



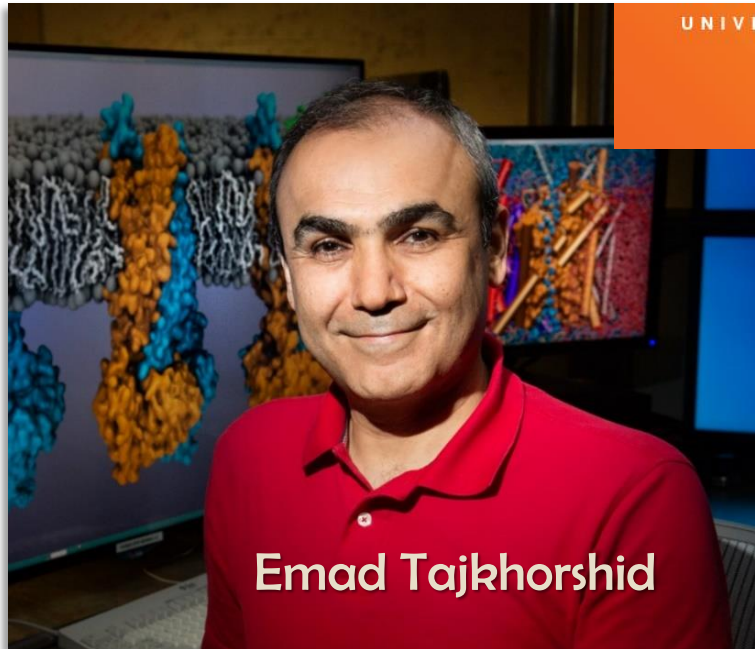
Hands-on Workshop on Computational Biophysics

Theoretical and Computational Biophysics Group (TCBG) and
National Center for Multiscale Modeling of Biological Systems (MMBioS)

June 28 -July 1, 2021

TCBG - Funded in 1989

“Bringing Physics to Life”



Emad Tajkhorshid

Professor of Biochemistry and Pharmacology,
Biophysics and Computational Biology,
Beckman Institute, U of Illinois, Urbana-
Champaign

UNIVERSITY OF

ILLINOIS

AT URBANA-CHAMPAIGN



Software	Citations
NAMD	15,000+
VMD	38,000+

MMBioS Funded in 2012

High Performance Computing BTRC for Multiscale Modeling of Biological Systems

Overarching biological theme:

- Spatial organization
- Temporal evolution

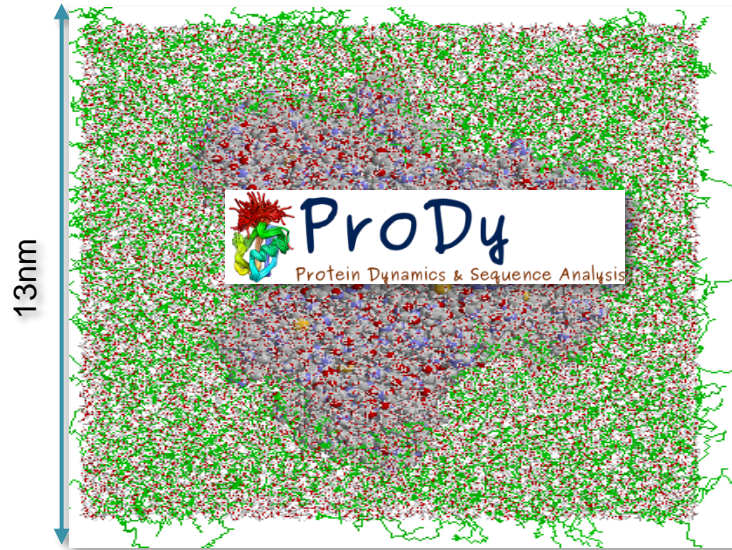
of

Synaptic Signaling & Regulation

NIGMS



Software development at multiple scale



to cellular architecture,



Ivet Bahar



Pemra Doruker



Burak Kaynak

Organization

Rozita Laghaei



Research Scientist, Pittsburgh
Supercomputing Center rlagha@psc.edu

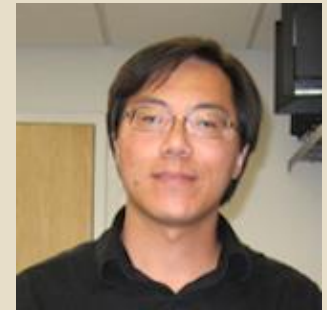


Ivan Cao-Berg
Research Software
Specialist, PSC



Anna Reinhard
Dept Comp & Sys Biol, Pitt
ANR189@pitt.edu

Administrators



Gengkon Lum, BS
Senior System Administrator
gengkon@pitt.edu



Afshin E Nasrabad, PhD
System Administrator
afe9@pitt.edu



Adam Kohlhaas
Dept Comp & Sys Biol, Pitt
KohlhaasA@pitt.edu



2015

Welcome!



2016



2018

Agenda

Day 1: Collective Dynamics of Proteins Using Elastic Network Models

9:30-10:00	Welcome and Brief Overview by Ivet Bahar
10:00-11:20	Elastic Network Models (ENMs) and Collective Motions. Basic Theory and Methods, <i>Ivet Bahar</i>
11:20-11:40	Break / Social
11:40-1:00	Ensemble Analysis of Structures for Inferring Functional Mechanisms, DynOmics Server, <i>Pemra Doruker</i>
1:00-1:20	Q & A and Lunch / Dinner break
2:00-3:30	ProDy API Basics , <i>Burak Kaynak</i> GNM Analysis of Equilibrium Dynamics, <i>Burak Kaynak</i>
3:30-3:50	Break / Social
3:50-5:45	ANM Analysis of Collective Motions and Visualization by NMWiz, <i>Burak Kaynak</i> Ensemble Analysis by ProDy, <i>Bentley Wingert</i>

Agenda

Day 2: Bridging Sequence, Structure and Function, Allostery, Druggability

10:00-11:20	Evol: Bridging Sequence and Structure,; Signature Dynamics of Protein Families, ENM for Membrane Proteins <i>Pemra Doruker</i>
11:20-11:40	Break / Social
11:40-1:00	Allostery and Druggability Simulations, <i>Burak Kaynak</i> and <i>Pemra Doruker</i> ,
1:00-1:20	Q & A and Lunch / Dinner break
2:00-3:30	Evol and SignDy , <i>Daniel Penaherrera</i> and <i>Bentley Wingert</i> Membrane ANM , <i>Daniel Penaherrera</i>
3:30-3:50	Break / Social
3:50-5:45	ESSA , <i>Burak Kaynak</i> Druggability Suite , <i>Jiyoung Lee</i>