SCHRÖDINGER USER SYMPOSIUM

Schedule of Talks

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

Applications of Prime in Lead Optimization: MM-GBSA calculations and loop predictions on Src and Abl Kinases
  David Dalgarno & Anna Kohlmann
  Ariad Pharmaceuticals

Inhibiting Class II viral fusion: a small molecule approach
  Aaron Schmidt
  Harvard Medical School

CONFIRM: Connecting Fragments in Receptor Molecules
  Eric Feyfant
  Wyeth

Exploring the Moenomycin binding site with CombiGlide and Core Hopping
  Piotr Sliz
  Harvard Medical School

Modeling of protein complexes by docking and low resolution experiments
  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
Woodo Sherman, Schrödinger