

Carlos Xavier Hernández

Research Scientist

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I am a Research Scientist at Facebook Research. Previously, I worked on neural interfaces at CTRL-labs. Prior to that, I did my thesis at Stanford with Vijay Pande on statistical modeling and representation learning of complex biomolecular dynamics.

Experience

Facebook Research Research Scientist	New York, NY, USA 2019 - PRESENT
CTRL-labs Research Scientist	New York, NY, USA 2018 - 2019
Stanford University NSF Graduate Research Fellow w/ Vijay S. Pande	Stanford, CA, USA 2013 - 2018

Education

Stanford University Ph.D. in Biophysics Thesis: "Towards a Deeper Understanding of Molecular Mechanics"	Stanford, CA, USA 2013 - 2018
Columbia University in the City of New York B.S. in Applied Mathematics	New York, NY, USA 2009 - 2013

Publications

Kinetic Machine Learning Unravels Ligand-Directed Conformational Change of μ -Opioid Receptor EN Feinberg, AB Farimani, CX Hernández, and VS Pande bioRxiv (doi: 10.1101/170886)	2018
Variational Encoding of Complex Dynamics CX Hernández*, HK Wayment-Steele*, MM Sultan*, BE Husic, and VS Pande Phys. Rev. E (doi: 10.1103/PhysRevE.97.062412)	2018
Using Deep Learning for Segmentation, and Counting within Microscopy Data CX Hernández, MM Sultan, and VS Pande arXiv (arXiv: 1802.10548)	2018
Markov State Models Provide Insights into Dynamic Modulation of Protein Function D Shukla, CX Hernández, JK Weber, and VS Pande Accounts of Chemical Research (doi: 10.1021/ar5002999)	2015
Structure-based Network Analysis of An Evolved G-Protein Coupled Receptor Homodimer Interface SE Nichols*, CX Hernández*, Y Wang, and JA McCammon Protein Science (doi: 10.1002/pro.2258)	2013
Understanding the Origins of a Pandemic Virus CX Hernández, J Chan, H Khiabani, and R Rabadan arXiv (arXiv: 1104.4568)	2011

Software

- MolEncoder: Molecular Autoencoder in PyTorch** 2017
 CX Hernández
[GitHub](https://github.com/cxhernandez/molencoder) (url: <https://github.com/cxhernandez/molencoder>)
 ● Python ★ 56 📄 10
- MDEntropy: Information-Theoretic Analyses for Molecular Dynamics** 2017
 CX Hernández and VS Pande
[The Journal of Open Source Software](https://doi.org/10.21105/joss.00427) (doi: 10.21105/joss.00427)
 ● Python ★ 19 📄 8
- MSMExplorer: Data Visualizations for Biomolecular Dynamics** 2017
 CX Hernández, MP Harrigan, MM Sultan, and VS Pande
[The Journal of Open Source Software](https://doi.org/10.21105/joss.00188) (doi: 10.21105/joss.00188)
 ● Python ★ 12 📄 13
- MSMBuilder: Statistical Models for Biomolecular Dynamics** 2017
 MP Harrigan, MM Sultan, CX Hernández, BE Husic, P Eastman, CR Schwantes, KA Beauchamp, RT McGibbon, and VS Pande
[Biophysical Journal](https://doi.org/10.1016/j.bpj.2016.10.042) (doi: 10.1016/j.bpj.2016.10.042)
 ● Python ★ 95 📄 69
- Osprey: Hyperparameter Optimization for Machine Learning** 2016
 RT McGibbon, CX Hernández, MP Harrigan, S Kearnes, MM Sultan, S Jastrzebski, BE Husic, and VS Pande
[The Journal of Open Source Software](https://doi.org/10.21105/joss.00034) (doi: 10.21105/joss.00034)
 ● Python ★ 71 📄 22
- MDTraj: A Modern, Open Library for the Analysis of Molecular Dynamics Trajectories** 2015
 RT McGibbon, KA Beauchamp, MP Harrigan, C Klein, JM Swails, CX Hernández, CR Schwantes, LP Wang, TJ Lane, and VS Pande
[Biophysical Journal](https://doi.org/10.1016/j.bpj.2015.08.015) (doi: 10.1016/j.bpj.2015.08.015)
 ● Python ★ 253 📄 163

Posters & Presentations

- Convolutional Neural Networks for Visual Recognition (CS231N)** Stanford, CA, USA
 Invited Presentation 2017
 “Using Deep Learning for Segmentation and Counting within Microscopy Data”
- Biophysical Society Meeting** Los Angeles, CA, USA
 Poster 2016
 “Intrinsic Disorder in the P53 C-Terminal Regulatory Domain Yields Multiple Pathways for Folding-Upon-Binding”
- Workshop on Molecular and Chemical Kinetics** Berlin, DEU
 Poster 2015
 “Inferring Causality Along Transition State Pathways”

Honors & Awards

- 2013 ADVANCE Summer Research Fellowship, Stanford University
- 2013 Graduate Research Fellowship, National Science Foundation
- 2012 EXROP Undergraduate Research Fellowship, Howard Hughes Medical Institute
- 2011 Genentech Summer Undergraduate Research Fellowship, Columbia University - Genentech

Press

- 2012 Pandemic Flu Risk Raised by Lax Hog-Farm Surveillance, Wired Magazine
- 2011 The Origin and Evolution of a Pandemic Virus, MAGNet Newsletter