CSCI5070 Advanced Topics in Social Computing

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Outline

- Scale-Free Networks
 - Generation
 - Properties
 - Analysis
- Dynamic Networks
 - Emergence
 - Epidemics
 - Synchrony
 - Influence Networks



SCALE-FREE NETWORKS



Scale-Free Networks

- A scale-free network is a network G with degree sequence distribution g' obeying a power law of the form $h(k) \sim k^{-q}$, where k is degree ($1 < k < \infty$) and q is an exponent (typically 2 < q < 3)
- Power-law distributions are not exponential distributions
 - The tail of an exponential distribution vanishes much faster

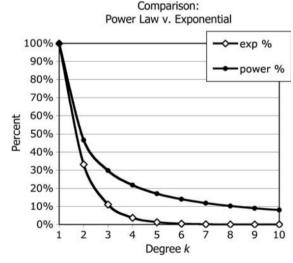


Figure Comparison of $y = 3 \exp(-qk)$, versus $y = 1/K^q$, for q = 1.1, shows the fat tail of the power law.



Generation

- The Barabasi–Albert (BA) Network
 - Sample from the degree sequence distribution of network G at timestep t
 - The probability of a high-degreed node obtaining a subsequent link continues to rise, and vice versa

Procedure

(1) *Growth*: Starting with a small number (m_0) of vertices, at every time step we add a new vertex with $m(\leq m_0)$ edges (that will be connected to the vertices already present in the system).

(2) *Preferential attachment*: When choosing the vertices to which the new vertex connects, we assume that the probability *P* that a new vertex will be connected to vertex *i* depends on the connectivity k_i of that vertex, such that $P(k_i) = \frac{k_i}{\sum_i k_i}$.

(3) After *t* time steps the model leads to a random network with $N = t + m_0$ vertices and *mt* edges."



BA Generative Procedure

- 1. Inputs: Δm = number of links to add to each new node; n = network size.
- 2. Initialize: Designate nodes by enumerating them as $0, 1, 2, \ldots, (n 1)$.
 - a. Given the ultimate number of nodes n > 3, initially construct a complete network with nNodes $= n_0 = 3$ nodes and nLinks = 3 links. The degree sequence of this complete network is g = [2,2,2], and the degree sequence distribution is g' = [1] because each node is connected to the other two.
- 3. While nNodes $\leq n$:
 - a. New node: Generate a new tail node v.
 - b. #New links: Set n_links = minimum (Δm , *n*). Cannot add more links than existing nodes.
 - c. Repeat n_links times:
 - i. *Preferential attachment*—select an existing head node u by sampling from the degree sequence cumulative distribution function CDF(i) defined by

$$CDF(i) = \sum_{j}^{i} \frac{\text{degree}(n_j)}{k_{\text{total}}}, \text{ where } k_{\text{total}} = 2n\text{Nodes}$$

ii. Let *r* be a uniform random number from [0,1). Then *u* is a random variate sampled from CDF(i) as follows:

$$CDF(u - 1) \le r \le CDF(u); \ u = r \text{ nNodes}$$

iii. Connect $v \sim u$, taking care to avoid duplicate links.



BA Network Entropy

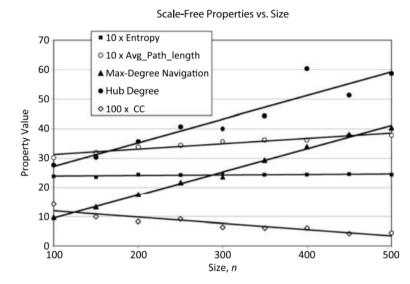


Figure Comparison of BA scale-free network properties versus size, n, for $\Delta m = 2$. Note the scale factors on entropy, average path length, and cluster coefficient.

Entropy remains constant as network size increases

$$I = \int_{\Delta m}^{\infty} h(k) \log_2(h(k)) \delta x = -1.443 \int_{\Delta m}^{\infty} h(k) \ln(h(k)) \delta x$$

Substitution of $h(x) = Ax^{-3}$, where $A = 2\Delta m (\Delta m + 1)$ into the integral,

$$I = -1.443A \int_{\Delta m}^{\infty} \left(\frac{\ln(A) - 3\ln(x)}{x^3}\right) \delta x = \Delta m + 1 \frac{1.1645 - \log_2(\frac{\Delta m + 1}{\Delta m^2})}{\Delta m}$$



Hub Degree versus Density

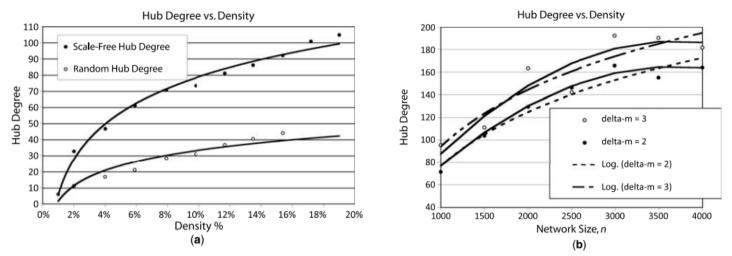


Figure Comparison of hub degree of networks for fixed and variable *n*: (a) scale-free and random networks versus density, for constant size, n = 200, (b) variable number of links *m*; and hub degree with variable density and size, *n*, but fixed Δm .

Degree(hub) ~
$$O(\log_2(\text{density})); 1\% \le \text{density} \le 20\%$$

Density(scale-free) =
$$2\frac{m}{n(n-1)} = 2\frac{\Delta m}{n}$$

- Extremely high hub degree is the predominant property of a scale-free network
- Hub degree grows logarithmically with density for both random and scale-free networks, but the rate of increase is much greater for a scale-free network



Average Path Length

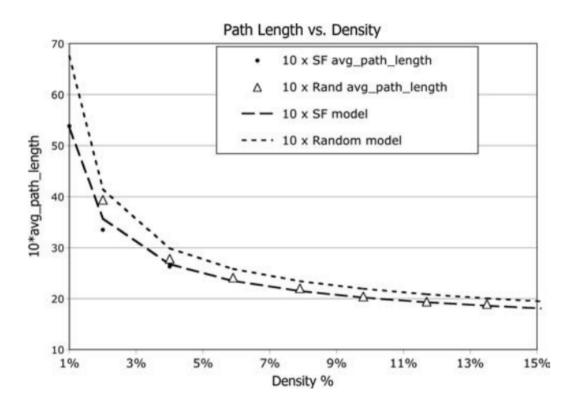


Figure Comparison of average path length of BA scale-free and ER random networks versus density, n = 200. The average path length of a scale-free network is slightly less then that of an equivalent random network.

avg_path_length(BA network) =
$$\frac{A \log(n)}{\log(n) + \log(C(\text{density}))}$$



Average Path Length

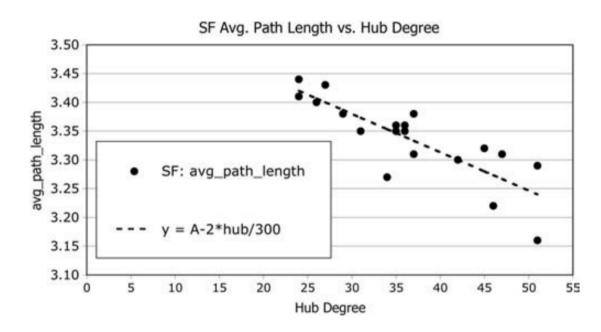


Figure Average path length declines linearly as hub degree increases in a scale-free network of size n = 200 and density 2% ($\Delta m = 2$). Least-squares curve fit to these data yields A = 3.58 and slope equal to $\frac{-2}{300}$.

avg_path_length(BA network) =
$$A - 2 \frac{\text{Hub_degree}}{300}$$



Closeness

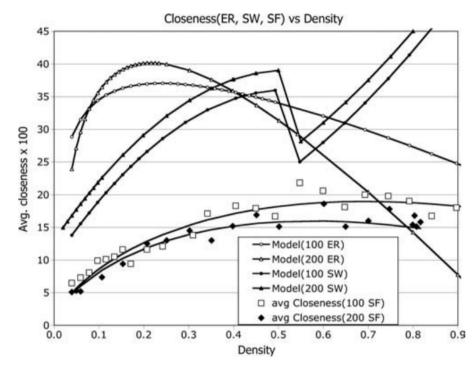


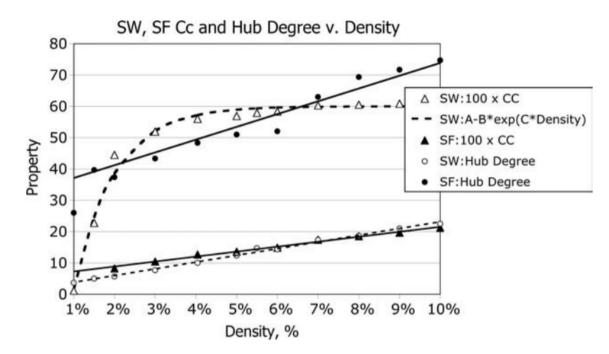
Figure Average closeness versus density for random, small-world, and scale-free networks shows that scale-free networks have weaker intermediaries on average.

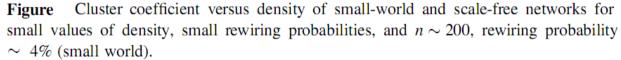
TABLE	Number of Paths	per Node Through Closest Node
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Network	Number of Paths <i>n</i>
Random Small world	0-10 0-20
Scale-free	0-38



Cluster Coefficient





- Scale-Free CC. $100(CC(\text{scale-free})) \sim O(\text{density}), \quad 1\% \leq \text{density} \leq 10\%.$ Alternatively, $100(CC(\text{scale-free})) \sim O(\Delta m)$ because density $= 2(\Delta m/n); n \gg 1.$
- Small-World CC. $100(CC(\text{small-world})) = A B \exp(C \text{ density}); 1\% \le \text{density} \le 10\%$, where A = 60; B = 158.5; C = -100, for n = 200.



Comparison

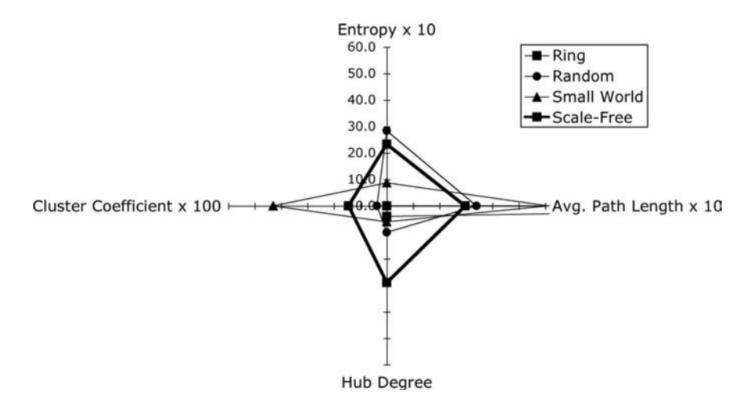


Figure Kiviat graph comparing ring, random, WS small-world, and BA scale-free network properties.



DYNAMIC NETWORKS



Definitions

- A dynamic network, G(t) = [N(t), L(t), f(t):R], is a timevarying 3-tuple consisting of a set of nodes N(t), a set of links L(t), and a mapping function $f(t): L(t) \rightarrow N(t) \times N(t)$, which maps links into pairs of nodes
- If G reaches a final state, $G(t_f)$, and remains there for all $t \ge t_f$, G is said to be convergent
- $G(t_f)$ is a fixed point
- If G(s) converges for all initial states s, G is strongly convergent



Emergence

- In a static network, the properties of nodes, links, and mapping unction remain unchanged over time
- In a dynamic network, the number of nodes and links, the shape of the mapping function, and other properties of the graph change over time
- Time-varying changes leading to structural reorganization in a network—is called emergence
- An emergent network is formed by starting at some predefined initial state and then transitioning through a series of small changes into an end state



Open-loop & Feedback-loop

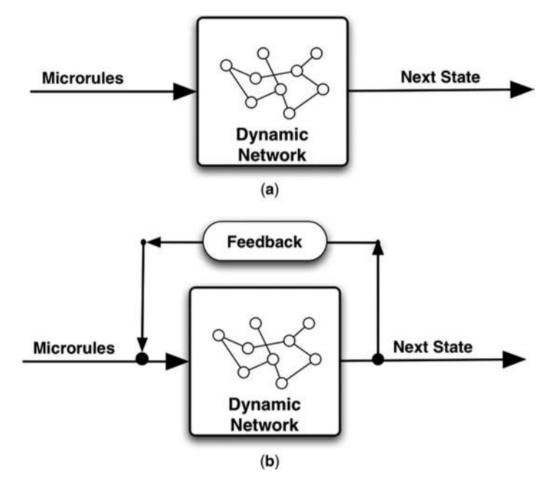


Figure Emergence in a dynamic network: (a) open-loop emergence, typical of intrinsic or genetic evolution; (b) feedback-loop emergence, typical of external or environmental forces shaping a network.



Hub Emergence

- Initially, G(0) is a random network with n nodes and m links
- At each time-step, select a node and link at random
 - The randomly selected node is selected if its degree is higher than that of the randomly selected link's head node
 - Rewire the link

```
public void NW_doIncreaseDegree(){
//Rewire network to increase node degrees
int random_node = (int) (nNodes * Math.random()); //A random node
int random_link = (int) (nLinks * Math.random()); //A random link
int to_node = Link[ random_link] .head; //Link' s to_node
int from_node = Link[ random_link] .tail; //Anchor node
int from_node = Link[ random_link] .tail; //Anchor node
if (node[ random_node] .degree > node[ to_node] .degree){
    if (NW_doAddLink(node[ from_node] .name, node[ random_node] .name))
        NW_doCutLink(random_link); //Replace random link
    else Message = "Warning: Duplicate Links Not Allowed.";
    }
}//NW_doIncreaseDegree
```



Hub Emergence

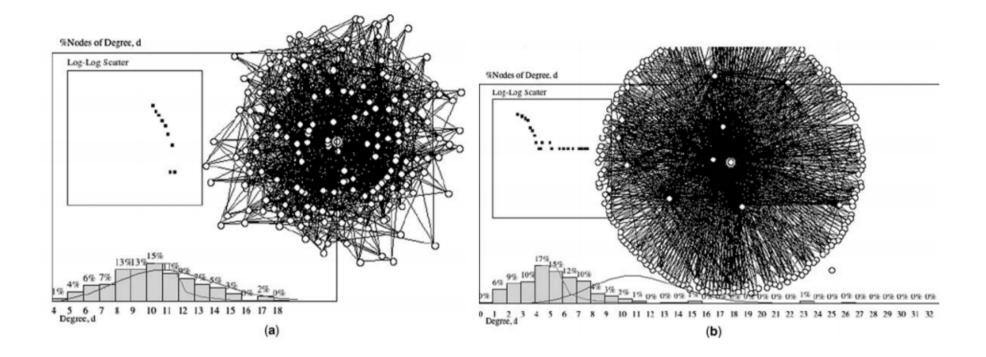


Figure Emergence of a network with high-degreed hubs from a random network: (a) random network, G(0)—the degree sequence distribution before emergence; (b) hub emerged network, G(160,000)—the degree sequence distribution after emergence. The initial network, G(0), n = 200, m = 1000, was generated by the Erdos–Renye procedure.

Cluster Emergence

- Select a random link and random node
- Rewire the link to point to the new (random) node, if the overall cluster coefficient remains the same or is increased
- If the cluster coefficient decreases as a result of rewiring, revert to the topology of the previous time-step
- Repeat this microrule indefinitely, or until stopped



Cluster Emergence

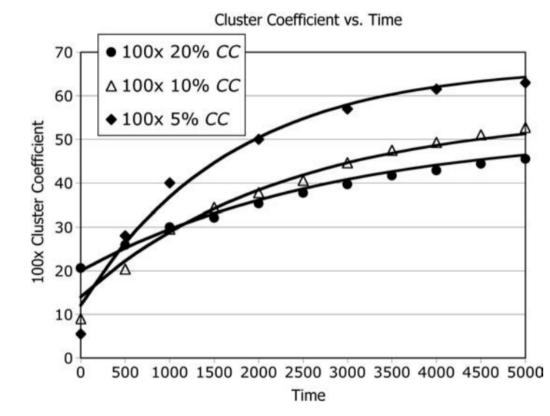


Figure Cluster coefficient versus time for n = 100, m = 249 (5%), m = 500 (10%), and m = 1000 (20%).



Epidemics

- A network epidemic is a process of widespread and rapid propagation of a contagion through a network
 - Typically, the contagion is a condition of network nodes (working, failing, dormant, active, etc.) brought about by adjacent nodes through propagation along one or more links
 - Infection rate: the probability that propagation of the contagion at node v successfully infects adjacent nodes



Kermack–McKendrick Model

- Assumes homogeneous and a very large population
 - Everyone has an equal chance of contracting the disease from anyone else
 - *n* is unbounded
- Each node is classified to one of four states
 - Susceptible: it can possibly become infected
 - Infected: it has contracted the contagion
 - Recovered: it has recovered from the infection and is now immune to future infection
 - *Removed*: it has died from the effects of the infection



Kermack–McKendrick Equations

We define γ as the rate of infection, Δ as the rate of removal or recovery, and t_r as the duration of a node's infected state before transition to a removed or recovered state.

Let S(t) be the number of susceptible actors, I(t) the number of infected actors, and R(t) the number of actors removed from the population as a result of death or immunity after recovering from the illness, at time *t*. In a finite population, n = S(t) + I(t) + R(t). The Kermack–McKendrick equations relate *S*, *I*, and *R* to one another through their time rate of change and initial conditions, as follows:

Infection % vs. Time
$$\frac{\delta S(t)}{\delta t} = -\gamma S(t)I(t); \ S(0) = S_0 \tag{I}$$

100% 80%

> 60% 40% 20% 0%

1 2 3

0

5 6

4 5 Time

Infection i(t)

$$\frac{\delta I(t)}{\delta t} = \gamma S(t)I(t) - \Delta I(t); \ I(0) = I_0$$
(II)

$$\frac{\delta R(t)}{\delta t} = \Delta I(t); \ R(0) = R_0$$
(III)

$$S(t) + I(t) + R(t) = n \tag{IV}$$



Other Models

- Susceptible–Infected–Removed (SIR) Model
- Susceptible—Infected—Susceptible (SIS) Model

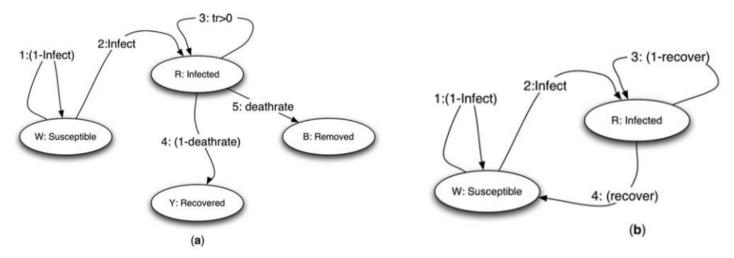


Figure Models of epidemics: (a) SIR; (b) SIS (color code : W—susceptible = white; R—infected = red; B—removed = black; Y—recovered = yellow).



Synchrony

- A network is said to synchronize if the values of all of its nodes converge to some constant as the time rate of change of all of its node values approaches zero
- A dynamic network is
 - stable if the value of its nodes synchronize
 - *transient* if its node values oscillate
 - bistable if its nodes oscillate between fixed values
 - *unstable* or *chaotic*, otherwise



Chaotic Maps

- A dynamic network is said to sync if, starting at some initial state G(0), it evolves in finite time to another state, G(t*), and stays there, forever.
- G(t*) is called a strange attractor
- Networks that appear to bounce around from one state to another in no apparent pattern are considered *chaotic*
- Networks that oscillate between two or more strange attractors are called oscillators

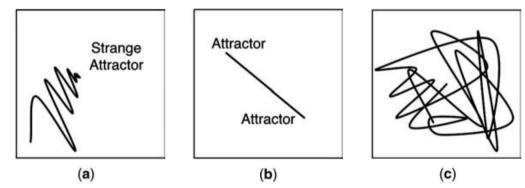


Figure Chaotic maps: trajectories of (a) a stable node as it reaches its strange attractor, (b) a bistable node as it oscillates between two attractors, and (c) a chaotic node as it wanders around in state space.

Network Stability

- Stability describes nodes and networks that recover from disruptions in their state
- A stable network will recover—perhaps very slowly while an unstable network will not
- Two general techniques used for analyzing the stability of a dynamic system
 - The Laplace or Z-transform (Lyapunov) method (Lago-Fernandez, 1999)
 - The Laplacian eigenvalue (spectral decomposition) method (Wang, 2002)



Network	Bistable?	Comments
Barbell	No	Bistable oscillator
Line	No	50%-50% oscillations
2-Regular	Yes	Odd-sized cycles
Complete	Yes	Odd-sized cycles
Toroid	Yes	Odd-sized cycles
Binary tree	No	No cycles
Hypercube	No	Even-sized cycles
Random	Nearly always	Likely has odd-sized cycle
Small-world	Yes	Has clusters
Scale-free	Nearly always	Likely has odd-sized cycle
Star	No	No cycles
Ring	N = odd number	Odd-sized cycle

TABLE Sync Results for Regular and Nonregular Networks



Influence Networks

- An influence network is a (directed or undirected) network, G = { N, L, f } containing nodes N; directed, weighted links L; and mapping function f : N × N that defines the topology
- Nodes are called actors
- Links are called influences
- Two major types
 - Undirected networks such as the buzz network (Rosen, 2000)
 - directed networks such as the social networks formed by negotiating parties in a human group



A Buzz Network

- Let S(t) be the state vector representing an actor's position (a product, idea, political belief, etc.)
 - S(t) = -1 if the position is negative
 - S(t) = 0 if neutral
 - S(t) = +1 if positive $-1 \le S(t) \le 1$ $S(t) = [s_1, s_2, s_3, \dots, s_n]^T = \text{state vector of } G; S_j(t) = \operatorname{row} j \text{ of } S(t)$

Buzz State Equation

$$S(t+1) = \phi S(t) + (1-\phi) \sum_{j \sim i} \frac{S_j(t)}{\text{degree}(i)}$$

where the summation $\sum_{i \sim i}$ is over the neighbors of node *i*, specifically, $j \sim i$ and $s_i = [-1, 1], i = 1, 2, ..., n$. $0 \le \phi \le 1$ is the *stubbornness factor*.



Properties

- Each seed actor spreads his or her product endorsement to adjacent neighbors, which in turn is spread to their neighbors, and so forth, much like the spread of an infection
- The state of each actor depends on the strength of the individual's convictions and ϕ_{1} e positions of adjacent neighbors
- Stability
 - Buzz networks reach a consensus that is influenced by the dominant (hub) node
 - Consensus: all nodes reach the same final state



Two-Party Negotiation

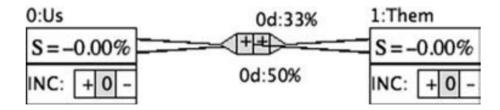


Figure I-net representation of a two-party negotiation: actor Us versus actor Them, and influences of $\frac{1}{3}$ and $\frac{1}{2}$, respectively.

- Two-party negotiation is a process of interactive compromise
 - The two actors must narrow the difference in positions as perceived by each other if the network is to reach a consensus

For example, after one timestep:

Us(1) = Us(0) + 50%(Them(0) - Us(0)) = 1 + 50%(-2) = 0

Them(1) = Them(0) + 33%(Us(0) - Them(0)) = -1 + 33%(2) = -33%

Two-Party Negotiation

$$s_i(t+1) = s_i(t) + \sum_j [s_j(t) - s_i(t)]\phi^T i, j;$$

Us(2) = Us(1) + 50%(Them(1) - Us(1)) = 0 + 50%(-0.33) = -0.167Them(2) = Them(1) + 33\%(Us(1) - Them(1)) = -0.33 + 33\%(0.33) = -22\% Us(3) = Us(2) + 50%(Them(2) - Us(2)) = 0.167 + 50%(-0.056) = -19%Them(3) = Them(2) + 33\%(Us(2) - Them(2)) = -22 + 33\%(0.056) = -20\%

- The two actor's states converge in the end
- Differences drop to zero and both actors reach a consensus state equal to negative 20%, which favors the initial position of Them
- Therefore, I:Them is more influential than actor 0:Us.



References

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