

Elan Ness-Cohn, Ph.D.

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EDUCATION

Ph.D. (Biomedical Research)

Northwestern University, Evanston

2022

- ▶ Dissertation: *Topology Inspired Methods for the Design and Analysis of Transcriptomic Time-Series Data in Circadian Biology Research*
- ▶ Advisor: Prof. R. Braun, Molecular Biosciences (Comp. Bio and Applied Math Group)

Graduate Certificate in Teaching

Northwestern Searle Center for Advancing Learning and Teaching

2021

B.Sc. (Biology), Concentration (Education)

Massachusetts Institute of Technology, Cambridge

2017

- ▶ Thesis: *Regenerative Medicine and Synthetic Biology: Working Toward the Clinic*
- ▶ Research Advisor: Prof R. Jaenisch, Biology (Stem Cell Group)
- ▶ Research Advisor: Prof D. Kim, Biology (Genetics Group)

EXPERIENCE

Chicago Biomedical Consortium (CBC)

Entrepreneurial Fellow

2022–date

- ▶ Diligenced early-stage therapeutic proposals for the allocation of \$250K grants to accelerate the translation of transformative biomedical technologies across Chicagoland's flagship Universities.
- ▶ Developed robust investment theses and project plans defining competitive positionings, timelines, budgets and resource allocations, mitigation plans, competitive landscapes analyses, and clinical/regulatory strategies to accelerate the commercialization of CBC funded projects.

Braun Lab

Ph.D. Candidate, Northwestern University

2017–2022

- ▶ *Topology Inspired Methods for the Design and Analysis of Transcriptomic Time-Series Data in Circadian Biology Research*
- ▶ Developed novel reference-free, non-parametric methods using principles from dynamical system theory, topological analysis, and inferential statistics to identify circadian genes with chronotherapeutic potential.
- ▶ Lead multiple collaborative projects leveraging inter-disciplinary expertise resulting in 3+ first author publications in high profile academic journals including *Science* and *Bioinformatics*.
- ▶ Trained/mentored undergraduate and graduate students in various computational research and data management techniques.

Jaenisch Lab

Undergraduate Research Assistant, Massachusetts Institute of Technology

2016–2017

- ▶ *Engineering Human Pluripotent Stem Cells With Insulin Reporter to Model Type 1 Diabetes*
- ▶ Designed/cloned fusion gene plasmid and accompanying CRISPR/Cas9 guide RNAs for targeted tagging of gene of interest
- ▶ Supported in the *in vitro* differentiation of pluripotent stem cells to insulin producing beta-like cells

Kim Lab

Undergraduate Research Assistant, Massachusetts Institute of Technology

2015–2016

- *Regulation of Stress Physiology and Longevity by the EIF3 Translation Initiation Complex in C. Elegans*
- Designed/cloned fusion gene plasmid and accompanying CRISPR/Cas9 guide RNAs for targeted tagging of gene of interest
- Generated CRISPR/Cas9 transgenic lines of *C. Elegans* through microinjection

COMPUTATIONAL & TECHNICAL SKILLS

Programming: R • Python • Bash • R Shiny • L^AT_EX • git • SQL • HPC (SLURM)

Machine Learning: Tidymodels • Keras • Tensorflow

Data Mining: statistical analysis • dimensionality reduction • clustering • visualization

Modeling: dynamical systems • topological analysis • toy model development

Research: algorithm & pipeline development • inter-disciplinary collaboration • software development

Laboratory: CRISPR/Cas-9 genome editing • plasmid design • molecular cloning • qPCR

Languages: English • Hebrew

SOFTWARE (HIGHLIGHTS)

Fasano-Franceschini Test - R Package

An open-source implementation of the Fasano and Franceschini test – a 2-D Kolmogorov-Smirnov (KS) two-sample test – downloaded by 6K+ in the R community.

R Package Website: [\[Link\]](#)

TimeCycle - R Package

A non-parametric method that leverages results from dynamical systems theory and topology to test whether a dynamical variable (gene expression) exhibits cycling dynamics.

R Package Website: [\[Link\]](#) • **Video Tutorial:** [\[Link\]](#)

TimeTrial - R Shiny Web Application

An interactive software suite that enables circadian researchers to perform head-to-head comparisons of four leading cycle detection methods using both synthetic and biological data.

R Shiny Synthetic Data: [\[Link\]](#) • **R Shiny Biological Data:** [\[Link\]](#) • **Video Tutorial:** [\[Link\]](#)

Additional work can be found on my GitHub profile: [🔗 github/nesscoder](#)

PUBLICATIONS

- [1] **Ness-Cohn, Elan** and Rosemary Braun. TimeCycle: Topology Inspired Method for the Detection of Cycling Transcripts in Circadian Time-Series Data. *Bioinformatics*, jun 2021.
- [2] **Ness-Cohn, Elan**, Ravi Allada, and Rosemary Braun. Comment on “Circadian rhythms in the absence of the clock gene Bmal1”. *Science*, 372(6539), 2021.
- [3] **Ness-Cohn, Elan**, Marta Iwanaszko, William L Kath, Ravi Allada, and Rosemary Braun. TimeTrial: An interactive application for optimizing the design and analysis of transcriptomic times-series data in circadian biology research. *J Biol Rhythms*, 35:439–451, 2020.
- [4] Sophia B. Gibson, **Ness-Cohn, Elan**, and Erik C. Andersen. Benzimidazoles cause lethality by inhibiting the function of *Caenorhabditis elegans* neuronal beta-tubulin. *International Journal for Parasitology: Drugs and Drug Resistance*, 20:89–96, dec 2022.
- [5] Douglas J. Cattie, Claire E. Richardson, Kirthi C. Reddy, **Ness-Cohn, Elan**, Rita Droste, Mary K. Thompson, Wendy V. Gilbert, and Dennis H. Kim. Mutations in nonessential eIF3k and eIF3l genes confer lifespan extension and enhanced resistance to ER stress in *caenorhabditis elegans*. *PLoS Genetics*, 12(9):e1006326, 2016.

MANUSCRIPTS SUBMITTED / IN PREPARATION

- [1] **Ness-Cohn, Elan** and Rosemary Braun. Fasano-Franceschini Test: an Implementation of a 2-Dimensional Kolmogorov-Smirnov test in R. *arXiv*, page 2106.10539, sept 2021.
- [2] **Ness-Cohn, Elan**, Dae-Sung Hwangbo, Yong-Jae Kwon, Marta Iwanaszko, William L Kath, Ravi Allada, and Rosemary Braun. TimeChange: Topology Inspired MMethod for the detection of differential cycling dynamics in circadian transcriptomic time-series data. *In Preparation*, 2021.

TEACHING

Northwestern University – Searle Teaching Certificate Program 2020 – 2021

A year long sequence of seminars, special-topics workshops, and guidance provided by peer and faculty mentors focused on improving student learning and inclusion in their disciplines.

Designed/developed an *Intro to Data Science Lab Course* complete with a course outline, sample lesson plan, sample assessments, grading scheme, evaluation plan, and interpretation of teaching effectiveness.

Northwestern University

What Do Your Data Say? (DataScience Bootcamp) , Teaching Assistant	Spring 2020
IGP 486: Advanced Bioinformatics/Genomics , Teaching Assistant	Winter 2020
Collabrative Learning & Intergrated Mentoring in the BioSciences (CLIMB)	
IGP 405: Cell Biology, Teaching Mentor	Winter 2021
IGP 484: Biostatitics, Teaching Mentor	Fall 2020
IGP 484: Biostatitics, Teaching Mentor	Fall 2019

Massachusetts Institute of Technology

7.012: Introductory Biology , Teaching Assistant	Spring 2017
18.05: Introduction To Probability and Statistics , Teaching Assistant	Spring 2016
Department of Biology Tutoring Program , Tutor	2016–2017
Introductory Biology, Introductory Experimental Biology and Communication, Genetics, Biochemistry, Cell Biology	

PRESENTATIONS

<i>Topological Strategies for the Analysis of Rhythmic Dynamics in Transcriptomic Time-Series Data</i>	
Talk , International Conference on Intelligent Systems for Molecular Biology (ISMB), Virtual	2021
<i>TimeCycle: Topology Inspired MMethod for the Detection of Cycling Transcripts In Circadian Time-Series Data</i>	
Microtalk , 3rd Annual Conference on Quantitative Approaches in Biology, Northwestern	2020
<i>TimeCycle: Topology Inspired MMethod for the Detection of Cycling Transcripts In Circadian Time-Series Data (Canceled Covid-19)</i>	
Minisymposium , Conference on the Life Sciences (LS20), California	2020
<i>TimeCycle: Topology Inspired MMethod for the Detection of Cycling Transcripts In Circadian Time-Series Data</i>	
Poster , 3rd Annual Southeaster Center for Mathematics & Biology Symposium, Georgia Tech	2020
<i>TimeCycle: Topology Inspired MMethod for the Detection and Direct Comparison of Cycling Transcripts Across Conditions</i>	
Poster , 2nd Annual Conference on Quantitative Approaches in Biology, Northwestern	2019
<i>TimeTrial: Interactive Application for the Design and Analysis of Transcriptomic Time-Series Data in Circadian Biology Research</i>	
Poster , Chicago Biomedical Informatics Data Jam, Chicago	2019

TimeCycle: Topology Inspired MMethod for the Detection and Direct Comparison of Cycling Transcripts Across Conditions
Poster, Chicago Biomedical Informatics Data Jam, Chicago 2019

TimeCycle: Topology Inspired MMethod for the Detection and Direct Comparison of Cycling Transcripts Across Conditions
Poster, 1st Annual Conference on Quantitative Approaches in Biology, Northwestern 2018

MENTORSHIP

May Nguyen, Undergraduate Research Assistant
 Quantitative Biology Undergraduate Summer Research Program Summer 2021
Coupling Environmental Inputs to Circadian Outputs in Drosophila

Kelly Paquin, Undergraduate Research Assistant
 Quantitative Biology Undergraduate Summer Research Program Summer 2020
Automating the Detection of Circadian Genes from Luciferase Reporter Construct in Drosophila

Carolina Clark, Ph.D. Candidate
 DGP Student Assisted Mentoring Program (STAMP) 2020–2022

Samuel Hamilton, Ph.D. Candidate
 DGP Student Assisted Mentoring Program (STAMP) 2019–2022

AWARDS

NSF-Simons Center for Quantitative Biology Travel Grant
 Northwestern University 2021

Boit Prize for Engineering Writing
 Massachusetts Institute of Technology 2017
 Senior Thesis - *Regenerative Medicine and Synthetic Biology: Working Toward the Clinic*

SERVICE

Professional Memberships and Leadership Roles

Driskill Graduate Program In the Life Sciences, Student Council Rep 2020–2022

NSF-Simons Center for Quantitative Biology Leadership Team, Graduate Student Rep 2019–2022