

Martin Karplus Celebration Symposium

October 1, 2014



On computational methods applied to chemical and biological systems

The Mark Hopkins Hotel, 999 California Street, San Francisco, CA, USA

Preface

This symposium recognizes the scientific accomplishments of Martin Karplus and his students. It also celebrates Martin's receipt of the 2013 Nobel Prize in Chemistry for the development of multiscale models for complex chemical systems. Driven by a desire to discover new things and to understand biological processes at a fundamental level in chemical and physical terms, Martin has contributed in inestimable ways to the field of theoretical chemistry and to the development of molecular dynamics simulation of biomolecules. Early on, Martin concentrated on chemical physics and provided critical insights on topics including magnetic resonance and dynamics of fundamental reactions like $H + H_2$. At Harvard, he charted a new path that directed his efforts to biology, thus returning to what first attracted him to science as a young boy. As well as his own scientific work, Martin's contributions include his influence on the nearly 250 students and postdocs who have been members of his research group. He provided an environment with highest scientific standards and a need for rigorous, critical thinking. This symposium is the third such gathering to bring together past group members to meet, discuss, and this year, to celebrate together the well deserved Nobel Prize.



Karplusians :**1955-2014**

Ivana Adamovic
Yuri Alexeev
David H. Anderson
Ioan Andricioaei
Yasuhide Arata
Georgios Archontis
Gabriel G. Balint-Kurti
Christian Bartels
Paul Bash
Donald Bashford
Mark Bathe
Oren M. Becker
Robert Best
Anton Beyer
Robert Birge
Ryan Bitetti-Putzer
Arnaud Blondel
Stefan Boresch
John Brady
Bernard Brooks
Charles L. Brooks III
Thomas H. Brown
Robert E. Bruccoleri
Paul W. Brumer
Axel T. Brünger
Rafael P. Brüschweiler
Matthias Buck
Amedeo Caflisch
William J. Campion
William Carlson
David A. Case
Leo Caves
Thomas C. Caves
Marco Cecchini
John-Marc Chandonia
Ta-Yuan Chang
Xavier Chapuisat
Sergei Chekmarev
Rob D. Coalson
François Colonna-Cesari
Michael R. Cook
Qiang Cui
Tara Prasad Das
Annick Dejaegere
Philippe Derreumaux
Aaron Dinner
Uri Dinur
Roland L. Dunbrack, Jr.
Chizuko Dutta
Nader Dutta
Claus Ehrhardt
Ron Elber
Marcus Elstner
Byung Chan Eu
Jeffrey Evanseck
Erik Evensen
Jeffrey Evenson
Thomas C. Farrar
Martin J. Field
Stefan Fischer
David L. Freeman
Thomas Frimurer
Kevin Gaffney
Jiali Gao
Yi Qin Gao
Bruce Gelin
R. Benny Gerber
Paula M. Getzin
Debra A. Giammona
Martin Godfrey
Andrei Golosov
David M. Grant
Daniel Grell
Peter Grootenhuis
Hong Guo
Ogan Gurel
Robert Harris
Karen Haydock
Russell J. Hemley
Jeffrey C. Hoch
Milan Hodoscek
Gary G. Hoffman
L. Howard Holley
Barry Honig
Victor Hruby
Rod E. Hubbard
Robert P. Hurst
Vincent B.-H. Huynh
Toshiko Ichiye
K. K. Irikura
Alfonso Jaramillo
Tom Jordan
Diane Joseph-McCarthy
Sun-Hee Jung
C. William Kern
William Kirchhoff
Burton S. Kleinman
Gearld W. Koeppl
H. Jerrold Kolker
Yifei Kong
Lewis M. Koppel
J. Kottalam
Felix Koziol
Christoph Kratky
Sergei Krivov
Olga Kuchment
Krzysztof Kuczera
John Kuriyan
Joseph N. Kushick
Peter W. Langhoff
Antonio C. Lasaga
Frankie T. K. Lau
Themis Lazaridis
Fabrice LeClerc
Angel Wai-mun Lee
Irwin Lee
Sangyoub Lee
Ming Lei
Ronald M. Levy
Xiaoling Liang
Carmay Lim
Xabier Lopez
Guobin Luo
Paul D. Lyne
Jianpeng Ma
Alexander D. MacKerell, Jr.
Christoph Maerker
Paul Maragkakis
Marc Martí-Renom
Jean-Louis Martin
Carla Mattos
J. Andrew McCammon
H. Keith McDowell
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Morten Meeg
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Stephen Michnick
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Andrew Miranker
Keiji Morokuma
A. Mukherji
Adrian Mulholland
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Setsuko Nakagawa
Kwango Nam
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Neil Ostlund
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Emanuele Paci
Yuh-Kang Pan
C.S. Pangali
Richard W. Pastor
Lee Pedersen
David Perahia
Robert Petrella
B. Montgomery Pettitt
Ulrich Pezzeca
Richard N. Porter
Jay M. Portnow
Carol B. Post
Lawrence R. Pratt
Martine Prévost
Blaise Prod'hom
Jingzhi Pu
Dagnija Lazdins Purins
Lionel M. Raff
Mario Raimondi
Francesco Rao
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Swarna Yeturu Reddy
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Bruno Robert
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Benoît Roux
Andrej Sali
Daniel Saltzberg
Michael Schaefer
Michael Schlenkrich
David M. Schrader
John C. Schug
Klaus Schulten
Eugene Shakhnovich
Moshe Shapiro
Ramesh D. Sharma
Isaiah Shavitt
Henry H.-L. Shih
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Balvinder Singh
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Sung-Sau So
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Collin Stultz
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S. Swaminathan
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Wei Yang
Robert Yelle
Darrin York
Hsiang-ai Yu
Guishan Zheng
Yaoqi Zhou
Vincent Zoete

Program

Wednesday, October 1, 2014

The Mark Hopkins Hotel, 999 California Street, San Francisco

7:30 - 8:55 AM Registration and Breakfast
Location: Peacock Court

8:55 - 9:00 AM **Andrej Sali**, University of California, San Francisco, CA, USA
Welcome Remarks
Location: Room of the Dons

Chair

Toshiko Ichiye, Georgetown University, Washington, DC, USA

9:00 - 9:25 AM **Andy McCammon**, University of California, San Diego, CA, USA
Molecular Dynamics and Drug Discovery

9:25 - 9:55 AM **Peter Rosky**, University of Texas, Austin, TX, USA
Back to the Future: QM/MM Simulation of Exciton Dynamics in Conjugated Molecular Materials

9:55 - 10:25 AM **Iwao Ohmine**, Institute for Molecular Science, Okazaki, Japan
Water Dynamics; fluctuation, phase transitions and reactions

10:25 - 10:45 AM Coffee break

Chair

Tom Simonson, Ecole Polytechnique, Palaiseau, France

10:45 - 11:15 AM **Adrian Mulholland**, University of Bristol, Bristol, UK
Enzyme catalytic mechanisms from combined quantum mechanics/molecular mechanics (QM/MM) modeling

11:15-11:45 AM **Marco Cecchini**, Université de Strasbourg, France
Computational approaches to chemomechanical transduction: the myosin molecular motor

11:45 -12:15 PM **Bernard R. Brooks**, National Institutes of Health, Bethesda, MD, USA
35 years of CHARMM

12:15 - 1:40 PM Lunch
Location: Peacock Court

Chair

Amedeo Caflisch, University of Zurich, Switzerland

- 1:40 - 2:10 PM **YiQin Gao**, Peking University, Beijing, P.R. China
DNA allostery and its sequence dependence
- 2:10 - 2:40 PM **Collin Stultz**, Massachusetts Institute of Technology, Cambridge, MA, USA
Modeling Intrinsically Disordered Proteins: Application to Amyloid-beta
- 2:40 - 3:10 PM **Paul Maragakis**, DE Shaw Research, New York, NY, USA
Targeting “undruggable” proteins using long molecular dynamics simulations
- 3:10 - 3:30 PM Coffee break

Chair

John E. Straub, Boston University, Boston, MA, USA

- 3:30 - 4:00 PM **Carol B. Post**, Purdue University, West Lafayette, IN, USA
New structural features of kinases as potential targets to alter protein-protein interactions
- 4:00 - 4:30 PM **Dennis Vitkup**, Columbia University, New York, NY, USA
Systems Biology of Psychiatric Disorders
- 4:30 - 5:00 PM **Axel Brunger**, Stanford University, Stanford, CA, USA
Molecular mechanisms of SNARE-mediated membrane fusion, calcium triggering, and recycling
- 5:00 - 5:15 PM Summing up
David Chandler, University of California, Berkeley, CA, USA
- 6:15 - 7:30 PM Reception
Location: Room of the Dons
- 7:30 - 10:00 PM Dinner
Location: Peacock Court

Moderator

John Kuriyan, University of California, Berkeley, CA, USA

- 9:10 - 10:00 PM Address
Greg Petsko, Weill Cornell Medical School, New York, NY, USA

Short Talks

Tuesday, September 30, 2014

Helen Diller HD 160, UCSF Cancer Center, 3rd Street, San Francisco, CA 94158

12:00 - 5:00 PM

Name	Title
Ogan Gurel	<i>Protein Electrodynamics and Terahertz Medicine</i>
Matthias Buck	<i>An overview of simulations on the plexin and Eph transmembrane receptor systems.</i>
Michael Weiss	<i>How insulin binds: from structure of a model hormone-receptor complex to next-generation analog design.</i>
Wei Yang	<i>Predictive Sampling of Protein Long-Timescale Motions</i>
Robert Best	<i>Binding mechanism of intrinsically disordered proteins from molecular simulation</i>
Peter Langhoff	<i>On the Question of Atoms and Bonds in Molecules</i>
Carla Mattos	<i>Ras GTPase: Effects of oncogenic mutants</i>
Yuri Alekseev	<i>Ab initio dynamics of proteins by using FMO</i>
Veronica Vaida	<i>Reactions at water surfaces: aerosols, vesicles and bubbles</i>
Wonmuk Hwang	<i>Simulating biomolecular processes at multiple scales</i>
Kwangho Nam	<i>Multi-scale QM/MM model for the study of catalytic mechanisms of protein tyrosine kinase</i>
Ioan Andricioaei	<i>Multiscale Motions in Nucleic Acids</i>
John-Marc Chandonia	<i>The SCOPe database (http://scop.berkeley.edu)</i>
Krzysztof Kuczera	<i>Uncovering the complex pathway of helix formation</i>
Hon Guo	<i>QM/MM investigation of substrate-assisted catalysis</i>
Jingzhi Pu	<i>How ABC-transporters hydrolyze ATP: development and application of QM/MM methods</i>

Symposium Organizing Committee

Andrej Sali, University of California, San Francisco
David Chandler, University of California, Berkeley
Matt Jacobson, University of California, San Francisco
John Kuriyan, University of California, Berkeley
Susan Marqusee, University of California, Berkeley
Carol Post, Purdue University

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