Daniel L. Parton

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Technical Skills

Python 7 years experience; numerical/scientific computing, data handling, parallel computing, bioin-

formatics, scalable code design; packages—numpy, pandas, scikit-learn, theano (some),

SQLAlchemy, lxml, Flask, jinja, matplotlib, mpi4py, nosetests, setuptools, conda.

Full-stack web development

2 years experience; JavaScript, HTML5, CSS3, knockout.js, Bootstrap, Python Flask, MySQL,

SQLite, RESTful APIs, nginx, Apache.

Other languages bash, SQL, XPath. Some experience: C, C++, R, MATLAB, CUDA.

General GitHub, git, svn, Travis CI, Sphinx, Docker, Amazon AWS, vim, LATEX, OS X, Linux, Windows.

Experience

2012-pres

MEMORIAL SLOAN KETTERING CANCER CENTER, NEW YORK, NY
Postdoctoral Research Fellow, Computational Biology Center | Advisor: John D. Chodera

• Investigation of conformational networks in cancer-associated proteins using GPU-accelerated simulations, clustering techniques and Markov models.

- Developed MSMSeeder—pipeline for generating diverse structural ensembles as seeds for massively parallel simulations of proteins.
- Developed TargetExplorer and kinomeDB—database application for aggregating biological data for a target protein family, with RESTful API and frontend web client (in development).
- Data-driven selection of kinase sequences suitable for bacterial expression.
- 3 first-author papers in preparation.
- Closely involved with design and purchase of a HPC cluster (960 cores, 120 GPUs) and an integrated robotic platform for biological experiments.
- Attended Hacker School—a three month, self-driven, collaborative programming course. I worked
 primarily on my research projects, particularly focusing on scalable software design. I also completed
 online courses in machine learning and algorithms, a Kaggle competition, and a JavaScript game.

2011-2012

University of Chicago

Postdoctoral Scholar, Department of Chemistry | Advisor: Gregory A. Voth

• Applied advanced computer simulation techniques and free energy calculations to the study of the actin cytoskeleton. One paper as first author (in press).

2007-2011

University of Oxford

Graduate Student Researcher, Department of Biochemistry

Advisors: Mark S. P. Sansom (Oxford) and Ross Nobes (Fujitsu Laboratories of Europe)

 Studied complex multi-protein systems using physics-based molecular simulations, developed software tools for simulation analysis, contributed to MDAnalysis, and led organic chemistry tutorials.
 Co-authored 5 papers, 3 as first author. Results presented at many national and international conferences, and in national scientific magazines.

2005-2007

UNIVERSITY OF BRISTOL AND GLAXOSMITHKLINE, UK

Undergraduate Researcher / Industrial Placement Student | Advisor: Chris Willis

• Conducted the design and synthesis of drug candidates, as a member of medicinal chemistry teams at Bristol (6 months) and GSK (1 year). Co-authored one paper.

Education

2007–2011 D.PHIL. in Biochemistry, University of Oxford, UK
2003–2007 M.Sci. in Chemistry with Industrial Experience, University of Bristol, UK