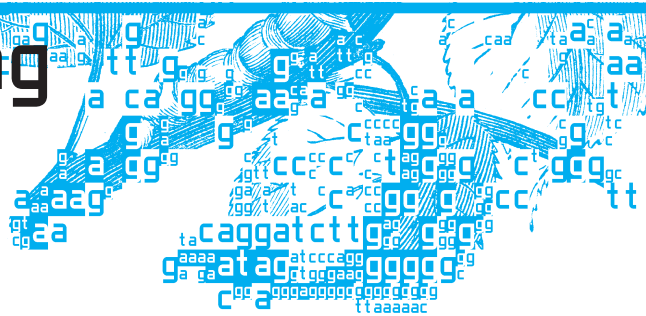


nbic thematic meeting



PROTEIN STRUCTURE BIOINFORMATICS

March 1st 2010, 10:00—17:00
UMC St Radboud, Nijmegen

PROGRAMME

- 10:00 Reception / registration / coffee
10:25 Welcome by G. Nicolaes, University of Maastricht
- 10:30 Dr. Elmar Krieger, Yasara; Graz, Austria
"Homology modelling with YASARA"
10:45 Discussion
- 11:30 Dr. Jürgen Haas, Swiss Institute of Bioinformatics (SIB)/Biozentrum;
Basel, Switzerland
"Preparing structures for bioinformatics, i.e. GROMACS"
12:15 Discussion
- 12:30 Lunch
13:30 Prof. dr. Bert de Groot, Max Planck Institute for Biophysical Chemistry;
University of Goettingen, Germany
"Using GROMACS in the modelling world"
14:15 Discussion
- 14:30 Dr. Sander Nabuurs, CMBI; UMC St Radboud Nijmegen
"Homology models and dynamics for drug design"
15:15 Discussion
- 15:30 Closing remarks by Ad van Gorp, Lead Pharma
- 15:35 Tea
16:00 Discussion about the organisation of protein structure bioinformatics
in the Netherlands (and its neighbours)
17:00 Drinks

LOCATION

UMC St Radboud
NCMLS Building, Colloquium Room,
8th Floor, Route 289
Geert Grooteplein Zuid 26-28
6525 GA Nijmegen

REGISTRATION

Please register via the NBIC website:
www.nbic.nl/about-nbic/events/nbic-events/