

<b>Scott Clark</b> scott@scottclark.io @DrScottClark		<b>June 30, 2016</b> www.scottclark.io Not looking for new job opportunities
<b>Education</b>		
<ul style="list-style-type: none"> <li> <b>Cornell University</b>  <i>Ph.D. Applied Mathematics, M.S. Computer Science</i> </li> </ul>		Ithaca, NY 2008 - 2012
<ul style="list-style-type: none"> <li> <ul style="list-style-type: none"> <li>Department of Energy Computational Science Graduate Fellow (Full Scholarship, 4 years)</li> <li>Machine Learning, Data Science, Data Mining in bioinformatics and global optimization</li> </ul> </li> </ul>		
<ul style="list-style-type: none"> <li> <b>Oregon State University</b>  <i>B.Sc. Mathematics, B.Sc. Computational Physics, B.Sc. Physics</i> </li> </ul>		Corvallis, OR 2004 - 2008
<ul style="list-style-type: none"> <li> <ul style="list-style-type: none"> <li>Graduated Magna Cum Laude with minors in Actuarial Sciences and Mathematical Sciences</li> </ul> </li> </ul>		
<b>Research and Work Experience</b>		
<ul style="list-style-type: none"> <li> <b>SigOpt Inc - <a href="https://sigopt.com">https://sigopt.com</a></b>  <i>Co-founder and CEO</i> </li> </ul>		San Francisco, CA November 2014 - current
<ul style="list-style-type: none"> <li> <ul style="list-style-type: none"> <li>Helping lead a world class team to eliminate expensive trial and error from every experts workflow. Using cutting edge optimization behind a simple API to help tune machine learning models and build better products in a variety of fields.</li> </ul> </li> </ul>		
<ul style="list-style-type: none"> <li> <b>Yelp Inc</b>  <i>Data Mining Engineer and Lead on Ad Targeting Team</i> </li> </ul>		San Francisco, CA July 2012 - December 2014
<ul style="list-style-type: none"> <li> <ul style="list-style-type: none"> <li><b>Optimization:</b> Co-developed and led team for MOE: the Metric Optimization Engine (<a href="https://github.com/Yelp/MOE">github.com/Yelp/MOE</a>, an <b>open source</b> optimization framework), found significant gains in different metrics across the organization using Bayesian Global Optimization algorithms.</li> <li><b>Targeting:</b> Implemented multi-armed bandit strategies for ad selection, sole targeting engineer on mobile app ads rollout, developed new location-based targeting algorithms, advised and helped develop other machine learning and math based targeting projects.</li> <li><b>Recruiting:</b> Created, implemented, and directed <a href="https://yelp.com/dataset_challenge">yelp.com/dataset_challenge</a>, gave tech talks across the country, led events, gave hundreds of technical interviews, closed candidates.</li> </ul> </li> </ul>		
<ul style="list-style-type: none"> <li> <b>Bloomberg LP</b>  <i>Financial Software Development Intern</i> </li> </ul>		New York, NY Summer 2011
<ul style="list-style-type: none"> <li> <ul style="list-style-type: none"> <li>Implemented statistical models to perform forward and backward portfolio analysis</li> </ul> </li> </ul>		
<ul style="list-style-type: none"> <li> <b>DOE Joint Genome Institute (Lawrence Berkeley National Lab)</b>  <i>Researcher in Analysis Group under Dr. Zhong Wang</i> </li> </ul>		Walnut Creek, CA Summer 2010

- Used machine learning to mine TBs of genome data efficiently using novel likelihood function

- **Los Alamos National Laboratory** Los Alamos, NM  
*Researcher in Metagenomics Group under Dr. Nick Hengartner* Summer 2009

- Used statistical models to discover sequence alignments using parallel algorithms on GPUs

- **Oregon State University** Corvallis, OR  
*Research Assistant under Prof. Malgorzata Peszynska and Prof. Rubin Lau-* 2005-2008  
*dau*

- Finite element analysis with uncertainty and web-based teaching in Java

- **Max Plank Institute for the Physics of Complex Systems** Dresden, Germany  
*NSF REU Research Assistant under Prof. Steven Tomsovic* Summer 2007

- Research on extreme value statistics in MATLAB and FORTRAN

- **University of California: Davis** Davis, CA  
*NSF REU Research Assistant under Prof. Daniel Cox* Summer 2006

- Computational biophysics research as applied to protein folding in Java

## Writing and Awards

- **2016 Forbes 30 Under 30:** Enterprise Tech. <http://onforb.es/10ILpBZ>
- **Department of Energy Computational Science Graduate Fellow:** Four year full fellowship. ~20 awarded nationally per year. Won the Communicating Science award ([bit.ly/VbcTZK](http://bit.ly/VbcTZK)).
- **SigOpt Blog:** Posts talking about using SigOpt to optimize everything ([blog.sigopt.com](http://blog.sigopt.com)).
- **Yelp Blog:** Wrote several posts announcing the open sourcing of MOE, the Yelp Dataset Challenge and more. [bit.ly/1x73xdr](http://bit.ly/1x73xdr), [bit.ly/1oCCZvv](http://bit.ly/1oCCZvv), [bit.ly/1s0sEBS](http://bit.ly/1s0sEBS), [bit.ly/1p1X7Hk](http://bit.ly/1p1X7Hk)
- **Press:** WSJ: [on.wsj.com/Va0vqQ](http://on.wsj.com/Va0vqQ), Cornell: [bit.ly/1oB2dzm](http://bit.ly/1oB2dzm), DIEXIS: [bit.ly/1oofb14](http://bit.ly/1oofb14)

## Skills

- **Numerical Analysis and Computer Science:** Machine Learning, Data Mining, Optimization, Computational Science, Artificial Intelligence, Linear Algebra, Monte Carlo Methods, ODEs, PDEs, Iterative Methods, Parallel Programming, Distributed Systems, Data Structures
- **Tech Stack:** Python, numerical libraries, linux, git, vim
- **Public Speaking:** I've given several hundred technical talks to audiences at machine learning conferences, Fortune 500 boards, and beyond.
- Exploring and implementing ideas. Give me an API/dataset and a problem and I will figure it out.

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## Selected Open Source Projects

- **SigOpt Examples** ([github.com/sigopt/sigopt-examples](https://github.com/sigopt/sigopt-examples)) Python  
*Examples of using SigOpt to tune ML algorithms.* 2014 - current
  - Examples of using SigOpt to tune everything from sklearn to beating Vegas and beyond.
- **MOE: Metric Optimization Engine** ([github.com/Yelp/MOE](https://github.com/Yelp/MOE)) Python, C++, CUDA  
*A global, black box optimization engine for real world metric optimization* 2010 - 2015
  - Implemented throughout Yelp, optimizing ad metrics. 2nd most popular open source project.
  - Talk: [bit.ly/1p1YZA2](https://bit.ly/1p1YZA2), Slides: [slidesha.re/1z0r0Jy](https://slidesha.re/1z0r0Jy), Blog: [bit.ly/1x73xdr](https://bit.ly/1x73xdr)
  - Presented to executives, universities, conferences and companies around the country.
- **ALE: Assembly Likelihood Estimator** ([github.com/sc932/ALE](https://github.com/sc932/ALE)) C, Python  
*Probabilistic evaluation of genome assemblies* 2010 - 2013
  - Uses statistical function to score and rank genome assemblies, published in Bioinformatics