# Nathaniel Budijono

Resumé

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#### Education

2018–2022 Bachelor of Science, University of Minnesota, Computer science and Mathematics.

Focuses: machine learning, computational biology, computational chemistry.

 Reviewer, Minnesota Undergraduate Research Academic Journal; Education Director, App Developers Club; Lab Director, Society of Asian Scientists and Engineers; Mentor, Honors program.

### Industry Experience

Sep 2021 - Intern, Loyal Cellular Longevity, Computational biology team.

May 2021 - Al Research Intern, Smart Information Flow Technologies (SIFT).

Aug 2021 • Developed decoding scheme for BERT-based language model to improve joint entity-relation classification. Huggingface, PyTorch.

• Statistically analyzed sentiment and word embeddings of online groups of interest. spaCy.

May 2019 - Software Engineer Intern, Rally Health, Data engineering team.

Aug 2019 • Created a utility for data pipeline to gauge data throughput, monitor job duration, and track which rows failed which data validation. *Scala, Datadog API, Databricks*.

• Created a comprehensive and responsive web dashboard with pipeline health and performance metrics for data pipeline. *Airflow, Flask, SQLAlchemy, nvd3.js.* 

• Created an extract-transform-load (ETL) job to process ledger data from incentives program and performed end-to-end testing. *Apache Spark, SQL*.

#### **Publications**

- Ludus: An Optimization Framework to Balance Auto Battler Cards. Nathaniel Budijono, Phoebe Goldman, Jack Maloney, Joseph Mueller, Phillip Walker, Jack Ladwig, and Richard Freedman. AAAI Symposium on Educational Advances in Artificial Intelligence, 2021.
- Linguistic Factors in Incel Messaging. Sonja Schmer-Galunder, Ruta Wheelock, Joan Zheng, Claire Yang, Ian Magnusson, Jeremy Gottlieb, Scott Friedman, and Nathaniel Budijono. Special Issue for the Workshop on Online Abuse and Harms, 2021.

## Research Experience

May 2020 - Research Assistant, University of Minnesota, Department of Computer Science.

Aug 2020 Advisor: Prof. Chad Myers

- Created a pipeline for visualizing genetic interaction networks from raw combinatorial CRISPR screen data. *Networkx, Scipy.*
- Investigated machine learning algorithms for predicting gene function from genetic interaction networks. *PyTorch, Scikit-learn.*

#### Awards

Oct 2021 Winner (1/20, \$500), Acceleration Consortium Hackathon

Jul 2021 David E. Shaw Research Science and Engineering Undergraduate fellowship

Sep 2021 Honorable mention (top 5+/1300), 3blue1brown Summer of Math Exposition

Oct 2020 Finalist (top 10/n, \$1000), Wells Fargo Campus Analytics Challenge

Mar 2020 Best ML, AI, and Social Good Hacks, Hacktech (Caltech hackathon)