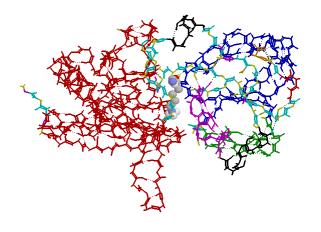
Workshop on:

Computational and Theoretical Biology



Michigan State University

Saturday, April 24, 1999

8:30 a.m. – 3:10 p.m. Room 224 Physics & Astronomy Building

Workshop on Computational and Theoretical Biology April 24, 1999 — Michigan State University

Saturday, April 24, 1999

8:30 – 9:00 Introductions by George Leroi, P.K. Wong and Tony Wojcik

Chair: Bill Punch

Session A in room 224 Physics and Astronomy Bldg.

9:00 – 9:10 <i>A1</i>	Robert J. Tempelman Generalized Linear Mixed Models for Genetic Evaluation of Livestock
9:10 – 9:20 <i>A2</i>	Chichia Chiu Numerical Methods for Pattern Formation Problems in Biology
9:20 – 9:30 <i>A3</i>	Sydney D'Silva How Honey Bees Make Decisions
9:30 – 9:40 <i>A4</i>	Frank B. Dazzo CMEIAS: A Tool for Computational Microbial Ecology
9:40 – 9:50 <i>A5</i>	Stuart H. Gage Computational Ecology and Visualization Technologies
9:50 – 10:00 <i>A6</i>	George Garrity Markup of Microbiological Data for Accelerated Publication in Print and Electronic Form
10:00 – 10:10 <i>A7</i>	Tom Getty Models of Signaling, Search, Discrimination and Selection
10:10 – 10:20 <i>A8</i>	Richard E. Lenski Genomic Complexity in Micro Organisms and Digital Organisms
10:20 - 10:50	Coffee Break

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Chair: Mike Thorpe

Session B in room 209 Physics and Astronomy Bldg.

9:00 – 9:10 <i>B1</i>	Simon J.L. Billinge Local Structure of Folded Proteins
9:10 – 9:20 <i>B2</i>	James R. Cole The Ribosomal Database Project: Providing an Evolutionary Framework
9:20 – 9:30 <i>B3</i>	Tien Yien Li Solving Polynomial Systems
9:30 – 9:40 <i>B4</i>	Shelagh S. Ferguson-Miller How Proteins Get Together and Electrons Get Transferred: Mutational, Spectroscopic, Kinetic and Computational Analysis of Cytochrome c Docking with Cytochrome c Oxidase
9:40 – 9:50 <i>B5</i>	Jay I. Goodman Altered DNA Methylation: An Epigenetic Mechanism Involved in Carcinogenesis
9:50 – 10:00 <i>B6</i>	Erik Goodman Evolutionary Algorithms for Biological Science
10:00 – 10:10 <i>B7</i>	Katharine Hunt Optical Trapping of Biomolecules
10:10 – 10:20 <i>B8</i>	Michael Kron Structure Based Design of Aminoacyl-tRNA Synthetase Inhibitors as Anti-Parasitic Drugs
10:20 - 10:50	Coffee Break

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Chair: P.K. Wong

Session C in room 224 Physics and Astronomy Bldg.

10:50 – 11:00 CI	Wolfgang Bauer Cancer Detection via Determination of Fractal Cell Dimension
11:00 – 11:10 C2	Andre Benard <i>Image-Based Analysis of Heat Transfer in Biological Systems</i>
11:10 – 11:20 C3	Raoul LePage Outline of Research Statistics Component
11:20 – 11:30 C4	Jack Deller BioSignal Processing Activities in MSU's Department of Electrical and Computer Engineering
11:30 – 11:40 C5	Nicolae Duta Learning Biological Shape Models
11:40 – 11:50 C6	Michael J. Harrison The Role of Thermally Excited Eardrum Pressure: Fluctuations in Establishing Primate Auditory Thresholds
11:50 – 12:00 <i>C7</i>	Robert Hubbard Biomechanical Models for Seating Design
12:00 – 12:10 <i>C8</i>	Fathi Salam Bio-Engineering: An Integrated Systems Approach
12:10 – 1:10	Lunch

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Chair: Tony Wocjik

Session D in room 209 Physics and Astronomy Bldg.

10:50 – 11:00 <i>D1</i>	Leslie Kuhn How Proteins Fold, Flex and Bind Other Molecules
11:00 – 11:10 D2	S.D. Mahanti Simulation on Coarse-Grained Models of Amphiphiles
11:10 – 11:20 D3	Sakti Pramanik Computational Challenges for Discovering Homologies between Genome Sequences
11:20 – 11:30 D4	William Punch Self-Assembly, Folding and Unfolding of Polymers
11:30 – 11:40 D5	M.F. Thorpe Protein Flexibility
11:40 – 11:50 <i>D6</i>	Eric Torng Incremental Update of Phylogenetic Trees Using Hierarchical Modeling
11:50 – 12:00 D7	Joseph White Bioinformatics for the Seed EST Functional Genomics Project
12:00 – 12:10 <i>D8</i>	Jack Preiss Previous Funding Support for Computational Biology from the REF Center of Protein Structure, Function and Design
12:10 – 1:10	Lunch

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Chair: Leslie Kuhn

Session E in room 224 Physics and Astronomy Bldg.

1:10 - 1:20	James E. Trosko
E1	Gene Regulation in Pluripotent Human Stem Cells
1:20 - 1:30	C.Y. Wang
E2	Simulation of Biological Growth
1:30 – 1:40	Tim Zacharewski
<i>E3</i>	Toxicogenomics
1:40 – 1:50	Milan Miklavcic
E4	Stability for Discrete Velocity Models of the Extended Boltzmann Equation
1:50 - 2:00	Mark Dykman
E5	Selective Control of Diffusion of Biological Systems
2:00 - 2:10	William M. Hartmann
E6	Perceptually Relevant Models of Neural Excitation in the Auditory System

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Chair: Sakti Pramanik

Session F in room 209 Physics and Astronomy Bldg.

1:10 – 1:20	Jianguo Liu
F1	Systems Modeling Laboratory: An Integrated Approach to Landscape and Biodiversity Study
1:20 – 1:30	David W. Hyndman
F2	Integrating Social Drivers and Environmental Impacts Using a
	Geographic Information System: The Land Transformation Modeling Project
1:30 - 1:40	Peter M. Saama
F3	Mixed Model Inference in the Analysis of cDNA Array Data
1:40 - 1:50	Tim Lilburn
F4	The Calculation of Large Phylogenetic Trees
1:50 - 2:00	Kim Scribner
F5	Novel Applications of Molecular Genetic Markers and Population Genetic Theory in Population Ecology and Resource Management
2:00 - 2:10	Mark Worden
F6	Role of Microbial Chemotaxis in Bioremediation of Microbial Ecology

2:10 – 2:40 Group discussions

2:40 – 3:10 Final wrap-up, list of recommendations, further activities etc.