# **Niklas Smedemark-Margulies**

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#### Current

#### **Northeastern University**

Sept. 2018 – Present

PhD in Computer Science, Probabilistic Modeling Lab. GPA: 3.9 / 4.0

Boston, MA

- · Variational inference for parameter estimation in network model of COVID-19 spread.
- Multi-task autoencoders for few-shot representation learning.
- Training generative models as signal priors for inverse imaging problems.
- Course work in algorithms, machine learning, compilers, distributed systems, deep learning, and inverse imaging problems.

## **Education**

#### **Harvard Medical School**

Sept. 2014 - May 2016

MMSc, Immunology. GPA: 3.9 / 4.0.

• Thesis: Next-Generation Roadmap for Patient-Centered Genomics

# **Amherst College**

Sep 2009 - May 2013

BA, Neuroscience. GPA: 3.7 / 4.0.

· Senior Research: Tools, Methods, and Applications for Optophysiology in Neuroscience

# Work Experience

# **Software Engineer**

March 2017 - April 2018

Genuity Science (formerly WuXi NextCODE)

Cambridge, MA

- Wrote batch and realtime analyses for DNA variant scoring in over 100,000 whole genomes. (GORpipe) (bash)
- Designed and implemented ingestion pipelines for germline and somatic exomes. (Docker) (Nextflow) (Bash)
- Created documentation and conducted training for internal developers and external clients.

#### **Software Engineer**

May 2016 – March 2017

Claritas Genomics

Cambridge, MA

- Developed pipelines for exome data and validated results using NIST gold standard datasets. (Java) (GORpipe) (AWS)
- Performed processing and support for production samples. (Python)

#### **Research Experience**

# **Research Intern**

June 2019 – August 2019

Q-State Biosciences

Cambridge, MA

- Increased SNR and achieved nearly 50-fold compression in detection of single-cell activity from fluorescence microscopy video data.
- Corrected for photobleach-induced exponential decay using spline detrending, followed by iterative rank-1 SVD for denoising, and non-negative matrix factorization for signal demixing. (MATLAB) (Python) (Docker)

## Research Associate, Data Coordinator

May 2015 - May 2016

Timothy Yu Lab, Boston Children's Hospital

Boston, MA

- Evaluated and curated potential disease-causing variants in clinical cohorts. (GORpipe) (Bash)
- Prototyped shared infrastructure for forming cohorts and variant interpretation. (AWS) (GORpipe) (Bash)

#### **Research Fellow**

May 2013 – May 2014

Adam Cohen Lab, Harvard University

Cambridge, MA

- Assisted development of voltage-sensitive fluorescent transmembrane protein for high-resolution measurement of activity in electrically active cells.
- Dissected and cultured mouse hippocampal and cortical neurons for functional analysis of protein candidates.

# **Teaching**

#### **Undergraduate Research Mentor**

Spring 2020

Probabilistic Modeling Lab

Northeastern University

• Supervised research on graph-based classifiers in the latent space of a deep neural network. (PyTorch)

**Teaching Assistant** 

Spring 2019

Algorithms

Northeastern University

• Helped create and grade homeworks, held office hours and review sessions. Python

## **Head Teaching Assistant**

Fall 2018

Database Design

Northeastern University

Created homeworks, held office hours, helped create exams, and helped organize other TAs. (SQL) (Python)

# **Projects**

DQN.

Summer 2020

• Reimplemented Deep Q-Networks for reinforcement learning in Atari. PyTorch Gym Jupyter Tensorboard

Raft.

Fall 2019

• Reimplemented Raft algorithm for distributed consensus in Golang. (Golang) (net/rpc) (Docker)

#### **Neural Topic Models for Lyrics.**

Spring 2019

• Classified song genre using features produced by neural topic modeling. PyTorch Pandas Gensim (NLTK) (Docker)

## **Technical Skills**

Languages: Python MATLAB Julia Bash Go OCaml C SQL GORpipe

Libraries: PyTorch (NumPy) (Matplotlib) (Jupyter) (Tensorboard) (Scikit-Learn) (Unittest)

**Developer Tools:** Git Docker Travis CI AWS Vim+ALE Linux

# Volunteering

Mentor, Big Brothers Big Sisters of MA

January 2016 - October 2018

#### **Selected Publications**

Kocanaogullari, A., **Smedemark-Margulies, Niklas**, Akcakaya, M., & Erdogmus, D. (Submitted). Geometric analysis of uncertainty sampling for dense neural network layer.

- Park<sup>1</sup>, J. Y., **Smedemark-Margulies**<sup>1</sup>, **Niklas**, Daniels, M., Yu, R., van de Meent, J.-W., & Hand, P. (2020). Generator surgery for compressed sensing.
- Smedemark-Margulies<sup>1</sup>, Niklas, Langton<sup>1</sup>, P., & Nguyen, H. L. (2020). Fair and useful cohort selection [https://arxiv.org/abs/2009.02207].
- Smedemark-Margulies, Niklas, Brownstein, C. A., Vargas, S., Tembulkar, S. K., Towne, M. C., Shi, J., Gonzalez-Cuevas, E., Liu, K. X., Bilguvar, K., Kleiman, R. J., et al. (2016). A novel *de novo* mutation in atp1a3 and childhood-onset schizophrenia [http://molecularcasestudies.cshlp.org/content/2/5/a001008.short]. *Molecular Case Studies*.
- Hochbaum, D. R., Zhao, Y., Farhi, S. L., Klapoetke, N., Werley, C. A., Kapoor, V., Zou, P., Kralj, J. M., Maclaurin, D., Smedemark-Margulies, Niklas, Saulnier, J. L., Boulting, G. L., Straub, C., Cho, Y. K., Melkonian, M., Wong, G. K.-S., Harrison, D. J., Murthy, V. N., Sabatiny, B. L., ... Cohen, A. E. (2014). All-optical electrophysiology in mammalian neurons using engineered microbial rhodopsins [https://www.nature.com/articles/nmeth.3000/]. *Nature methods*.
- Prilutsky, D., Palmer, N. P., **Smedemark-Margulies, Niklas**, Schlaeger, T. M., Margulies, D. M., & Kohane, I. S. (2014). Ipsc-derived neurons as a higher-throughput readout for autism: Promises and pitfalls [https://www.sciencedirect.com/science/article/abs/pii/S1471491413002062]. *Trends in molecular medicine*.
- **Smedemark-Margulies, Niklas**, & Trapani, J. G. (2013). Tools, methods, and applications for optophysiology in neuroscience [https://www.frontiersin.org/articles/10.3389/fnmol.2013.00018/full]. *Frontiers in molecular neuroscience*.