NETHERLANDS SOCIETY ON BIOMOLECULAR MODELLING
Fall meeting
November 29th 2017

Venue: Boothzaal, University Library De Uithof, Utrecht

The Netherlands Society on Biomolecular Modelling (NSBM) is an independent society that brings people together in The Netherlands that work on three dimensional structures of biomolecules and biomolecular simulations. The NSBM cordially invites you to participate in its 2017 fall meeting in Utrecht. During the meeting, the floor will be given to PhD students and post-docs who study protein structure and/or structural bioinformatics or related topics.

Location and dates

- Date: **November 29th**, 2017, 10:00-17:00
- Venue: Boothzaal, University Library De Uithof, Heidelberglaan 3, 3584 CS Utrecht
- http://www.uu.nl/universiteitsbibliotheek/locaties

Deadlines

- **Registration deadline to be considered for a short talk: November 17th**
- **Registration deadline (including lunch): November 22nd**

Application procedure

- Participants should send an email to register (required but free registration) to: g.nicolaes@maastrichtuniversity.nl
- ***Submission of an abstract is required if you want to present a poster.***
- Indicate in your email if you want be selected for a short oral presentation.

Note: The maximum number of participants is limited to 75! So please register on time.

Lead contact

- Dr. Gerry Nicolaes – g.nicolaes@maastrichtuniversity.nl

http://www.3dstructure-function.nl/nsbm-fall-meeting/

Preliminary Programme

- 09:30-10:00 Arrival and coffee
- 10:00-10:10 Welcome word
- 10:10-11:00 Keynote I
  - o **Prof. Jürgen Pleiss**, Institut für Technische Biochemie (ITB), Stuttgart University (D)
  - o *Data-integrated simulation of enzymes*
- 11:00-11:20 Short talk selected from abstracts
- 11:20-11:40 Short talk selected from abstracts
- 11:40-12:10 Lecture
  - o **Tom van den Bergh**, Bio-Prodict, Nijmegen (NL)

  12:10-12:30 Lecture
- o Short talk selected from abstracts
- 12:30-14:00 Lunch and **poster session**
- 14:00-14:50 Keynote II:
  - o **Prof. Chris Ulens**, KU Leuven (B).
  - o *Allosteric modulation of ligand-gated ion channels*
- 14:50-15:10 Short talk selected from abstracts
- 15:10-15:30 Short talk selected from abstracts
- 15:30-16:00 Coffee break
- 16:00-16:50 Keynote III:
  - o **Dr. Chris de Graaf**, VU Computational Medicinal Chemistry group, Amsterdam (NL)
  - o 3D-e-Chem: Structural cheminformatics tools for navigating protein-ligand interaction space 16:50 Closure

Costs

Attendance is free of charge but registration is required.
Lunch and coffee will be provided thanks to the financial support of the Nijmegen Centre for Molecular and Biomolecular Informatics.

Organized with the support of: