

## MICHAELA BREZINOVA

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### EDUCATION

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- **UNIVERSITY OF CAMBRIDGE**, PhD in Computational Chemistry (Vendruscolo Lab) **Oct 2023 - Present**
  - Focus on advancing computational approaches to drug discovery (primarily of small molecules) with an emphasis on neurodegenerative disorders, such as Alzheimer's and Parkinson's disease. See *Publications* section.
  - **Technical Skills and Tools:** *Python, TensorFlow, PyTorch, Bash, HPC (Slurm workload manager), AutoDock Vina, Chimera (X), OpenEye Suite, RDKit, Open Babel*
- **UNIVERSITY OF CAMBRIDGE**, MPhil in Computational Biology (**Distinction**) **Oct 2021 - Nov 2022**
  - **Relevant Coursework:** *Genomics I/II, Biological Imaging and Analysis, Deep Learning, Genome Sequence Analysis, Population Genetics Analysis*
- **UNIVERSITY OF EDINBURGH**, BSc Honours in Mathematics and Computer Science (**First Class**) **Sep 2016 - Jun 2020**
  - **Relevant Coursework:** *Honours Algebra, Probability, Bayesian Theory, Algorithms and Data Structures, Data and Analysis, Machine Learning Practical, Processing Formal and Natural Languages, Quantum Information*

### WORK EXPERIENCE

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- **UNIVERSITY OF CAMBRIDGE**, Research Assistant **(Cambridge, UK) May 2023 - Oct 2023**
  - Worked in the Vendruscolo Lab as a part of the global Aligning Science Across Parkinson's initiative (Team Wood). The main focus was unsupervised learning analysis of DAB staining microscopy images of large alpha-synuclein aggregates from Parkinson's patients and developing a kinetic model of aggregate growth.
  - **Technical Skills and Tools:** *Python, TensorFlow, FIJI (ImageJ)*
- **ILLUMINA**, Bioinformatics Intern **(Cambridge, UK) Aug 2022 - Nov 2022**
  - Worked as a part of Genome Quality Group, focused on data analysis of large amounts of genomic data to maximize the quality and breadth of information that can be learnt from a genome and in turn improve the output of the DRAGEN platform (a platform that enables analysis of next-generation sequencing data)
  - **Technical Skills and Tools:** *Nextflow, Make, Bash, BCFTools, R*
- **GOOGLE**, Software Engineer at Google Health **(London, UK) Sep 2020 - Oct 2021**
  - Worked on Care Studio, clinical software to unify healthcare data, with focus on desktop features and front-end latency optimizations
  - **Technical Skills and Tools:** *Soy (Closure templates), TypeScript, Java, C++*
    - Delivering new features to the platform that optimise the data shown to clinicians (required full-stack work)
    - Collaborating with clinicians, product managers as well as UX designers to make sure the features work exactly as intended and are improving the customer experience
    - Designing (under supervision), implementing and delivering the first version of a critical latency feature required to meet the release latency requirements
- **CERN**, Remote Summer Collaboration **(Remote) Jun 2020 - Aug 2020**
  - Replacement of a Summer Student Programme (cancelled due to COVID-19), worked on a profile customisation for ALICE (A Large Ion Collider Experiment) O2 InfoLogger (<https://cds.cern.ch/record/2728718>)
  - **Technical Skills and Tools:** *HyperScript, JavaScript, NodeJS, SQLite*
    - Planning and delivering a new profile customisation for ALICE O2 InfoLogger taking software robustness and durability (10+ years) into account (once the experiment resumes, the program has to be functioning for 10+ years)
- **GOOGLE**, SWE Intern at Google Travel NBU (Next Billion Users) **(Zurich, Switzerland) Jul 2019 - Sep 2019**
  - Worked on a new flexible dates feature for India Rail Travel App
  - **Technical Skills and Tools:** *Soy (Closure templates), TypeScript, Java*
- **GOOGLE**, STEP Intern at Google Account **(Munich, Germany) Jul 2018 - Sep 2018**
  - Worked on the redesign of the Takeout page ([takeout.google.com](https://takeout.google.com)). The new design is currently in production.
  - **Technical Skills and Tools:** *Soy (Closure templates), TypeScript, Java*

### PATENTS

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- Andrews, D.J., **Brezinova, M.**, 2024. Detecting variants in nucleotide sequences based on haplotype diversity (<https://patents.google.com/patent/WO2025090883A1/>)

## PUBLICATIONS

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- **Brezinova, M.**, Fuxreiter, M. and Vendruscolo, M., 2025. DropFit: Determination of the Critical Concentration for Protein Liquid-Liquid Phase Separation. *Journal of Molecular Biology*, p.169294. Server website: <https://www-cohsoftware.ch.cam.ac.uk/index.php/dropfit>
- (To be published in *EMBO Molecular Systems Biology*) **Brezinova, M.**, Brotzakis, Z.F., Horne, R.I., Chowdhury, V.R., Gregory, R.C., Gentile, F. and Vendruscolo, M., (2024). 'Identification of high-affinity secondary nucleation inhibitors of A $\beta$ 42 aggregation from an ultra-large chemical library using Deep Docking'. (<https://www.researchsquare.com/article/rs-4512167/v1>). GitHub link: [https://github.com/MichaelaBrezinova/open\\_source\\_deep\\_docking\\_protocol](https://github.com/MichaelaBrezinova/open_source_deep_docking_protocol)
- Amico, T., Dada, S., Lazzari, A., **Brezinova, M.**, Trovato, A., Vendruscolo, M., Fuxreiter, M., & Maritan, A. (2024) 'A scale-invariant log-normal droplet size distribution below the critical concentration for protein phase separation', *eLife*, 13, RP94214. doi: 10.7554/eLife.94214.2

## TALKS

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- Talk at In Silico #001 event series (<https://luma.com/dw9ijcrm>) (London, UK) Jul 2025
- Short Talk at AI in Drug Discovery and IRB Barcelona Biomedicine Conference (Barcelona, Spain) Apr 2025
- Talk at Bio2Brain Network's (<https://bio2brain.eu/>) AI Workshop (Online) Jun 2024

## SKILLS

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- **Programming Languages:** *Python* (experienced), *Bash* (experienced), *TypeScript* (experienced), *Soy (Closure Templates)* (experienced), *HTML* (experienced), *CSS* (experienced), *Java* (skilful), *R* (skilful), *C++* (skilful), *Haskell* (familiar)
- **Cheminformatics:** *RDKit* (skilful), *Open Babel (Obabel)* (skilful), *Chimera* (skilful), *AutoDock Vina* (skilful), *OpenEye Suite* (skilful)
- **Machine Learning and Data Science:** *TensorFlow* (skilful), *PyTorch* (familiar)
- **Computational Tools and Frameworks:** *Slurm* (experienced), *Nextflow* (skilful), *Make* (skilful)
- **Spoken languages:** *Slovak* (native proficiency), *Czech* (native proficiency), *English* (full professional proficiency), *German* (limited working proficiency), *Mandarin Chinese* (elementary proficiency)

## VOLUNTEERING AND EXTRACURRICULAR - ACADEMIC

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- **Lab Demonstrator at the University of Cambridge** (Cambridge, UK) Jan 2024 - Present
  - Lab demonstrator for computational practicals of Chemistry Part IB course. The main focus is the use of Python, Google CoLab, Avogadro and Orca
- **Code First: Girls, Introduction to Web Development course instructor** (Edinburgh, UK) Jan 2020 - Mar 2020
  - Teaching female students about the front end-development: HTML, CSS, Bootstrap, Git and GitHub
- **Director of Media for the Women in Tech Conference in Edinburgh** (Edinburgh, UK) May 2017 - Jun 2018
  - Organising Women in Tech Conference in Edinburgh, focusing on media and graphic design
- **MathPALS Student Leader at the University of Edinburgh** (Edinburgh, UK) Sep 2017 - Dec 2017
  - Facilitator at Mathematics Peer Assisted Learning program, holding study groups for first-year Mathematics students

## VOLUNTEERING AND EXTRACURRICULAR - OTHER

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- **Secretary at the Cambridge University Massage Society** Jun 2024 - Present

## COMPETITION ACHIEVEMENTS

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- 3rd place at International Team High School Internet Mathematical Olympiad
- 3rd place at National Scientific Research Paper's Competition for high school students in Mathematics and Physics category with paper on topic *Mathematics and Music*