KASIA KEDZIERSKA

I am a PhD student at the University of Oxford. As a computational biologist, I use Data Science and Machine Learning to answer biological questions. Specifically, I study cancer of the uterus and chromatin organisation in disease progression. Last summer, I joined Computational Biology Department at Novo Nordisk Research Centre in Oxford as an intern, where I was working with NLP methods and knowledge graphs. Currently, I am looking into adapting my experience from the internship to build models for investigating cancer evolution.

➡ SELECTED RESEARCH EXPERIENCE

2021

Intern

Novo Nordisk Research Centre Oxford

Oxford, United Kingdom

• I used NLP based methods to screen biomedical artciles and identify potential therapeutic targets. I built R Shiny App to allow colleagues within the company to investigate results from our pipeline.

present 2018

DPhil Candidate

Wellcome Centre for Human Genetics, Big Data Institute

Q University of Oxford, UK

- In my PhD project Functional and evolutionary characterisation of chromatin organisation in endometrial cancer I am looking at how chromatin organisation influences disease progression in cancer of the uterus.
- Mentored by D. Church and D. Woodcock
- Currently I'm working on building and refining ML models of cancer evolution.

2018 2017 **Visiting Graduate Student**

Ratan group

Q University of Virginia, USA

- Developed SONiCS a tool for genotyping short tandem repeats (STRs) profiled using capture assays.
- Worked on the Master thesis Analysis of the mutational burden across gene sets in cancer.



EDUCATION

present 2018

DPhil. Candidate, Genomic Medicine and Statistics

Nuffield Department of Medicine, Brasenose College

Our University of Oxford, UK

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

2018 2015 M. Sc. Eng., Biotechnology

Warsaw University of Technology

• Warsaw, Poland

• 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.

View this Resume online

kasia.codes/resume/

CONTACT

✓ kasia@well.ox.ac.uk

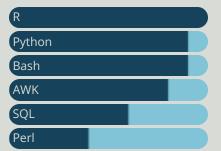
y kzkedzierska

github.com/kzkedzierska

6 kasia.codes

in kzkedzierska

CODING SKILLS



LANGUAGES

Polish	
English	
French	
Estonian	

Made with the package pagedown.

Based on the Nick Strayer's CV package; modified source code for this CV is available here.

Last updated on 2022-09-30.

■ SELECTED PUBLICATIONS

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

Full list of	public	cation on my Google Scholar profile scholar.google.com/citations?user=Yv6poTwAAAAJ.		
2021		Multi-omics analyses of early liver injury reveals cell-type-specific transcriptional and epigenomic shift M. Migdal, E. Tralle, K. A. Nahia, L. Bugajski, K. Z. Kedzierska, F. Garbicz, K. Piwocka, C. L. Winata, M. Pawlak ◆ BMC Genomics		
2020	•	The MLH1 polymorphism rs1800734 and risk of endometrial cancer with micros	satellite instability	
		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		
		Southey, N. E. Mille, I. Tominison, D. Charen, A. B. Sparaic, T. A. O Mara and A. Ecc	♥ Clinical Epigenetics	
2019		 Dynamics of cardiomyocyte transcriptome and chromatin landscape demarcates key events of he development 		
		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	♥ Genome Research	
2018		SONiCS: PCR stutter noise correction in genome-scale microsatellites		
		K. Z. Kedzierska, L. Gerber, D. Cagnazzi, M. Krützen, A. Ratan, L. Kistler	♀ Bioinformatics	
	@	SELECTED TALKS AND POSTERS		
2022	•	Systematic characterisation of chromatin modifiers in endometrial cancer	O Cardilla Carada	
		European Association for Cancer Research 2022 Congress	Seville, Spain	
		Poster presentation		
2019	•	Analysis of the mutational burden across gene sets in cancer		
		Polish Bioinformatics Society Symposium	• Cracow, Poland	
		• Invited talk		
2018	•	Differential mutation analysis across gene sets in cancers		
		Biology of Genomes 2018	Cold Spring Harbor, NY, USA	
		• Poster		
	Ö	SELECTED AWARDS AND HONOURS		
present 2021	•	Senior Hulme Scholarship	_	
		Brasenose College, University of Oxford		
		 Senior Hulme Scholarship is awarded by Brasenose College, University of Oxford to D academic performance is deemed to be exceptional. 	Phil students whose	
2022 2021		Graduate Prize in the 'Outstanding work outside degree' category	•	
		Nuffield Department of Medicine, University of Oxford	Oxford, UK	
		 Each year Nuffield Department of Medicine, based on nominations, awards selected F performance within and outside of their degree. 	PhD students based on their	

ATTENDED WORKSHOPS, SUMMER SCHOOLS

2019 Machine Learning Summer School

Imperial College London, University College London

O London, United Kingdom

♣ TEACHING EXPERIENCE

2022 • 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.

2021 • Data visualization in bioinformatics - hackathon mentor

Online hackathon NGSprint

Discord

• I led the hackathon in data viusalisation with emphasis on computational biology. Under my supervision, 3 teams of around 5 people each, created interactive and captivating visualisation. 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/.

2019 • Introduction to R

2020

2019

2022

2018

Wellcome Centre for Human Genetics

Oxford, United Kingdom

- 8 week course in Introduction to R, Data Manipulation, Data Visualisation and RNA-seq data analysis.
- · Materials available on github/kzkedziersa/r intro

2020 • Introduction to Managing Code with Git

Wellcome Centre for Human Genetics

Oxford, United Kingdom

• 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 Gran
Visegrad Fund

Visegrad Grant to organize #NGSchool2020 postponed until 2022

• 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 9 people and led the organisation of summer school, conference, online seminars and hackathon.

NON-PROFIT WORK

President

NGSchool Society

- The goal of the Society is to promote and support science, with emphasis on computational biology.
- President since 2019; Vice President 2018 2019, Founder