JAMES (JAMIE) T. MORTON

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EDUCATION

Graduate University of Colorado, Boulder 2014 - Present PhD student in Computer Science

Integrative Quantitative Biology Program

Undergraduate Miami University

2010 - 2014 $\,$ Four B.S. Degrees with majors in

Computer Science (Cum Laude)

Engineering (Cum Laude) Mathematics and Statistics

Engineering Physics GPA: 3.74/4.0

Study Abroad Hong Kong University

Spring 2012

Hong Kong University of Science and Technology

HONORS

- NSF Graduate Fellow, 2015 2018 (Started date deferred from Fall 2014 as requested)
- Integrated Quantitative Biology Fellowship, University of Colorado Boulder, 2014 2016
- National Barry Goldwater Scholar, 2013
- Benjamin Harrison Scholar, Miami University, 2010-2014
- First place, Institute of Navigation (ION) Autonomous Snowplow Competition, 2014
- NSF REU, Cold Spring Harbor Laboratories, Summer 2012
- Provost Academic Achievement Award, Miami University, 2012
- Ohio Space Grant Scholar Award, NASA, 2012 2014
- Dean's List, Miami University, 2010-13
- R.L. Edwards Scholarship, Department of Physics, Miami University, 2011, 2013
- Mary Jeannette and Clifford Harvey Scholarship, Department of Mathetmatics, Miami U., 2013
- Mary Jean and Joseph R. Priest Scholarship, Department of Physics, Miami University, 2012
- President List, Miami University, 2010-11
- Nestle Scholar, Computer Sci. and Software Eng. Dept, Miami University, 2011
- Faculty Prize, Department of Mathematics, Miami University, 2011
- Joseph A. Culler Award, Department of Physics, Miami University, 2010,2011
- NSF Travel Grant, Coupling, Energetics, & Dynamics of Atmospheric Regions workshop, 2010
- Wright Scholar, Air Force Research Laboratory, Wright Patterson Air Force Base, 2009

EXPERIENCE

Research Experience

Rotation student

University of Colorado, Boulder, CO, Fall 2014

- Worked with Dr. Christopher Lowry
- Developed software tools to visualize mouse microbial data
- Co-author on manuscript under preparation.

Rotation student

University of Colorado, Boulder, CO, Fall 2014

- Worked with Dr. Noah Fierer and Albert Barberan
- Developed a Random Forests classifier to classify homes with pets based on dust samples
- Co-author on manuscript under preparation.

Research Assistant

- Worked with Dr. Iddo Friedberg
- Developed software tool to identify bacteriocin associated gene clusters
- First author on conference oral presentation accepted at Rocky Mountains Bioinformatics Conference.
- First author on journal manuscript ready for submission.

Data Scientist Intern

Johns Hopkins University, MD, Summer 2013

- Worked with Dr. Benjamin Langmead to develop scalable RNAseq Analysis software
- Developed spliced alignment algorithm using the Hadoop Framework.
- Co-author on manuscript under preparation.

Undergraduate Research Program

Cold Spring Harbor Laboratories, NY, Summer 2012

- Worked with Dr. Thomas Gingeras and Dr. Alex Dobin
- Developed software that maps reads between the reference and personal genome
- Studied Allele Specific Expression in a personal genome
- Results incorporated into PIs NSF proposal

Research Assistant

Miami University, OH, Spring 2011 - Fall 2011

- Worked with Dr. John Karro and Dr. Chun Liang
- Designed Hidden Markov Model software to identify poly(A) tails in RNAseq data
- Designed Profile Hidden Markov Model software to identify adapter sequences in RNAseq data
- Contributed HMMER parser to Biopython
- Released a software package, presented poster at Genome Informatics 2011

Research Assistant

Miami University, OH, Summer 2010

- Worked with Dr. Qihou Zhou on processing incoherent scattering radar data
- Developed signal processing algorithms to extract atmospheric parameters from this data
- Awarded NSF travel grant to present a poster paper at an international workshop

Engineering Aide

Wright Patterson Air Force Base, OH, Summer 2010

- Designed and implemented a time difference of arrival localization algorithm
- Programmed USRP using GNU radio for signal transmission and receiving

Wright Scholar

Wright Patterson Air Force Base, OH, Summer 2009

• Studied cognitive radio, radar, and GPS concepts and techniques

Teaching Experience

Teaching Assistant

Miami University, OH, Spring 2011

- Assisted Professor Mostafa Modirrousta in teaching of two sections of Intro to Engineering labs
- Graded lab reports for a class of 32 students

Teaching Assistant

Miami University, OH, Spring 2008

• Assisted Professor Felice Marcus to teach a class of Chinese engineers English

National Competitions

Autonomous Snowplow Competition

St. Paul MN, January, 2014

• One of three team members, First Place Award, Best Report Award, Best Presentation Award Intelligent Ground Vehicle Competition Rochester, MI , Summer 2013

• One of four team members, 5th place in Design

SKILLS

Foreign Language Skills

• Chinese – Working Proficency in Mandarin and written Chinese

Technical Skills

- \bullet Python \bullet C/C++ \bullet Java \bullet Javascript \bullet LATEX \bullet ROS \bullet Hadoop
- ullet Matlab ullet R ullet Unix ullet SQL ullet OpenCL ullet CUDA ullet git

PUBLICATIONS

- Morton, J., Abrudan, P., Figuegoaa, N., Liang, C., Karro, J. SCOPE++: Sequence Classification Of homoPolymer Emissions, *Genomics*. 104.3 (2014) 157–162
- Morton, J., Freed, S. Lee, S. Friedberg, I Prediction of Bacteriocin Associated Operons, Near ready for submission
- Nellore A., Morton, J., Langmead B. Rail-RNA: A scalable spliced read aligner, Near ready for submission
- Insights into Indoor Microbial Communities Barberan, A., Dunn, R. Fierer, N., Morton, J., In preparation

PRESENTATIONS

- Morton, J., Freed, S. Lee, S. Friedberg, I. Prediction of Bacteriocin Associated Operons Rocky Mountain Bioinformatics Conference, 2014
- Morton, J., Freed, S. Lee, S. Friedberg, I. A pipeline for Identifying Bacteriocin-Associated Gene Clusters. ISMB Boston, 2014
- Morton, J., Freed, S. Lee, S. Friedberg, I. Discovering the Next Antibiotic Ohio Space Grant Consortium, Cleveland OH, 2014
- Morton, J., P., Abrudan, J. Karro, C. Liang , Sequence classification of homopolymer emissions (SCOPE), Great Lakes Bioinformatics Conference, Pittsburgh, PA, 2013
- Morton, J., P., Abrudan, J. Karro, C. Liang, Sequence classification of homopolymer emissions (SCOPE), Ohio Space Grant Consortium, Cleveland OH, 2013
- Morton, J., P., Abrudan, <u>J. Karro</u>, C. Liang, Sequence classification of homopolymer emissions (SCOPE), IEEE 2nd International Conference on Computational Advances in Bio and Medical Sciences, ICCABS 2012, Las Vegas, NV, February 2012
- Morton, J., J. Karro, C. Liang, A novel approach for identifying poly(A) tails in raw cDNA sequence data using General Hidden Markov Models, Genome Informatics Cold Spring Harbor, NY, November 2011.

OPEN SOURCE CONTRIBUTIONS

- Sci-kit Bio (Contributor)
- Emperor (Contributor)
- Scipy (Contributor)
- Biopython (Contributor)
- SCOPE++: Sequence Classification Of homoPolymer Emissions (Developer)

ACTIVITIES

- International Society of Computational Biology Student member, Summer 2014-Present
- Sigma Pi Sigma, Tau Beta Pi, Eta Kappa Nu Spring 2014-Present
- National Society of Collegiate Scholars, Fall 2012 Spring 2013
- Association for Computing Machinery Student member, Fall 2011-Present
- Institute of Electrical and Electronics Engineers Student member, Fall 2011-Present
- IEEE Miami Student Chapter Treasurer, Fall 2011- Spring 2012