



Session : **Bioinformatics**
Time: 15:10 – 16:30

Chair:	<i>Ugur Sezerman , Sabanci University</i>
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A COMPARATIVE STUDY OF CROSS-CORRELATION METHODS FOR ALIGNMENT OF DNA SEQUENCES CONTAINING REPETITIVE PATTERNS ([Abstract](#))

Arndzej Brodzik (MITRE Corporation, United States)



HANDLING UPDATES OF A BIOLOGICAL SEQUENCE BASED ON HIDDEN MARKOV MODELS ([Abstract](#))

Changjin Hong (University of Minnesota, United States)
Ahmed H. Tewfik (University of Minnesota, United States)
David H. C. Du (University of Minnesota, United States)

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ROBUST BICLUSTERING ALGORITHM (ROBA) FOR DNA MICROARRAY DATA ANALYSIS ([Abstract](#))

Alain Beaudelaire (University of Minnesota, United States)
Tchagang
Ahmed H. Tewfik (University of Minnesota, United States)



GRIDDING THE SPOT CENTERS OF MICROARRAY IMAGES ([Abstract](#))

Jinn Ho (Academia Sinica, Taiwan)
Wen-Liang Hwang (Academia Sinica, Taiwan)
Henry Horn-Shing Lu (National Chiao Tung University, Taiwan)
D. T. Lee (Academia Sinica, Taiwan)

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