# **Daniel Berenberg**

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Permanent Address: 7 Abingdon Avenue Medford, NJ 08055

#### Education

University of Vermont Graduate College, Burlington, Vermont

2016 - 2018

M.S., Computer Science — 3.9/4.0

Selected Courses: Advanced courses in Artificial Intelligence (DL, ML, RL), Complex Systems, Data Science, Software Engineering, Algorithms for Complex Networks

 ${\bf University\ of\ Vermont\ CEMS}, \ {\bf Burlington}, \ {\bf Vermont}$ 

2013 - 2017

B.S., Mathematics - 3.41/4.0

⇒ Minor: Computer Science; 40+ credits in Russian

St. Petersburg University, St. Petersburg, Russia CIEE Study Abroad Program

Fall 2015

### Professional Experience

University of Vermont Mathematics, Burlington, VT

Current

Teaching Assistant - Data Science

⇒ Asked to TA course by instructor. Duties include holding regular office hours and teaching statistical, computational, and programmatic concepts to students in Python.

 ${\bf Vermont} \ \, {\bf Complex} \ \, {\bf Systems} \ \, {\bf Center} \, \, {\bf -Bagrow} \ \, {\bf Lab}, \, {\bf Burlington}, \, {\bf Vermont} \, \,$ 

Fall 2017 - Current

Research Assistant

- ⇒ Intersecting graph theory and statistics with NLP/NLU to research complex networks.
- $\Rightarrow$  Studying cutting edge computational linguistics research.
- $\Rightarrow$  Developing deep language processing models and implementing Python-based research tools using NLTK, Keras, Tensorflow, and Stanford CoreNLP.

**Vermont Artificial Intelligence Laboratory (VAiL)**, Burlington, Vermont *Machine Learning Engineer* 

 $Summer\ 2018$ 

- ⇒ Designed and constructed artificially intelligent computer vision system capable of predicting heart rate and respiratory rate from a noisy dataset of smart phone videos.
- ⇒ Implemented advanced video preprocessing and manipulation software using various Python libraries including OpenCV, PIL, numpy, and scipy.
- ⇒ Developed high performance 3D convolutional network using Keras and Tensorflow.

The Flatiron Institute - Computational Biology Lab, Manhattan, New York Research Intern

Summer 2017

 $\Rightarrow$  Utilized advanced protein fold simulation codebase (Rosetta) and molecular visualization software (PyMOL) to research physically stable conformations of empirically unobserved protein structures. Duties included implementing various python and bash scripts to interface with the C++ codebase.

 ${\bf University\ of\ Vermont\ Computer\ Science},\ {\bf Burlington},\ {\bf Vermont}$ 

Spring 2017

Teaching Assistant - Introduction to Java

#### **Publications**

Efficient Crowd Exploration of Large Networks: The Case of Causal Attribution

D. Berenberg, J. P. Bagrow, In Proc. ACM Hum-Comput. Interact. (CSCW '18) (2018)

⇒ Honorable mention for best paper

Neural language representations predict outcomes of scientific research

J.P. Bagrow, D. Berenberg, and J. Bongard, Preprint (2018)

### Skills

Development/Scripting: Python, C/C++, Java, Unix Shell/Bash, Ocaml, HTML/CSS

Frameworks/Tools: Keras, Tensorflow Scikit-Learn, SciPy & NumPy, git, LATEX, Wordpress

Technical: OOP & functional programming, Agile development, cluster computing

**Languages**: Russian  $\sim 2.1$  on Language Proficiency Index

#### Leadership & Other Achievements

UVM Computer Science Fair, 3<sup>rd</sup> place Dean's List Fall 2017