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, at Novo Nordisk Research Centre in Oxford, I used 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.

RESEARCH 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

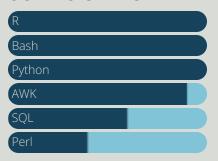
In my PhD project I looked at how **chromatin organization** influences **disease initiation and progression** in uterine cancer using multimodal data. I was also working on building and refining ML models of **cancer evolution**, specifically identyfing evolutionary trajectories in the cancer of the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

View this CV online kasia.codes/cv/

CONTACT

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

CODING SKILLS



Pvthon:

PyTorch, keras/Tensorflow, scverse/scanpy, scikit-learn, NumPy, Pandas, Seaborn, matplotlib and more

R:

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

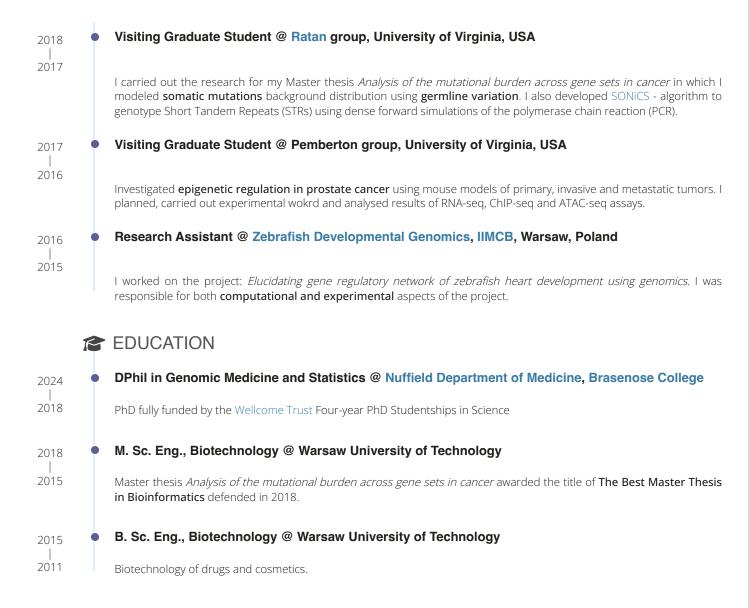
High Performance / Cloud Computing: SLURM, Sun Grid Engine (SGE), Microsoft Azure, Azure Blob Storage, AWS Cloud Storage and more

LANGUAGES



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

Last updated on 2024-10-07.



PUBLICATIONS

2023

Full list of publication is available through my Google Scholar profile scholar.google.com/citations?user=Yv6poTwAAAA|.

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 2023 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 Discordant prognosis of mismatch repair deficiency in colorectal and endometrial cancer reflects 2022 variation in antitumour immune response and immune escape @ The Journal of pathology M. A. Glaire, N. Al. Ryan, M. E. Ijsselsteijn, K. Kedzierska, S. Obolenski, R. Ali, E. J. Crosbie, T. Bosse, N. F. D. Miranda, D. N. Church Multi-omics analyses of early liver injury reveals cell-type-specific transcriptional and epigenomic 2021 shift @ BMC Genomics M. Migdal, E. Tralle, K. A. Nahia, L. Bugajski, K. Z. Kedzierska, F. Garbicz, K. Piwocka, C. L. Winata, M. Pawlak The MLH1 polymorphism rs1800734 and risk of endometrial cancer with microsatellite instability @ 2020 **Clinical Epigenetics** H. Russell, K. Kedzierska, D. D. Buchanan, R. Thomas, E. Tham, M. Mints, A. Keränen, G. G. Giles, M. C. Southey, R. L. Milne, I. Tomlinson, D. Church, A. B. Spurdle, T. A. O'Mara and A. Lewis Prognostic integrated image-based immune and molecular profiling in early-stage Endometrial 2020 **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 2019 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 Genomic analysis of DNA repair genes and androgen signaling in prostate cancer @ BMC Cancer 2018 K. Jividen, K. Z. Kedzierska, C.-S. Yang, K. Szlachta, A. Ratan and B. M. Paschal SONICS: PCR stutter noise correction in genome-scale microsatellites @ Bioinformatics 2018

POSTERS, AND TALKS

2024

K. Z. Kedzierska, L. Gerber, D. Cagnazzi, M. Krützen, A. Ratan, L. Kistler

Invited attendee @ Chan Zuckerberg Initiative AI in Single-Cell Biology Workshop, Redwood City, CA, USA

Chromatin modfiers in endometrial cancer, Poster @ Biology of Genomes 2023, Cold Spring 2023 Harbor, NY, USA Systematic characterisation of chromatin modifiers in endometrial cancer, Poster @ European 2022 Association for Cancer Research 2022 Congress, Seville, Spain Analysis of the mutational burden across gene sets in cancer, Invited talk @ Polish Bioinformatics 2019 Society Symposium, Cracow, Poland Differential mutation analysis across gene sets in cancers, Poster @ Biology of Genomes 2018, 2018 Cold Spring Harbor, NY, USA Epigenetic regulation of prostate cancer, Talk @ Visiting Graduate Traineeship Program Grantees 2017 Symposium, Charlottesville, VA, USA AWARDS JXTX + CSHL 2023 Biology of Genomes Scholarship @ JXTX Foundation, Cold Spring Harbor 2023 Laboratory Awarded to outstanding graduate students in genomics and data sciences. Graduate Prize in the 'Outstanding work outside degree' category @ Nuffield Department of 2022 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. Senior Hulme Scholarship @ Brasenose College, University of Oxford 2023 2021 Senior Hulme Scholarship is awarded by Brasenose College, University of Oxford to DPhil students whose academic performance is deemed to be exceptional. **Best Master Thesis in Bioinformatics** @ Polish Bioinformatics Society 2019 2018 Analysis of the mutational burden across gene sets in cancer - Best Master Thesis defended in Bioinformatics in 2018 in Poland Visiting Graduate Traineeship Program, now known as BioLAB @ Fulbright Poland 2017 2016 The Visiting Research Graduate Traineeship Program offered 12-month research traineeships for outstanding, qualified students from the life sciences at selected institutions in the United States. Grasz o Staz @ PwC Poland 2015 "Grasz o Staz" competition was a national, prestigious and highly competitive (1:25 success rate) scholarship program in Poland organized by PwC.

₽ TEACHING EXPERIENCE

2022

2021

2020

2019

2019

2019

2017

2022

2019

 Unsupervised learning @ NGSchool2022: Machine Learning in Computational Biology, Jablonna, Poland

I co-led, with Kaspar Märtens, lecture and tutorial sessions on unsupervised learning and its use cases in computational biology. All materials are availble at github.com/kzkedzierska/ngs22_unsupervised.

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

I led the hackathon in data viusalisation 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/.

 Unsupervised learning, Introduction to Python @ #NGSchool2019: Machine Learning for Biomedicine, Ostróda, Poland

Tutor for the Introduction to Python (3 h workshop) and for the Unsupervised learning (1,5 h lecture).

• 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/.

ATAC-seq workshop @ #NGSchool2017: Single-cell Sequencing, Jachranka, Poland

Invited speaker

ATTENDED WORKSHOPS, SUMMER SCHOOLS

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

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

€ GRANTS

2020

2019

2023 | 2018

2022

2018

2016

2015

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.

Visegrad Grant to organise NGSchool2019 @ Visegrad Fund

23,500 EUR awarded towards organising #NGSchool2019 allowed to keep the cost of attenting the school to the minimum and record the lectures for broader access.

Scientific grant as part of Wellcome Trust funded DPhil @ Wellcome Trust

30,000 GBP towards research expenses for the PhD project which allowed me to design and led pilot experiments, managing the grant for successful execution of research objectives.

NON PROFIT WORK

President @ NGSchool Society

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

President @ Warsaw Society of Biotechnology

Symbioza's aim is to integrate biotechnology community in Poland starting from the youngest generations, and promotion and popularization of biotechnology.