# Evan M. Cofer

# Computational Biology & Computer Science

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#### Education

September Ph.D. in Quantitative and Computational Biology, Princeton University, 2017 - Princeton, NJ. Present

2012 - 2017 Bachelor of Science in Computer Science, Trinity University, San Antonio, TX.

Computer Science GPA: 3.93/4.0 Trinity University Mach Research Fellow Trinity University Trustee's Scholarship

### Awards, Honors, and Scholarships

March 2017 Graduate Research Fellowship Program: Honorable Mention, National Science Foundation (NSF).

> "The program supports outstanding graduate students who are pursuing research-based master's and doctoral degrees in science, technology, engineering, and mathematics (STEM) or in STEM education. The NSF accords Honorable Mention to meritorious applicants who do not receive Fellowship awards. This is considered a significant national academic achievement."

February Trinity CS Department Outstanding Senior Research Award, Trinity University.

 $^{2017}$  "Not awarded every year, this honor is given to recognize that senior who has a particularly distinguished record of Computer Science research."

April 2016 Mach Research Fellowship, Trinity University.

Yearly, each academic department recommends a rising senior for the Mach, which recognizes outstanding undergraduate research achievement. From these nominees, the Faculty Research Committee selects five recipients. I was nominated by the computer science department, and subsequently chosen as a recipient.

April 2016 & No-Boundary Thinking Contest: First Place, Trinity University.

December "Entrants should propose a computational solution to an interdisciplinary problem. In addition 2015 to an essay, submissions should include a presentation at the department colloquium." I competed twice, winning both times. My first submission considered the use of authorship/topic networks for improving gene name disambiguation. For the second entry, I described a means of constructing lightweight ontologies to aid in the exploration of large datasets of leaked documents.

December **Dean's List**, *Trinity University*.

2015 – May Achieved a grade point average of 3.65 or higher while taking 15 or more hours. 2017

August 2012 - **Trustee's Scholarship**, *Trinity University*.

May 2017 Scholarship for academic merit.

## Research Experience

September Troyanskaya Lab, Princeton University.

2017 - Researching deep learning methods for analyzing biological big data.

 $Present \quad http://function.princeton.edu/$ 

May 2017 - Kostic Lab, Joslin Diabetes Center and Harvard Medical School.

September

Conducted research in metagenomics and type 1 diabetes. Developed scalable and costefficient tools for bioinformatics in the cloud. Assisted with interviews for computational research staff.

http://www.kosticlab.org/

January 2016 Hibbs Lab, Trinity University.

June 2017

Quantified the rate at which somatic mutations accumulate in healthy, aging mammals. Developed a deep learning model to distinguish somatic mutations from noise in high volume DNA sequencing data (>13 trillion paired-end reads).

http://www.cs.trinity.edu/mhibbs/HibbsHome/Research.html

April 2015 - **Jiang Lab**, *Trinity University*.

July 2015

Modelled the popular vote of battleground states in the 2012 United States presidential election. Predicted outcomes both nationally and state-by-state, using bootstrapping to counter poll sparsity.

http://www.cs.trinity.edu/xjiang/

May 2013 - Glawe Lab, Trinity University.

July 2013

Analyzed the use and accuracy of nutating disc mechanical flow meters for recycling water condensation from air conditioning systems. In addition to research, managed all equipment and developed graphical user interfaces for lab instruments.

https://new.trinity.edu/faculty/diana-glawe

# Relevant Work Experience

January 2017 **Teaching Assistant**, *Trinity University*.

May 2017 Worked as a grader, tutor, and teaching assistant for CSCI-1323 Discrete Structures.

#### Software

2016 **Reddigest**, https://github.com/evancofer/reddigest-scalafied.

Collaborated with two other developers to create Reddigest, a web application based around Reddit. Reddigest filters out links on Reddit that a user has already viewed. Initially built with PHP/javascript/Apache HTTP, but rewritten in Scala/Scala.js/Play to improve maintainability, scalability, and overall quality.

2015 Polling Data Retrieval Program.

A Java program to fetch and parse XML polling data from the Huffington Post. Retrieved data is processed with MATLAB. Source code available upon request.

Many Miscellaneous Code and Projects, https://github.com/evancofer.

Source code available upon request.

## Relevant Coursework

Completed at Princeton University, (\* denotes courses in progress).

Theoretical Machine Learning\*, Numerical Algorithms for Scientific Computing\*, Method and Logic in Quantitative Biology, Introduction to Genomics and Computational Molecular Biology

#### Completed at Trinity University.

CS Thesis I & II, Artificial Intelligence, Compilers, Software Engineering, Principles of Functional Languages, Theoretical Computer Science, Database Systems, UNIX Power Tools, Web Application Development, Advanced Algorithms, Principles of Computer Design, CS Thesis Reading, Data Abstraction, Computational Game Theory and Multi-agent Systems, Discrete Data Structures, Principles of Computer Science I & II, Probability & Statistics for Scientists & Engineers, Differential Equations & Linear Algebra, Calculus I, II, & III, Directed Studies in Computer Science, Competitive Programming, Electronic Circuits & Lab, Physics I & II, Thermodynamics I, Mass and Energy Balance

#### **Proficiencies**

#### Languages and Technologies.

C++, Python, R, Scala (Akka, Play, Scala.js), TensorFlow, bash, git, Linux, C, Keras/Elephas, SQL, JavaScript, Haskell, HPC (Torque, Slurm), MongoDB, LATEX, CSS/HTML, MIPS Assembly

#### Academic and Professional Affiliations

February 2017 Member, Upsilon Pi Epsilon.

Present International honor society for computer scientists.
http://upe.acm.org/

May 2016 - Member, International Society for Computational Biology (ISCB).

Present Scholarly society for researchers in computational biology and bioinformatics. https://www.iscb.org

August 2015 - Member, Association for Computing Machinery (ACM).

Present Professional society for computer scientists. https://www.acm.org

## **Publications**

(Random ordering where not marked by  $^{\dagger}$ ) Ching T, Himmelstein DS, Beaulieu-Jones BK, Kalinin AA, Do BT, Way GP, Ferrero E, Agapow P, Xie W, Rosen GL, Lengerich BJ, Israeli J, Lanchantin J, Woloszynek S, Carpenter AE, Shrikumar A, Xu J, **Cofer EM**, Harris DJ, DeCaprio D, Qi Y, Kundaje A, Peng Y, Wiley LK, Segler MHS, Gitter A $^{\dagger}$ , and Greene CS $^{\dagger}$ . Opportunities and obstacles for deep learning in biology and medicine [in press]. *Journal of the Royal Society Interface*.

JM Luber, BT Tierney, **Cofer, EM**, CJ Patel, and AD Kostic. Aether: Leveraging linear programming for optimal cloud computing in genomics. *Bioinformatics*, December 2017. doi:10.1093/bioinformatics/btx787.

#### **Preprints**

JM Luber, BT Tierney, **Cofer, EM**, CJ Patel, and AD Kostic. Aether: Leveraging linear programming for optimal cloud computing in genomics. *bioRxiv*, July 2017. doi:10.1101/162883.

(Random ordering where not marked by †) Ching T, Himmelstein DS, Beaulieu-Jones BK, Kalinin AA, Do BT, Way GP, Ferrero E, Agapow P, Xie W, Rosen GL, Lengerich BJ, Israeli J, Lanchantin J, Woloszynek S, Carpenter AE, Shrikumar A, Xu J, **Cofer EM**, Harris DJ, DeCaprio D, Qi Y, Kundaje A, Peng Y, Wiley LK, Segler MHS,

Gitter  $A^{\dagger}$ , and Greene CS $^{\dagger}$ . Opportunities and obstacles for deep learning in biology and medicine. *bioRxiv*, May 2017. doi:10.1101/142760.

## Conference Presentations, Posters, and Talks

**Cofer EM**. Determining the age-associated somatic mutation rate in diverse mouse tissues [talk]. *The Computational Systems for Integrative Genomics (CSIG) Conference*, June 2017. Princeton University. Princeton, NJ.

**Cofer EM**. The rate of somatic mutations in aging mice [undergraduate thesis]. *Trinity University Computer Science Undergraduate Honors Thesis*, April 2017. Trinity University. San Antonio, TX.

**Cofer EM**. Accurately measuring the mammalian somatic mutation rate using deep learning [talk]. *Computer Science Colloquium*, March 2017. Trinity University. San Antonio, TX.

**Cofer EM** and Hibbs MA. Accurately measuring the mammalian somatic mutation rate using deep learning [platform talk & poster]. *Conference for Science at the Forefront of Basic and Translational Research*, October 2016. University of Texas at San Antonio. San Antonio, TX.

**Cofer EM**. Using deep learning to classify mutations [talk]. *Computer Science Colloquium*, September 2016. Trinity University. San Antonio, TX.

**Cofer EM**, Kennedy RM, and Hibbs MA. Accurately measuring the mammalian somatic mutation rate [poster]. *International Conference on Intelligent Systems for Molecular Biology*, July 2016. Orlando, FL.

**Cofer EM** and Hibbs MA. Using deep learning to classify mutations [talk]. *Trinity University Summer Undergraduate Research Conference*, July 2016. Trinity University. San Antonio, TX.

**Cofer EM\*** and Witecki I\*. Ontology construction: a means of improving data-driven journalism [talk]. *Computer Science Colloquium*, April 2016. Trinity University. San Antonio, TX. (\*Equal contribution)

**Cofer EM**. Gene name disambiguation: a novel approach [talk]. *Computer Science Colloquium*, December 2015. Trinity University. San Antonio, TX.

**Cofer EM**, Kwessi EA, Nguyen HV, Nishikawa KA, and Jiang AX. Modeling the 2012 presidential election's battleground states [talk]. *Trinity University Summer Undergraduate Research Conference*, July 2015. Trinity University. San Antonio, TX.

**Cofer EM\***, Ybarra T\*, and Glawe DD. Positive displacement meter performance [talk]. *Trinity University Summer Undergraduate Research Conference*, July 2013. Trinity University. San Antonio, TX. (\*Equal contribution)