# Elan Ness-Cohn

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github.com/nesscoder in linkedin.com/nesscohn

### EDUCATION

# Ph.D. (Biology)

Northwestern University, Evanston

08/2017–12/2021 (Anticipated)

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

2020-2021

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

Massachusetts Institute of Technology, Cambridge

2017

Thesis: Regenerative Medicine and Synthetic Biology: Working Toward the Clinic

Research Advisor: Prof D. Kim, Biology (Genetics Group) Research Advisor: Prof R. Jaenisch, Biology (Stem Cell Group)

#### EXPERIENCE

#### **Braun Lab**

### Ph.D. Candidate, Northwestern University

08/2017-date

Topology Inspired Methods for the Design and Analysis of Transcriptomic Time-Series Data in Circadian Biology Research

- Developed a suite of open-source software packages for optimizing the design and analysis of \*omic time-series experiments in circadian biology research.
- Lead multiple collaborative projects with peers leveraging inter-disciplinary expertise to investigate hypotheses and problem-solve obstacles.
- Mentored undergraduate and graduate students through various computational research techniques and organizational skills.

#### Jaenisch Lab

### Undergraduate Research Assistant, Massachusetts Institute of Technology

2016-2017

Engineering Human Pluripotent Stem Cells With Insulin Reporter to Model Type 1 Diabetes

- $\bullet$  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 Tanslation 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

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### COMPUTATIONAL & TECHNICAL SKILLS

Programming: R • Python • Bash • R Shiny • LATEX• git • SQL • HPC (SLURM)

Machine Leaning: 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 – used by 1600+ 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: Q 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 Bmall". *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] 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 (in review at The R Journal).
- [2] Ness-Cohn, Elan, Dae-sung Hwangbo, William L Kath, Ravi Allada, and Rosemary Braun. TimeChange: Topology Inspired MEthod for the detection of differential cycling dynamics in circadian transcriptomic time-series data. *In Preparation*, 2021.

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## 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
Collaborative I compine to Interpreted Montaning in the DieSciences (CLIMD)	

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 -

Topological Strategies for the Analysis of Rhythmic Dynamics in Transcriptomic Time-Series Data	
Talk, International Conference on Intelligent Systems for Molecular Biology (ISMB), Virtual	2023

TimeCycle: Topology Inspired MEthod for the Detection of Cycling Transcripts In Circadian Time-Series Data
Microtalk, 3rd Annual Conference on Quantitative Approaches in Biology, Northwestern 2020

TimeCycle: Topology Inspired MEthod for the Detection of Cycling Transcripts In Circadian Time-Series Data (Canceled Covid-19)

Minisymposium, Conference on the Life Sciences (LS20), California

TimeCycle: Topology Inspired MEthod for the Detection of Cycling Transcripts In Circadian Time-Series Data

Poster, 3rd Annual Southeaster Center for Mathematics & Biology Symposium, Georgia Tech

2020

TimeCycle: Topology Inspired MEthod for the Detection and Direct Comparison of Cycling Transcripts Across Conditions

Poster, 2nd Annual Conference on Quantitative Approaches in Biology, Northwestern 2019

TimeTrial: Interactive Application for the Design and Analysis of Transcriptomic Time-Series Data in Circadian Biology Research

Poster, Chicago Biomedical Informatics Data Jam, Chicago

2019

2020

TimeCycle: Topology Inspired MEthod for the Detection and Direct Comparison of Cycling Transcripts Across Conditions

Poster, Chicago Biomedical Informatics Data Jam, Chicago

2019

TimeCycle: Topology Inspired MEthod for the Detection and Direct Comparison of Cycling Transcripts Across Conditions

Poster, 1st Annual Conference on Quantitative Approaches in Biology, Northwestern

2018

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# 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 Summer 2020 Quantitative Biology Undergraduate Summer Research Program Automating the Detection of Circadian Genes from Luciferase Reporter Construct in Drosophila Carolina Clark, Ph.D. Candidate DGP Student Assisted Mentoring Program (STAMP) 2020-date Samuel Hamilton, Ph.D. Candidate DGP Student Assisted Mentoring Program (STAMP) 2019-date AWARDS NSF-Simons Center for Quantitative Biology Training Grant Northwestern University 2018 Boit Prize for Engineering Writing Massachusetts Institute of Technology 2017

# Professional Memberships and Leadership Roles

SERVICE -

Driskill Graduate Program In the Life Sciences, Student Council Rep

NSF-Simons Center for Quantitive Biology Leadership Team, Graduate Student Rep

2020–date
2019–date

Senior Thesis - Regenerative Medicine and Synthetic Biology: Working Toward the Clinic

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