

A network science for complexity & society

Brennan Klein

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01 June, 2023

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France Regional Conference on Complex Systems



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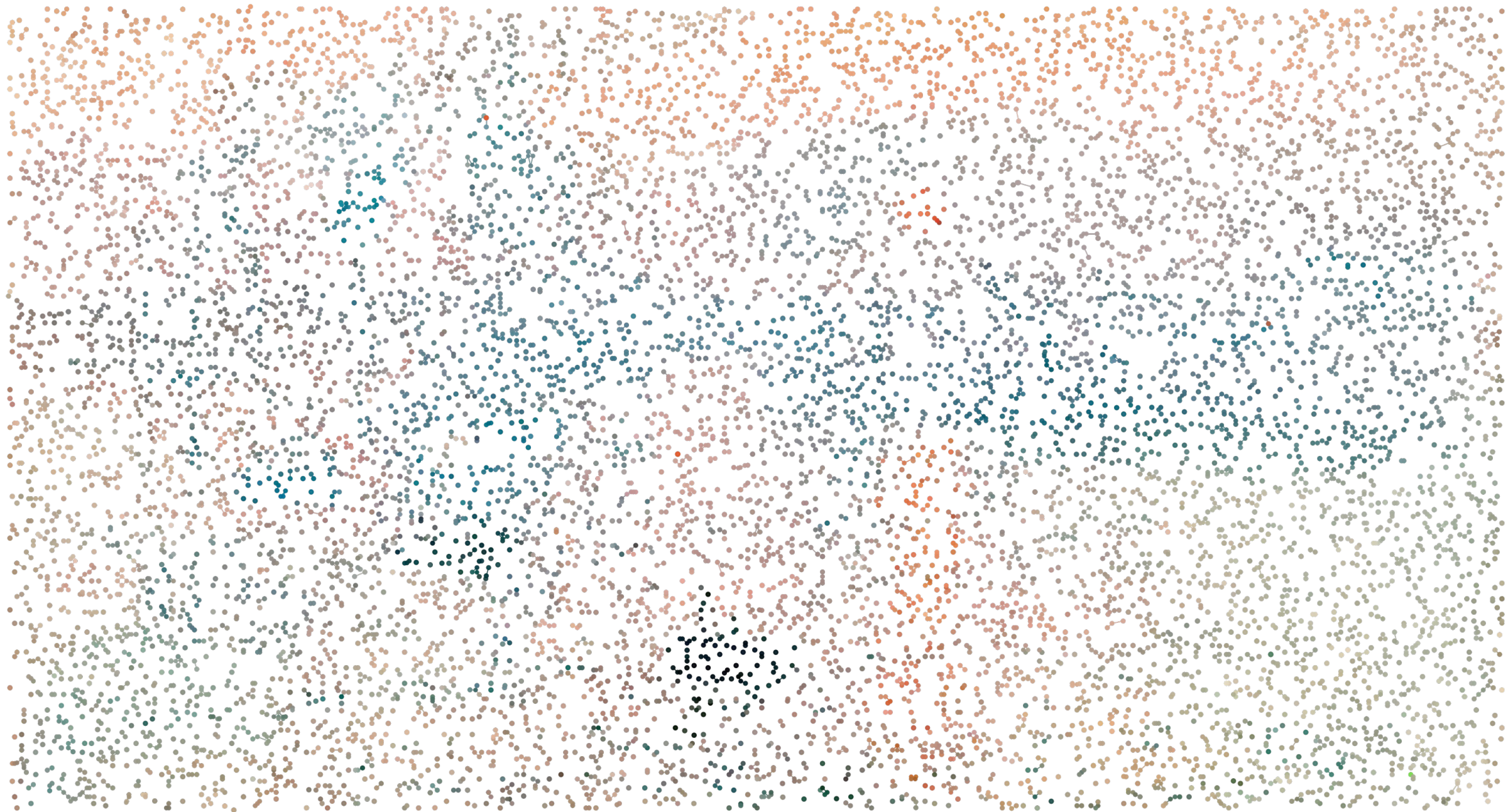


Complexity & Society Lab
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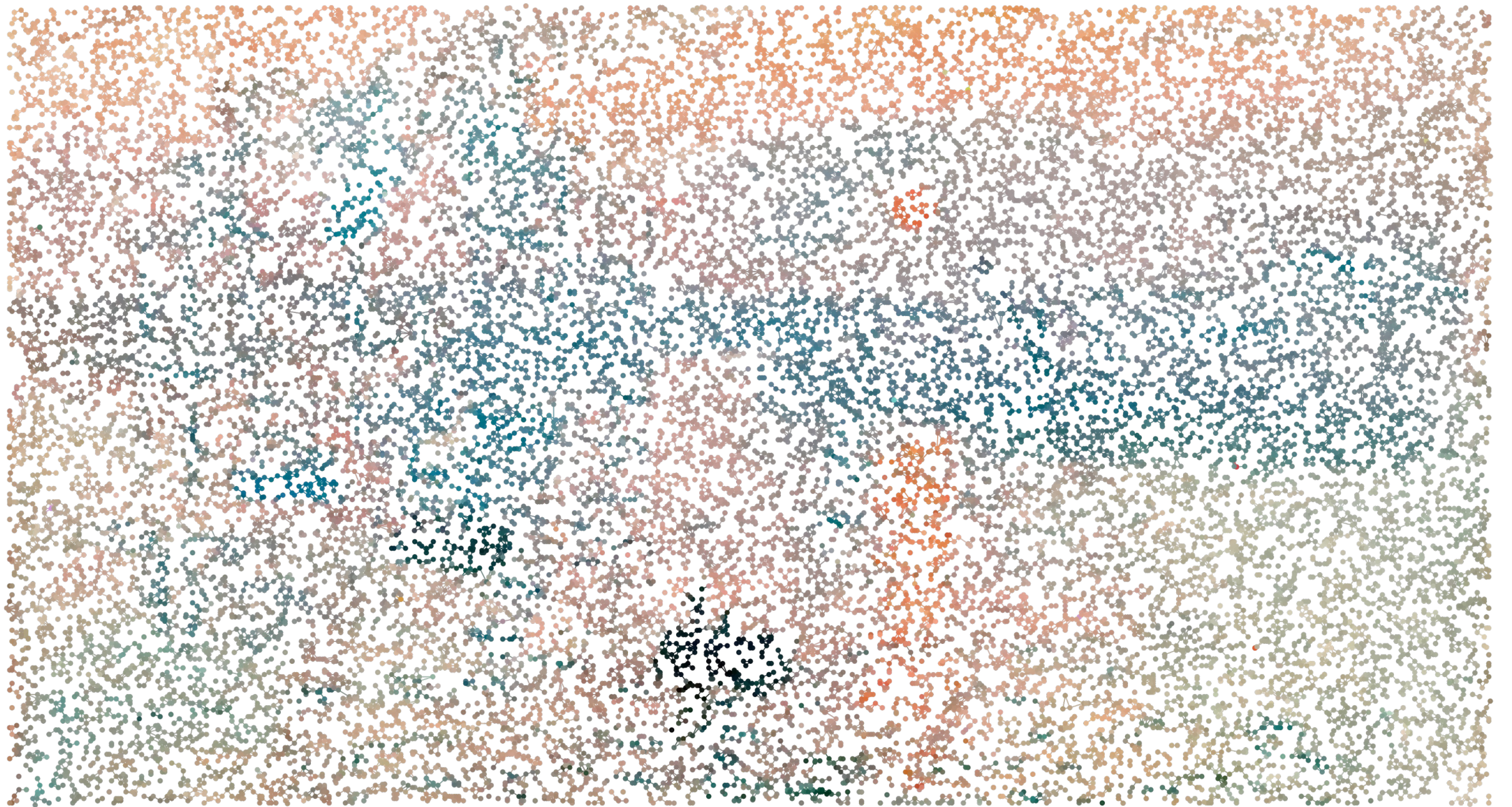


VERSES

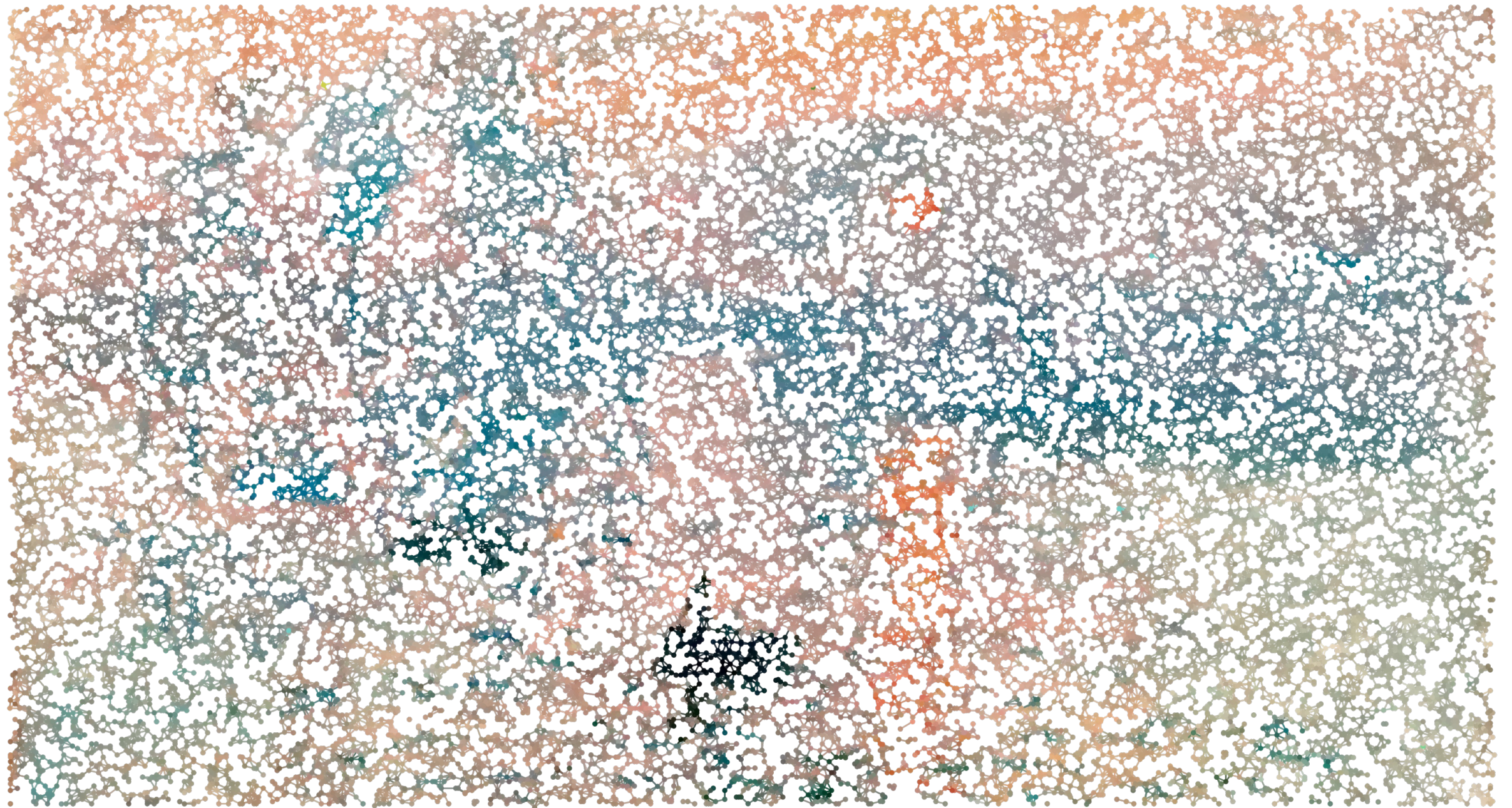




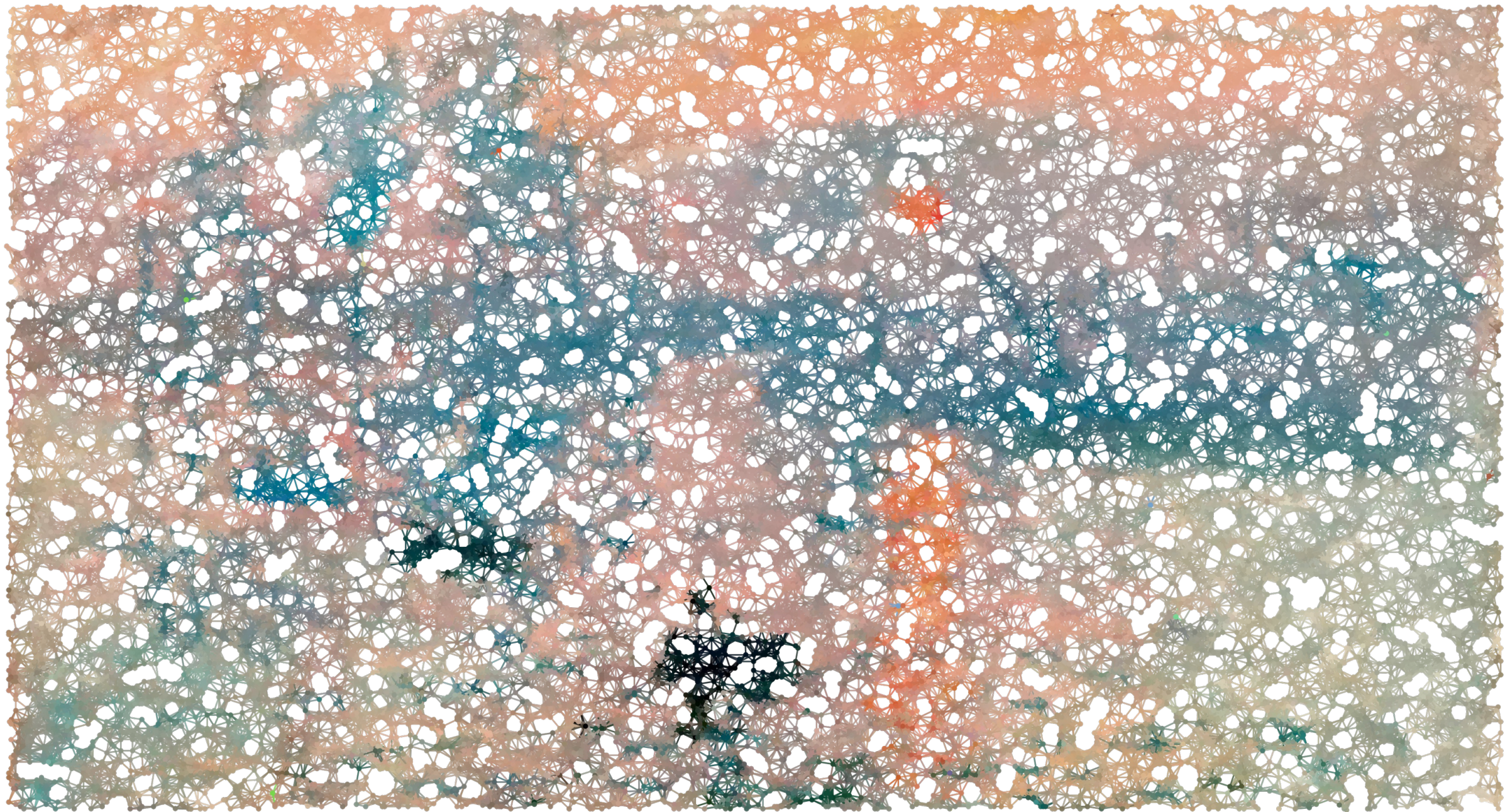
Network art based on "Impression, soleil levant" - Monet



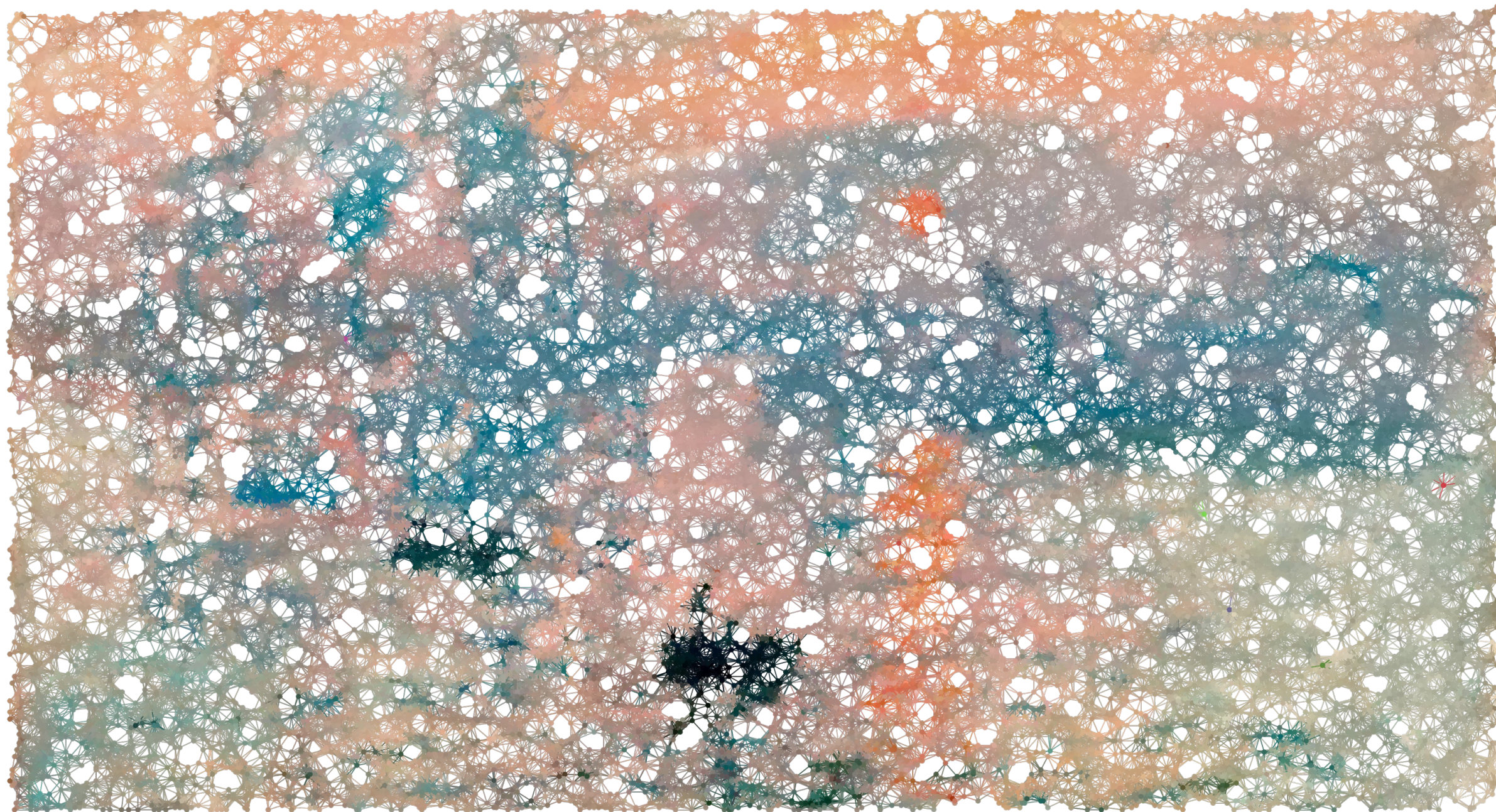
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VERSES

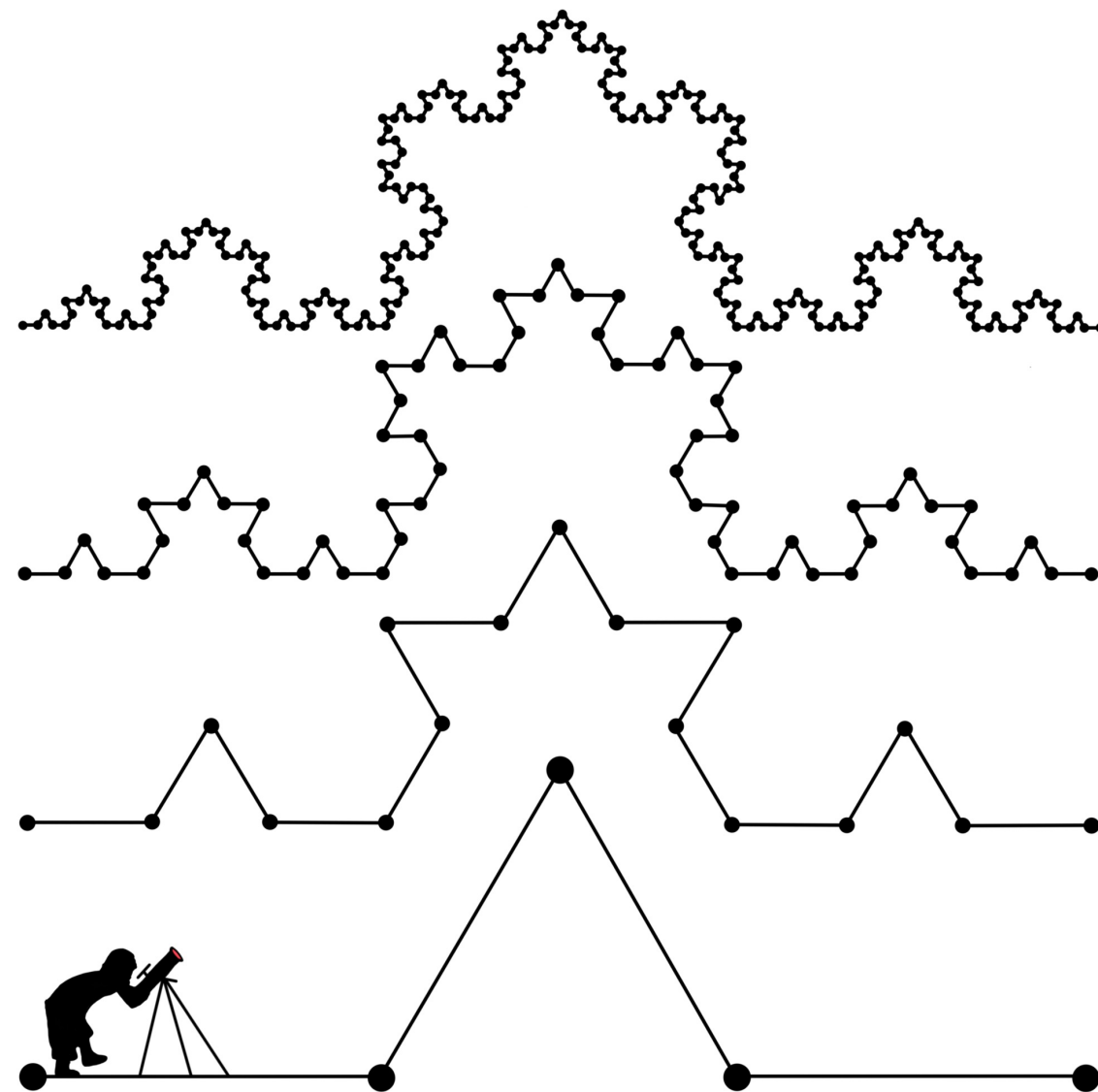


Thank you

FRCCS Organizing Committee
FRCCS Award Committee

Prof. Sam Scarpino
Prof. Alessandro Vespignani

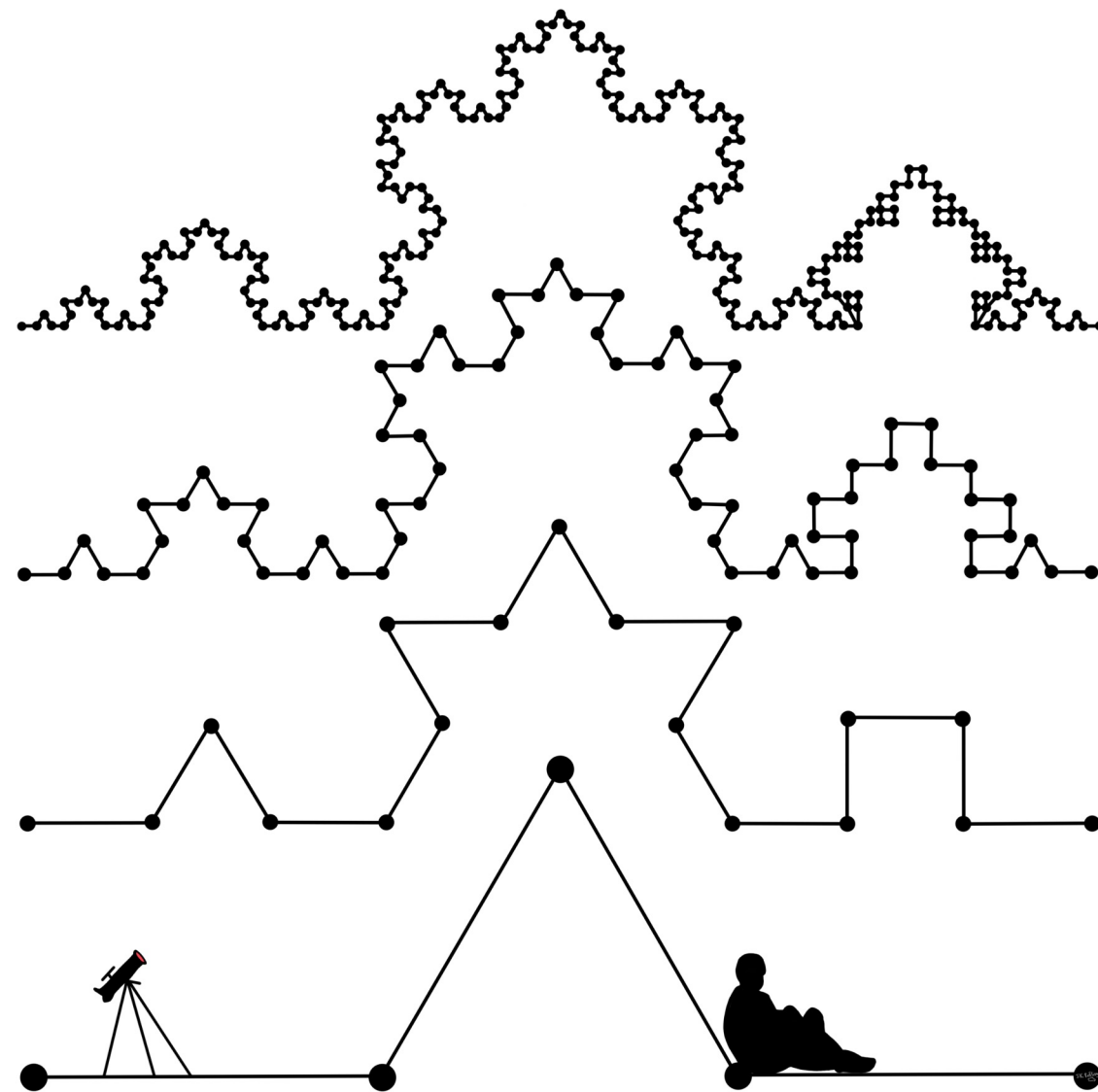
Many collaborators
You all!



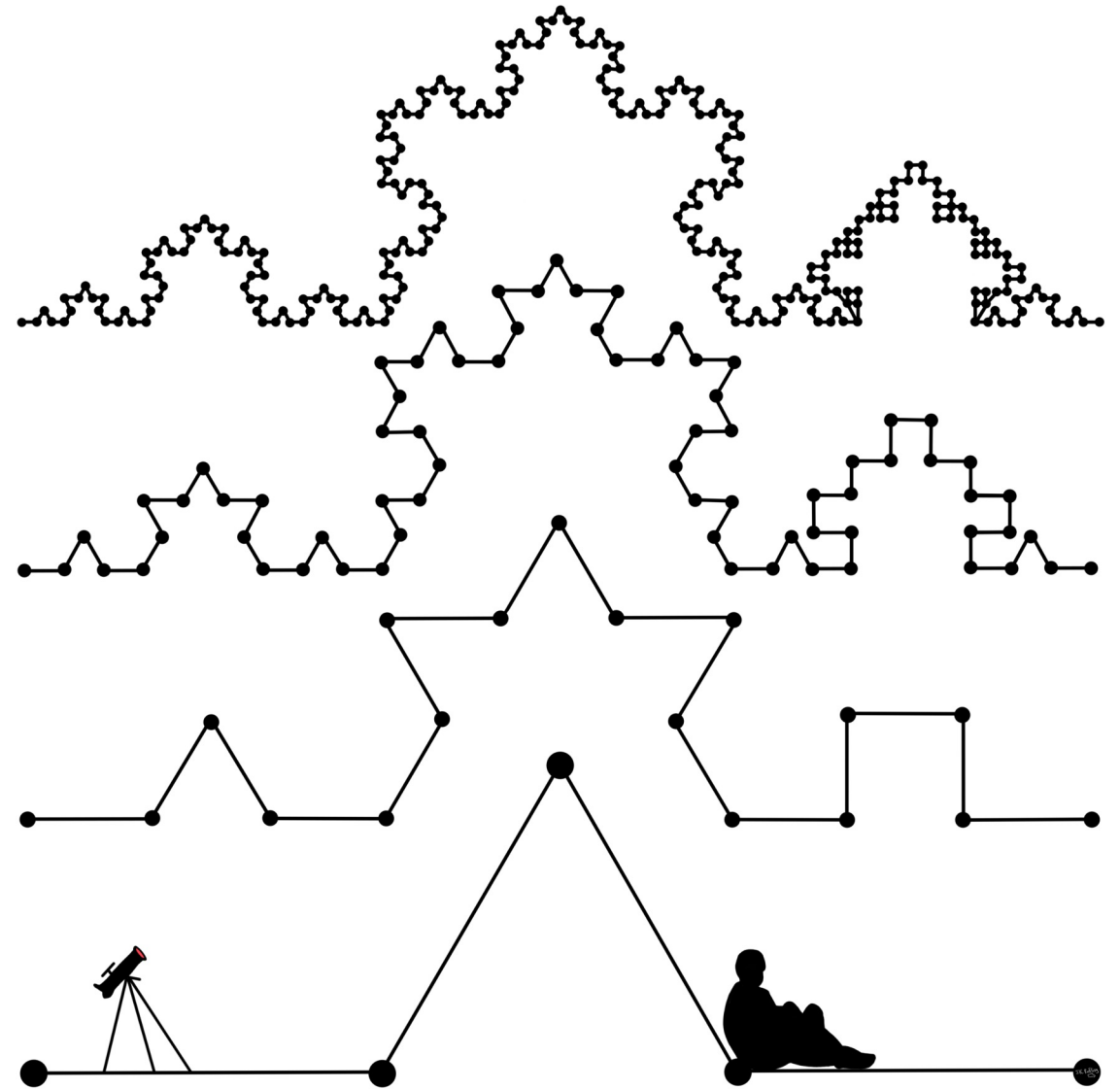
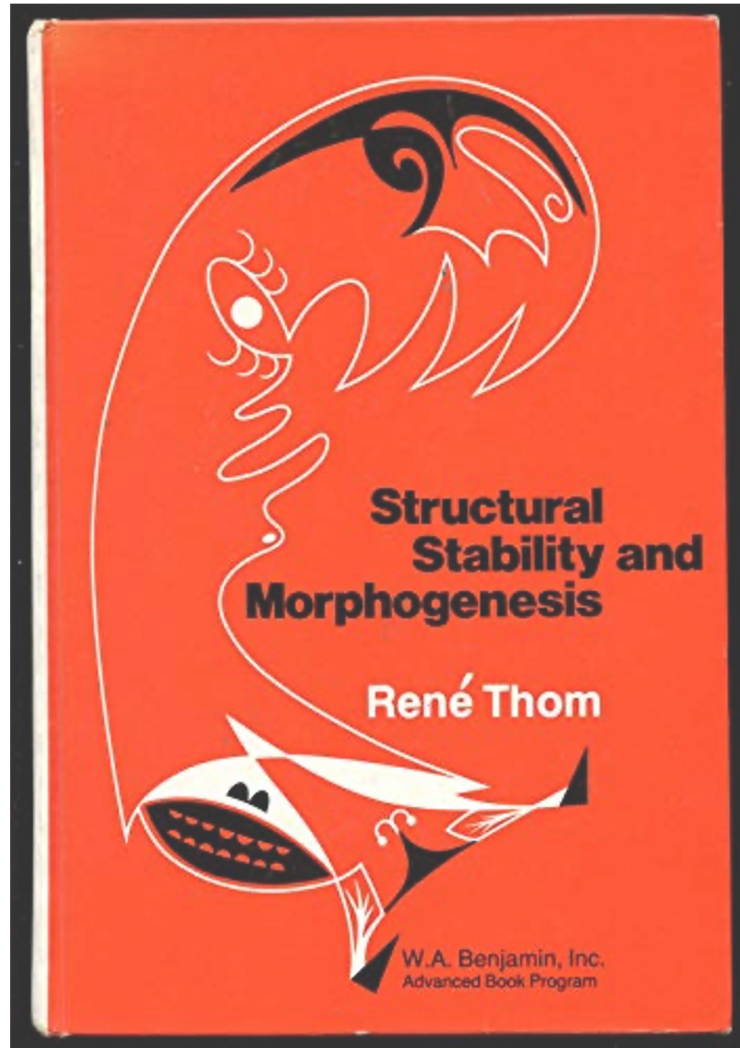
René Thom

“If one must choose between rigor and meaning, I shall unhesitatingly choose the latter.”

— René Thom

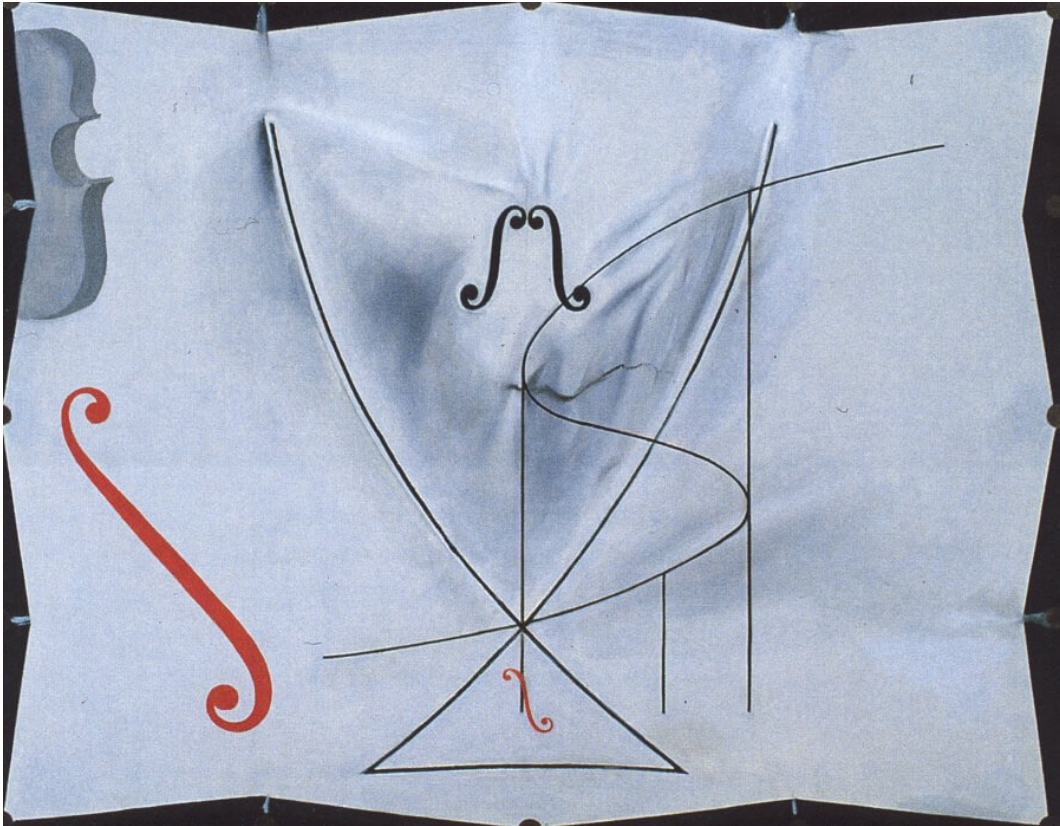


René Thom



René Thom

Inspired works of art from Salvador Dalí...



"The Swallow's Tail" from the Series on Catastrophes, 1983



"Topological Abduction of Europe: Homage to Rene Thom"

About me

- Background: Cognitive Science
(Swarthmore College: Philadelphia, USA)
- PhD: Network Science
(Northeastern University: Boston, USA)
- Current position: Postdoctoral Researcher
(Northeastern University: Boston, USA)
 - Also: Data for Justice Fellow, The Hutchins Center @Harvard
 - Also: Senior Research Scientist, VERSES Inc.



Prior work: Complexity

Research Article

WILEY Hindawi

The Emergence of Informative Higher Scales in Complex Networks

Brennan Klein^{1,2} and Erik Hoel³

communications biology

ARTICLE

<https://doi.org/10.1038/s42003-021-02867-8> OPEN

A computational exploration of resilience and evolvability of protein-protein interaction networks

Brennan Klein^{1,2,8}, Ludvig Holmér³, Keith M. Smith^{4,6}, Mackenzie M. Johnson⁵, Anshuman Swain⁶, Laura Stolp⁷, Ashley I. Teufel^{5,8,9} & April S. Kleppe^{10,11,8}

JOSS The Journal of Open Source Software

netrd: A library for network reconstruction and graph distances

Stefan McCabe¹, Leo Torres¹, Timothy LaRock¹, Syed Arefinul Haque¹, Chia-Hung Yang¹, Harrison Hartle¹, and Brennan Klein^{1, 2}

JOSS The Journal of Open Source Software

pymdp: A Python library for active inference in discrete state spaces

Conor Heins^{1,2,3,4}, Beren Millidge^{4,5}, Daphne Demekas⁶, Brennan Klein^{4,7,8}, Karl Friston⁹, Iain D. Couzin^{1,2,3}, and Alexander Tschantz^{1,10,11}

Received: 15 April 2021 | Accepted: 3 January 2022
DOI: 10.1111/2041-210X.13805

APPLICATION

Exploring noise, degeneracy and determinism in biological networks with the einet package

Brennan Klein^{1,2} | Anshuman Swain³ | Travis Byrum³ | Samuel V. Scarpino^{1,4,5,6} | William F. Fagan⁹

Evolution and emergence: higher order information structure in protein interactomes across the tree of life

Brennan Klein^{1,2,4}, Erik Hoel⁴, Anshuman Swain⁴, Ross Griesenow⁵ and Michael Levin³

¹Network Science Institute, Northeastern University, Boston, MA, USA
²Laboratory for the Modeling of Biological and Socio-Technical Systems, Northeastern University, Boston, MA, USA
³Allen Discovery Center, Tufts University, Medford, MA, USA
⁴Department of Biology, University of Maryland, College Park, MD, USA
⁵Department of Computer Science, Drexel University, Philadelphia, PA, USA
⁶Corresponding author. E-mail: b.klein@northeastern.edu

PROCEEDINGS A

royalsocietypublishing.org/journal/rspa

Network comparison and the within-ensemble graph distance

Harrison Hartle¹, Brennan Klein^{1,2}, Stefan McCabe¹, Alexander Daniels³, Guillaume St-Onge^{4,5}, Charles Murphy^{4,5} and Laurent Hébert-Dufresne^{3,4,6}

Research

Cite this article: Hartle H, Klein B, McCabe S, Daniels A, St-Onge G, Murphy C,

Experimental Economics (2021) 24:772–799
<https://doi.org/10.1007/s10683-020-09680-w>

ORIGINAL PAPER

Optimal design of experiments to identify latent behavioral types

Stefano Baldetti^{1,2} · Brennan Klein³ · Christoph Riedl^{3,4,5,6}

International Workshop on Active Inference

↳ IWAI 2022: **Active Inference** pp 75–98 | [Cite as](#)

Home > [Active Inference](#) > Conference paper

Spin Glass Systems as Collective Active Inference

Conor Heins¹, Brennan Klein, Daphne Demekas, Miguel Aguilera & Christopher L. Buckley

Conference paper | [First Online: 22 March 2023](#)

INTERFACE FOCUS

On Bayesian mechanics: a physics of and by beliefs

royalsocietypublishing.org/journal/rfs

Maxwell J. D. Ramstead^{1,2}, Dalton A. R. Sakhivadivel^{1,3,4,5,7}, Conor Heins^{1,6,7,8}, Magnus Koudahl⁹, Beren Millidge^{1,10}, Lancelot Da Costa¹¹, Brennan Klein^{1,12} and Karl J. Friston^{1,2}

under review (*Journal of Consciousness Studies*)

The inner screen model of consciousness: applying the free energy principle directly to the study of conscious experience

Maxwell J. D. Ramstead^{1,2,A}, Mahault Albarracín^{1,3,A}, Alex Kiefer^{1,4}, Brennan Klein^{1,5}, Chris Fields⁶, Karl Friston^{1,2}, and Adam Safran^{7,8}

under review (*Collective Intelligence*)

Designing Ecosystems of Intelligence from First Principles

Karl J. Friston^{1,2}, Maxwell J.D. Ramstead^{1,2}, Alex B. Kiefer^{1,3}, Alexander Tschantz¹, Christopher L. Buckley^{1,4}, Mahault Albarracín^{1,5}, Riddhi J. Pitliya^{1,6}, Conor Heins^{1,7,8,9}, Brennan Klein^{1,10}, Beren Millidge^{1,11}, Dalton A.R. Sakhivadivel^{1,12,13,14}, Toby St. Clere Smithe^{1,6,15}, Magnus Koudahl^{1,16}, Safae Essafi Tremblay^{1,17}, Capm Petersen¹, Kaiser Fung¹, Jason G. Fox¹, Steven Swanson¹, Dan Mapes¹, and Gabriel René¹

Prior work: Society

RESEARCH

CORONAVIRUS

The effect of human mobility and control measures on the COVID-19 epidemic in China

Moritz U. G. Kraemer^{1,2,3*}, Chia-Hung Yang⁴, Bernardo Gutierrez^{1,5}, Chieh-Hsi Wu⁶, Brennan Klein⁴, David M. Pigott⁷, Open COVID-19 Data Working Group[†], Louis du Plessis¹, Nuno R. Faria¹, Ruoran Li⁸, William P. Hanage⁹, John S. Brownstein^{2,3}, Maylis Layan^{3,10}, Alessandro Vespignani^{4,11}, Huaiyu Tian¹², Christopher Dye¹, Oliver G. Pybus^{1,13*}, Samuel V. Scarpino^{4*}

RESEARCH

CORONAVIRUS

Spatiotemporal invasion dynamics of SARS-CoV-2 lineage B.1.1.7 emergence

Moritz U. G. Kraemer^{1*}, Verity Hill^{2†}, Christopher Ruis^{3†}, Simon Dellicour^{4,5†}, Sumali Bajaj^{1†}, John T. McCrone², Guy Baele⁵, Kris V. Parag⁶, Anya Lindström Battle⁷, Bernardo Gutierrez¹, Ben Jackson², Rachel Colquhoun², Aine O'Toole², Brennan Klein⁸, Alessandro Vespignani⁸, COVID-19 Genomics UK (COG-UK) Consortium[†], Erik Volz⁶, Nuno R. Faria^{1,6,9}, David M. Aanensen^{10,11}, Nicholas J. Loman¹², Louis du Plessis¹, Simon Cauchemez¹³, Andrew Rambaut^{2*}, Samuel V. Scarpino^{8,14,15*}, Oliver G. Pybus^{1,16*}

ARTICLE

<https://doi.org/10.1038/s41467-021-22521-5> OPEN

The effect of eviction moratoria on the transmission of SARS-CoV-2

Anjalika Nande^{1,7}, Justin Sheen^{2,7}, Emma L. Walters³, Brennan Klein^{4,5}, Matteo Chinazzi^{4,5}, Andrei H. Gheorghe¹, Ben Adlam¹, Julianna Shinnick², Maria Florencia Tejada², Samuel V. Scarpino⁴, Alessandro Vespignani^{4,5}, Andrew J. Greenlee³, Daniel Schneider³, Michael Z. Levy^{2,8,9} & Alison L. Hill^{1,6,8,9*}

communications medicine

nature > communications medicine > articles > article

Article | [Open Access](#) | Published: 14 February 2023

Forecasting hospital-level COVID-19 admissions using real-time mobility data

Brennan Klein , Ana C. Zenteno , Daisha Joseph , Mohammadmehdi Zahedi , Michael Hu , Martin S. Copenhaver , Moritz U. G. Kraemer , Matteo Chinazzi , Michael Klompas , Alessandro Vespignani , Samuel V. Scarpino  & Hoijat Salmasian 

Clinical Infectious Diseases

CORRECTED PROOF

Examining the Robustness of 3 Versus 6 Feet of Physical Distancing in Schools: A Reanalysis of van den Berg et al

Brennan Klein , Daniel A Harris 

PLOS DIGITAL HEALTH

RESEARCH ARTICLE

Higher education responses to COVID-19 in the United States: Evidence for the impacts of university policy

Brennan Klein , Nicholas Genorou , Matteo Chinazzi , Zarana Bhadriraja , Rishab Gunashekar , Preeti Kori , Bodian Li , Stefan McCabe , Jon Green , David Lazer , Christopher R. Marsicano , Samuel V. Scarpino , Alessandro Vespignani 

nature

nature > articles > article

Article | [Open Access](#) | Published: 19 April 2023

COVID-19 amplified racial disparities in the US criminal legal system

Brennan Klein , C. Brandon Ogbunugafor , Benjamin J. Schafer , Zarana Bhadriraja , Preeti Kori , Jim Sheldon , Nitish Kaza , Arush Sharma , Emily A. Wang , Tina Eliassi-Rad , Samuel V. Scarpino  & Elizabeth Hinton 

under review (PNAS)

Spatial scales of COVID-19 transmission in Mexico

Brennan Klein^{1,2}, Harrison Hartle¹, Munik Shrestha¹, Ana Cecilia Zenteno³, David Barros Sierra Cordera⁴, José R. Nicolas-Carlock⁵, Ana I. Bento⁶, Benjamin M. Althouse^{7,8}, Bernardo Gutierrez^{9,10,11}, Marina Escalera-Zamudio^{9,11}, Arturo Reyes-Sandoval^{12,13}, Oliver G. Pybus^{9,14,18}, Alessandro Vespignani^{1,2}, Alberto Diaz Quiñonez^{*11,15}, Samuel V. Scarpino^{*11,16,17}, and Moritz U.G. Kraemer^{*9,18}

revision (PLOS Digital Health)

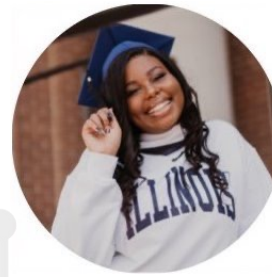
Characterizing collective physical distancing in the U.S. during the first nine months of the COVID-19 pandemic

Brennan Klein¹, Timothy LaRock¹, Stefan McCabe¹, Leo Torres¹, Lisa Friedland¹, Maciej Kos¹, Filippo Privitera¹, Brennan Lake¹, Moritz U.G. Kraemer², John S. Brownstein^{6,7}, Richard Gonzalez⁸, David Lazer¹, Tina Eliassi-Rad¹, Samuel V. Scarpino^{1,9,10}, Alessandro Vespignani^{1,3}, and Matteo Chinazzi^{1,2}

Complexity & Society Lab (the “And Lab”)



Harrison Hartle
PhD Candidate,
Network Science



Brein Mosely
PhD Student,
Education



Conor Heins
PhD Student,
Collective Behavior



Christina Steele
Research Assistant,
Data for Justice



Moritz Laber
PhD Student,
Network Science



Daphne Demekas
Research Assistant,
Artificial Intelligence



Oghenetega Ogodo
MS Student,
Urban Planning



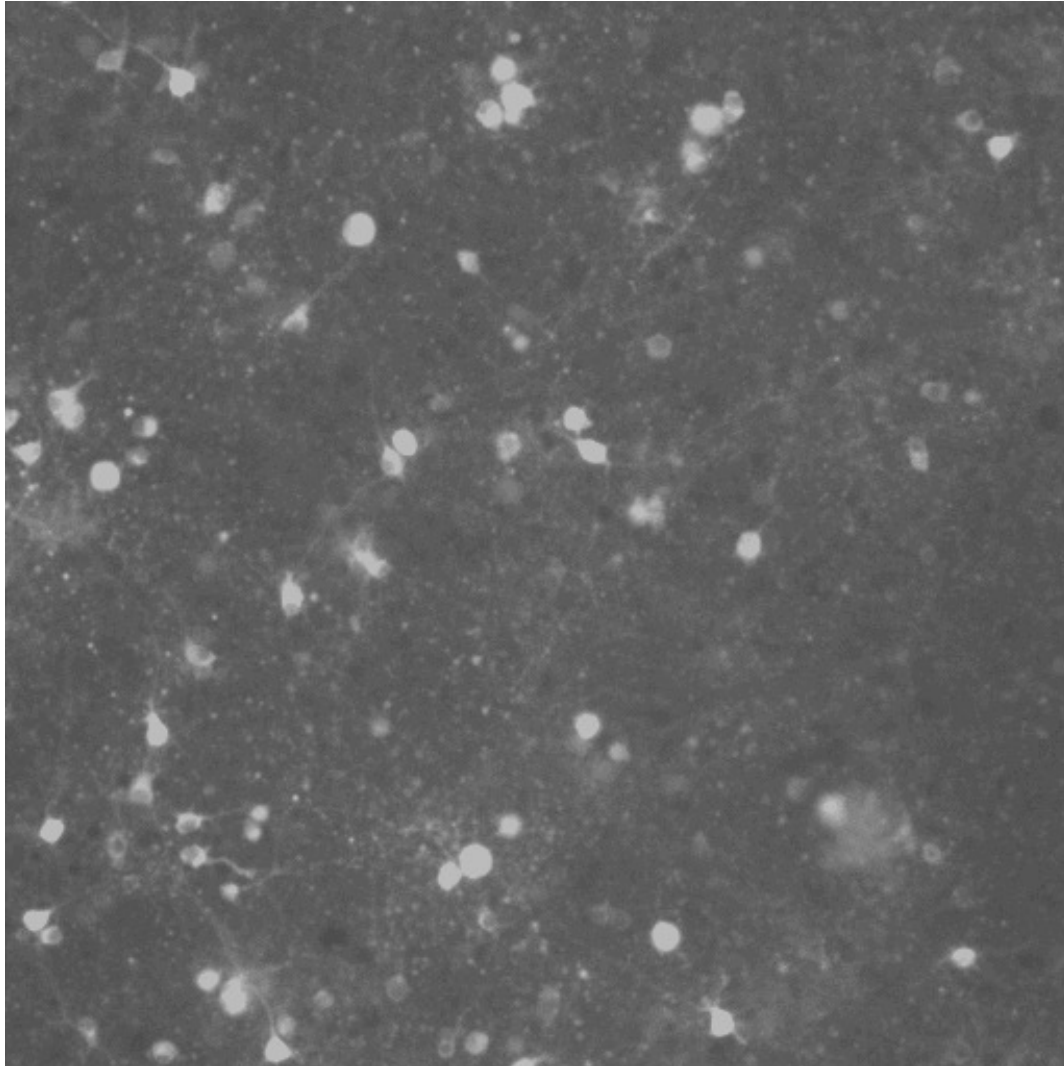
Arush Sharma
Research Assistant,
Data for Justice



Today

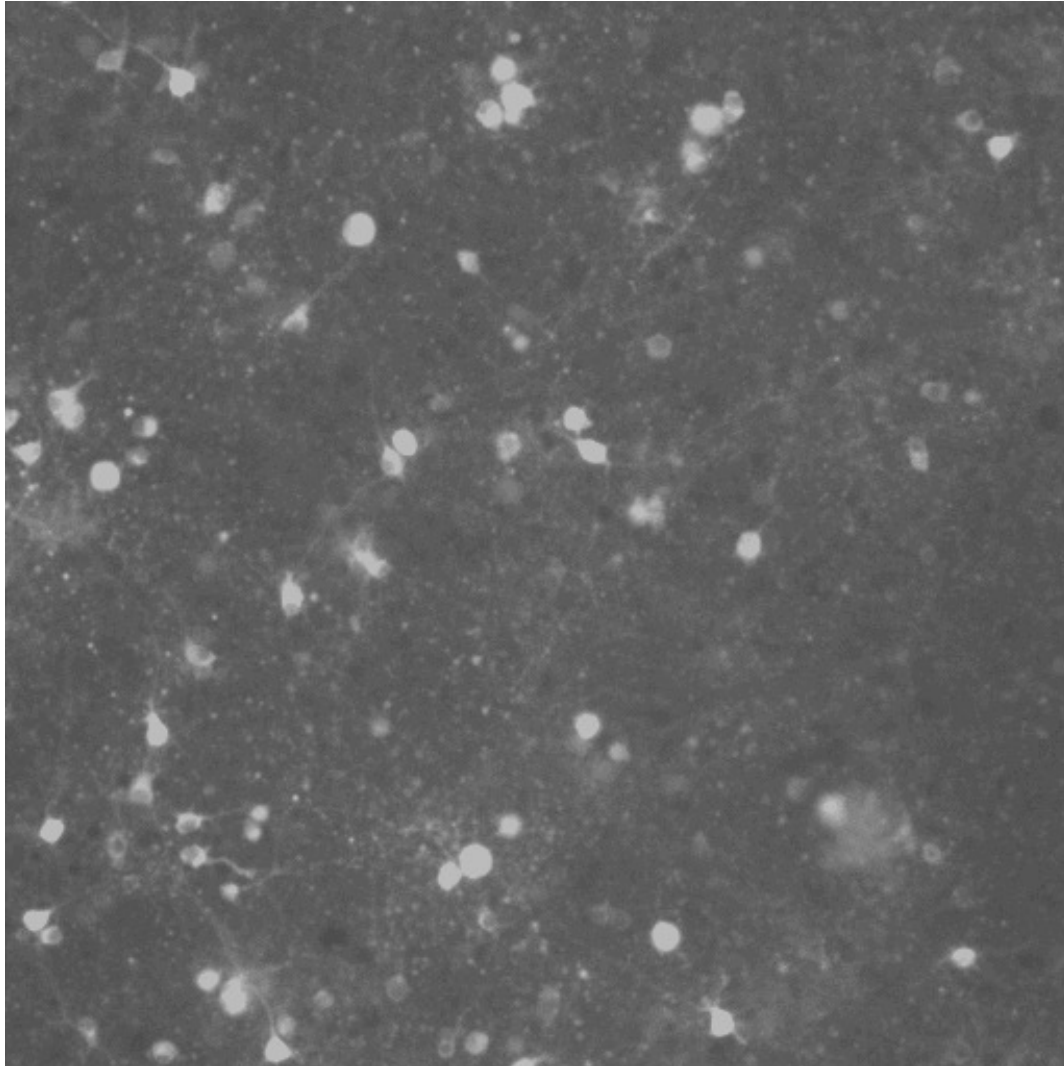
1. Introduction
2. Complexity
 - i. Representation & comparison*
 - ii. Informative scales in networks*
3. Society
 - i. Mobile device data for disease modeling*
 - ii. Assorted COVID-19 projects*
4. Research vision and outlook

Networks of thought



This is a recording of neurons from the hippocampus of a brain **suffering from Alzheimer's.**

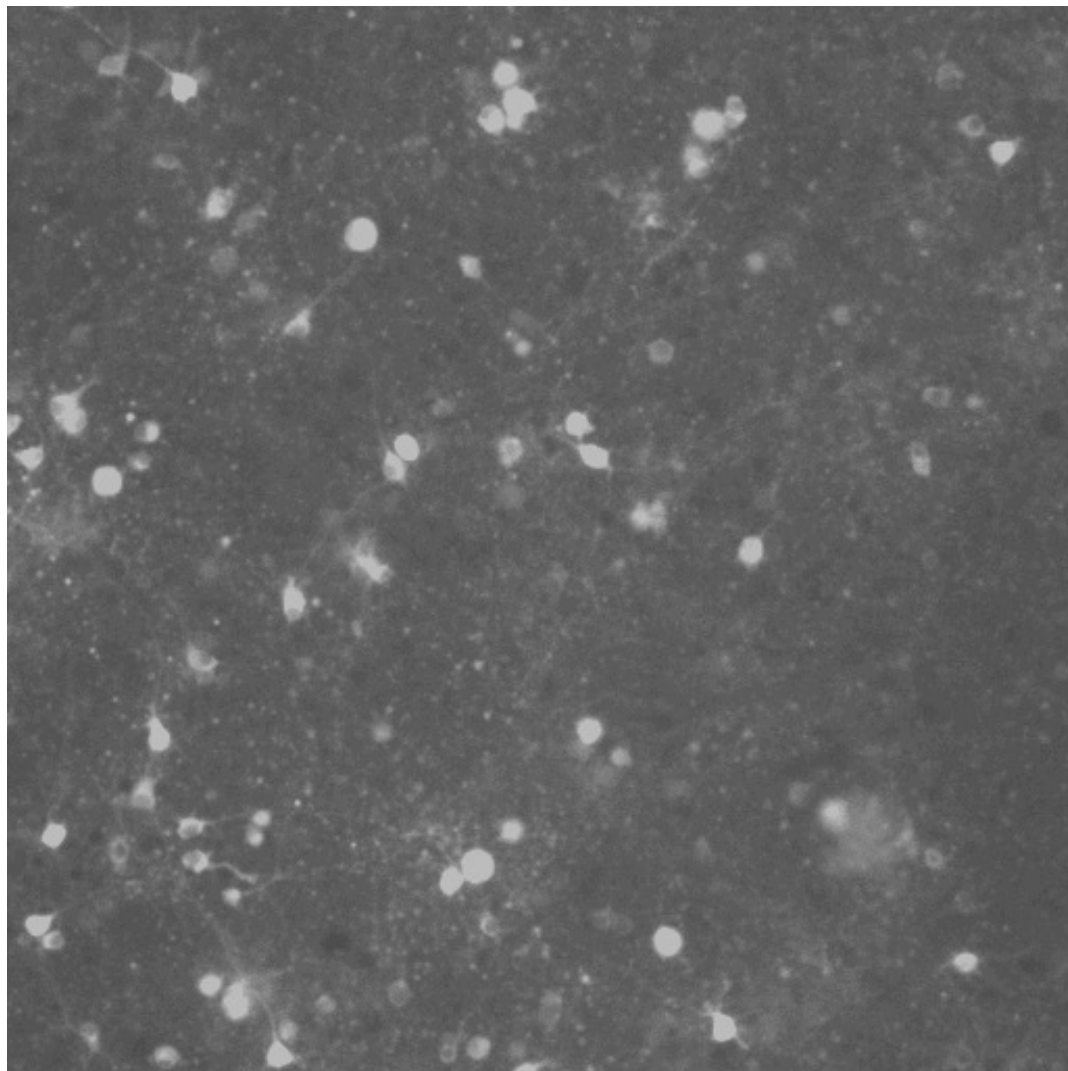
Networks of thought



This is a recording of neurons from the hippocampus of a brain suffering from Alzheimer's.

I want to cure this disease by studying **ecosystems**.

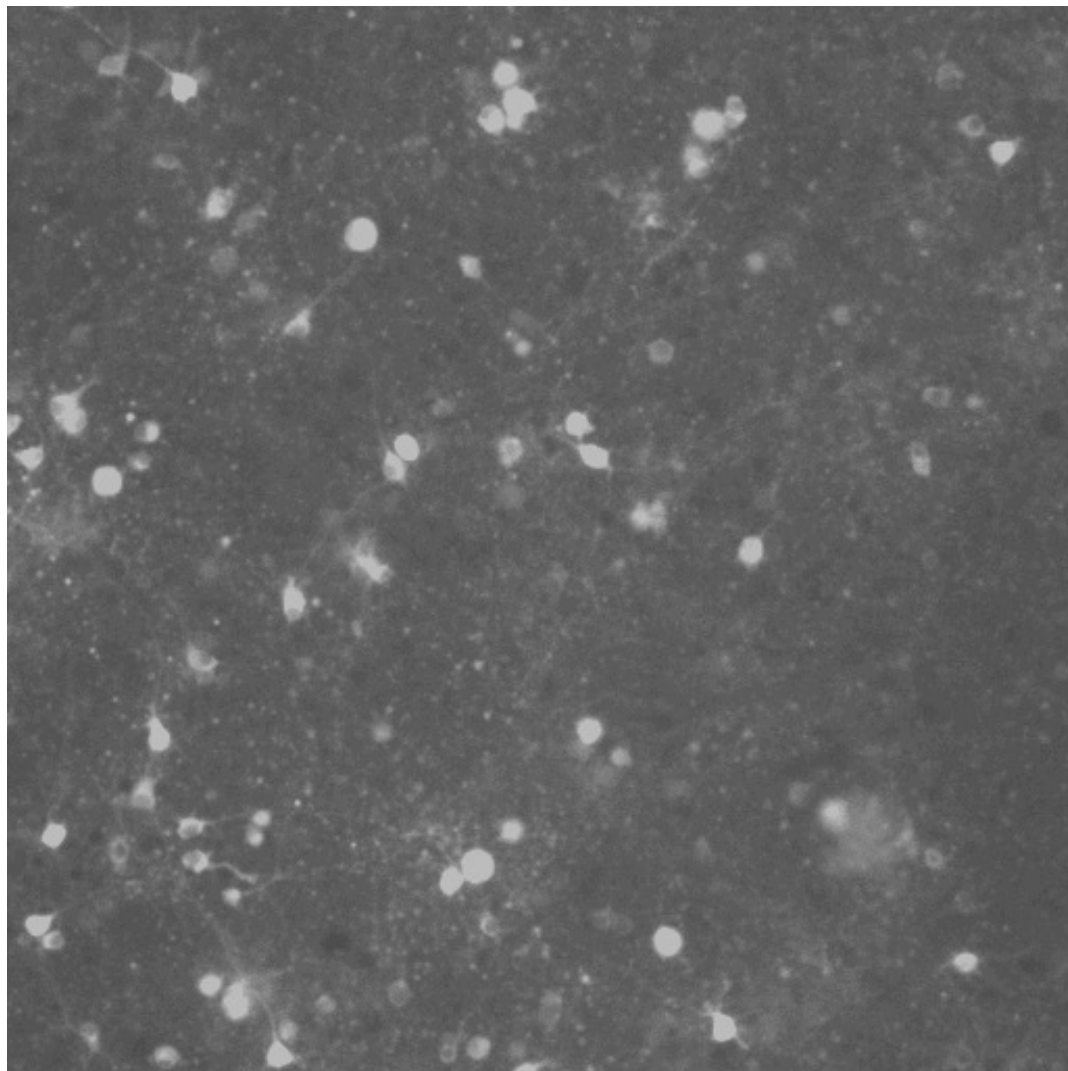
Networks of thought



This is a recording of neurons from the hippocampus of a brain suffering from Alzheimer's.

I want to cure this disease by studying **the economy**.

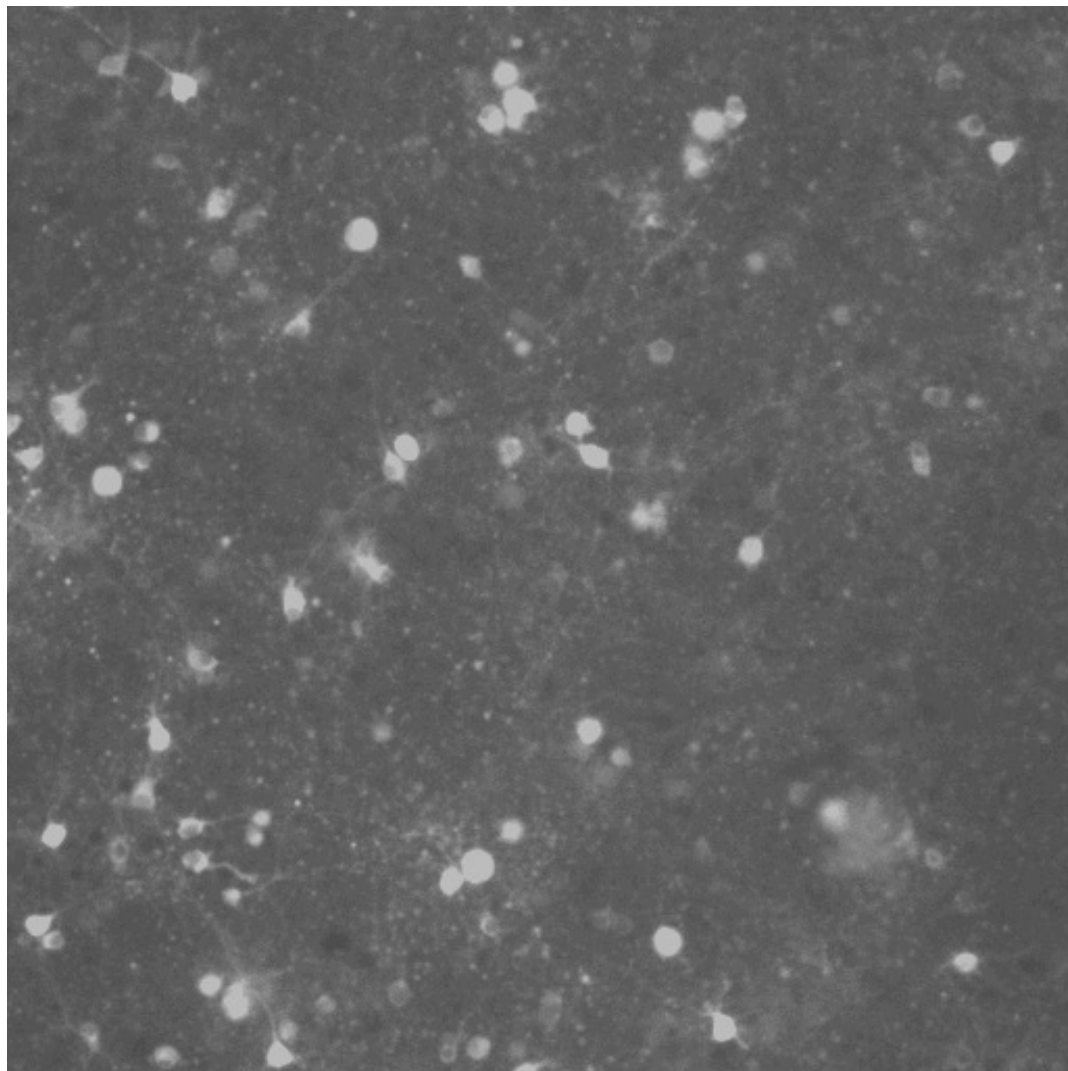
Networks of thought



This is a recording of neurons from the hippocampus of a brain suffering from Alzheimer's.

I want to cure this disease by studying **schools of fish**.

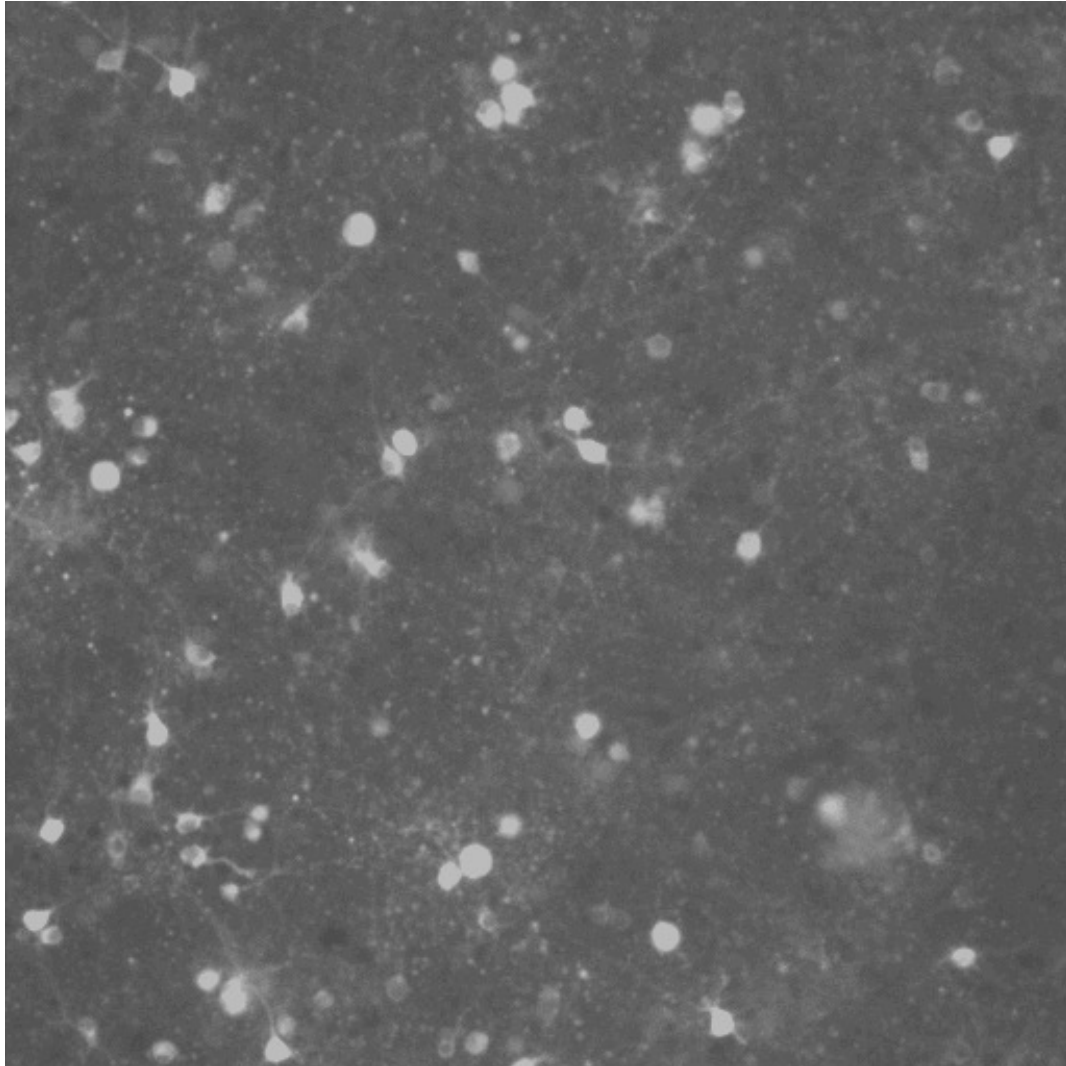
Networks of thought



This is a recording of neurons from the hippocampus of a brain suffering from Alzheimer's.

I want to cure this disease by studying **fractals**.

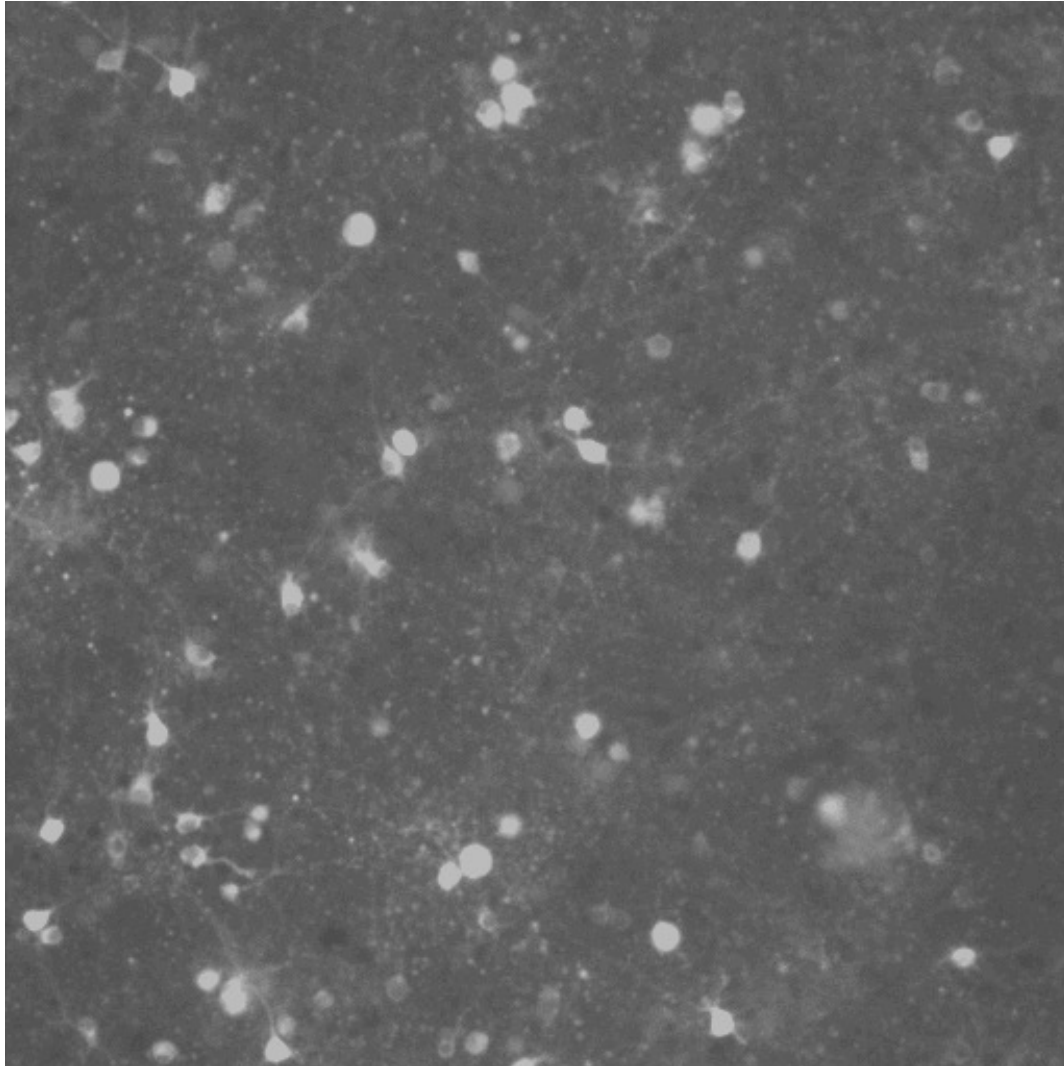
Networks of thought



This is a recording of neurons from the hippocampus of a brain suffering from Alzheimer's.

I want to cure this disease by studying **epidemics**.

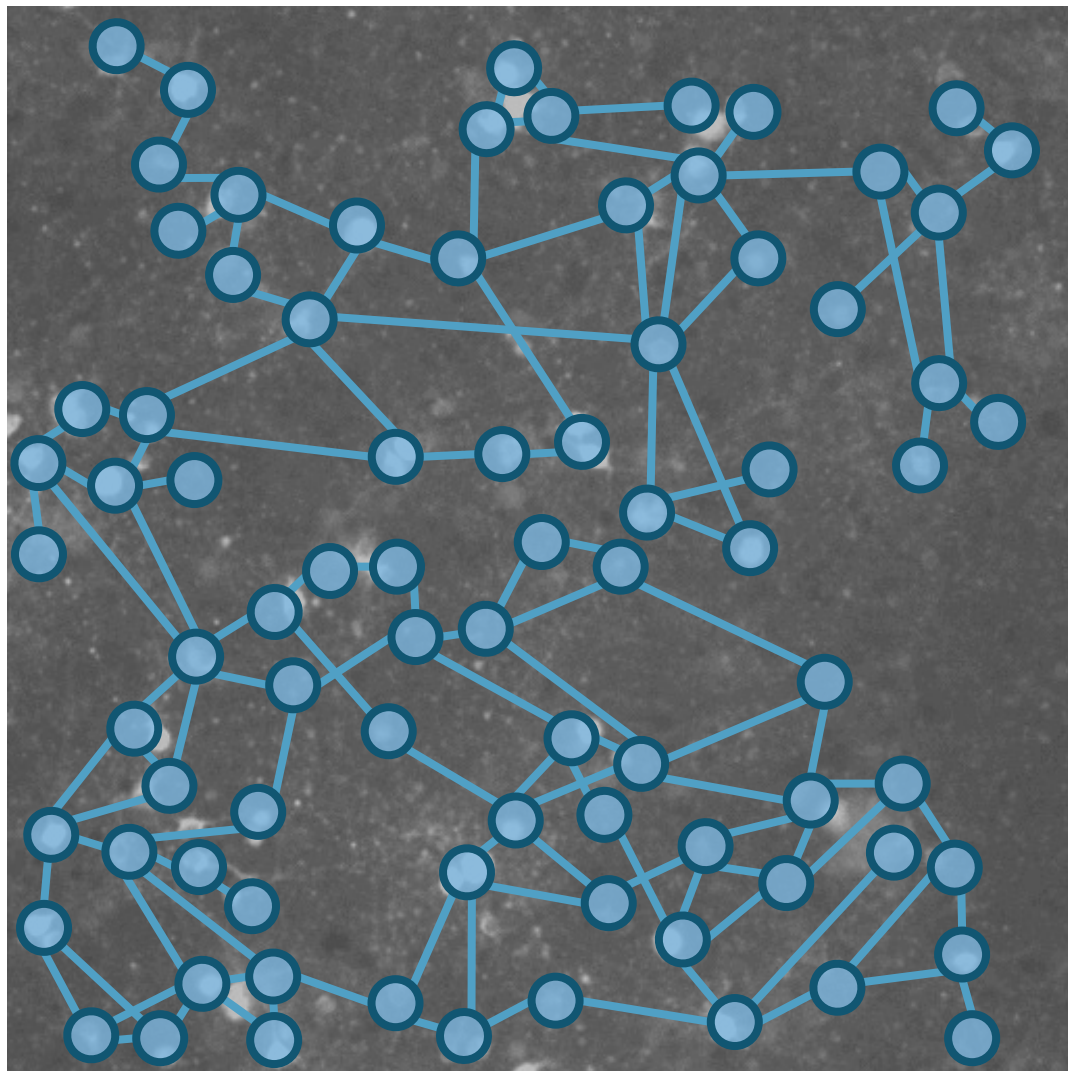
Networks of thought



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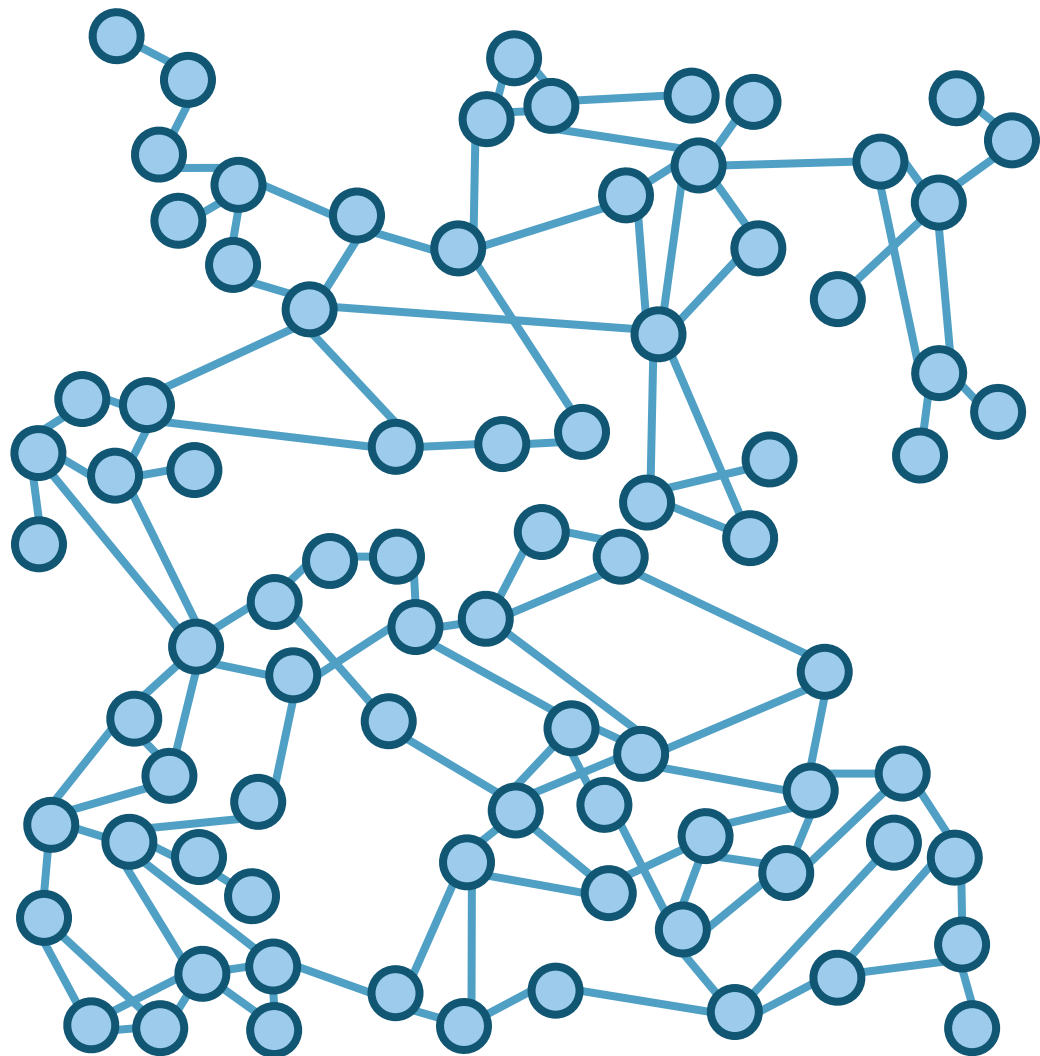
I want to cure this disease by studying **networks**.

The problem



We're good network scientists, so let's just draw some nodes ...and edges.

The problem

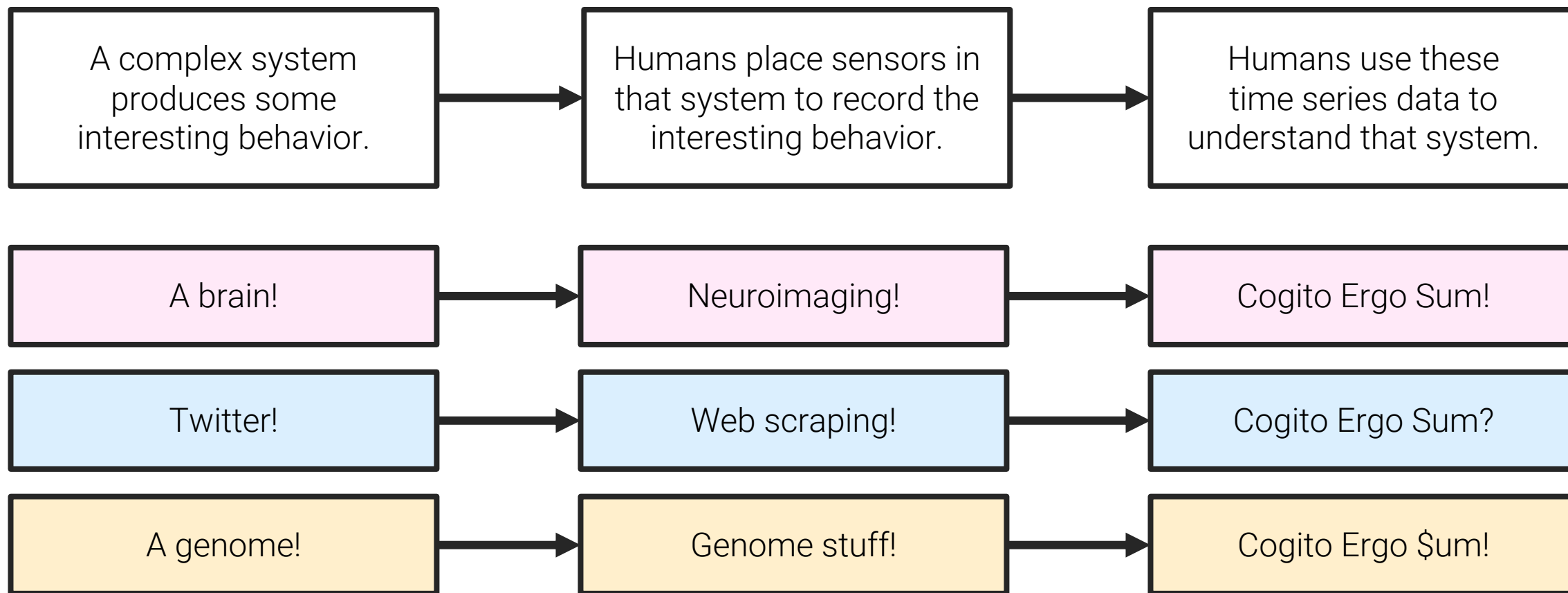


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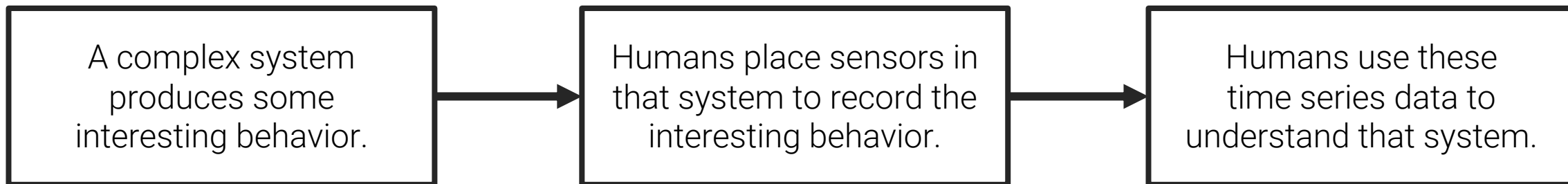
Easy, right?

Network (re)construction

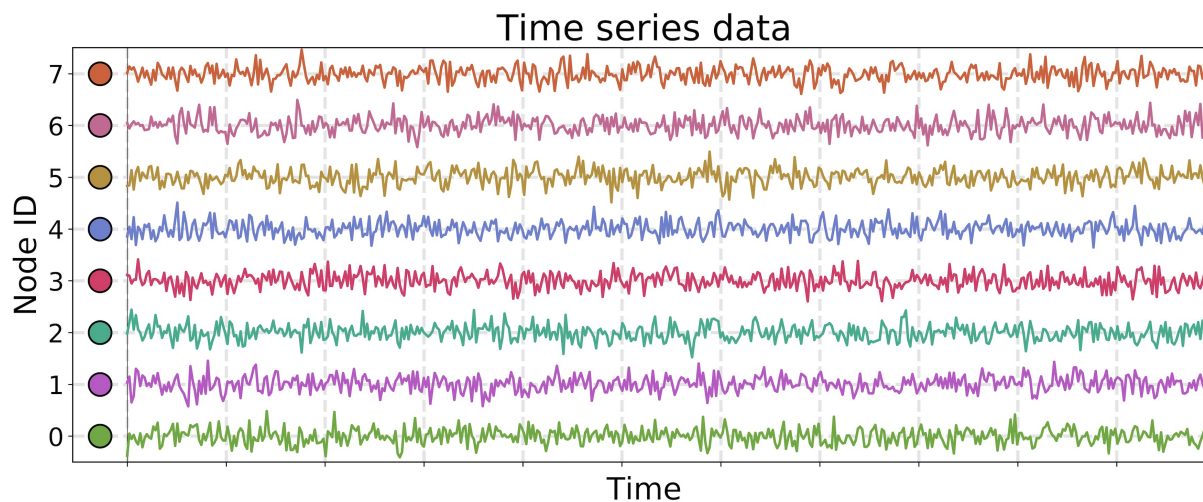
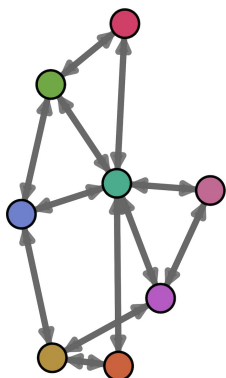
The problem



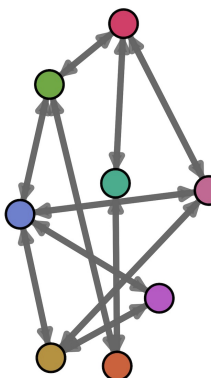
The problem



"True" network structure



Reconstructed network

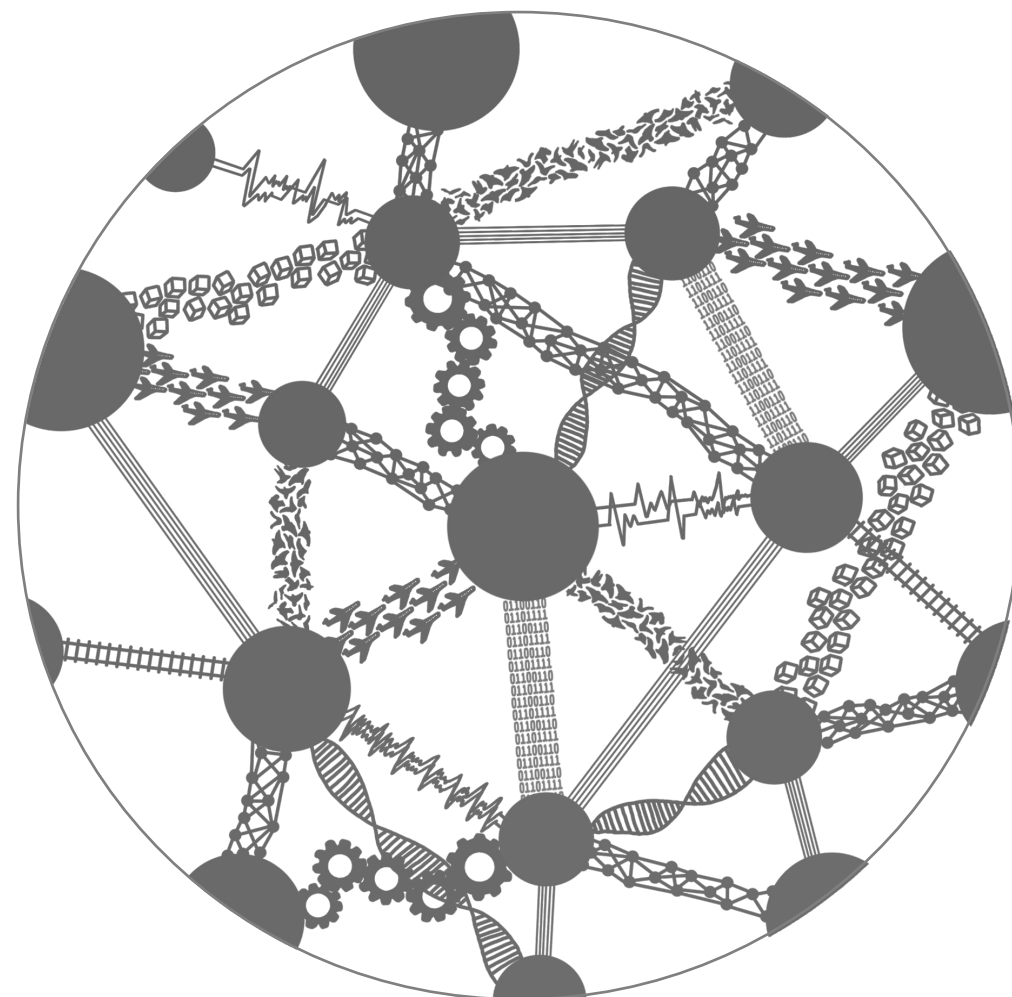


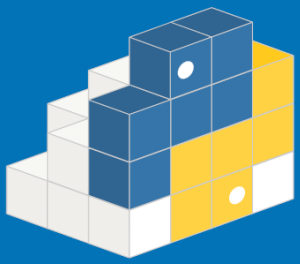
Networks as models, networks as maps

We want a **useful** network science.

A good model can help understand and predict the behavior of a system.

To make good models of systems, we need adequate toolkits, strong benchmarks, standardization, etc.



netrd 0.2.2

```
pip install netrd
```



PePy

netrd

Summary

PyPI link
<https://pypi.org/project/netrd>

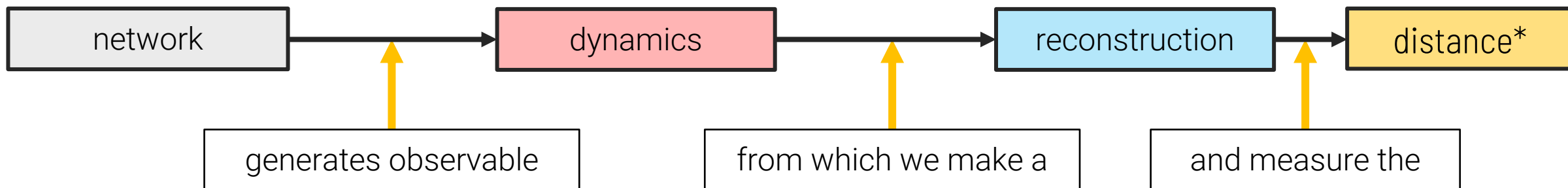
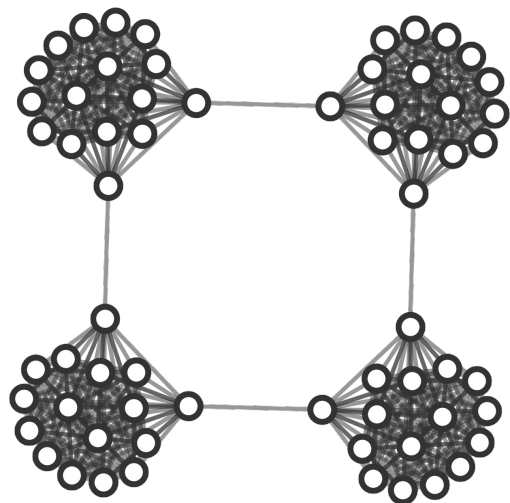
Total downloads
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Total downloads - 30 days
515

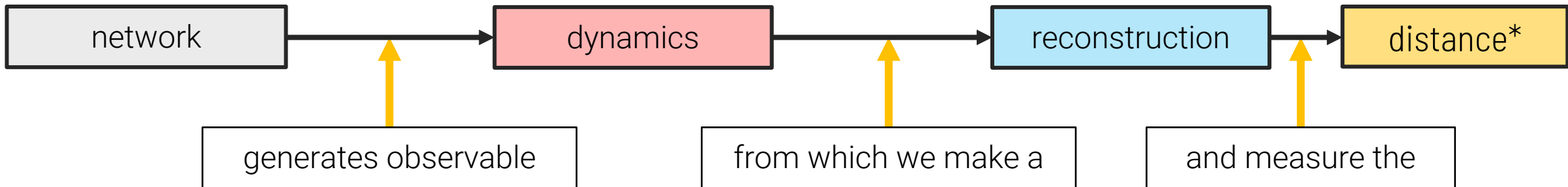
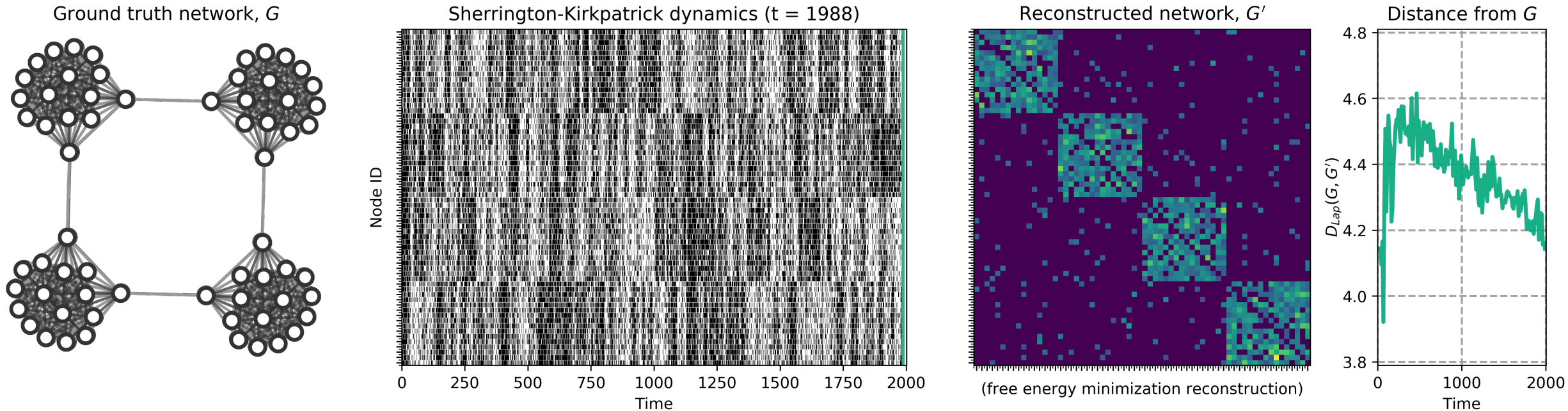
Total downloads - 7 days
78

netrd: A library for network {reconstruction, distances, dynamics}

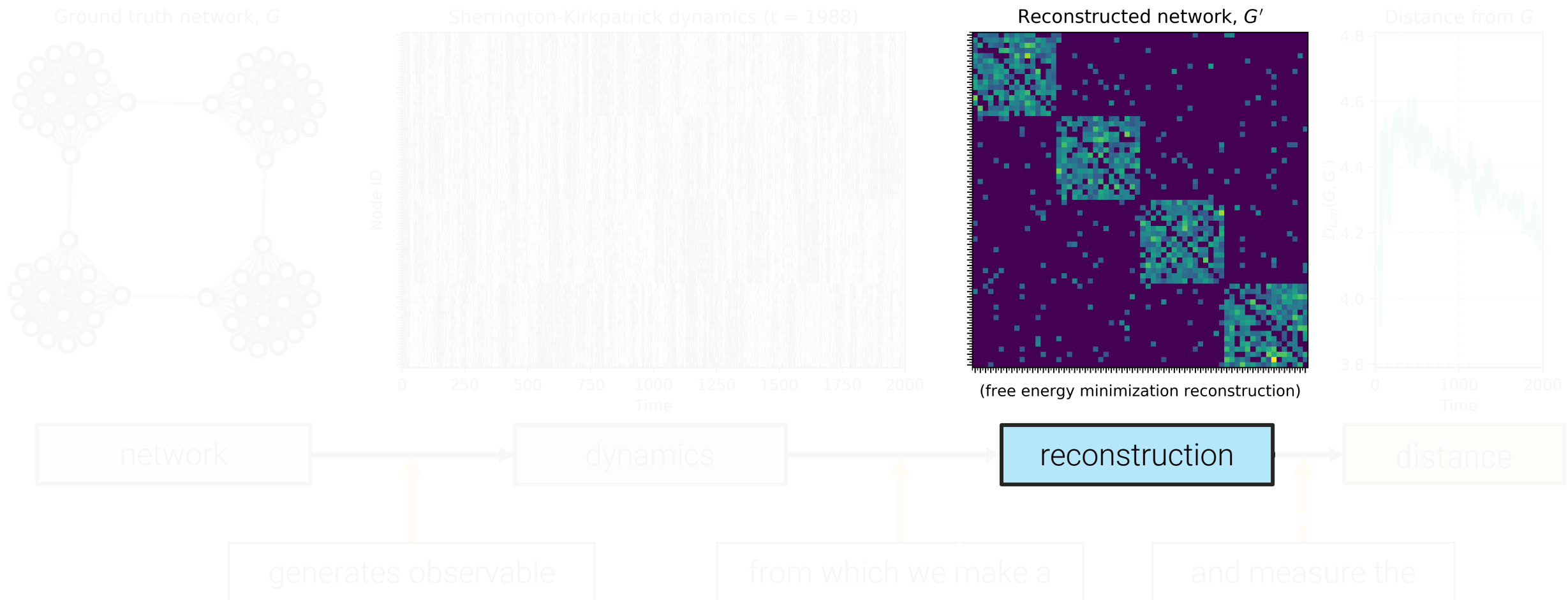
Basic pipeline



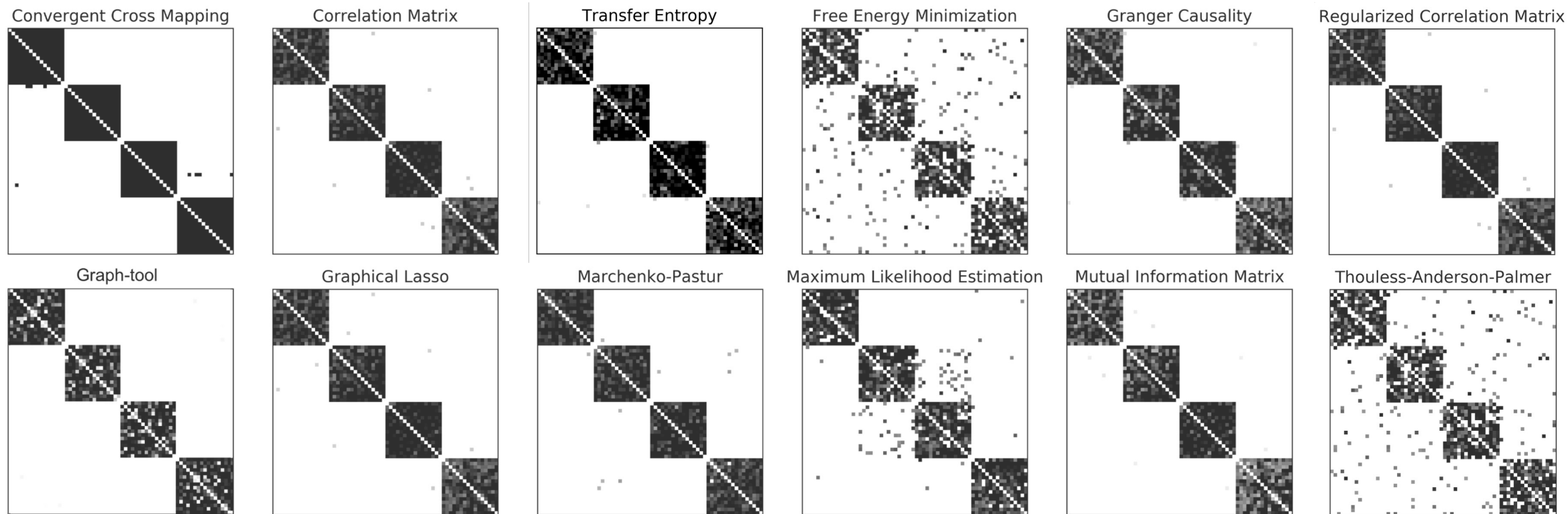
Basic pipeline



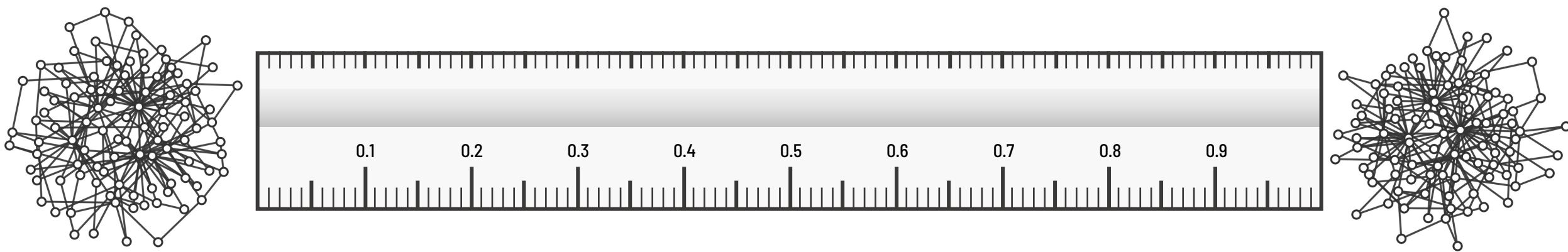
Basic pipeline



Systematic comparison of different tools

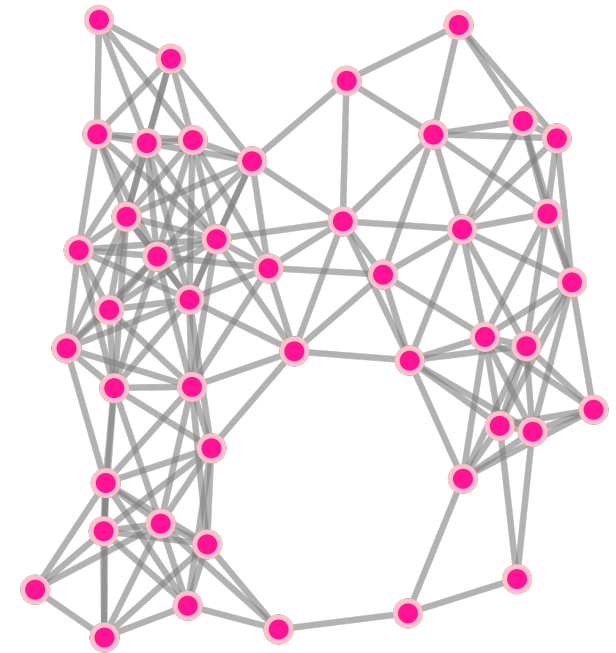
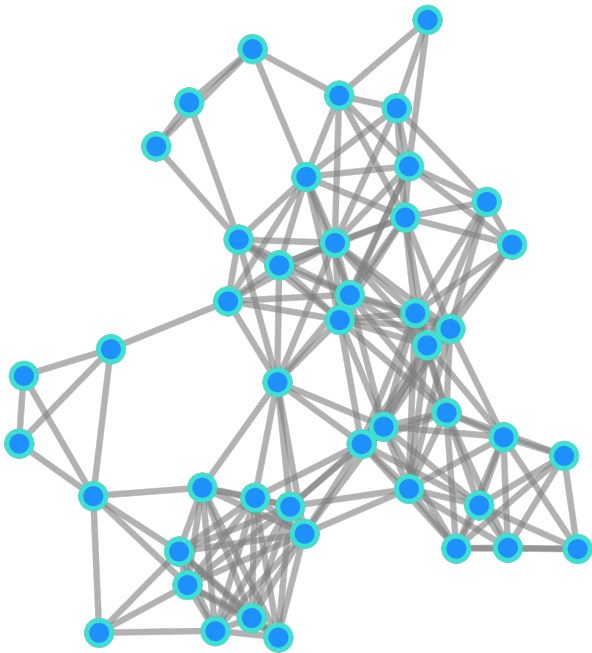


Systematic comparison of *graphs in general*



On measuring distances

How close (similar) are these two graphs?



On measuring distances

How close (similar) are these two mugs?



On measuring distances

How close (similar) are these two mugs?



How do their positions differ?
meters

How do their volumes differ?
liters

How do their temperatures differ?
degrees

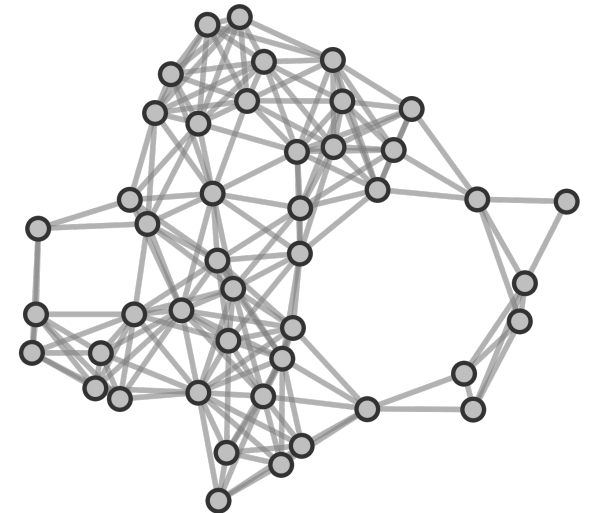
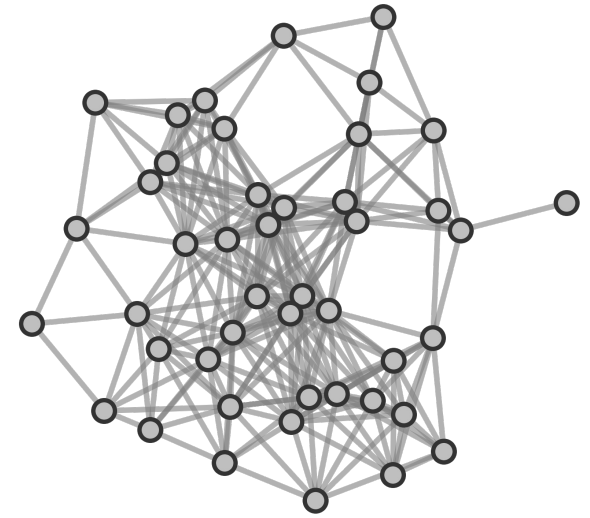
How do their functions differ?
...ask someone at a café?



Graph distances

We calculate the **dissimilarity** between pairs of graphs by measuring the **distance** between their **descriptors**.

- dissimilarity** → the graph distance in question
- descriptors** → a graph observable of interest
- distance** → a (well-motivated) distance metric



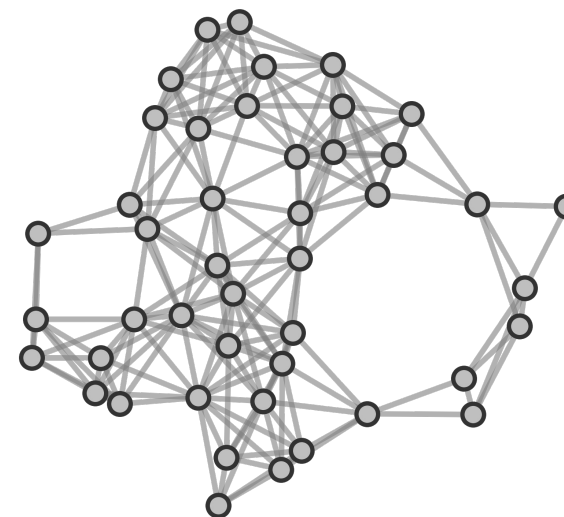
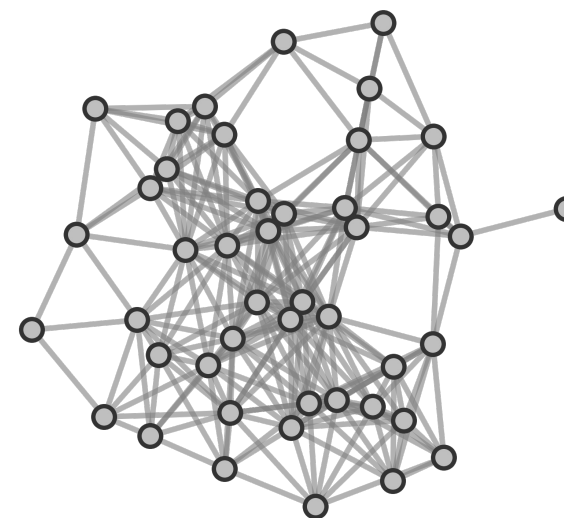
Graph distances

We calculate the **dissimilarity** between pairs of graphs by measuring the **distance** between their **descriptors**.

dissimilarity → the graph distance in question
descriptors → a graph observable of interest
distance → a (well-motivated) distance metric

Definition. Given a set of graphs $\mathcal{M} \subseteq \mathcal{G}$, a graph description Ψ , its descriptor space \mathcal{D} , and a distance d on \mathcal{D} , the associated graph distance measure $D : \mathcal{M} \times \mathcal{M} \rightarrow \mathbb{R}_+$ is a function defined by

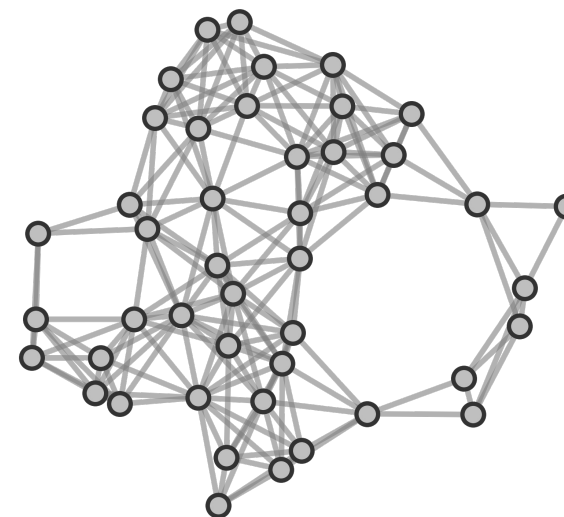
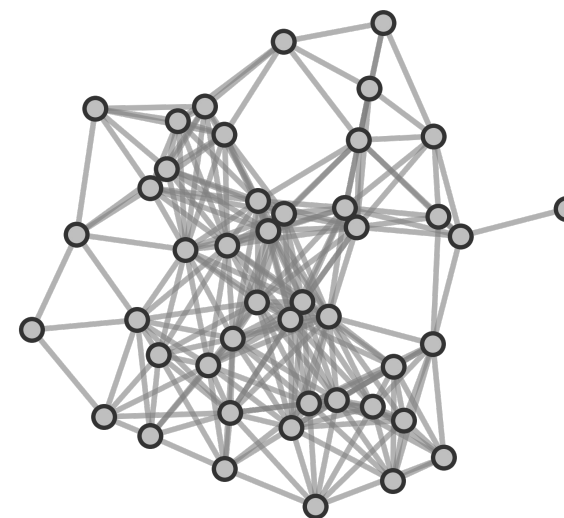
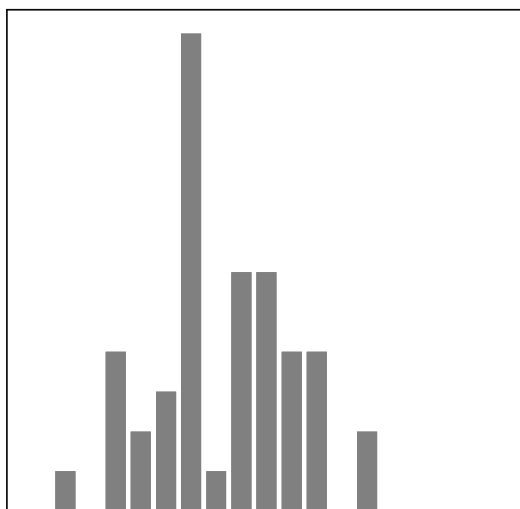
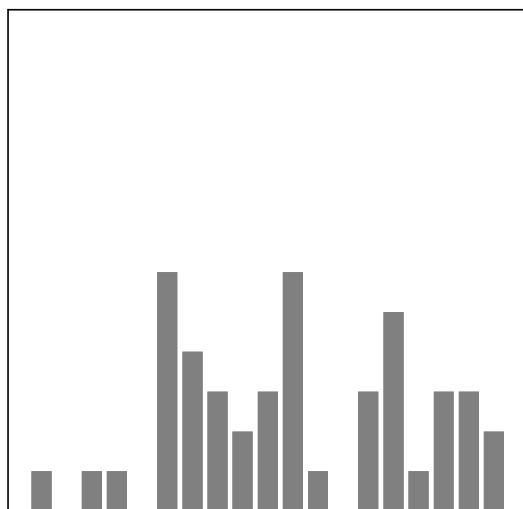
$$D(G, G') = d(\psi_G, \psi_{G'}).$$



Graph distances

We calculate the **dissimilarity** between pairs of graphs by measuring the **distance** between their **descriptors**.

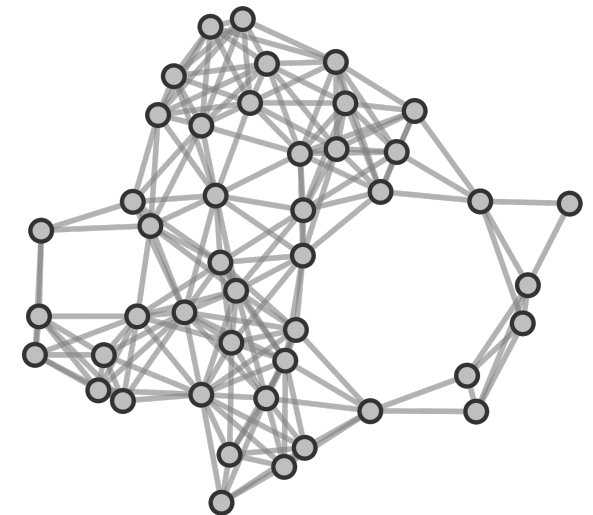
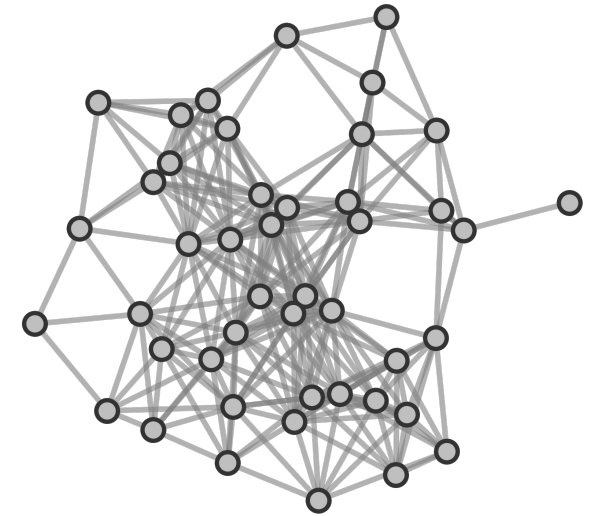
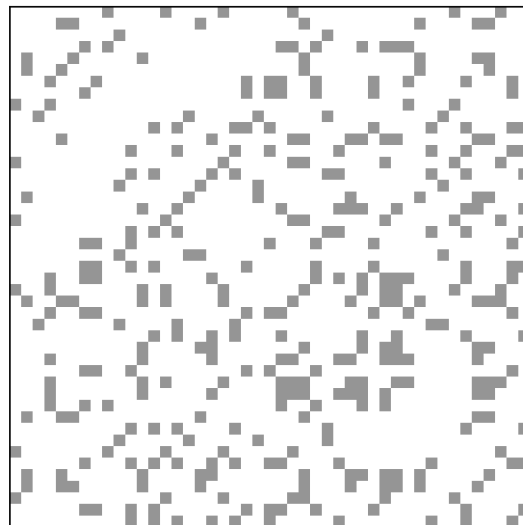
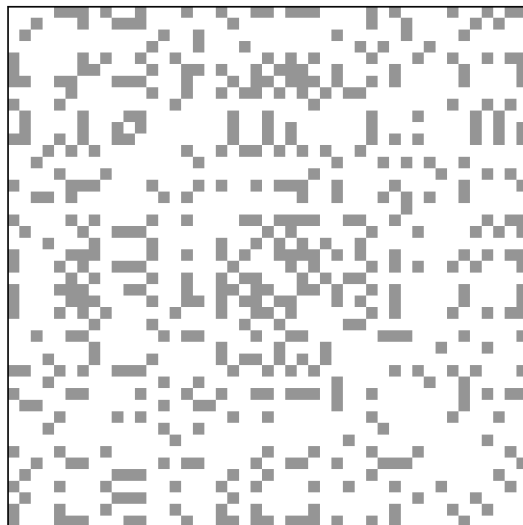
dissimilarity → Degree distribution JSD (DJS)
descriptors → the degree distribution
distance → Jensen-Shannon divergence



Graph distances

We calculate the **dissimilarity** between pairs of graphs by measuring the **distance** between their **descriptors**.

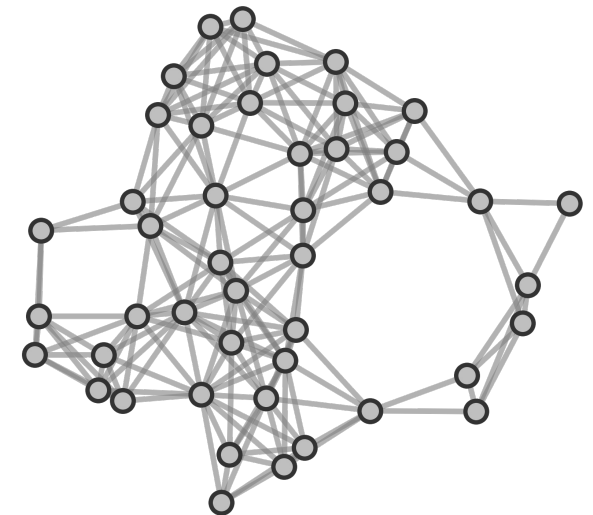
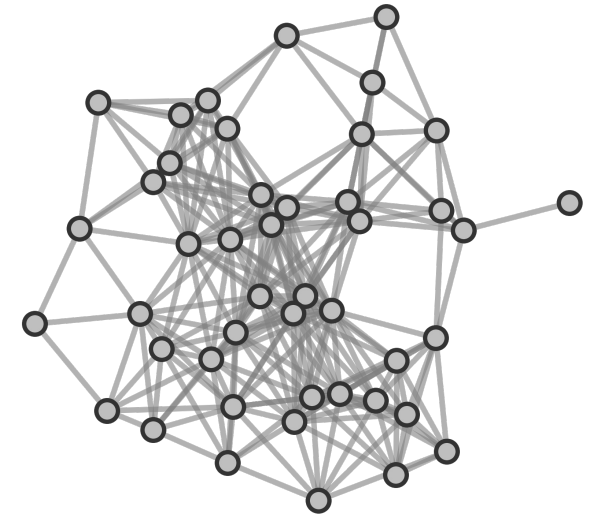
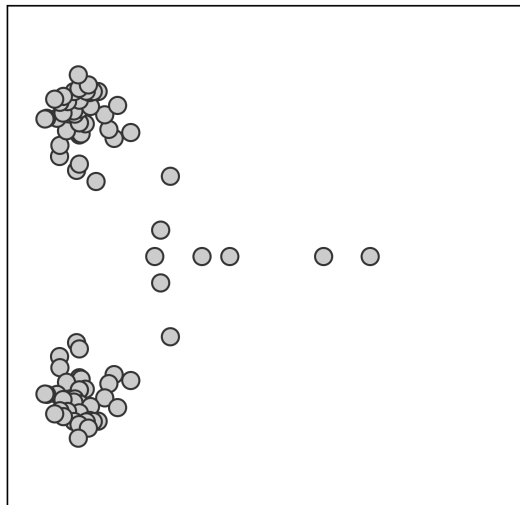
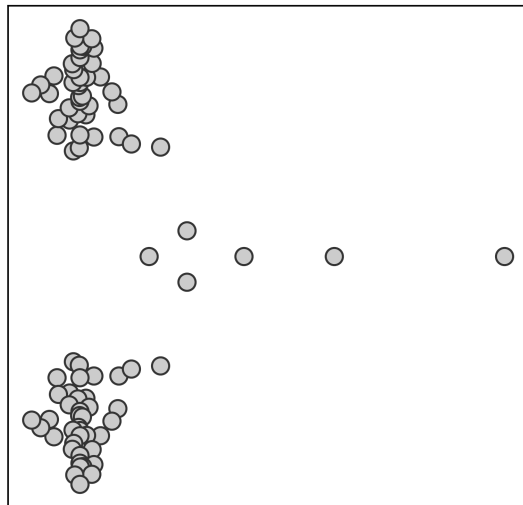
dissimilarity → Hamming distance (HAM)
descriptors → the adjacency matrix
distance → Euclidean distance (squared)



Graph distances

We calculate the **dissimilarity** between pairs of graphs by measuring the **distance** between their **descriptors**.

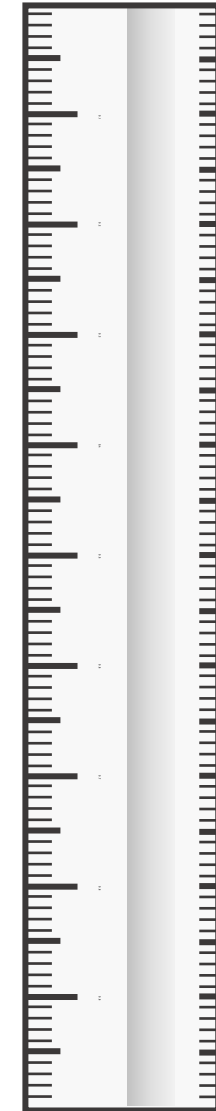
dissimilarity → Non-backtracking Spectral (NBD)
descriptors → eigenvalues of the NBM
distance → Earth-mover's distance



Graph distances

We calculate the **dissimilarity** between pairs of graphs by measuring the **distance** between their **descriptors**.

	Graph distance	Label
1	Jaccard (Jaccard, 1901)	JAC
2	Hamming (Hamming, R.W., 1950)	HAM
3	Hamming-Ipsen-Mikhailov (Jurman, Visintainer, Filosi, Riccadonna, & Furlanello, 2015)	HIM
4	Frobenius (Golub & van Loan, 2013)	FRO
5	Polynomial dissimilarity (Donnat & Holmes, 2018)	POD
6	Degree JSD (Carpi, Rosso, Saco, & Ravetti, 2011)	DJS
7	Portrait divergence (Bagrow & Bollt, 2019)	POR
8	Quantum spectral JSD (De Domenico & Biamonte, 2016)	QJS
9	Communicability sequence (Chen, Shi, Qin, Xu, & Pan, 2018)	CSE
10	Graph diffusion distance (Hammond, Gur, & Johnson, 2013)	GDD
11	Resistance-perturbation (Monnig & Meyer, 2018)	REP
12	NetLSD (Tsitsulin et al., 2018)	LSD
13	Lap. spectrum; Gauss. kernel, JSD (Jurman et al., 2011)	LGJ
14	Lap. spectrum; Loren. kernel, Euc. (Jurman et al., 2011)	LLE
15	Ipsen-Mikhailov (Ipsen & Mikhailov, 2002)	IPM
16	Non-backtracking eigenvalue (Torres et al., 2019)	NBD
17	Distributional Non-backtracking (Mellor & Grusovin, 2019)	DNB
18	D-measure distance (Schieber et al., 2017)	DMD
19	DeltaCon (Koutra et al., 2016)	DCN
20	NetSimile (Berlingerio et al., 2012)	NES



The within-ensemble graph distance (WEGD)

Definition:

the average distance between pairs of graphs with the same parameterization independently sampled from a given ensemble.

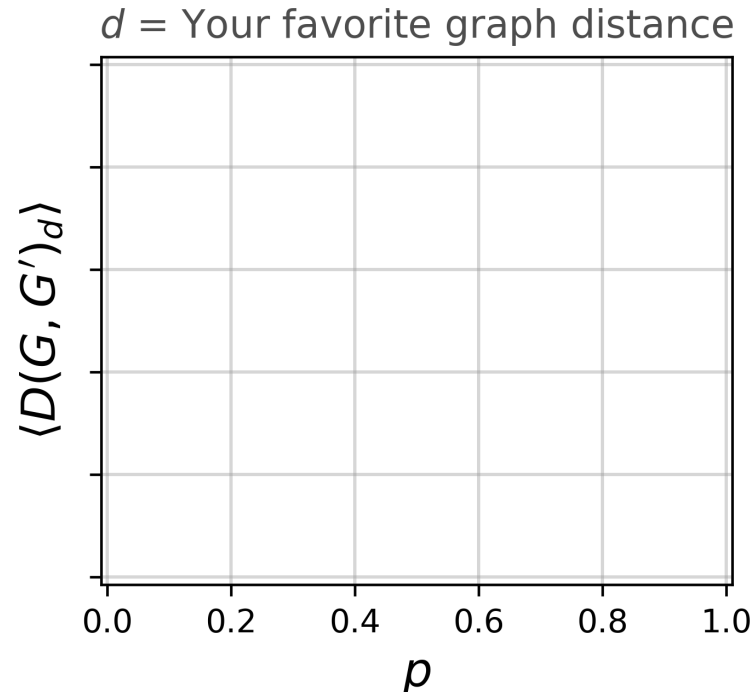
Random graph ensembles are the ideal setting to define a network comparison benchmark.

1. They are well-studied objects on their own.
2. We understand key properties quite well.
3. We can tune them to suit our questions.

The within-ensemble graph distance (WEGD)

Definition:

the average distance between pairs of graphs with the same parameterization independently sampled from a given ensemble.



For example, $G(n, p)$

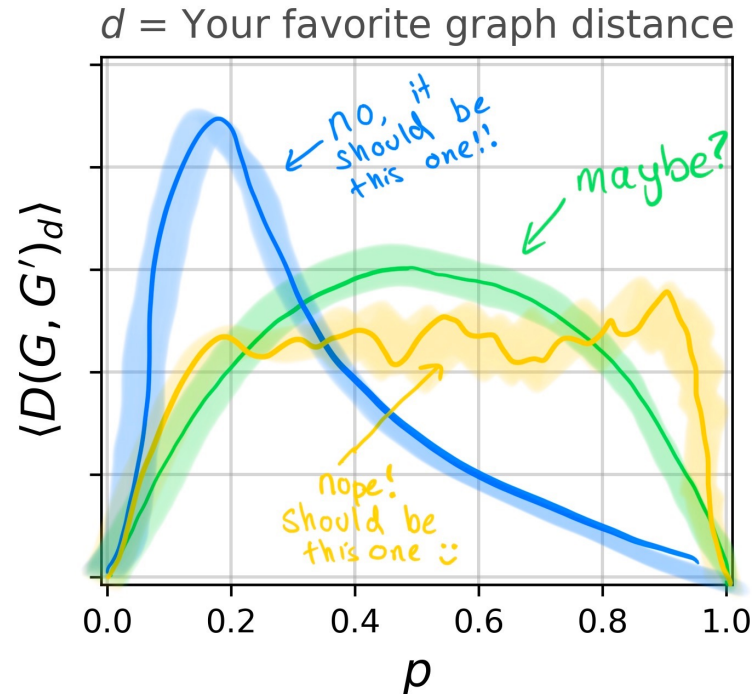
- fixed value for n
- vary p

What should a graph distance capture?
Sparsity? Symmetry around $p=0.5$? Etc.

The within-ensemble graph distance (WEGD)

Definition:

the average distance between pairs of graphs with the same parameterization independently sampled from a given ensemble.

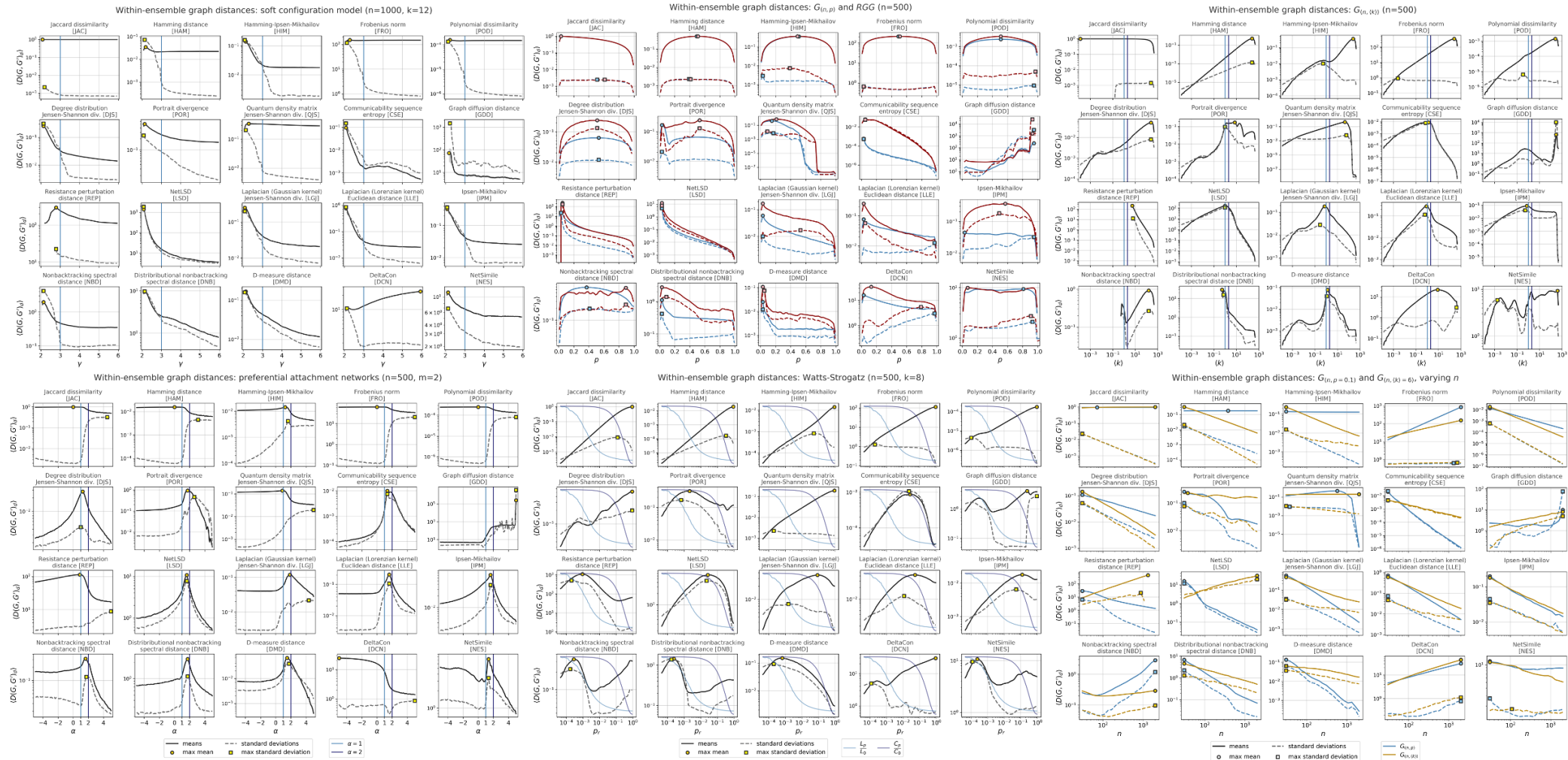


For example, $G(n, p)$

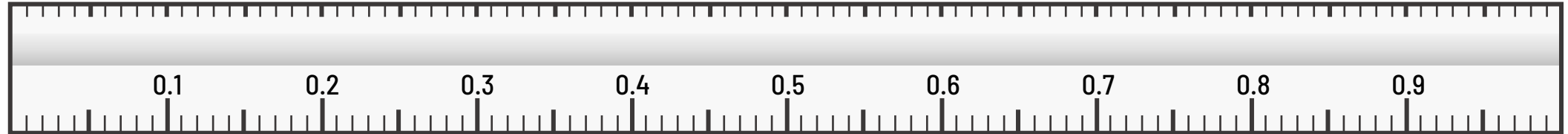
- fixed value for n
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What should a graph distance capture?
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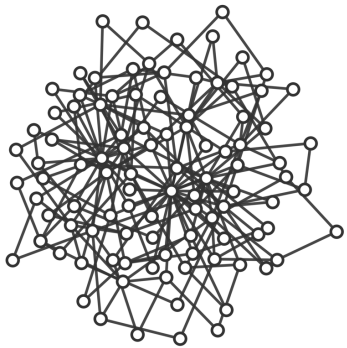
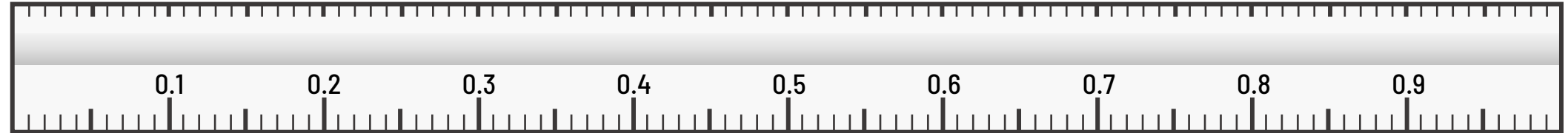
The within-ensemble graph distance (WEGD)



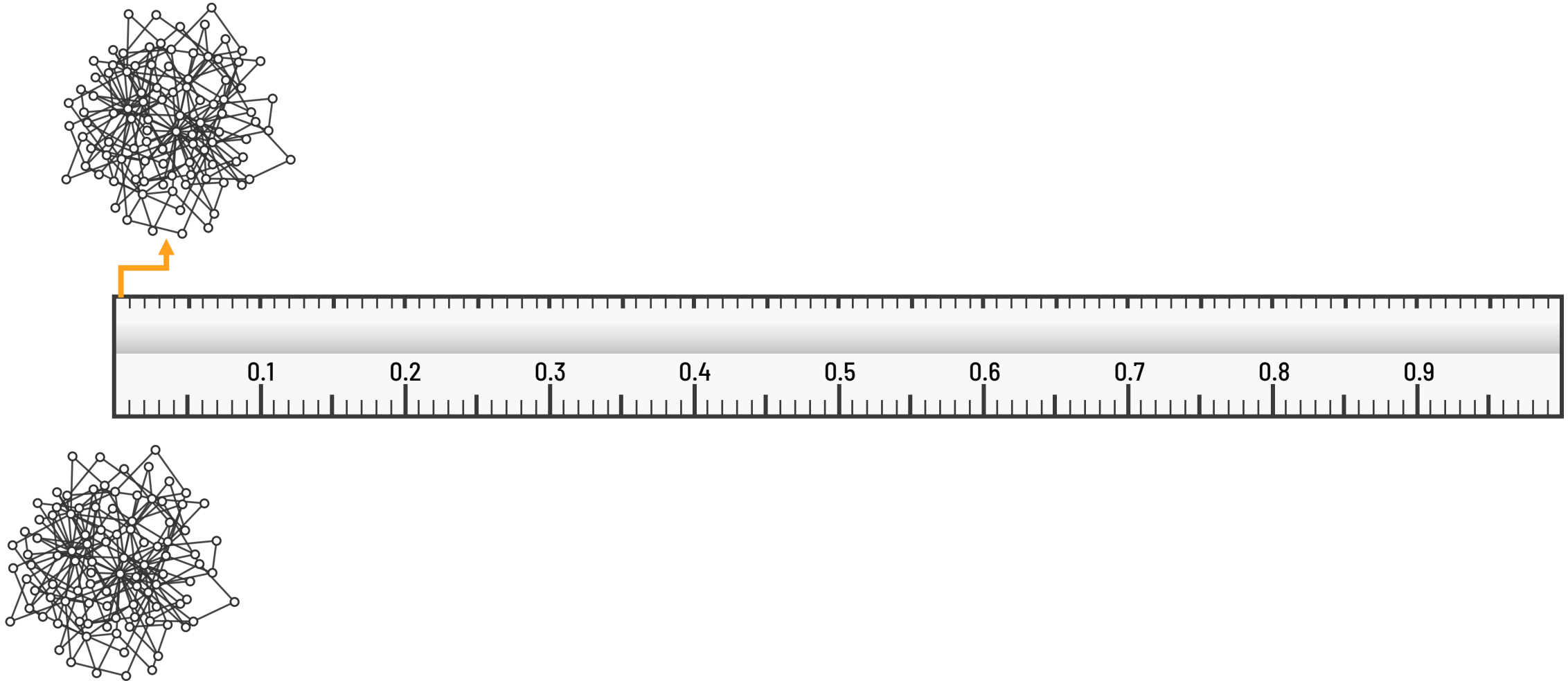
Toward a *standardized* graph distance



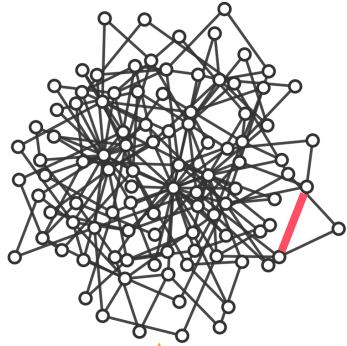
Toward a *standardized* graph distance



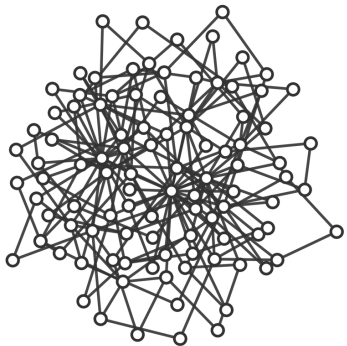
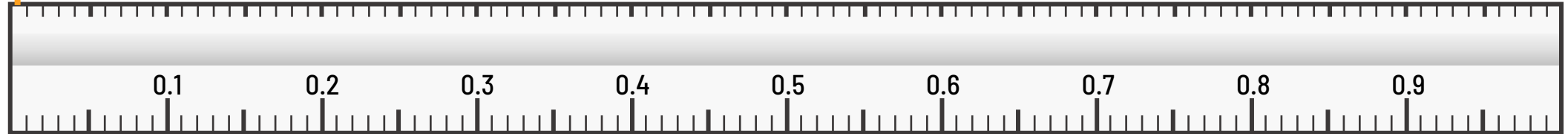
Toward a *standardized* graph distance



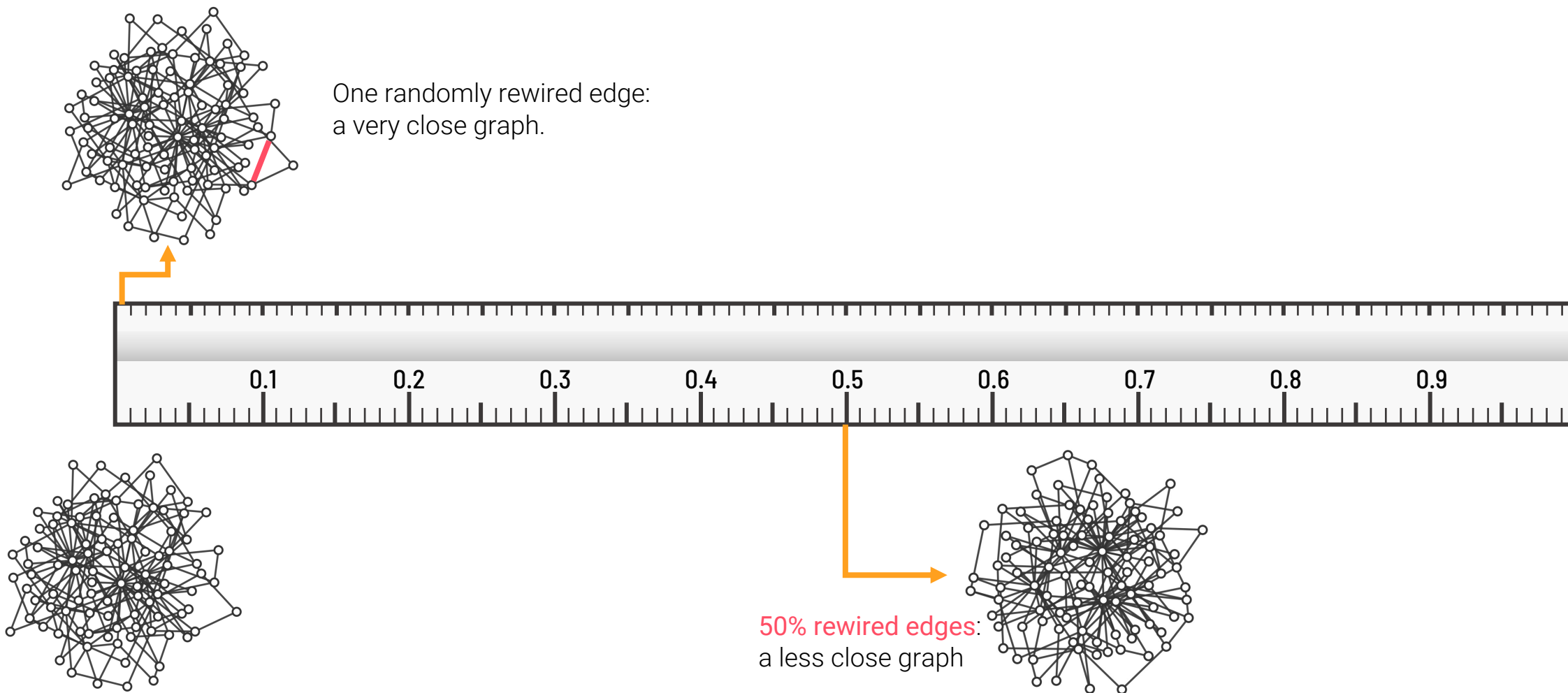
Toward a *standardized* graph distance



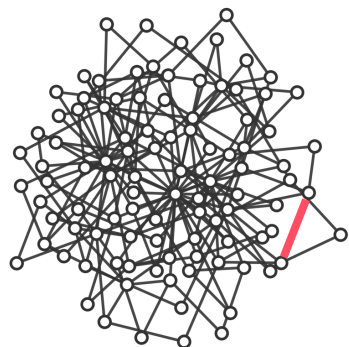
One randomly rewired edge:
a very close graph.



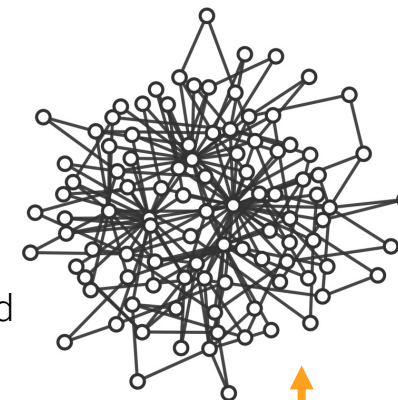
Toward a *standardized* graph distance



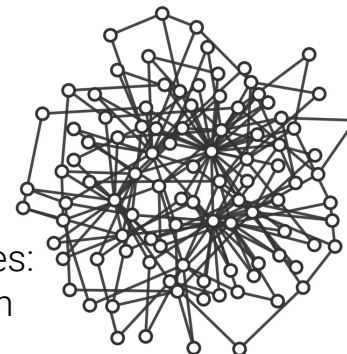
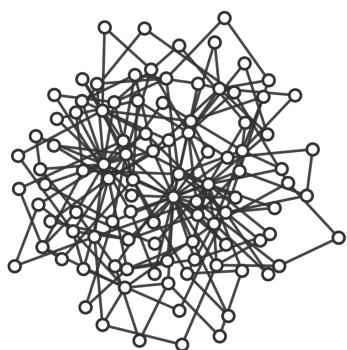
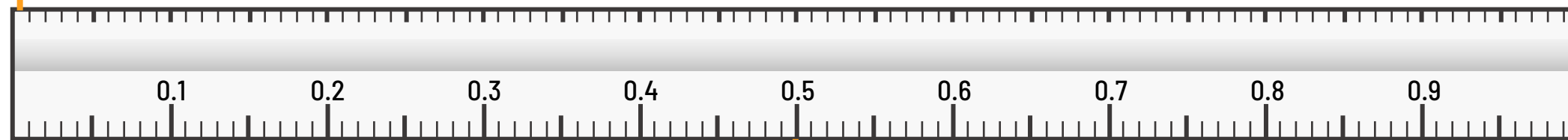
Toward a *standardized* graph distance



One randomly rewired edge:
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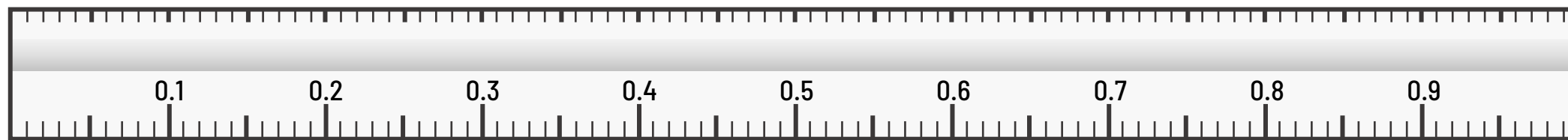
100% rewired edges:
a notion of standardized
graph distance



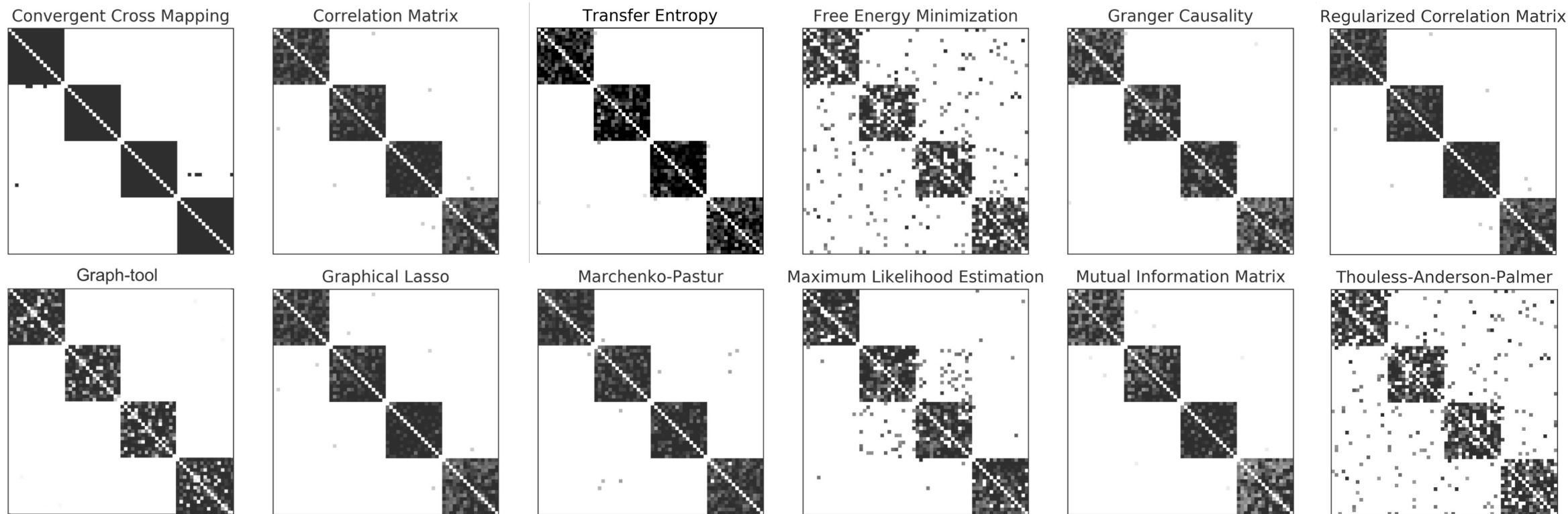
50% rewired edges:
a less close graph

Toward a *standardized* graph distance: the metER

$$D_{d,s}(G_1, G_2) = \frac{D_d(G_1, G_2)}{\langle D_d(G_{n,p}, G'_{n,p}) \rangle}$$



Systematic comparison of different tools

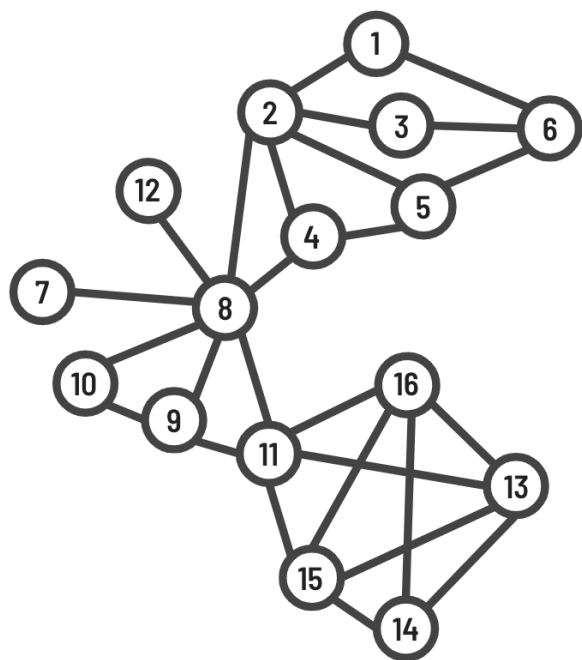


Models, representation, theory

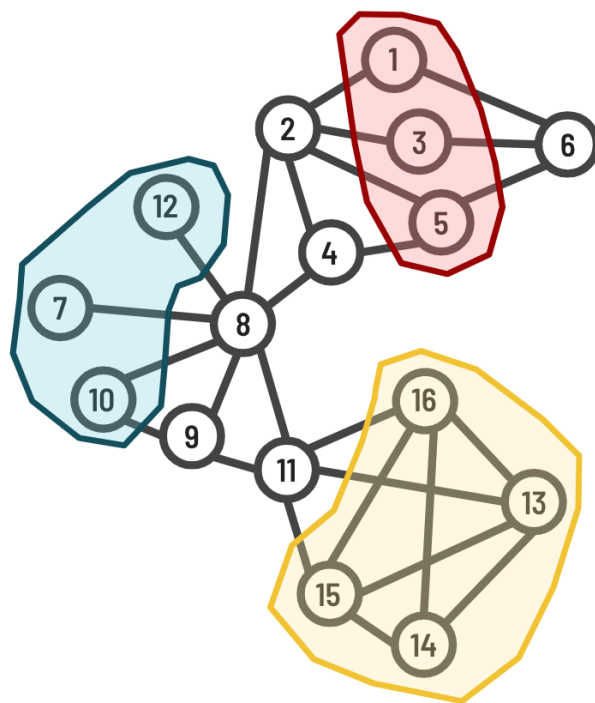
- Can we quantify how informative a given network structure is?
- Are certain networks more or less informative? If so, why?
- Are there principled ways to identify the right scale to model a system?
- Do networks in nature and society differ in information or scale?



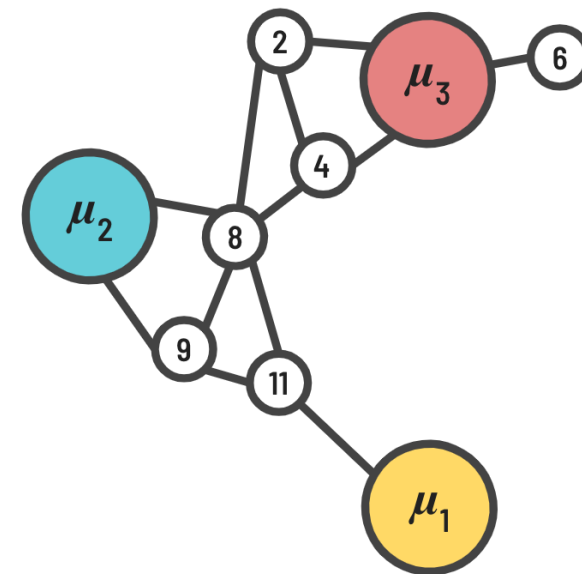
Informative *higher scales* in complex networks



microscale



grouping

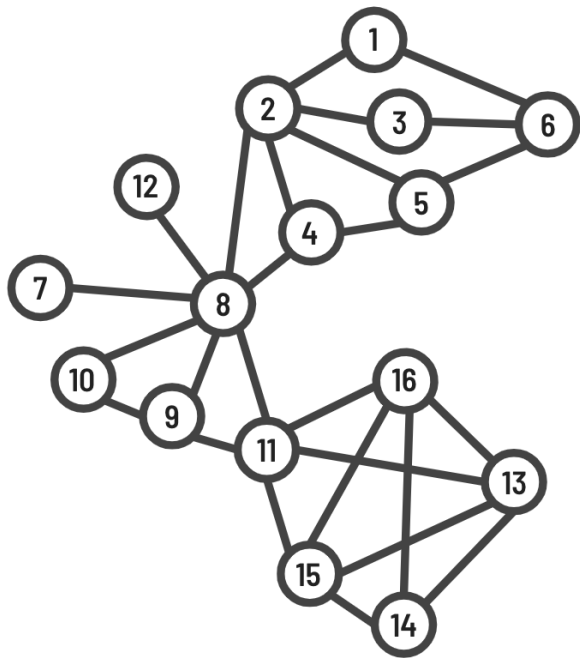


macroscale

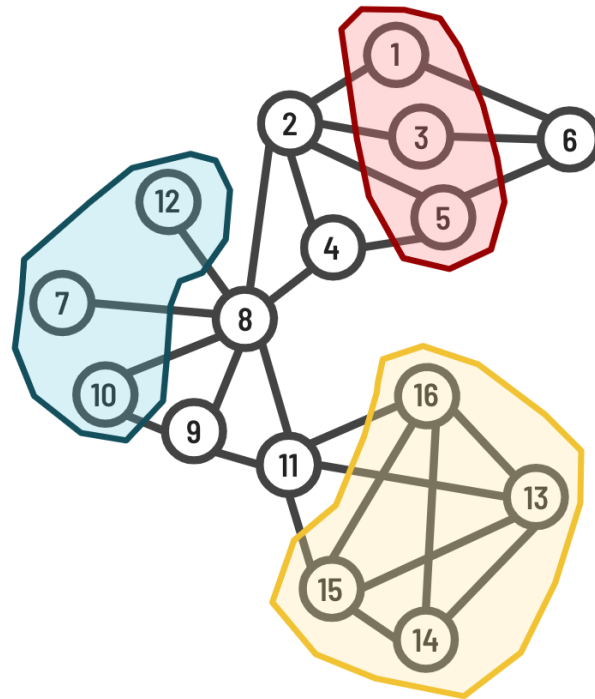
Causal emergence

a phenomenon where a coarse-grained “macroscale” network has more effective information than its corresponding “microscale” network

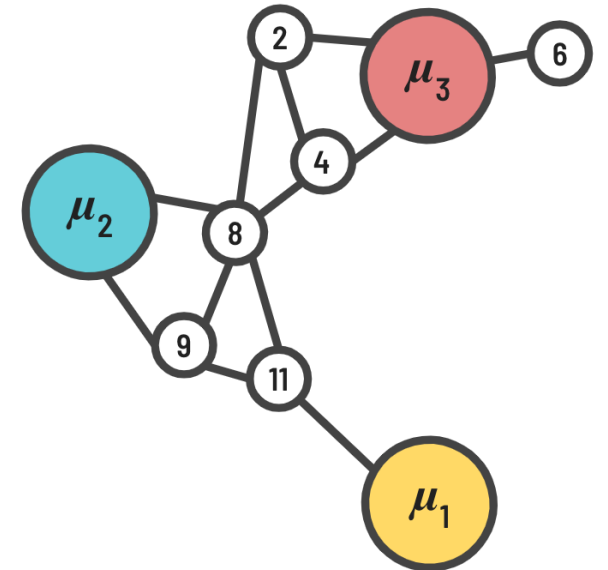
$$EI = H(\langle W_i^{out} \rangle) - \langle H(W_i^{out}) \rangle$$



microscale



grouping



macroscale

Causal emergence

a phenomenon where a coarse-grained “macroscale” network has more effective information than its corresponding “microscale” network

$$EI = H(\langle W_i^{out} \rangle) - \langle H(W_i^{out}) \rangle$$

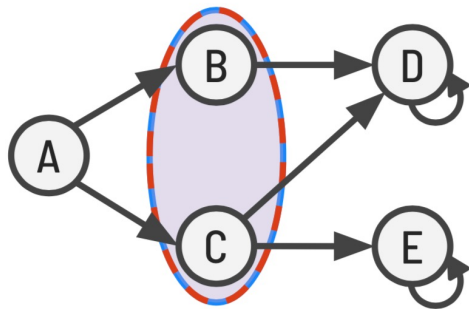
	A	B	C	D	E
A	w_{AA}	w_{AB}	w_{AC}	w_{AD}	w_{AE}
B	w_{BA}	w_{BB}	w_{BC}	w_{BD}	w_{BE}
C	w_{CA}	w_{CB}	w_{CC}	w_{CD}	w_{CE}
D	w_{DA}	w_{DB}	w_{DC}	w_{DD}	w_{DE}
E	w_{EA}	w_{EB}	w_{EC}	w_{ED}	w_{EE}

	A	B	C	D	E
A	w_{AA}	w_{AB}	w_{AC}	w_{AD}	w_{AE}
B	w_{BA}	w_{BB}	w_{BC}	w_{BD}	w_{BE}
C	w_{CA}	w_{CB}	w_{CC}	w_{CD}	w_{CE}
D	w_{DA}	w_{DB}	w_{DC}	w_{DD}	w_{DE}
E	w_{EA}	w_{EB}	w_{EC}	w_{ED}	w_{EE}



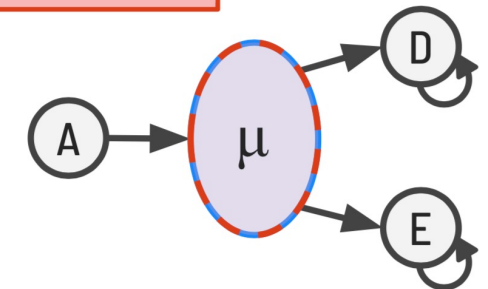
	A	B	C	D	E
A	w_{AA}	w_{AB}	w_{AC}	w_{AD}	w_{AE}
μ	$w_{\mu A}$	$w_{\mu B}$	$w_{\mu C}$	$w_{\mu D}$	$w_{\mu E}$
D	w_{DA}	w_{DB}	w_{DC}	w_{DD}	w_{DE}
E	w_{EA}	w_{EB}	w_{EC}	w_{ED}	w_{EE}

	A	μ	D	E
A	w_{AA}	$w_{A\mu}$	w_{AD}	w_{AE}
μ	$w_{\mu A}$	$w_{\mu\mu}$	$w_{\mu D}$	$w_{\mu E}$
D	w_{DA}	$w_{D\mu}$	w_{DD}	w_{DE}
E	w_{EA}	$w_{E\mu}$	w_{ED}	w_{EE}



Group the out-weights of B and C together to form a new W_{μ}^{out}

Sum the in-weights to B and C to create the in-weights of μ



microscale

macroscale

Causal emergence

a quantity that indicates how much higher-scale structural information a network contains.

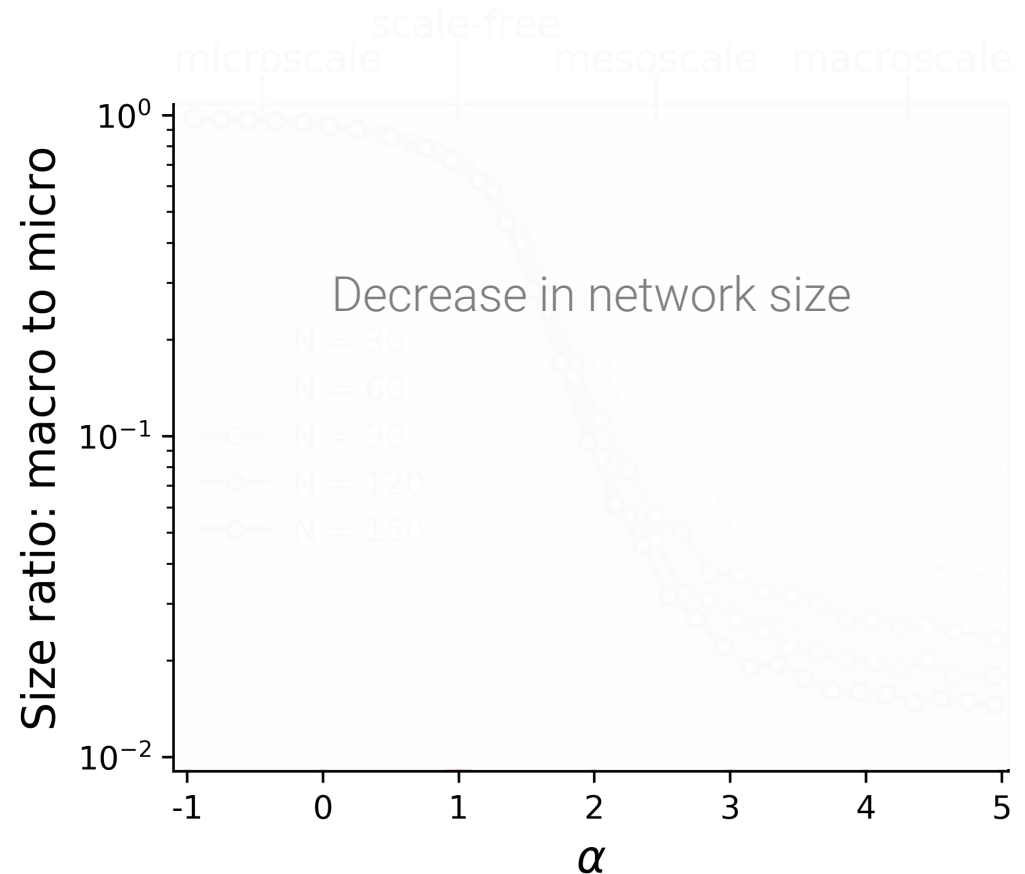
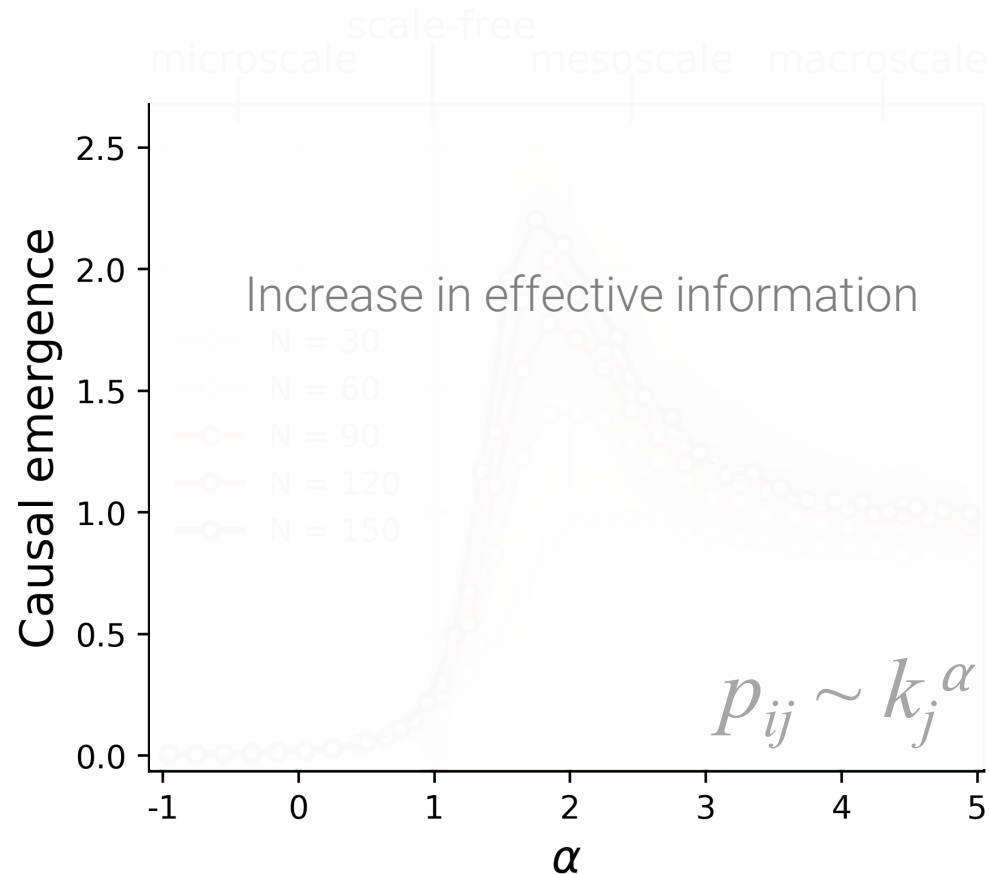
Increase in effective information

Decrease in network size

63

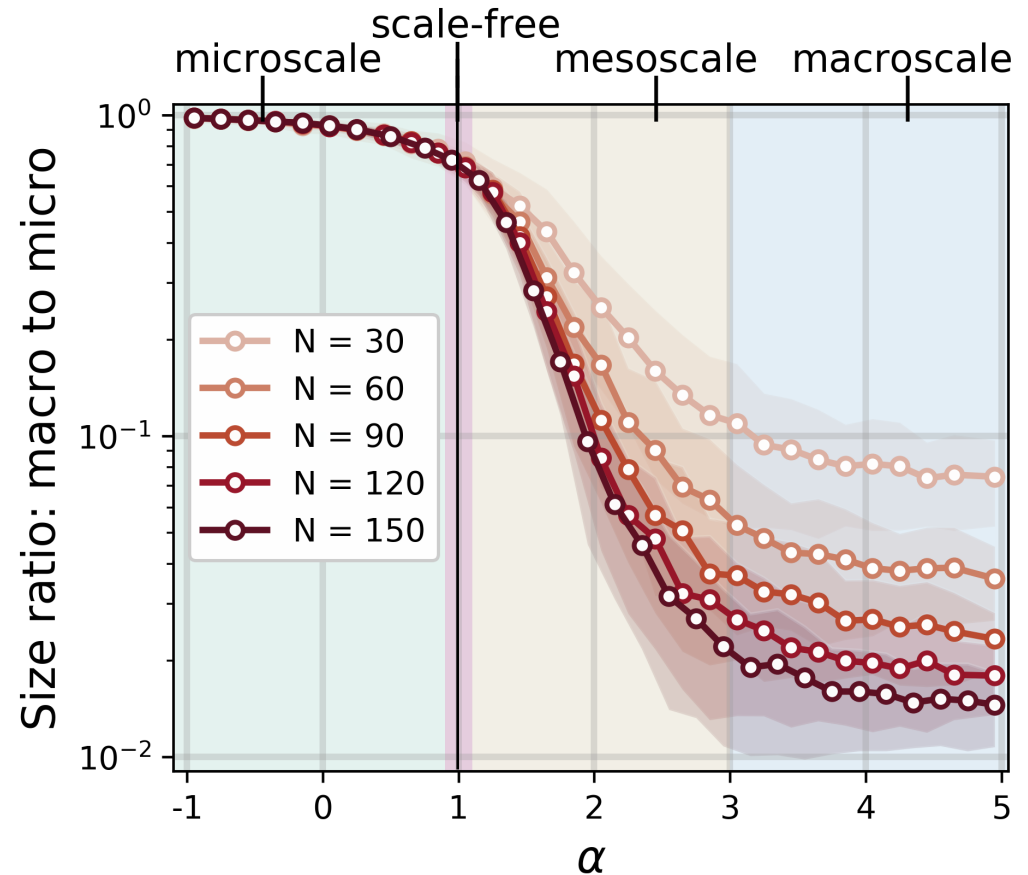
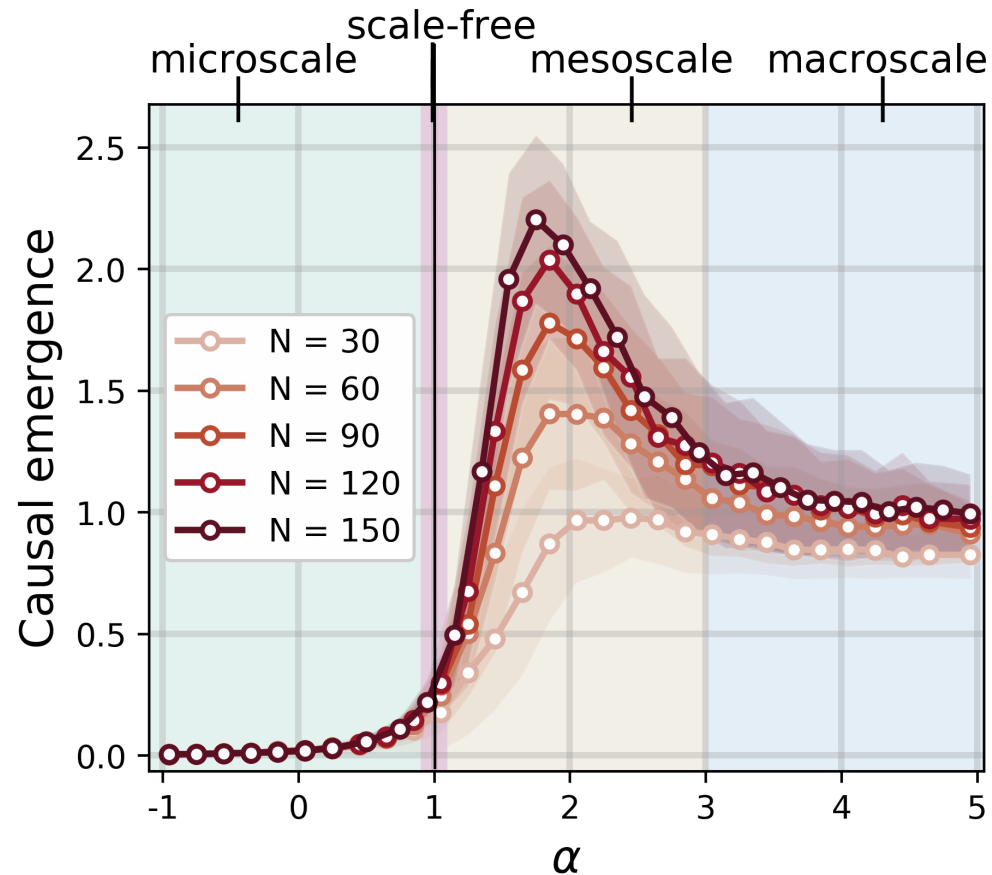
Causal emergence

a quantity that indicates how much higher-scale structural information a network contains.



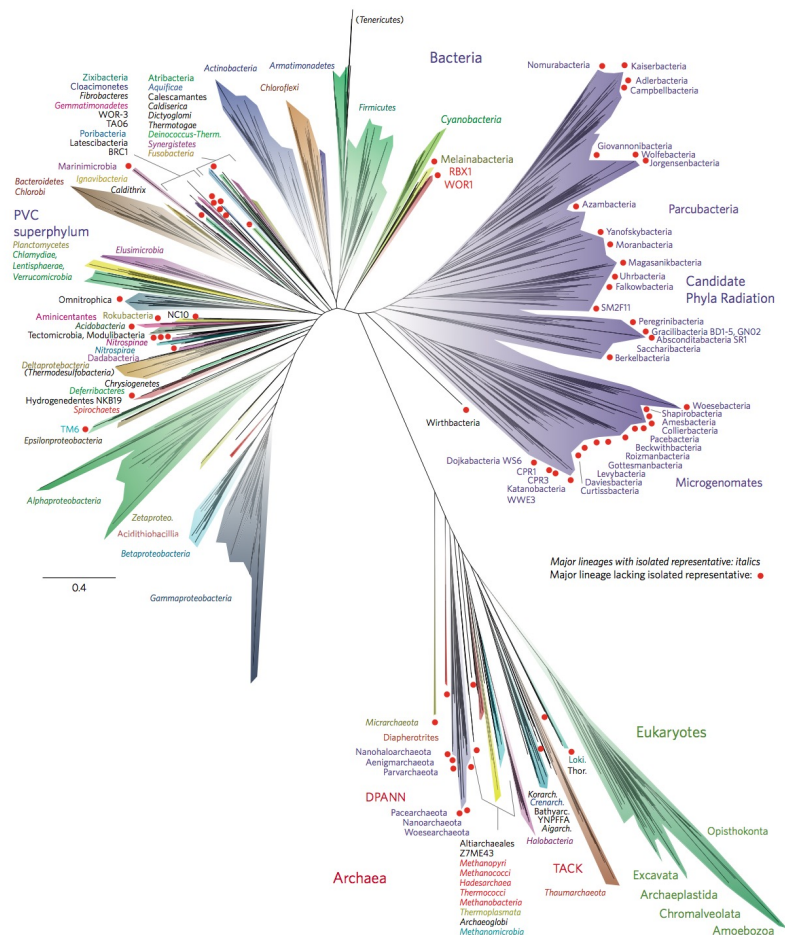
Causal emergence

a quantity that indicates how much higher-scale structural information a network contains.



Causal emergence in biological systems

a quantity that indicates how much higher-scale structural information a network contains.

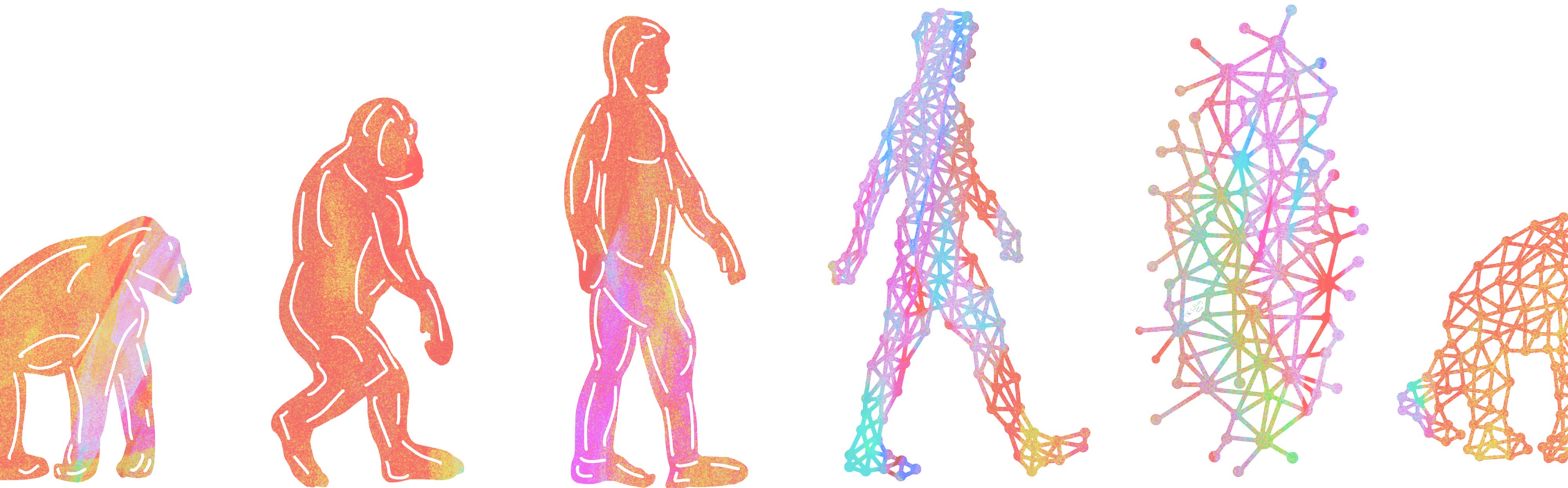


Biological systems are characterized by noisy microscale interactions leading to emergent macroscale behavior.

- Carlile, M. (1982). Prokaryotes and eukaryotes: Strategies and successes. *Trends in Biochemical Sciences*. 7.4, pp. 128–130.
- Edelman G.M. & Gally, J.A. (2011). Degeneracy and complexity in biological systems. *PNAS*. 98.24, pp. 13763–13768.
- Lukeš, J., et al. (2011). How a neutral evolutionary ratchet can build cellular complexity. *IUBMB Life*. 63.7, pp. 528–537.
- Tsimring, L.S. (2014). Noise in biology. *Reports on Progress in Physics*. 77.2.
- Brunet, T.D.P. & Doolittle, W.F. (2018). The generality of constructive neutral evolution. *Biology and Philosophy*. 33.1-2.

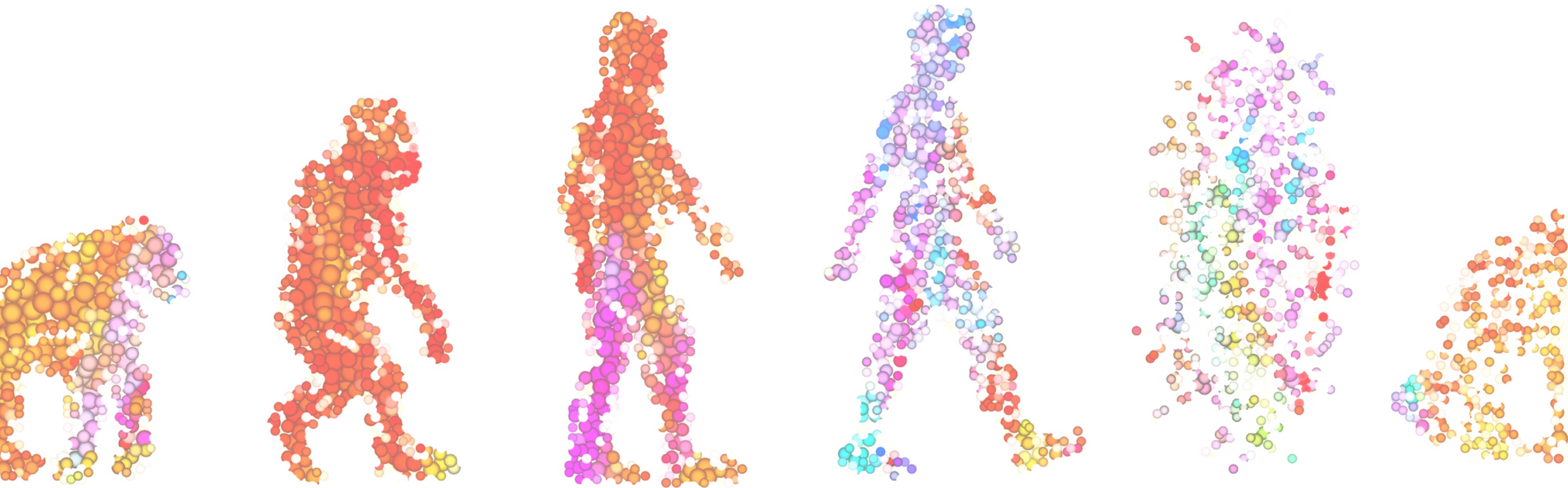
With a dataset of 1800+ species of protein-protein interaction networks, we find that **eukaryotic species** have more causal emergence than **prokaryotic species**.

Higher-order structure in networks...



Higher-order structure in networks...

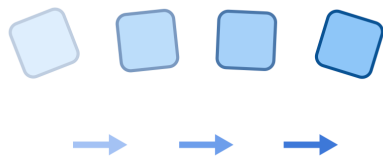
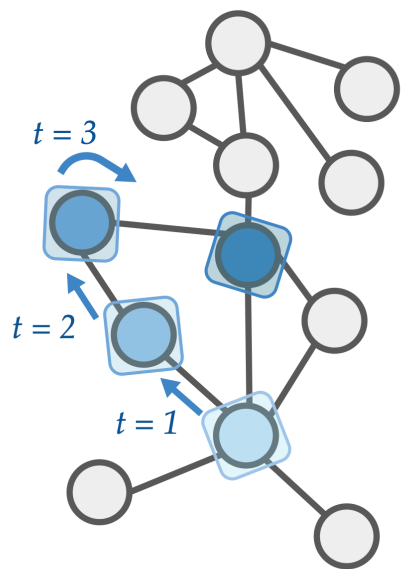
...what about higher-order dynamics?



Higher-order structure in networks...

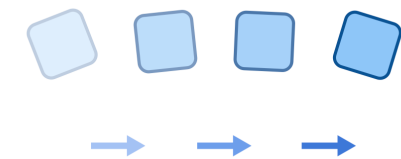
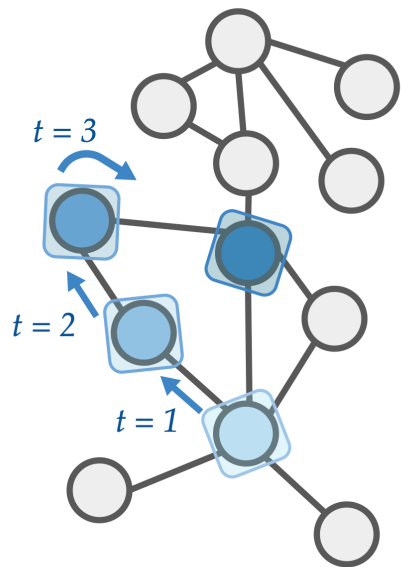
...what about higher-order dynamics?

Single random walker



Higher-order dynamical process: “Blob walks”

Single random walker
(blob of size $b = 1$)



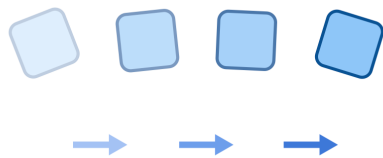
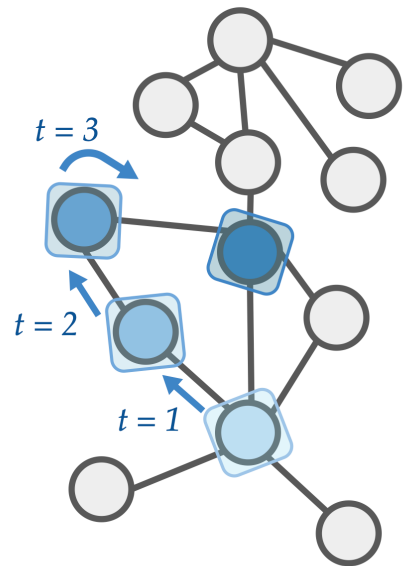
Higher-order random walks:

- Diffusion of a connected subgraph (a “blob”), B_t of size b on a graph.
 - A blob of size b occupies an entire connected subgraph of size b . At each timestep, the blob transitions into a new configuration B_{t+1} subject to the constraint that the new configuration retains $b-1$ nodes in B_t while again forming a connected subgraph.

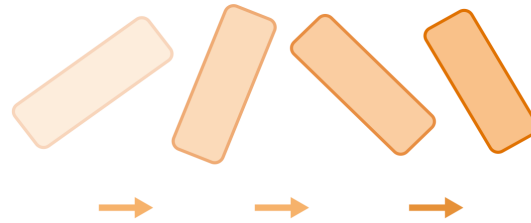
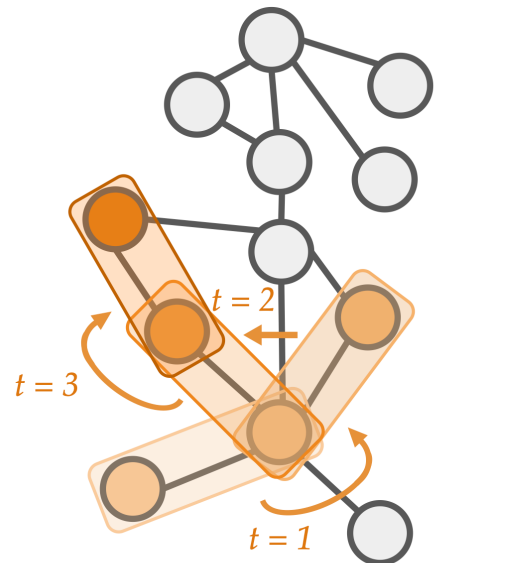
A classic random walker in this framework is simply a blob of size $b=1$.

Higher-order dynamical process: “Blob walks”

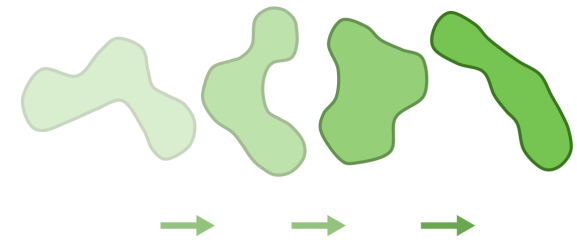
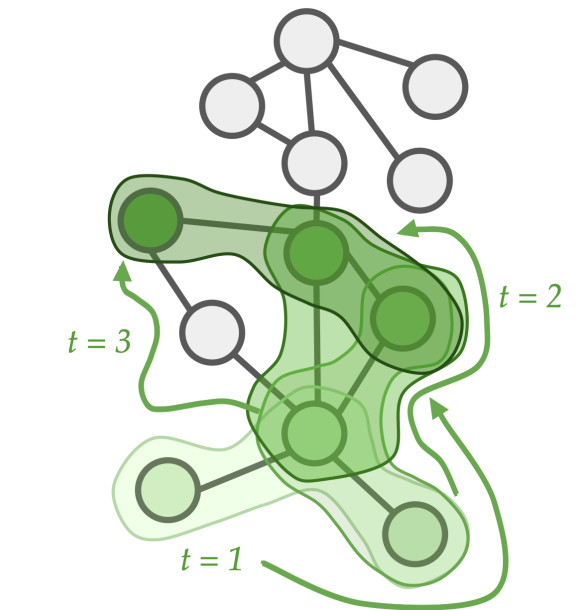
Single random walker
(blob of size $b = 1$)



Blob diffusion
(blob of size $b = 2$)

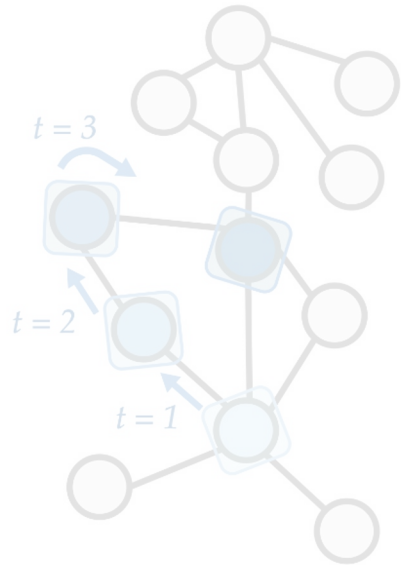


Blob diffusion
(blob of size $b = 3$)



Higher-order dynamical process: “Blob walks”

Single random walker
(blob of size $b = 1$)

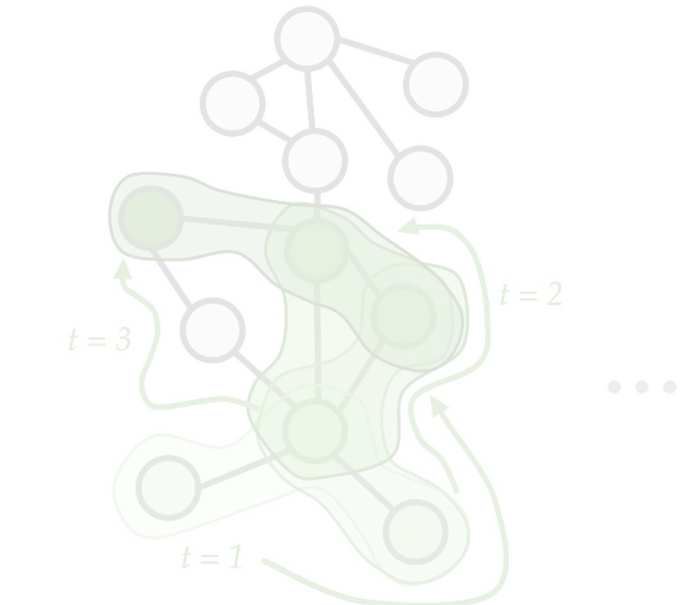


Moritz Laber
PhD Student,
Network Science

Blob diffusion
(blob of size $b = 2$)



Blob diffusion
(blob of size $b = 3$)



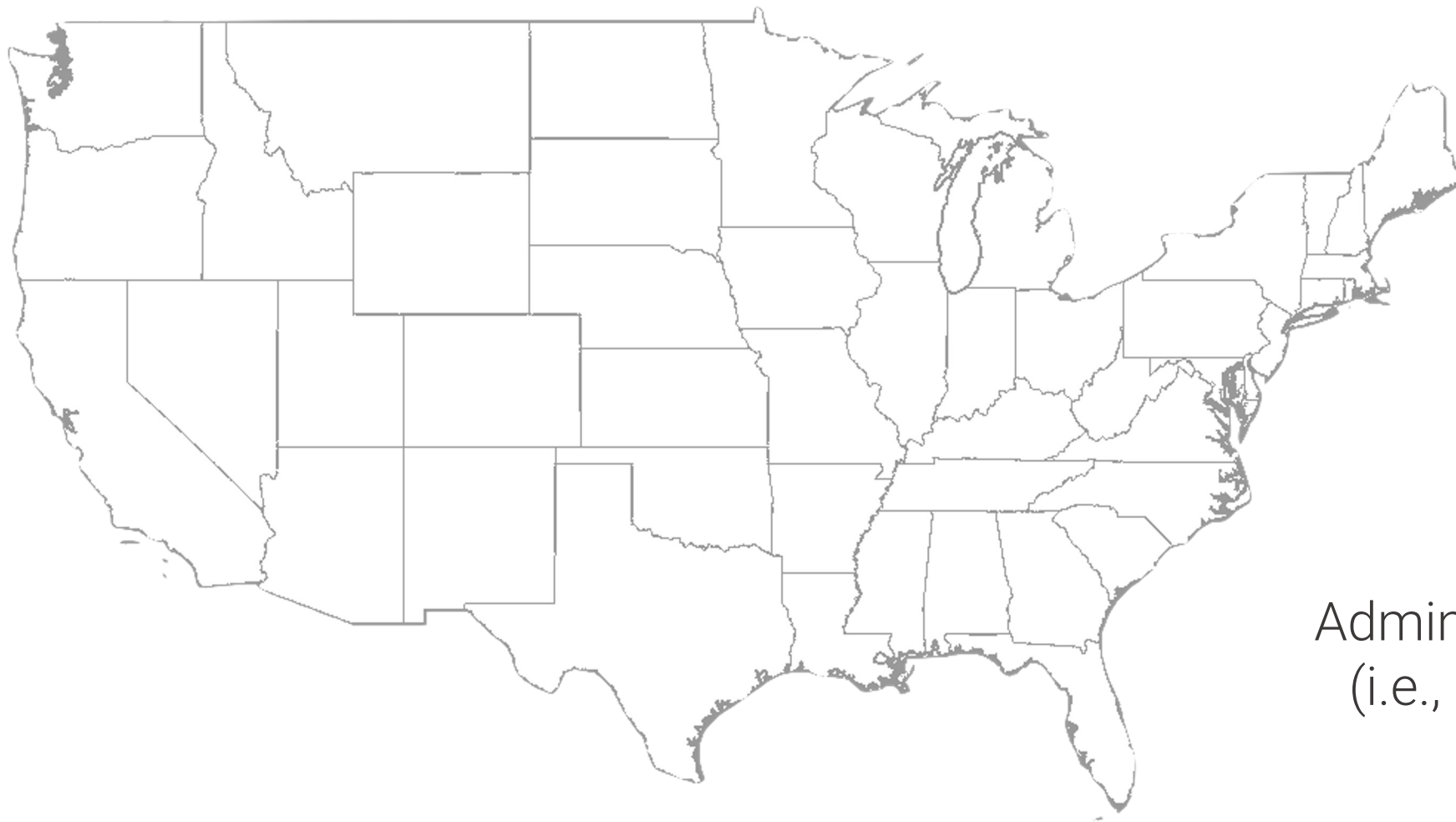
Led by Northeastern University PhD student, **Moritz Laber**, who will discuss these results at NetSci 2023 in Vienna.



Today

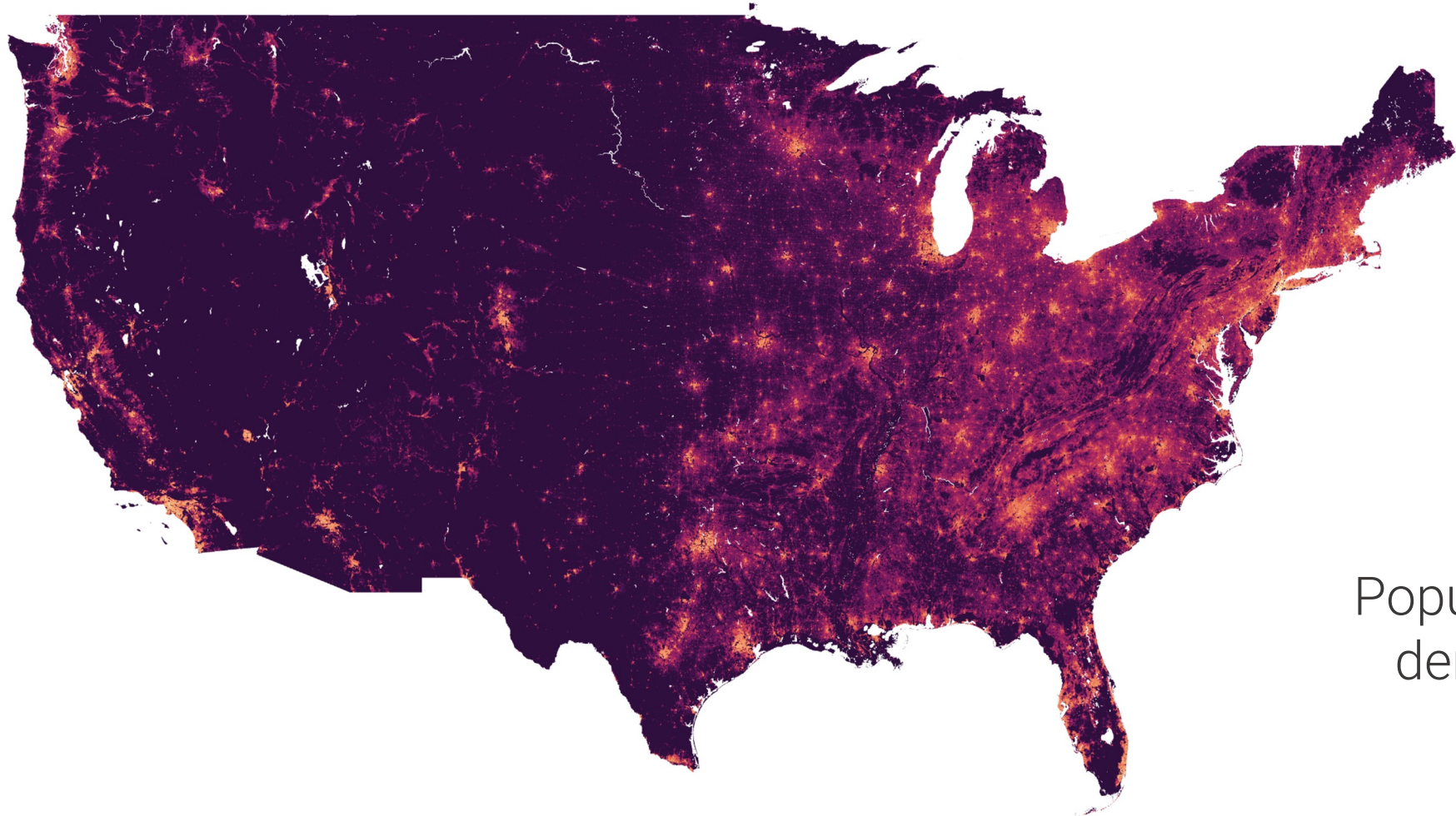
1. Introduction
2. Complexity
 - i. Reconstruction & comparison*
 - ii. Informative network scales & **representation***
3. Society
 - i. Mobile device data for disease modeling*
 - ii. Assorted COVID-19 projects*
4. Research vision and outlook

Models, representation, theory



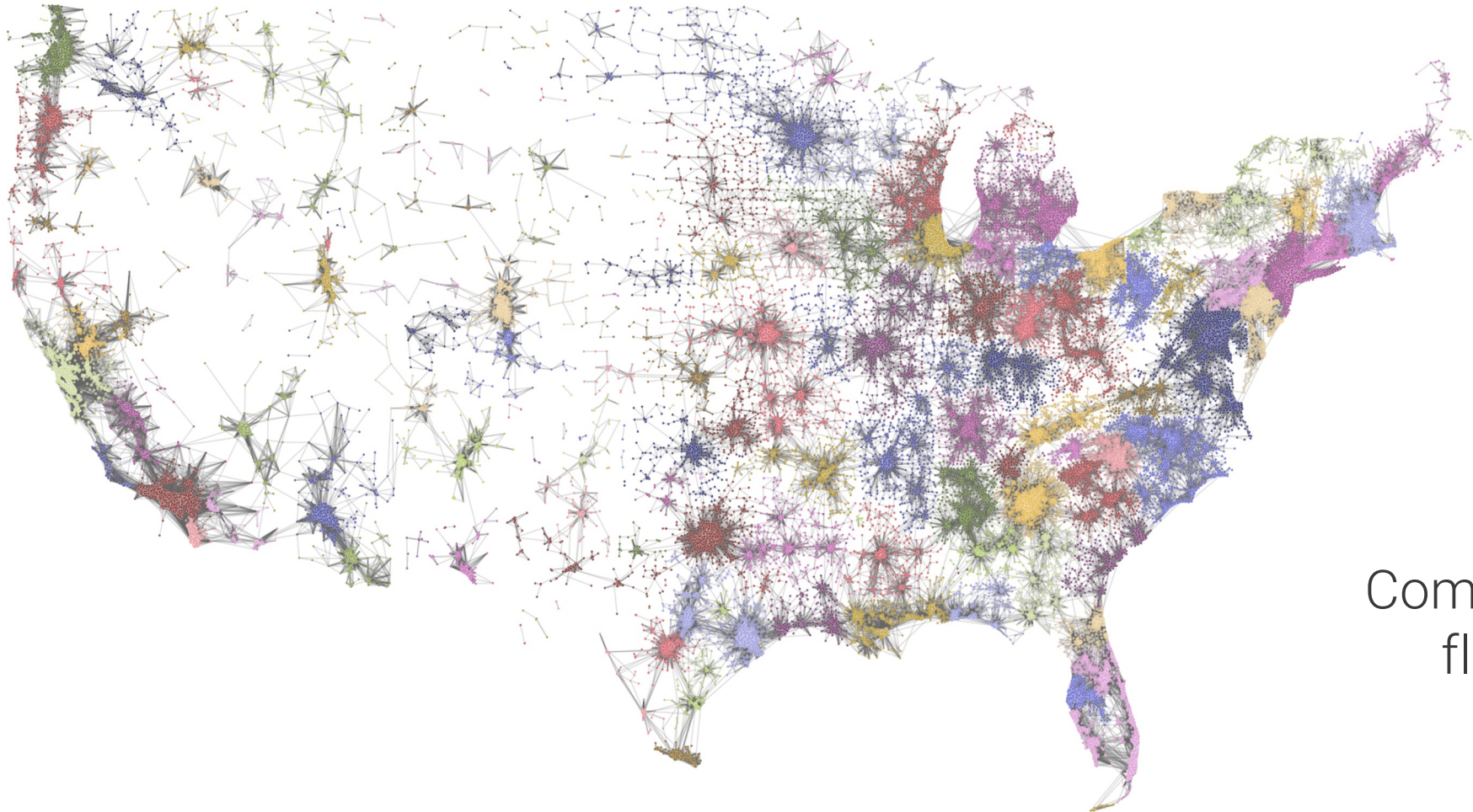
Administrative
(i.e., states)

Models, representation, theory



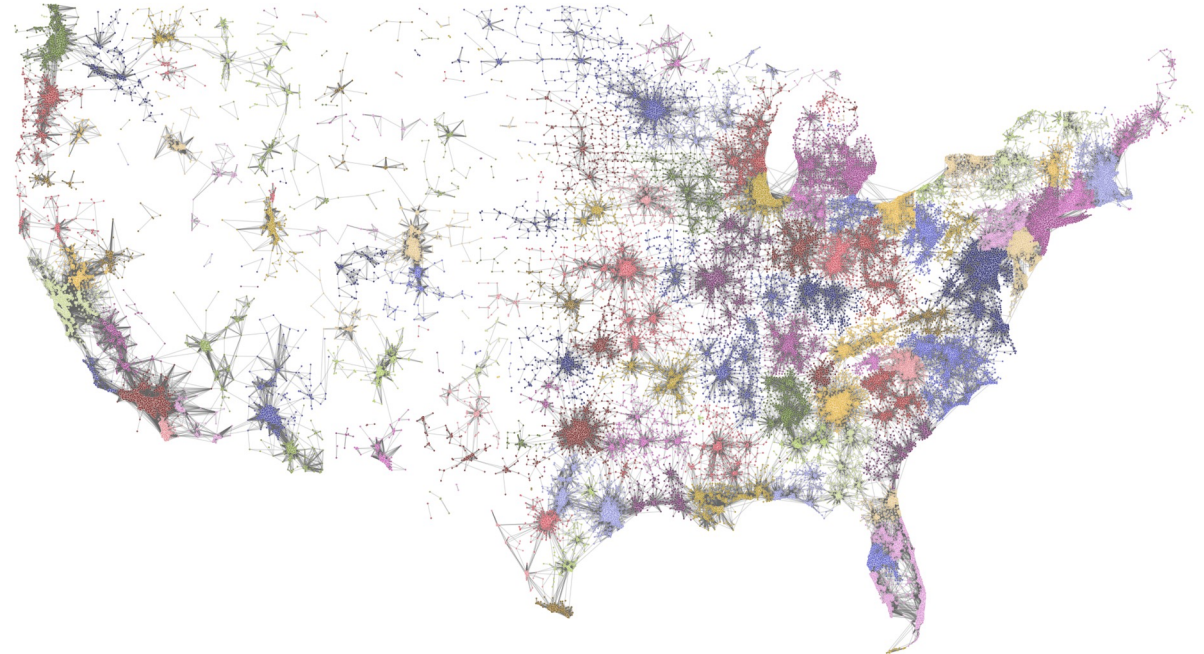
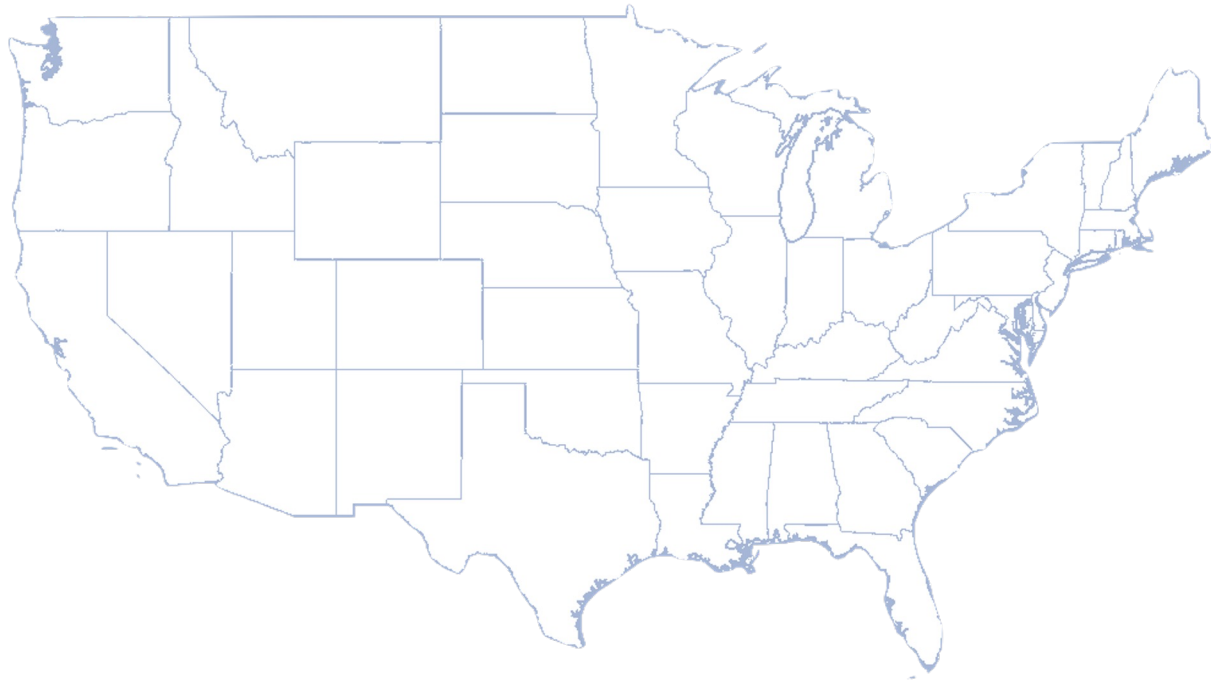
Population
density

Models, representation, theory

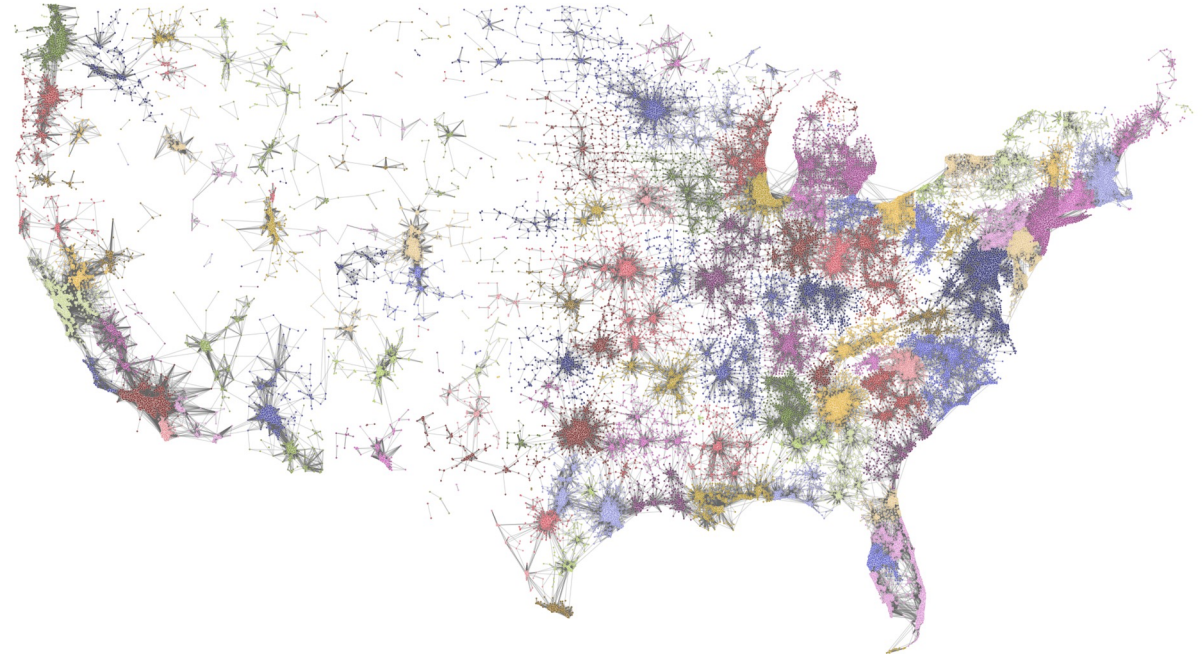
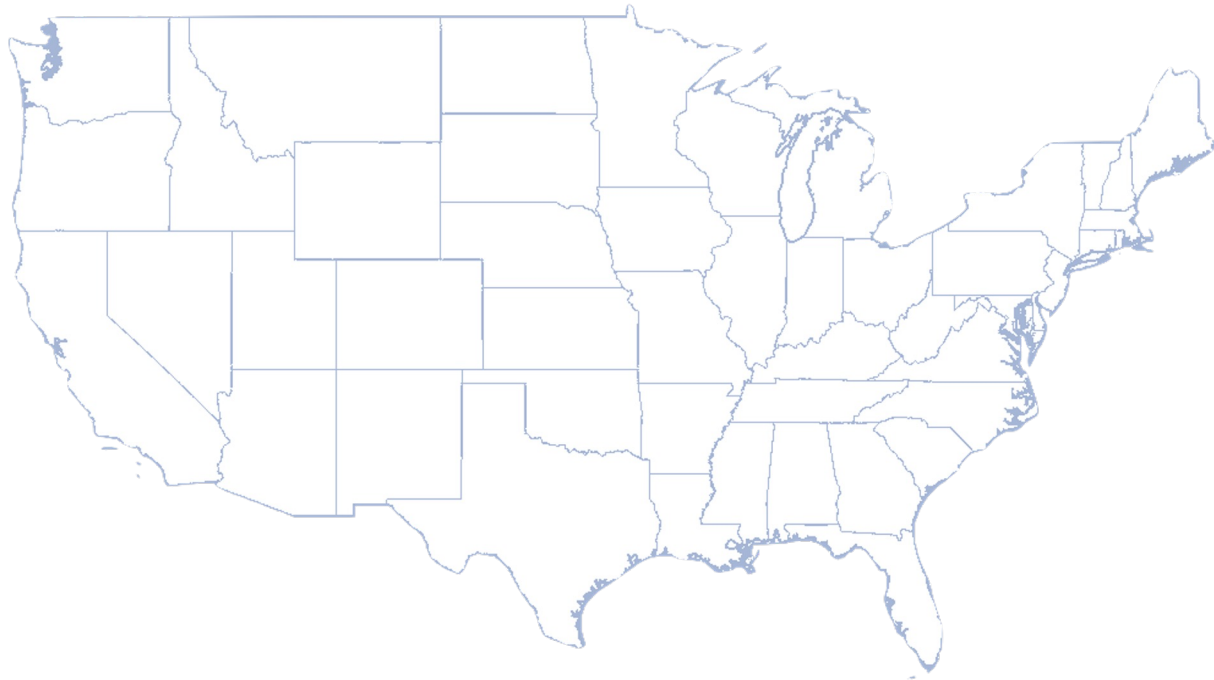


Commuter
flow

Which is better for comparing policy differences?



Which is better to model spread of diseases?

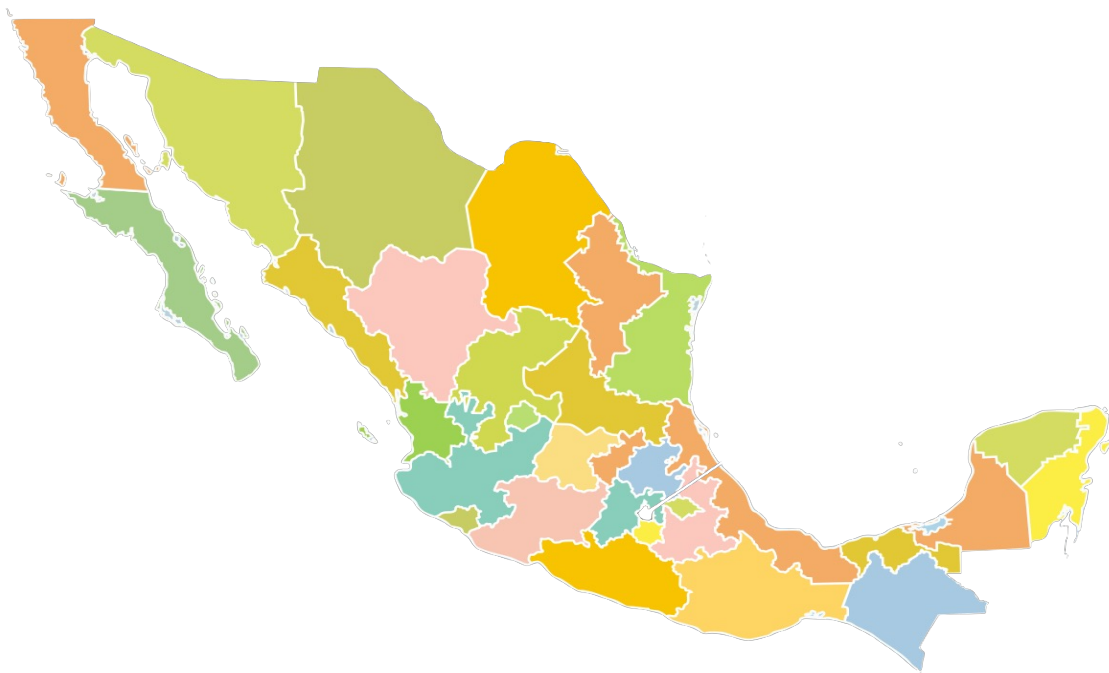


Networks, scale, and disease dynamics

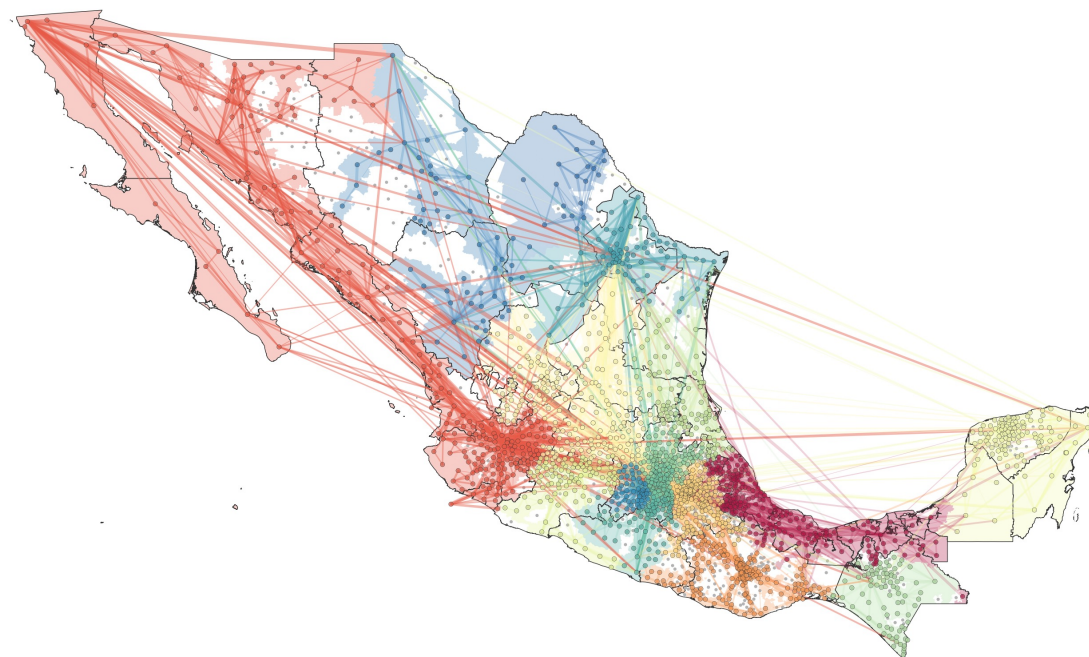
Spatial scales of COVID-19 transmission in Mexico

Brennan Klein^{*1,2}, Harrison Hartle¹, Munik Shrestha¹, Ana Cecilia Zenteno³,
David Barros Sierra Cordera⁴, José R. Nicolas-Carlock⁵, Ana I. Bento⁶,
Benjamin M. Althouse^{7,8}, Bernardo Gutierrez^{9,10,11},
Marina Escalera-Zamudio^{9,11}, Arturo Reyes-Sandoval^{12,13},
Oliver G. Pybus^{9,14,18}, Alessandro Vespignani^{1,2},
Jose Alberto Diaz-Quinonez^{*†15}, Samuel V. Scarpino^{*‡1,16,17}, and
Moritz U.G. Kraemer^{*§9,18}

Networks, scale, and disease dynamics

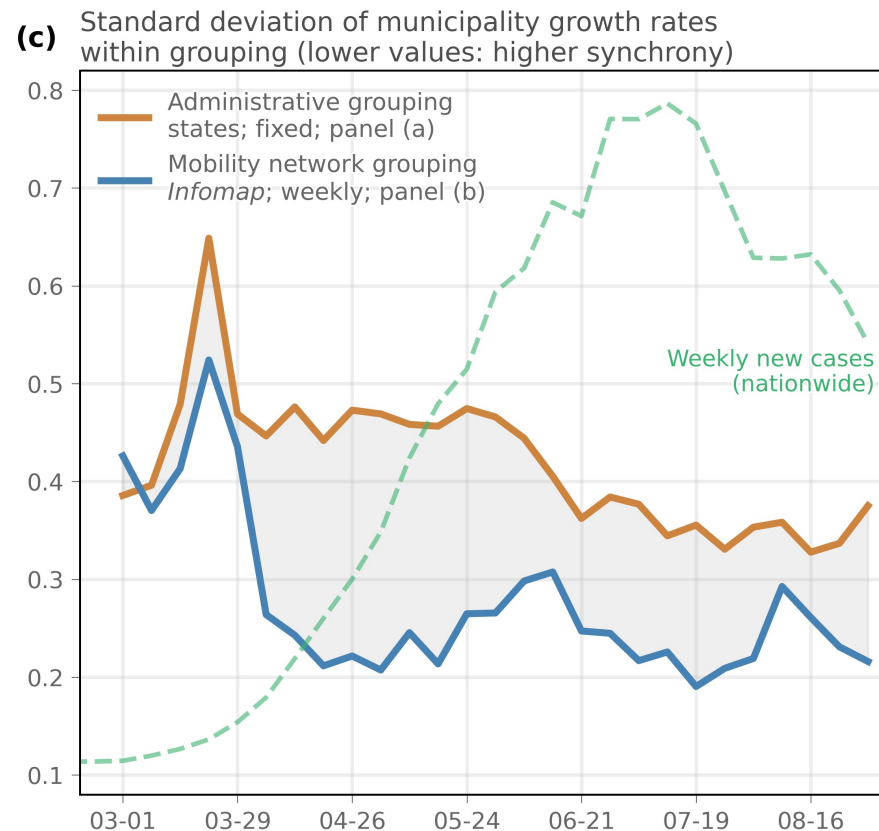
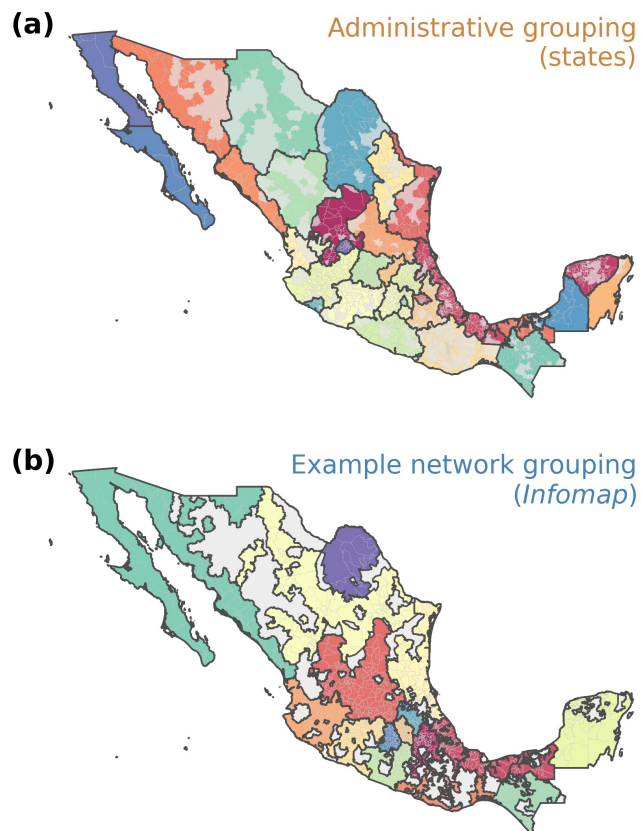


Policy operates on this scale
(**administrative**)

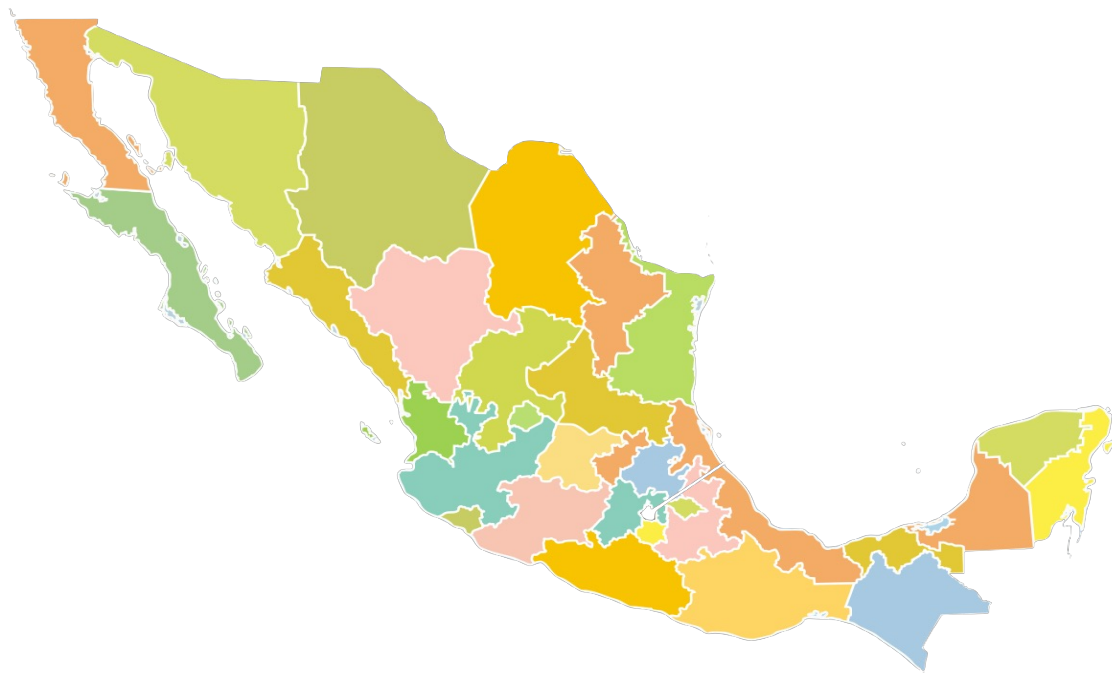


People interact at this scale
(**network**)

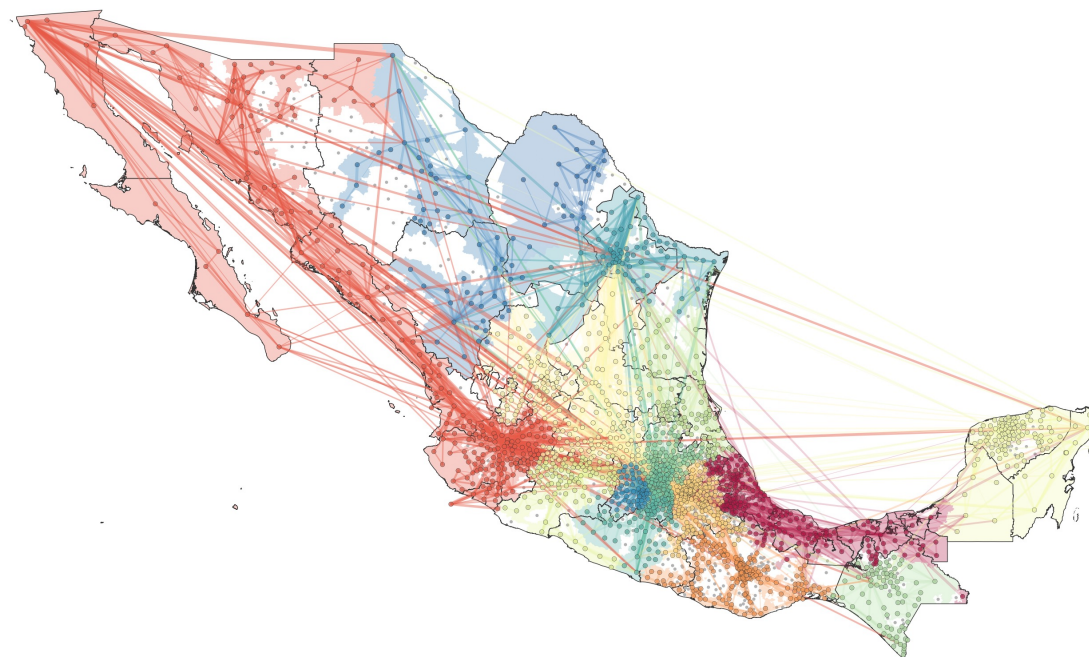
Networks, scale, and disease dynamics



Networks, scale, and disease dynamics



Higher within-community variance
of infection growth rates

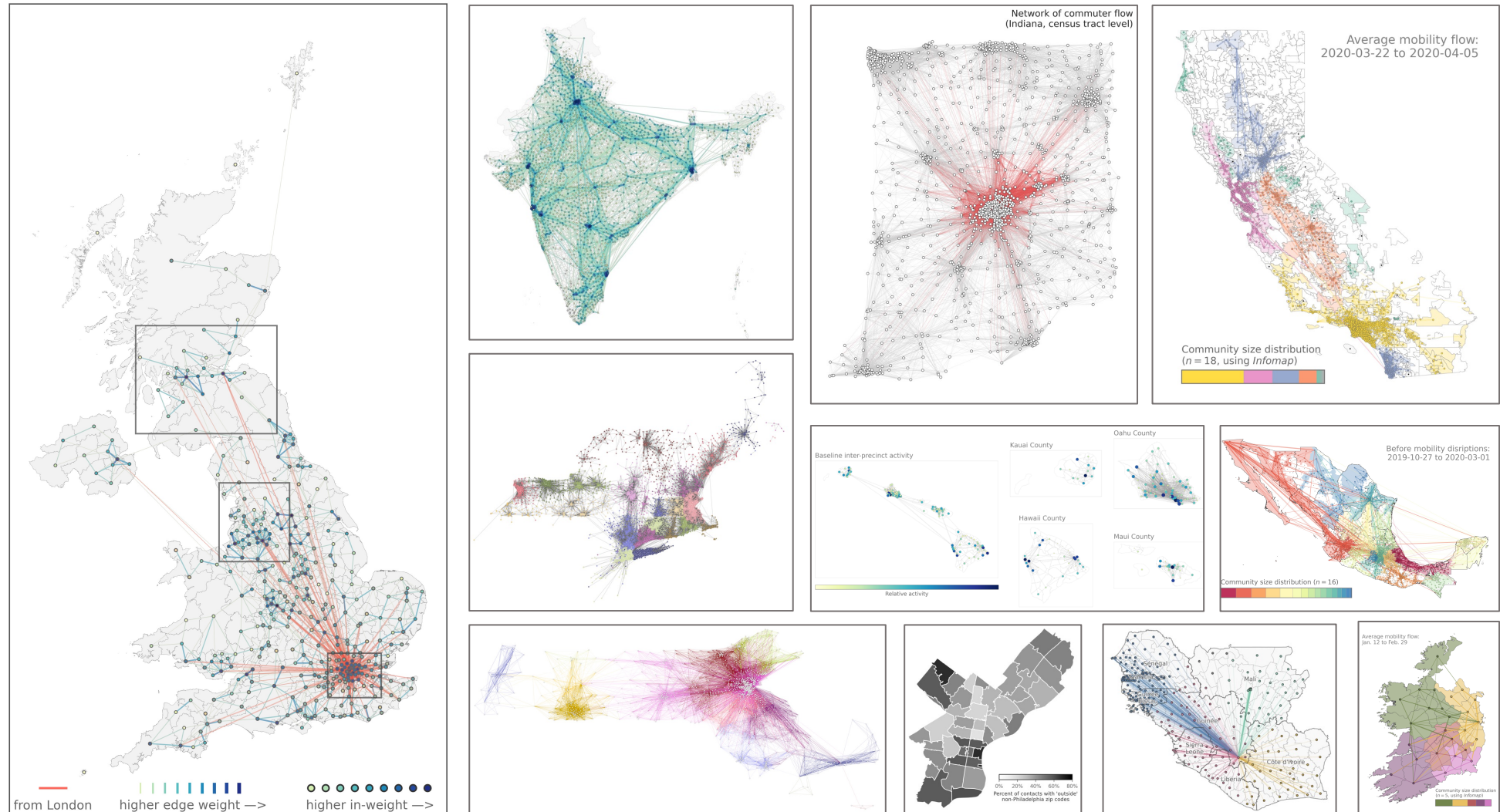


More *synchronized*
outbreaks

Today

1. Introduction
2. Complexity
 - i. Reconstruction & comparison*
 - ii. Informative network scales & representation*
3. Society
 - i. Mobile device data for disease modeling*
 - ii. Assorted COVID-19 projects*
4. Research vision and outlook

(The right) network data can improve modeling



But we don't always *need* networks...

RESEARCH

CORONAVIRUS

The effect of human mobility and control measures on the COVID-19 epidemic in China

Moritz U. G. Kraemer^{1,2,3*}, Chia-Hung Yang⁴, Bernardo Gutierrez^{1,5}, Chieh-Hsi Wu⁶, Brennan Klein⁴, David M. Pigott⁷, Open COVID-19 Data Working Group[†], Louis du Plessis¹, Nuno R. Faria¹, Ruoran Li⁸, William P. Hanage⁹, John S. Brownstein^{2,3}, Maylis Layan^{9,10}, Alessandro Vespignani^{4,11}, Huaiyu Tian¹², Christopher Dye¹, Oliver G. Pybus^{1,13*}, Samuel V. Scarpino^{4*}

RESEARCH

CORONAVIRUS

Spatiotemporal invasion dynamics of SARS-CoV-2 lineage B.1.1.7 emergence

Moritz U. G. Kraemer^{1*}, Verity Hill^{2,†}, Christopher Ruis^{3,†}, Simon Dellicour^{4,5,†}, Sumali Baja^{1,†}, John T. McCrone², Guy Baele⁵, Kris V. Parag⁶, Anya Lindström Battle⁷, Bernardo Gutierrez¹, Ben Jackson², Rachel Colquhoun², Aine O'Toole², Brennan Klein⁸, Alessandro Vespignani⁹, COVID-19 Genomics UK (COG-UK) Consortium[†], Erik Volz⁶, Nuno R. Faria^{1,6,9}, David M. Aanensen^{10,11}, Nicholas J. Loman¹², Louis du Plessis¹, Simon Cauchemez¹³, Andrew Rambaut^{2*}, Samuel V. Scarpino^{8,14,15*}, Oliver G. Pybus^{1,16*}

ARTICLE

<https://doi.org/10.1038/s41467-021-22521-5> OPEN

The effect of eviction moratoria on the transmission of SARS-CoV-2


Anjalika Nande^{1,7}, Justin Sheen^{2,7}, Emma L. Walters³, Brennan Klein^{4,5}, Matteo Chinazzi^{4,5}, Andrei H. Gheorghe¹, Ben Adlam¹, Julianna Shinnick², Maria Florencia Tejada², Samuel V. Scarpino⁴, Alessandro Vespignani^{4,5}, Andrew J. Greenlee³, Daniel Schneider³, Michael Z. Levy^{2,8,9} & Alison L. Hill^{1,6,8,9}

communications medicine

nature > communications medicine > articles > article

Article | [Open Access](#) | Published: 14 February 2023


Forecasting hospital-level COVID-19 admissions using real-time mobility data

Brennan Klein , Ana C. Zenteno , Daisha Joseph, Mohammadmehdi Zahedi, Michael Hu, Martin S. Copenhaver, Moritz U. G. Kraemer, Matteo Chinazzi, Michael Klompas, Alessandro Vespignani, Samuel V. Scarpino  & Hoijat Salmasian 

Clinical Infectious Diseases

CORRECTED PROOF









Examining the Robustness of 3 Versus 6 Feet of Physical Distancing in Schools: A Reanalysis of van den Berg et al

Brennan Klein , Daniel A Harris

PLOS DIGITAL HEALTH

RESEARCH ARTICLE

Higher education responses to COVID-19 in the United States: Evidence from the impacts of university policy





Brennan Klein , Nicholas Generous , Matteo Chinazzi^{1,2}, Zarana Bhadracha , Rishab Gunashekar^{1,4}, Preeti Kori , Bodian Li^{1,5}, Stefan McCabe , Jon Green , David Lazer¹, Christopher R. Marsicano , Samuel V. Scarpino^{1,9,10}, Alessandro Vespignani 

nature

nature > articles > article

Article | [Open Access](#) | Published: 19 April 2023

COVID-19 amplified racial disparities in the US criminal legal system

Brennan Klein , C. Brandon Ogbunugafor , Benjamin J. Schafer, Zarana Bhadracha, Preeti Kori, Jim Sheldon, Nitish Kaza, Arush Sharma, Emily A. Wang, Tina Eliassi-Rad, Samuel V. Scarpino  & Elizabeth Hinton 

under review (PNAS)

Spatial scales of COVID-19 transmission in Mexico

Brennan Klein^{1,2}, Harrison Hartle¹, Munik Shrestha¹, Ana Cecilia Zenteno³, David Barros Sierra Cordera⁴, José R. Nicolas-Carlock⁵, Ana I. Bento⁶, Benjamin M. Althouse^{7,8}, Bernardo Gutierrez^{9,10,11}, Marina Escalera-Zamudio^{9,11}, Arturo Reyes-Sandoval^{12,13}, Oliver G. Pybus^{9,14,18}, Alessandro Vespignani^{1,2}, Alberto Diaz Quiñonez^{*15}, Samuel V. Scarpino^{*11,16,17}, and Moritz U.G. Kraemer^{*9,18}

revision (PLOS Digital Health)

Characterizing collective physical distancing in the U.S. during the first nine months of the COVID-19 pandemic

Brennan Klein¹, Timothy LaRock¹, Stefan McCabe¹, Leo Torres¹, Lisa Friedland¹, Maciej Kos¹, Filippo Privitera¹, Brennan Lake¹, Moritz U.G. Kraemer², John S. Brownstein^{6,7}, Richard Gonzalez⁸, David Lazer¹, Tina Eliassi-Rad¹, Samuel V. Scarpino^{1,9,10}, Alessandro Vespignani^{1,3}, and Matteo Chinazzi^{1,2}

But we don't always *need* networks...

- Modeling collective physical distancing during the pandemic
 - Klein et al. (2020); Klein et al. (2020); Klein et al. (2022).
- Predicting hospital admissions 21 days out
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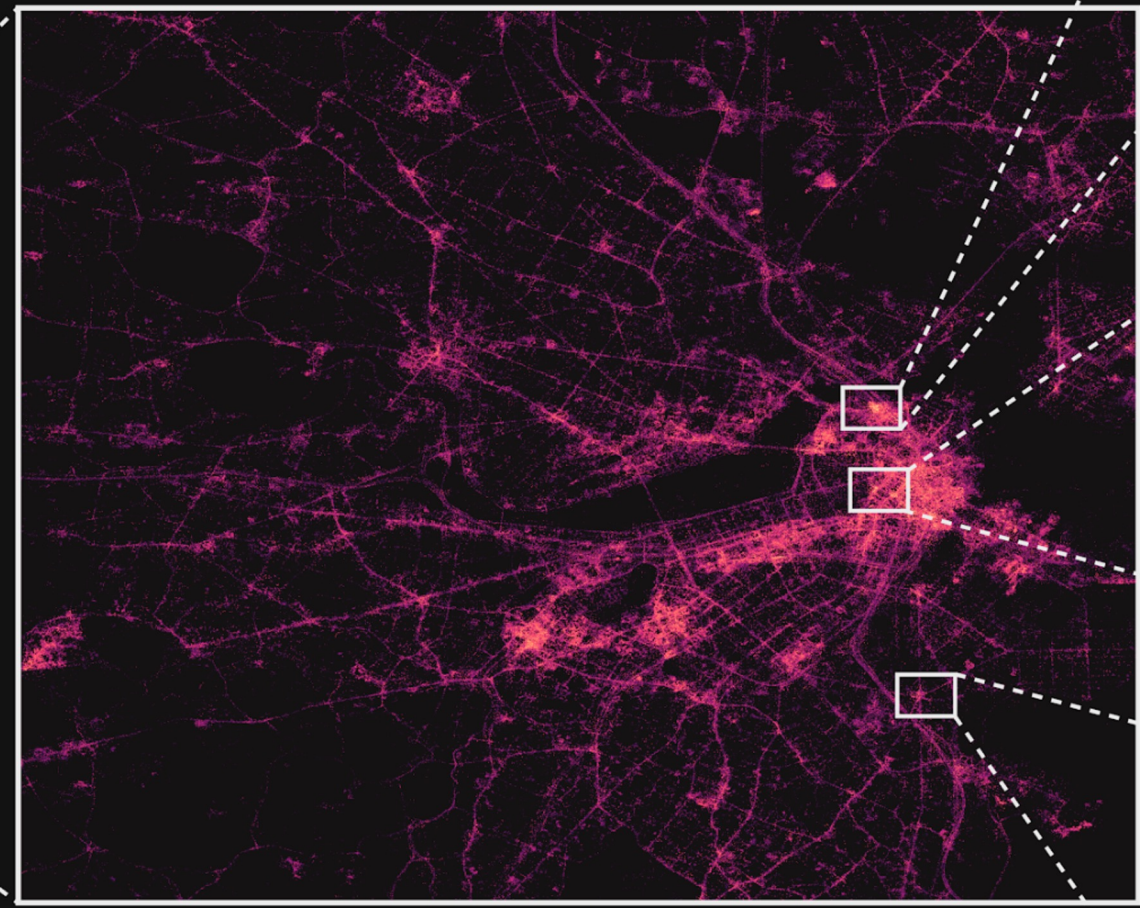
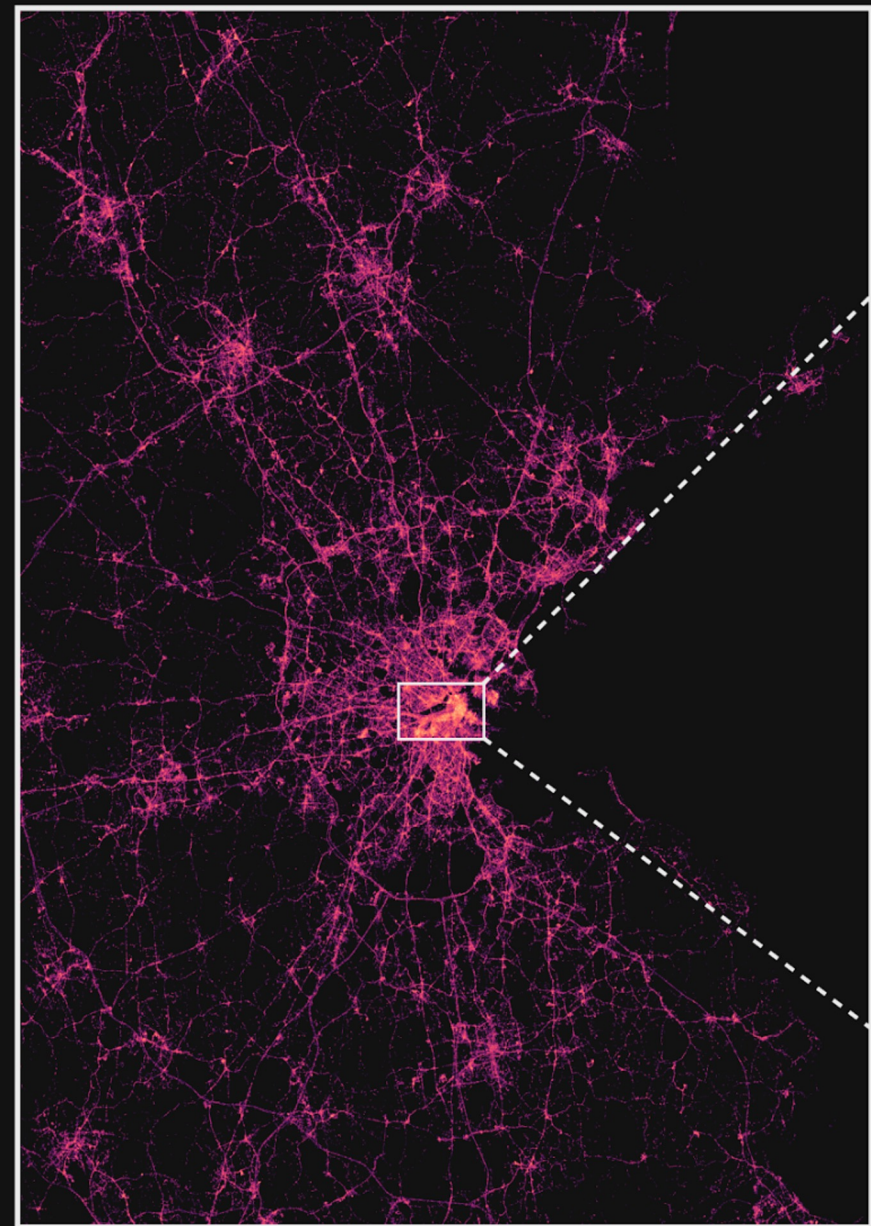
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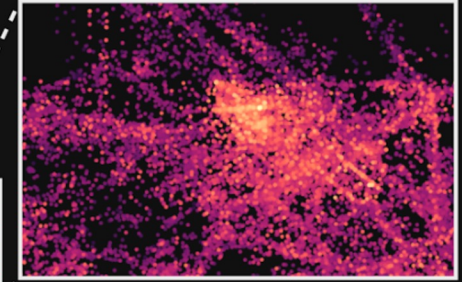
Data from Cuebiq Inc.

Through its [Data for Good](#) program, Cuebiq Inc. provides access to privacy-enhanced mobility data for academic / humanitarian initiatives.

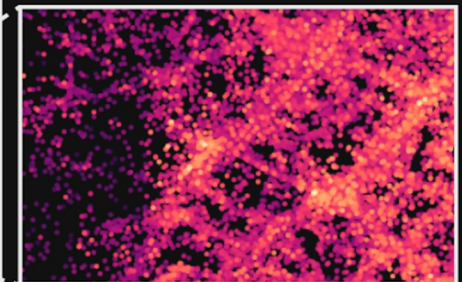
- Opt-in, first-party GPS data
- Obscured personal areas (home and work)
 - Up-leveled to the census block group
- Data from 40+ million total users
 - We select a panel of users from this
- High temporal and spatial resolution.
 - Heavy-tailed location accuracy



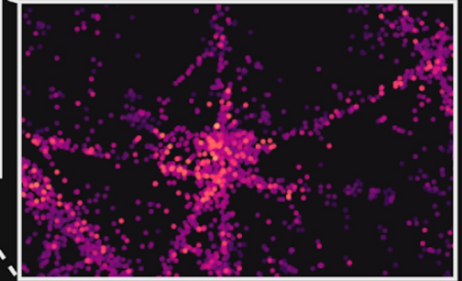
Large sporting arena



Indoor shops/restaurants

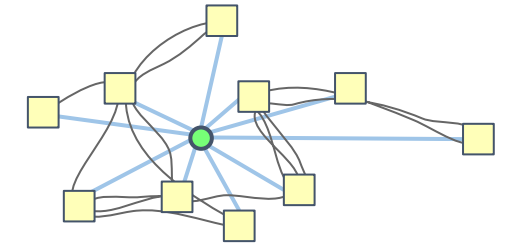


Large transit hub



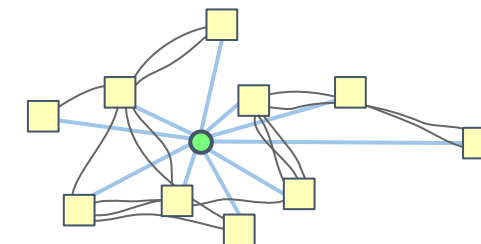
What can Cuebiq data tell us?

- Mobility behavior of users:
 - Measured through the radius of gyration¹
- Commuting patterns of users:
 - If one user visits their two (up-leveled) “personal areas” in a given 24hr period
- Estimated contact between users:
 - If users are close to each other within the same 5-minute time window (and thus, present an opportunity for transmission)



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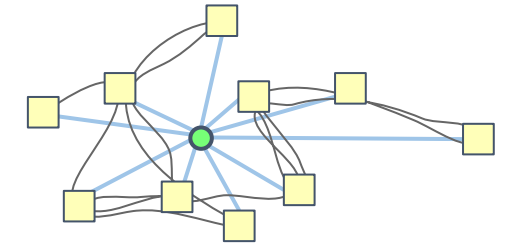


$$r = \sqrt{\frac{1}{n} \sum_{i=1}^n \|\vec{r}_i - \vec{r}_{cm}\|^2}$$

¹González, Hidalgo, & Barabási. Understanding individual human mobility patterns. *Nature* 453, 779–782 (2008).

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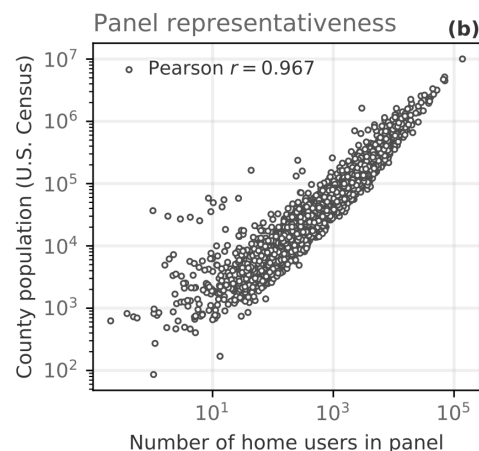
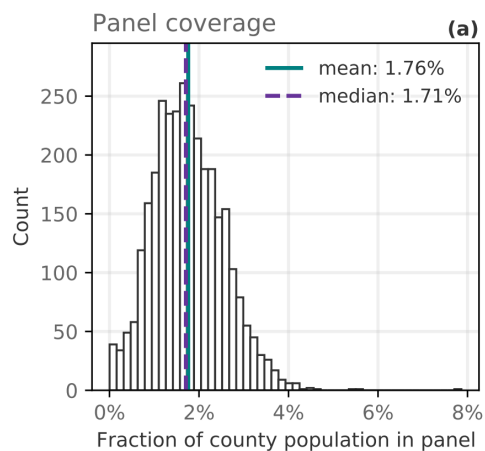
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What can Cuebiq data tell us?

But! We can't just take all 40+ million users and compute measures based on their data.

We need rigorous statistical controls to account for potential over/under sampling.

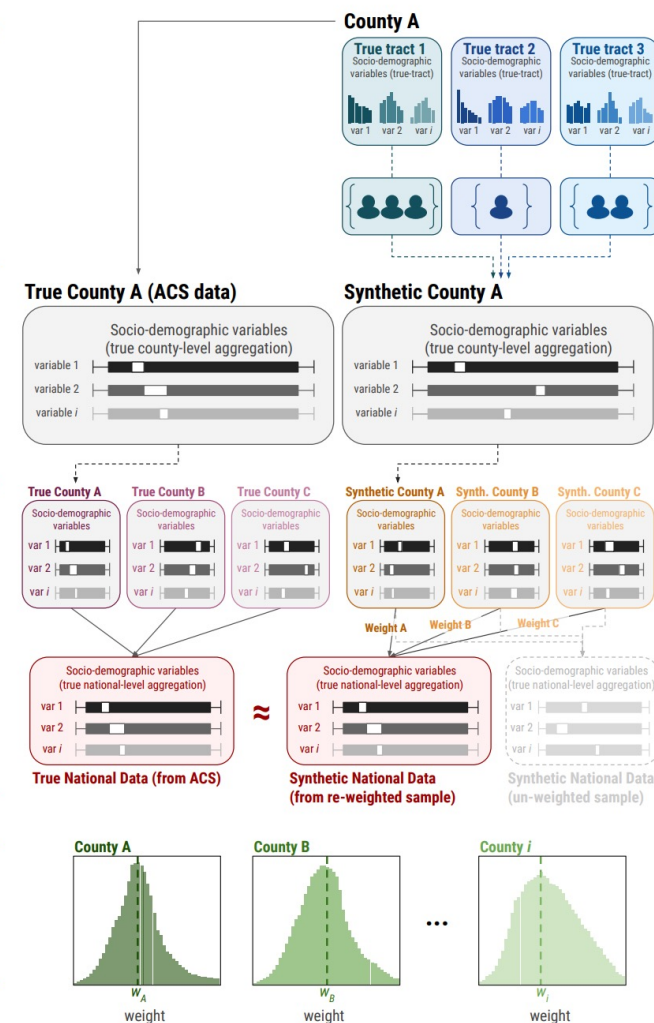


1. Sampling
Sample socio-demographic characteristics from the ACS Survey data.
For each tract, draw the same number of samples (*synthetic users*) as there are Cuebiq users.

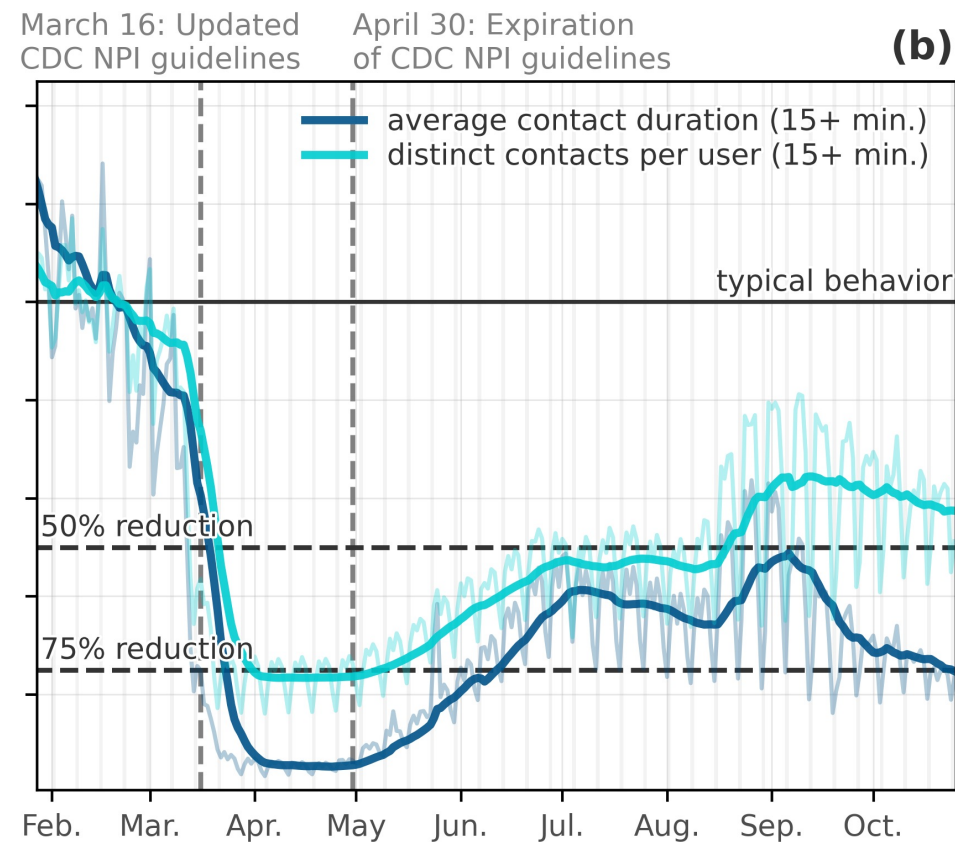
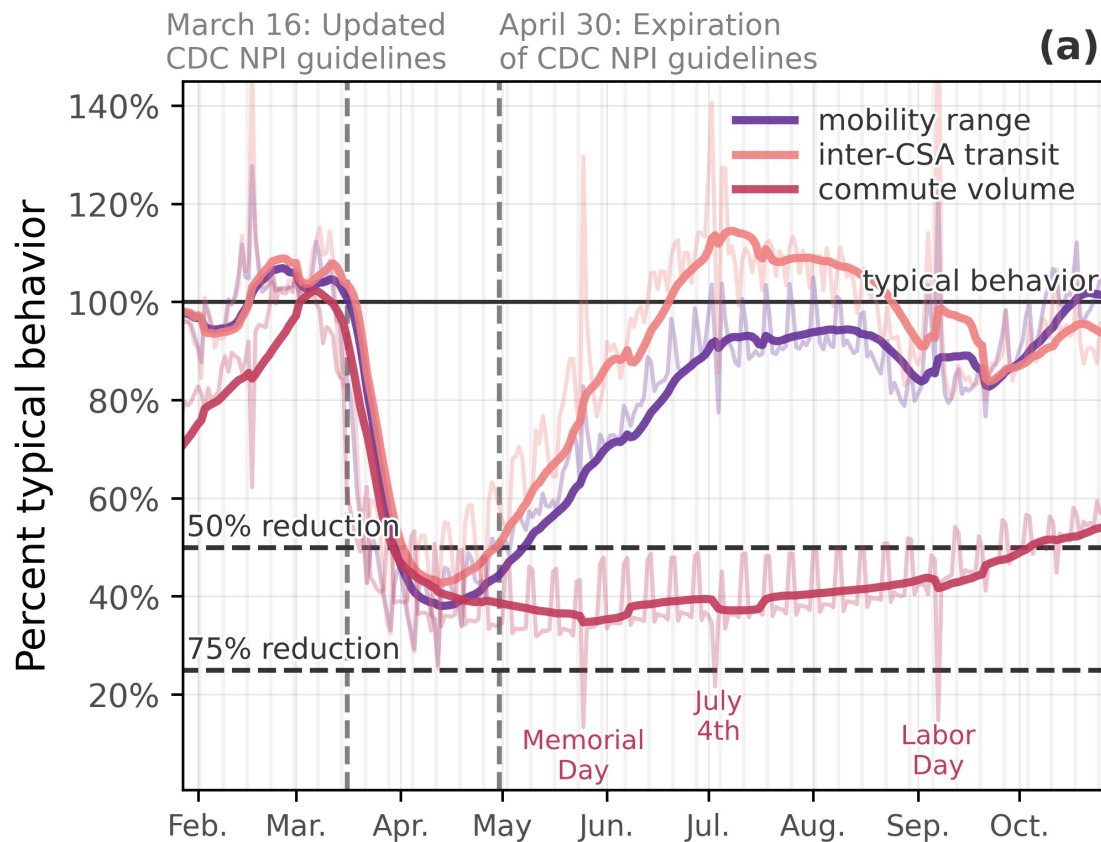
2. Aggregation
Aggregate samples on a county-level to create a *synthetic* county population.

3. Weights estimation
Estimate weights for each county using a generalized linear model (GLM logit) to reduce differences in the distributions of socio-demographics between synthetic and true counties when aggregated at a national level.

4. Bias reduction
Repeat steps 1-3 10,000 times for each county. Select average weight, w_i , for each county.



What can Cuebiq data tell us?



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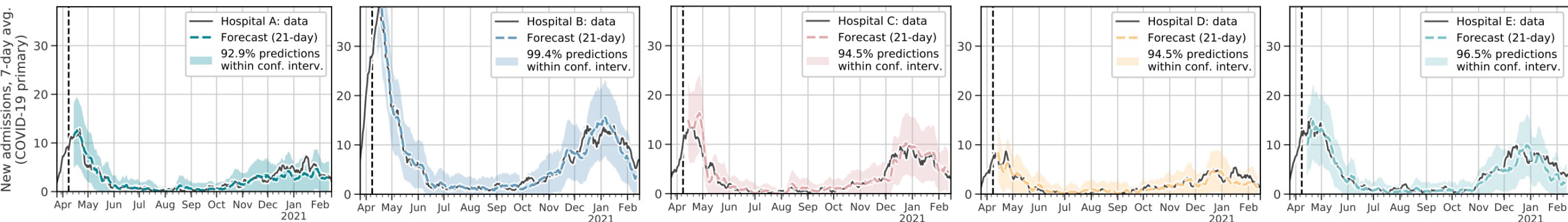
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Forecasting primary COVID-19 admissions

- Large healthcare network in Massachusetts approached us:
 - “We have pretty good in-house forecast accuracy for individual-hospital admissions **14 days out.**”
 - “What about 21 days out?”
- Our approach: combine exogenous mobility data with current admissions data in a *k-step ahead nonlinear autoregressive forecasting model*.

$$y(t+k) = f\left(y(t), y(t-1), \dots, y(t-p-1), \mathbf{X}(t), \mathbf{X}(t-1), \dots, \mathbf{X}(t-p-1)\right) + \epsilon(t)$$

k-days ahead forecast Target time series, $y(t)$ Exogenous (mobility) variables, $\mathbf{X}(t)$



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Sloppy data, sloppy policy

Effectiveness of 3 Versus 6 ft of Physical Distancing for Controlling Spread of Coronavirus Disease 2019 Among Primary and Secondary Students and Staff: A Retrospective, Statewide Cohort Study ^{FREE}

Polly van den Berg, Elissa M Schechter-Perkins, Rebecca S Jack, Isabella Epshtein, Richard Nelson, Emily Oster, Westyn Branch-Elliman ✉

Clinical Infectious Diseases, Volume 73, Issue 10, 15 November 2021, Pages 1871–1878, <https://doi.org/10.1093/cid/ciab230>

Published: 10 March 2021 [Article history](#) ▼

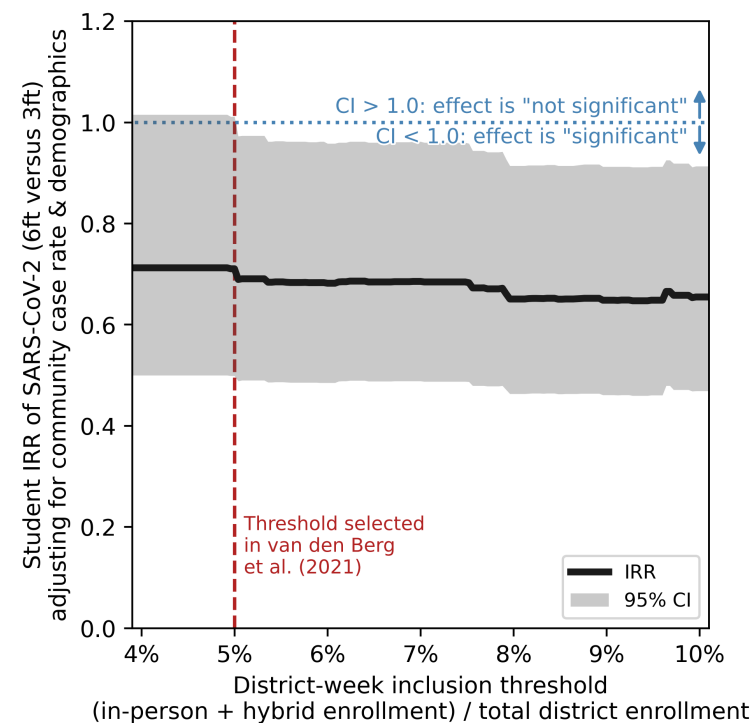
Received: 23 February 2021

Editorial decision: 08 March 2021

Published: 10 March 2021

Corrected and typeset: 27 April 2021

A correction has been published: *Clinical Infectious Diseases*, ciab1049, <https://doi.org/10.1093/cid/ciab1049>



Clinical Infectious Diseases

CORRECTED PROOF

Examining the Robustness of 3 Versus 6 Feet of Physical Distancing in Schools: A Reanalysis of van den Berg et al.

Brennan Klein ✉, Daniel A Harris

Clinical Infectious Diseases, ciac187, <https://doi.org/10.1093/cid/ciac187>

Published: 05 March 2022 [Article history](#) ▼

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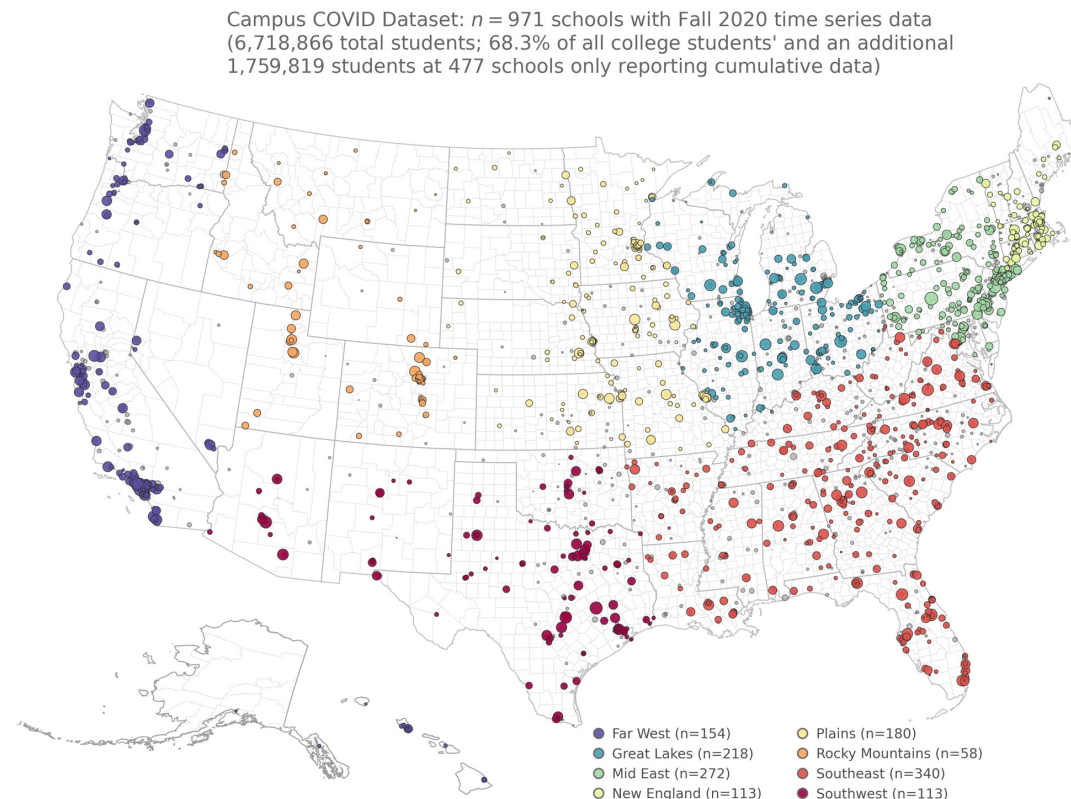
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Higher education COVID-19 policy

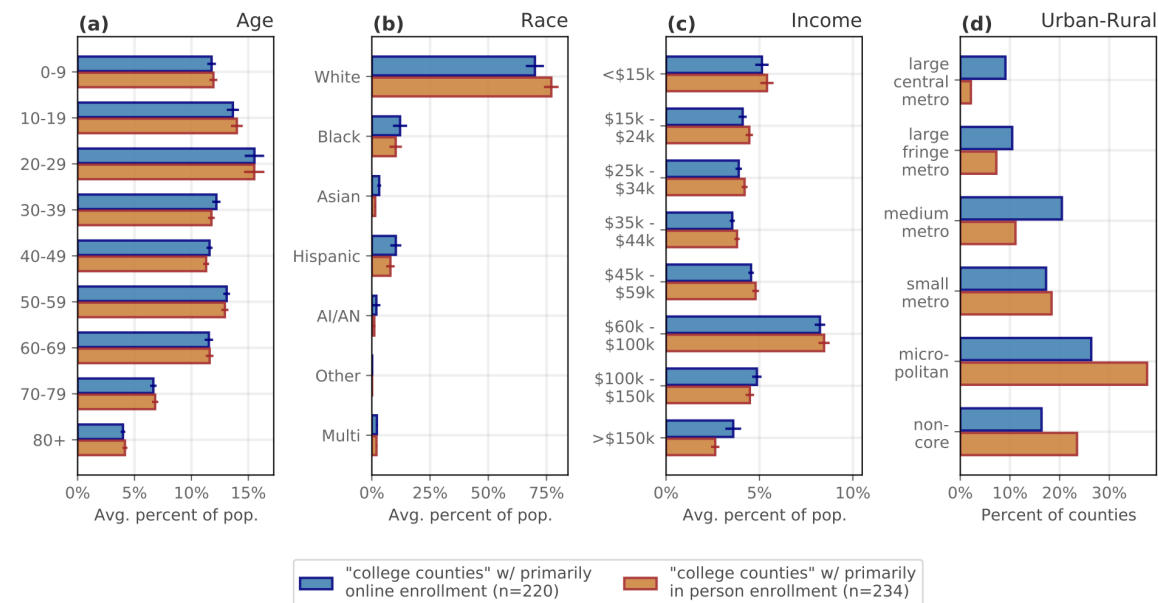
- Gather the largest dataset of reported tests and cases at institutes of higher education (IHE) for Fall 2020.

All data available on github
<https://github.com/jkbren/campus-covid/>



Higher education COVID-19 policy

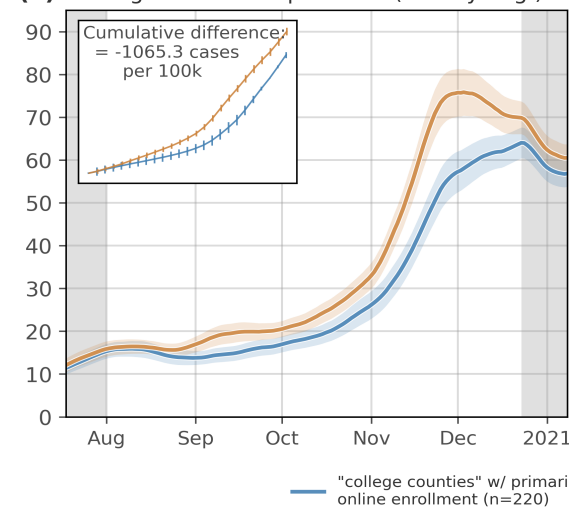
- Gather the largest dataset of reported tests and cases at institutes of higher education (IHE) for Fall 2020.
- Ask whether IHE policy (on-campus testing, online/in-person) has benefits for the surrounding community.
 - Create statistically-matched groups of college counties and compare reported cases / deaths at the county level.



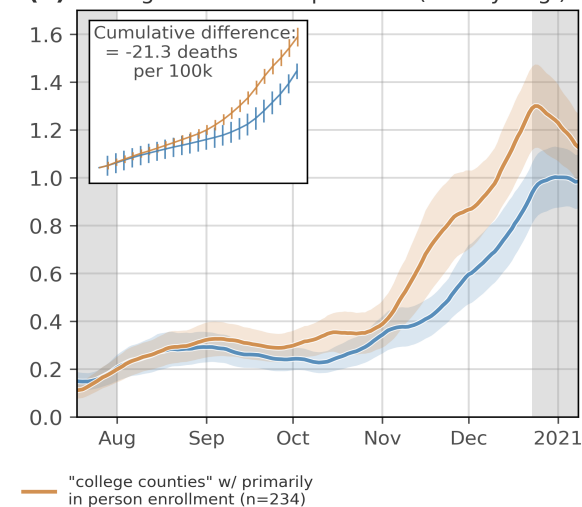
Higher education COVID-19 policy

- Gather the largest dataset of reported tests and cases at institutes of higher education (IHE) for Fall 2020.
- Ask whether IHE policy (on-campus testing, online/in-person) has benefits for the surrounding community.
 - Create statistically-matched groups of college counties and compare reported cases / deaths at the county level.
- Prudent campus policy has a positive impact on the surrounding county.

(a) Average new cases per 100k (21-day avg.)



(b) Average new deaths per 100k (21-day avg.)



"college county": counties where total IHE enrollment is at least 3.68% of the total county population

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- 4. Problems that excite me lately**

Someday (soon)

1. Higher-order dynamical processes
2. New (principled) graph distances
3. Connectomics of artificial intelligence
4. What is a *typical* graph in model?
5. Individuation and the network science of network mergers
6. The free energy principle and active inference

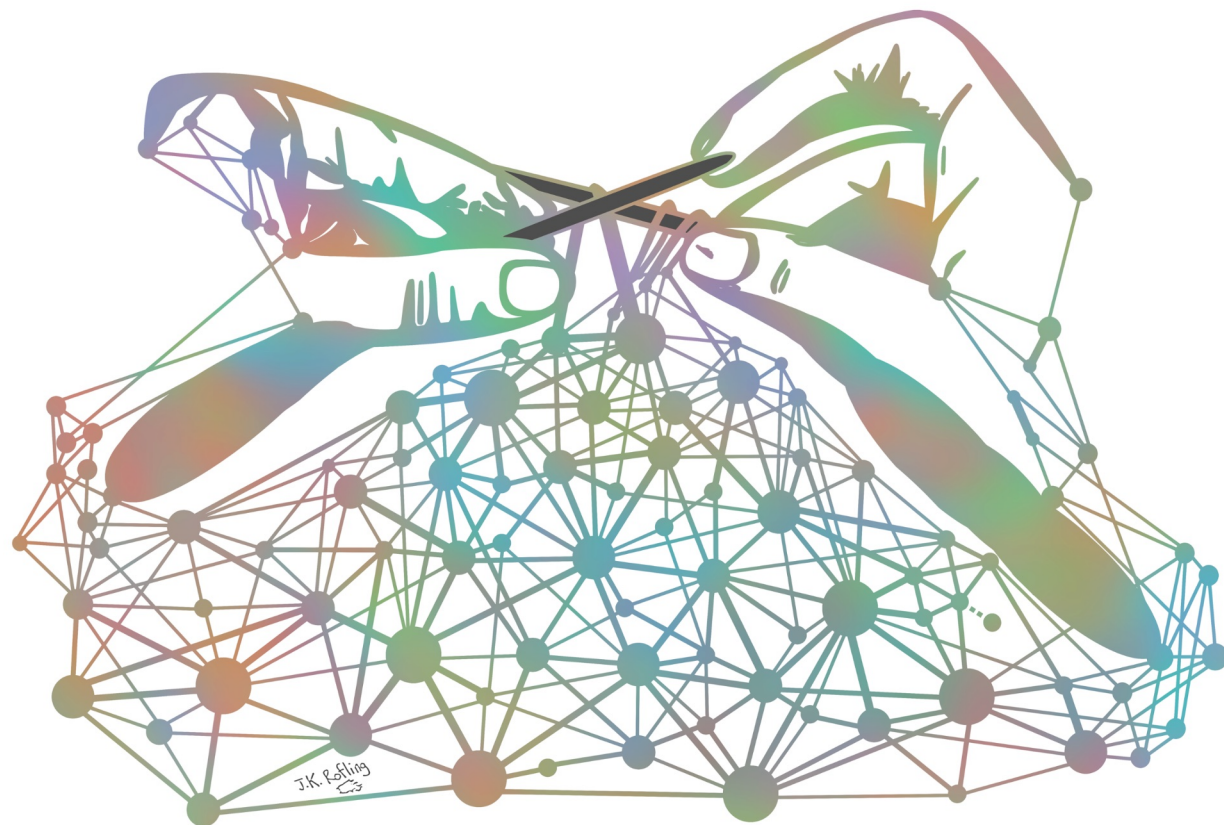


René Thom

“At a time when so many scholars calculate throughout the world, is it not desirable for some, if they can, to dream?”

— René Thom

Au moment où tant de savants calculent de par le monde, n'est-il pas souhaitable que d'aucuns, s'ils le peuvent, rêvent?



Questions, contact, follow-up

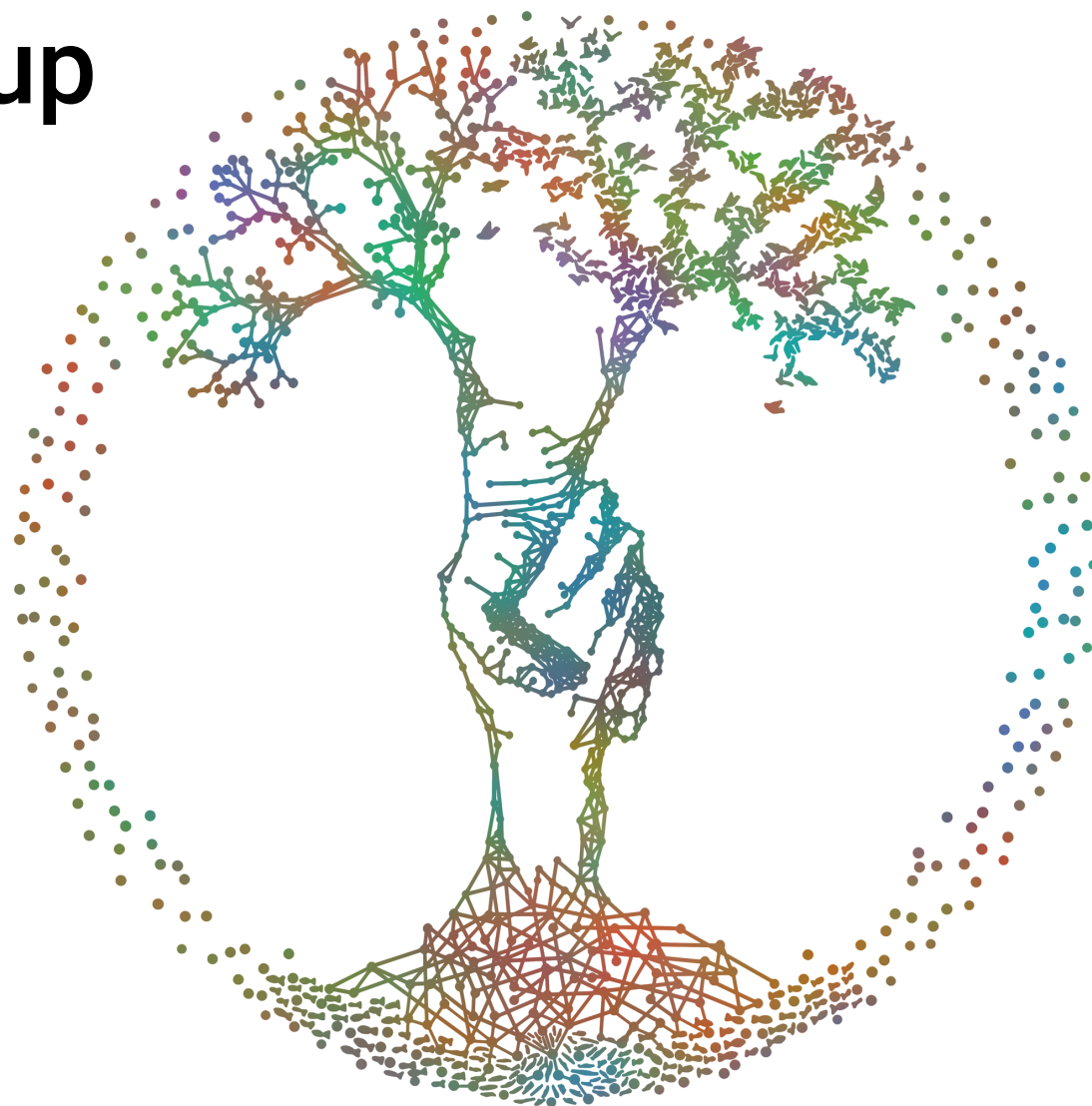
A network science for complexity & society

Brennan Klein

Northeastern University
Network Science Institute

b.klein@northeastern.edu || [@jkbren](https://twitter.com/jkbren) || brennanklein.com

France Regional Conference on Complex Systems
René Thom Young Researcher Award



Northeastern University
Network Science Institute



Complexity & Society Lab
@ Northeastern University



HARVARD
UNIVERSITY



Hutchins Center
for African &
African American
Research

VERSES



FRCCS