# MICHELLE GILL, PH.D.

# Data scientist, biophysicist

■ New York, NY

**\** 203-901-4028

nlgill ?

themodernscientist.com

in michellelynngill

# **EDUCATION**

**Ph.D. Molecular Biophysics & Biochemistry** 2006, Yale University, *New Haven, CT* 

**B.S. Biochemistry**, Summa Cum Laude 2001, University of Kansas, Lawrence, KS

# **SKILLS**

Programming: Python, Matlab, C, R, Shell

Big data: Spark, Dask

Machine learning: Supervised and unsupervised algorithms, Clustering, Natural language processing, Principal component analysis, Recommender systems, Computer vision, Neural networks, Deep learning, Compressed sensing

**Statistics:** Regression (linear, logistic, non-linear), Monte Carlo simulations

Databases: PostgreSQL, MongoDB

Other: Unix, Git, AWS, Docker, Flask, LaTeX

## **AWARDS**

- Ruth L. Kirschstein National Research Service Postdoctoral Fellowship
- NSF Graduate Research Fellowship
- Barry M. Goldwater Scholar
- Kansas Board of Regents full-tuition merit scholarship

# **EXPERIENCE**

### **Senior Data Scientist**

2016 - Present

Metis

- Designed and created Hadoop and Spark machine learning and NLP curriculum using self-made Docker containers
- Conducted corporate trainings focused on Python, Spark, Hadoop, and Hive
- Developed 12-week machine learning course for F100 company
- Co-instructed 12-week data science bootcamps
- Developed and conducted take home coding exercise to assist with interview preparation

### **Research Scientist**

2014 - 2016

National Cancer Institute

- Developed <u>NESTA-NMR</u>, which enables up to 100X faster acquisition of large (~10 GB) experimental data sets
- Created website and documentation for NESTA-NMR
- Elucidated mechanisms of cancer pathway using highresolution experimental techniques

### **Postdoctoral Research Fellow**

2008 - 2014

Columbia University Medical Center

- Used Monte Carlo simulations to model effect of physical changes on enzyme activity
- Developed <u>MFOutParser</u>, a Python library that parses a challenging text format, enabling 10X faster analysis times
- Member of team that studied anti-oncogenic associated mechanisms of substrate binding to AlkB, a DNA repair enzyme

### **Postdoctoral Research Fellow**

2007 - 2008

University of Kansas

 Developed principal component analysis-based method to visualize changes in vaccine structure that are critical for efficacy

### Consultant

2006 - 2007

The Boston Consulting Group

- Developed Excel-based statistical tools and Access database for organizational streamlining of pharmaceutical client
- Part of team that developed municipal bond investment strategy for financial services client
- Member of team awarded 2007 Global Strategy Olympics Prize for pharmaceutical client work

# PRESENTATIONS & PROJECTS

- "Learning from Text: Natural Language Processing with Python", Tutorial, ODSC East, Boston
- Created wine label recognition application computer vision
- <u>pdLSR</u> is a library for performing linear and non-linear least squares regression in a dataframe-aware fashion

Updated: 6/6/2017

Current version: http://resume.michellelynngill.com