

The 9th International Conference on Computational Methods in Systems Biology (CMSB 2011) is organized by Inria at Institut Henri Poincaré, 11 rue Pierre et Marie Curie, 75005 Paris, France, on September 21-23, 2011.

The HiBi workshop on High-Performance Computational Biology is merged in CMSB. The HiBi track is composed of the High Performance computing session plus a few papers in other sessions.

Wednesday, September 21, 2011

13.00-13.50 Registration

13.50-14.00 **Welcome address: François Fages**

14.00-15.00 **Invited talk: Denis Thieffry.** Logical modelling of haematopoietic cell fate decision.

15.00-15.30 **Student best paper award talk:** Jimmy Omony, Astrid R. Mach-Aigner, Leo H. De Graaff, Gerrit Van Straten and Anton J.B. Van Boxtel. Evaluation of design strategies for time course experiments in genetic networks: the *XlnR regulon* in *Aspergillus niger*

15.30-16.00 Coffee break

Stochastic models session - Chair Jérôme Féret

16.00-16.30 Sucheendra K. Palaniappan, S. Akshay, Blaise Genest and P.S. Thiagarajan. A Hybrid Factored Frontier Algorithm for Dynamic Bayesian Network models of Biopathways

16.30-17.00 Alessio Angius and Andras Horvath. Analysis of stochastic reaction networks with Markov reward models

17.00-17.30 Jacques Demongeot and Sylvain Sené. The singular power of the environment on stochastic nonlinear threshold Boolean automata networks

Thursday, September 22, 2011

9.00-10.00 **Invited talk: Pedro Mendes.** Kinetic modelling of large-scale metabolic networks.

10.00-10.30 Coffee break

High performance computing session - Chair Tommaso Mazza

10.30-11.00 Diego Darriba, Guillermo L. Taboada, Ramón Doallo and David Posada. HPC selection of models of DNA substitution

11.00-11.30 Martin Falk, Michael Klann, Michael Ott, Heinz Koepl and Thomas Ertl. Parallelized Agent-based Simulation on CPU and Graphics Hardware for Spatial and Stochastic Models in Biology

11.30-12.00 Hiroyuki Kuwahara and Xin Gao. An Efficient Parallel Stochastic Simulation Method for Analysis of Nonviral Gene Delivery Systems

12.00-12.30 Irene Sánchez-Linares, Horacio Pérez-Sánchez, Ginés D. Guerrero, José M. Cecilia and José M. García. Accelerating multiple target drug screening on GPUs

12.30-14.00 Lunch break

14.00-14.30 **Nvidia best paper award talk:** Eugen Czeizler, Vladimir Rogojin and Ion Petre. The phosphorylation of the heat shock factor as a modulator for the heat shock response

Simulation session - Chair Adelinde Uhrmacher

14.30-15.00 Ezio Bartocci, Elizabeth M. Cherry, James Glimm, Radu Grosu, Scott A. Smolka and Flavio Fenton. Toward Real-time Simulation of Cardiac Dynamics

15.00-15.30 Xuefeng Gao, Sabin Tabirca and Mark Tangney. Computer Simulation of Salmonella typhimurium Accumulation within Tumors

15.30-16.00 Coffee break

Discrete models session - Chair Olivier Roux

16.00-16.30 Hannes Klarner, Heike Siebert and Alexander Bockmayr. Parameter inference for asynchronous logical networks using discrete time series

16.30-17.00 Luca Cardelli and Cosimo Laneve. Reversible Structures

18.30-21.00 **Social dinner cruise - On Captain Fracasse, Île aux cygnes, metro Bir Hakeim**

Friday, September 23, 2011

9.00-10.00 **Invited talk: Thomas Henzinger.** Propagation Models for Computing Biochemical Reaction Networks

10.00-10.30 Coffee break

Spatial models session - Chair Monika Heiner

10.30-11.00 Mark Burkitt, Dawn Walker, Daniela Romano and Alireza Fazeli. Modelling Sperm Behaviour in a 3D Environment

11.00-11.30 Abhishek Murthy, Ezio Bartocci, Flavio Fenton, James Glimm, Scott Smolka and Radu Grosu. Towards Wave Curvature based Prediction of Onset of Atrial Fibrillation

11.30-12.00 Arne T. Bittig, Fiete Haack, Carsten Maus and Adelinde M. Uhrmacher. Adapting Rule-based Model Descriptions for Simulating in Continuous and Hybrid Space

12.00-12.30 Gordon Plotkin and Nicolas Oury. Coloured Stochastic Multi-Level Multiset Rewriting

12.30-14.00 Lunch break

Multi-scale models session - Chair Gordon Plotkin

14.00-14.30 Nicola Paoletti, Pietro Liò, Emanuela Merelli and Marco Viceconti. Osteoporosis: a multiscale modeling viewpoint

14.30-15.00 Andrea Degasperi and Muffy Calder. Multi-Scale Modelling of Biological Systems in Process Algebra with Multi-Way Synchronisation

15.00-15.30 Qian Gao, Fei Liu, David Gilbert, Monika Heiner and David Tree. A Multiscale Approach to Modelling Planar Cell Polarity in *Drosophila* Wing using Hierarchically Coloured Petri Nets

15.30-15.45 Closing remarks

15.45-16.30 Steering committee meeting

Invited talks

Thomas Henzinger - IST, Austria

Pedro Mendes - University of Manchester, UK

Denis Thieffry - Ecole Normale Supérieure, Paris, France

Program committee

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Carolyn Talcott SRI International, Menlo Park CA, USA

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Program

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