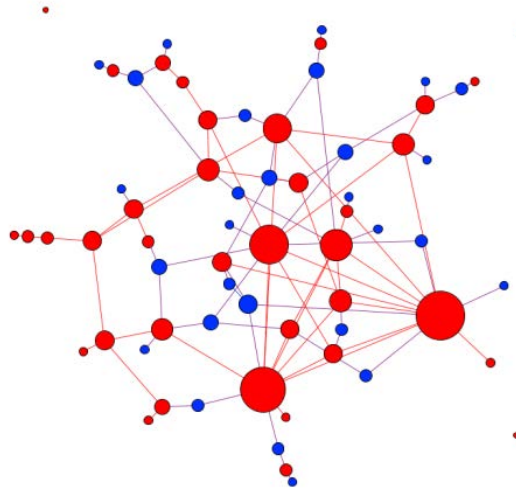
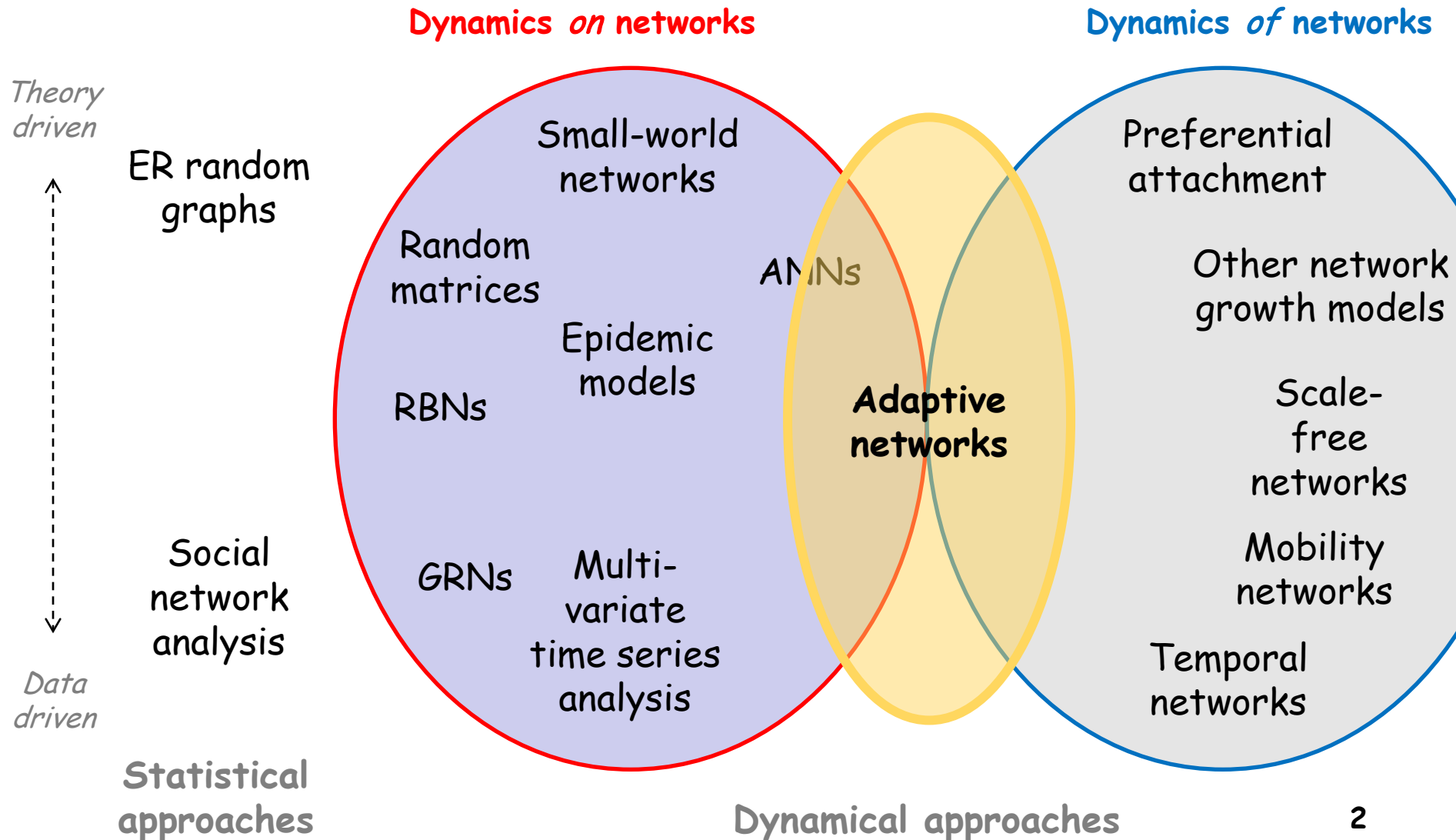


# Simulation III: Adaptive Networks



**Hiroki Sayama**  
sayama@binghamton.edu

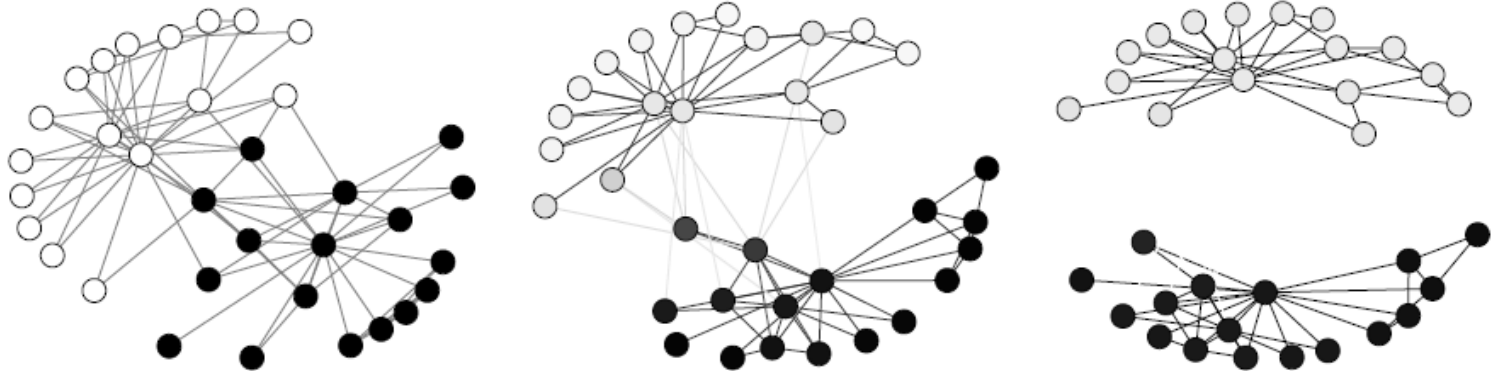
# A map of network science



# Adaptive networks

---

- Complex networks whose states and topologies co-evolve, often over similar time scales
  - Link (node) states adaptively change according to node (link) states



# Adaptive networks in action

---

- Many real-world complex systems show coupling between “dynamics *of* networks” and “dynamics *on* networks”
- 

| System                | Nodes           | Edges                                | States of nodes  | Topological changes  |
|-----------------------|-----------------|--------------------------------------|--|--|
| Organism              | Cells           | Intercellular communication channels | Gene/protein activities  | Fission and death of cells during development                        |
| Ecological community  | Species         | Interspecific relationships          | Population   | Speciation, invasion, extinction of species                          |
| Human society         | Individual      | Conversations, social relationships  | Social, professional, economical, political, cultural statuses | Changes in social relationships, entry and withdrawal of individuals |
| Communication network | Terminals, hubs | Cables, wireless connections         | Information stored and transacted                              | Addition and removal of terminal or hub nodes                        |

# Simulation of Adaptive Networks

# Simulating state-topology coevolution

---

- Technically, very easy; not so much different from other network simulation models
- One minor problem:  
How to handle topological changes while state changes are also ongoing?  
→ Asynchronous updating

# Example: Epidemics on adaptive networks

---

- Original epidemic network model  
+ adaptive changes of links
- A susceptible node that has a link to an infected node will cut the link and reconnect it to another susceptible node with probability  $p_c$
- Does the disease stay in the network?

# Exercise

---

- Study the effects of rewiring probability on the disease fixation on and the global network structure of an initially random social network
  - In what condition will the disease remain within society?
  - How will the topology of the network be reformed through the disease propagation process?



# Example: Adaptive voter model

- Original voter model
  - + adaptive disconnection of links
- A link that connects two nodes with different opinion states may be cut with probability  $p_c$
- How will the social network and opinions evolve?

# Exercise

---

- Study the effects of the link disconnection probability on the consensus formation in the adaptive voter model
  - Plot the final number of opinions as a function of  $p_c$
  - How will the topology of the network be changed by the diversity of opinions?

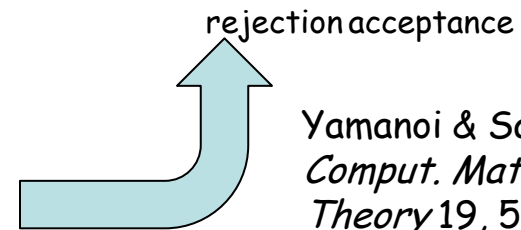
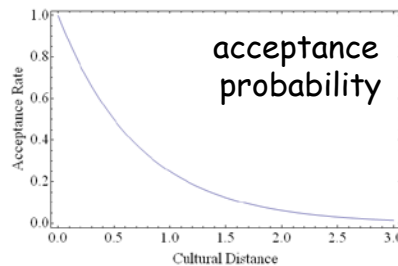
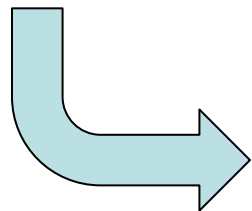
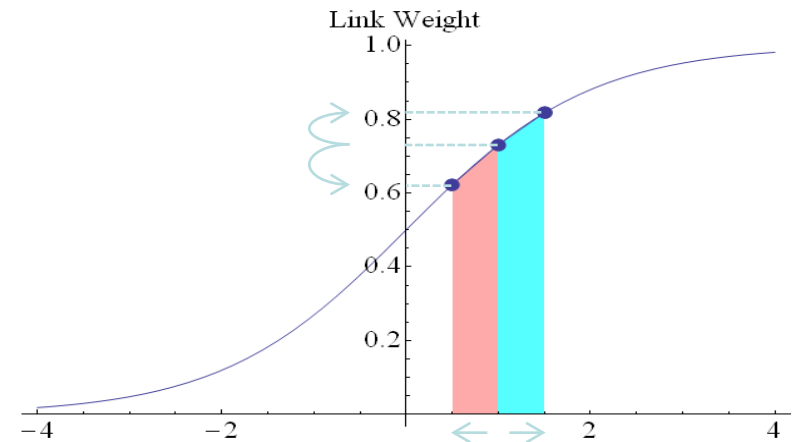
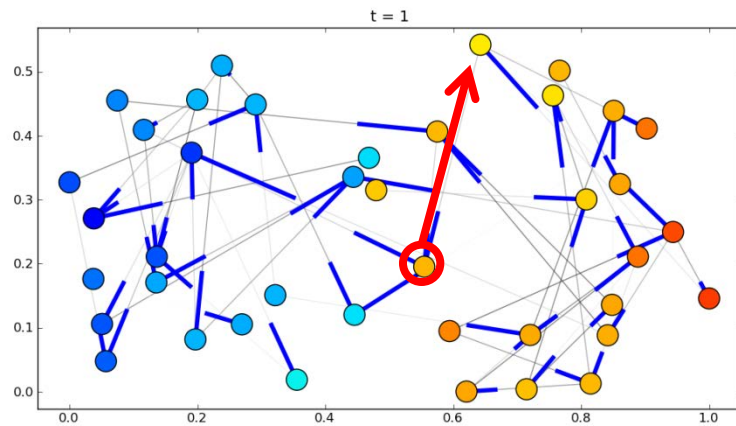
# Example: Adaptive diffusion model

---

- Original diffusion model
  - + adaptive disconnection of links
- Link weights will increase or decrease based on the similarity/dissimilarity of node states across the links
  - Conceptually similar to the adaptive voter model

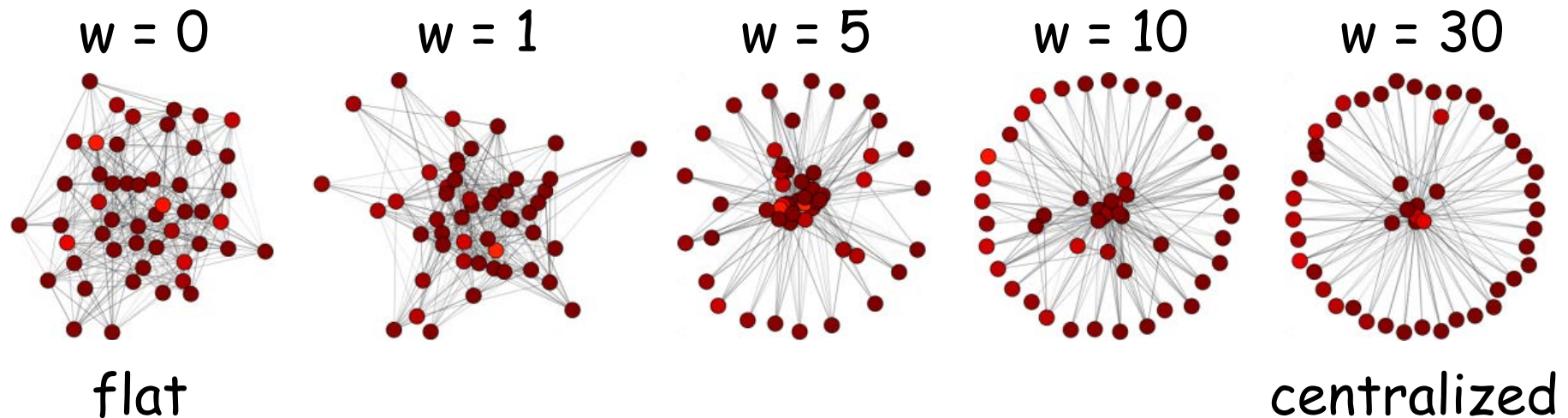
# Application 1: Corporate merger

- Modeling and simulation of cultural integration in two merging firms



Yamanoi & Sayama,  
*Comput. Math. Org.  
Theory* 19, 516-537,  
2013.

# “Within-firm” concentration (w)

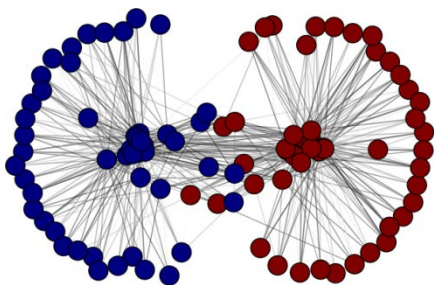


- Prob. for node  $i$  to become an info source:

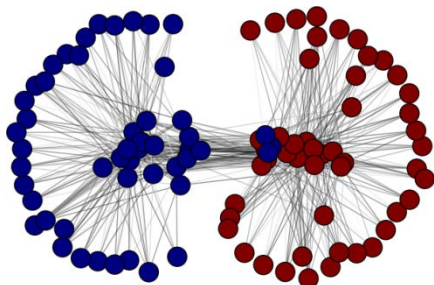
$$P_w(i) \sim (i/n)^w \quad (i = 1, 2, \dots, n; \quad n = \text{firm size})$$

# “Between-firm” concentration (b)

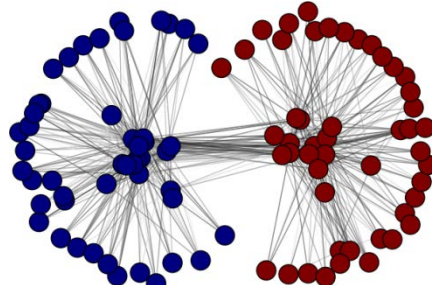
$b = 0.1$



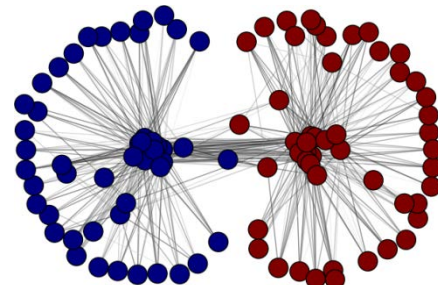
$b = 1$



$b = 3$



$b = 5$



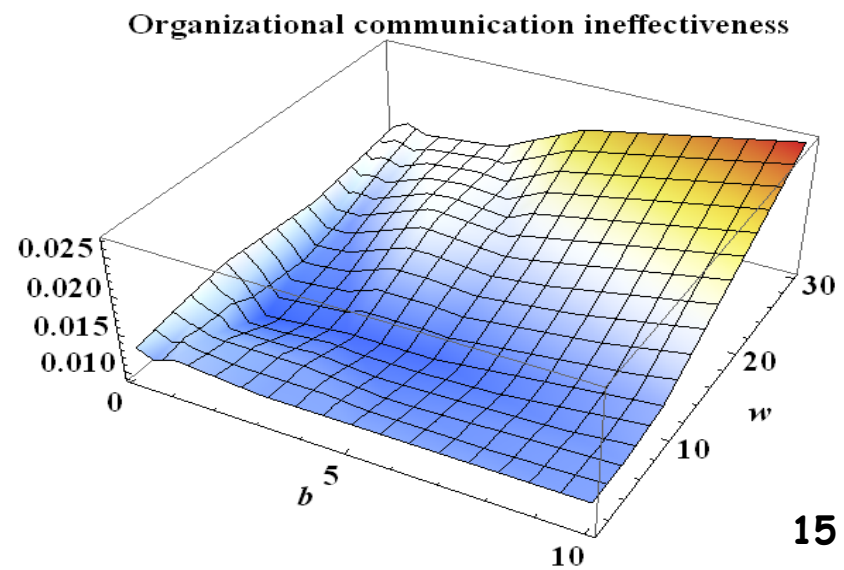
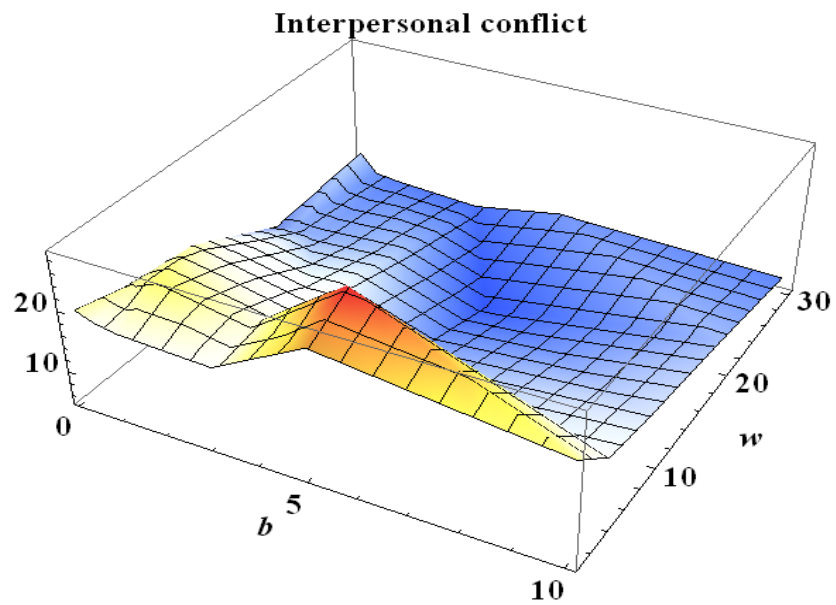
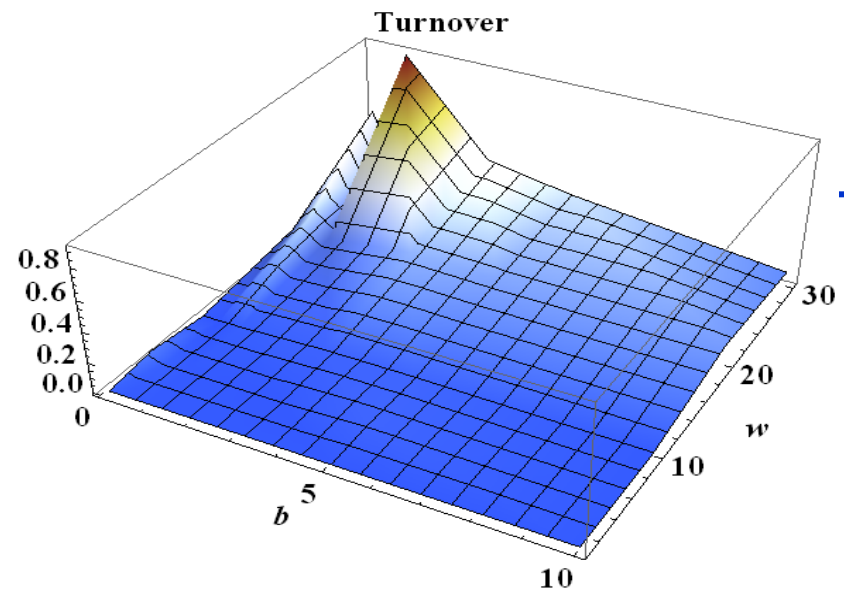
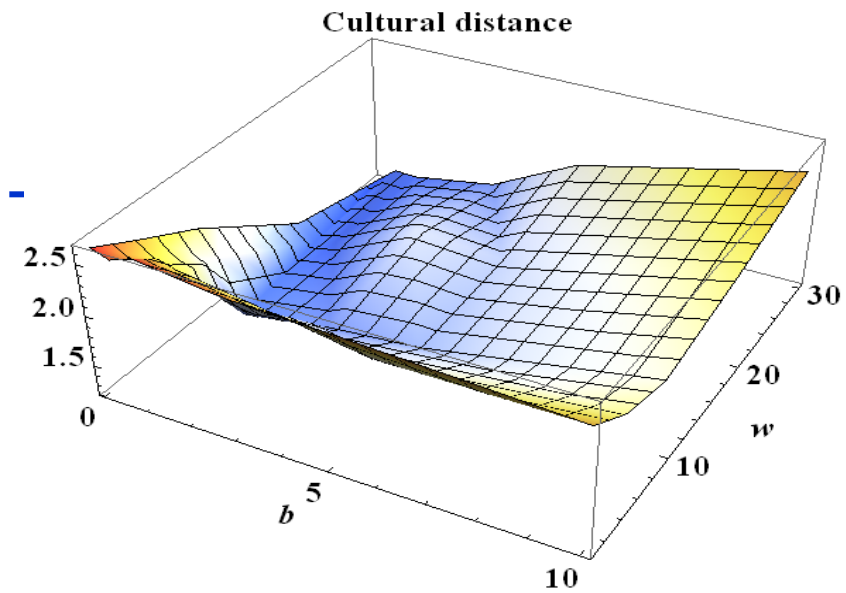
nearly random

executive-level

- Prob. for node  $i$  to have an inter-firm tie:

$$P_b(i) \sim c_i^b$$

( $c_i$  = within-firm closeness centrality of  $i$ )





# Application 2: Social diffusion and global drift

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- Sayama & Sinatra, PRE 91, 032809, 2015

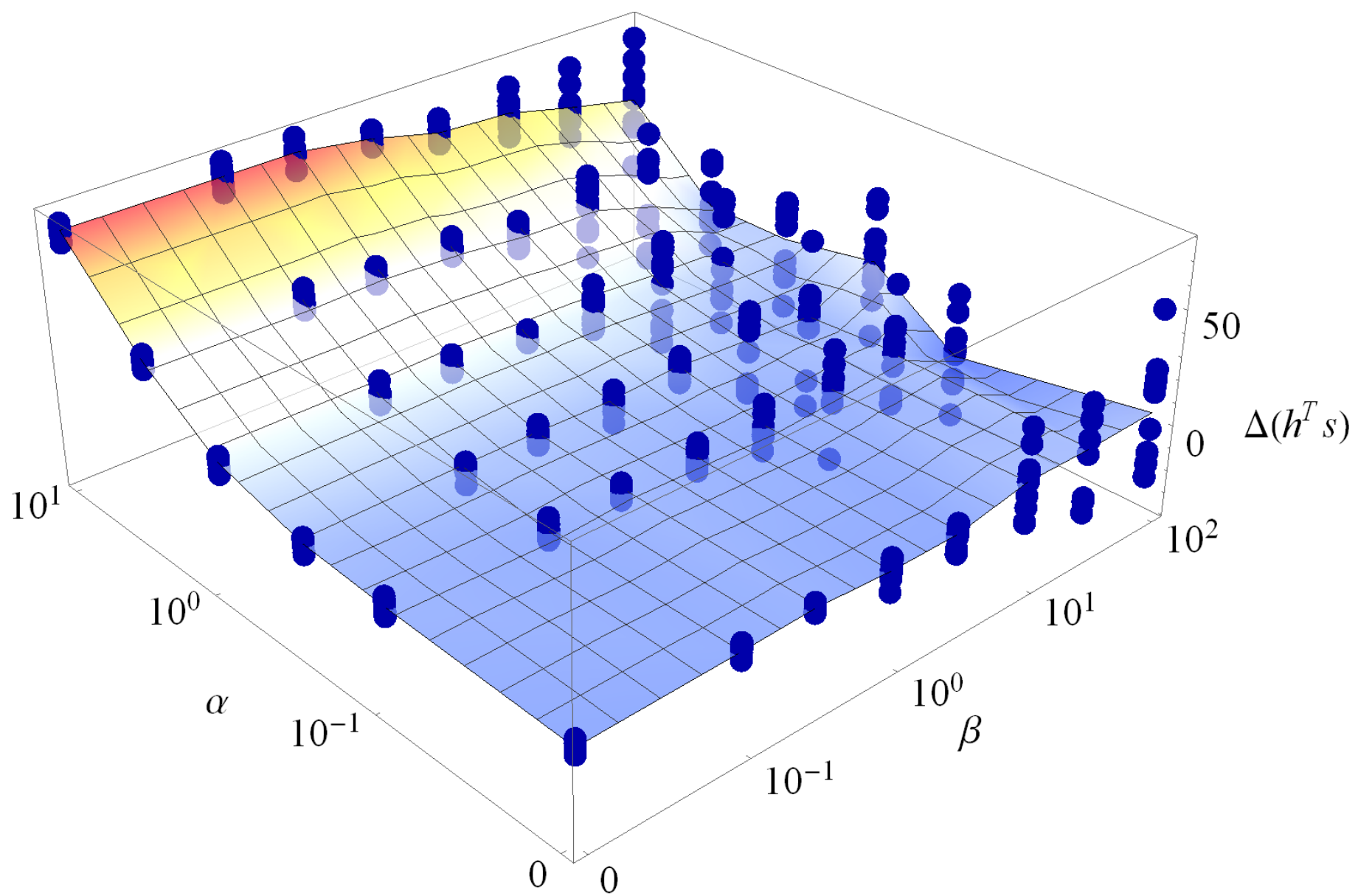
$$\frac{ds_i}{dt} = c(\langle s_j \rangle_j^i - s_i)$$



**Adaptive link weight adjustment:**

$$\frac{da_{ij}}{dt} = a_{ij} \left[ \alpha \frac{s_i + s_j - 2\langle s \rangle}{2\sigma_s} - \beta \frac{(k_i - \langle k \rangle)(k_j - \langle k \rangle)}{\sigma_k^2} \right]$$





# Exercise

---

- Change the rule of link weight adjustment in the adaptive diffusion model
  - E.g., Sayama & Sinatra (2015)
- Simulate the revised model and see how the network topology and state co-evolve

# Theoretical Framework:

## Generative Network Automata

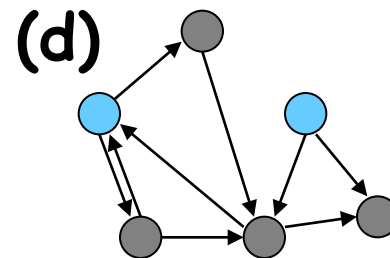
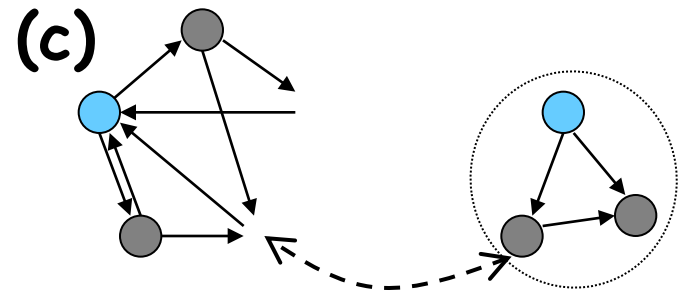
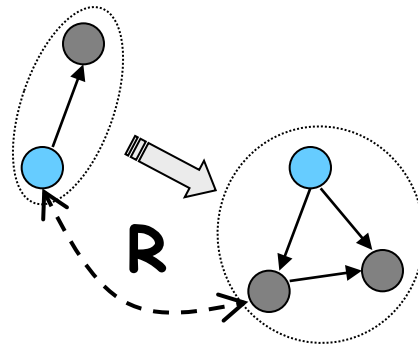
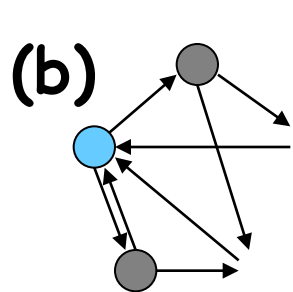
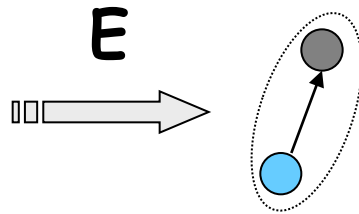
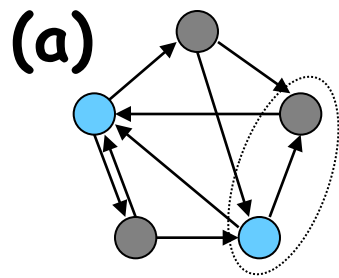
# Generative network automata

---

- Unified representation of dynamics *on* and *of* networks using graph rewriting
- Defined by  $\langle E, R, I \rangle$ :
  - E : Extraction mechanism — When, Where
  - R : Replacement mechanism — What
  - I : Initial configuration

*Sayama, Proc. 1st IEEE Symp. Artif. Life, 2007, pp.214-221.*

# GNA rewriting example

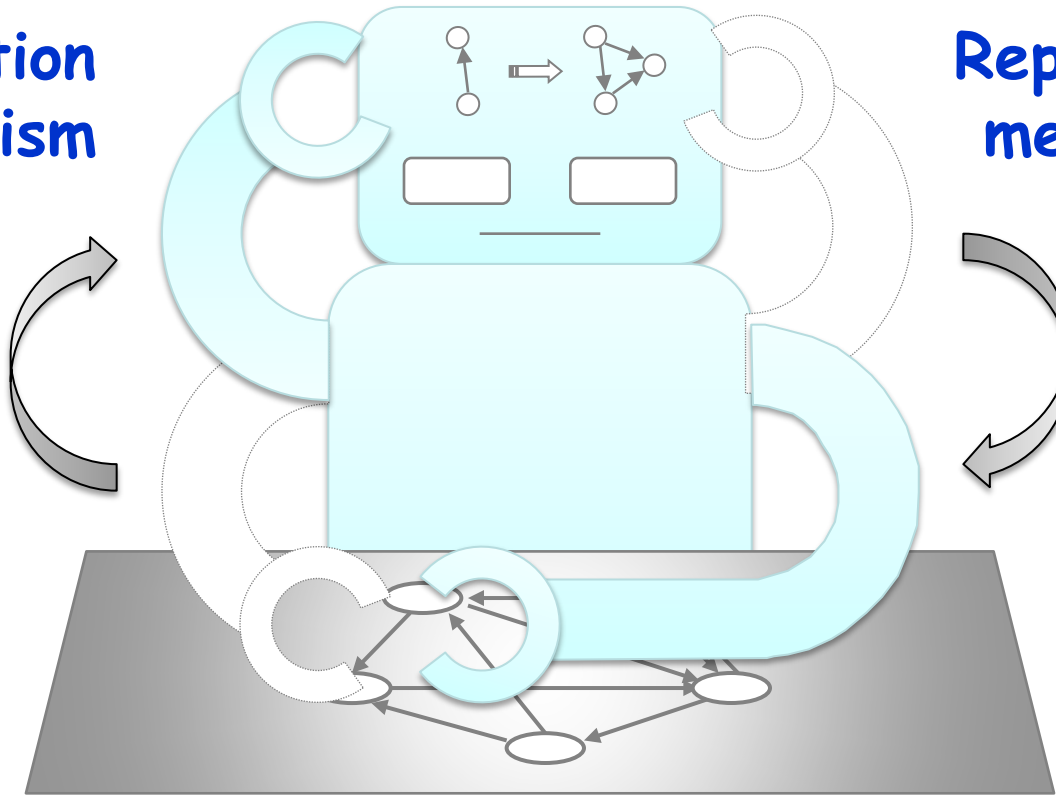


# Actually, it's a generative network automata~~x~~-on

---

**E :**  
**Extraction**  
**mechanism**

**R:**  
**Replacement**  
**mechanism**

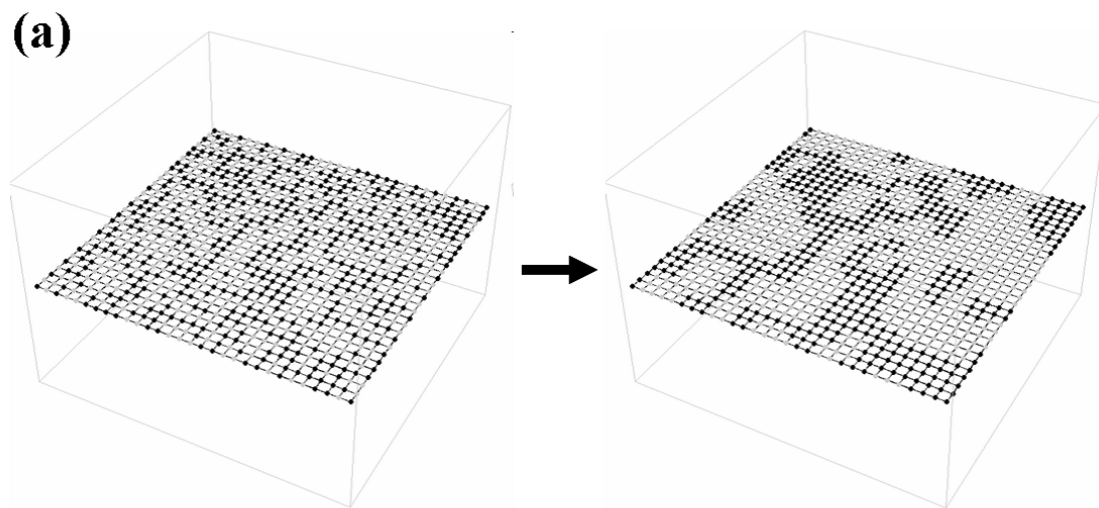


# Generality of GNA

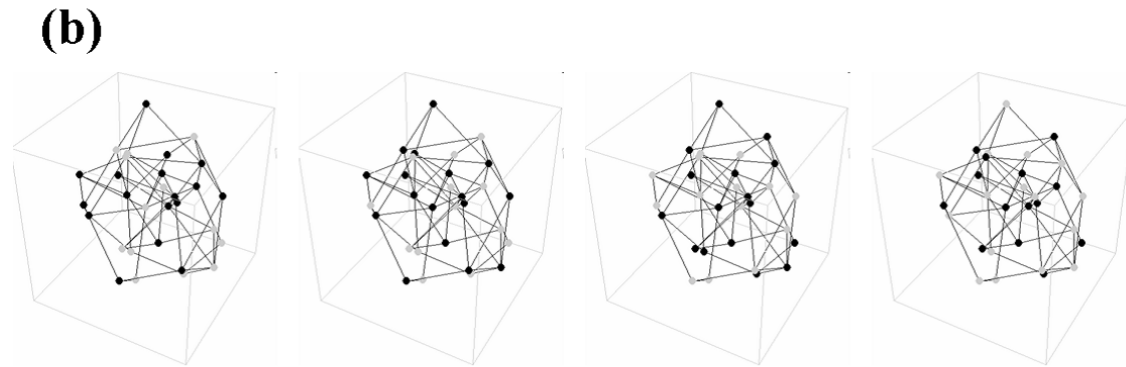
---

- GNA can uniformly represent in  $\langle E, R, I \rangle$ :
  - Conventional dynamical systems models
    - If R always conserves local network topologies and modifies states of nodes only
    - E.g. cellular automata, Boolean networks
  - Complex network generation models
    - If R causes no change in local states of nodes and modifies topologies of networks only
    - E.g. small-world, scale-free networks

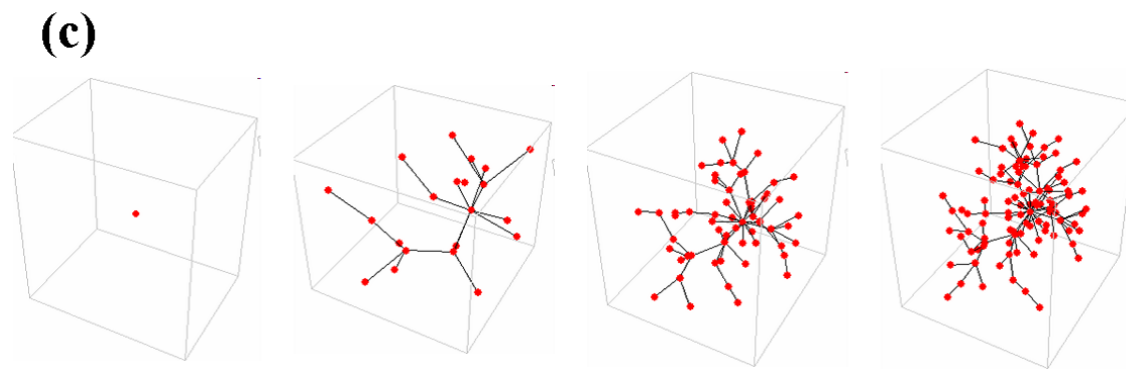
**Cellular  
automata**



**Random  
Boolean  
network**



**BA  
scale-free  
network**



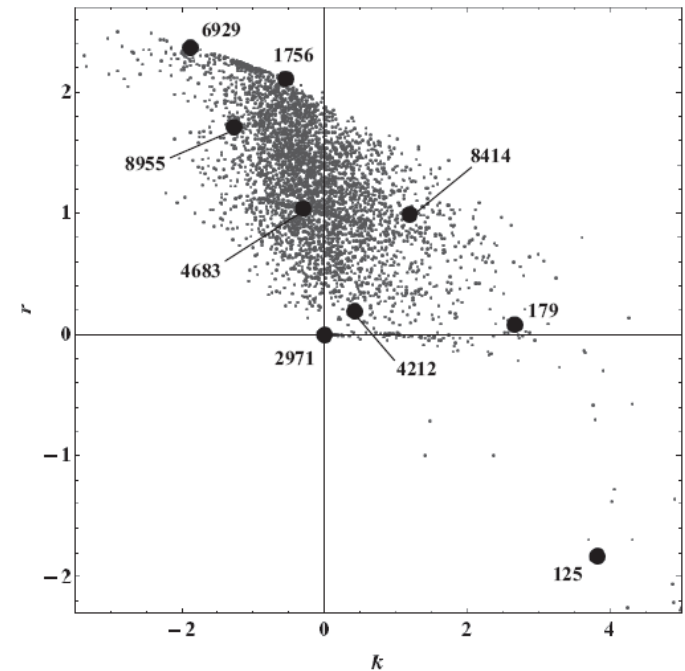
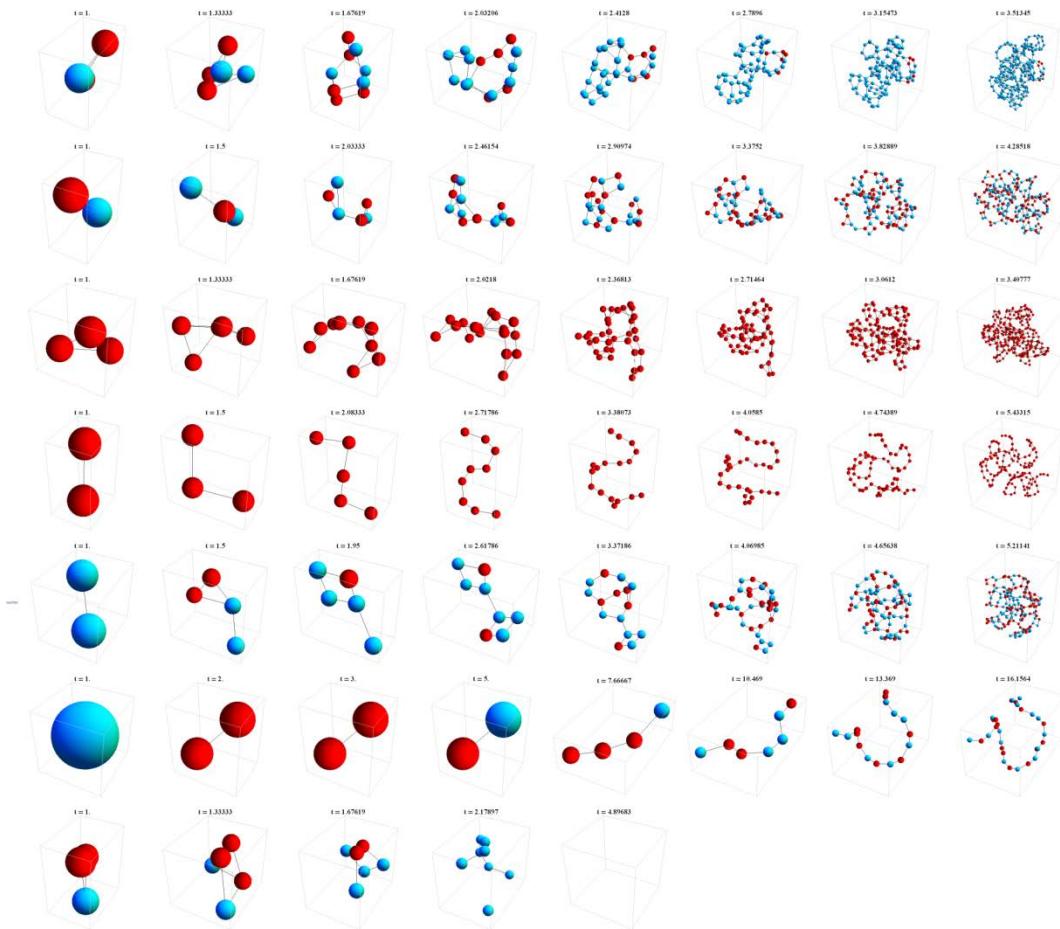


# Exhaustive search of rules

---

- E samples a node randomly and then extracts an induced subgraph around it
- R takes 2-bit inputs (states of the node and neighbors) and makes 1-out-of-10 decisions
  - Total number of possible R's:  $10^{2^2} = 10,000$
- “Rule Number”  $rn(R)$  is defined by
$$rn(R) = a_{11} 10^3 + a_{10} 10^2 + a_{01} 10^1 + a_{00} 10^0$$
  - $a_{ij} \in \{0, 1, \dots, 9\}$  : Choices of R when state of u is i and local majority state is j

# Exhaustive search of rules



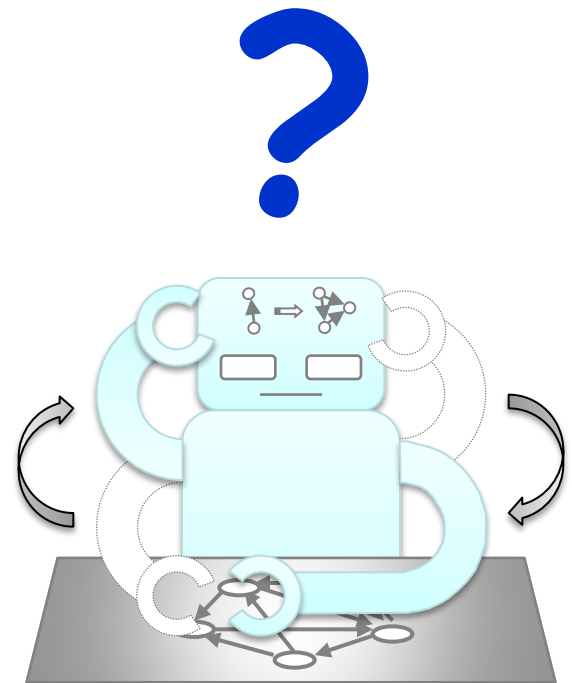
Sayama & Laramée, *Adaptive Networks*, Springer, 2009, pp.311-332.

# Developing Adaptive Network Models from Empirical Data

# A challenge

---

- How to derive a set of dynamical rules directly from empirical data of network evolution?
- Separation of extraction and rewriting in GNA helps the rule discovery



Pestov, Sayama, & Wong, *Proc. 9th Intl. Conf. Model. Simul. Visual. Methods*, 2012.

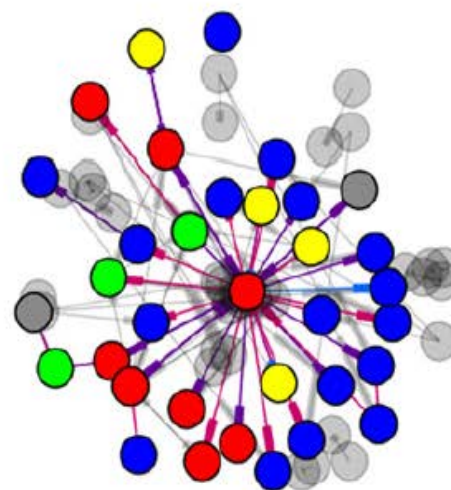
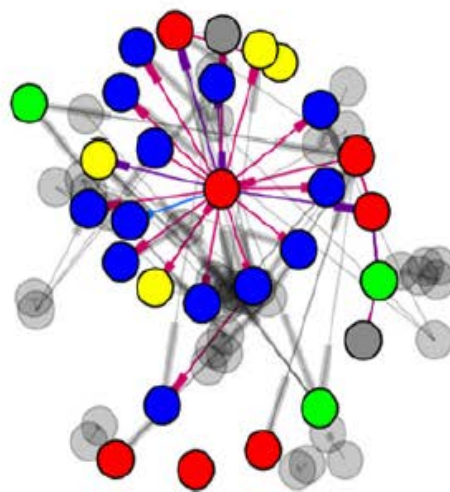
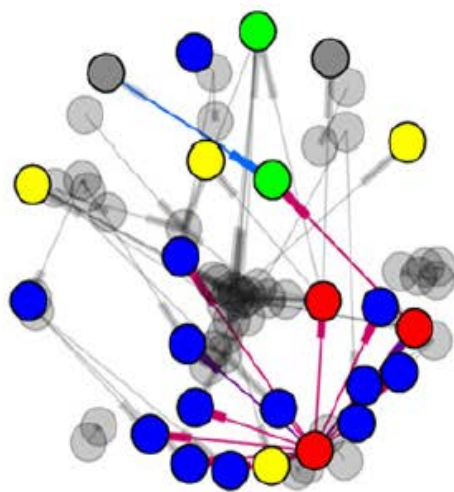
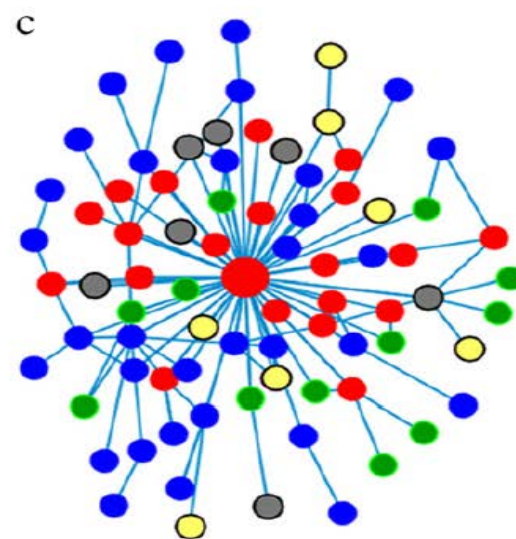
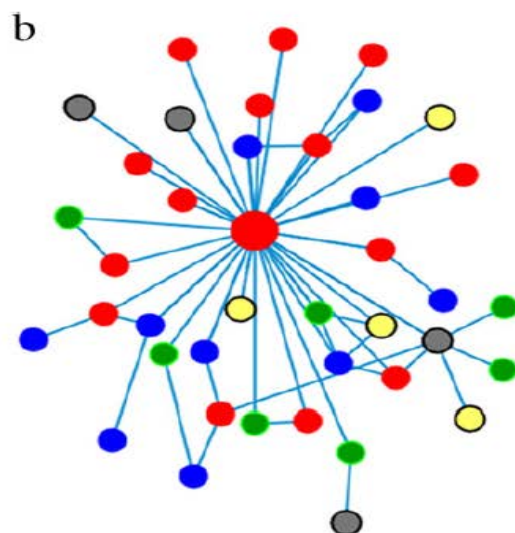
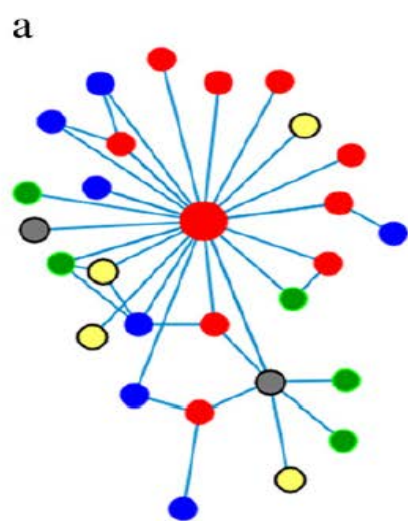
Schmidt & Sayama, *Proc. 4th IEEE Symp. Artif. Life*, 2013, pp.27-34.

# Application to operational network modeling

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- Canadian Arctic SAR (Search And Rescue) operational network
  - Rewriting rules manually built directly from actual communication log of a December 2008 SAR incident
  - Developed a simulator for hypothetical SAR operational network development



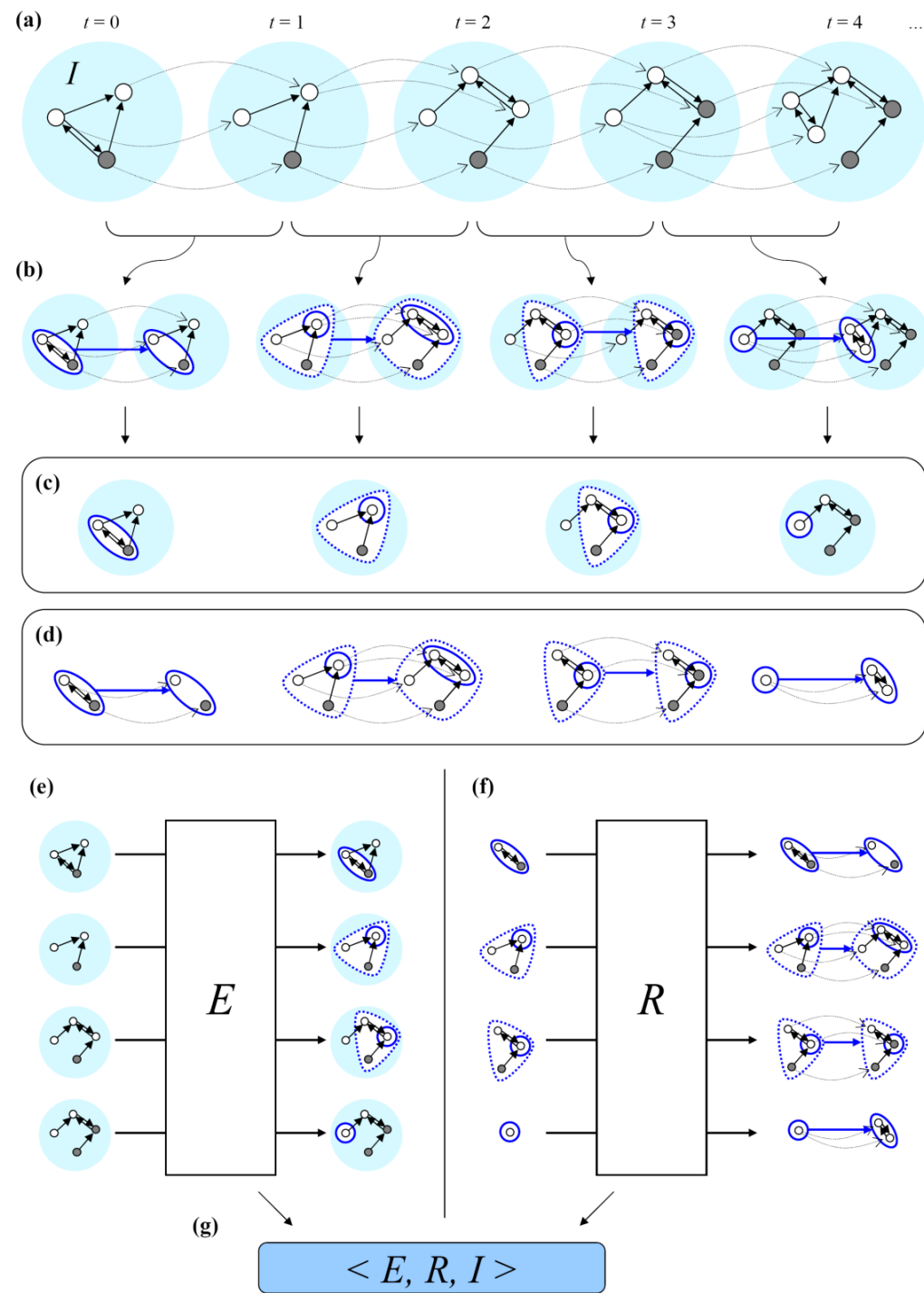


# Automation of model discovery from data: PyGNA

---

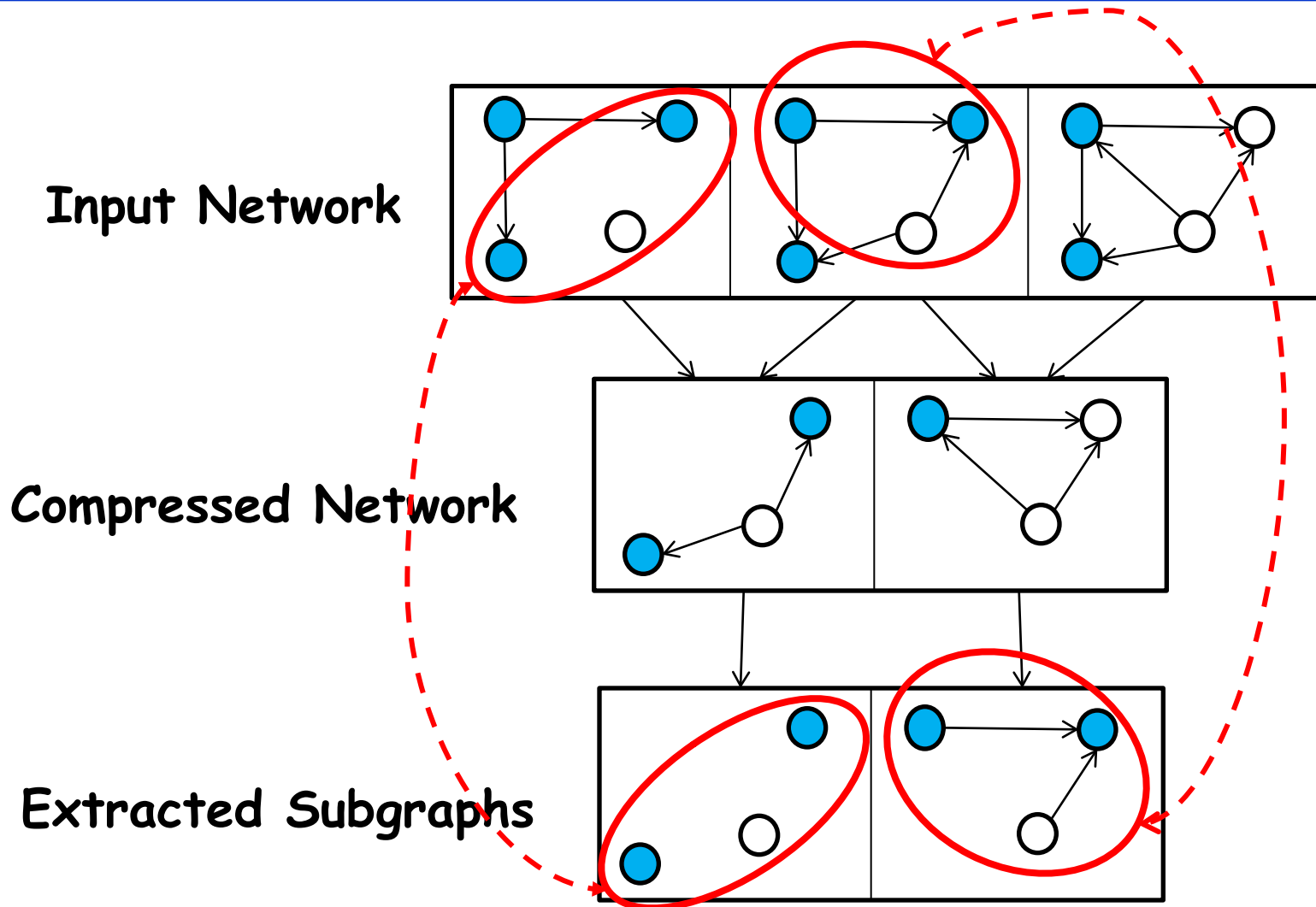
- Adaptive network rule discovery and simulation implemented in Python
  - <https://github.com/schmidtj/PyGNA>
- Input: Time series of network snapshots
- Output: A GNA model that best describes given data







# Extracted subgraphs

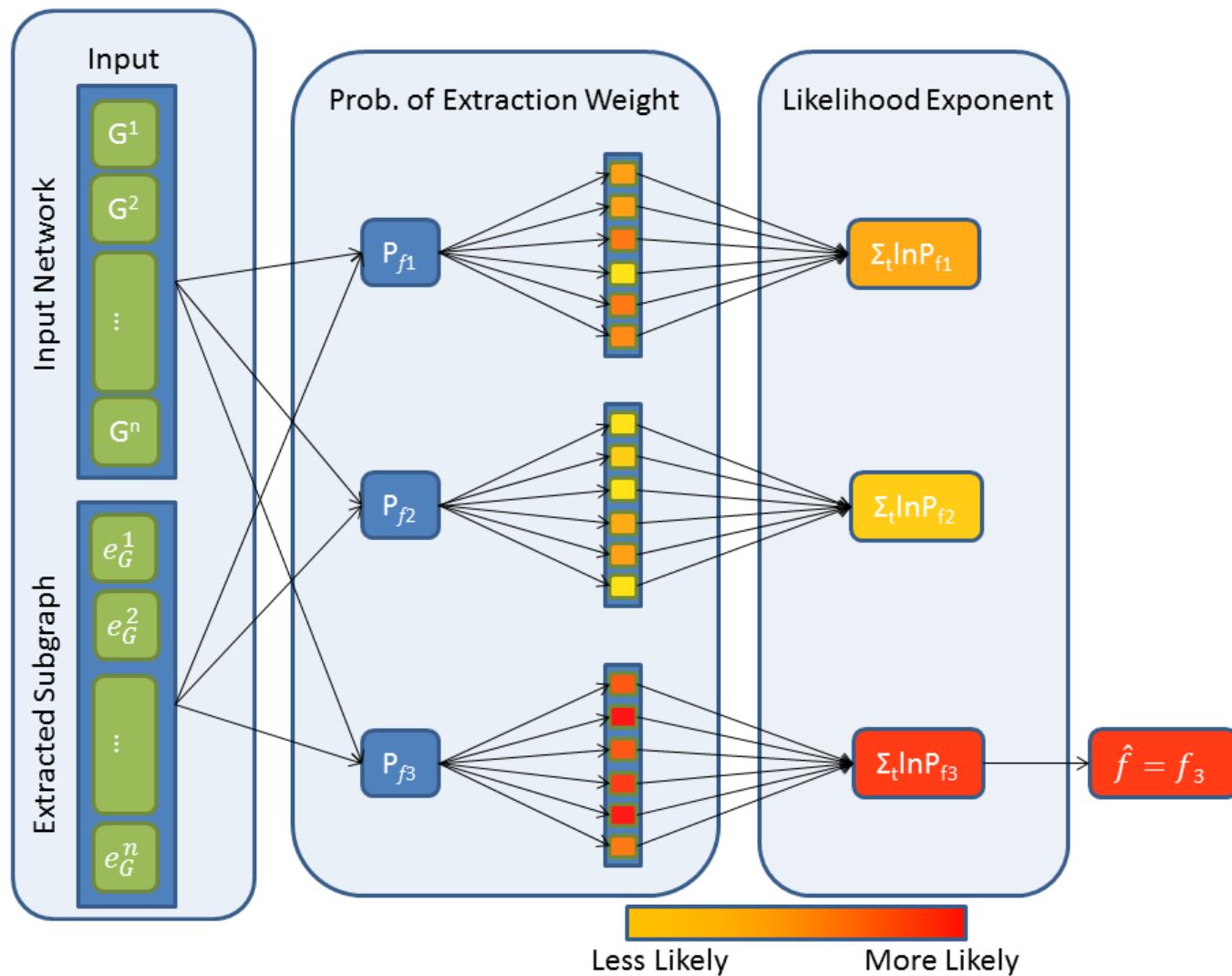


# Extraction mechanism

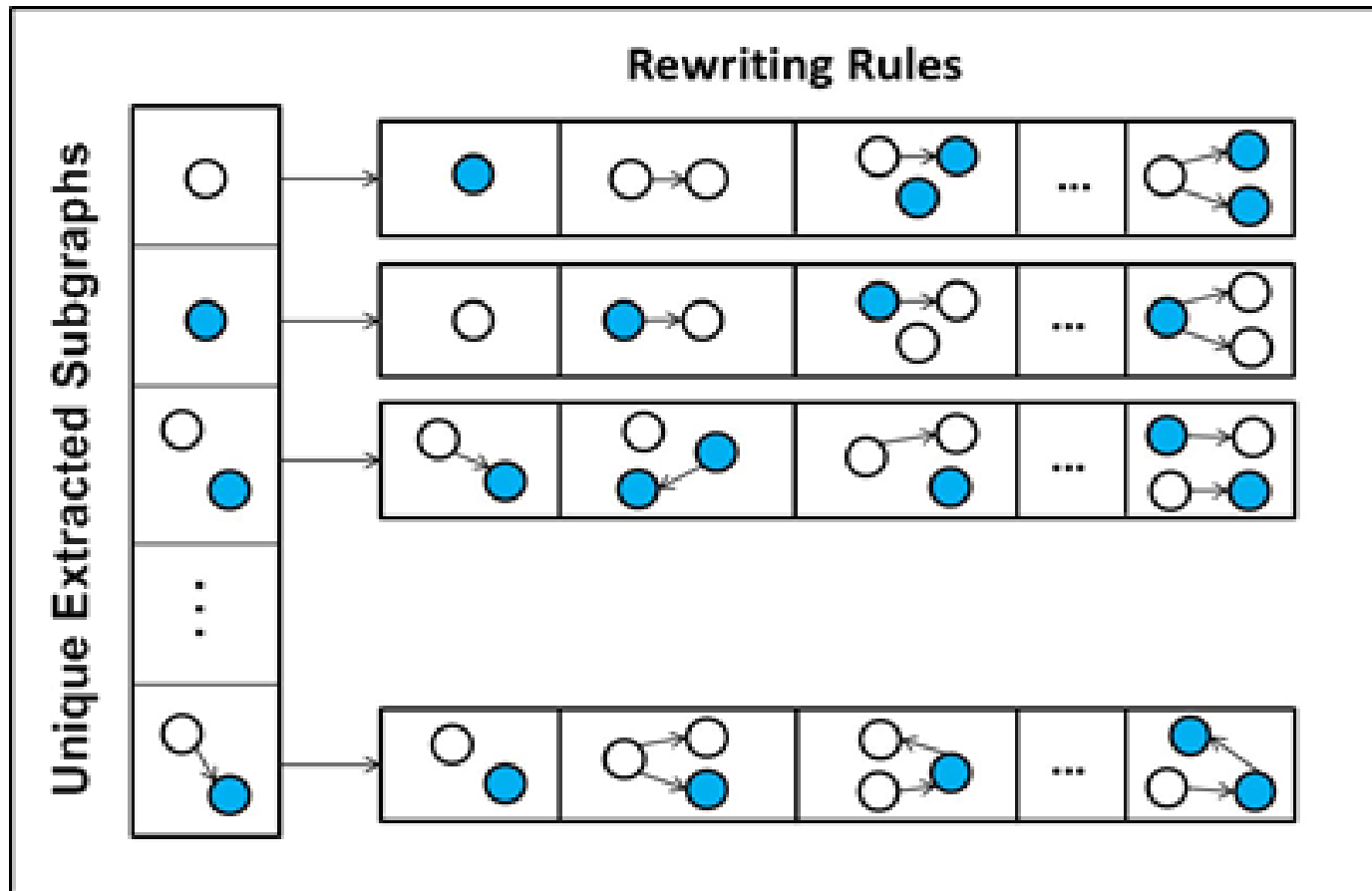
## identification: “Where, when”

- **Candidate models provided by user**
  - Degree-based preferential selection
  - State-based preferential selection
  - Degree & State-based etc...
- **Maximum likelihood method**
  - Computes likelihood using each hypothetical model & accumulates log likelihood over time
  - Chooses the model with maximum likelihood

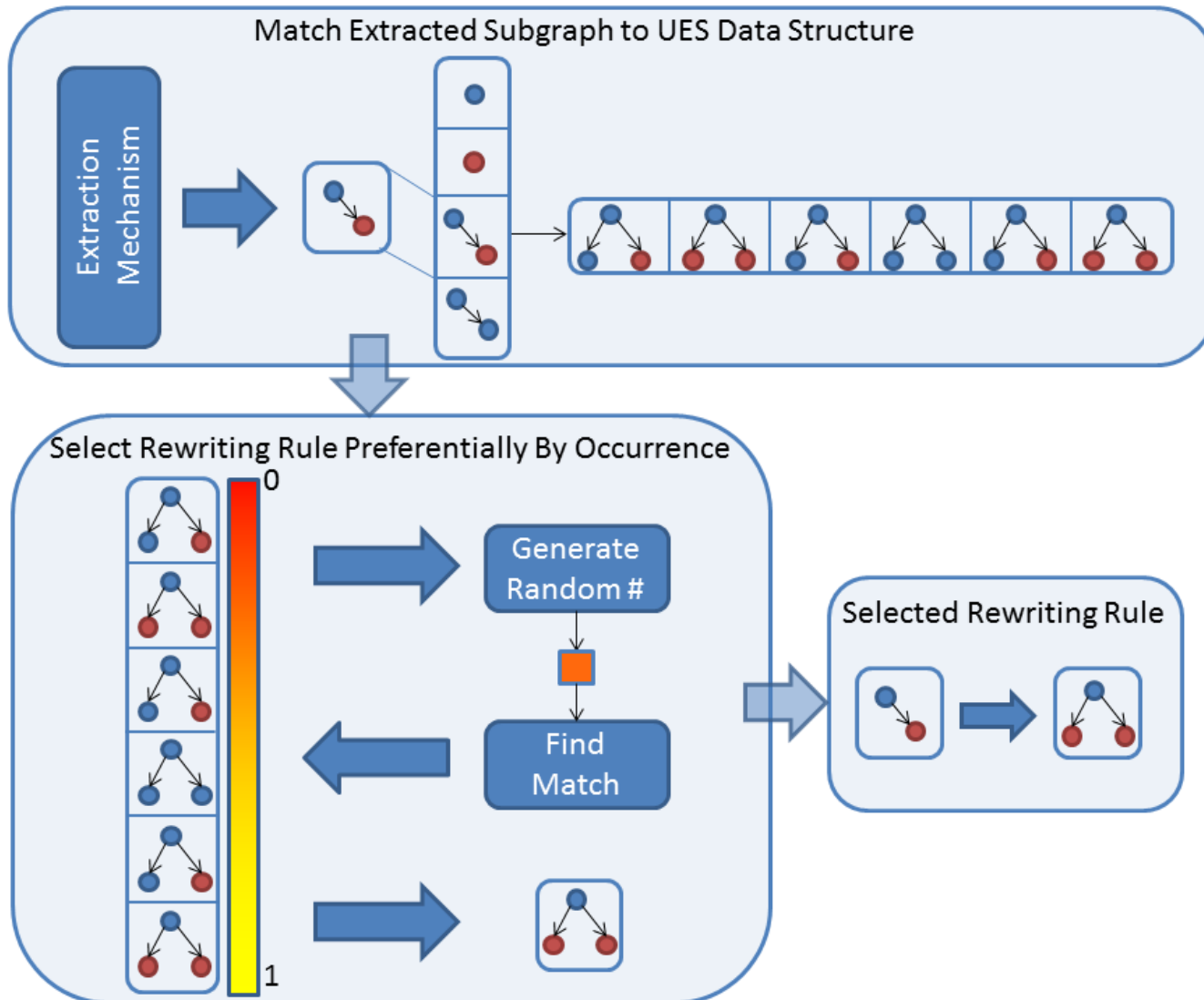
# Algorithm



# Replacement mechanism identification: "What"

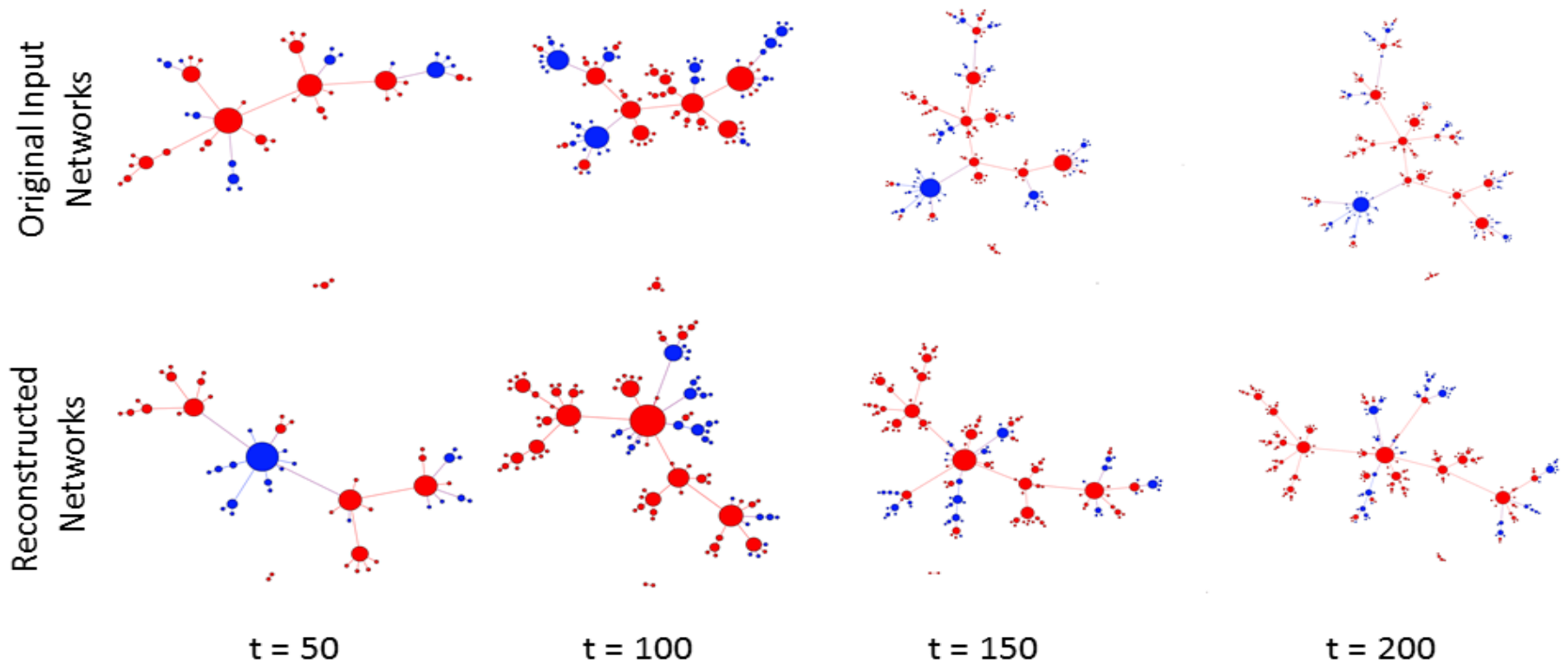


# Algorithm

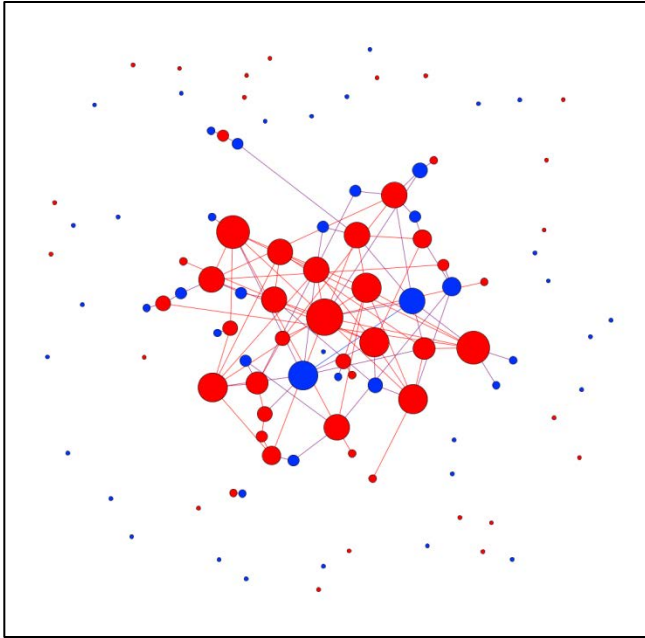


# Results

- Example: “Degree-state” networks

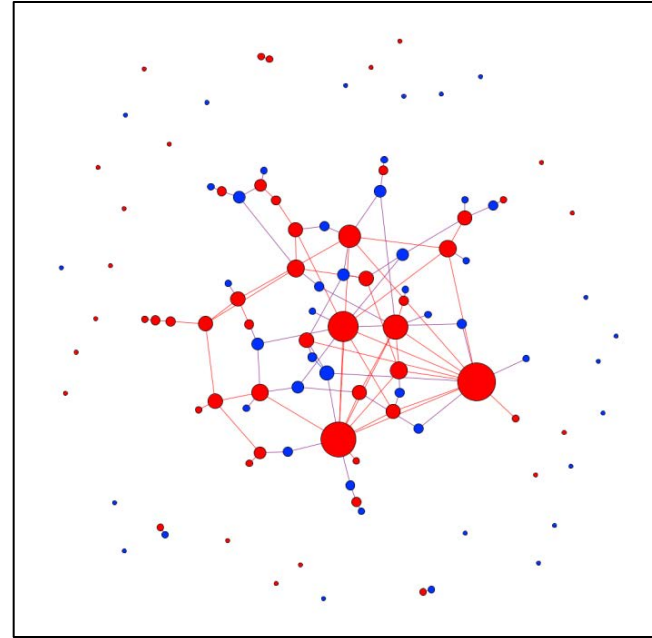
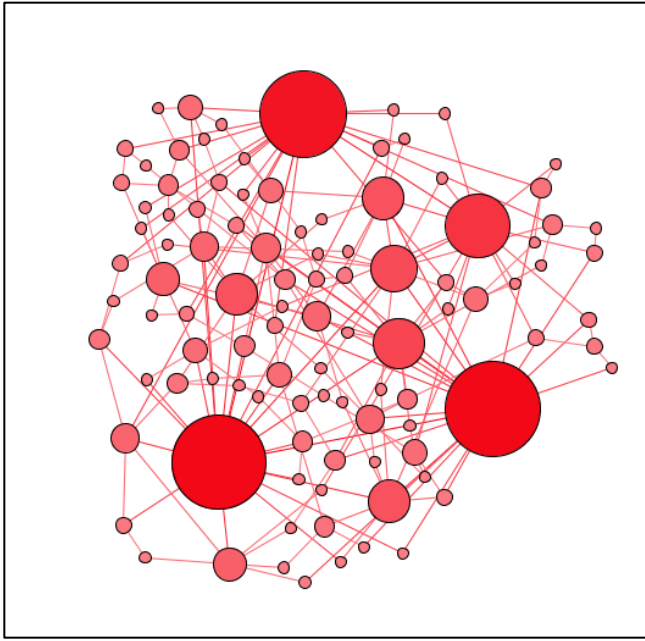


State-based

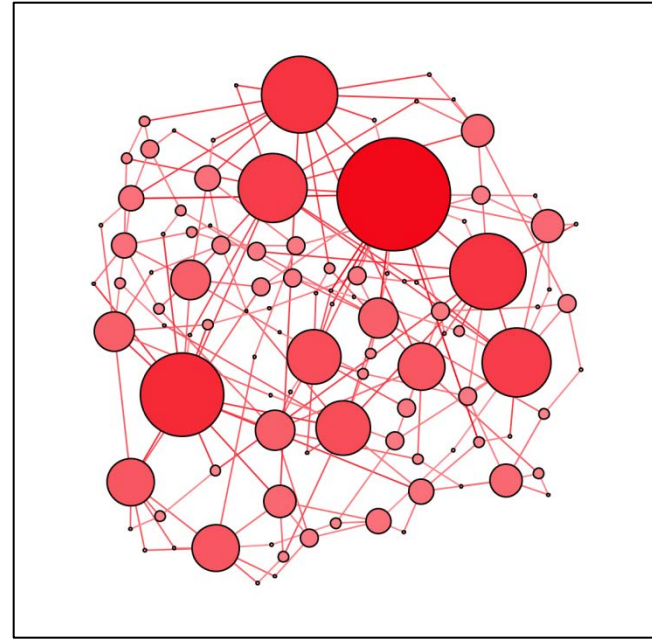


Input

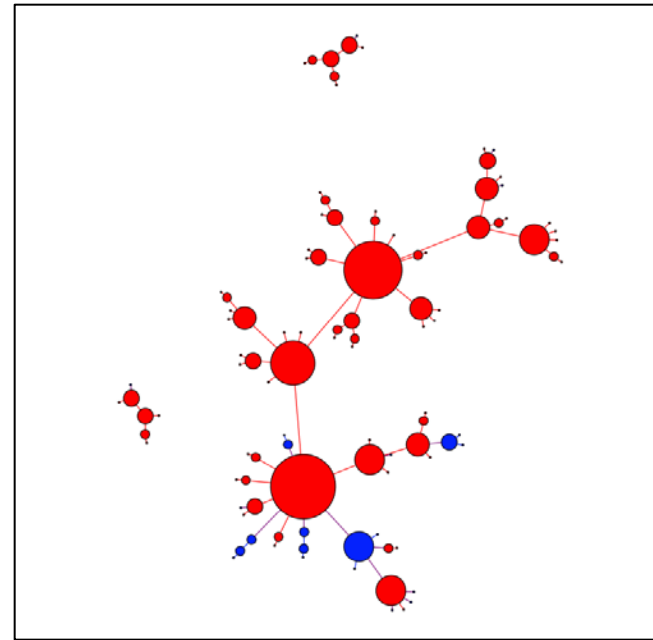
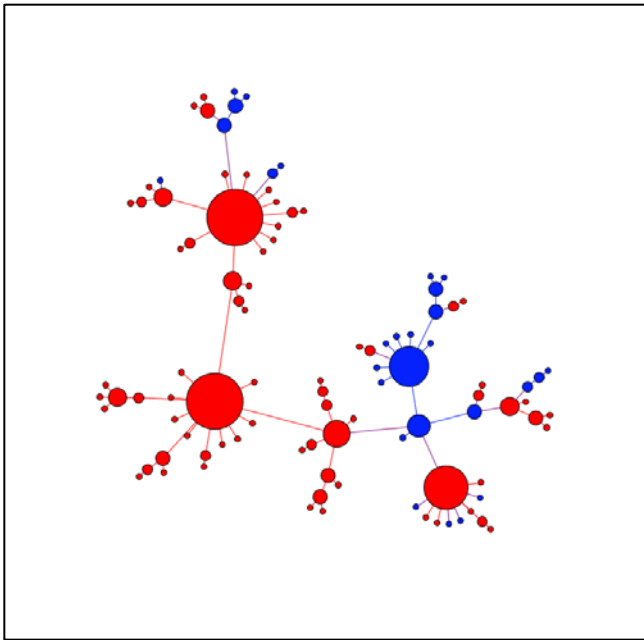
Barabási-Albert



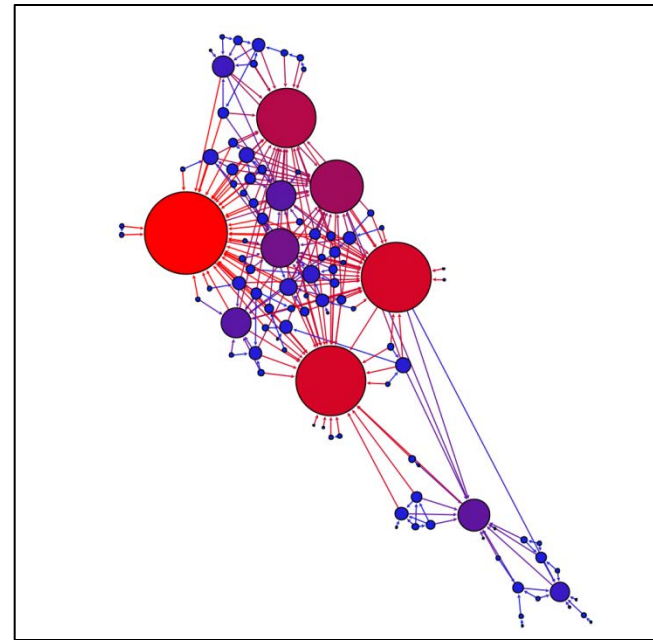
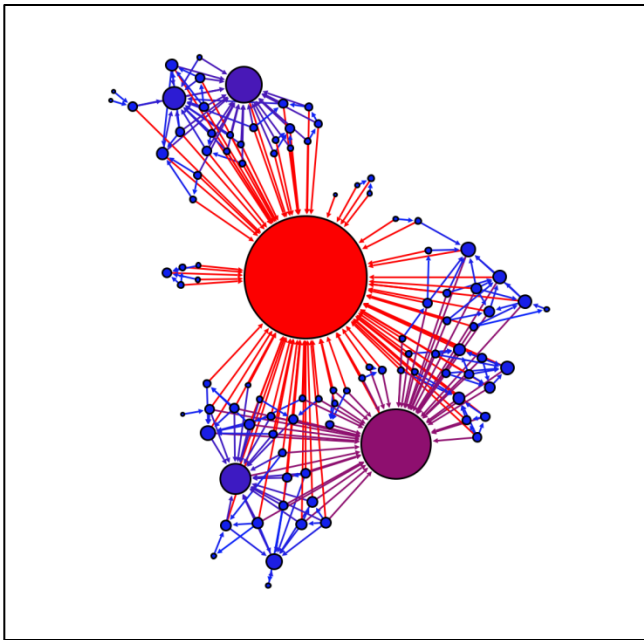
Simulated



Degree-State



Forest Fire

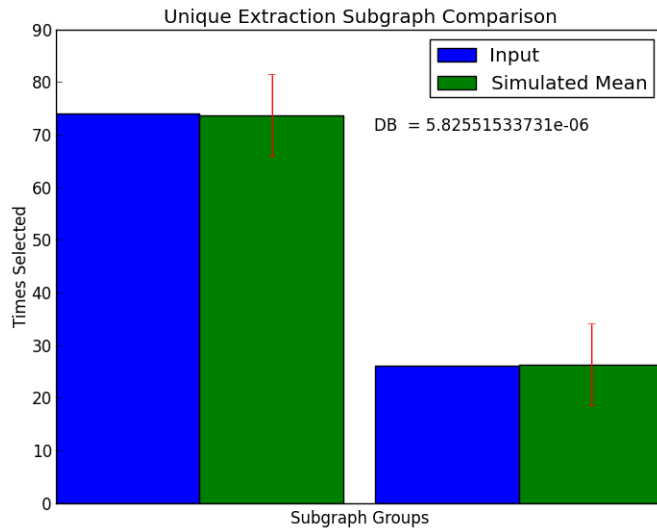


Input

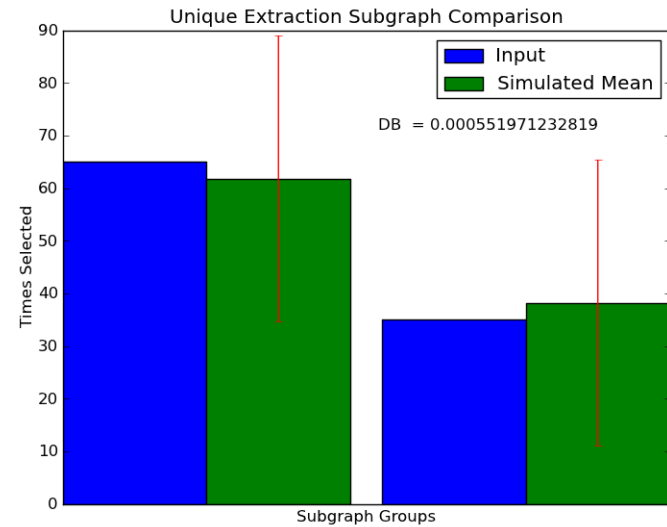
Simulated



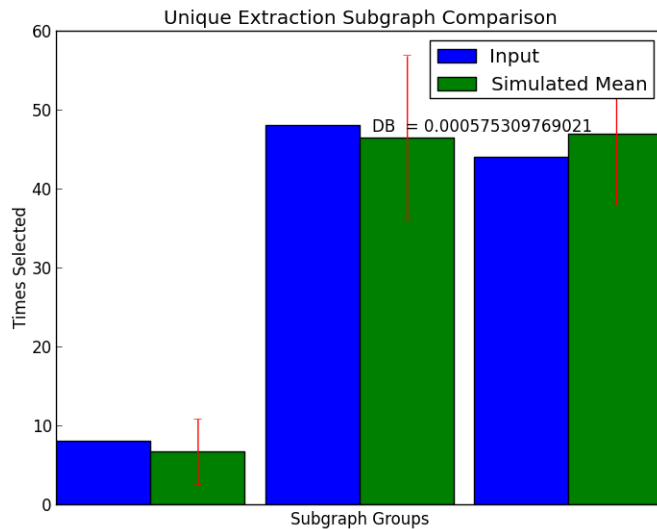
# Barabási-Albert



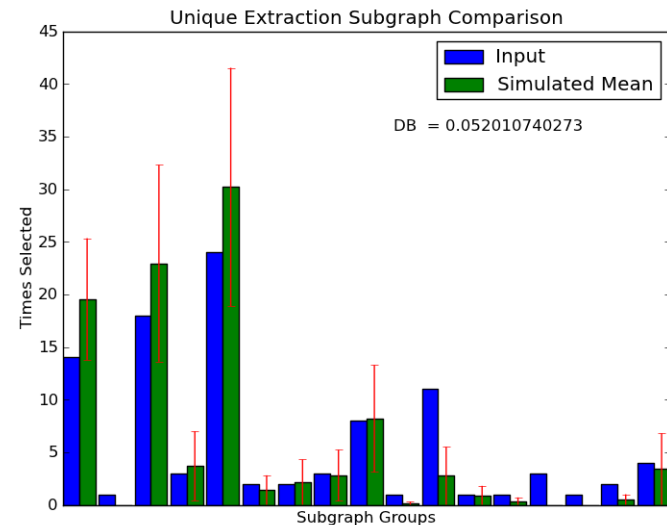
# Degree-state



# State-based



# Forest Fire



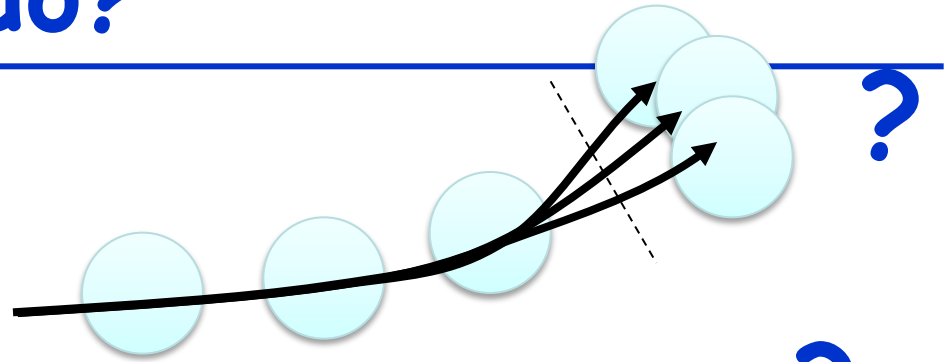
# Comparison with other methods

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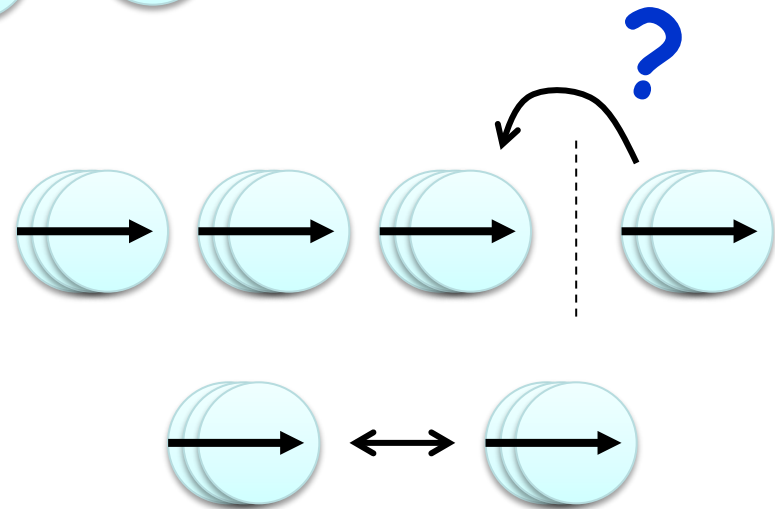
- PyGNA produces *generative models* using detailed state-topology information
  - Capable of generative simulation that is not available in statistical approaches (e.g., Rossi et al. 2013)
- PyGNA models extraction and replacement as *explicit functions*
  - More efficient and flexible than graph-grammars (e.g., Kurth et al. 2005)

# What can we do?

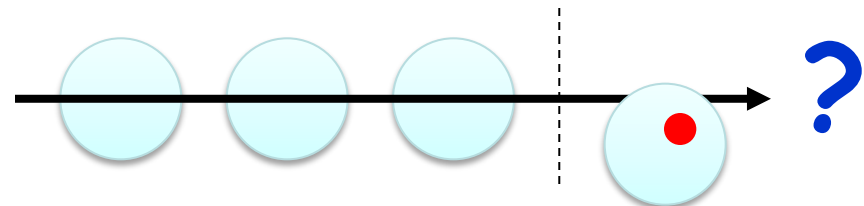
- Prediction



- Classification



- Anomaly detection



# Summary

---

- State-topology coevolution of adaptive networks is a promising, unexplored area
  - Theory-driven approaches
    - Dynamical modeling, exhaustive rule search
    - Applications to social sciences etc.
  - Data-driven approaches
    - Application to operational network modeling
    - Automatic rule discovery from data

<http://coco.binghamton.edu/NSF-CDI.html>

**Additional Topic:**  
**Analysis of Adaptive Networks**

# How to analyze adaptive network dynamics?

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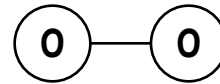
- Non-trivial coupling between network states and topologies are not easily handled in a simple analytical formula
- But such couplings could be partially incorporated in analysis by considering densities of node “pairs”

# Pair approximation

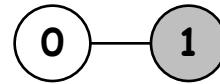
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- Considers densities of every pair of nodes with states & connectivity (in addition to individual state densities)

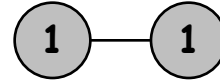
$\rho_{00c}$  = density of



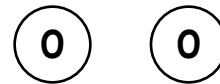
$\rho_{01c}$  = density of



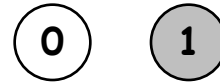
$\rho_{11c}$  = density of



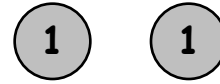
$\rho_{00n}$  = density of



$\rho_{01n}$  = density of



$\rho_{11n}$  = density of

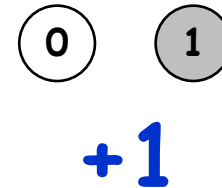
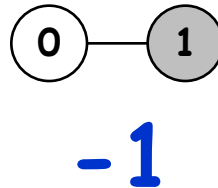
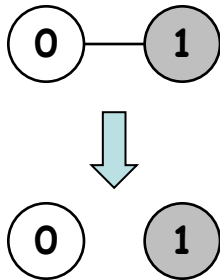


Describes  
how these  
densities  
change over  
time

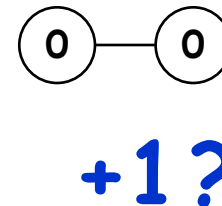
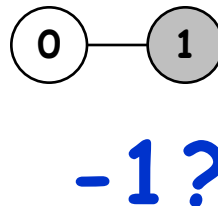
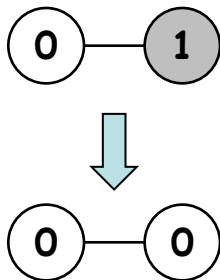
# Example: Adaptive voter model

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- Disconnect of a link:



- Change of an opinion:



(Any other densities affected too?)



# Exercise

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- Complete the number of changes in each pair density for the adaptive voter model on a random network
- Calculate how often each transition occurs
- Make a prediction using the pair-approximation-based model

# Exercise

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- Conduct pair approximation of the adaptive SIS model and study its dynamics

# FYI: Moment closure

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- Similar approximations are possible for densities of higher-order motifs
- Approximation techniques (including MFA, PA and higher-order ones) is called the “moment closure method”
  - Predicting the change of a “moment” ( $\rho_{00}$ ) would require higher-order “moments” ( $\rho_{000}$ ), but you “close” this open chain by assuming  $\rho_{000} = \rho_{00} \rho_{00} / \rho_0$ , etc.