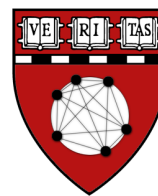


HARVARD MEDICAL SCHOOL
CENTER FOR MOLECULAR AND CELLULAR DYNAMICS
SBGRID NSF RESEARCH COORDINATION NETWORK



SCHRÖDINGER USER SYMPOSIUM

Schedule of Talks

Application of interaction fingerprints in Maestro for virtual screening and structure based design

Applications of Prime in Lead Optimization: MM-GBSA calculations and loop predictions on Src and Abl Kinases

Inhibiting Class II viral fusion: a small molecule approach

CONFIRM: Connecting Fragments in Receptor Molecules

Exploring the Moenomycin binding site with CombiGlide and Core Hopping

Modeling of protein complexes by docking and low resolution experiments

Claudio Chuaqui
AstraZeneca

David Dalgarno & Anna Kohlmann
Ariad Pharmaceuticals

Aaron Schmidt
Harvard Medical School

Eric Feyfant
Wyeth

Piotr Sliz
Harvard Medical School

Sandor Vajda
Boston University

August 15, 2008

8:30 AM – 1:30 PM

Ballard Room

Countway Library of Medicine, Harvard Medical School

Registration is required and space is extremely limited. To register, email cmcdadmin@crystal.harvard.edu by **August 13**. Your registration will be confirmed on a **first-come, first-served basis** as space allows.

Hosts: Piotr Sliz, Harvard Medical School
Woody Sherman, Schrödinger