

## ***7<sup>th</sup> Workshop on Computational Advances for Next Generation Sequencing (CANGS)***

in conjunction with

*7<sup>th</sup> IEEE International Conference on Computational Advances in Bio and Medical Sciences (ICCABS)*  
October 19-21, 2017 in Orlando, FL, USA

Workshop Co-Chairs:

Ion Mandoiu (University of Connecticut, [ion@engr.uconn.edu](mailto:ion@engr.uconn.edu))

Pavel Skums (Georgia State University, [pskums@gsu.edu](mailto:pskums@gsu.edu))

Alex Zelikovsky (Georgia State University, [alexz@cs.gsu.edu](mailto:alexz@cs.gsu.edu))

Massively parallel DNA and RNA sequencing have become widely available, reducing the cost by several orders of magnitude and placing the capacity to generate gigabases to terabases of sequence data into the hands of individual investigators. These next-generation technologies have the potential to dramatically accelerate biological and biomedical research by enabling the comprehensive analysis of genomes and transcriptomes to become inexpensive, routine and widespread. The exploding volume of data has spurred the development of novel algorithmic approaches for primary analyses of sequence data in such areas as error correction, de novo genome assembly, novel transcript discovery, virus quasispecies assembly, etc. This workshop will bring together specialists to discuss the various mathematical and computational challenges presented by next-generation sequencing technologies.

Workshop topics of interest include but are not limited to:

- Cloud-based NGS analysis
- Long single molecule read analysis
- Cancer sequencing
- 3D genome architecture
- NGS error correction
- Read mapping
- Variant detection and genotyping
- Characterization of structural variants
- De novo genome and transcriptome assembly
- Haplotype assembly
- Genome-assisted transcriptome reconstruction
- Transcriptome quantification
- Small RNA analysis
- Scaffolding and genome finishing
- Epigenomics
- Metagenomics and metatranscriptomics
- Single cell sequencing
- Population genomics

This meeting is by invitation only. If you would like to inquire about the possibility of being invited, please contact the workshop chairs as soon as possible, but no later than September 15, 2017. One page abstracts of invited talks will be included in the ICCABS proceedings published in the IEEE Xplore Digital Library. As in previous years, following the workshop invited speakers will be invited to submit full length articles to special issues of *BMC Bioinformatics* and *BMC Genomics*.

Workshop Registration Fees:

- IEEE Members: \$400
- IEEE Non-members: \$500