

A TOPOLOGICAL VIEW OF COLLECTIVE BEHAVIOR

Chad Topaz (Williams College)
NSF DMS-1813752

MY FIRST CONFERENCE

Fifth SIAM Conference on APPLICATIONS of DYNAMICAL SYSTEMS

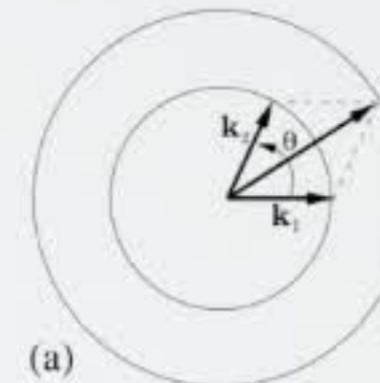
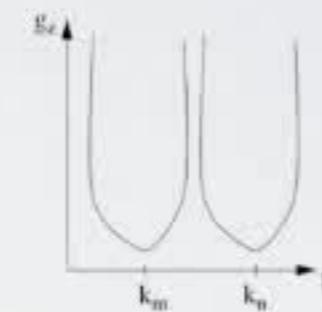
May 23 - 27, 1999

Snowbird Ski and Summer Resort
Snowbird, Utah

Q: Something something co-dimension blah blah blah?

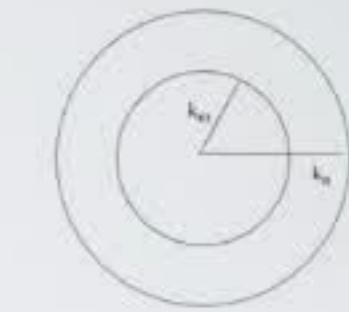
A: $\bar{\backslash}(\times)\bar{/}$

Pattern formation background:
Resonant triad interactions



$$k_3 = k_1 + k_2$$

(a)



$$k_3 = k_1 - k_2$$

(b)

Example (a) $\rightarrow h = z_1 e^{i\vec{k}_1 \cdot \vec{x}} + z_2 e^{i\vec{k}_2 \cdot \vec{x}} + z_3 e^{i\vec{k}_3 \cdot \vec{x}} + c.c.$

$$\dot{z}_1 = \lambda z_1 + \bar{z}_2 z_3 + (a|z_1|^2 + b|z_2|^2 + c|z_3|^2)z_1$$

$$\dot{z}_2 = \lambda z_2 + \bar{z}_1 z_3 + (a|z_2|^2 + b|z_1|^2 + c|z_3|^2)z_2$$

$$\dot{z}_3 = \mu z_3 + \bar{z}_1 z_2 + (d|z_1|^2 + d|z_2|^2 + e|z_3|^2)z_3$$

$$\ddot{z}_1 = \eta z_1 + \bar{z}_1 \dot{z}_3 + (\alpha|\dot{z}_1|_3 + \beta|\dot{z}_2|_3 + \gamma|\dot{z}_3|_3)z_1$$

$$\ddot{z}_2 = \eta z_2 + \bar{z}_2 \dot{z}_3 + (\alpha|\dot{z}_2|_3 + \beta|\dot{z}_1|_3 + \gamma|\dot{z}_3|_3)z_2$$

$$\ddot{z}_3 = \eta z_3 + \bar{z}_3 \dot{z}_3 + (\alpha|\dot{z}_3|_3 + \beta|\dot{z}_1|_3 + \gamma|\dot{z}_2|_3)z_3$$

TOPOLOGICAL DATA ANALYSIS

Persistent Homology – a Survey

Herbert Edelsbrunner and John Harer

ABSTRACT. Persistent homology is an algebraic tool for measuring topological features of shapes and functions. It casts the multi-scale organization we frequently observe in nature into a mathematical formalism. Here we give a record of the short history of persistent homology and present its basic concepts. Besides the mathematics we focus on algorithms and mention the various connections to applications, including to biomolecules, biological networks, data analysis, and geometric modeling.

TDA AT WORK



PHYSICAL REVIEW E 94, 032909 (2016)

ARTICLE

Received 15 Aug 2014; revised 17 Dec 2015; accepted 17 Jan 2016; published 1 Mar 2016
J. Math. Biol. (2018) 76:1559–1587
<https://doi.org/10.1007/s00285-017-1184-8>

Topological data analysis of contagion maps for examining spreading processes on networks
Chad Giomi^a
^aDepartment of Physics, University of Pennsylvania, Philadelphia, Pennsylvania, USA

Mathematical Biology

Dane Taylor^{1,2}, Florian Klimm^{3,4}, Mason A. Porter^{5,6} & Peter J. Mucha⁷
A top SPEC
Department of Biomedical Engineering
David I.

Persistent Homology: An Introduction and a New Text Representation for Natural Language Processing

Xiaojin Zhu

Department of Computer Sciences, University of Wisconsin-Madison
Madison, Wisconsin, USA 53706

jerryzhu@cs.wisc.edu

AMS NOTICES, MAY 2019

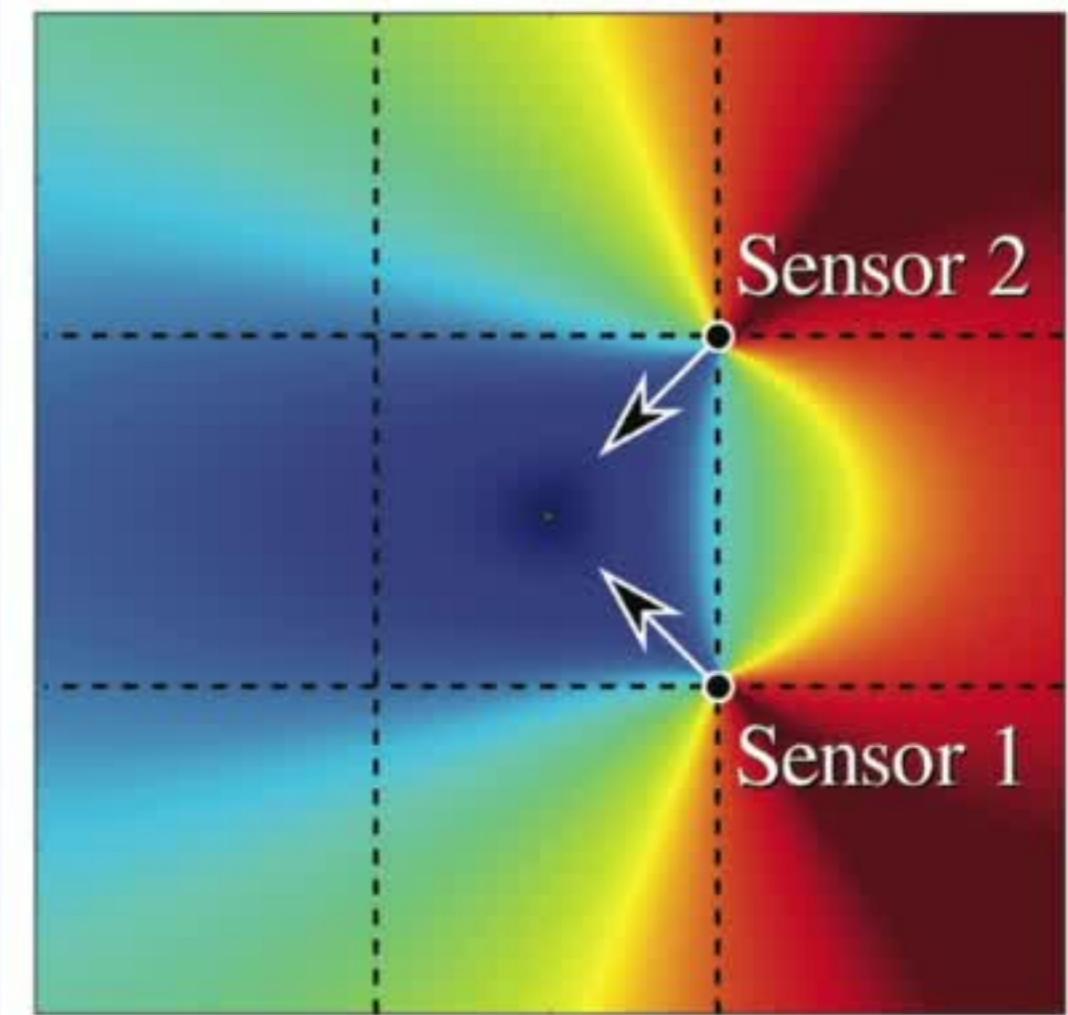
Topological Time Series
Analysis



Jose A. Perea

1026 A. 2019

Hunting for Foxes with Sheaves

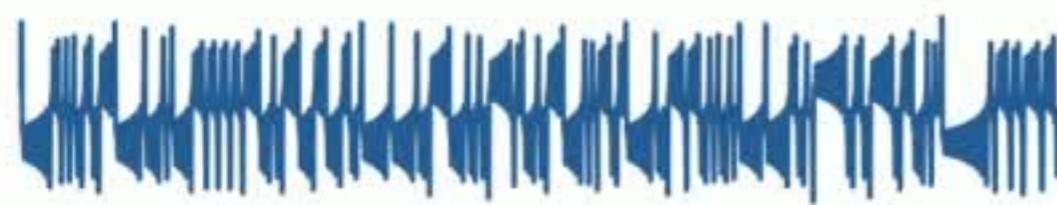


Michael Robinson

1026 A. 2019

AMS NOTICES, MAY 2019

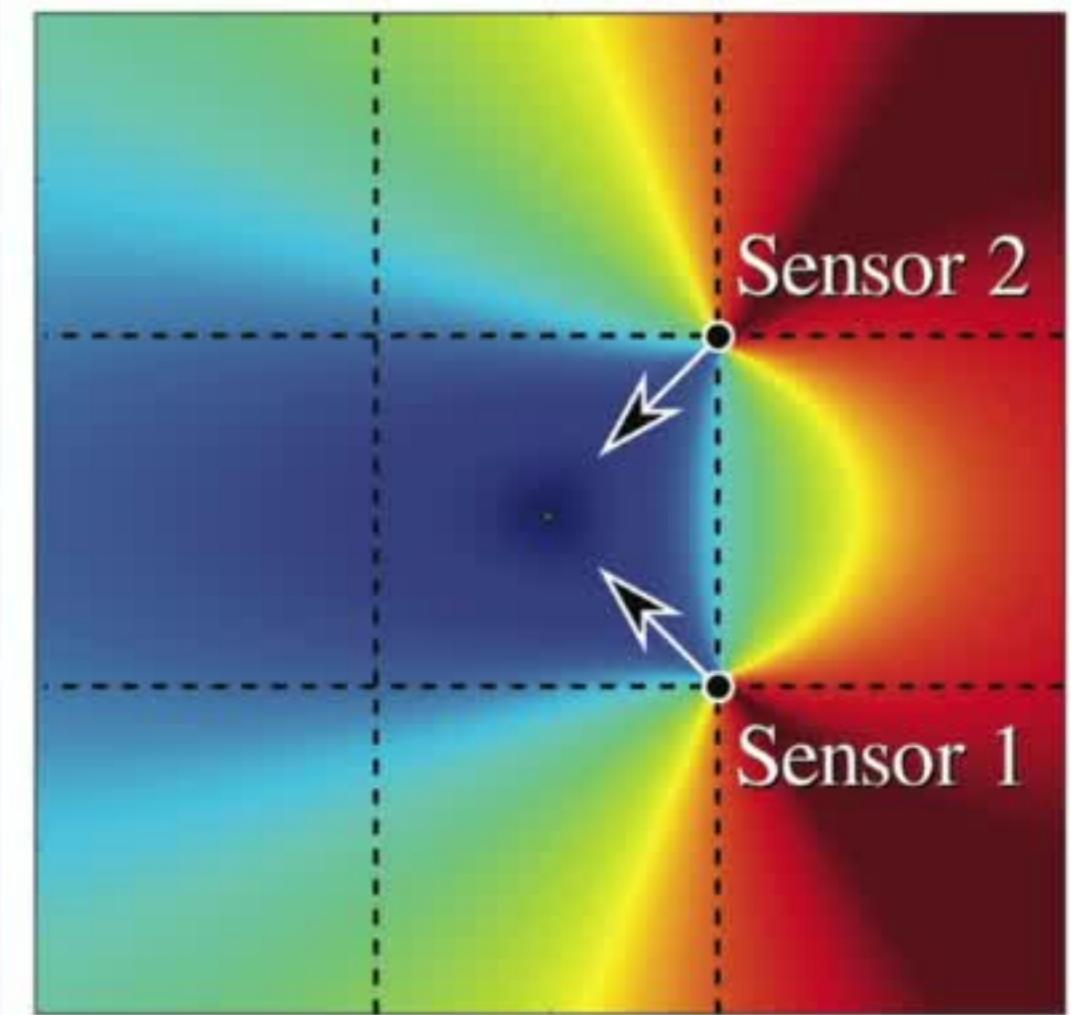
Topological Time Series
Analysis



Jose A. Perea

1026 A. 1939

Hunting for Foxes with Sheaves

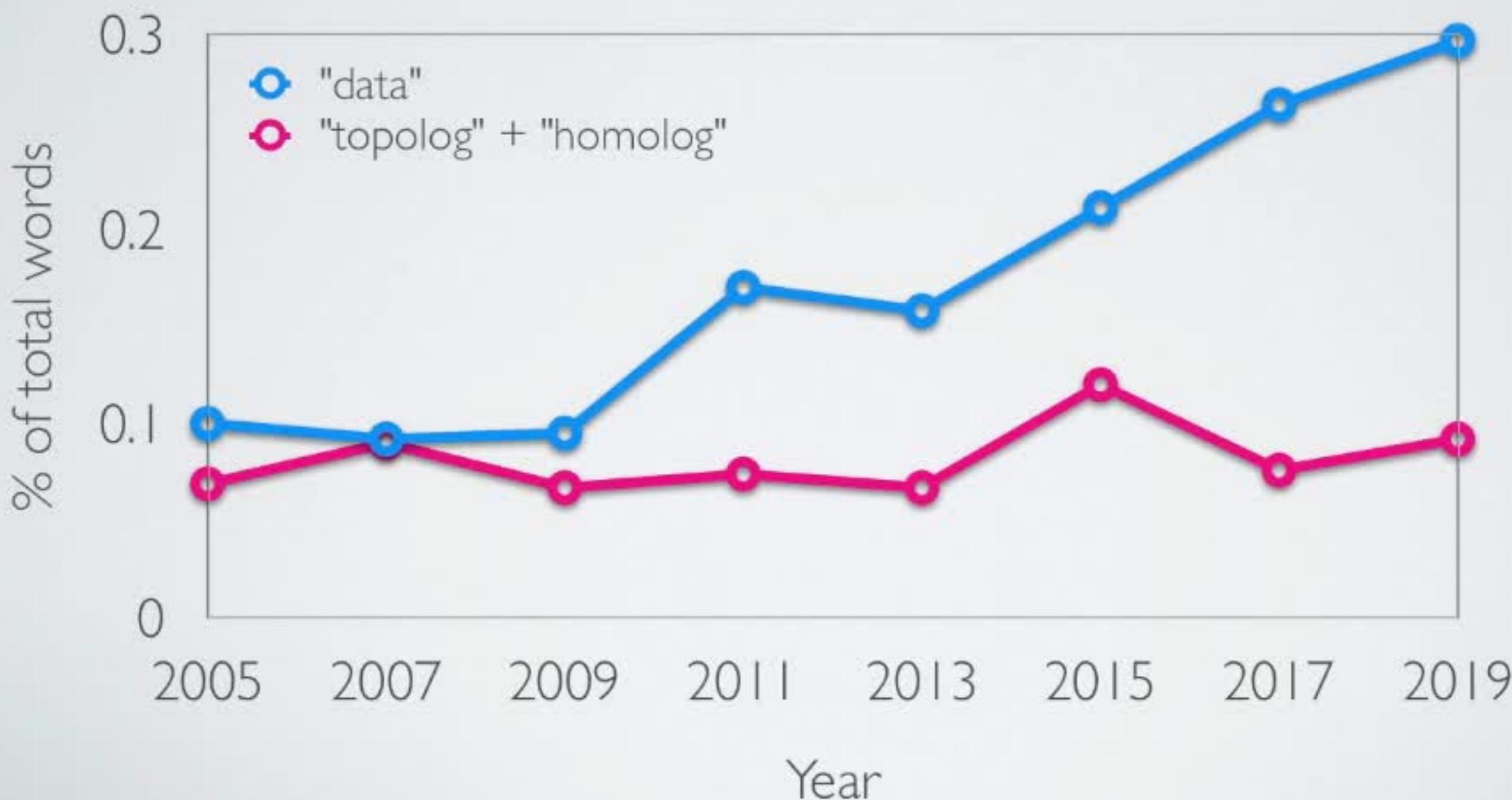


Michael Robinson

1026 A. 1939

TOPOLOGY AT SIAM DS

Occurrences of specific words as percentage of total words in SIAM DS abstract booklet



BACK TO 2009

Theorem 2.1 (structure) If D is a PID, then every finitely generated D -module is isomorphic to a direct sum of cyclic D -modules. That is, it decomposes uniquely into the form

$$D^\beta \oplus \left(\bigoplus_{i=1}^m D/d_i D \right), \quad (1)$$

for $d_i \in D$, $\beta \in \mathbb{Z}$, such that $d_i | d_{i+1}$. Similarly, every graded module M over a graded PID D decomposes uniquely into the form

$$\left(\bigoplus_{i=1}^n \Sigma^{\alpha_i} D \right) \oplus \left(\bigoplus_{j=1}^m \Sigma^{\gamma_j} D/d_j D \right), \quad (2)$$

where $d_j \in D$ are homogeneous elements so that $d_j | d_{j+1}$, $\alpha_i, \gamma_j \in \mathbb{Z}$, and Σ^α denotes an α -shift upward in grading.

BACK TO 2009

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TOPOLOGY I, CHAD 0

Theorem 2.1 (structure) If \mathbb{R}^n is a \mathbb{R}^m then every \mathbb{R}^m is \mathbb{R}^n to a \mathbb{R}^m of \mathbb{R}^n . That is, it \mathbb{R}^m into the form

$$\boxed{\text{[Redacted]}} \quad (1)$$

for \mathbb{R}^m such that \mathbb{R}^m . Similarly, every \mathbb{R}^m over a \mathbb{R}^n into the form

$$\boxed{\text{[Redacted]}} \quad (2)$$

where \mathbb{R}^m are \mathbb{R}^m so that \mathbb{R}^m denotes an \mathbb{R}^m in \mathbb{R}^n .

MY OUTLOOK, CHANGED

“Think of a bird flock as a bunch of data points moving in six-dimensional position-velocity space. At every moment in time, build an object by forming connections between points that are close. Ask about the topology of that object, and how it changes as your notion of closeness changes.”

- Lori Ziegelmeier, 2014
Assistant Professor of Mathematics
Macalester College



(T)APOLOGY

Main idea: Betti numbers of simplicial complexes are useful topological invariants that you can calculate with linear algebra.

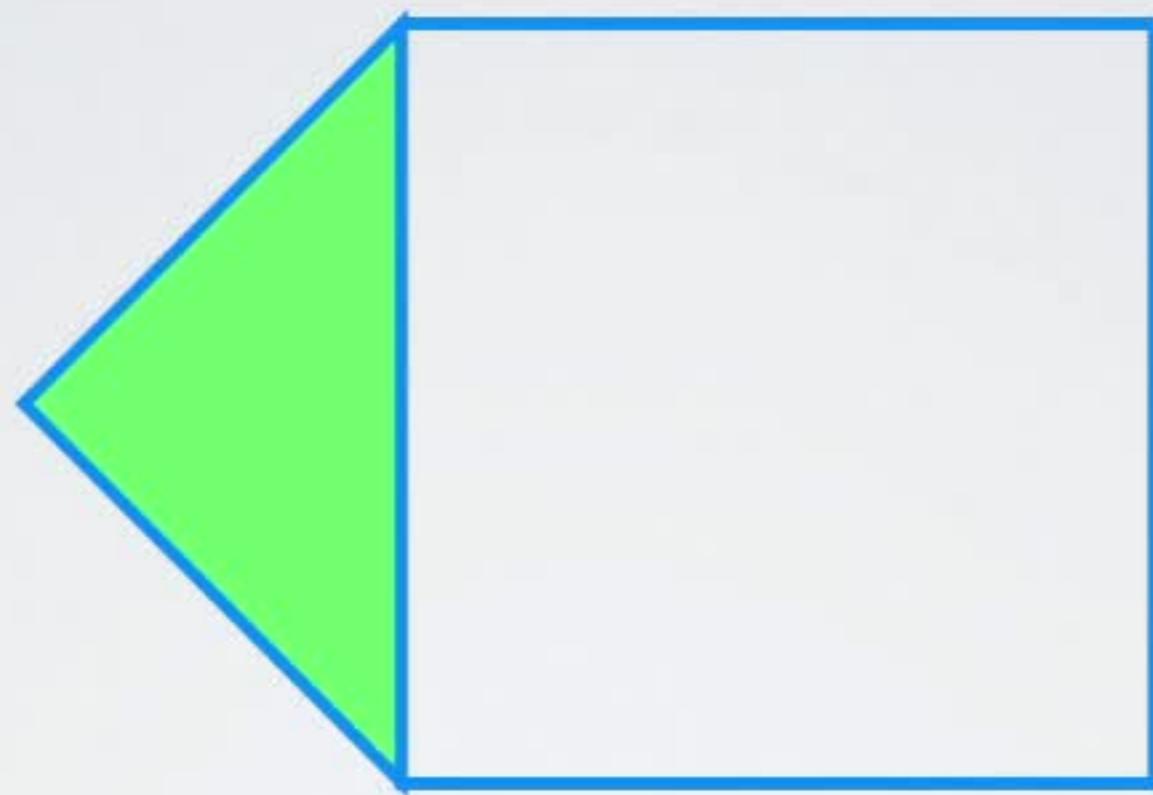
TOPOLOGY



TOPOLOGY



SIMPLICIAL COMPLEXES



$$\# \text{ loops } b_1 = 3 - l(\text{homologous}) - l(\text{boundary}) = 1$$

$$\# \text{ loops } b_1 = \dim \text{Ker } \partial_{1 \rightarrow 0} - \dim \text{Im } \partial_{2 \rightarrow 1}$$

IT'S LINEAR ALGEBRA

Chad's Self-Help
Homology Tutorial

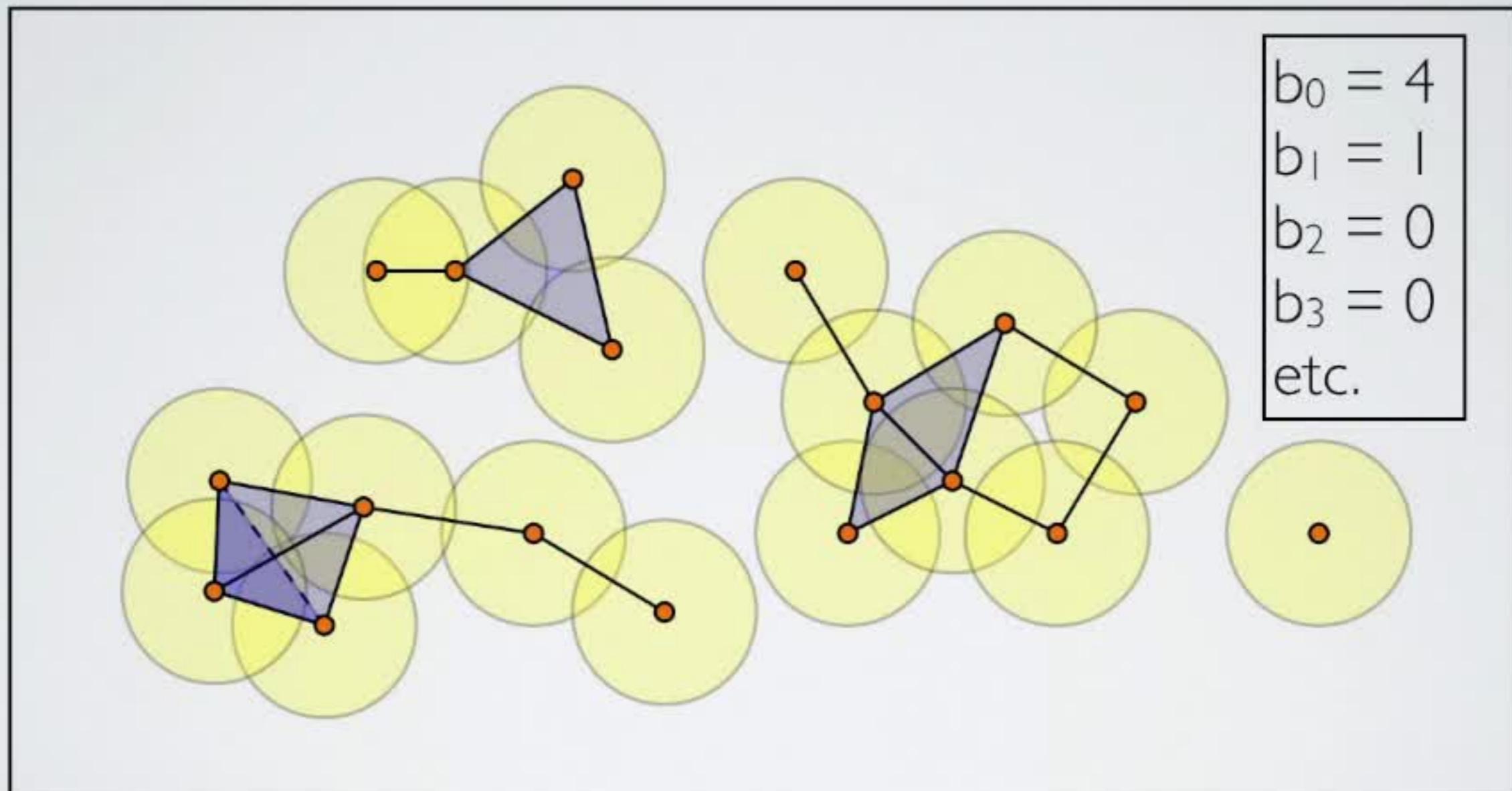
For the Simple(x)-Minded

A Full-Color Extravaganza

If you don't love topology
then how in the hell are you
gonna love yourself?

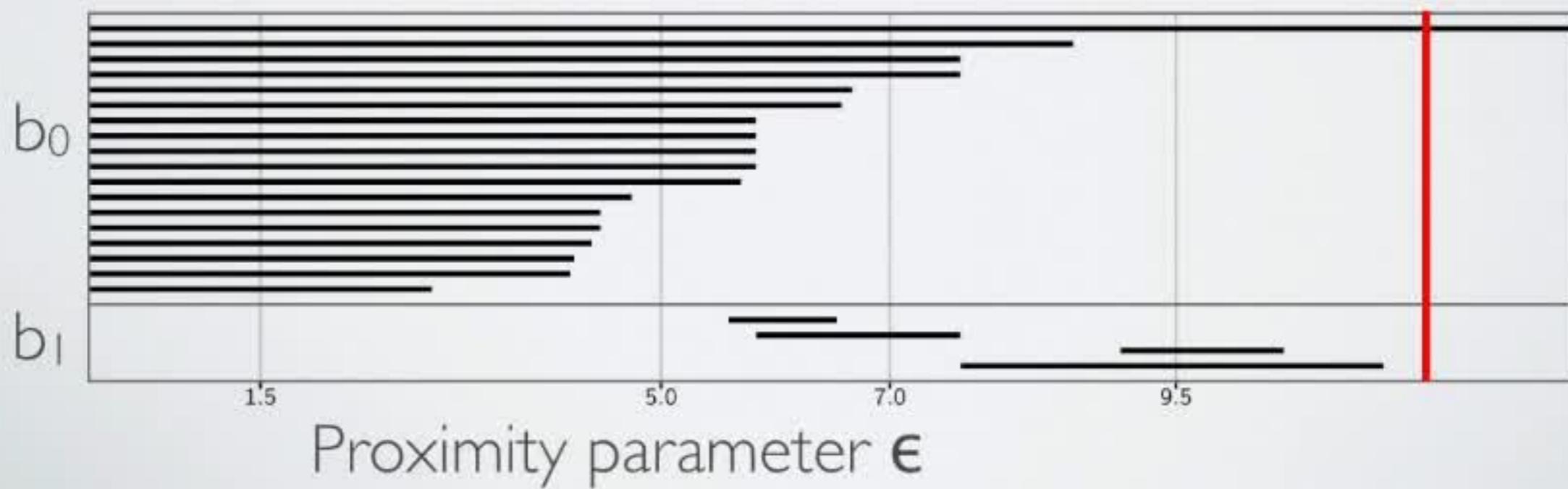
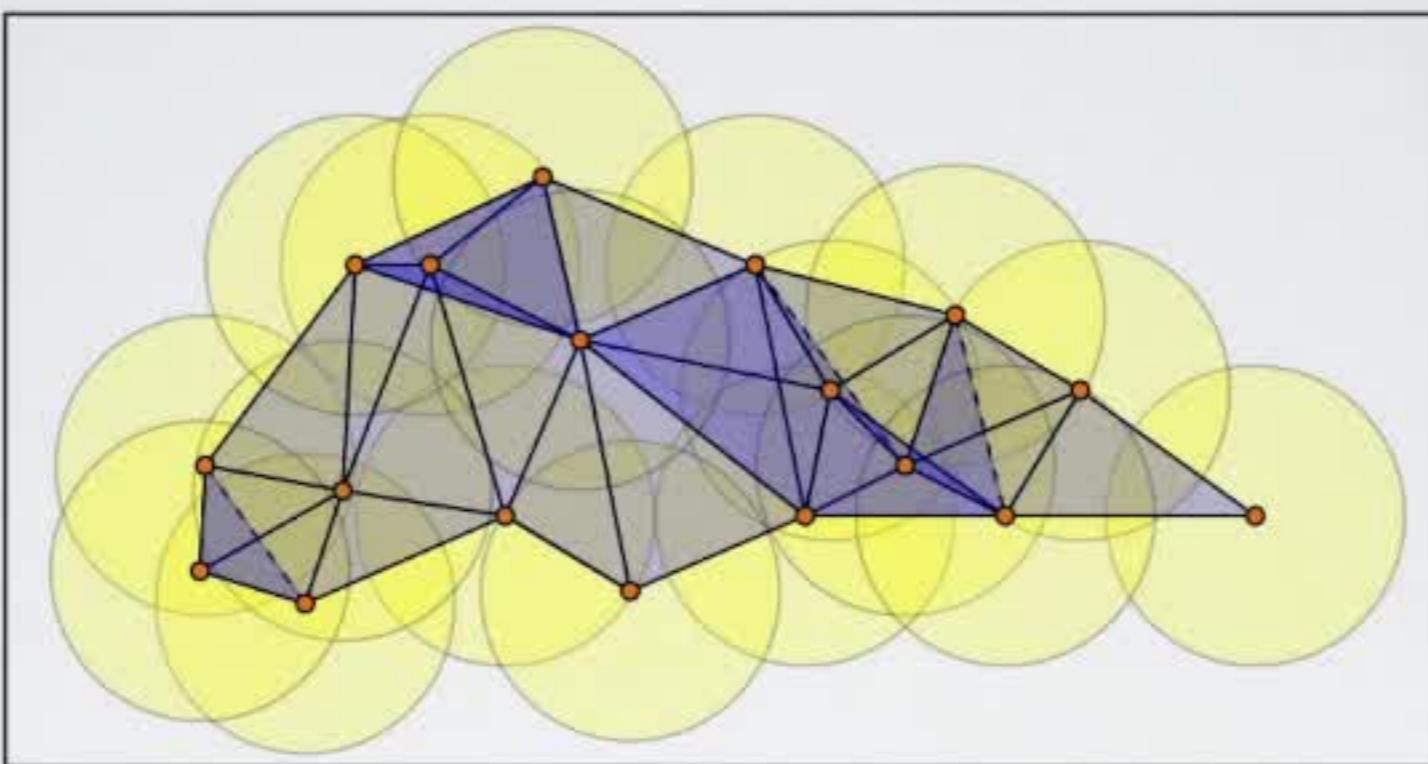


CALCULATE BETTI NUMBERS



Vietoris-Rips Complex

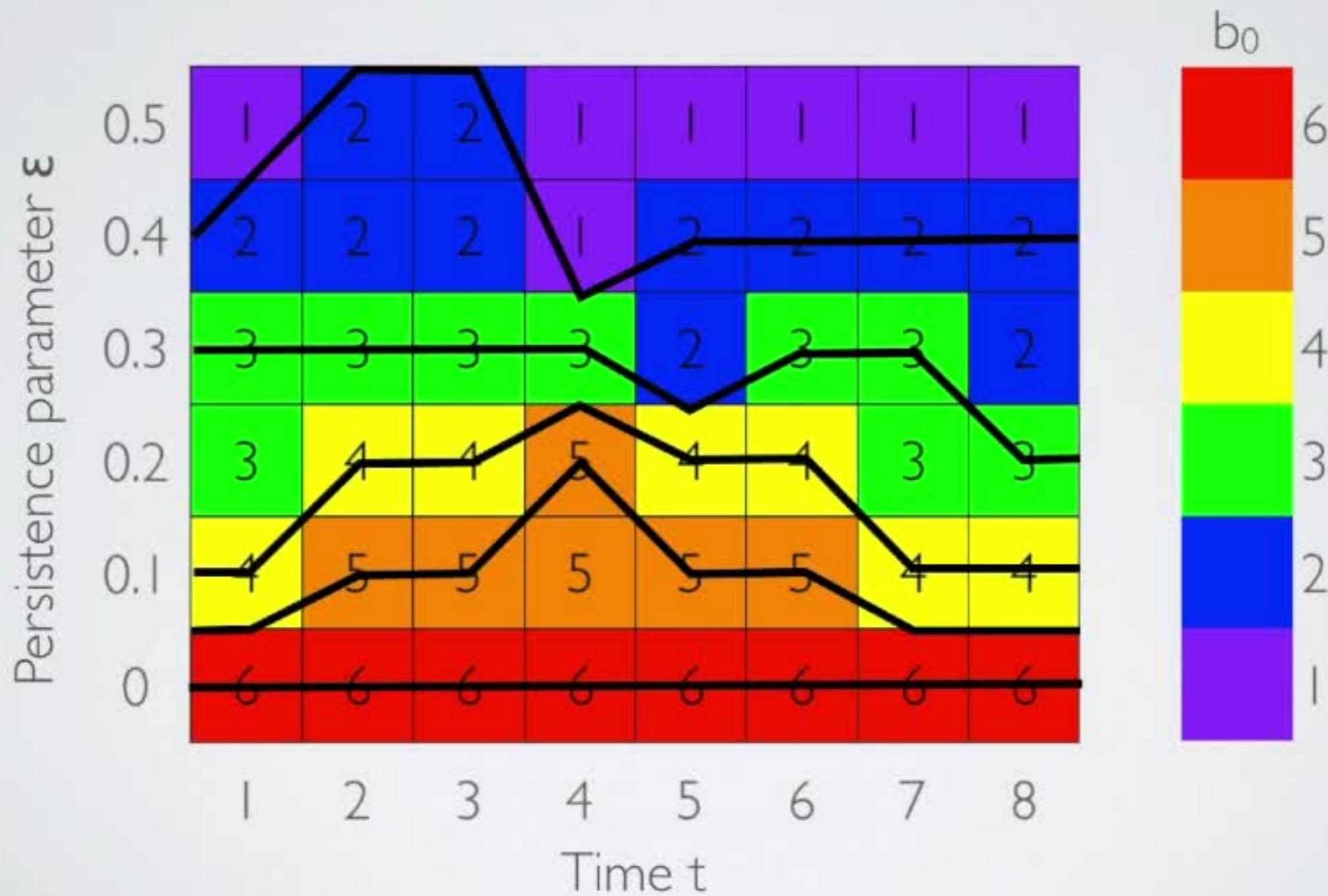
FIND PERSISTENT HOMOLOGY



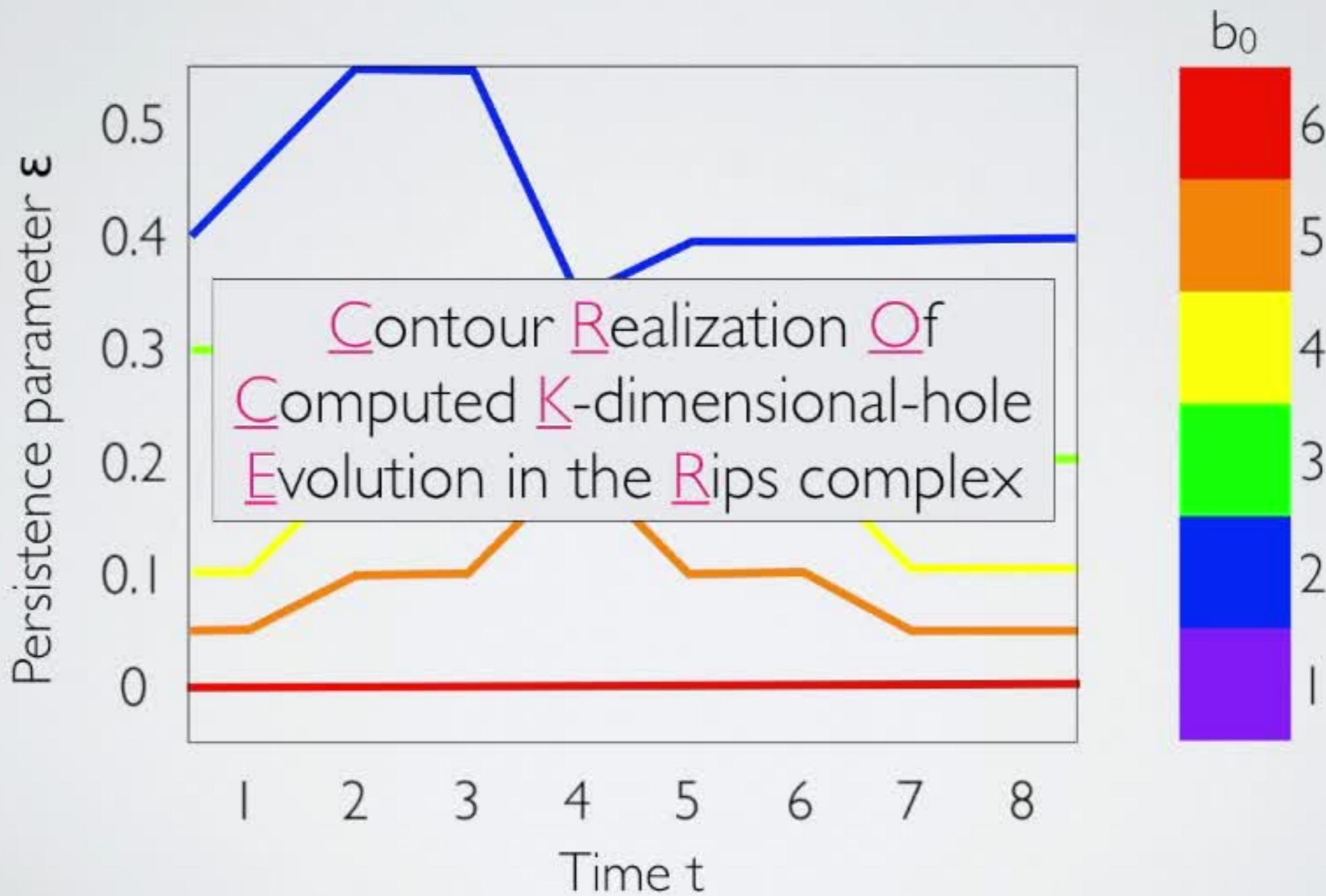
START AGAIN WITH PERSISTENT HOMOLOGY



EVOLVE IN TIME



EVOLVE IN TIME

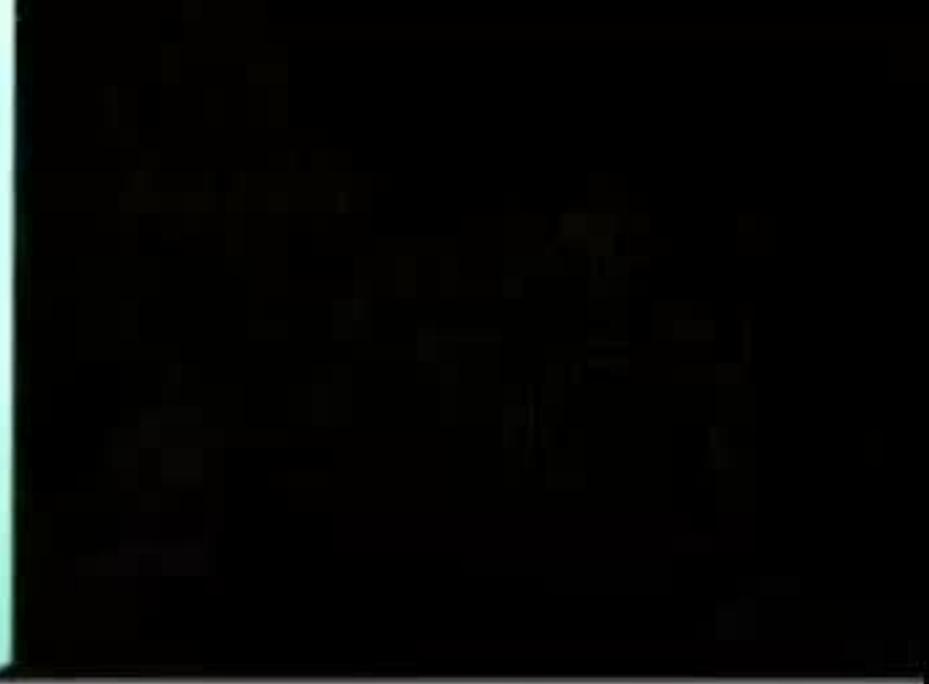


OTHER TOPOLOGICAL SIGNATURES MAY INCLUDE

- Persistence diagrams
- Persistence landscapes
- Persistence images
- Vineyards
- Crocker videos
- Who knows what else

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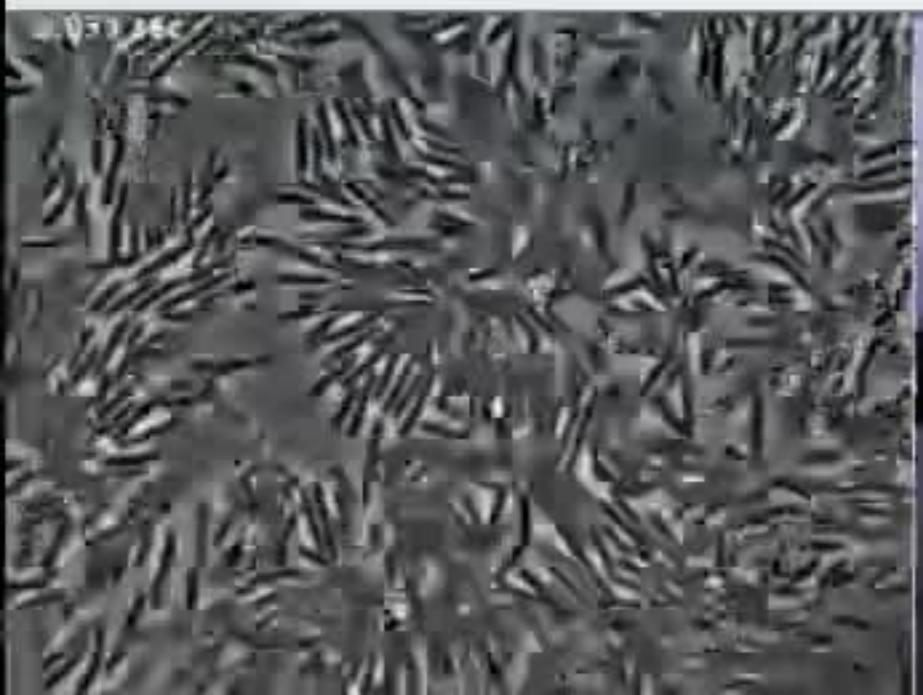


COLLECTIVE MOTION





COLLECTIVE MOTION





COLLECTIVE MOTION



A SEMINAL MODEL OF ALIGNING AGENTS

Novel type of phase transition in a system of self-driven particles

T Vicsek, A Czirók, E Ben-Jacob, I Cohen, O Shochet - Physical review letters, 1995 - APS

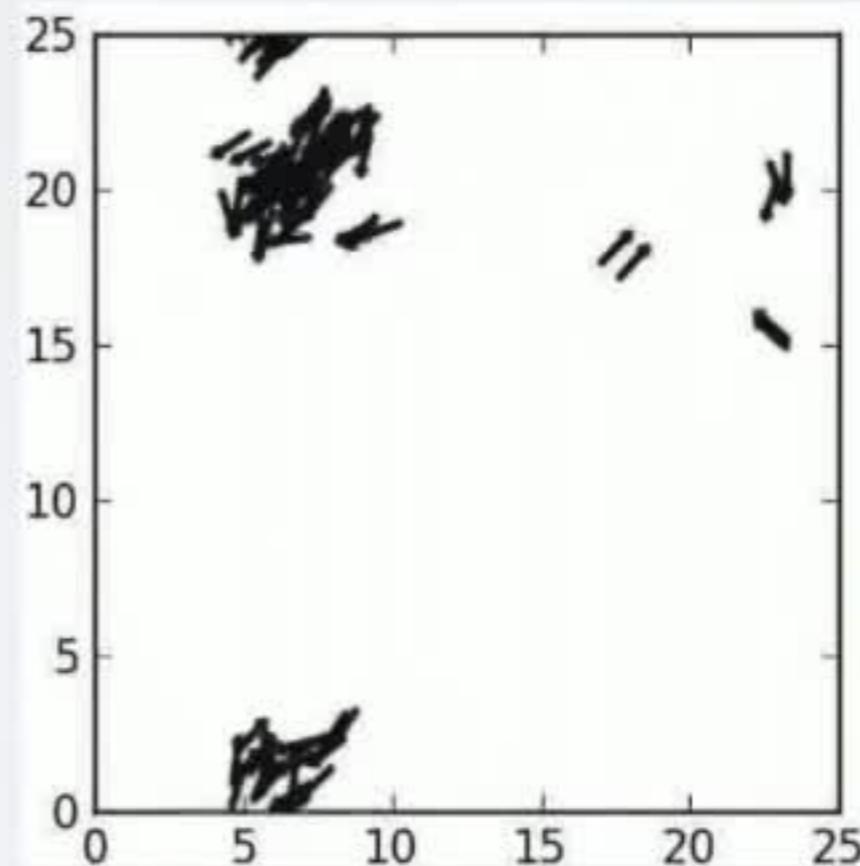
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$$\theta_i \rightarrow \langle \theta_j \rangle_{|\mathbf{x}_i - \mathbf{x}_j| \leq R} + U(-\eta/2, \eta/2)$$

alignmentnoise

$$\mathbf{v}_i \rightarrow v_0 (\cos \theta_i, \sin \theta_i)$$

$$\mathbf{x}_i \rightarrow \mathbf{x}_i + \mathbf{v}_i \Delta t$$



<http://youtube/phRZV3oCal>

A SEMINAL MODEL OF ALIGNING AGENTS

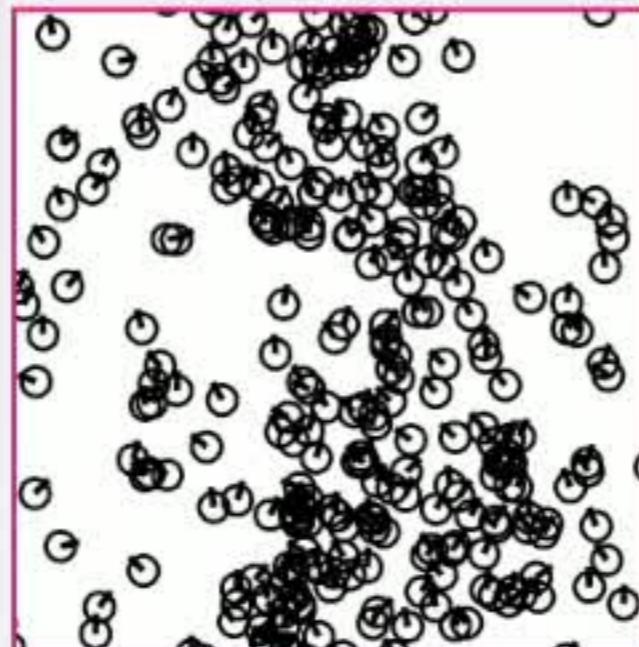
Novel type of phase transition in a system of self-driven particles

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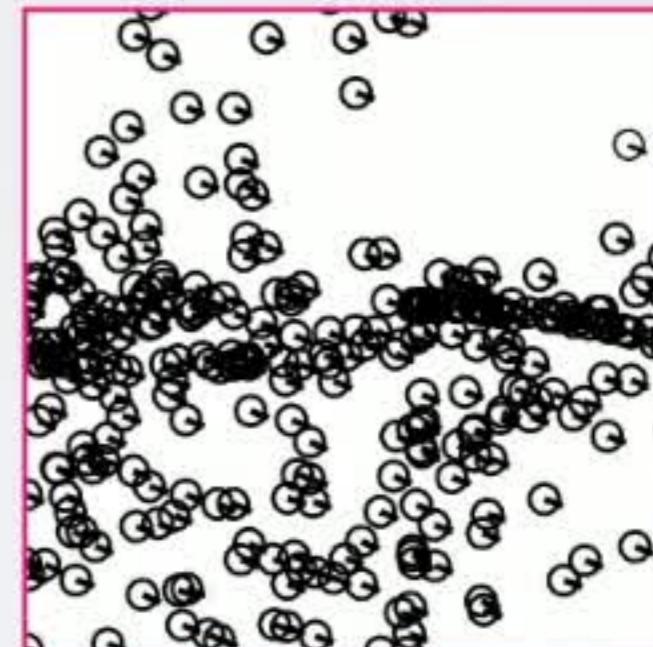
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clusters



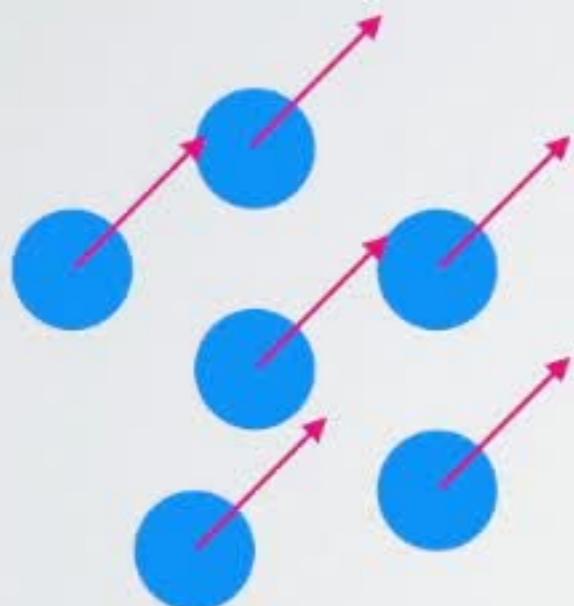
loose alignment



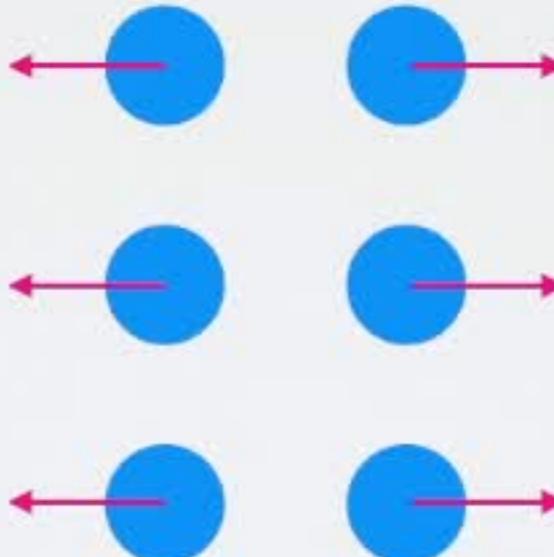
polarization

SUMMARIZE DATA WITH ORDER PARAMETER

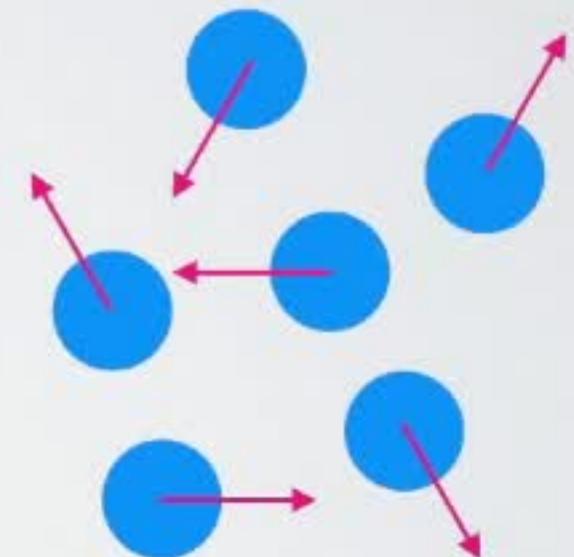
$$\text{Alignment } \phi(t) = \frac{1}{Nv_0} \left| \sum_{i=1}^N \mathbf{v}_i(t) \right|$$



$$\phi = 1$$

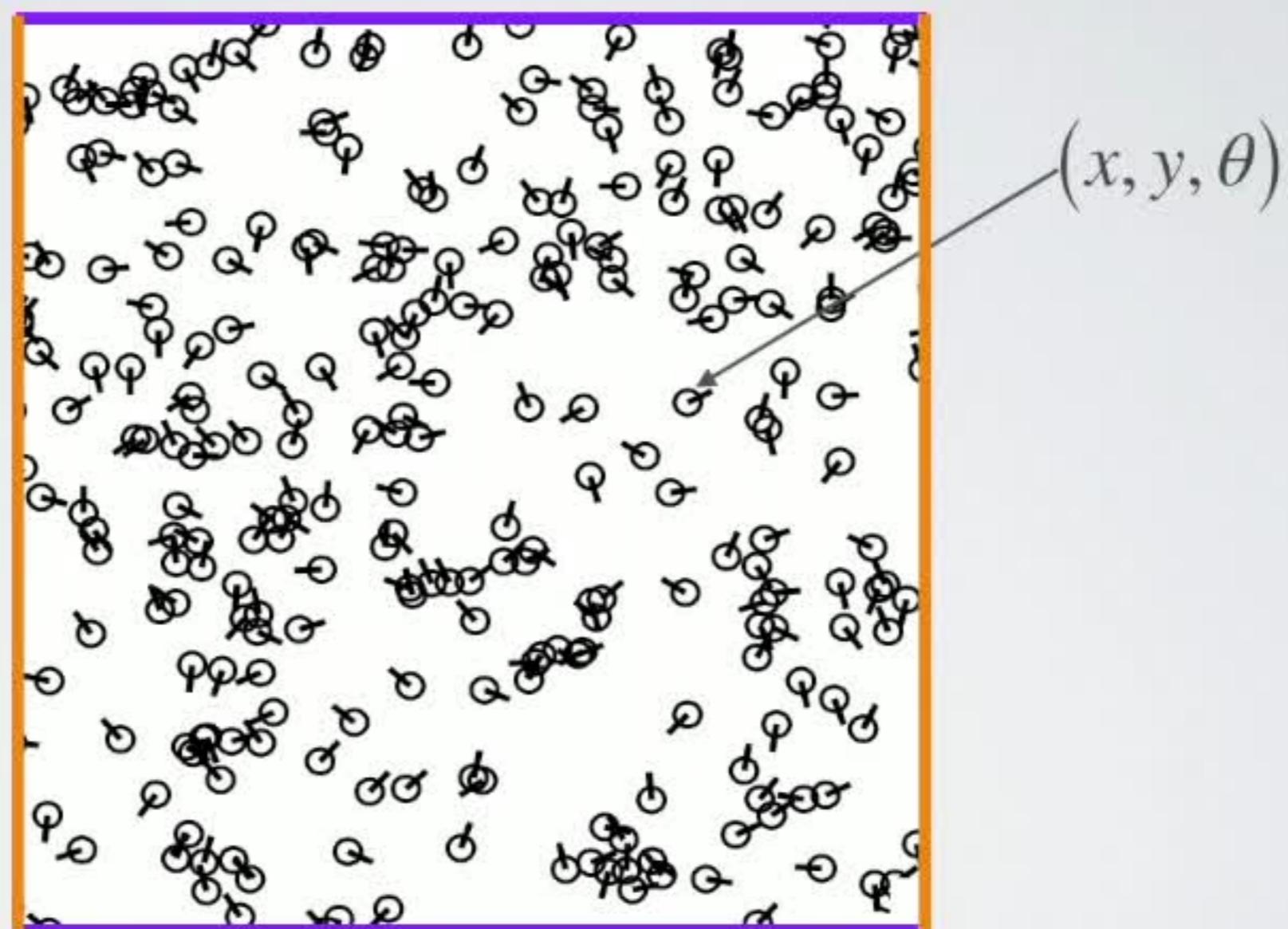


$$\phi = 0$$



$$\phi = 0$$

INITIAL CONDITION COVERS A THREE TORUS



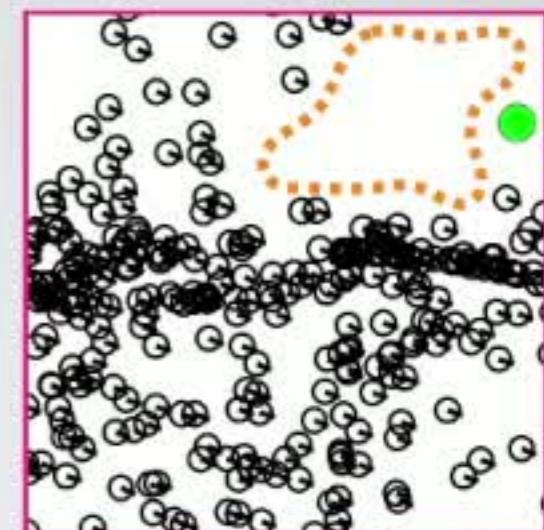
$$b = (1, 3, 3, 1, 0, \dots)$$

TOPOLOGICAL ORDER PARAMETER

Parameter Set #1
Clusters?

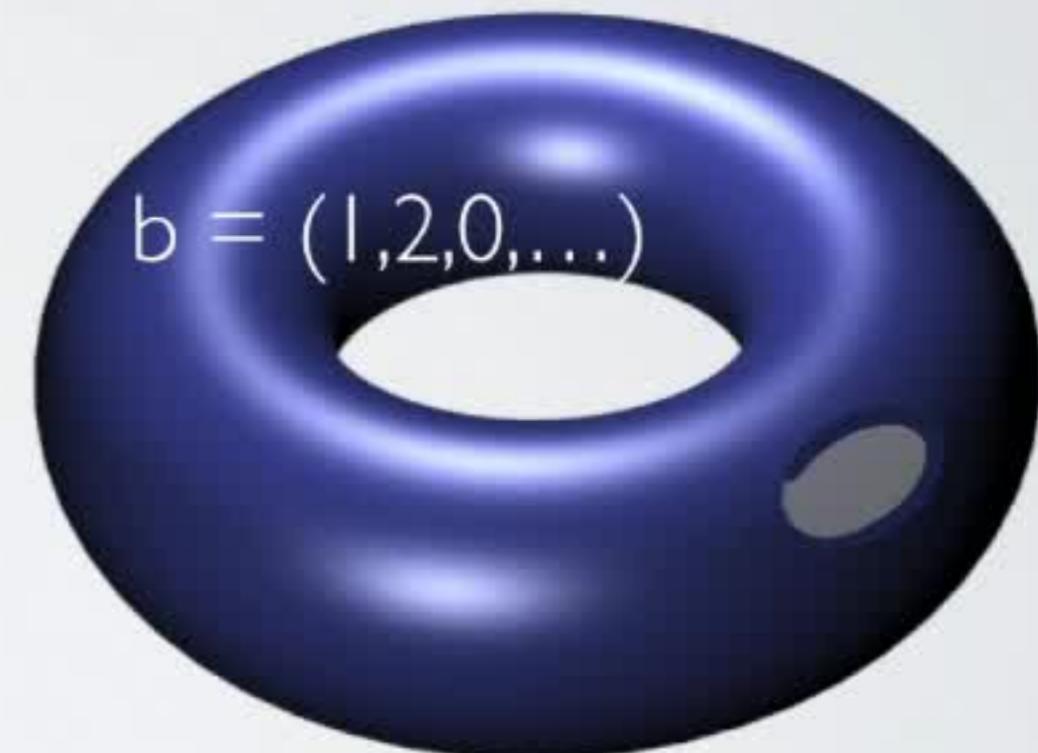
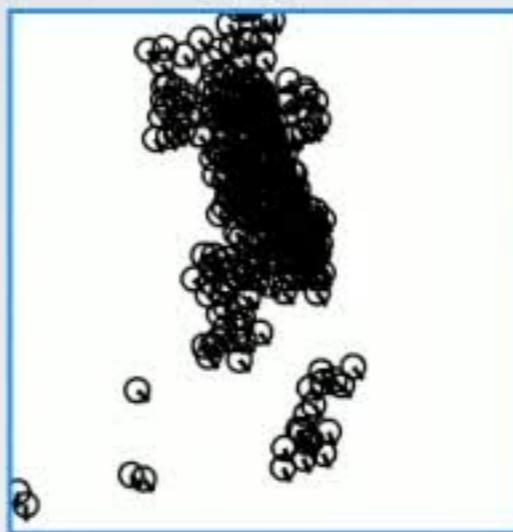


Parameter Set #2
Polarization?



TOPOLOGICAL ORDER PARAMETER

Parameter Set #1
Clusters?

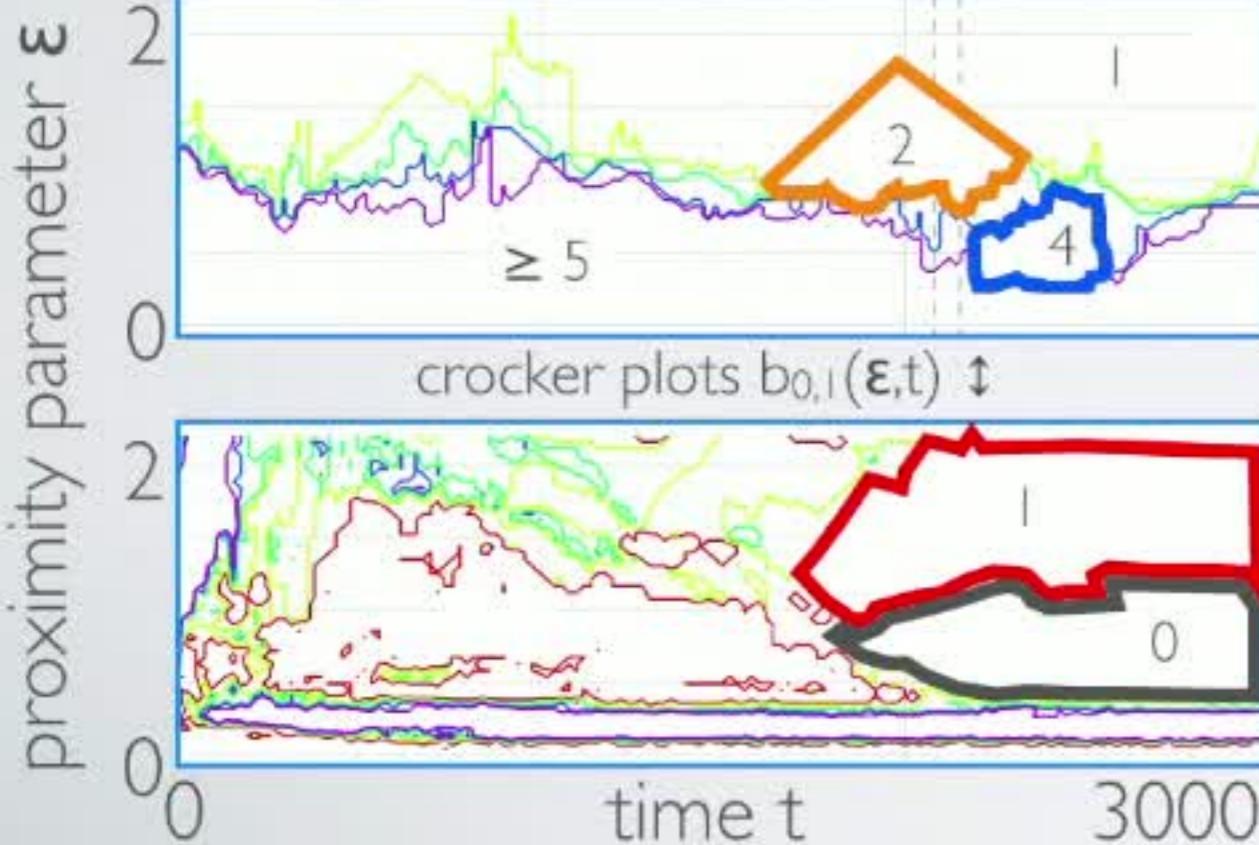
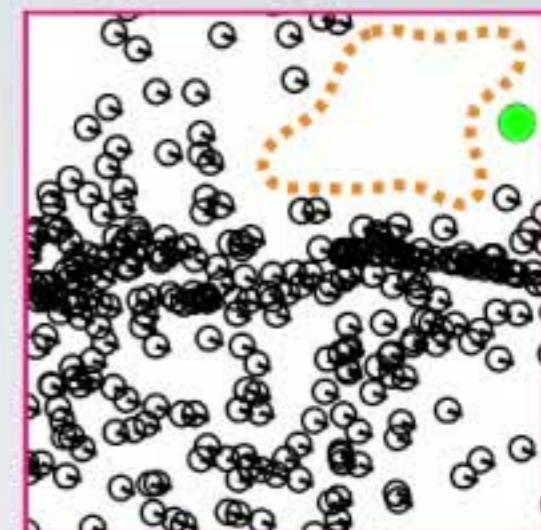


TOPOLOGICAL ORDER PARAMETER

Parameter Set #1
Clusters?

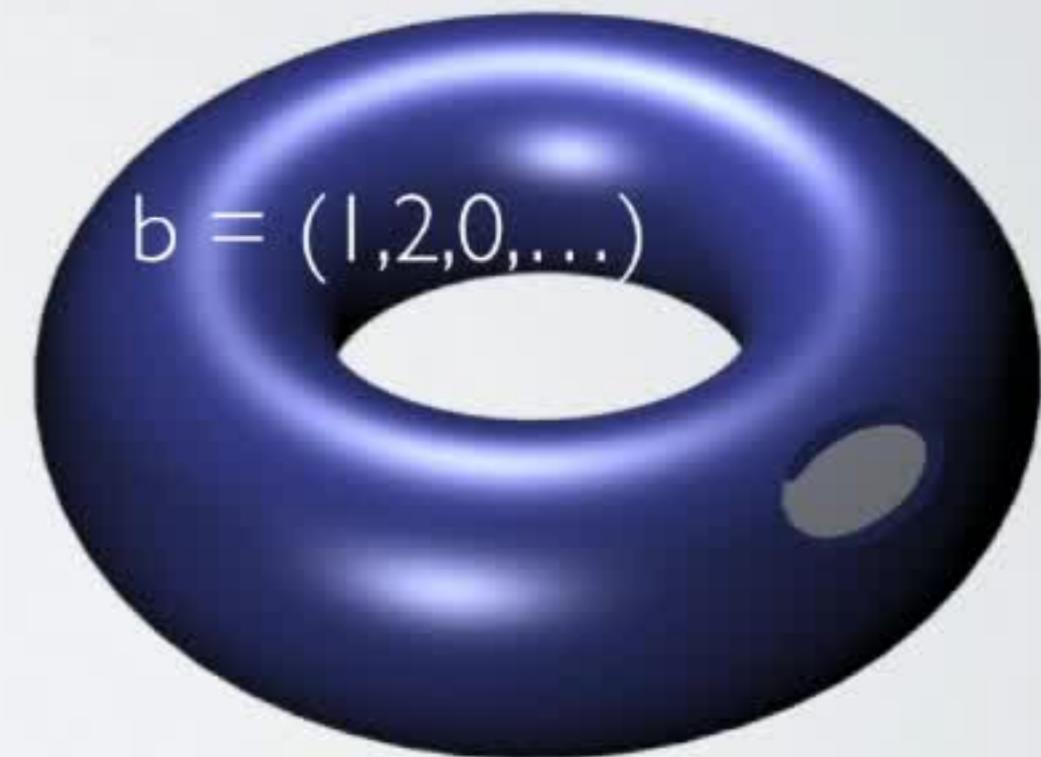


Parameter Set #2
Polarization?



TOPOLOGICAL ORDER PARAMETER

Parameter Set #1
Clusters?





all the data

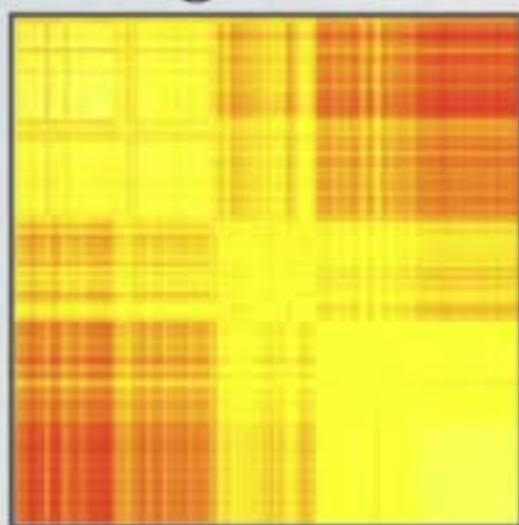
order parameter

topology

DATA EXPLORATION

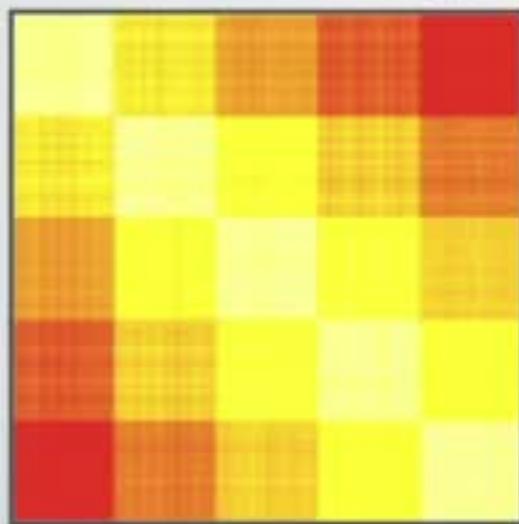
MACHINE LEARNING RESULTS

Alignment



far

Betti #'s $b_{0,1}$



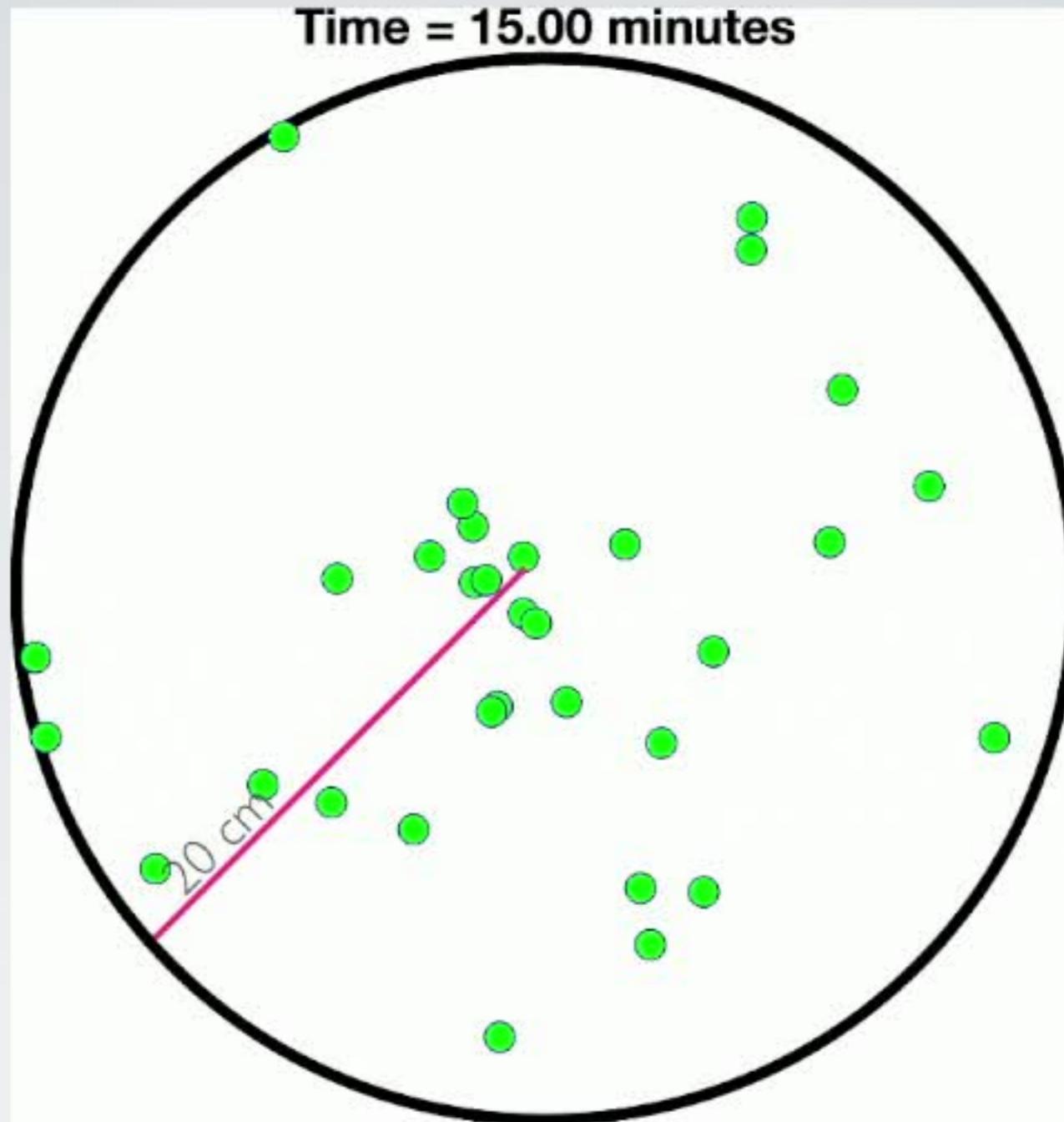
close

k-Medoids Clustering

Metric	PCA 3d?	Accuracy
Alignment	✓	51%
$b_{0,1}$	✓	100%
b_0	✓	100%
b_1	✓	99%

COLLECTIVE MOTION

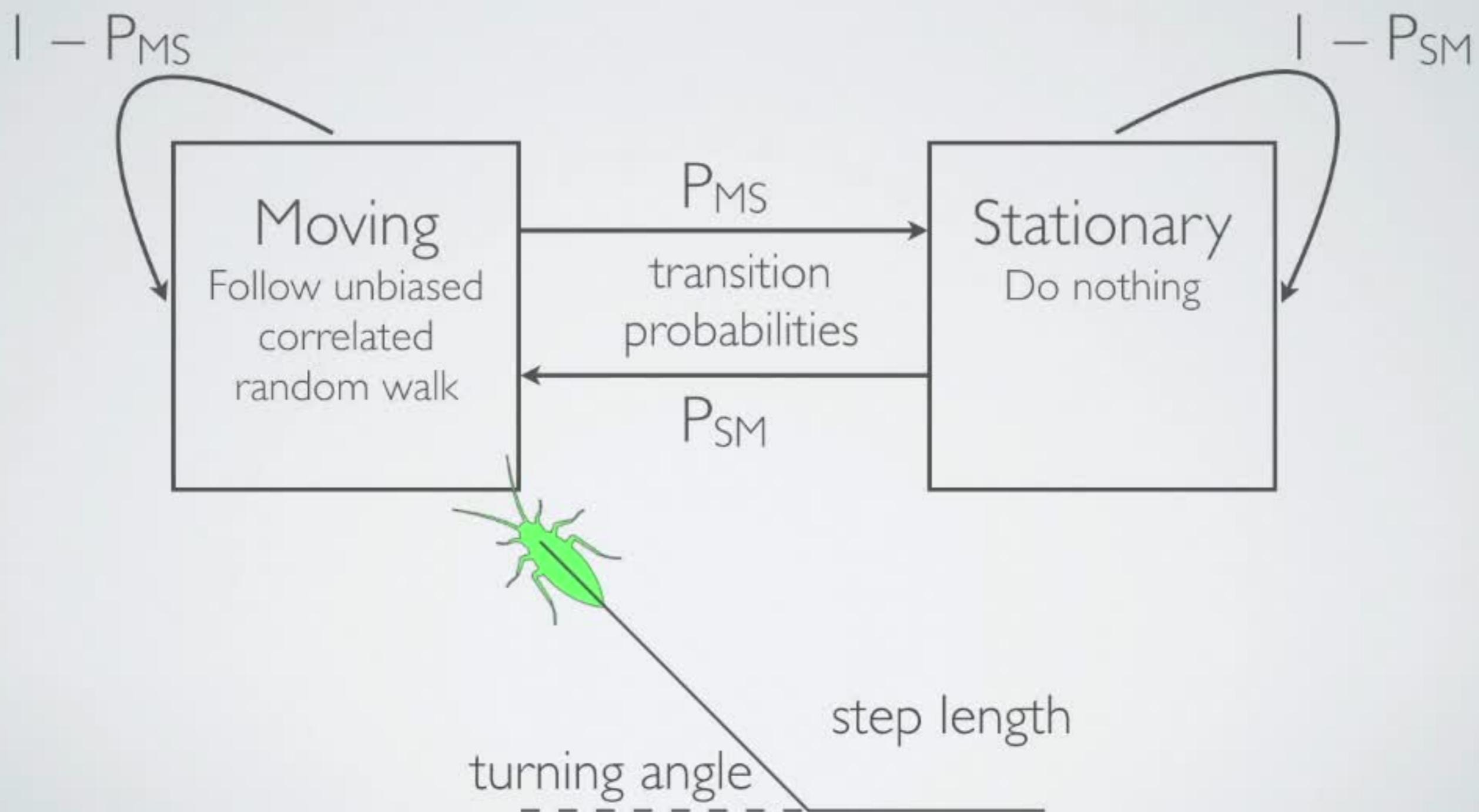
Main idea: Use topology as a metric for model selection.
Ziegelmeier, Ulmer, Topaz, PLOS One (2019)



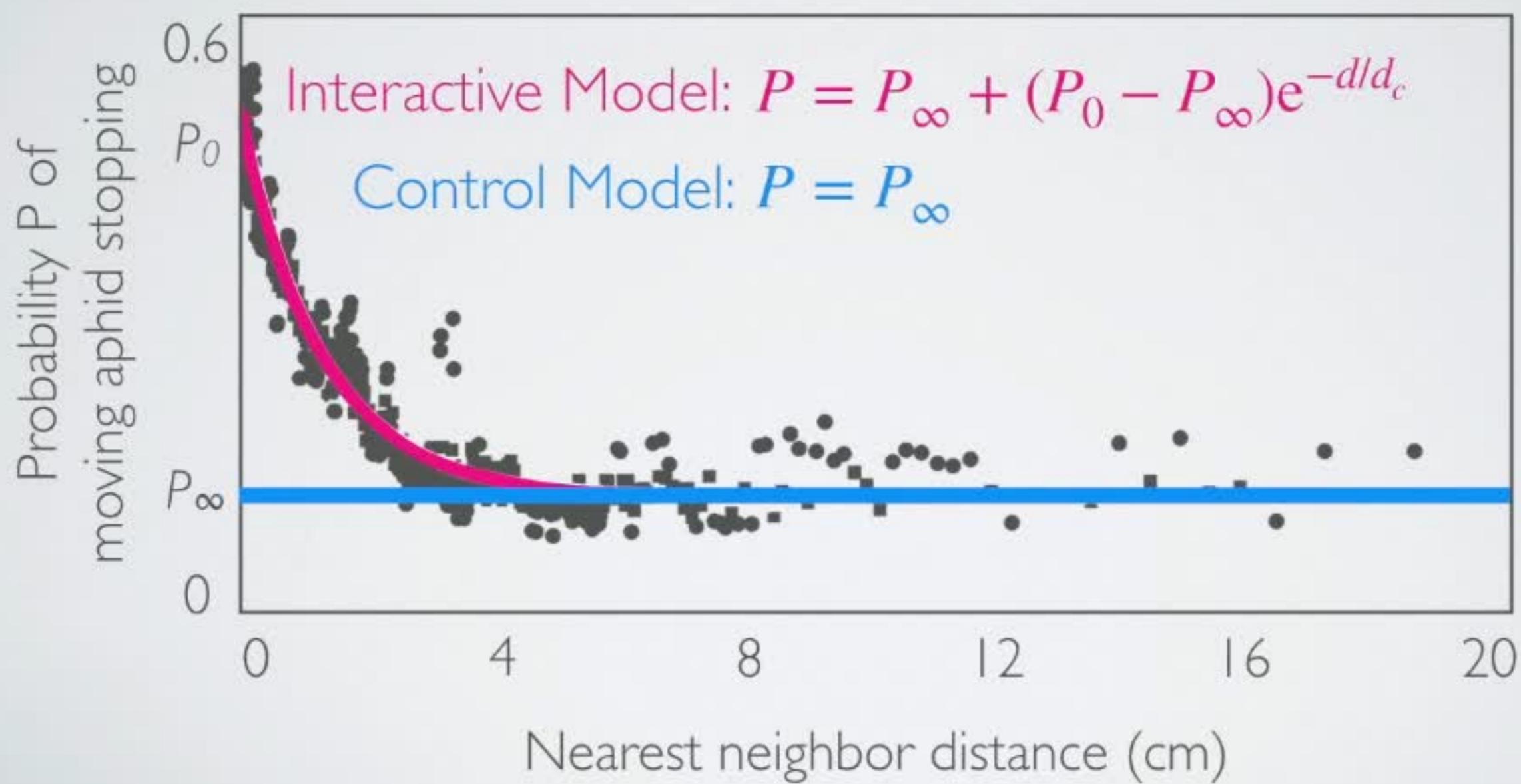
PEA APHID EXPERIMENT

Nilsen, Paige, Warner, Mayhew, Sutley, Lam, Bernoff, Topaz, PLOS One (2013)

TWO-STATE RANDOM WALK MODEL

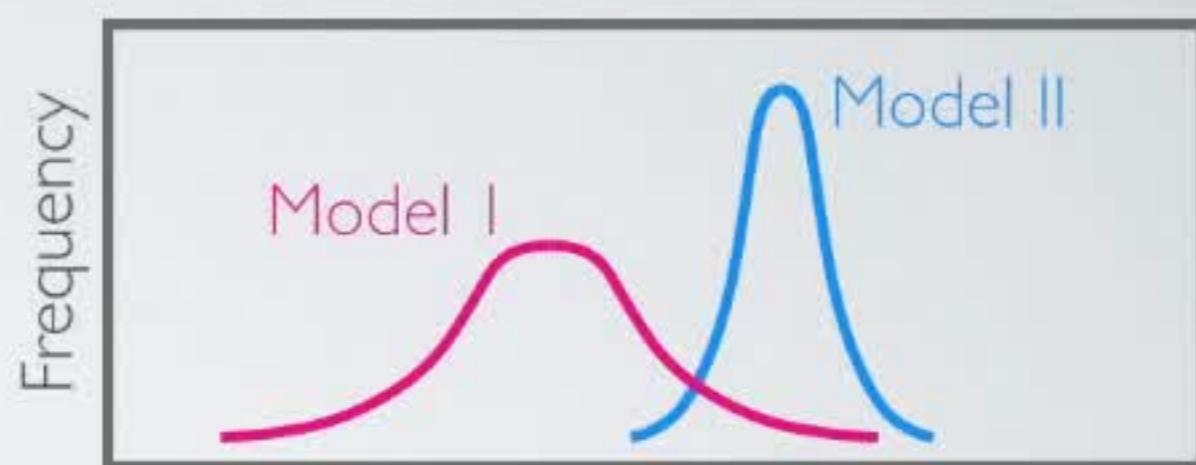


TWO POSSIBLE MODELS



EVALUATE MODEL FITNESS

Experiment }
Interactive model $\times 1000$ } $\times 9$
Control model $\times 1000$



Distance between simulations and experiment according to some metric

TRADITIONAL	A PRIORI, A.K.A. CHEATING	TOPOLOGY
Polariz. Angular Mom.	Abs. Angular Mom.	Avg. Buddy Dist. % Moving

EVALUATE MODEL FITNESS

A PRIORI, A.K.A. CHEATING

EVALUATE MODEL FITNESS

A PRIORI, A.K.A. CHEATING

EVALUATE MODEL FITNESS

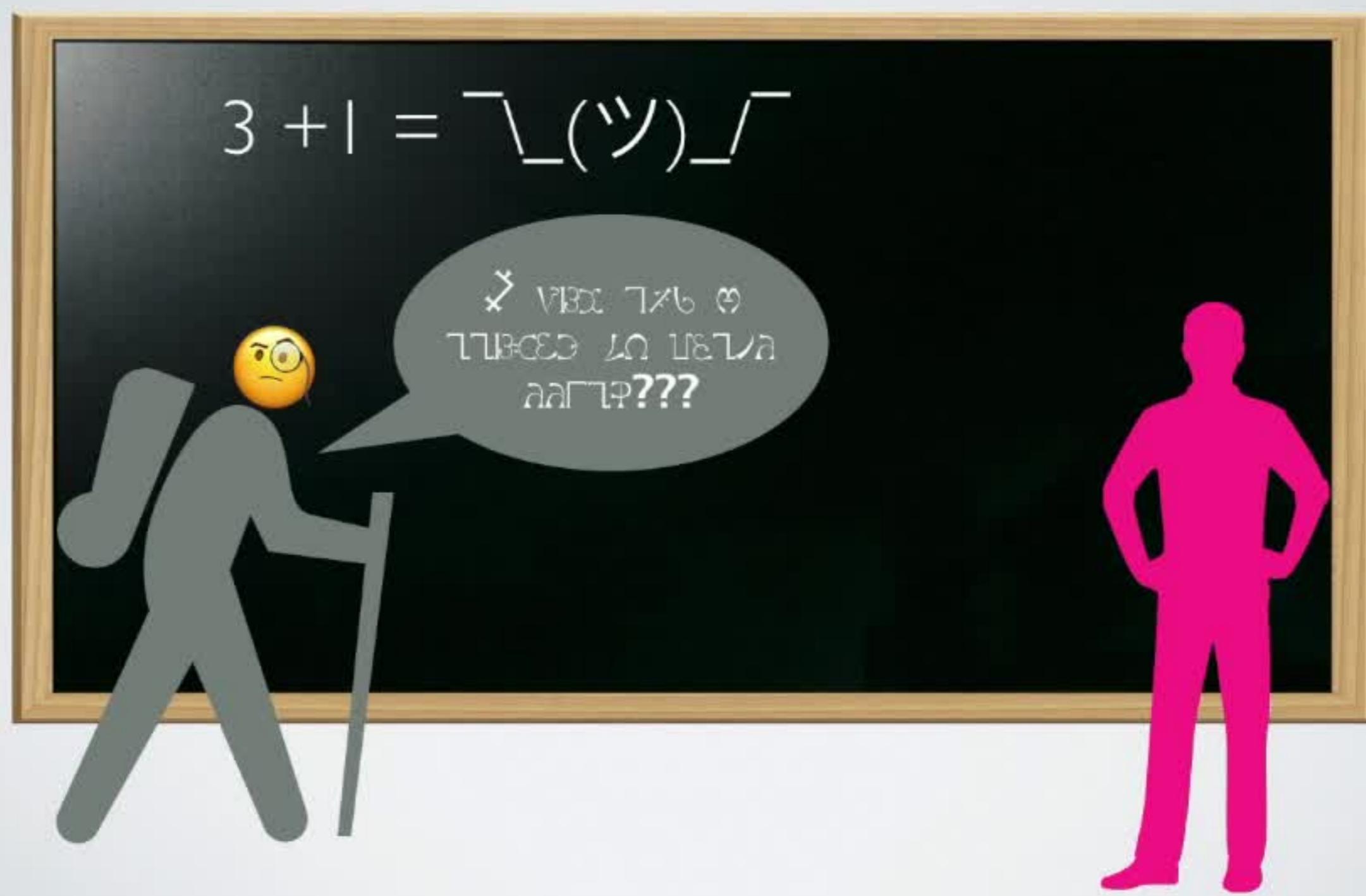
A PRIORI, A.K.A. CHEATING



MODEL SELECTION

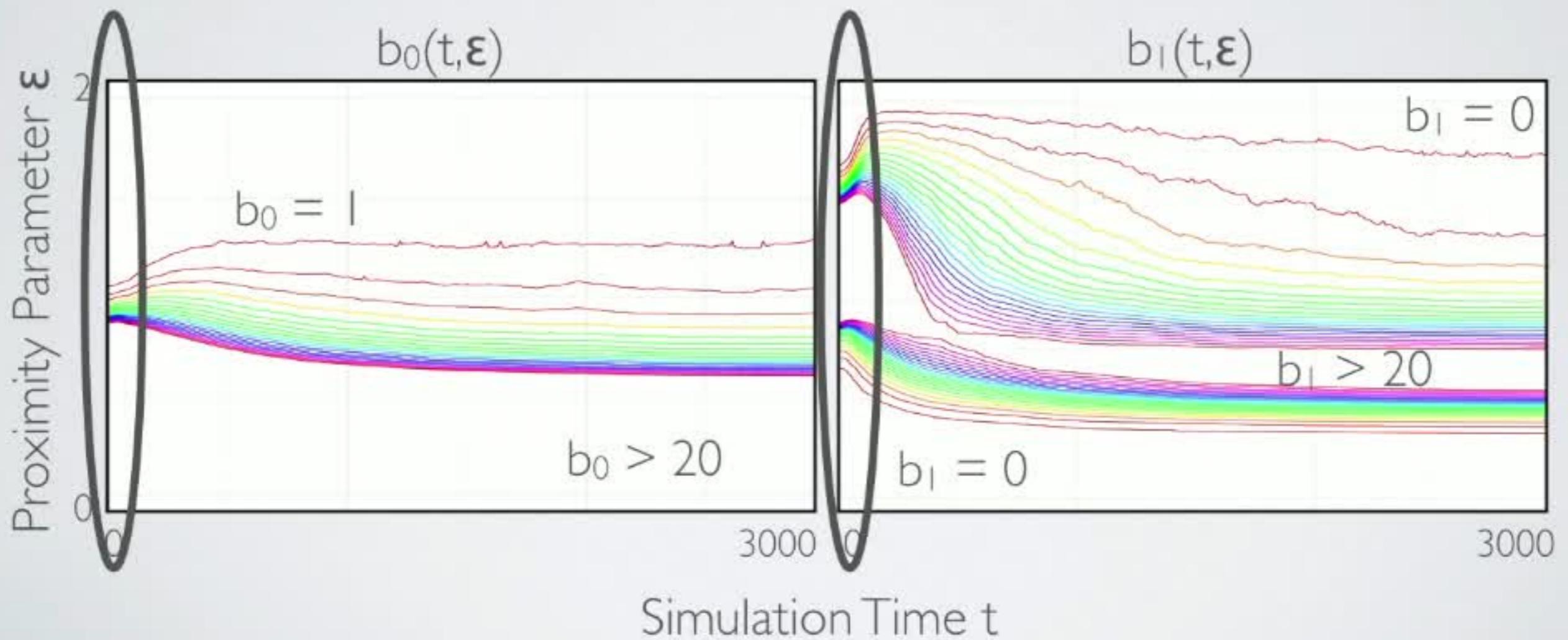
ONCE UPON A TIME

ONCE UPON A TIME



AVERAGE HOMOLOGY?

Vicsek model (naive) average over $n = 1000$ simulations

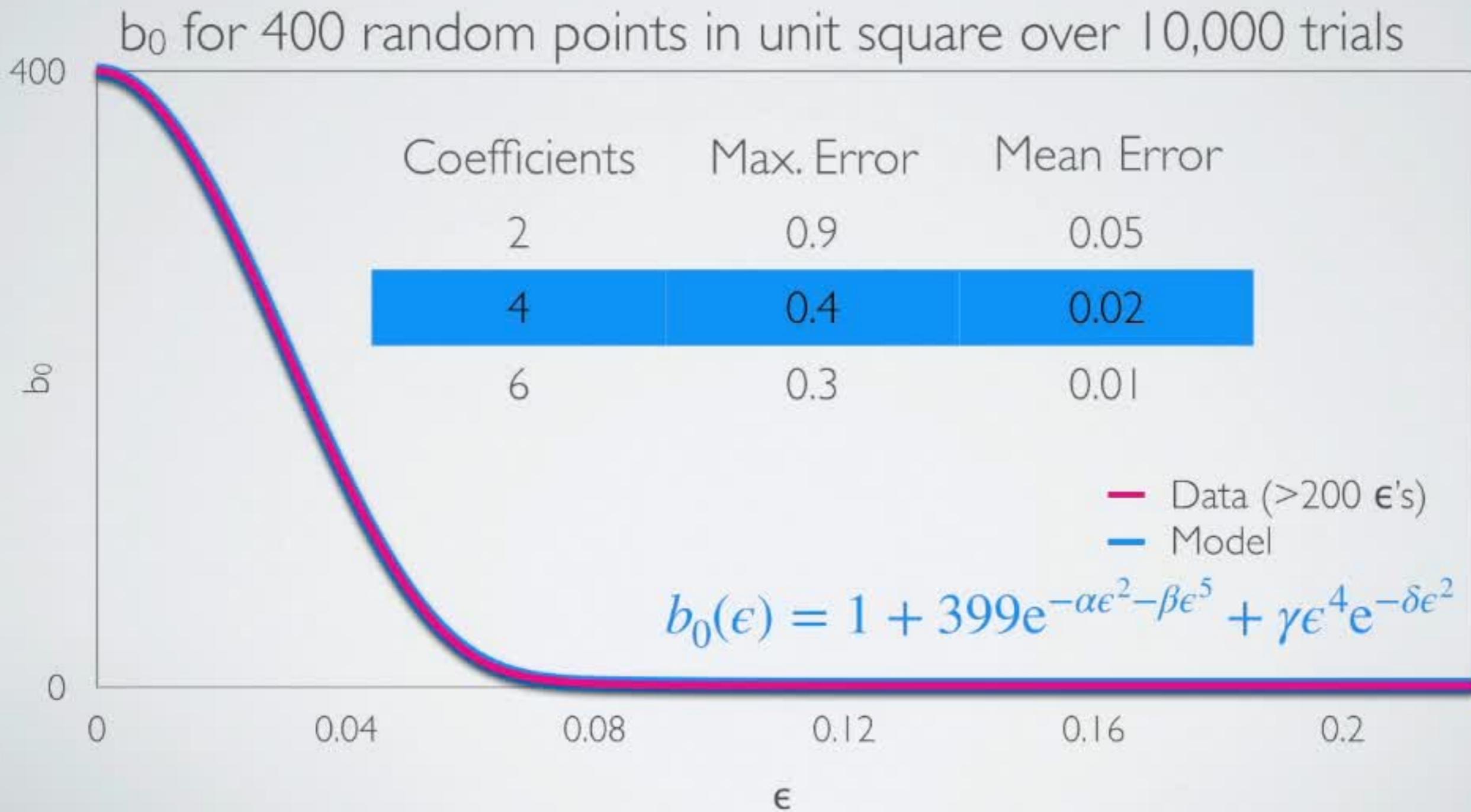


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SYMBOLIC REGRESSION

SIAM DS15, IP7, Hod Lipson

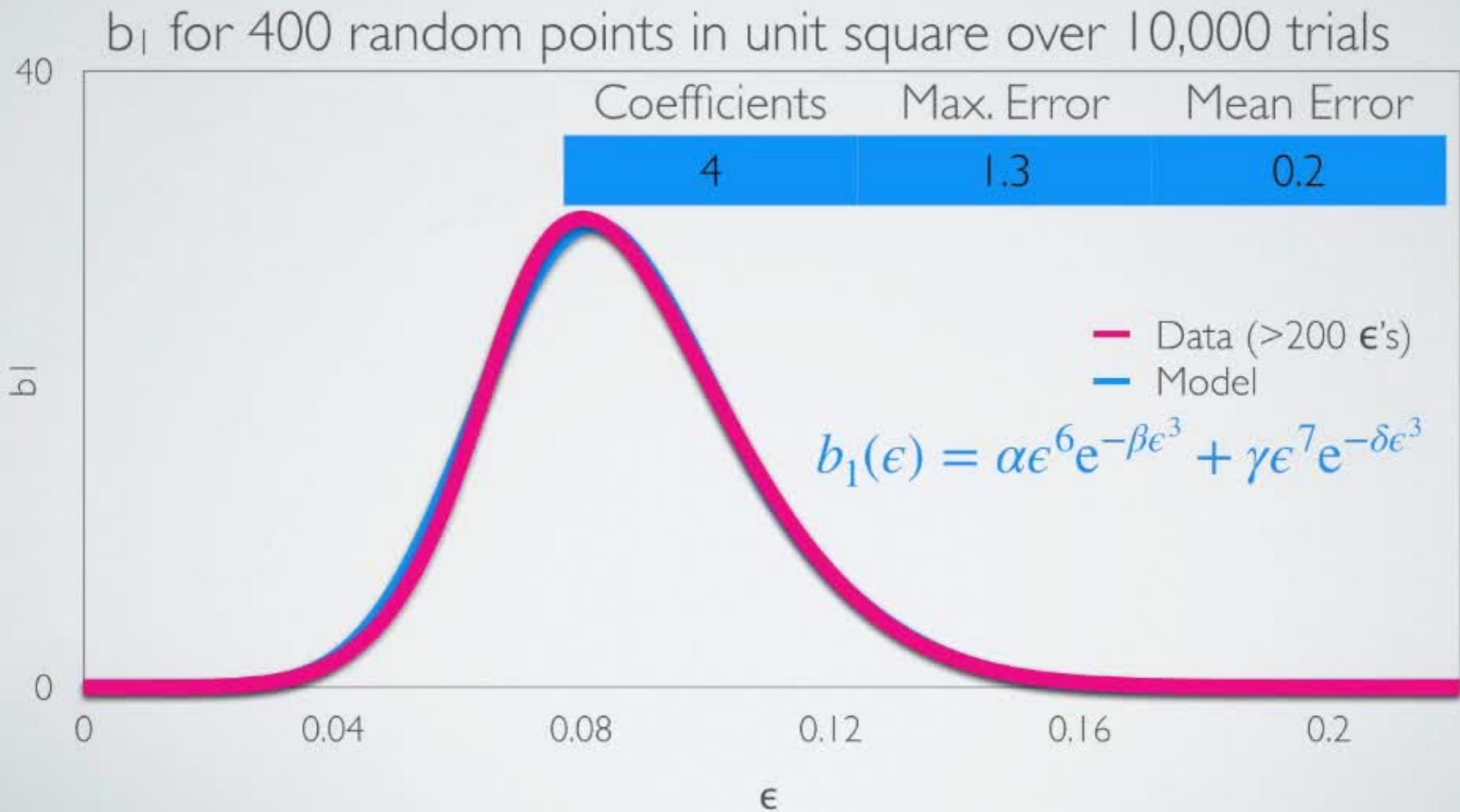
Distilling Natural Laws from Raw Data, from Robotics to Biology and Physics



SYMBOLIC REGRESSION

SIAM DS15, IP7, Hod Lipson

Distilling Natural Laws from Raw Data, from Robotics to Biology and Physics



TDA

- I have found TDA valuable for
 - Exploring large data sets
 - Recovering parameters
 - Choosing between models of given data
 - Constructing reduced theoretical models??
- We may benefit from topological tools
- Topologists may benefit from understanding our needs
- ...and from applied approaches to theoretical questions