



# Linking Local Decisions with Global Outcomes in Networks: Case Studies in Behavior and Population Health

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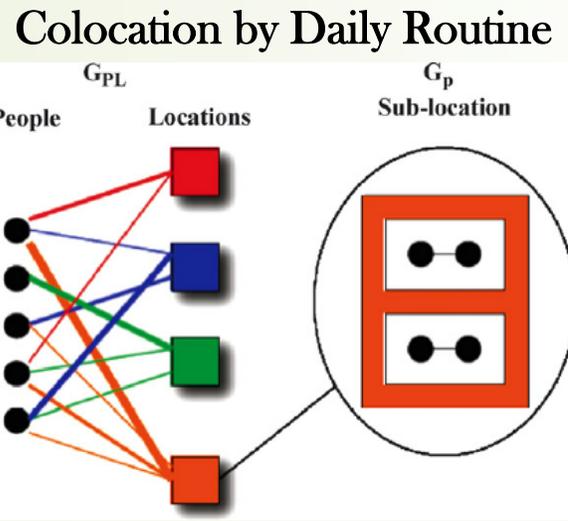
# Networks



# Self-Organizing Networks

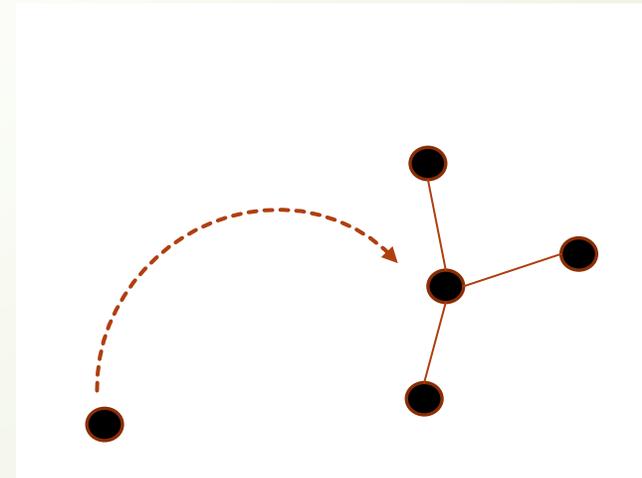
Emerge from Individual behavioral choices

- Not merely “emergent”
- Choices reflect decisions based on the network



Emergent  
(image from EpiSim)

Make Friends with Popular People



Emergent *and* Self-Organizing

# Self-Organizing Animal Social Networks

Emerge from individual behavioral choices that depend on the behaviors of others

- Proximity
- Grooming
- Aggression
- Mating
- Communicating (harder to tell)



# Individual Benefits from Being in a Group

## Group Success

Benefit to the individual, but achieved collaboratively

- Diffusion of risk from predators
- Increased foraging success
- Better engineering



From [www.amentsoc.org](http://www.amentsoc.org)



From [telegraph.co.uk](http://telegraph.co.uk)

# Individual Benefits from Being in a Network

Very cool studies have looked at evolutionary fitness associated with social network position

- Can genes determine position in a network?
- Do particular positions lead to better survival/more reproduction?

These focus on direct, individual fitness outcomes



From sci-news.com

# Individual Costs to Being in a Group/Network

Group Participation is not without costs

- Attract predators
- Competition for food/mates
- *Disease transmission*



# Group Benefits, Individual Success

This sets up a system of feedback control

- Individual-scale self-organization
- Direct and Indirect fitness components
- Selection includes group benefits and costs
- Individuals pass on their genes or not



# Bio Language: Multilevel Selection

## Math Language: Multiscale Feedback Control

		ORG	
		ORG>0	ORG<0
IND	IND>0	Selected <u>For</u>	If $ \text{IND}  >  \text{ORG} $ : <u>For</u>  <u>Against</u>
	IND<0	If $ \text{IND}  <  \text{ORG} $ : <u>For</u>  <u>Against</u> If $ \text{IND}  >  \text{ORG} $ :	Selected <u>Against</u>

Reproduced from Hock, Ng and Fefferman, 2010



To explore, we build a mathematical abstraction and use it as a computational experimental system

Assumption: Individuals make genetically determined “selfish” social affiliation choices (with no regard for group-level effects)

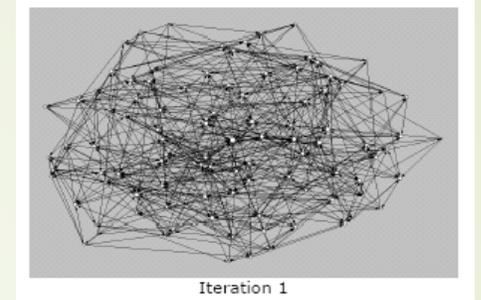
Initial hypothetical proxies of 3 measurements of social status from social network theory:

- Degree - non-transitive dominance hierarchy
- Closeness - genetic relatedness
- Betweenness - much more complicated, assumed much harder to detect

## Centrality has useful built-in features:

	Individual Outcome	Global Outcome
degree	$D(v_i) = \frac{d_{in}(v_i)}{n-1}$	$\frac{\sum_{i=1}^n (D^* - d_{in}(v_i))}{(n-1)(n-2)}$
closeness	$C(v_i) = \frac{n-1}{\sum_{j \neq i} d(v_i, v_j)}$	$\frac{\sum_{i=1}^n C(v_i)}{(n-1)(n-2)}$
betweenness	$B(v_i) = \frac{2\text{count}(v_i)}{(n-1)(n-2)}$	$\frac{\sum_{i=1}^n B(v_i)}{(n-1)(n-2)}$

## Experimental setup:



- Initialize a digraph with  $n$  vertices
- Randomly generate arcs such that each vertex has out-degree = 5

Then we iterate the following steps

- Compute the 3 centrality measures for all of the vertices and for the entire network
- In each step, each vertex **drops two** (how on next slide) of its existing out-neighbors and **replaces them with two new ones**

# 3 Different types of Populations - How to choose which social contacts to drop

- *Each vertex **drops two** of its existing out-neighbors and **replaces them with two new ones***

All individuals in a population use the same measure (Degree, Closeness or Betweenness) to evaluate others

Each individual drops the two out arcs to the two affiliates with the **worst centrality measure** (among the five neighbors) and picks up two new ones (ratios are arbitrary, results are robust)

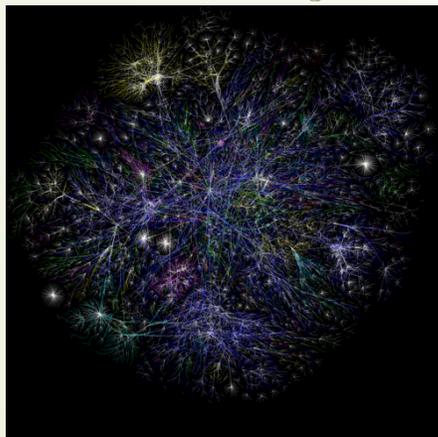
In all types of populations, we record all of the centrality measures for each individual and for the entire network over time

# Unimportant Note: Not Just a Model System

A bunch of real-world human networks are actually  
Centrality self-organizing:

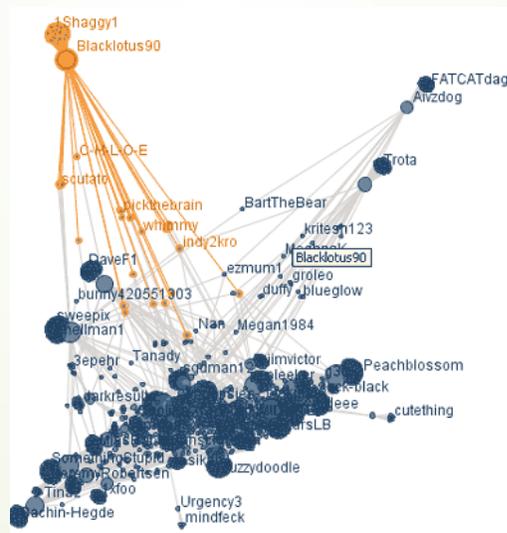
Degree

From Lumeta.org



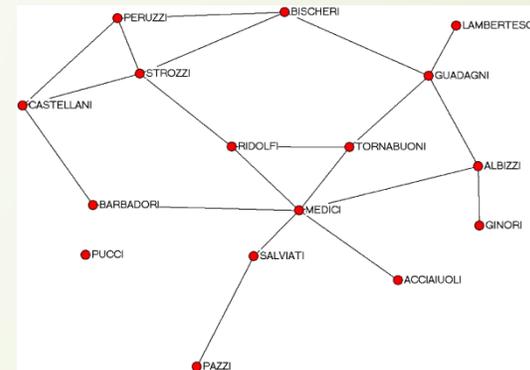
HTML links  
on the Web

Closeness



Top StumbleUpon  
Users

Betweenness



15<sup>th</sup> Century  
Marriages between  
Florentine Families

# How to choose which social contacts to pick up

## Three different ways:

Incomplete knowledge - individuals know centrality of only their current contacts, so two new contacts are **chosen at random** from all the rest

Complete knowledge - individuals know the centrality measure of everyone and **choose the two best**

Zero Knowledge - individuals have no centrality measure preference - so they drop two connections at **random** (and then add back two new ones at random)

*Note: We don't need to be able to calculate centrality to have good proximate ways to estimate it*



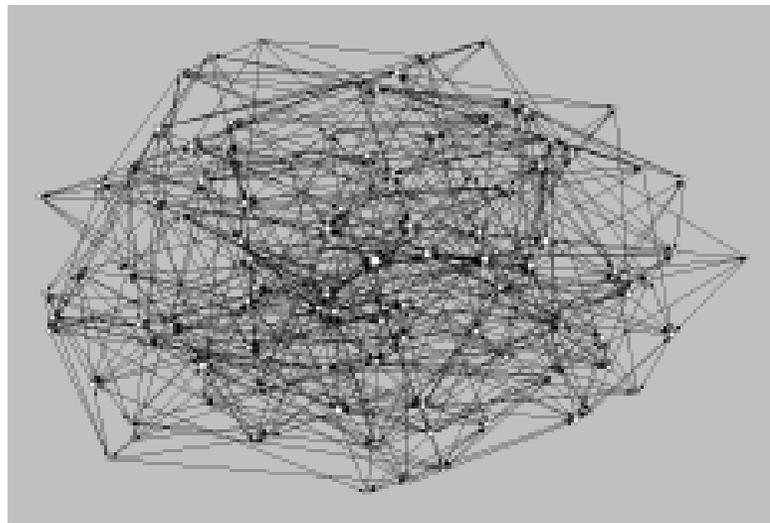
# The evolutionary interpretation of the levels of knowledge

Zero-knowledge : before to the evolution of individual social choice or individuals are terrible at evaluating each other's status (no good proximate mechanism)

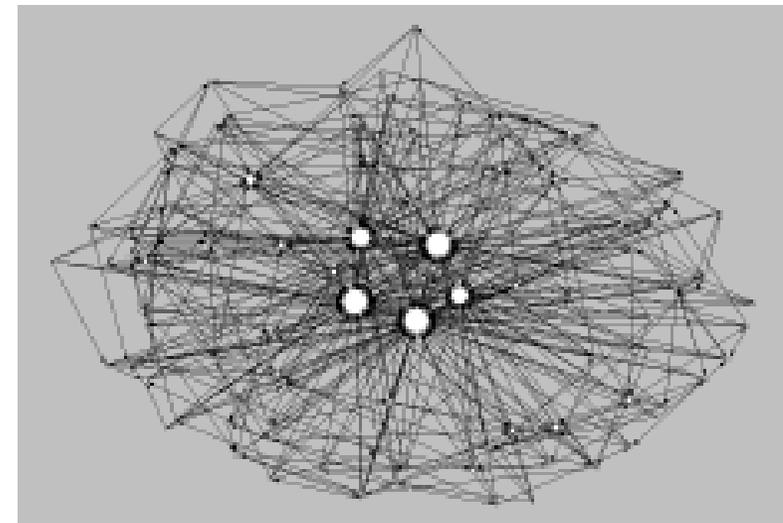
Complete-knowledge : social choice in smaller populations where you can evaluate everyone

Incomplete-knowledge : social choice in larger populations where you can only evaluate your friends

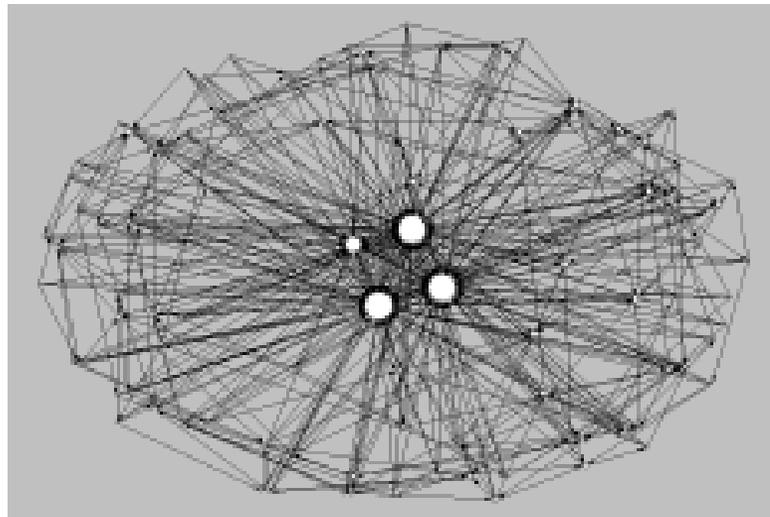
# Convergence!



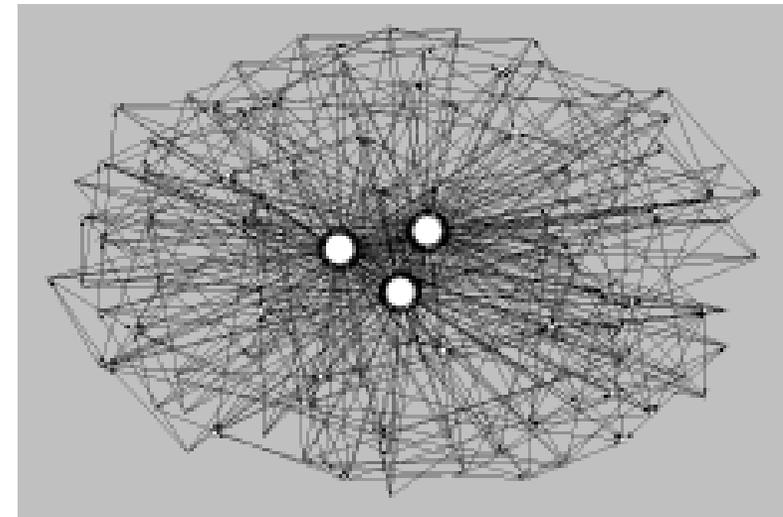
Iteration 1



Iteration 50



Iteration 100



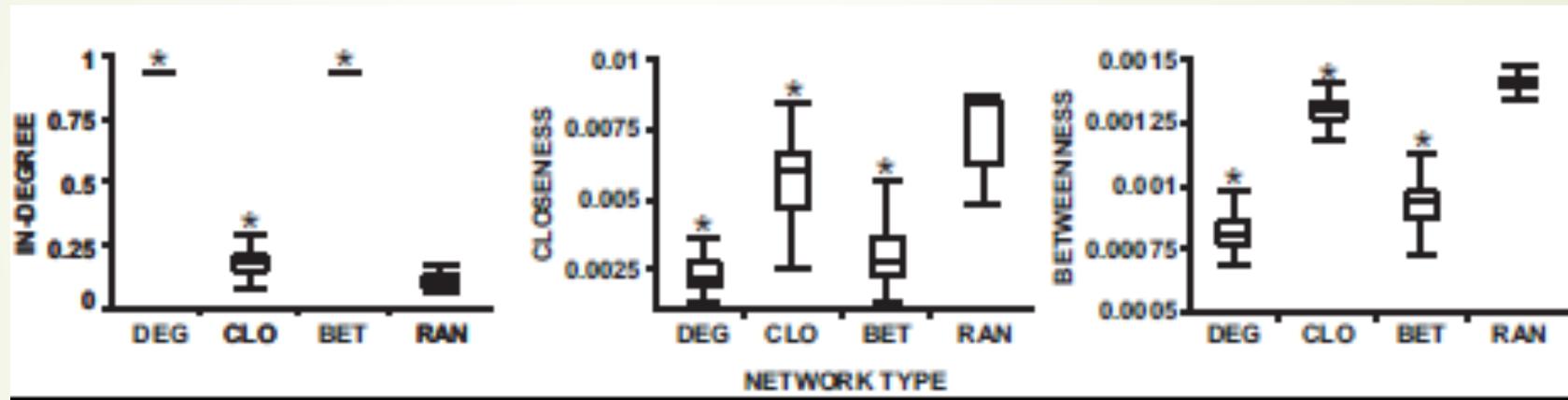
Iteration 200

# Result: Different Individual Strategies Work to Accomplish Different Group Outcomes

D ? B > C > R

R > C > B > D

R > C > B > D



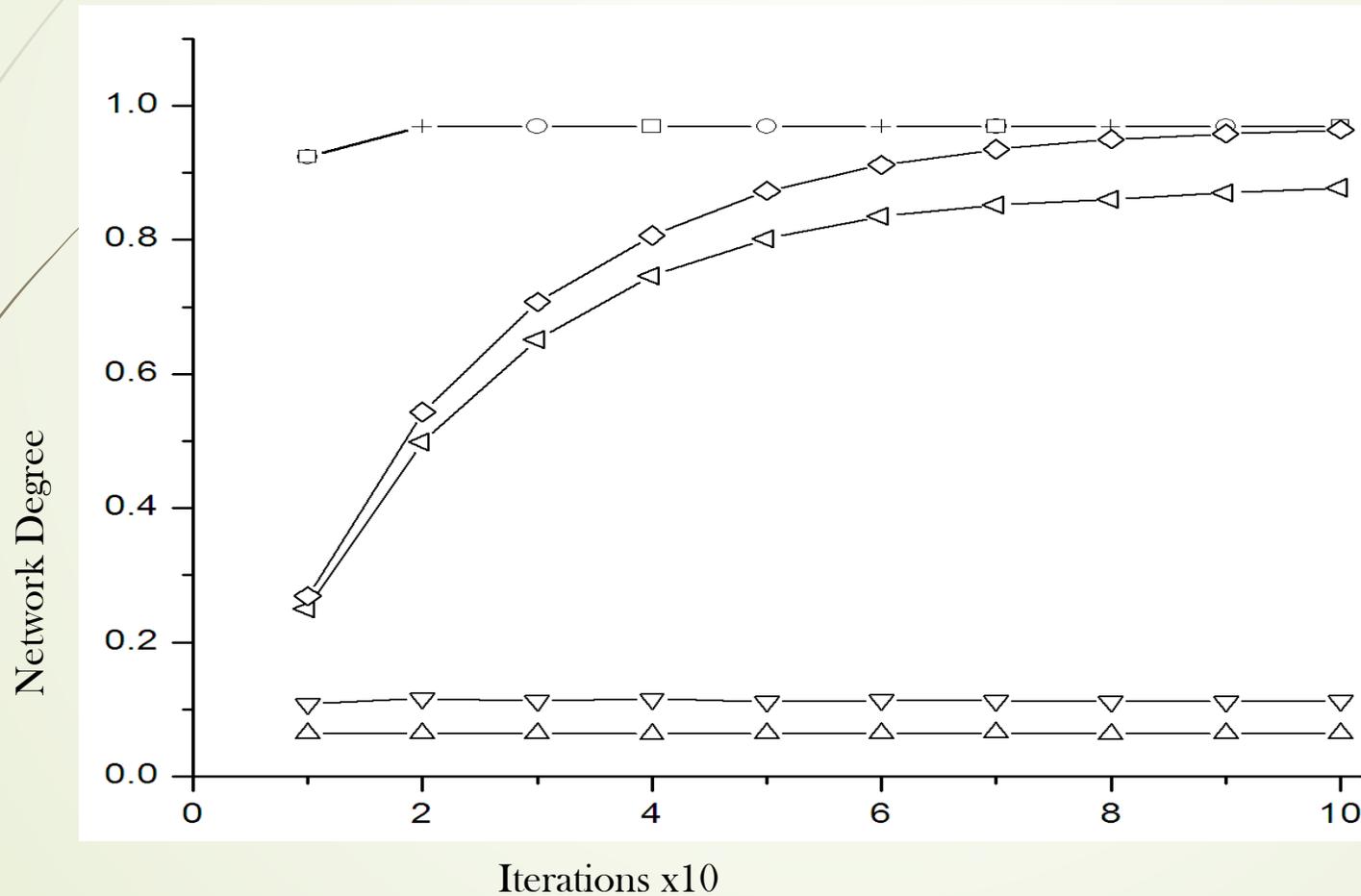
Already gives us some insight into evolutionary pressures on self-organizing social behaviors

*Discussed in Fefferman and Ng, 2007a  
and Hock and Fefferman 2011, and 2012*

Good abstract experimental system: separation in success across metrics and among populations with these traits

# What happens over time?

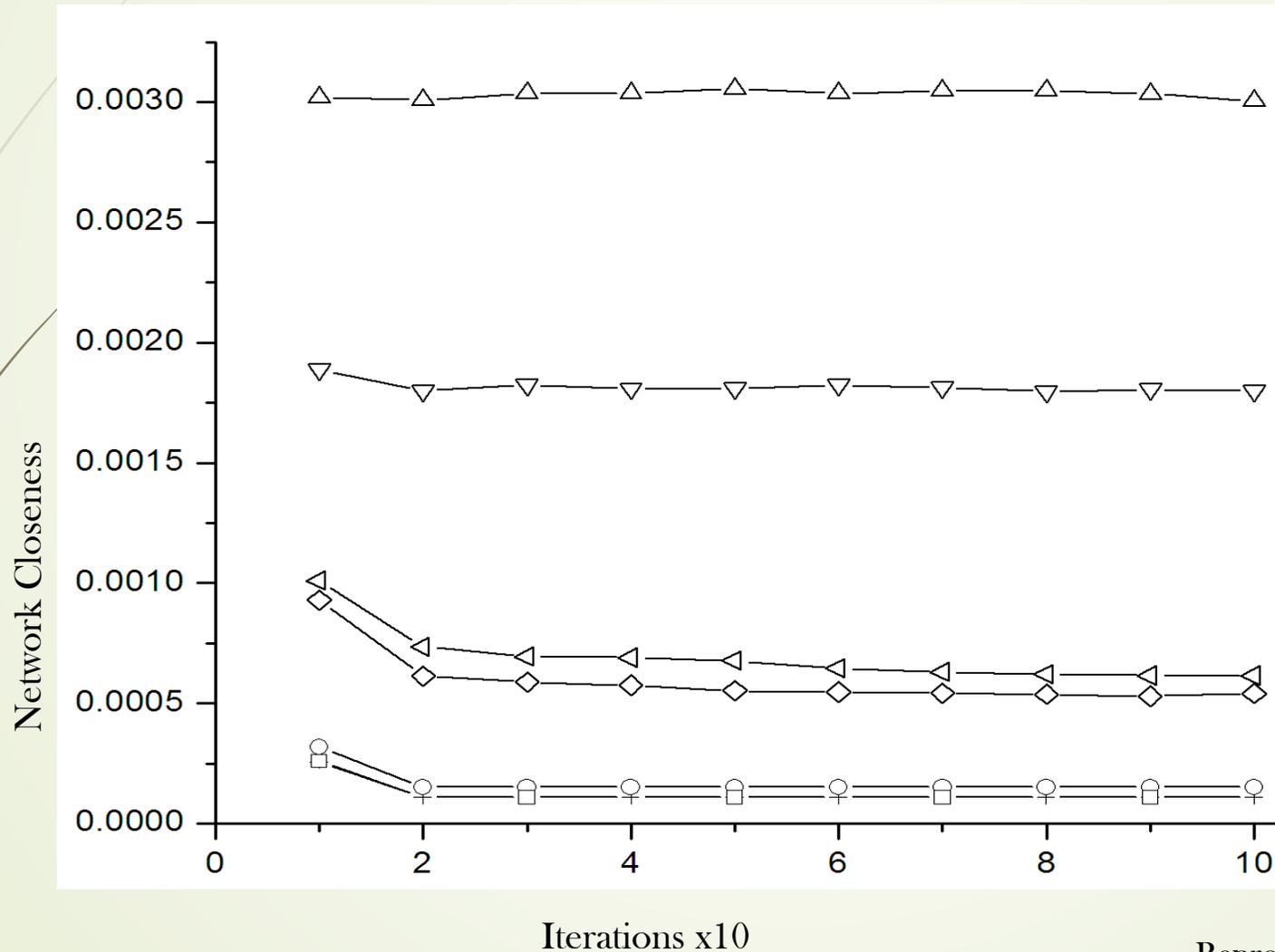
Measured Metric is Degree



- △ R
- ◁ B Incomplete
- ▽ C Incomplete
- ◇ D Incomplete
- B Complete
- C Complete
- +
- +

# And the stability and success under Closeness?

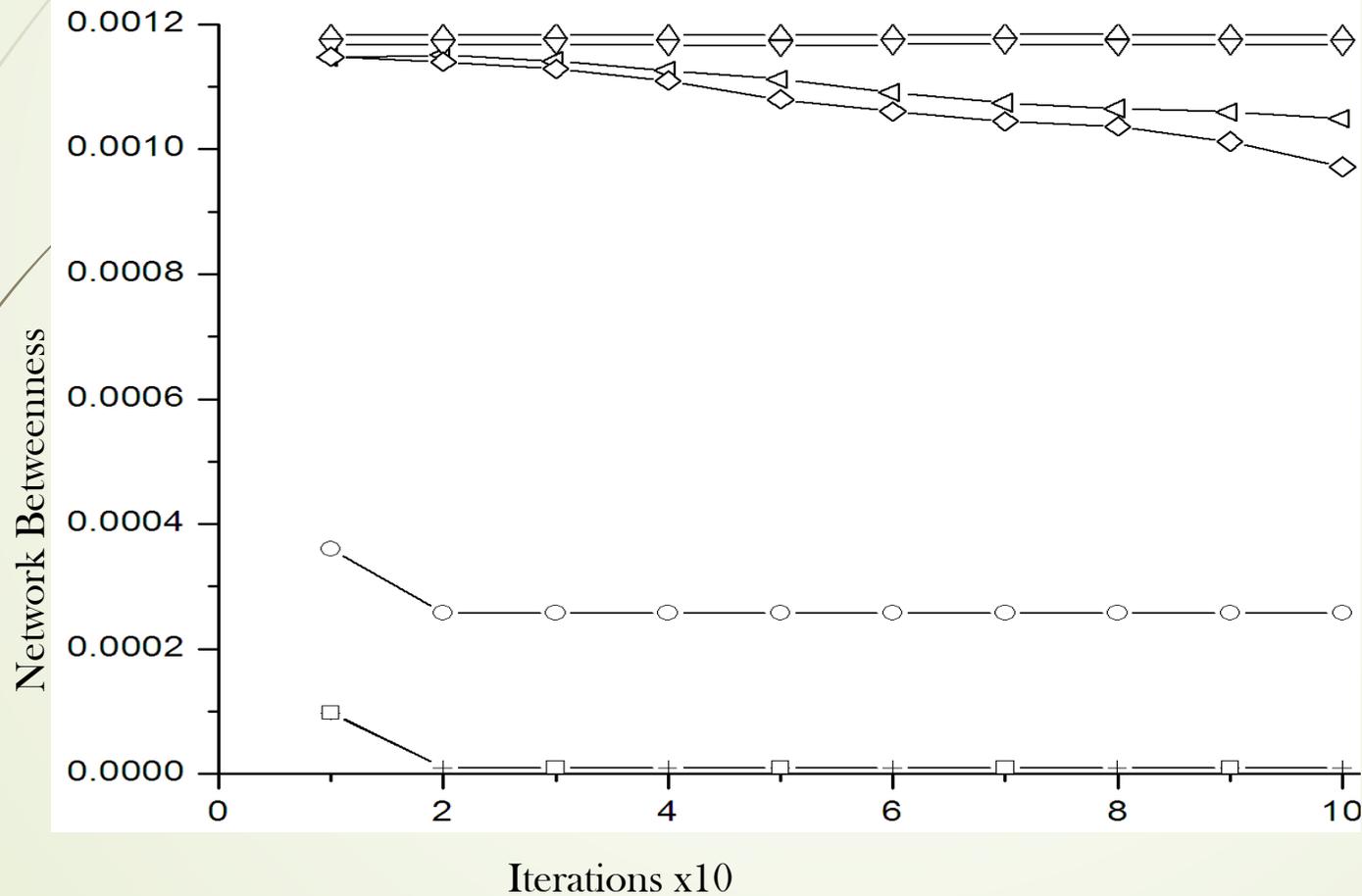
Measured Metric is Closeness



- △ R
- ◁ B Incomplete
- ▽ C Incomplete
- ◇ D Incomplete
- B Complete
- C Complete
- ⊕ D Complete

# And the stability and success under Betweenness?

Measured Metric is Betweenness

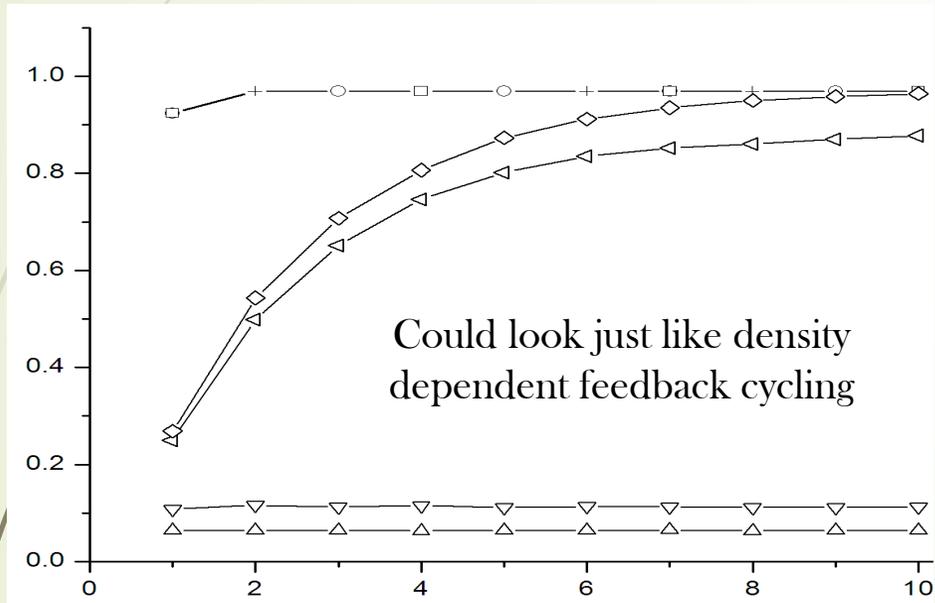


- △ R
- ◁ B Incomplete
- ▽ C Incomplete
- ◇ D Incomplete
- ◻ B Complete
- C Complete
- + D Complete

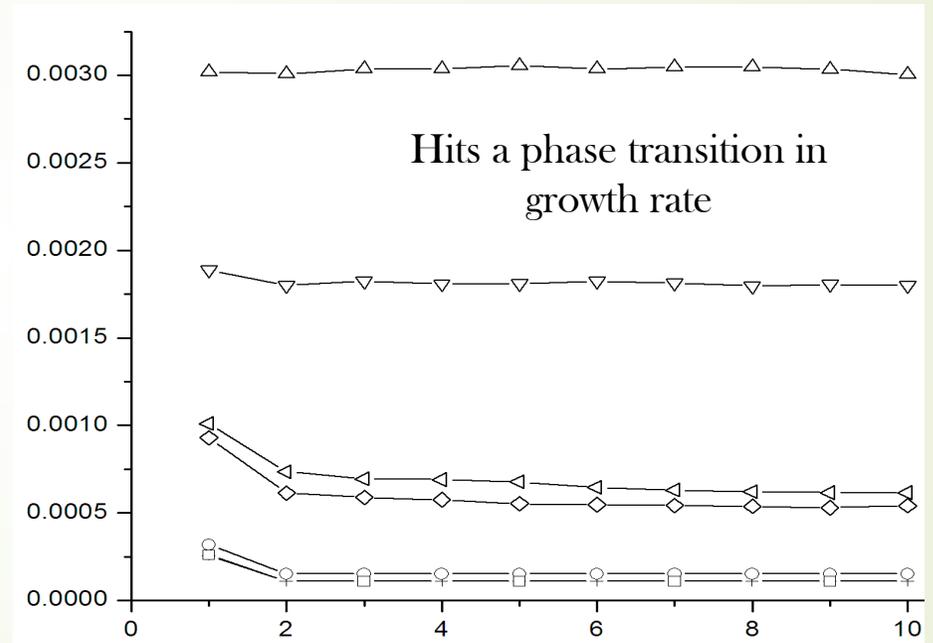
# Selection and Population Size

- △ R
- ◁ B Inc
- ▽ C Inc
- ◇ D Inc
- B Com
- C Com
- + D Com

## Degree-Driven Fitness



## Closeness-Driven Fitness



Already gives us some insight into evolutionary pressures on self-organizing social behaviors in populations of different sizes!

# Can this happen?

We would need at least one of the following:

- (1) Neutral outcomes - no net impact
- (2) Individuals do well by choosing things that accidentally are best for the group
- (3) Individuals may or may not benefit, but any costs are recouped by the distributed effects of group benefits

Do any of these 3 things happen?

In math language: Do global feedback rules to local decisions prohibit this type of emergence?

In bio language: Can this type of system evolve by natural selection?

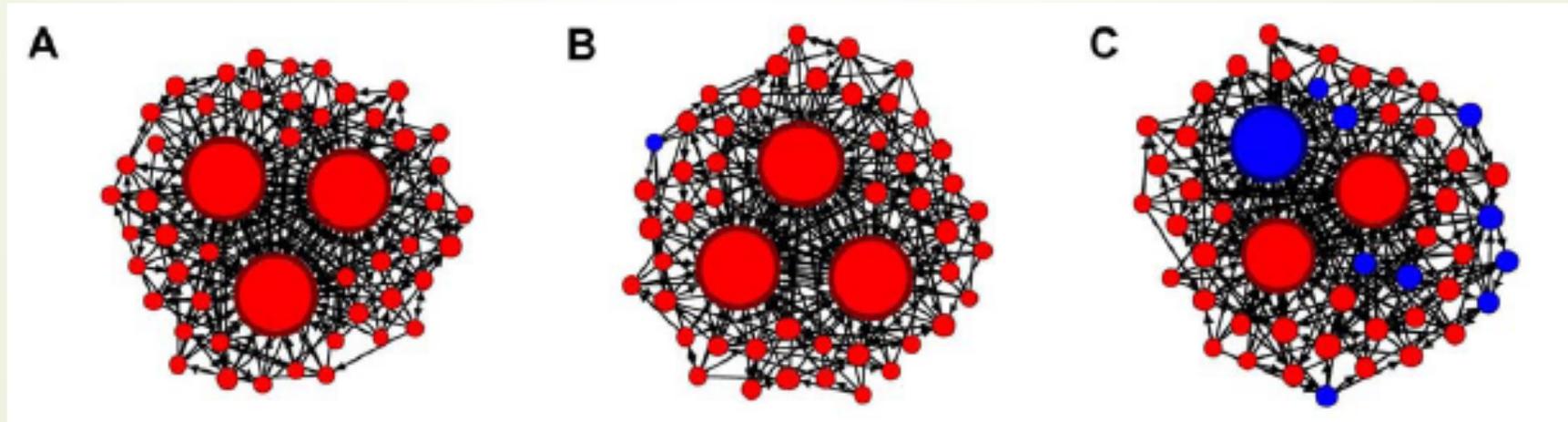
# Answers Come from Studying Rule-Breakers

How do individuals do if they follow the Degree self-organizing rule or not?

0% Rule Breakers

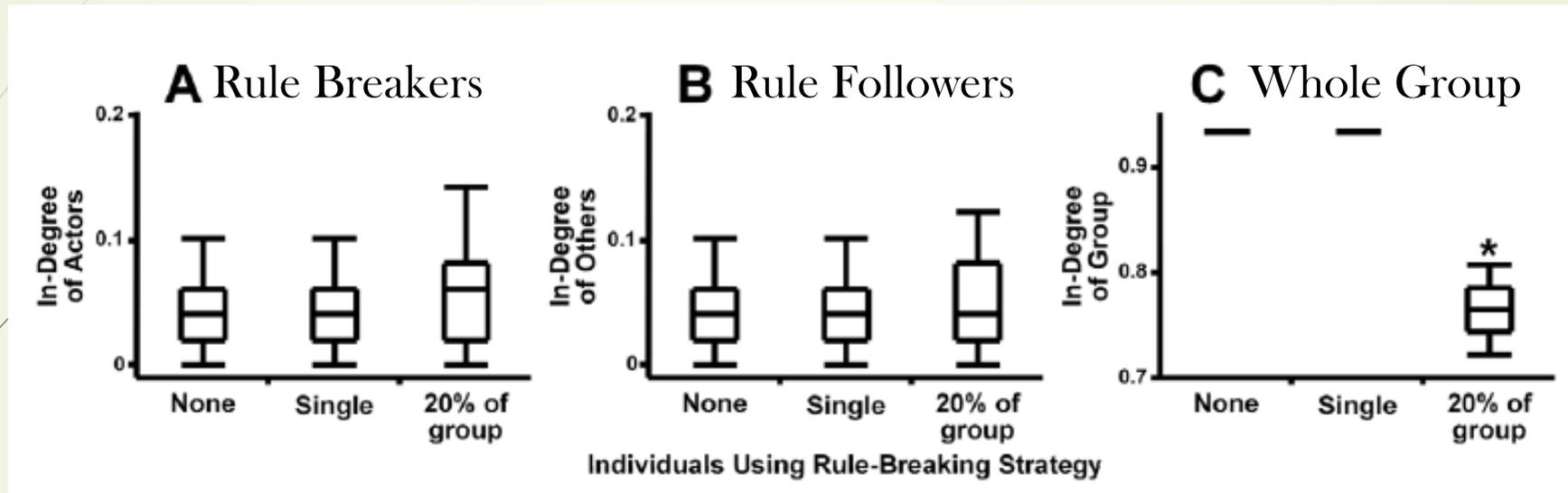
2% Rule Breakers

20% Rule Breakers



Size is Scaled by Degree

## Answer for Degree: Neutral



Individuals do just as well if they do or do not participate in the self-organizing rule of Degree

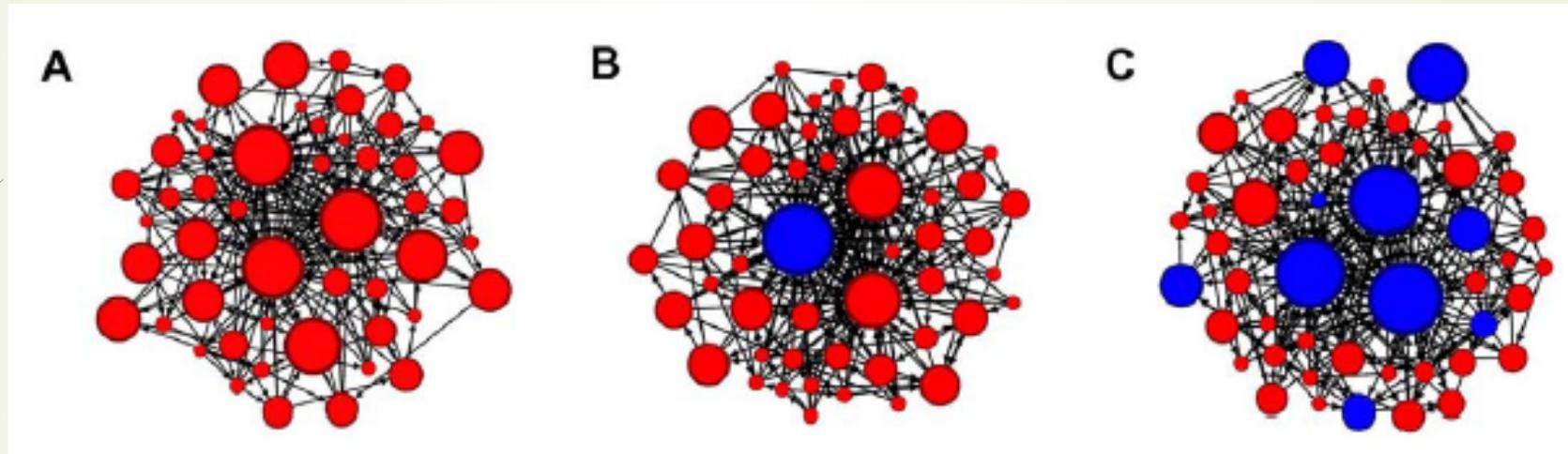
**HOWEVER**, rule-breakers change overall group organization *even though* individual values don't change (*means evolutionary incentive to participate*)

# What About Following the Betweenness Self-Organizing Rule?

0% Rule Breakers

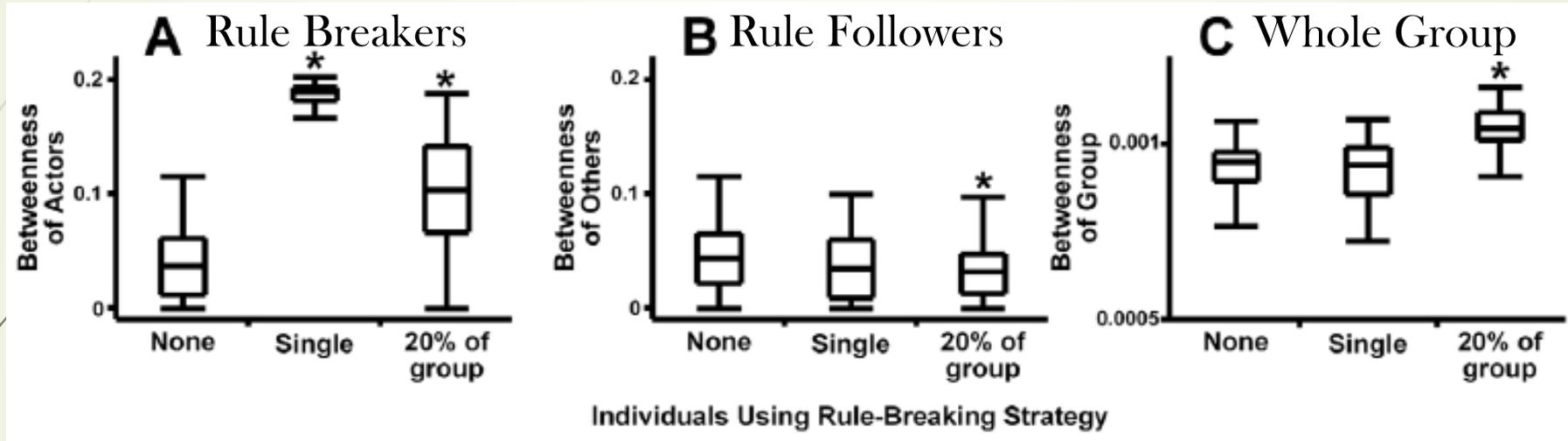
2% Rule Breakers

20% Rule Breakers



Size is Scaled by Betweenness

## Answer for Betweenness: **Net Benefit**



Small numbers of rule breakers do better, large numbers of rule breakers mean only some do better

Consistent with the previous group-level analyses, with more rule-breakers, the whole population did better - evolution of this system may be unstable

# What have we built? Not a model of a real system

Experimental system with nice properties

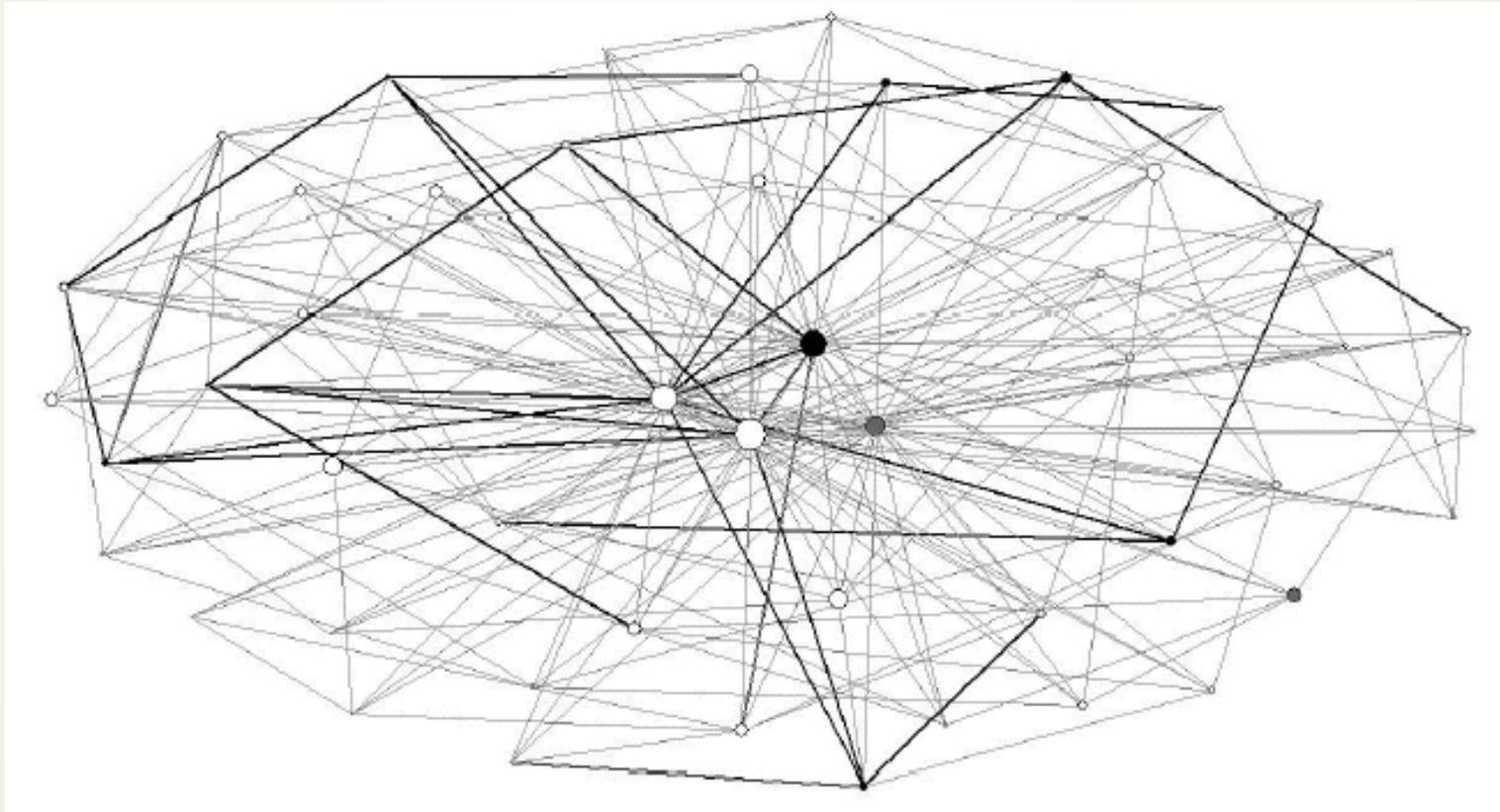
- Self-organizing behaviors
- Individual benefits
- Group benefits

Global outcomes provide mechanism of control on evolution of social traits

- Insights into organizational success and population size
- Evolution of cooperation
- *Quantitative predictions about stability of self-organizing systems*

# Group Success May Not Be Our Network-Level Outcome - Lessons in Epidemiology!

Simulated Disease Process (SEIIS) on All Population Types:





# A Few Quick Cool Things I Won't Explain Today:

Can be found in Fefferman and Ng, 2007; Hock and Fefferman, 2011.

Super-spreaders aren't always high degree individuals in networks with ongoing self-organization!

Computational experiments on coupled stochastic processes are tricky - use a complete graph as a way to cheat the number of model realizations needed for network evolution AND epidemic spread

Different self-organizing strategies increase population robustness against disease at different probabilities of per-contact transmission

Increased network centrality doesn't always correlate with increased disease burden

## One Thing I Will Talk About Quickly:

Details in Fefferman and Ng, 2007

Ongoing self-organized rewiring dynamics were *protective* against disease

# Disease on Self-Organizing Networks

	3-Way Test	B-population		C-population		D-population	
		Dynamic	Static	Dynamic	Static	Dynamic	Static
<b>B</b>	Dynamic <sup>†</sup>	B static > B dynamic <sup>†</sup>		< <sup>*</sup>	< <sup>*</sup>	>	> <sup>*</sup>
<b>C</b>	Static <sup>†</sup>			C static > C dynamic <sup>†</sup>		> <sup>*</sup>	> <sup>*</sup>
<b>D</b>						D static > D dynamic <sup>†</sup>	
		Overall: Dynamic C>B>D; Static C>B>D					

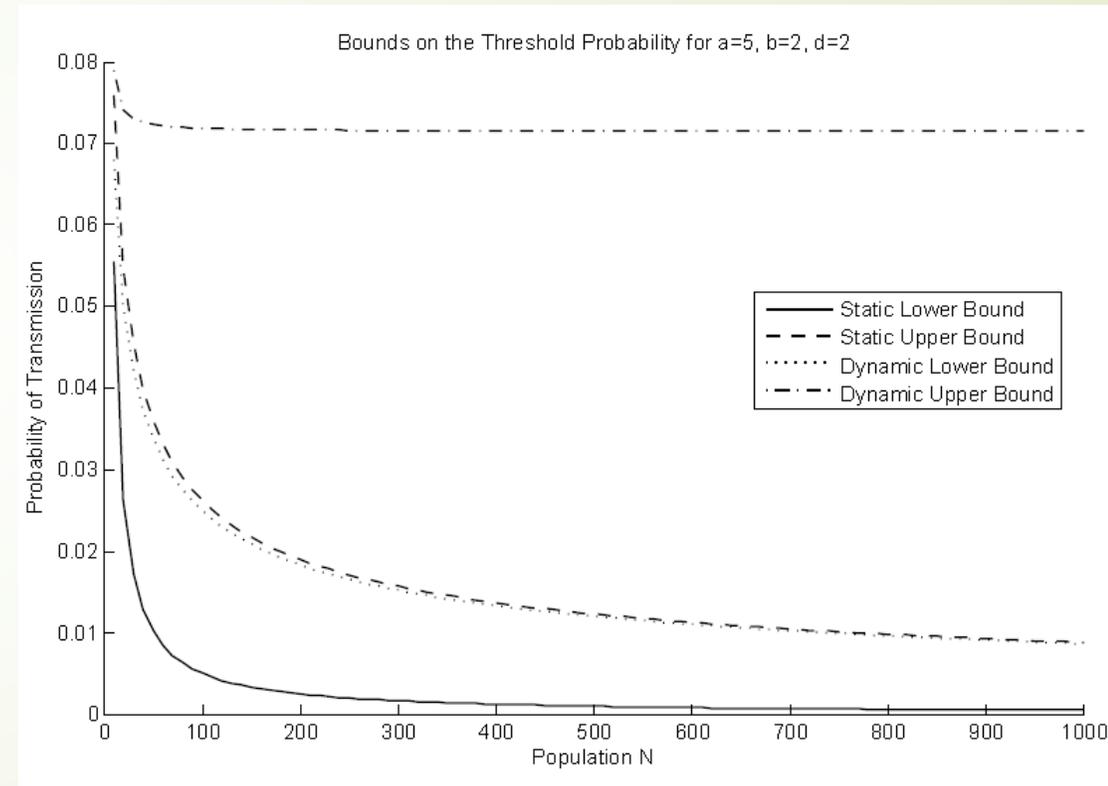
*Reproduced from Fefferman and Ng, 2007b*

Finally! We discovered computationally something we can prove analytically!

# I'm actually not going to go through the proof

## The Intuition for how

- Instead of estimating numbers of infections, focus on threshold in transmissibility that implies transmission will occur on the current network state
- Calculate limit of recurrence relation of shifting states
- Compare boundaries of that threshold for static and dynamic network recurrences





# These systems are analytically challenging

(even in the simple, deterministic cases)

- Bi-directionally coupled
  - Coupling occurs at multiple scales
  - Scale of coupling can (and usually is) asymmetric
  - Frequently discrete in individual action, though continuous in global dynamics
- 



# When should we even bother looking for analytic approximations?

- Can we reduce the dynamics or shift the outcome variable to something with a well-defined recurrence relation?
  - Is temporal order critical to individual dynamics?
  - Can we approximate the global behavior without finding a good approximation for individual dynamics under any circumstances?
- 



## More Importantly:

There are reasons we can discuss analytic solutions for anything about this:

- We constructed the simulation to be a mathematically controllable abstraction
- It has minimal complexity for the features we need
- The observed behaviors come from single differences in action/assumption



# Potential Recommendation for Studying Self-Organization

(definitely not for everyone or every problem)

We all have a tendency to model actual biological systems

Instead, sometimes we might want to try modeling the simplest systems with isolated, abstract behaviors in mathematically controllable ways and *then* gradually put back the realism as we need it to gain further insight



# Where some of these things (and more on this topic) are published:

- Brooks, H.Z., M.E. Hohn, C. Price, A.E. Radunskaya, S.S. Sindi, N.D. Williams, S.N. Wilson, N.H. Fefferman. 2018. Springer.
- Williams, N.D., H.Z. Brooks, M.E. Hohn, C. R. Price, A.E. Radunskaya, S.S. Sindi, S.N. Wilson, and N. H. Fefferman. 2018. Springer.
- Gallos, L., and N.H. Fefferman. 2015. *PLoS One*.
- Greening, B. and N.H. Fefferman. 2014. *Nature Sci Rep*.
- Hock, K. and N.H. Fefferman. 2012. *Ecological Complexity*.
- Hock, K. and N.H. Fefferman. 2011. *PLoS One*.
- Hock, K. and N.H. Fefferman. 2011. *Ann Zoo Fen*.
- Hock, K., K.L. Ng, and N.H. Fefferman. 2010. *PLoS One*.
- Fefferman, N.H. and K.L. Ng. 2007. *Physical Review E*.
- Fefferman, N.H. and K.L. Ng. 2007. *Ann Zoo Fen*.

# Talented Researchers of the Fefferman Lab:



**Post docs:** Dr. Erick Chastain, Dr. Jing Jiao (starting soon!), Dr. Kellen Myers, Dr. Nourridine Siewe, Dr. Gonzalo Suarez, and Dr. Oyita Udiani (Former post docs pictured because their work was presented: Dr. Karlo Hock and Dr. Kah Loon Ng)

**Grad Students:** J. Beck, J. DeSalu, A. Redere

**Funders to the lab:** NSF, NIH, DHS, DoD, USDA, USFWS