



# Strasbourg Summer School on Cheminformatics

Obernai, France, 22 - 25 June 2008

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## Sunday, 22 June

**14:00-17:00** Registration

**17:00-17:10** *Opening Remarks (A. Varnek, D. Rognan)*

**17:10-17:40** **T. Fujita**

QSAR: discovery and first steps

**17:40-18:20** **J. Gasteiger**

History and challenges of cheminformatics.

**18:20-19:00** **W. Warr**

Current trends in cheminformatics

**19:00** *Welcome party*

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## Monday, 23 June (morning session)

### *Molecular Similarity Analysis in Virtual Screening*

*Chairman: J. Gasteiger*

**8:30-09:30** **J. Bajorath**

Fingerprint Design and Molecular Complexity Effects

**9:30-10:30** **V. Gillet**

Diversity Analysis and Library Design

**10:30 10:45** *Coffee break*

**10:45-11:45** **G. Schneider**

*De novo* Design

## Monday, 23 June (after-noon session)

*Chairman: J. Bajorath*

**14:00 – 15:00**

**R. Glen**

Lessons learned from modelling bioactivity - what works and what doesn't

**15:00 – 16:00**

**T. Oprea**

Exploring novel estrogen receptors and more...

**16:00 – 16:15**

*Coffee break*

**16:15 – 17:30**

**N. Weill, G. Marcou, D. Horvath, A. Varnek and D. Rognan**

Tutorial: Impact of dataset composition on models performance

**17:45-19:00**

Meeting SFCi (*Albert Schweitzer hall*)

**17:45-19:00**

Software demonstration (*Hansi hall*)

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## Tuesday, 24 June (morning session)

*QSAR/QSPR strategies, ADME/Tox predictions*

*Chairwoman: W. Warr*

**8:30-09:30**

**R. Todeschini**

Molecular descriptors: an overview

**9:30-10:30**

**A. Tropsha**

The good, the bad and the ugly practices of QSAR modelling

**10:30 10:45**

*Coffee break*

**10:45-11:45**

**T. Langer**

Pharmacophore Approach in Drug Discovery

## Tuesday, 24 June (after-noon session)

*Chairman: R. Glen*

**14:00 – 15:00**

**J. Aires-de-Sousa**

Classification of chemical reactions

**15:00 – 16:00**

**I. Tetko, I. Baskin**

Machine learning methods in QSAR

**16:00 – 17:00**

**I. Baskin, I. Tetko and A. Varnek**

Tutorial on application of non-linear methods in chemistry (neural networks, support vector machines)

**17:00 17:15**

*Coffee break*

**17:00 - 18:00**

*Poster session (Bertoldi hall)*

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## Wednesday, 25 June (morning session)

*Bio- and Chemoinformatics: Key Technologies within the Drug Discovery Process*

*Chairman: R. Todeschini*

**8:30-09:15**

**J. Mestres**

*In silico* target profiling

**9:15--10:15**

**D. Rognan**

Docking and post-docking strategies

**10:15 10:30**

*Coffee break*

**10:30- 11:15**

**M. Stahl**

What Crystal Structure Databases Tell us about Conformational Preferences of Drug-like Molecules

**11:15 – 12:00**

**P. Ertl (Novartis)**

The Role of Cheminformatics in the Modern Drug Discovery Process

**Wednesday, 25 June (after-noon session)**

**14:00 – 16:00**      **Round Table** (*A. Tropsha, P. Vayer, L. Morin-Allory*)

- *Hot topics in chemoinformatics*
- *Teaching chemoinformatics*

**16:00**            **Closing Remarks** (*A. Varnek, D. Rognan*)