

Carlos Xavier Hernández

Δ	А	Ы	r	6	c	e	

Genentech

Room S296 318 Campus Drive, Stanford, CA 94305 Telephone: (973) 280-3860 Email: cxh@stanford.edu URL: cxhernandez.com

Experience	Graduate Fellow in the Biophysics Program with Vijay Pande @ Stanford University	2013 - Present
	Undergraduate Researcher at the Center for Computational Biology and Bioinformatics with Raul Rabadan @ Columbia University Medical Center	2010 - 2013
	Undergraduate Fellow at the Department of Chemistry and Biochemistry with James Andrew McCammon @ University of California, San Diego	2012
	Undergraduate Fellow at the Department of Biochemistry and Molecular Biophysics with Stephen Goff @ Columbia University Medical Center	2011
Education	Stanford University Biophysics, PhD	Present
	Columbia University in the City of New York Applied Mathematics, BSc	2013
Skills	 Python Java MATLAB R UNIX HTML CSS JavaScript SQL Computational Biochemistry Web Mobile Development Machine Learning Data Mining 	
Awards	Graduate Research Fellowship National Science Foundation	2013
	ADVANCE Summer Institute Fellowship Stanford University	2013
	Exceptional Research Opportunities Program Fellowship Howard Hughes Medical Institute	2012
	Summer Undergraduate Research Fellowship	2011



Publications	MSMBuilder: Data Visualizations for Biomolecular Dynamics Hernández CX, Harrigan MP, Sultan MM, and Pande VS The Journal of Open Source Software. doi: 10.21105/joss.00188	2017
	MSMBuilder: Statistical Models for Biomolecular Dynamics Harrigan MP, Sultan MM, Hernández CX , Husic BE, and Pande VS Biophysical Journal. doi: 10.1016/j.bpj.2016.10.042	2017
	Osprey: Hyperparameter Optimization for Machine Learning McGibbon RT, Hernández CX , Harrigan MP, Kearnes S, Sultan MM, Jastrzebski S, Husic BE, and Pande VS The Journal of Open Source Software. doi: 10.21105/joss.00034	2016
	MDTraj: a modern, open library for the analysis of molecular dynamics trajectories McGibbon RT, Beauchamp KA, Schwantes CR, Wang LP, Hernández CX, Harrigan MP, Lane TJ, Swails JM, and Pande VS Biophysical Journal. doi: 10.1016/j.bpj.2015.08.015	2015
	Markov State Models Provide Insights into Dynamic Modulation of Protein Function Shukla D, Hernández CX , Weber JK, and Pande VS Accounts of Chemical Research. doi: 10.1021/ar5002999	2015
	Structure-based Network Analysis of an Evolved G-Protein Coupled Receptor Homodimer Interface Nichols SE*, Hernández CX *, Wang Y, and McCammon JA Protein Science. doi: 10.1002/pro.2258	2013
	Understanding the Origins of a Pandemic Virus Hernández CX, Khiabanian H, Chan J, and Rabadan R arXiv. arXiv: 1104.4568v1 [q-bio.PE]	2011
Press	Pandemic Flu Risk Raised by Lax Hog-Farm Surveillance Wired Science	2012
	The Origin and Evolution of a Pandemic Virus MAGNet Newsletter, 4:15-18	2011
Mentorship	Irfan Kherani Global Health and Health Policy Student @ Princeton University Thesis: Cross-Hemispheric Phylogenetic Analysis of 2009 H1N1 Pandemic Influenza	2011
	Ravi Vaidya High School Student Thesis: Distinct Circulation of Influenza Virus in Geographically Diverse U.S. Air Force Installations	2010