KASIA KEDZIERSKA

I am a PhD student at the University of Oxford. I am a computational biologist, i.e., I use Data Science and Statistical Machine Learning to answer biological questions. Specifically, I study cancer of the uterus and chromatin organisation in disease progression. Currently, I joined Computational Biology Department at Novo Nordisk Research Centre in Oxford as an intern, where I am working with NLP methods and knowledge graphs.

RESEARCH EXPERIENCE

2021

Intern

Novo Nordisk Research Centre Oxford

Oxford, United Kingdom

• I am applying NLP based methods to screen biomedical artciles and identify potential therapeutic targets.

present 2018

DPhil Candidate

Wedge group and Church group

Q University of Oxford, UK

• PhD project: Functional and evolutionary characterisation of chromatin organisation in endometrial cancer

2018 2017 **Visiting Graduate Student**

Ratan group

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.

2017 2016 **Visiting Graduate Student**

Pemberton group

Q University of Virginia, USA

- Worked on Epigenetic regulation in prostate cancer.
- Performed experiments and analyzed data from RNA-seg, ATAC-seg, and ChIP-seq assays.

2016 2015 Research Assistant

Zebrafish Developmental Genomics

Q IIMCB, Warsaw, Poland

- I worked on the project: Elucidating gene regulatory network of zebrafish heart development using genomics.
- I was responsible for both computational and experimental aspects of the project.



EDUCATION

present 2018

DPhil. Candidate, Genomic Medicine and Statistics

Nuffield Department of Medicine, Brasenose College

• University of Oxford, UK

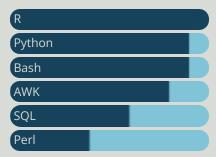
• PhD fully funded by Wellcome Trust Four-year PhD Studentships in Science

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CONTACT

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- in kzkedzierska

CODING SKILLS



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 2021-10-03.

2018 I		M. Sc. Eng., Biotechnology	• Warsaw, Poland	
2015		Warsaw University of Technology	•	
		 Thesis: Analysis of the mutational burden across gene sets in cancer. Thesis awarded the best Master thesis in Bioinformatics defended in 2018 title. 		
2015 		B. Sc. Eng., Biotechnology	• Warsaw, Poland	
2011		Warsaw University of Technology	• • • • • • • • • • • • • • • • • • • •	
		PUBLICATIONS		
2020		The MLH1 polymorphism rs1800734 and risk of endometrial cancer with microsatellite 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, R. L. Milne, I. Tomiinson, D. Church, A. B. Spurdie, T. A. O'Mara and A. Le	wis ♀ Clinical Epigenetics	
2020		Prognostic integrated image-based immune and molecular profiling in early-stage Endometrial Cancer		
2020	Ĭ	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, T. Bosse, H. W. Nijman, V. H. Koelzer and D. N. Church	, V. T. Smit, C. L. Creutzberg,	
			Cancer Immunology Research	
2019		Dynamics of cardiomyocyte transcriptome and chromatin landscape demarcates key events of heart 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		Genomic analysis of DNA repair genes and androgen signaling in prostate cancer	er	
2016		K. Jividen, K. Z. Kedzierska , CS. Yang, K. Szlachta, A. Ratan and B. M. Paschal	♥ BMC Cancer	
2018		SONiCS: PCR stutter noise correction in genome-scale microsatellites		
2010		K. Z. Kedzierska , L. Gerber, D. Cagnazzi, M. Krützen, A. Ratan, L. Kistler	Bioinformatics	
		DOCTEDS AND TALKS		
		POSTERS, AND TALKS		
2019	•	Analysis of the mutational burden across gene sets in cancer	♀ Cracow, Poland	
		Polish Bioinformatics Society Symposium • Invited talk	♥ Cracow, r olarid	
2018		Differential mutation analysis across gene sets in cancers	Cold Spring Harbor, NY, USA	
		The Biology of Genomes 2018 • Poster	•	
2017		Epigenetic regulation of prostate cancer Visiting Graduate Traineeship Program Grantees Symposium	• Charlottesville, VA, USA	
		• Talk		

AWARDS AND HONOURS

2022 Senior Hulme Scholarship

2021

2018

2017

2016

2021

2020

2019

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

2019 **Best Master Thesis in Bioinformatics**

• Analysis of the mutational burden across gene sets in cancer - Best Master Thesis defended in Bioinformatics in 2018 in Poland.

Visiting Graduate Traineeship Program

• 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

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

♣ TEACHING EXPERIENCE

Data visualization in bioinformatics - hackathon mentor

Discord

• I led the hackathon in data viusalisation with emphasis on computational biology. Under my supervision, 3 teams of areound 5 people each, created interactive and captivating visualisation to guide audience through results for selected cancer type. Teaching materials

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).
- · Materials for the Introduction to Python are available on github

2019 • Introduction to R

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/.

2017 • ATAC-seq workshop

#NGSchool2017: Single-cell Sequencing

Q Jachranka, Poland

- Invited speaker
- Materials for the course can be available on gitub.com/kzkedzierska/ATACseq_workshop

ATTENDED WORKSHOPS, SUMMER SCHOOLS

2019

Machine Learning Summer School

Imperial College London, University College London

O London, United Kingdom

€ GRANTS

present | 2020 Visegrad Grant to organize #NGSchool2020 postponed until 2022

Visegrad Fund

• 32,190 EUR awarded towards organising #NGSchool2020 and #NGSymposium. Due to COVID-19 pandemic we organised a virtual events in 2020 & 2021 and are planning in person summer school and conference in 2022.

2019

Visegrad Grant to organize #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

2021 | 2018 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