

# Kristina ULICNA

PhD in Computational Biology

📍 London, United Kingdom

✉ [kristina.smith.ulicna@gmail.com](mailto:kristina.smith.ulicna@gmail.com)

🐦 [Twitter](#) | [in LinkedIn](#) | [GitHub](#)

🔍 [Google Scholar](#) | [My Webpage](#)

## PhD Project Summary

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

### 1. Trajectory reconstruction & lineage

- 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 & temporal landscape visualisation of diverse cell fates

*Directly supervised* a Masters student's project to develop an AI-driven cell segmentation tool from microscopy datasets

## Computational Skills

Strongest programming language: **Python**

Experience with deep learning strategies for image analysis & computer vision apps

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

Skilled in "image2seq" representations & time-sequence trajectory data analysis

- **Machine learning libraries:** skimage, sklearn, torch, pytorch lightning, pytorch geometric, btrack, arboretum
- **Standard & scientific libraries:** numpy, scipy, pandas, matplotlib, seaborn, h5py, networkx, napari, dtaiDistance

Experienced in **conda** envs, **GitHub** (git), **LaTeX** (overleaf), **iPython** (jupyter & colab)

## 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-developed 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
- Apr 2021 – Sep 2021 **Research Intern @ Microsoft Research Cambridge | Cambridge, UK**
- Developed an AI-based end-to-end pipeline to classify subcellular protein localisation in single cells from Human Protein Atlas' [Kaggle dataset](#) of weakly labelled microscopy images using Azure computing & [InnerEye Deep Learning](#) OS toolkit
  - Trained a competitively-performing model (MIL / SimCLR & BYOL methodology) in collaboration with competition [organisers](#) evaluated as best approach off-chart
- 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, USA**
- R. Weinberg's Lab: Investigated determinants of cancer cell invasion, metastasis & tumour stroma immunomodulation upon epithelial-to-mesenchymal transition

## Teaching & Outreach Experience

- Mar 2020 – Mar 2022 **Graduate Teaching Assistant @ UCL BIOC0016 module | London, UK**
- Co-designed iPython [practical sessions](#) for 70+ undergraduates to introduce concepts in bioimage analysis & bioinformatics to train an ML cell state classifier
- Jan – Dec 2021 **Google Certified Trainer for AI Tech & Tools | Bratislava, Slovakia**
- Empowering non-tech professionals by leveraging their expertise & leadership in AI tech 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 (bio-)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

## Leadership & Teamwork Experience

- Sep 2019 – Feb 2023 **Student Ambassador, LIDo PhD Programme @ UCL | London, UK**
- Outlined programme structure & shared own experience with new student cohorts
  - Guided individual students through responsibilities with changing rotation projects
- Sep 2015 – Aug 2017 **Jury Member, LEAF Award @ LEAF | Bratislava, Slovakia**
- Shortlisted self-driven, talented students with community involvement in jury team
- Sep 2016 – Aug 2018 **University Mentor, Talent Guide @ LEAF | Bratislava, Slovakia**
- Counselling college choices with gifted high-schoolers & edited personal statements

- 2021 **30 under 30** @ [\*Forbes Slovakia\*](#)
- 2019 **European Union** Council Delegate
- 2019 **Travel Grant:** *Students to the World*
- 2019 **Travel Grant:** *Talents of New Europe*
- 2017 **United Nations** Assembly Delegate
- 2017 **GSK Healthcare** STEM Awardee
- 2014 'Absolute Winner at the Festival of Science & Technology' @ AMAYET
- 2013 'The Special Award by the Dean of the Faculty of Natural Sciences'

- 2023 **Pint of Science Event** | London, UK
- 2022 **Reflect Festival** | Limassol, Cyprus
- 2015 **Universal Expo Milano** | Milan, Italy
- 2014 **Intel ISEF Finals** | Los Angeles, USA
- 2013 **Global Scholars Programme @ ALA**  
| Johannesburg, South Africa
- 2013 **International Congress of Young Investigators** | Zaragoza, Spain

The infographic displays four horizontal bars representing language proficiency levels. Each bar is composed of 10 segments. The first four segments are dark blue, and the remaining six are light blue. The bars are labeled as follows:

- Slovak**: native proficiency
- Czech**: bilingual proficiency
- English**: full professional proficiency
- German**: limited working proficiency

Below the German bar, the text "IELTS & iBT TOEFL language test certificates" is written in a smaller font.

- Ballroom dancing; standard & Latin style (UK national partner competitions level)
- Bachata & salsa social dancing (int/adv)
- Long-distance running (half-marathons)
- Playing tennis (competitively at school)

- 2022 **Reflect Festival** "AI for Science" [talk](#)
- 2022 **Forbes Slovakia** laureate [survey](#)
- 2021 **Forbes Slovakia** 30 under 30 [chart](#)
- 2021 "Zijem Vedu" researcher [interview](#)
- 2020 **PyLadies Dublin** key note [interview](#)
- 2020 **UCL Cancer Domain** [masterclass](#)
- 2020 **StartItUp** research profile [interview](#)

- Member of the University of London DanceSport Society (active competitor)
- Member of London Stem Cell Network
- Member of British Pharmacol Society
- EU / Slovak driving licence | *B type*

Dec 2022	<b>Moroccan Advanced Science Institute</b>	Rabat, Morocco
	• 'AI for Science' Workshop with poster presentation to research industry sponsors	
Sep 2022	<b>Weizmann Institute of Science</b>	Rehovot, Israel
	• 'International School of Biological Physics of Cells', part of EMBO PhysCell 2022	
Apr 2022	<b>Institut Curie &amp; Sorbonne Université</b>	Paris, France
	• 'International course of Cell Biology & Cancer' with the 'Science & Life' workshop	
Dec 2020	<b>Jumping Rivers Ltd.</b>	London, UK
	• Two intermediate courses: 'Machine Learning with Python', 'Python & Tensorflow'	
Aug 2019	<b>University of Hong Kong</b>	Hong Kong
	• Practical course 'Advanced Imaging: Deep Learning in Live Imaging & Cell Biology'	
Jun 2019	<b>University of Genova</b>	Genova, Italy
	• 'Machine Learning Crash Course' covering theoretical foundations & core concepts	
Sep 2016	<b>University of Cambridge</b>	Cambridge, UK
	• Two beginners courses: 'Solving Biological Problems', 'Statistical Analysis' using R	
Jun - Jul 2012	<b>Johns Hopkins University</b>	Baltimore, USA
	• Interactive class 'The History of Disease' by Centre for Talented Youth (CTY JHU)	

2023	Speaker	Crick BioImage Analysis Symposium (CBIAS) @ The Francis Crick Institute
	Speaker	Network of EU BioImage Analysts Symposium (NEUBIAS) @ Porto, Portugal
2021	Speaker	Machine Learning Methods Advances @ Recursion CytoData Society Meeting
	Speaker	AI Microscopy Symposium   The Allen Institute for Brain Science @ Seattle, US
	Panelist	"See the Hidden": Future of AI in Microscopy Workshop @ Leica Microsystems
	Panelist	"Women in AI for Global Health" Session @ Mozilla Festival Tech Conference
2020	Speaker	UCL BioImage Analysis Interest Group @ Laboratory of Molecular Cell Biology
	Speaker	Imperial College Brain Sciences Seminar @ UK Dementia Research Institute
	Speaker	Python Software Foundation Pro Network Meetup @ #PyLadies Dublin

2023	Speaker	'Work-in-progress' spotlight @ CVPR CVMI Workshop @ Vancouver, Canada
2022	Poster	'AI for Science' Workshop @ Advanced Science Institute @ Rabat, Morocco
	Poster	EMBO Workshop: Physics of Cells (PhysCell Conf 2022) @ Ein Gedi, Israel
	Poster	Women in Machine Learning (WiML) Panel @ ICML Conference Workshop
	Speaker	The Crick Annual PhD Student Symposium @ The Francis Crick Institute
2021	Speaker	Health Intelligence Lab   Internship project summary @ Microsoft Research
	Poster	Women in Machine Learning (WiML) Panel @ NeurIPS Conference Workshop
	Poster	<u>C</u> rick <u>B</u> io <u>I</u> mage <u>A</u> nalysis <u>S</u> ymposium (CBIAS) @ The Francis Crick Institute
	Poster	Women in Data Science (WIDS) Worldwide Conference @ Stanford University
	Speaker	Virtual Seminars in Biomedical Science @ Imperial College London
2020	Poster	London Stem Cell Network Annual Symposium @ The Francis Crick Institute
	Poster	Society of Biomolecular Imaging & Informatics High Content Conference
	Speaker	UCL Institute of Structural & Molecular Biology Postgraduate Symposium
2019	Speaker	UCL Institute of Structural & Molecular Biology Friday Wrap @ Birkbeck Univ.
	Speaker	Quantitative Systems Biology Workshop @ King's College London
	Poster	UCL Graduate Student Symposium @ The Francis Crick Institute

**Ulicna, K.,** Kelkar, M., Soelistyo, C.J., Charras, G.T. & Lowe, A.R. (2023). *Learning dynamic image representations for self-supervised cell cycle annotation*. bioRxiv 2023.05.30.542796; doi: <https://doi.org/10.1101/2023.05.30.542796> | [BioRxiv](#) | [GitHub](#) repository (under review)

Soelistyo, C.J., **Ulicna, K.** & Lowe, A.R. (2023). *Perspective: Machine learning enhanced cell tracking*. (under review)

**Ulicna, K.,** Ho, L.T.L., Soelistyo, C.J., Day, N.J. & Lowe, A.R. (2022). *Convolutional neural networks for classifying chromatin morphology in live cell imaging*. Methods in Molecular Biology, Chromatin Architecture, Springer Nature Protocols | [Springer](#) | [GitHub](#) repository

**Ulicna, K.,** Vallardi, G., Charras, G. & Lowe, A.R. (2021). *Automated deep lineage tree analysis using a Bayesian single cell tracking approach*. Frontiers in Computer Science, Computer Vision: Methods & Tools for Bioimage Analysis | [Frontiers](#) | [BioRxiv](#) | [GitHub](#) repository