
3. WORKSHOP

COMPUTATIONAL BIOLOGY IN SAXONY: PROBLEMS AND PERSPECTIVES

December 02, 2004

Dresden University of
Technology



OBJECTIVE

The geographical vicinity of established and new research groups in Saxony in the field of computational biology offers a superb chance to exploit synergetic effects. The workshop aims at mutual information about ongoing activities and the identification of key topics for future common research projects. In particular, the Center for High Performance Computing (ZHR, Technische Universität Dresden) as organizer of the workshop offers its expertise and support for the solution of sophisticated problems in computational biology.

ORGANIZERS

Prof. Dr. Wolfgang E. Nagel
Dr. Andreas Deutsch
Zentrum für Hochleistungsrechnen
Technische Universität Dresden

WORKSHOP VENUE

TU Dresden
Zellescher Weg 12
01069 Dresden
Willers-Bau, Room C 207

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REPLY CARD

3. WORKSHOP COMPUTATIONAL BIOLOGY IN SAXONY

Name

First Name

Institution/Address

Phone

Fax

E-mail

- I take part in the workshop
- I do not take part in the workshop, but I would like to be informed about further workshops

**THURSDAY,
DECEMBER 02, 2004**

- | | | | |
|--------------|--|--------------|--|
| 09:00 | Welcome
Andreas Deutsch, ZHR, TU Dresden | 14:00 | Depolymerization of microtubules by motor proteins
Gernot Klein, Frank Jülicher, MPI-PKS, Dresden |
| 09:15 | Structural and game-theoretical analysis of metabolic networks
Stefan Schuster, Bioinformatics, Uni Jena | 14:30 | Kinetic models for chemosensitive movement
Angela Stevens, MPI-MIS, Leipzig |
| 10:00 | Genome annotation and protein folds
Mayte Pisabarro, BIOTEC, TU Dresden | 15:00 | Coffee |
| 10:30 | Coffee | 15:30 | Interacting particle systems modelling cell-cell adhesion and persistence
Anja Voss-Böhme
Mathematical Institute, TU Dresden |
| 11:00 | APArT: an automated protein annotation tool based on weak sequence similarities
Bianca Habermann, MPI-CBG, Dresden | 16:00 | A new model for myxobacterial pattern formation based on cell shape
Jörn Starruß, Thomas Bley
Bioprocess Engineering, TU Dresden |
| 11:30 | The small world of structural protein interactions and beyond
Michael Schröder, BIOTEC, TU Dresden | 16:30 | Multiple intracellular particles tracking: individual endosome behavior and pool statistic
Yannis Kalaidzidis, Marino Zerial
MPI-CBG, Dresden |
| 12:00 | Lunch | 17:00 | Morphogen gradient formation by transcytosis
Tobias Bollenbach, Karsten Kruse
MPI-PKS, Dresden |
| 13:00 | Data-intensive computing: perspectives for computational biology
Wolfgang E. Nagel, TU Dresden | 17:30 | Wrap up |
| 13:30 | Modelling mechanical oscillations in mitosis
Joe Howard, MPI-CBG, Dresden | | |