

Aalto University School of Science and Technology

CS-E5745 Mathematical Methods for Network Science

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Dynamical models of/on networks

Learning goals this week:

- Gain insights on the big picture on how to approach network dynamic problems analytically
- Ability to recognize and formulate approximations for these problems
- Ability to write down master/rate equations
- Newman: Networks, An Introduction Sections 14 and 17

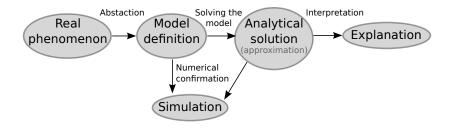


Dynamical models of/on networks

- Models of networks: Network evolves according to some rules
- Models on networks: States of nodes evolve according to some rules
- Co-evolution models: Both network and states of nodes evolve simultaneously
- Similar techniques can be used to solve all of these

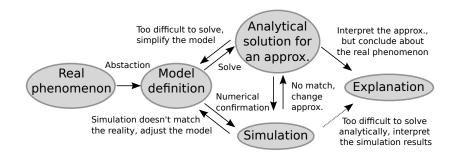


Pipeline, as presented in a publication





Pipeline, a more realistic picture





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- Phenomena: Networks are born via growth, new nodes are more likely to connect to already popular nodes
- Data: Probability of link creation can be measured as a function of degree, power-law degree distributions
- Model: A variation of preferential attachment models (e.g., Price model):
 - 0. Start with t_0 seed nodes with some connections, and set time to $t = t_0$
 - 1. Increase time t = t + 1, add node with index t
 - 2. Add *m* links from *t* to other nodes, each link to node *i* prob. $p_i = \frac{k_i}{\sum_j k_j}$
 - 3. GOTO 1.
- There could be multi-links (or the probabilities are different), but we ignore this

- We can solve the BA model *exactly* if we so wish
- Idea: Think about the BA model as a Markov chain where the state space is (a subset of) all graphs
 - Easy to write the transition probabilities between graphs
 - The result give us a probability distribution over all graphs for each time t



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- Idea: Think about the BA model as a Markov chain where the state space is (a subset of) all graphs
 - Easy to write the transition probabilities between graphs
 - The result give us a probability distribution over all graphs for each time t
- Problem: the size of the state space explodes
 - > $2^{T(T-1)/2}$ graphs with T nodes, but not all reachable
 - For every state at time t, there are $\binom{t}{m}$ possible transitions
 - ► Total number of states at time *T* is $N_T = \prod_{t=t_0}^T {t \choose m} \ge e^{mT}$ (for *T* large enough)
- This solution is not practical



Approximate solutions

- Idea: Define a simple system that mimics the (expected) behavior of a complex system
- The state of the system at each time is reduced to a property or a vector of properties
 - The choise of properties is usually guided by the process
- The evolution rules are written in terms of average behavior (master/rate equation)
 - Either as difference or differential equation(s)
- Only mean behavior matters: statistical fluctuations are ignored
 - The properties are assumed to be independent
 - Dependencies can be added by including combinations of properties
- The resulting system is usually much easier to solve!



Typical procedure in approximate solutions

- ► The "real" model for the system *S* is defined using stochatic rules *F* such that S(t) = F(S(t-1))
- Calculate some properties X of the system $X(S) = \mathbf{X}$
 - The properties can be discrete in the system S but evolve to be continuous in X: e.g., probability that an edge exists
- ► Define deterministic rules *f* for the evolution of the properties $\mathbf{X}(t) = f(\mathbf{X}(t-1))$, s.t. $f(\mathbf{X}) \approx \langle F(S) | X(S) = \mathbf{X} \rangle$

• Usually defined in elementwise way: $X_i(t) = f_i(\mathbf{X}(t-1))$

- Set $X(S(t_0)) = \mathbf{X}(t_0)$ and solve $\mathbf{X}(t) = f(\mathbf{X}(t-1))$ for any value of t or for $t \to \infty$
 - We now hope that $X(\langle F^t(S(t_0)) \rangle) \approx f^t(\mathbf{X}(t_0))$
 - For continuous time one defines $\frac{d}{dt}\mathbf{X}(t) = f(\mathbf{X}(t))$



- Idea: map each graph to a vector where each element corresponds to an edge, s.t. X_i = 1 if edge exists, and 0 otherwise
 - Expected value of binary variable is the probability of getting 1
 - Notation: A_{ij} is the probability that link between i and j exists
 - Set m = 1 for simplicity
- ► The update rule for **X**(*t*):
 - Calculate expected degree for each node $\langle k_i \rangle = \sum_i A_{ij}$

• Set
$$A_{it} = \frac{\langle k_i \rangle}{\sum_i \langle k_j \rangle}$$

The final state X(T) gives the probability of each edge existing at time T



Practical guidelines for selecting variables for approximations

- Anything that can be measured from the state of the system could be selected as a property we track
 - Some selections make it difficult to come up with the update function
 - The selecting variables appearing in the update rule is a good idea
- The above approximation discards dependencies between the variables, but one can define a combination variables that keep track of these
 - Example: Probability for each triangle to exists, instead of a probability each link to exists



Idea 1: map each graph into a vector of degrees of nodes
The expected degree of each node is tracked
Note: expected degrees are continuous variables
X_i(t) is the expected degree of node *i* at time step t
Now we should write the equations X_i(t) = f_i(X(t-1))



 Idea 2: map each graph into a vector of counts of nodes of each degree (or a degree distribution)

The expected number of nodes of each degee is tracked

- X_k(t) is the expected number of nodes of degree k at time step t
- ▶ Now we should write the equations $X_k(t) = f_k(\mathbf{X}(t-1))$



Practical guidelines for writing master equations

- When the tracked variables are not binary variables, it might be more difficult to write the update equations
- Think about each element $X_i(t)$ separately, divide it to
 - growth $X_i^+(t) = f_i^+(\mathbf{X}(t-1))$ term and
 - decline $X_i^-(t) = f_i^-(\mathbf{X}(t-1))$ term
- The total is $X_i(t) X_i(t-1) = X_i^+(t) X_i^-(t) = f_i^+(\mathbf{X}(t-1)) + f_i^-(\mathbf{X}(t-1))$
- Are there any conserved quantities? (If something grows, others decline)
- Same applies for rate equations



 X_k(t) is the expected number of nodes of degree k at time step t

Addition of nodes to degree categories:

•
$$X_k^+(t) = m \frac{(k-1)}{\sum_k k X_k(t-1)} X_{k-1}(t-1)$$
, when $k > m$

•
$$X_k^+(t) = 1$$
, when $k = m$

Removal of nodes from degree categories:

$$X_k^-(t) = m_{\frac{k}{\sum_k k X_k(t-1)}} X_k(t-1)$$

• Note that $\sum_k kX_k(t) = 2mN(t)$

In total we have:

•
$$X_k(t) = X_k(t-1) + \frac{(k-1)}{2N(t-1)}X_{k-1}(t-1) - \frac{k}{N(t-1)}X_k(t-1),$$

when $k > m$

•
$$X_m(t) = X_m(t-1) + 1 - \frac{m}{N(t-1)}X_m(t-1)$$



Models on networks

- Above approach can also be used to solve models on networks
- The state of the systems is not a graph, a vector of states for the nodes
- Examples:
 - Infection spreading: Nodes are either infected or not infected
 - Opinion formation: Each nodes state represent its opinion (discreet or continuous value)



Spreading models on networks

- Network spreading models follow the classic literature on mathematical epidemiology
- Models are determined by the possible states and transition rates between them
- States:
 - S: Susceptible
 - E: Exposed
 - I: Infected
 - R: Recovered (or Removed)
- Naming convention: state changes from left to right
- Typical models include: SI, SIS, SIRS, SEIR



Example: SIR on networks

$\blacktriangleright S \xrightarrow{\beta} I \xrightarrow{\gamma} R$

- Classic result (no network): Epidemic if $R_0 = \frac{\beta}{\gamma} > 1$
 - i.e., when rate or infectino is larger than rate of recovery
- Degree-based approximation: Epidemic if $\frac{\beta}{\gamma} > \frac{1}{\langle q \rangle}$

where \langle q \rangle is the expected excess degree

• Node-based approximation: Epidemic if $\frac{\beta}{\gamma} > \frac{1}{\lambda_1}$

• where λ_1 is the largest eigenvalue of an adjacency matrix



SIR can be mapped to percolation

- Modification: all individuals stay infected for time *τ* (instead of I → R)
- Infected node: probability that an *edge* leading out is activated during infection is Φ = 1 − e^{−βτ}
- Set edges occupied with probability $\Phi \to$ components give you the possible sizes of epidemics
 - We can now calculate the statistics of epidemics using percolation theory on networks



- A general theory for approximating processes where each node can be in one of two states¹
 - Examples: SI, SIS, voter model, ...
 - Assumption: network is produced by configuration model
- Probabilities that node of degree k is susceptible (infected) and has m infected neighbors: s_{k,m} (i_{k,m})
- Dynamics determine the rates $S \rightarrow I$: $F_{k,m}$ and $I \rightarrow S$: $R_{k,m}$
 - Example: in the SIS model $F_{k,m} = m\beta$ and $R_{k,m} = \gamma$

¹Phys. Rev. Lett. 107, 068701 (2011)



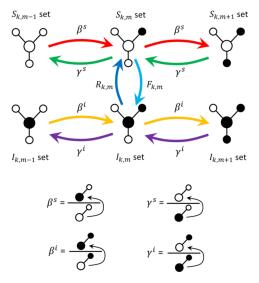
The master equations become complicated:

$$\frac{ds_{k,m}}{dt} = -F_{k,m}s_{k,m} + R_{k,m}i_{k,m}$$
$$-\beta^{s}(k-m)s_{k,m} + \beta^{s}(k-m+1)s_{k,m-1}$$
$$-\gamma^{s}ms_{k,m} + \gamma^{s}(m+1)s_{k,m+1}$$

• Where
$$\beta^{s} = \frac{\langle \sum_{m=0}^{k} (k-m)F_{k,m}s_{k,m} \rangle}{\langle \sum_{m=0}^{k} (k-m)s_{k,m} \rangle}$$
, $\gamma^{s} = \frac{\langle \sum_{m=0}^{k} (k-m)R_{k,m}i_{k,m} \rangle}{\langle \sum_{m=0}^{k} (k-m)i_{k,m} \rangle}$
• Similar equation for i_{k}

Similar equation for i_{k,m}







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- ► The $(k_{max} + 1)(k_{max} + 2)$ master equations can be solved numerically, or further approximations can be made
- Pair approximation (PA):
 - ρ_k : Probability that node of degree k is infected
 - *p_k*: Probability that neighbor of a node of degree k is infected

• $s_{k,m} \approx (1 - \rho_k) B_{k,m}(p_k)$, where $B_{k,m}(p) = {\binom{k}{m}} p^m (1 - p)^{k-m}$

- Mean-field approximation (MF):
 - *ρ_k*: Probability that node of degree k is infected

•
$$\mathbf{s}_{k,m} \approx (1 - \rho_k) \mathbf{B}_{k,m}(\langle \frac{k}{z} \rho_k \rangle)$$



