

#### **PERSONAL INFO**

- San Diego, California, USA
- kyetokikuchi@gmail.com
- in linkedin.com/in/kaitokikuchi/

## TECHNICAL SKILLS

- Python: Proficient in the data science stack including Pandas, NumPy, SciPy, Scikit-learn, Matplotlib, Streamlit
- ML/AI: Experienced in Pytorch, MLOps with Valohai, CNNs, clustering,
- Image Analysis: ImageJ/FIJI, OpenCV, QuPath
- Biological data: Confocal & fluorescence 2D/3D microscopy, histopathology whole slide images, protein-protein interaction networks, metabolic time-series data, Neuropixel activity readings, DNA/RNA sequences
- Other: Linux/UNIX, Google Cloud Platform (GCP), Clojure, Julia, R, **MATLAB**

#### VOLUNTEERING

• UJA, Inc: Advocation group for overseas Japanese researchers. Currently serving as the the translational research working group chair. Organized monthly online seminars, participated in fundraising, and chaired panel discussions.

### LANGUAGES

• English: Native fluency • Japanese: Native fluency

• German: Elementary

#### **INTERESTS**

Pottery/Cooking/Biking/Baking/Kintsugi

# Kaito Kikuchi Computational Biology & Al Scientist

## WORK EXPERIENCES

Manager, **Computational Biology** Reveal Biosciences, USA Aug 2023 - Present

- Leading the computational biology/image analysis team with 5 direct reports.
- Developed automated analysis pipelines and train machine learning models for digital pathology.
- · Built an internal data QC webapp in Python for outlier detection and data visualization.

**Image Analysis Scientist** Sep 2022 - Aug 2023

- Conducted comprehensive data analysis for product
- Acting as the technical expert on Al pipeline development within the company.
- Serving as the product owner of flagship WSI viewer software, imageDX.
- **Graduate Research Assistant** University of California San Diego, USA

Sep 2016 - Jun 2022

- Investigated how bacterial spores use electrochemical charges to return to life by combining molecular genetics, single-cell imaging, microfluidics, and data
- Designed experimental methods for imaging spore germination within a microfluidics device.
- · Analyzed image data by stabilize image drift, tracking spores, and extracting image features.
- · Performed data analysis in Python, including data parsing, time series analysis, statistical validations, and publication-ready data visualizations.
- **Research Assistant** Systems Biology Institute, Japan

Apr 2013 - Aug 2016

- Participated in bioinformatics projects aiming to locate drug-targets from protein-protein interaction (PPi) networks.
- Developed a novel network mining method by sequentially applying clustering algorithms.
- Applied clustering technique to MRSA, refining the module analysis step to use an ensemble voting method with 10 classification algorithms to predict drug targets.

## **EDUCATION**

University of California San Diego

Sep 2016 - Jul 2022

- University of Tokyo Apr 2014 - Mar 2016
- International Christian University Apr 2010 - Mar 2014

**Ph.D. Biology** with specialization in quantitative biology

M.A. Biophysics with a focus in singlecell live imaging

**B.A. Biology** with a focus in microbial genetics and DNA repair

## **PUBLICATIONS**

Kikuchi K, Galera-Laporta L, Weatherwax C, Lam J, Moon E, Theodorakis E, Garcia-Ojalvo J, Süel GM. Electrochemical potential enables dormant spores to integrate environmental signals. Science. Oct 2022.

Zhai X, Larkin JW, Kikuchi K, Redford SE, Roy U, Süel GM, Mugler A. Statistics of correlated percolation in a bacterial community. PLoS Comp Bio. Dec 2019.

Larkin JW, Zhai X, Kikuchi K, Redford SE, Prindle A, Liu J, Greenfield S, Walczak AM, Garcia-Ojalvo J, Mugler A, Süel GM. Signal percolation within a bacterial community. Cell Systems. Aug 2018.

Hill S, et al. [HPN-DREAM Consortium, including Kikuchi K]. Inferring causal molecular networks: empirical assessment through a community-based effort. Nature Methods. Apr 2016.

Hase T, **Kikuchi K**, Ghosh S, Kitano H, Tanaka H. *Identification of drug-target modules* in the human protein-protein interaction network. Artificial Life and Robotics. Dec 2014.