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 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.

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

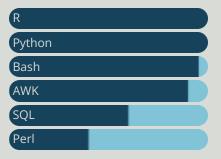
Investigated **epigenetic regulation in prostate cancer** using mouse models of primary, invasive and metastatic tumors. I planned, carried out experimental wokrd and analysed results of RNA-seq, ChIP-seq and ATAC-seq assays.

View this CV online kasia.codes/cv/

CONTACT

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

CODING SKILLS

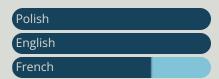


R: tidyverse, data.table, Shiny, plotly, tidymodels, caret and others

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

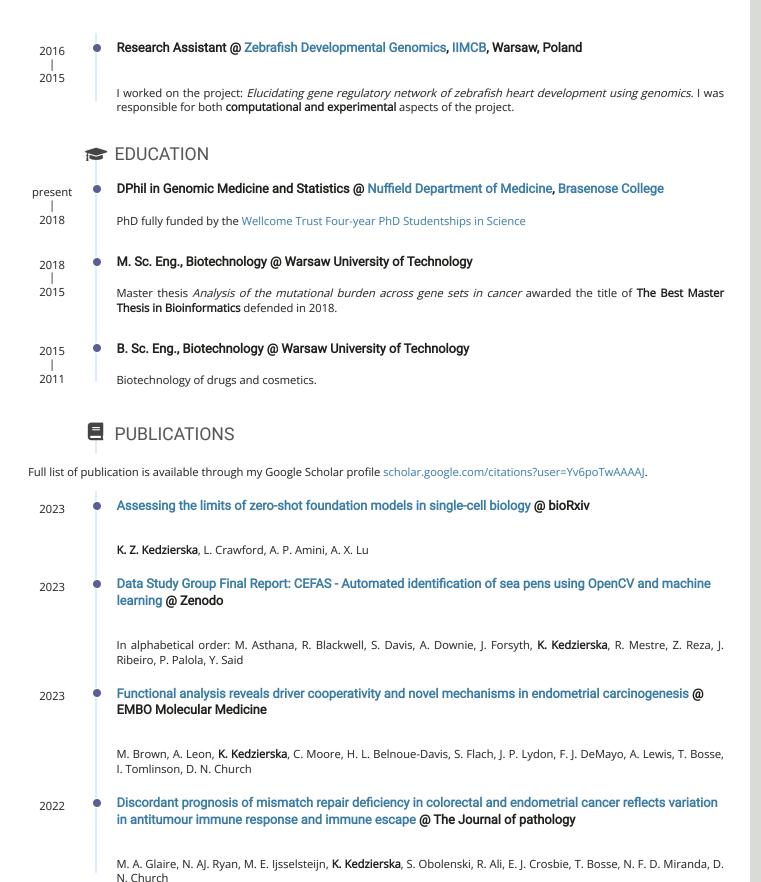
keras/Tensorflow, PyTorch, Seaborn, matplotlib and more

LANGUAGES



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

Last updated on 2023-10-20.



2021	•	Multi-omics analyses of early liver injury reveals cell-type-specific transcriptional and epigenomic shift @ BMC Genomics
		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

Epigenetic regulation of prostate cancer, Talk @ Visiting Graduate Traineeship Program Grantees 2017 Symposium, Charlottesville, VA, USA **Q** AWARDS 2023 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, 2022 **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 Unsupervised learning @ NGSchool2022: Machine Learning in Computational Biology, Jablonna, Poland 2022 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 2021

I led the hackathon in data viusalisation with emphasis on computational biology. Teaching materials are available at

github.com/kzkedzierska/NGSprint data viz.

2020		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/.
2019	•	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).
2019	•	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.
2019	•	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/.
2017	•	ATAC-seq workshop @ #NGSchool2017: Single-cell Sequencing, Jachranka, Poland
		Invited speaker
		ATTENDED WORKSHOPS, SUMMER SCHOOLS
2022	•	Sea pen identification from video footage challenge @ Turing Data Study Group @ The Alan Turing Institute, London, UK
2019	•	Machine Learning Summer School @ Imperial College London, University College London, London, UK
	€	GRANTS
2022	•	Visegrad Grant to organise NGSchool2022 @ Visegrad Fund
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

2022 | 2018 President @ NGSchool Society

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

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.