Maulik Nariya

Curriculum Vitae

Postdoctoral fellow

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Education

2012–2018 **Ph.D. Physics**

University of Kansas, Lawrence

Dissertation: Mathematical Model of Length Control in the Type III Secretion

System

Advisors: Eric Deeds and Jack Shi

2007–2010 M.S. Physics

University of Pune, Pune

Thesis: Review of Neutrino Oscillations

Advisor: S. Uma Sankar

2004–2007 B.S. Physics

St. Xavier's College, Mumbai

Experience

2018–present Postdoctoral fellow

Department of Systems Biology, Laboratory of Systems Pharmacology

Harvard Medical School Advisor: Peter Sorger

Skills

Proficient with data mining and machine learning in Python

Experience with other languages Fortran, C++

Proficient with Linux-based research computing

Capable of software development and version control using GitHub

Experience with containerizing software tools

Research projects

2019—present Improved estimate of machine learning predictor performance in presence of known confounders

In this project we highlight useful properties of leave-pair-out cross-validation method for biological datasets (RNA-seq data for breast cancer cells and Alzheimer's patient data), in particular we propose a method that helps improve

the estimate of predictor performance. (manuscript in preparation)

2018–present Predicting drug response from baseline omics profiles of breast cancer cells

NIH LINCS funded project at the Laboratory of Systems Pharmacology, Harvard Medical School. In this work we used baseline datasets, RNA-seq, proteomics and phosphoproteomics, of 69 breast cancer cells to build predictive models using a Random Forest Regressor to predict drug response across a variety of breast cancer drugs. (manuscript in preparation)

2014–2018 Mathematical modeling for length control in type III secretion system

Ph.D. dissertation defended at Department of Physics and Astronomy in collaboration with Center for Computational Biology, University of Kansas. This work involves making mathematical models and performing stochastic simulation of biochemical reactions involved in the growth of the type II

secretion injectisome.

2016–2018 Comparative characterization and biosimilarity assessment of drug samples using data mining and machine learning techniques

FDA funded project at the Macromolecule and Vaccine Stabilization Center and Center for Computational Biology, University of Kansas. The goal of this project

was to model the assessment of biosimilar drugs based on the data available from biological, chemical and physical assays.

Publications

- 1. **Nariya M.K.**, Sorger P.K., Sokolov A., Leave-pair-out cross-validation allows for a robust evaluation of model performance in presence of outliers and known confounders (manuscript in preparation)
- 2. Subramanian K., Mills C.E., **Nariya M.K.**, Chen C., Hafner M., Sokolov A., Boswell S., Everley R.A., Berberich M.J., Kalocsay M., Gaudio B., Victor C., Chung M., Bradshaw G., Sorger P.K., Predicting drivers of drug response from baseline omics data across breast cancer cells and models (manuscript in preparation)
- 3. Kalocsay M., Berberich M.J., Everley R.A., **Nariya M.K.**, Chung M., Gaudio B., Victor C., Bradshaw G.A., Hafner M., Sorger P.K., Mills C.E., Subramanian K. Data Descriptor: Proteomic profiling across breast cancer cells and models (in review with *Scientific Data*)
- 4. Nariya M.K., ShiJ.J., Mallela A., and Deeds E.J., Robustness and Evolution of Length Control Strategies in the Type III Secretion System and Flagellar Hook (in review at *Biophysical Journal*)
- 5. Mallela A., **Nariya M. K.**, and Deeds E. J. Crosstalk and Ultra-sensitivity in Protein Degradation Pathways
- 6. **Nariya M. K.**, Israeli J., Shi J. J., and Deeds E. J. Mathematical Model for Length Control by the Timing of Substrate Switching in the Type III Secretion System, *PLoS Comp Biol* 12(4): e1004851. DOI: 10.1371/journal.pcbi.1004851
- 7. Nariya M. K., Kim J. H., Xiong J., Kleindl P. A., Hewarathna A. N., Joshi S. B., Schöneich C., Forrest M. L., Middaugh C. R., Volkin D. B., and Deeds E. J. Comparative Characterization of Crofelemer Samples Using Data Mining and Machine Learning Approaches with Analytical Stability Data Sets, *J Pharm Sci*, 106(11): 3270–3279, DOI: 10.1016/j.xphs.2017.07.013
- 8. Hewarathna A., Mozziconacci O., **Nariya M. K.**, Kleindl P. A., Xiong J., Fisher A., Joshi S. B., Middaugh C. R., Forrest M. L., Volkin D. B., and Deeds E. J. Chemical Stability of the Botanical Drug Substance Crofelemer: A Model System for Comparative Characterization of Complex Mixture Drugs, *J Pharm Sci*, 106(11): 3257–3269, DOI: 10.1016/j.xphs.2017.06.022
- 9. Kleindl P. A., Xiong J., Hewarathna A., Mozziconacci O., **Nariya M. K.**, Fisher A., Deeds E. J., Joshi S. B., Middaugh R. C., Volkin D. B., and Forrest M. L. The Botanical Drug Substance as a Model System for Comparative Characterization of Complex Mixture Drug, *J Pharm Sci*, 106(11): 3242–3256, DOI: 10.1016/j.xphs.2017.07.012

Conference presentations

2018	Predicting drug response from baseline omics profile of breast cancer cells. Poster presentation at HiTS Annual Symposium, Boston, MA
2017	Mathematical Modeling of Ruler Mechanism for Length Control in Type III Secretion System. Poster presentation at $11^{\rm th}$ Annual q-bio Conference, Rutgers University, New Brunswick, NJ
2016	Mathematical Modeling for Type III Secretion System. Poster presentation at 10 th Annual q-bio Conference, Vanderbilt University, Nashville TN

Awards

2018	NIH LINCS fellowship — Harvard Medical School
2017	Graduate Scholarly Presentation Travel Fund — University of Kansas
2014	E. E. Slossen Award for Outstanding Graduate Teaching Assistants — University of Kansas

Teaching Experience

Statistical Physics — University of Kansas

Lead discussion, graded homework assignments, and designed exams for undergraduate level course on Statistical Physics

General Physics Laboratory — University of Kansas

Taught several general physics laboratory course (including one honors course), delivered instructions for conducting the laboratory experiment, graded laboratory reports and conducted final exams

References

Peter Sorger, Ph.D.

Otto Krayer Professor of Systems Biology
Director of Harvard Program in Therapeutic
Sciences
Harvard Medical School, Boston MA
peter_sorger@hms.harvard.edu

Eric Deeds, Ph.D.

Professor Integrative Biology and Physiology University of California Los Angeles, Los Angeles deeds@ucla.edu

Artem Sokolov, Ph.D.

Instructor in Biomedical Informatics
Director of Informatics and Modeling at the
Laboratory of Systems Pharmacology
Harvard Medical School, Boston MA
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