

Kristina ULICNA

PhD in Computational Biology

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PhD Project Summary

Theme: Quantitative labelling of single-cell trajectories in time-lapse microscopy

1. Trajectory reconstruction & lineaging

- Co-developed a robust, supervision-free, deep learning-based cell tracking pipeline for deep lineage analysis of live-cell microscopy 2D cell lines data
- Analysed multi-generational lineage trees of >20k single-cell trajectories to interpret proliferation characteristics predisposing cells to fast divider rates

2. Track representation & interpretation

- Generated an explainable AI model to learn dynamic image representations & interpretable latent space features to map similarities of cell cycle continuity
- Transformed a sequence of image representations into an unsupervised trajectory annotation, classifying cell cycle phases & quantitative confidence scoring over time

Computational Skills

Strongest programming language: **Python**

Experience with **deep learning** strategies, **bioimage analysis** & computer vision apps

- **Fully supervised:** U-Net, ConvNet, TCN
- **Weakly supervised:** multiple-instance
- **Self-supervised:** generative (VQ-) VAEs
- **Unsupervised:** hierarchical clustering
- **Dimensionality reduction:** PCA, UMAP

Skilled in **image representation learning** & time-sequence trajectory data analysis

- **Machine learning libraries:** skimage, sklearn, pytorch (lightning & geometric)
- **Standard & scientific libraries:** numpy, scipy, matplotlib, networkx, napari

Experienced in **software design** (virtual envs, git-based version control, iPython)

Laboratory Skills

- DNA sequencing data readout analysis
- Molecular biology & vector construction
- Cell-based assays & imaging platforms
- Cell / gene engineering & tissue culture
- Gene expression detection methods

Summary

As a **Research Associate** at The Alan Turing Institute, I apply my **deep learning & bioimage analysis skills** across multi-disciplinary projects combining cell biology, computational single-cell tracking & **interpretable AI/ML**. I focus on **Python development** for time series image data analysis to identify **meaningful biological patterns** controlling cell cycle & fate. As a practical & detail-oriented scientist with **biomedical background**, I have demonstrated my **research & leadership skills** in individual & collective settings, gained via work experience in **academic biomedical research groups**, leading **technological companies** & through **community projects**.

Education

Oct 2018 – Dec 2022 **PhD in Biosciences, BBSRC LIDo DTP Programme, UCL | London, UK**

- Fully-funded [doctoral thesis](#): 'Machine Learning for Single-Cell Trajectory Analysis'
- Advisors: Drs [Alan Lowe](#) & [Guillaume Charras](#) | Defended Feb 2023; no corrections
- 3x merit-based scholarships: Yale School of Medicine, Tatra Bank Research Grants

Sep 2014 – Jul 2018 **BSc Biomedical Science (Hons), King's College London | London, UK**

- First Class Hons (**76%**) Biomedical Science with Molecular Biology Extramural Year
- Awards & Scholarships: Desmond Tutu Scholarship '14, Best Lay Article Award '15

Industrial & Academic Experience

Feb 2023 – Present **Research Associate @ The Alan Turing Institute | London, UK**

- Co-developing [graph representation analysis](#) for [connected embeddings](#) ([GRACE](#)) for automated object identification of structural patterns in (bio-)imaging datasets
- Built an explainable, autoencoder-driven image representation learning [framework](#) for dynamic single-cell trajectory analysis for self-supervised cell cycle annotation
- **"Best Poster Award"** at 2023 ICML Comp Bio [Workshop](#), sponsored by CZI & GSK

Apr 2021 – Sep 2021 **Research Intern @ Microsoft Research Cambridge | Cambridge, UK**

- Developed an AI-based end-to-end pipeline to classify subcellular protein localisation from Human Protein Atlas' [Kaggle dataset](#) of weakly labelled microscopy images
- Trained a competitively-performing model, co-evaluated as best off-chart approach

Sep 2016 – Aug 2017 **Industrial Trainee @ Crescendo Biologics Ltd. | Cambridge, UK**

- Engineered a novel, universal tool cell line for early drug discovery, i.e. phage display selection & functional screening of antibody fragment onco-therapeutics

Jun 2016 – Sep 2016 **Cancer Research UK Intern @ Cambridge University | Cambridge, UK**

- S. Bohndiek Lab: Characterised anti-angiogenic drug effects via breast cancer cell-based growth, viability assays to evaluate oxygen role in cancer progression

Jun 2015 – Aug 2015 **Visiting Scholar, Whitehead Institute @ MIT | Cambridge, MA, USA**

- R. Weinberg's Lab: Investigated determinants of cancer cell invasion, metastasis & tumour stroma immunomodulation upon epithelial-to-mesenchymal transition

Teaching & Outreach Experience

Nov 2023 **Guest Lecturer, Data Science @ UCL STAT0042 module | London, UK**

- Co-designed iPython [practical sessions](#) for 70+ undergraduates to introduce common bioinformatics concepts in bioimage analysis to train a cell state classifier

Mar 2020 – Mar 2022 **Graduate Teaching Assistant @ UCL BIOC0016 module | London, UK**

- Co-designed iPython [practical sessions](#) for 70+ undergraduate students to introduce concepts in bioimage analysis & bioinformatics to train an ML cell state classifier

Sep 2021 – May 2022 **Master's Student Daily Supervisor @ UCL | London, UK**

- Directly supervised a Biochemistry Masters student's to develop an AI-driven cell segmentation tool from microscopy datasets via research guidance & references

Jan – Dec 2021 **Google Certified Trainer for AI Tech & Tools | Bratislava, Slovakia**

- Empowering non-tech professionals by leveraging their expertise & leadership in AI technical field via [AI-basics talks & workshops](#) with Google Slovakia outreach team

May – Jun 2021 **"Smart Microscopy" Workshop @ ZEISS | Gothenburg, Sweden**

- Trained 30+ intermediate-level attendees at ["Train Your Own Model"](#) hands-on workshop session to use python image analysis tools to annotate cell imaging data

May - Jun 2020 **"Introduction to Deep Learning" @ UCL Cancer Domain | London, UK**

- Delivered beginner-friendly [masterclass series](#) for 100+ interdisciplinary scientists with real-life examples of deep learning-based bioimage analysis from PhD project

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certificates

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