SHAKH-50 SYMPOSIUM PROGRAM DEC 8, 2007 Pfizer Lecture Hall, 12 Oxford St., Cambridge, MA 02138

9:00 - 9:30 a.m.	Coffee/Tea/Registration	Outside of Pfizer Lecture Hall
9:30 -11:20 a.m.	Arup Chakraborty	Understanding Adaptive Immunity:
Folding and	MIT	A crossroad of the physical and life sciences
theory	Nikolay Dokolyan	Dynamics in Computational Protein Engineering
	UNC Chapel Hill	
	Estelle Pitard	Thermodynamics of histories for models of glassy dynamics
	CNRS, Montpellier, France	
	Phillip Geissler	Dynamics of organization, and disorganization, at the nanometer scale
	UC Berkeley	
11:20 -1:40 a.m.	Coffee Break	Outside of Pfizer Lecture Hall
11:40 a.m1 p.m.	William Eaton	Single molecule FRET studies of protein folding
Folding and	NIH	
drug design	Guido Tiana	Breaking the toy: folding inhibition of lysozyme
	University of Milano, Italy	
	Jun Shimada, NYU and	Some Keys to Building Bridges between Theory
	Alexey Ishchenko, <i>Vitae</i>	and Experiment in Drug Discovery
	Pharmaceuticals	
1:00 - 2:00 p.m.	Lunch	Chemistry Department Center, 12 Oxford St., 2 nd Floor Mallinckrodt
2:00 - 3:15 p.m.	Eric Deeds	Coupling signal responses through shared components
Molecular and	Harvard Medical School	
cellular	Leonid Mirny	Cooperation among transcription factors: united they stand
biophysics	MIT	
	Boris Shakhnovich	TATA box is not enough: function and conservation of TBP bound
	Harvard University	promoters
3:15 - 3:45 p.m.	Coffee Break/Discussions	Outside of Pfizer Lecture Hall
3:45 - 5:30 p.m.	Edo Kussell	How to tell if a bacterium is sensing (without doing genetics.)
Evolution	NYU	
	Konstantin Zeldovich	Evolution, protein stability, and bacterial competition assays.
	Harvard University	
	Emmanuel Tannenbaum	A comparison of various asexual and sexual replication strategies
	GeorgiaTech & Ben-Gurion Univ.,	
	Israel	
	Igor Berezovsky	On the uniqueness of viral proteins.
	University of Bergen, Norway	
5:30 - 6:00 p.m.	Closing remarks	