Network Dynamics: an Overview

Giacomo Como Department of Automatic Control, LTH www.control.lth.se/Staff/GiacomoComo/ giacomo.como@control.lt<u>h.se</u>

October 1, 2013

Network Dynamics, Ph.D. course, 9 ECTS

- ▶ 18 two-hours meetings (including today)
- \blacktriangleright 1 lecture + 1 exercise session per week
- ▶ (bi-)weekly homeworks + small final presentation
- reference books:
- 0 notes and specific research papers
- 1 Easley & Kleinberg, 'Networks, crowds, and markets', 2010
- 2 Newman, 'Networks', Oxford U.P., 2010
- 3 Levine, Peres & Wilmer, 'Markov chains and mixing', 2008
- 4 Draief & Massoulie, 'Epidemics and rumors in complex networks'
- 5 Aldous& Fill, 'Reversible Markov chains and random walks'
- today: broad intro, course overview

Similar courses around the world

- MIT grad: http://stellar.mit.edu/S/course/6/sp11/6.986/
- MIT undergrad: http://stellar.mit.edu/S/course/6/fa09/6.207J/
- Cornell undergrad (by D.Easley, J.Kleinberg, & E.Tardos):: http://www.infosci.cornell.edu/courses/info2040/2011fa/
- Berkeley grad 1 (by D.Aldous): http://www.stat.berkeley.edu/ aldous/260-FMIE/index.html
- Berkeley grad 2 (by E.Mossel): http://www.stat.berkeley.edu/ mossel/teach/SocialChoiceNetworks10/index.html
- North-Easter U. (by A.Barabasi): http://barabasilab.neu.edu/courses/phys5116/
- Michigan U. (by L.Adamic): http://open.umich.edu/education/si/si508/fall2008/materials

(Complex) networks

(Large-scale) systems of (simple) interacting units

- infrastructure networks: transportation, power, gas, and water distribution, sewer, Internet
- informational networks: WWW, citation networks
- ▶ social networks: friendships, family ties, Facebook etc.
- economic networks: supply chains
- financial networks: borrowing-lending nets
- biological networks: neural networks, gene/protein interactions
- ecological networks: food webs, flocks, ...

Studying (complex) networks

network structure + interaction mechanism

 \Downarrow

emerging behavior

- spread of epidemics and information
- design of distributed algorithms
- opinion formation, social influence, and learning
- network robustness,
- cascaded failures, systemic risk

Mathematical representation of network structure



 $\begin{array}{l} (\text{un}) \text{directed (weighted) graph } \mathcal{G} = (\mathcal{V}, \overline{\mathcal{E}}) \\ \mathcal{V} = \text{set of vertices (or nodes)} \qquad n = |\mathcal{V}| < +\infty \\ \mathcal{E} \subseteq \mathcal{V} \times \mathcal{V} = \text{ set of edges (or links)} \end{array}$

Examples 1



- Internet: nodes=routers, edges=direct physical links (und.)
- traffic networks: nodes=junctions, links=roads (directed)
- ▶ actors collaboration: nodes=actors, link⇔ same movie (und.)
- scientific collab.: nodes=researchers, link=coauthors (und.)

Example 2: political blogs before 2004 US elections



from Adamic and Glance, 'The Political Blogosphere and the 2004 U.S. Election: Divided They Blog', 2005

Example 3: Family ties in 15th century Florence



from Padgett and Ansell, 'Robust action and the rise of the Medici, 1400-1434', 1993

Example 4: High school friendships



from Moody, 'Race, school integration, and friendship segregation in America', $2002\,$

Example 5: Sexual contacts



from Newman, 'The structure and function of complex systems', 2003

Example 6: protein network in yeast nucleus



from Maslov and Sneppen 'Specicity and stability in topology of protein networks', 2002

Example 7: Freshwater food web



from Martinez, 'Artifacts or attributes? Effects of resolution on the Little Rock Lake food web', 1991

Random networks

network structure + interaction mechanism ↓ emerging behavior

structure of large-scale networks is difficult to describe exactly: huge or non directly accessible data

Random networks

$\begin{array}{l} \textbf{network structure} + \textbf{interaction mechanism} \end{array}$

↓ emerging behavior

structure of large-scale networks is difficult to describe exactly: huge or non directly accessible data

aggregate properties:

- connectivity, diameter / average distance
- frequency of subgraphs
- degree distribution

statistical approach:

- ensemble of graphs
- typical properties as $n = |\mathcal{V}| \to +\infty$

Properties widely observed in empirical studies

- 1. small world \leftrightarrow diameter $\approx \log n$
- 2. high clustering \leftrightarrow many triangles
- 3. scale free \leftrightarrow power law degree distribution

Small world

- Milgram's experiment ('67): randomly selected group of few hundreds of people from Omaha (NE). A letter given to each of them to be delivered to a stock broker living in Boston (MA). Letter can only be handed to a person know directly.
 35% letters reach destination, median # of steps: 5.5
 "6 degrees of separation"
- ► Albert, Jeong, and Barabasi ('99): WWW network, n ~ 800M average distance of webpages ~ 0.35 + 2.06 log n = 18.59

Power laws



 $p_k \sim C k^{-\gamma}$

show up in quite different contexts:

- percentage of words in a book
- percentage of cities of a given size
- percentage of people having a certain income

Power law \Longrightarrow heavy tails: lots of large cities, lots of rich people

Typically explained by rich-gets-richer mechanisms

Power law networks

$$d_v := \#\{ ext{neighbors of } v\}$$
 $p_d := rac{1}{n} \#\{v : d_v = d\}$
 $p_d \sim Cd^{-\gamma}$

Empirical studies:

- Barabasi and Albert ('99): WWW has $\gamma_{in} \sim 2.1$, $\gamma_{out} \sim 2.7$
- Faloutos ('99): Internet $\gamma \sim 2.16$
- actor collaborations: $\gamma \sim 2.3$
- ▶ Redner ('98): citation network: $\gamma_{in} \sim 2.6$, $p_d^{out} \sim C \exp(-Kd)$
- ▶ Liljeros ('01): # sexual partners per year (in Sweden) $\gamma_{male} \sim 3.3$, $\gamma_{female} \sim 3.5$

$$\begin{array}{rcl} 2 < \gamma \leq 3 & \Longrightarrow & \langle d \rangle < +\infty & \langle d^2 \rangle = +\infty \\ \gamma \geq 3 & \Longrightarrow & \langle d \rangle < +\infty & \langle d^2 \rangle < +\infty \end{array}$$

power law \leftrightarrow scale free

 $\overline{\mathcal{G}}(n,p) = (\mathcal{V},\mathcal{E})$ $|\mathcal{V}| = n$ $\mathbb{P}(\{v,w\} \in \mathcal{E}) = p$ mutually independent

$$\mathcal{G}(n,p)=(\mathcal{V},\mathcal{E})$$
 $|\mathcal{V}|=n$
 $\mathbb{P}\left(\{v,w\}\in\mathcal{E}
ight)=p$ mutually independent



n = 100p = 0.15

$$\mathcal{G}(n,p)=(\mathcal{V},\mathcal{E})$$
 $|\mathcal{V}|=n$
 $\mathbb{P}\left(\{v,w\}\in\mathcal{E}
ight)=p$ mutually independent



n = 100p = 0.2

$$\mathcal{G}(n,p)=(\mathcal{V},\mathcal{E}) \qquad |\mathcal{V}|=n$$
 $\mathbb{P}\left(\{m{v},m{w}\}\in\mathcal{E}
ight)=p \qquad ext{mutually independent}$

If $p = \lambda/n$, with high probability as $n \to \infty$,

$$\begin{array}{c} \mbox{phase} \\ \mbox{transition} \end{array} \left\{ \begin{array}{c} \lambda < 1 \Longrightarrow \mbox{size}(\mbox{largest component}) \asymp \log n \\ \lambda > 1 \Longrightarrow \begin{array}{c} \mbox{size}(\mbox{largest component}) \asymp n \\ \mbox{diam}(\mbox{giant component}) \asymp \log n \end{array} \right\} \begin{array}{c} \mbox{small} \\ \mbox{world} \end{array}$$

► Poisson degree distribution:
$$p_d \sim \frac{\lambda^d}{e^\lambda d!} \Rightarrow \text{NO power law}$$

▶ limited #(triangles) \Rightarrow NO clustering

Random graphs 2: preferential attachment

Barabasi-Albert ('99)

- 1. start from a small given graph n_0
- 2. add a vertex and connect it with *d* older vertices randomly chosen with conditional probability \propto their current degree
- 3. repeat step 2 $n n_0$ times



http://ccl.northwestern.edu/netlogo/models/run.cgi?PreferentialAttachm

Random graphs 2: preferential attachment

Barabasi-Albert ('99)

- 1. start from a small given graph n_0
- 2. add a vertex and connect it with *d* older vertices randomly chosen with conditional probability \propto their current degree
- 3. repeat step 2 $n n_0$ times

 $\implies \begin{cases} p_d \sim Cd^{-3} \Rightarrow \text{power law} \\ diam \asymp \log n \Rightarrow \text{small world} \\ (\text{suitably modified}) \Rightarrow \text{high clustering} \end{cases}$

Dynamics over networks

```
network structure + interaction mechanism

\downarrow \downarrow

emerging behavior
```

- random walks
- linear interactions: averaging / voter model
- epidemics: SI, SIR, SIS (contact model)
- other: majority model, evolutionary dynamics, games
- monotone dynamical systems



▶ P stochastic matrix on V, e.g., $P_{ij} = 1/d_i$

$$\triangleright \mathbb{P}(V(t+1) = i | V(t) = i) = P_{ij}$$

▶ network connected ⇒ unique stationary distr. π ($\pi_v = \frac{d_v}{\sum_u d_u}$)



▶ P stochastic matrix on V, e.g., $P_{ij} = 1/d_i$

$$\triangleright \mathbb{P}(V(t+1) = i | V(t) = i) = P_{ij}$$

▶ network connected ⇒ unique stationary distr. $\pi (\pi_v = \frac{d_v}{\sum_u d_u})$



▶ P stochastic matrix on V, e.g., $P_{ij} = 1/d_i$

$$\triangleright \mathbb{P}(V(t+1) = i | V(t) = i) = P_{ij}$$

▶ network connected \Rightarrow unique stationary distr. π $(\pi_v = \frac{d_v}{\sum_u d_u})$



▶ P stochastic matrix on V, e.g., $P_{ij} = 1/d_i$

$$\triangleright \mathbb{P}(V(t+1) = i | V(t) = i) = P_{ij}$$

▶ network connected ⇒ unique stationary distr. $\pi (\pi_v = \frac{d_v}{\sum_u d_u})$



▶ P stochastic matrix on V, e.g., $P_{ij} = 1/d_i$

$$\triangleright \mathbb{P}(V(t+1) = i | V(t) = i) = P_{ij}$$

▶ network connected \Rightarrow unique stationary distr. π $(\pi_v = \frac{d_v}{\sum_u d_u})$



> P stochastic matrix on \mathcal{V} , e.g., $P_{ij} = 1/d_i$

$$\triangleright \mathbb{P}(V(t+1) = i | V(t) = i) = P_{ij}$$

▶ network connected ⇒ unique stationary distr. π $(\pi_v = \frac{d_v}{\sum_u d_u})$



- ▶ how fast does $||\mathbb{P}(V(t) = \cdot) \pi||$ go to 0 ?
- ▶ when will V(t) hit some other $w \in \mathcal{V}$?
- ▶ when will independent V(t) and $\tilde{V}(t)$ meet for the first time?

Distributed averaging



Gossip model:

- every node v has a state $x_v(t) \in \mathbb{R}$
- nodes get activated at independent Poisson times
- ▶ when a node v is activated, it choses a neighbor w at random and updates its value to $x_v(t) = (1 \omega)x_v(t^-) + \omega x_w(t^-)$
- network connected \implies convergence to consensus

Distributed averaging



How is the limit consensus value related to the initial states of the nodes?

How does the network structure affect the speed of convergence? What is the effect of heterogeneity of the agents behavior?

Consensus vs disagreement

most mathematical models: connected network ⇒ (asymptotic) consensus



"Since universal ultimate agreement is an ubiquitous outcome of a very broad class of mathematical models, we are naturally led to inquire what on earth one must assume in order to generate ..." (Abelson '64)



"If people tend to become more alike in their opinions, attitudes, and behavior as they interact, why do not such differences eventually disappear?" (Axelrod '97)
Gossip model with stubborn agents



model propaganda: political parties, media sources, advertising, ...

Gossip model with stubborn agents (cont'd)

 $\mathcal{G} = (\mathcal{V}, \mathcal{E})$ connected opinions: $X_{v}(t) \in \mathbb{R}$

 $\mathcal{V}=\mathcal{A}\cup\mathcal{S}$

 $\mathcal{A} = \{ \textbf{regular} \text{ individuals} \}$



 $\omega \in]0,1]$ 'trust' $T^a :=$ random clock (rate-1 Poisson)

 $T_k^a = t \implies a \text{ chooses } b \sim a \text{ at random}$

 $X_a(t^-) = x, \ X_b(t^-) = y \implies X_a(t) = x + \omega(y - x)$

Typical sample-path behavior

 $X_a(t)$



 $\exists s, s' \in S : x_s \neq x_{s'} \implies \mathbb{P}(\mathsf{NO \ convergence, \ NO \ consensus}) = 1$

Typical sample-path behavior (cont'd)





 $\mathbb{P}(Z_a(t) \text{ converges}) = 1$

Voter model

 $\omega = 1 \Longrightarrow$ voter model: $X_a(t) = X_b(t^-)$



Dual process: $(V_i(t))_{i \in \mathcal{V}}$ coalescing random walk, absorbing set S

$$(X_i(t))_{i\in\mathcal{V}} \stackrel{d}{=} (X_{V_i(t)}(0))_{i\in\mathcal{V}}$$



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



 \triangleright S=susceptible I=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



 \triangleright S=susceptible I=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



► *S*=susceptible *I*=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)



$\blacktriangleright X_v(t) \in \{S,I\}$

▶ $x_v(t) \rightarrow I$ if and only if v is in the same connected component of some w with $x_w(0) = I$

how fast will every node become infected?



- ▶ S=susceptible I=infected R=recovered
- $\triangleright X_{\nu}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- ▶ susceptible node becomes infected when meeting infected node
- random nodes get recovered (if infected)



- ▶ S=susceptible I=infected R=recovered
- $\triangleright X_{\nu}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times



- ▶ S=susceptible I=infected R=recovered
- $\triangleright X_{\nu}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times



- > S=susceptible I=infected R=recovered
- \triangleright $X_v(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times



- ▶ S=susceptible I=infected R=recovered
- $\triangleright X_{\nu}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times



- ▶ S=susceptible I=infected R=recovered
- $\triangleright X_{\nu}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times



- > S=susceptible I=infected R=recovered
- $\triangleright X_{\nu}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times



- ▶ S=susceptible I=infected R=recovered
- $\blacktriangleright X_{\nu}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times



- ▶ S=susceptible I=infected R=recovered
- $\blacktriangleright X_{v}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- ▶ infected nodes recover (die) at random times



- ▶ S=susceptible I=infected R=recovered
- \triangleright $X_v(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times


- ▶ S=susceptible I=infected R=recovered
- $\blacktriangleright X_{v}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times



- ▶ S=susceptible I=infected R=recovered
- $\blacktriangleright X_{v}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times



- ▶ S=susceptible I=infected R=recovered
- $\blacktriangleright X_{v}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times



- ▶ S=susceptible I=infected R=recovered
- $\blacktriangleright X_{v}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times



- \triangleright S=susceptible I=infected R=recovered
- $\blacktriangleright X_{\nu}(t) \in \{S, I, R\}$
- ▶ a random link gets activated at t (meeting)
- susceptible node becomes infected when meeting infected node
- infected nodes recover (die) at random times



 \triangleright S=susceptible I=infected R=recovered

 $\triangleright X_{v}(t) \in \{S, I, R\}$

▶ the process stops in finite T with $x_v(T) \in \{I, R\}$

▶ how big is *T*?

▶ what is the fraction of $x_v(T) = R$?

Epidemics 3: SIS (a.k.a. contact model)



► S=susceptible I=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)

susceptible node becomes infected when meeting infected node



► S=susceptible I=infected

 $\blacktriangleright X_v(t) \in \{S,I\}$

▶ a random link gets activated at t (meeting)

> susceptible node becomes infected when meeting infected node



► S=susceptible I=infected

 $\blacktriangleright X_v(t) \in \{S,I\}$

▶ a random link gets activated at t (meeting)

> susceptible node becomes infected when meeting infected node



► S=susceptible I=infected

 $\blacktriangleright X_v(t) \in \{S,I\}$

▶ a random link gets activated at t (meeting)

> susceptible node becomes infected when meeting infected node



► S=susceptible I=infected

 $\blacktriangleright X_v(t) \in \{S,I\}$

▶ a random link gets activated at t (meeting)

> susceptible node becomes infected when meeting infected node



► S=susceptible I=infected

► $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)

> susceptible node becomes infected when meeting infected node



► S=susceptible I=infected

► $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)

> susceptible node becomes infected when meeting infected node



► S=susceptible I=infected

 $\blacktriangleright X_v(t) \in \{S,I\}$

▶ a random link gets activated at t (meeting)

> susceptible node becomes infected when meeting infected node



► S=susceptible I=infected

► $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)

> susceptible node becomes infected when meeting infected node



► S=susceptible I=infected

 $\blacktriangleright X_v(t) \in \{S,I\}$

▶ a random link gets activated at t (meeting)

> susceptible node becomes infected when meeting infected node



► S=susceptible I=infected

► $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)

> susceptible node becomes infected when meeting infected node



 \triangleright S=susceptible I=infected

 \triangleright $X_v(t) \in \{S, I\}$

▶ a random link gets activated at t (meeting)

▶ in finite (very very large) $T x_v(T) = S$ for all v

• metastable state with nontrivial fraction of $x_v = S$

More dynamics



majority voter

threshold models

▶ Moran process: evolution

games

dynamic flows

Analysis techniques

- Perron-Frobenius theory of positive systems
- monotone dynamical systems
- fast-mixing
- coupling
- duality
- martingale arguments
- mean-field limits
- branching process approximations
- density evolution

Next meetings: conference room M-building 1st floor

▶ Wed, October 2, 10:15-12:00: lecture ▶ Tue. October 8. 13:15-15:00: exercise ▶ Wed, October 9, 10:15-12:00: lecture ▶ Tue, October 15, 13:15-15:00: exercise ▶ Wed, October 16, 10:15-12:00: lecture ▶ Tue, October 22, 13:15-15:00: exercise ▶ Wed, October 23, 10:15-12:00: lecture ▶ Tue, October 29, 13:15-15:00: exercise ▶ Wed. October 30, 10:15-12:00: lecture ▶ Tue, November 5, 13:15-15:00: exercise ▶ Wed, November 6, 10:15-12:00: lecture ▶ Tue, November 12, 13:15-15:00: exercise ▶ Wed, November 13, 10:15-12:00: lecture Tue. November 19, 13:15-15:00: exercise ▶ Wed, November 20, 10:15-12:00: lecture ▶ Tue, November 26, 13:15-15:00: exercise ▶ Wed, November 27, 10:15-12:00: lecture