# Tom Hartvigsen

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# **EDUCATION**

# WORCESTER POLYTECHNIC INSTITUTE

PH.D. IN DATA SCIENCE
Expected May 2021 | Worcester, MA
Cum. GPA: 3.83 / 4.0
Advisor: Dr. Elke Rundensteiner

#### **SUNY GENESEO**

BA IN APPLIED MATHEMATICS Minor in BioMathematics May 2016 | Geneseo, NY Cum. GPA: 3.07 / 4.0

### LINKS

LinkedIn://in/thartvigsen Github://thartvigsen

# COURSEWORK

#### GRADUATE

Knowledge Discovery and Data mining Introduction to Data Science Statistical Learning Database Management Systems Big Data Management\* Business Intelligence\* Deep Learning\*\* \*to be completed Dec 2017 \*\*to be completed May 2018

#### **UNDERGRADUATE**

Modeling Biological Systems Combinatorics Differential Equations Probability and Statistics Linear Algebra Calculus I-III

# SKILLS

#### **PROGRAMMING**

Python - Deep Learning: TensorFlow, Machine Learning: Scikit-Learn, Numpy
• R - Statistical Learning, Graph mining, ggplot2 • Shell • LATEX • SQL - PostgreSQL, SQLPLUS

# **AWARDS**

2016 GAANN Research Fellowship U.S. Department of Education

## **EXPERIENCE**

#### WPI | GAANN RESEARCH FELLOW

August 2016 - present | Worcester, MA

• Working with Dr. Elke Rundensteiner on Deep Learning for time series classification.

#### UNIVERSITY OF ARIZONA | RESEARCH EXPERIENCE FOR

#### **UNDERGRADUATES INTERN**

June 2015 - Aug 2015 | Tucson, AZ

- School of Natural Sciences and the Environment advised by Dr. Shirley Papuga.
- Used MATLAB to design Machine Learning algorithms to segment sequences of images of Creosote Bushes to record changes through time in regards to drought.
- Presented findings at the Undergraduate Research Opportunities Consortium (UROC) in August 2015.

## RESEARCH

#### DATA SCIENCE RESEARCH GROUP Aug 2016 - Pres. | Worcester, MA

- Currently focusing on **early time series classification** with application to infection prediction.
- Developing deep recurrent models for using TensorFlow, moving towards Deep Reinforcement Learning.
- Developed CREST, a python-driven machine learning tool for infection detection in hospitals using SVMs, Random Forests, and Logistic Regression.
- Used the publicly available MIMIC III Database, Python, and PostgreSQL.

# **BIOMATHEMATICS INNOVATION GROUP** Jan 2013 - May 2016 | Geneseo, NY

- Used R to build population networks driven by differential equation models. Used Python to collect lyrical data from song lyric websites. Used Python to scrape IMDB and build graphs relating movies.
- Built interdisciplinary projects, mentored younger students, and brought 6 projects to undergraduate research conferences.

# **PUBLICATIONS**

- Hartvigsen, T., Sen, C., Brownell, S., Teeple, E., Rundensteiner, E. (January, 2018). Early Prediction of MRSA Infections using Electronic Health Records . HealthInf 2018.
- Sen, C., Hartvigsen, T., Claypool, K., Rundensteiner, E. (2017, September).
   CREST Risk Prediction for Clostridium Difficile Infection Using Multimodal Data Mining. ECML/PKDD 2017.
- Teeple, E., Hartvigsen, T., Sen, C., Rundensteiner, E. (2018). Risk Stratification and Diagnostic Performance of a Machine Learning Algorithm for Clostridium Difficile Detection Using Electronic Health Records Data. Preparing for submission the the New England Journal of Medicine in December 2017.