdulahad Bayraktar, Tugba Önal-Süzek, <u>Barış Ethem Süzek</u> and Ömür Baysal
- (Lightning Talk) Systematic Normalization with Multiple Housekeeping Genes for the Discovery of Genetic Dependencies in Cancer by Oliver Bonham-Carter and Yee Mon Thu

16:20-16:30 Student Travel Awards & Closing Remarks – CNB-MAC 2019 Workshop Chairs

16:30-17:15 Poster Session – Session chair: Dr. Byung-Jun Yoon

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