KASIA KEDZIERSKA

For my PhD @ University of Oxford, I studied cancer of the uterus and chromatin organisation in disease progression. Over the summer, I joined the bioML team at the Microsoft Research New England as an intern where I worked on Foundation Models in single-cell biology.

Previously, I worked with **NLP** methods and knowledge graphs during my internship @ Novo Nordisk Research Centre in Oxford. Using **Transfer Learning** and **YOLO model** I built a framework to identify sea pens from a video footage of an ocean floor @ Turing Data Study Group.

With extensive domain knowledge in **Computational Biology** complemented by my experience in **Data Science** & **Machine Learning** I am eager to drive advancements at the crossroads of these disciplines.

RESEARCH EXPERIENCE

 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.

present | 2018

2023

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

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

2018 | 2017 Visiting Graduate Student @ Ratan group, University of Virginia, USA

I carried out the research for my Master thesis *Analysis of the mutational burden across gene sets in cancer* in which I modeled **somatic mutations** background distribution using **germline variation**. I also developed SONiCS - algorithm to genotype Short Tandem Repeats (STRs) using dense forward simulations of the polymerase chain reaction (PCR).

2017 | 2016 Visiting Graduate Student @ Pemberton group, University of Virginia, USA

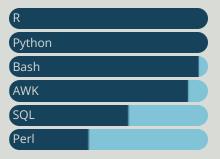
Investigated **epigenetic regulation in prostate cancer** using mouse models of primary, invasive and metastatic tumors. I planned, carried out experimental wokrd and analysed results of RNA-seq, ChIP-seq and ATAC-seq assays.

View this CV online kasia.codes/cv/

CONTACT

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

CODING SKILLS

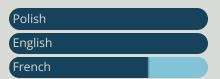


R: tidyverse, data.table, Shiny, plotly, tidymodels, caret and others

Python: NumPy, Pandas, SciPy, scikit-learn,

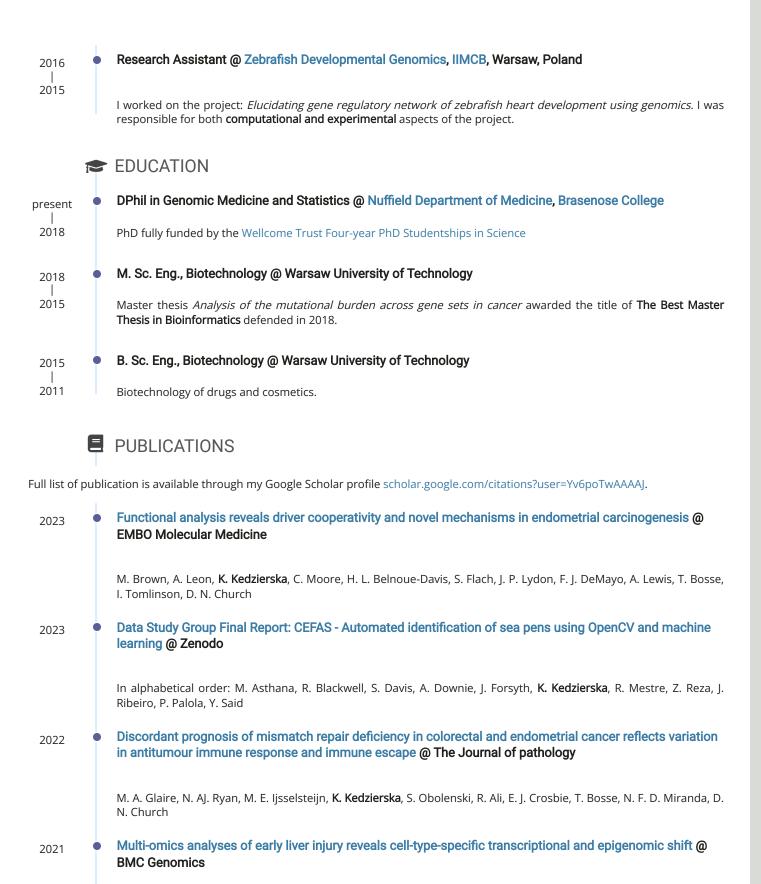
keras/Tensorflow, PyTorch, Seaborn, matplotlib and more

LANGUAGES



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

Last updated on 2023-09-13.



M. Migdal, E. Tralle, K. A. Nahia, L. Bugajski, K. Z. Kedzierska, F. Garbicz, K. Piwocka, C. L. Winata, M. Pawlak

2020	•	The <i>MLH1</i> polymorphism rs1800734 and risk of endometrial cancer with microsatellite instability @ 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
2020	•	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
2019	•	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
2018	•	Genomic analysis of DNA repair genes and androgen signaling in prostate cancer @ BMC Cancer
		K. Jividen, K. Z. Kedzierska , CS. Yang, K. Szlachta, A. Ratan and B. M. Paschal
2018	•	SONICS: PCR stutter noise correction in genome-scale microsatellites @ Bioinformatics
		K. Z. Kedzierska, L. Gerber, D. Cagnazzi, M. Krützen, A. Ratan, L. Kistler
	@	POSTERS, AND TALKS
2023	•	Chromatin modfiers in endometrial cancer, Poster @ Biology of Genomes 2023, Cold Spring Harbor, NY, USA
2022	•	Systematic characterisation of chromatin modifiers in endometrial cancer, Poster @ European Association for Cancer Research 2022 Congress, Seville, Spain
2019	•	Analysis of the mutational burden across gene sets in cancer, Invited talk @ Polish Bioinformatics Society Symposium, Cracow, Poland
2018	•	Differential mutation analysis across gene sets in cancers, Poster @ Biology of Genomes 2018, Cold Spring Harbor, NY, USA
2017	•	Epigenetic regulation of prostate cancer, Talk @ Visiting Graduate Traineeship Program Grantees Symposium, Charlottesville, VA, USA

Q AWARDS

2023

2021

2019

2018

2017

2016

2022

2021

2020

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

Awarded to outstanding graduate students in genomics and data sciences.

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.

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.

Best Master Thesis in Bioinformatics
@ Polish Bioinformatics Society

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

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.

2015 Grasz o Staz @ PwC Poland

"Grasz o Staz" competition was a national, prestigious and highly competitive (1:25 success rate) scholarship program in Poland organized by PwC.

♣☐ TEACHING EXPERIENCE

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 available 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/.

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 2019 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 2019 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 2017 Invited speaker ATTENDED WORKSHOPS, SUMMER SCHOOLS Sea pen identification from video footage challenge @ Turing Data Study Group @ The Alan Turing Institute, 2022 London, UK Machine Learning Summer School @ Imperial College London, University College London, UK 2019 **€** GRANTS Visegrad Grant to organise NGSchool2022 @ Visegrad Fund 2022 2020 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. 2019 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. NON PROFIT WORK

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

President @ NGSchool Society

2022

2018

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

2019

2016 | 2015

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.