

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.

SELECTED WORK EXPERIENCE

- 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.
- present | 2018 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 identifying 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).

EDUCATION

- present | 2018 DPhil in Genomic Medicine and Statistics @ Nuffield Department of Medicine, Brasenose College**
PhD fully funded by the **Wellcome Trust Four-year PhD Studentships in Science**
- 2018 | 2015 M. Sc. Eng., Biotechnology @ Warsaw University of Technology**
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.

SELECTED AWARDS AND HONORS

- 2023 JXTX + CSHL 2023 Biology of Genomes Scholarship @ JXTX Foundation, Cold Spring Harbor Laboratory**
Awarded to outstanding graduate students in genomics and data sciences.
- 2022 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.
- 2023 | 2021 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.

SUMMER SCHOOLS & HACKATHONS

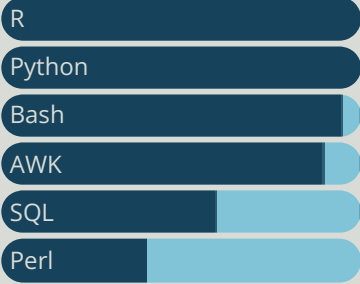
- 2022 Sea pen identification from video footage challenge @ Turing Data Study Group @ The Alan Turing Institute, London, UK**
- 2019 Machine Learning Summer School @ Imperial College London, University College London, London, UK**

View this Resume online
kasia.codes/resume/

CONTACT

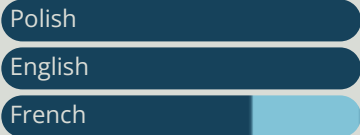
- ✉ kasia@well.ox.ac.uk**
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- 🔗 kasia.codes**
- in kzkedzierska**

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





SELECTED PUBLICATIONS

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

- 2023 • **Functional analysis reveals driver cooperativity and novel mechanisms in endometrial 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
- 2023 • **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
- 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 • **SONiCS: PCR stutter noise correction in genome-scale microsatellites @ Bioinformatics**
K. Z. Kedzierska, L. Gerber, D. Cagnazzi, M. Krützen, A. Ratan, L. Kistler



SELECTED CONFERENCE PRESENTATIONS

- 2023 • **Chromatin modifiers 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



SELECTED 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 visualisation 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 • **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/.



SELECTED GRANTS

- 2022 | 2020 • **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](#).



NON-PROFIT WORK

- 2022 | 2018 • **President @ NGSchool Society**
The goal of the Society is to promote and support science, with emphasis on computational biology.