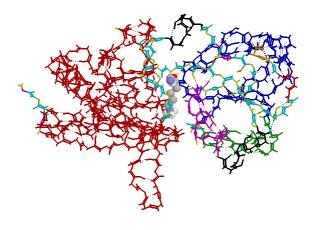
# Workshop on:

# Computational and Theoretical Biology



**Michigan State University** 

Saturday, April 24, 1999

8:30 a.m. -3:10 p.m.
Room 110 Radiology Building
(Near Clinical Center on the MSU Campus)

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

# Saturday, April 24, 1999

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

Chair: Bill Punch

#### Session A in room 110 Radiology Building

<b>9:00 – 9:10</b> <i>A1</i>	<b>Robert J. Tempelman</b> Generalized Linear Mixed Models for Genetic Evaluation of Livestock		
<b>9:10 – 9:20</b> <i>A2</i>	Chichia Chiu Numerical Methods for Pattern Formation Problems in Biology		
<b>9:20 – 9:30</b> <i>A3</i>	Sydney D'Silva How Honey Bees Make Decisions		
<b>9:30 – 9:40</b> <i>A4</i>	Frank B. Dazzo  CMEIAS: A Tool for Computational Microbial Ecology		
<b>9:40 – 9:50</b> <i>A5</i>	Stuart H. Gage Computational Ecology and Visualization Technologies		
<b>9:50 – 10:00</b> <i>A6</i>	George Garrity  Markup of Microbiological Data for Accelerated Publication in Print and Electronic Form		
<b>10:00</b> – <b>10:10</b> <i>A7</i>	Tom Getty Models of Signaling, Search, Discrimination and Selection		
<b>10:10 – 10:20</b> <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	
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# Session B in room 139 Radiology Building

9:00 – 9:10 B1	Simon J.L. Billinge Local Structure of Folded Proteins
<b>9:10 – 9:20</b> <i>B2</i>	James R. Cole The Ribosomal Database Project: Providing an Evolutionary Framework
<b>9:20 – 9:30</b> <i>B3</i>	Tien Yien Li Solving Polynomial Systems
9:30 – 9:40 B4	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
<b>9:40 – 9:50</b> <i>B5</i>	Jay I. Goodman  Altered DNA Methylation: An Epigenetic Mechanism Involved in Carcinogenesis
<b>9:50</b> – <b>10:00</b> <i>B6</i>	Erik Goodman Evolutionary Algorithms for Biological Science
<b>10:00 – 10:10</b> <i>B7</i>	Katharine Hunt Optical Trapping of Biomolecules
<b>10:10 – 10:20</b> <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 110 Radiology Building

10:50 – 11:00 CI	Wolfgang Bauer Cancer Detection via Determination of Fractal Cell Dimension		
11:00 – 11:10 C2	Andre Benard Image-Based Analysis of Heat Transfer in Biological Systems		
11:10 – 11:20 <i>C3</i>	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		
<b>11:30 – 11:40</b> <i>C5</i>	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		
<b>12:00 – 12:10</b> <i>C</i> 8	Fathi Salam Bio-Engineering: An Integrated Systems Approach		
12:10 – 1:10	Lunch		

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Chair: A. Jain	
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# Session D in room 139 Radiology Building

<b>10:50 – 11:00</b> <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	
<b>11:30 – 11:40</b> <i>D5</i>	M.F. Thorpe Protein Flexibility	
<b>11:40 – 11:50</b> <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	
<b>12:00 – 12:10</b> <i>D8</i>	Jack Preiss  Previous Funding Support for Computational Biology from the REA  Center of Protein Structure, Function and Design	
12:10 – 1:10	Lunch	

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Chair: Leslie Kuhn

# Session E in room 110 Radiology Building

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
E3	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
2:10 - 2:20	J. Potchen
<i>E7</i>	CT and MR Fly Through Images in Humans and ROC of Observer Performance

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Chair: Sakti Pramanik

# Session F in room 139 Radiology Building

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

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2:10-2:40	Group	disci	issions

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