# Finn Kuusisto

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## RESEARCH

My research is in machine learning and data mining, with a primary focus in biomedical applications. In particular, I work on employing clinical, genetic, and text data in predicting drug effects, treatment assignment, toxicity prediction, integration of expert knowledge, and precision medicine in general.

# TECHNICAL SKILLS

## **LANGUAGES AND TOOLS**

Python • Java • JS • C# • C++ • C ATEX• SciKit-Learn • TensorFlow Git • MySQL • Flask Framework

# **COMMUNITY WORK**

## **PEER REVIEWING**

NIPS • ICML • AAAI • AIStats • AMIA Informatics Summit • AMIA Annual Symposium • Machine Learning • KAIS • DAMI

#### **EDITORIAL WORK**

ACM - XRDS MAGAZINE | JAN 12 - DEC 15 Editor for Back, a then-now comparison of technology fitting the issue theme.

## **MENTORING**

Mentor for local startups and undergraduates at Gustavus and UW-Madison.

#### **BADGERLOOP**

Member of UW-Madison's team for the 2018 Hyperloop Pod Competition.

# LINKS

LinkedIn://finn-kuusisto Github://finnkuusisto

# **EDUCATION**

## UNIVERSITY OF WISCONSIN | MADISON, WI

PHD | COMPUTER SCIENCE | JAN 10 - AUG 15 MS | COMPUTER SCIENCE | SEP 08 - DEC 09

#### **GUSTAVUS ADOLPHUS COLLEGE** | St. Peter, MN

BA | COMPUTER SCIENCE, CLASSICS | AUG 03 - MAY 07

Magna Cum Laude • Phi Beta Kappa • Eta Sigma Phi • Cross Country • Track & Field

# **EXPERIENCE**

## MORGRIDGE INSTITUTE FOR RESEARCH | MADISON, WI

POSTDOCTORAL FELLOW | JUL 17 - PRESENT

POSTDOCTORAL RESEARCH ASSOCIATE | SEP 15 - JUN 17

Research with the Regenerative Biology lab building predictive models of toxicity from gene expression, and doing text mining for cell reprogramming and drug repurposing. Includes mentoring graduate and undergraduate research assistants.

## UNIVERSITY OF WISCONSIN | MADISON, WI

RESEARCH ASSISTANT | MAY 11 - AUG 15

Research with Jude Shavlik, David Page, and Elizabeth Burnside in uplift modeling, breast cancer prediction, adverse drug event prediction, and precision medicine.

## LECTURER | JUN 10 - AUG 10

Lecturer for 8-week summer session of the intro CS course. Included development of course syllabus, preparing/grading assignments/exams, and supervising a TA.

#### TEACHING ASSISTANT | SEP 08 - MAY 11

Four semesters as an instructor, and two as a lab TA for the intro CS course.

## CONSULTING | MADISON, WI

## DATA SCIENCE CONSULTANT | NOV 14 - AUG 15

Consulting on data collection and processing, design and implementation of predictive modeling pipelines, and model evaluation and selection.

#### **REMUGIO** | MADISON, WI

CO-FOUNDER | SEP 14 - APR 18

Second-by-second feedback from viewers on video content. gBeta Accelerator Summer 2016 • Madworks Accelerator Fall 2016

# **OPEN SYSTEMS INTERNATIONAL | PLYMOUTH, MN**

SOFTWARE ENGINEER | AUG 07 - AUG 08

# SELECTED PUBLICATIONS

- [1] F. Kuusisto, D. Page, and R. Stewart. Word embedding mining for SARS-CoV-2 and COVID-19 drug repurposing. F1000Research [Awaiting Review], 2020.
- [2] F. Kuusisto, V. Santos Costa, Z. Hou, J. Thomson, D. Page, and R. Stewart. Machine learning to predict developmental neurotoxicity with high-throughput data from 2d bio-engineered tissues. In *IEEE International Conference on Machine Learning Applications*, 2019.
- [3] A. Movaghar, D. Page, M. Brilliant, J. Greenberg, J. Hong, L. S. DaWalt, K. Saha, F. Kuusisto, R. Stewart, E. Berry-Kravis, and M. R. Mailick. Data-driven phenotype discovery of fmr1 premutation carriers in a population-based sample. *Science Advances*, 2019.
- [4] R. Kleiman, F. Kuusisto, I. Ross, R. Stewart, D. Page, and J. Weiss. Machine learning assisted discovery of novel predictive lab tests using electronic health record data. In AMIA Informatics Summit, 2019.
- [5] F. Kuusisto, J. Steill, Z. Kuang, J. Thomson, D. Page, and R. Stewart. A simple text mining approach for ranking pairwise associations in biomedical applications. In AMIA Joint Summits on Translational Science, 2017.
- [6] F. Kuusisto, I. Dutra, M. Elezaby, E. Mendonca, J. Shavlik, and E. Burnside. Leveraging expert knowledge to improve machine-learned decision support systems. In AMIA Joint Summits on Translational Science, 2015.
- [7] F. Kuusisto, V. Santos Costa, H. Nassif, E. Burnside, D. Page, and J. Shavlik. Support vector machines for differential prediction. In *European Conference on Machine Learning (ECML-PKDD)*, 2014.