HENRIK ÅHL, PHD Postdoctoral Research Associate

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EXPERIENCE

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Ongoing December 2021	Research Associate, Sainsbury Laboratory, University of Cambridge, Cambridge, UK Developed computer simulation frameworks with a strong integrative focus, combining 3/4D experimental data, computational analysis, and theoretical models to elucidate how plants grow. Applied state-of-the-art data-driven methodologies, with an emphasis on high-throughput pipelines, computer vision, statistical analysis, and data-driven mathematical modelling. Deterministic GRN modelling FEM mechanical modelling Computer vision Numerical simulations Confocal imaging Machine learning: Support vector machines High-performance computing Spatiotemporal patterning analysis
September 2016 August 2016	Research Assistant, Sainsbury Laboratory, University of Cambridge, Cambridge, UK Established a stochastic partial differential equation framework to simulate the regulatory network and resulting gene expression patterns which govern stem cell identity in plant shoots. [Numerical simulations] Optimisation Stochastic GRN modelling High-performance computing Itô calculus
June 2016 July 2015	Supplemental Instructor, DEPT. OF PHYSICS, LUND UNIVERSITY, Lund, Sweden Led weekly supplemental exercise sessions for undergraduate students in physics. Groups of up to 25 people. Lecturing Scientific communication
June 2015 August 2015	Research Assistant, DEPT. OF THEORETICAL PHYSICS AND ASTRONOMY, LUND UNIVERSITY, Lund, Sweden Investigated the consequences for evolvability of artificial genomes due to various mutation operations. Strong code optimisation focus in C++.

Simulated evolution (Genetic algorithms Linux Sequence alignment High-performance computing

EDUCATION

2021 2017	PhD in Applied Mathematics and Theoretical Physics, UNIVERSITY OF CAMBRIDGE, Cambridge, UK Thesis: Integrative high-throughput analyses of aerial morphodynamics in plants. 3D Segmentation (Numerical methods) (Confocal imaging) (GRN modelling) (FEM mechanical modelling) (Support vector machines) (Quantitative phenotyping) (Single-cell tracking) (Image registration)
2017	MPhil in Computational Biology, UNIVERSITY OF CAMBRIDGE, Cambridge, UK
2016	Thesis: In vivo single cell dynamics of the Arabidopsis thaliana aerial stem cell niche. With distinction.
	Computer vision Functional genomics Genome informatics Structural biology Systems biology
2016 2014	BSc in Theoretical Physics, LUND UNIVERSITY, Lund, Sweden Thesis: Linking the dynamics of genetic algorithms to the encoding of information. With distinction Artificial neural networks Mechanics Quantum physics Genetic algorithms Scientific programming
2013	BA in Theoretical Philosophy (Unfinished), LUND UNIVERSITY, Lund, Sweden
2011	Courses in Philosophy, Computer Science, Mathematics, Economics, and Latin.
	Scientific & Philosophical methodology Logics Analysis & Algebra Software development & design principles

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PUBLICATIONS AND SELECT PRESENTATIONS

- Åhl, H., Willis, L, & Jönsson, H. (2022). Regulation of floral phyllotaxis by auxin transport proteins. *Preprint*; in preparation.
- 2022 Åhl, H., Zhang, Y, & Jönsson, H. (2022). POM2/CSI1 modulates the shoot dome shape through mechanical regulation. Preprint; in preparation.
- Scandinavian Plant Physiology Symposium. Longyearbyen, Svalbard. Oral presentation 2022
- Åhl, H., Zhang, Y., & Jönsson, H. (2022). High-throughput 3D phenotyping of plant shoot apical meristems. 2022 Frontiers in Plant Science, p.712.
- 2021 Sainsbury Laboratory Seminar. Sainsbury Laboratory at the University of Cambridge, Cambridge, UK. Oral presentation
- Åhl, H. (2021). Integrative high-throughput analyses of aerial morphodynamics in plants (Doctoral disser-2021 tation). University of Cambridge.
- Morphogenesis seminar, Remote. Oral presentation 2020
- 2019 Bhatia, N., Åhl, H., Jönsson, H., & Heisler, M. G. (2019). Quantitative analysis of auxin sensing in leaf primordia argues against proposed role in regulating leaf dorsoventrality. Elife, 8, e39298.
- 12th Plant Development Conference, Retzbach, Germany. Poster presentation 2019
- Merlevede, A., Åhl, H., & Troein, C. Homology and linkage in crossover for linear genomes of variable length 2019 (2019). PLoS ONE, 14(1), e0209712.
- Sainsbury Laboratory Seminar. Sainsbury Laboratory at the University of Cambridge, Cambridge, UK. Oral 2018 presentation
- Montpellier Morphogenesis Spring School. Montpellier, France. Oral presentation 2017

COMPETENCIES

Scientific computing Git, SVN, Eclipse, Spyder, Linux (Ubuntu, OpenSUSE), Slurm, Sh/Bash/Zsh

> Adobe (Photoshop, Illustrator), Inkscape Graphical

Vim, ETFX, Microsoft Office (Word, Excel, PowerPoint), WPS Writing and presentation



LANGUAGES

Swedish				
English				
German		\bigcirc	\bigcirc	\bigcirc



Python	••••
R	
Java	
C++	

VOLUNTEERING

2020-	Reviewer, PLoS ONE
2020-2021	Instructor, St Edmund's College Tabata Club
2019-2022	Pro se litigation, EW High Court & CFC
2017-2021	Founder, St Edmund's College Poetry Society
2015	Coordinator, Fjällräven Classic
2013-2014	Coordinator, Ordkonst
2013-2014	Producer and Show host, Lyriklådan, Radio AF

AWARDS

2021	Crane's Charity Award
2020	University of Cambridge Support Award
2020	St Edmund's College Monetary Award
2019	St Edmund's College Monetary Award
2017	Montpellier Spring School Travel Award
2017	St Edmund's College Sports Award
2014	Akademiska Föreningen Annual Award
<u>Total</u>	£6000

Personal interests

LEISURE: Chess, Photography, Nature & Wildlife

SPORT: Climbing, Ultimate frisbee, Powerlifting, Tabata

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