KASIA KEDZIERSKA

I am a Computational Biologist and Machine Learning Researcher dedicated to advancing methods development and application within biology and healthcare.

Currently, I lead an R&D project focused on biological Foundation Models (FMs). During my internship with the bioML team at Microsoft Research New England, I evaluated FMs for single-cell biology applications. Previously, @ Novo Nordisk Research Centre in Oxford, I utilized NLP techniques and knowledge graphs to enhance biomedical research. Additionally, with the Turing Data Study Group, I developed a framework using Transfer Learning and the YOLO model to identify sea pens from ocean floor video footage.

With my extensive expertise in Computational Biology, coupled with a robust background in Data Science and Machine Learning, I am enthusiastic about driving innovations at the intersection of these disciplines.

\square SFI ECTED WORK EXPERIENCE

present 2024

Deep Learning Researcher @ DeepLife, Remote

Leading an R&D project on Foundation Models in biology, including mentoring and supervising a Master's student.

2023

Intern @ Microsoft Research New England, Cambridge, Massachusetts, USA During the summer I investigated the potential of the Foundation Models in the space of

single cell biology. I was mentored by Alex Lu, Ava Amini, and Lorin Crawford.

2021

Intern @ Novo Nordisk Research Centre Oxford, Oxford, United Kingdom I worked with NLP and knowledge graphs to screen biomedical articles to identify and prioritise therapeutic targets. To increase the impact of the analysis and increase reach I built and deployed an **interactive dashboard** (using **R Shiny**) to allow colleagues within the company to investigate and visualise the results directly in real-time.

2024 2018 DPhil Researcher @ Wellcome Centre for Human Genetics, Big Data Institute, University of Oxford, UK

In my PhD project I looked at how **chromatin organization** influences **disease initiation** and progression in uterine cancer using multimodal data. I was also working on building and refining ML models of **cancer evolution**, specifically identyfing evolutionary trajectories in the cancer of the uterus.

EDUCATION

2024 2018 DPhil in Genomic Medicine and Statistics @ Nuffield Department of Medicine, **Brasenose College**

PhD fully funded by the Wellcome Trust Four-year PhD Studentships in Science

2018 2015 M. Sc. Eng., Biotechnology @ Warsaw University of Technology

Master thesis Analysis of the mutational burden across gene sets in cancer awarded the title of The Best Master Thesis in Bioinformatics defended in 2018.

SELECTED AWARDS AND HONORS

2023

JXTX + CSHL 2023 Biology of Genomes Scholarship @ JXTX Foundation, **Cold Spring Harbor Laboratory**

Awarded to outstanding graduate students in genomics and data sciences.

2022

Graduate Prize in the 'Outstanding work outside degree' category @ Nuffield Department of Medicine, University of Oxford

Each year Nuffield Department of Medicine, based on nominations, awards selected PhD students based on their performance within and outside of their degree.

2023 2021

2019

Senior Hulme Scholarship @ Brasenose College, University of Oxford

Senior Hulme Scholarship is awarded by Brasenose College, University of Oxford to DPhil students whose academic performance is deemed to be exceptional.

SUMMER SCHOOLS & HACKATHONS

Sea pen identification from video footage challenge @ Turing Data Study 2022 **Group @ The Alan Turing Institute, London, UK**

Machine Learning Summer School @ Imperial College London, University College London, London, UK

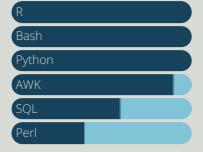
View this Resume online

** kasia.codes/resume/

CONTACT

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- **y** kzkedzierska
- github.com/kzkedzierska
- **@** kasia.codes
- **in** kzkedzierska

CODING SKILLS



Python:

PyTorch, keras/Tensorflow, scikitscverse/scanpy, Pandas, learn, NumPy, Seaborn, matplotlib and more

R:

tidyverse, data.table, Shiny, plotly, tidymodels, caret and others

High Performance / Cloud Computing:

SLURM, SGE, Google Cloud, Scaleway, Microsoft Azure, Azure Blob Storage, AWS Cloud Storage and more

LANGUAGES



CV source code. Made with pagedown, based on nstrayer/cv.

■ SELECTED PUBLICATIONS

Full list of publication is available through my Google Scholar profile scholar.google.com/citations?user=Yv6poTwAAAAJ.

• Assessing the limits of zero-shot foundation models in single-cell biology @ bioRxiv

K. Z. Kedzierska, L. Crawford, A. P. Amini, A. X. Lu + This work was featured in The New York Times +

Data Study Group Final Report: CEFAS - Automated identification of sea pens using OpenCV and machine learning @ Zenodo

In alphabetical order: M. Asthana, R. Blackwell, S. Davis, A. Downie, J. Forsyth, **K. Kedzierska**, R. Mestre, Z. Reza, J. Ribeiro, P. Palola, Y. Said

Functional analysis reveals driver cooperativity and novel mechanisms in endometrial carcinogenesis @ EMBO Molecular Medicine

M. Brown, A. Leon, K. Kedzierska, C. Moore, H. L. Belnoue-Davis, S. Flach, J. P. Lydon, F. J. DeMayo, A. Lewis, T. Bosse, I. Tomlinson, D. N. Church

Prognostic integrated image-based immune and molecular profiling in early-stage Endometrial Cancer @ Cancer Immunology Research

N. Horeweg, M. de Bruyn, R. A. Nout, E. Stelloo, **K. Kedzierska**, A. León-Castillo, A. Plat, K. D. Mertz, M. Osse, I. M. Jürgenliemk-Schulz, L. C.H.W. Lutgens, J. J. Jobsen, E. M. van der Steen-Banasik, V. T. Smit, C. L. Creutzberg, T. Bosse, H. W. Nijman, V. H. Koelzer and D. N. Church

Dynamics of cardiomyocyte transcriptome and chromatin landscape demarcates key events of heart development @ Genome Research

M. Pawlak, **K. Z. Kedzierska**, M. Migdal, K. A. Nahia, J. A. Ramilowski, L. Bugajski, K. Hashimoto, A. Marconi, K. Piwocka, P. Carninci and C. L. Winata

SONICS: PCR stutter noise correction in genome-scale microsatellites @ Bioinformatics K. Z. Kedzierska, L. Gerber, D. Cagnazzi, M. Krützen, A. Ratan, L. Kistler

SELECTED CONFERENCE PRESENTATIONS

- Systematic characterisation of chromatin modifiers in endometrial cancer, Poster @ European Association for Cancer Research 2022 Congress, Seville, Spain
- Analysis of the mutational burden across gene sets in cancer, Invited talk @ Polish Bioinformatics Society Symposium, Cracow, Poland

♣ SELECTED TEACHING EXPERIENCE

- Data visualization in bioinformatics hackathon mentor @ Online hackathon NGSprint, Discord

 I led the hackathon in data visualisation with emphasis on computational biology. Teaching materials are available at github.com/kzkedzierska/NGSprint_data_viz.
- Online tutorials: Python for Data Science and Introduction to Python @ NGSeminars, YouTube

 I led two Python tutorials: Introduction to Python kasia.codes/talk/intro_to_python/ and Python for Data Science kasia.codes/talk/py4ds/.
- Introduction to R @ Wellcome Centre for Human Genetics, Oxford, UK
 8 week course in Introduction to R, Data Manipulation, Data Visualisation and RNA-seq data analysis.
- Introduction to Managing Code with Git @ Wellcome Centre for Human Genetics, Oxford, UK
 I led a 2-hour introduction to working with Git. Materials, including slides and exercises are available at kasia.codes/talk/into_to_git/.

€ SELECTED GRANTS

2022 Visegrad Grant to organise NGSchool2022 @ Visegrad Fund

32,190 EUR awarded towards organising affordable training and conference focusing on ML application in Computational Biology. During this project I managed an international team of volunteers and led the organisation of summer school, conference, online seminars and hackathon.

NON-PROFIT WORK

2020

2018

President @ NGSchool Society

The goal of the Society is to promote and support science.

The goal of the Society is to promote and support science, with emphasis on computational biology.