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Statistical Mechanics of Complex Networks

Lecture 5: Evolution (Scale-free and Structural balance)

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Networks can be disordered



- Positive interaction
- -- Negative interaction
 - Thickness indicates strength of interaction

Many naturally occurring networks have links with heterogeneously distributed properties

Differences in links can be
Quantitative: distribution of degree and/or link weights
Qualitative: nature of interactions (+ve or -ve)



"Scale free" networks

Barabasi and Albert (1999): In many large networks node connections follow a scale-free distribution \Rightarrow degree distribution has a power law tail

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P(k) \sim k^{-\alpha}
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In contrast,

Nodes in Erdos-Renyi random networks, e.g., G(N,p), exhibit Poisson degree distrn: P(k) = $e^{-\lambda} (\lambda^k/k!)$

Power laws, Pareto distributions and Zipf's law

"When the probability of measuring a particular value of some quantity varies inversely as a power of that value, the quantity is said to follow a power law, also known variously as Zipf's law or the Pareto distribution"

Mark E J Newman Contemporary Physics, **46** (2005) 323



$$P(w) = C w^{-\alpha} exponent P(w)$$

$$\Rightarrow \log P(w) = -\alpha \log(w) + \log(C)$$

$$Y = -\alpha X + B \log P(w)$$

$$slope:$$

Appears as a linear relation in a log-log graph (i.e., the axes of the graph are expressed logarithmically) -α

log w

Contrast with Distributions having a characteristic scale

length scale)

Example: Exponential distribution

$$P(w) = C e^{-w/w^{*}}$$

Mean : w* Standard deviation: w*

For instance, Radioactive decay

 $P(t) = C e^{-t/\tau} \tau$: half-life

Or, waiting time between buses arriving

Essentially, exponential distribution of times between events seen when events are independent and occur at a constant average rate (i.e., a Poisson process)



Power law distribution \Rightarrow "Long tails"

High probability of having values that are extremely large deviations from the mean, much larger than that expected from the variance



Networks with power-law degree distribution characterized by

Hubs

highly connected nodes which hold the network together



Random networks with arbitrary degree distributions

The configuration model

Generating random networks with any desired degree distribution (instead of Poisson, as in ER graphs), or rather, desired degree sequence

If the exact degree k_i of each individual node i=1,2,...,N in the network is specified a random network with these degrees is created by matching "stubs"



$k_{a} = 3$ $k_{b} = 2$ $k_{c} = 1$ $k_{c} = 1$

Recipe

- Assign to each node i a total of k_i stubs of edges ("half-edges") such that $\sum k_i = 2L$ where L is the total number of links
- Now choose a pair of stubs at random and connect to form a link
- Continue by choosing another pair of stubs randomly from the remaining 2L – 2 stubs, and so on until all stubs are used
- Results in a network in which each node i has exactly the assigned degree k_i

analogous to the G(N,L) random graph model, where the number of links is fixed

How can scale-free networks evolve ?

The Price-Barabasi-Albert preferential attachment scheme:

- (A) Networks expand continuously by addition of new nodes
- (B) New nodes attach preferentially to nodes already wellconnected, i.e., probability that a new node is connected to a node of degree k_i is $\Pi(k_i)=k_i/\Sigma_jk_j$ ("linear" scheme)

Resulting network degree distribution: $P(k) \sim k^{-3}$

The birth of a BA scale-free network (H Jeong)





10[°]

The Mechanism: A Cumulative Advantage Process

Derek J de Solla Price : the earliest mathematically detailed mechanism by which scale-free degree distribution can arise in the context of networks of citations between scientific papers



Derek J de Solla Price

Journal of the American Society for Information Science—September-October 1976 A General Theory of Bibliometric and Other Cumulative Advantage Processes*

Derek de Solla Price

Department of History of Science and Medicine Yale University New Haven, CT 06520

A Cumulative Advantage Distribution is proposed which models statistically the situation in which success breeds success. It differs from the Negative Binomial Distribution in that lack of success, being a non-event, is not punished by increased chance of failure. It is shown that such a stochastic law is governed by the Beta Function, containing only one free parameter, and this is approximated by a skew or hyperbolic distribution of the type that is widespread in bibliometrics and diverse social science phenomena. In particular, this is shown to be an appropriate underlying probabilistic theory for the Bradford Law, the Lotka Law, the Pareto and Zipf Distributions, and for all the empirical results of citation frequency analysis. As side results one may derive also the obsolescence factor for literature use. The Beta Function is peculiarly elegant for these manifold purposes because it yields both the actual and the cumulative distributions in simple form, and contains a limiting case of an inverse square law to which many empirical distributions conform.

The Rich get Richer

Distribution of citations to scientific papers



The stem cell wars

© NewScientist

The most influential players in cellular reprogramming are revealed by recording how many times the scientists have referred to each other's work. Each link shows where one researcher cited another four or more times in papers in leading journals (for analysis, see "The strongest link", below right)



Image: Chatterjee et al, PLoS One (2016)

Matthew Effect

Image: ThoughtCo



Robert Merton



Already well-known scientists receive disproportionate credit for their contributions, while less-known scientists receive less credit than their works merit

Why the degree distribution is scale-free

Start with a small number (m_0) of nodes, at every time step, add a new node with $m(\leq m_0)$ links to nodes already present in the system.

From

Statistical mechanics of complex networks

Reka Albert and Albert-Laszlo´Barabasi Rev Mod Phys 74 (2002) 47-97

Continuum theory: The continuum approach introduced by Barabási and Albert (1999) and Barabási, Albert, and Jeong (1999) calculates the time dependence of the degree k_i of a given node *i*. This degree will increase every time a new node enters the system and links to node *i*, the probability of this process being $\Pi(k_i)$. Assuming that k_i is a continuous real variable, the rate at which k_i changes is expected to be proportional to $\Pi(k_i)$. Consequently k_i satisfies the dynamical equation

$$\frac{\partial k_i}{\partial t} = m \Pi(k_i) = m \frac{k_i}{N-1}.$$

$$\sum_{j=1}^{N-1} k_j$$
(79)

The sum in the denominator goes over all nodes in the system except the newly introduced one; thus its value is $\sum_{j} k_{j} = 2mt - m$, leading to

$$\frac{\partial k_i}{\partial t} = \frac{k_i}{2t}.$$
(80)

The solution of this equation, with the initial condition that every node *i* at its introduction has $k_i(t_i) = m$, is

$$k_i(t) = m \left(\frac{t}{t_i}\right)^{\beta}$$
 with $\beta = \frac{1}{2}$. (81)

Equation (81) indicates that the degree of all nodes evolves the same way, following a power law, the only difference being the intercept of the power law.

Using Eq. (81), one can write the probability that a node has a degree $k_i(t)$ smaller than k, $P[k_i(t) < k]$, as

$$P[k_i(t) < k] = P\left(t_i > \frac{m^{1/\beta}t}{k^{1/\beta}}\right).$$
(82)

Assuming that we add the nodes at equal time intervals to the network, the t_i values have a constant probability density

$$P(t_i) = \frac{1}{m_0 + t}.$$
(83)

Substituting this into Eq. (82) we obtain

$$P\left(t_i > \frac{m^{1/\beta}t}{k^{1/\beta}}\right) = 1 - \frac{m^{1/\beta}t}{k^{1/\beta}(t+m_0)}.$$
(84)

The degree distribution P(k) can be obtained using

$$P(k) = \frac{\partial P[k_i(t) < k]}{\partial k} = \frac{2m^{1/\beta}t}{m_0 + t} \frac{1}{k^{1/\beta + 1}},$$
(85)

predicting that asymptotically $(t \rightarrow \infty)$

$$P(k) \sim 2m^{1/\beta} k^{-\gamma} \quad \text{with} \quad \gamma = \frac{1}{\beta} + 1 = 3 \tag{86}$$

being independent of m, in agreement with the numerical results.

Importance of "hubs"



However targeting the highest-degree nodes (hubs) has devastating effect on the network – most nodes become isolated on removing a few hubs: *Vulnerability to targeted removal of hubs*

How does a heterogeneous degree distribution affect dynamical processes on the network – e.g., epidemic propagation?

The Kermack-McKendrick S-I-R Model (1927)



 $= N \beta \tau > I$

 R_0

 \Rightarrow Condition for epidemic S > 1 / β τ

As in the initial stage of an epidemic S = N, total population... An epidemic will occur if N $\beta \tau > 1$

Minimum immunization coverage required to stop epidemic (obvious policy implications)

SIR model equation: dl/dt = β SI – 1/ τ \Rightarrow To stop epidemic we need to make dl/dt < 0, i.e., S(t=0) < 1 / $\beta\tau$ where

- $\boldsymbol{\beta}$: rate of infection spreading
- $\boldsymbol{\tau}$: average infectious period

Let total population be N

Thus, proportion of the population that is susceptible, s = S(t=0)/N needs to be made smaller than $I/(N\beta\tau) = I/R_0$ (because $R_0 = N \beta\tau$)

 \Rightarrow The fraction of population that needs to be immunized to stop the epidemic (assuming homogeneous mixing) is p > 1- (1/R₀) For R₀ \approx 2, p_{min} \approx 50%, while for R₀ \approx 3, p_{min} \approx 66%

No threshold for epidemics in scale-free networks

Networks of sexual relations have been claimed to be scale-free !

VOLUME 90, NUMBER 2

PHYSICAL REVIEW LETTERS

week ending 17 JANUARY 2003

A few highly promiscuous individuals act as "hub" nodes

plays crucial role in spreading sexually transmitted diseases !

Absence of Epidemic Threshold in Scale-Free Networks with Degree Correlations

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> Random scale-free networks have the peculiar property of being prone to the spreading of infections. Here we provide for the susceptible-infected-susceptible model an exact result showing that a scale-free degree distribution with diverging second moment is a sufficient condition to have null epidemic threshold in unstructured networks with either assortative or disassortative mixing. Degree correlations result therefore irrelevant for the epidemic spreading picture in these scale-free networks. The present result is related to the divergence of the average nearest neighbor's degree, enforced by the degree detailed balance condition.

If the contact structure of a disease is network with heterogeneous degree distribution, the condition for occurrence of an epidemic is: $R = N\beta\tau > \langle k \rangle / \{\langle k^2 \rangle - \langle k \rangle\}$

For a scale-free network having degree exponent $2 < \alpha \le 3$, $k^2 > \rightarrow \infty$ \Rightarrow There is no epidemic threshold !

Even diseases with extremely low transmission probabilities are likely to cause a major outbreak involving a significant fraction of population

But are most real networks "scale free" ?

- Scale-free networks characterized by long-tailed degree distribution (power laws) have been proposed as unifying concept for biological complex systems – have been reported in metabolic, protein, and gene interaction networks.
- But many of these reports of scale-free networks are possibly just a result of bad statistics (a combination of extremely limited data and faulty analysis) !
- Almost any distribution seen over a small enough range in a double logarithmic scale would appear linear – and wrongly interpreted as power law



Guelzim et al., 2002

- To establish power laws from finite data one has to use unbiased techniques such as maximum likelihood estimation.
- Rigorous re-analysis of many of the data sets used by earlier studies that claimed power-law degree distributions have shown little evidence for scale-free nature!
 (E.g., R Khanin & EWit, / Comp Biol 13

(E.g., R Khanin & E Wit, J Comp Bio (2006) 810)

Can other processes yield scale-free degree distribution? The Case of Duplication & Divergence

preferential attachment appropriate for explaining scale-free character of WWW Less clear how it might play a role in biological systems, e.g., protein-protein interaction network that has been claimed to be scale-free

As most biological systems have emerged through a long history of evolution, can evolutionary processes give rise to a network with scale-free property ?

Vazquez et al, ComPlexUs (2003)

In the Duplication-Divergence mechanism, a node along with all its interactions are duplicated with probability p and then some of the interactions mutated with probability $q \Rightarrow$ claimed to yield networks with scale-free degree distribution but is it true?



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Heider's "Three-Body Problem"

A basic characterization of relationships between mutual acquaintances





Fritz Heider (1896-1988)

F Heider (1946) Attitudes and cognitive organization. J Psychol 21:107–112.

aminoapps.com

- (I) Om and Xena are friends \Rightarrow (2) Pradeep and Xena are friends \Rightarrow PX: +ve interaction (link) (3) Om and Pradeep are enemies \Rightarrow
 - OX: +ve interaction (link) OP: -ve interaction (link)



Structural Balance



Relationship triangles containing exactly 2 friendships are prone to transition to triangles with either 1 or 3 friendships \Rightarrow friend of my enemy is my enemy

A single friendship may appear in a relationship triangle that initially had none \Rightarrow enemy of my enemy is my friend

Structural Balance from triads to networks



Dorwin Cartwright Frank Harary (1915-2008) (1921-2005)

Carwright & Harary (1956)

Generalization of Heider's theory to network of N nodes Psychol Rev 63:277–293

In a balanced network, every cycle (closed loop) is *balanced*, i.e., product of the signs of the links in the loop is +ve

A <u>complete</u> graph (a network where all pairs of nodes are connected) is balanced if each constituent triad is balanced

The local concept of balance results in non-trivial network structure

Any balanced network can be partitioned into two communities such that all edges inside each community are positive and all edges between nodes in opposite communities are negative (one of these communities may be empty)



In absence of any external influence or noise, the two communities are unified and opposed in their response to any issue

Balance and International Relations



Alliance network of nations in 1962 As bipartite relations among countries that comprise major alliances change through events such as war, triads become unbalanced \Rightarrow creates tension \Rightarrow Reorganization into a balanced state involving new blocs and alliances (Evolution to balance) For physicists

Structural balance \equiv No Frustration

E.g., Ising spin systems with exchange interactions of FM or AFM type



-ve -ve

Absence of structural balance would result in a rugged energy landscape, with the system trapped in any one of a large number of local minima

A balanced network would have smooth energy landscape



e.g., Spin Glasses

Structural balance in the brain ?

On the basis of spontaneous correlations & anti-correlations of fluctuations in fMRI between different brain regions, two "diametrically opposed" widely distributed brain regions identified.

One network consists of regions routinely exhibiting task-related activations; the other of regions routinely exhibiting task-related deactivations.

Fox et al, PNAS 102 (2005) 9673



But how is balance achieved ?

Most studies on structural balance have been carried out in the context of social networks

Can other kinds of networks, in particular those that occur in biology, exhibit balance ?

And if so, what is the mechanism of evolution to balance ?

In particular, can balance be achieved as an outcome of link adaptation dynamics that depends on the state dynamics of the nodes of the network



"Learning" to Balance

http://thebrain.mcgill.ca



Donald O Hebb (1904-85)

Hebb's hypothesis (1949)

Neurons that fire together, wire together

Intuitive interpretation

Agents behaving alike have their ties strengthened, while those behaving differently gradually develop antagonistic relations.

Long-term potentiation

First empirical observation (Lomo, 1966) supporting Hebb's hypothesis

Persistent increase in synaptic strength after high-freq stimulation



Hebb-like rule results in evolution to structural balance through Coevolution of coupling strength & spin dynamics



Structural balance: A tool for seeing some order behind the chaos of financial markets ?



Image: Wikipedia

Robust signatures of systemic crises

Financial markets undergo fluctuations at all times – but only occasionally these spill over into the real economy resulting in economic collapse

E.g., 1929 Great Recession and the 2008 Great Recession

Can we distinguish such crises events from day-to-day ups and downs of the market ?

For this purpose look at the long-term evolution of a market over almost a century: the daily closing prices of all stocks in New York Stock Exchange between 31 Dec 1925 and 1 Feb 2012

[CRSP (Center for Research in Security Prices) database]

Identifying the network of relations between different stocks in terms of how similar their prices movements are over time

By spectral analysis of cross-correlation matrix

Correlation Analysis

I. Construct the correlation matrix C composed of correlation values between every pair of stocks

Correlation between returns for stocks i and j:

$$C_{ij} = \langle r_i r_j \rangle$$
 where $r_i = [R_i - \langle R_i \rangle]/\sigma_i$

Data set:

Data split into 85 overlapping periods of 1000 days (labeled Period 0 to Period 84), the temporal window being shifted by 260 days Stocks with > 50 missing days not considered To compare different periods, we consider 300 or 500 stocks in each Sector to which a stock belongs identified by SIC (Standard Industrial Classification) code

Correlation Matrix

Distribution of magnitudes of crosscorrelations varies with time over the period considered.

Distributions are highly asymmetric, skewed towards +ve values.





Degree of asymmetry varies with time, with the distributions in the 1930s and early 1940s and the 1990s onwards era having the largest degree of **skewness** – coincides with periods of marked upheaval in economy.

The problem with analyzing raw correlations

Cross-correlations of infinitely long time series of different variables will reflect actual inter-relations between them

But in reality, we only get data over a finite time !

Due to stochastic fluctuations, even the output of uncorrelated random processes may exhibit spurious correlations, if we calculate C over finite-length time-series !

Question: Can we identify true interactions between variables by filtering out the effect of cross-correlations generated because of such finite time effects ?

Solution: Look at the spectral properties of C and compare with that of an ensemble of random cross-correlation matrices

The spectrum of "Wigner-ium"

Nuclear physics: What is the energy spectrum of a complex nucleus ?



Image: O. Bohigas, R.U. Haq and A. Pandey, in Nuclear Data for Science and Technology (1983)



Wigner: Instead of focusing on specific energy levels, look at the spectral properties (eigenvalues & eigenvectors) of an ensemble of random (Hamiltonian) matrices

Distribution of spacing between neighboring energy levels

$$P(s) = s \exp(-[\pi/4]s^2)$$

II. Obtain eigenvalues of the correlation matrix

If all stocks are uncorrelated, C will be a random (Wishart) matrix with Marchenko-Pastur distribution (1967)

P (
$$\lambda$$
) = $[Q/2\pi] \sqrt{[(\lambda_{max} - \lambda) (\lambda - \lambda_{min})]}$ where Q = T/N
 λ

Bounds of random distrn : $\lambda_{max} = [I + (I/\sqrt{Q})]^2$ and $\lambda_{min} = [I - (I/\sqrt{Q})]^2$



A small fraction of eigenvalues (~ 3%) deviate from random behavior

The largest eigenvalue is more than 28 times larger than the predicted max. random bound + a few "intermediate" eigenvalues

Deviating eigenvalues ⇒Information about interaction structure of the market

Look at eigenvectors *u* of the largest few eigenvalues



All stocks contribute (almost) uniformly to largest eigenvalue ⇒Market mode

Common component affecting all stocks with same bias

Intermediate eigenvalues should reflect group structure in market if eigenvectors are *localized*

However ...

largest eigenmode (*market*) dominates all intra-group correlations (if existing). \Rightarrow no straightforward detection of significantly related groups of stocks. For this purpose, use

Matrix Decomposition Technique

Aim: removing the effect of (i) market mode & (ii) random noise

Expanding correlation matrix as $C = \sum_{i} \lambda_{i} u_{i}^{T} u_{i}$

Allows decomposition of C into contributions due to

- market, common for all stocks
- groups of co-moving stocks (identified with various business sectors)
- random, idiosyncratic effects for each stock

$$C = C_{market} + C_{sector} + C_{random}$$
$$= \lambda_0 u_0^T u_0 + \Sigma_{i=1,...,N_{group}} \lambda_i u_i^T u_i + \Sigma_{i=N_{group}+1,...,N-1} \lambda_i u_i^T u_i$$
Largest eigenvalue Intermediate eigenvalues Random bulk eigenvalues



Higher incidence of larger values of positive correlations in certain periods is representative of higher correlation between groups of stocks.

Reconstructing the interaction network

Method: Use C^{group} to generate an adjacency matrix A, such that • $A_{ij} = I$ if $|C^{group}_{ij}| > 3 \times std$ dev of C^{random} distribution • $A_{ii} = 0$ otherwise

Node colors represent sector



Dec, 2002 - Dec, 2006

Great recession Feb, 2008 - Feb, 2012

Blue links: +ve interactions, **Red** links: – ve interactions

During a major crisis, network structure shows increased
Connection density,
Number of negative edges, and

- Clustering

Kuyyamudi, Chakrabarti & Sinha, PRE (2019)





An even more robust indicator of economic crisis is the loss of structural balance

i.e., the emergence of frustrated triads in the interaction network

A closer look at triangles in the NYSE stock interaction networks

Red: empirical networks, **Gray**: degree-preserved randomized networks

Kuyyamudi, Chakrabarti & Sinha, PRE (2019) 10⁰ 0 +++ : Balanced Triads $\mathbf{0}$ ---: Balanced Triads 10⁰ ±10 -: Unbalanced Triads 10⁻ 10⁻ **Unbalanced Triads** 10⁻⁵ 20 30 60 70 80 50 0 40 Period 1929 Great Crash 2007-08 Crisis

Frustration \Rightarrow **Systemic Risk**

Frustrated triads appear in individual eigenmodes in many periods, but in the network only prior to/during systemic crisis in the economy (e.g., negative growth rate of GDP per capita)



Qualitatively similar behaviour shown by NASDAQ and the FOREX markets

Assignment

Consider a fully connected network of 4 nodes shown on the right, the nodes being labelled A,B,C,D and the links designated AB,BC,CD, etc. (i.e., indicating the pair of nodes that each link connects).

[Note that each of the links are distinct, i.e., AB being negative in a network where all other links are positive is a distinct configuration compared to one in which AD (for instance) is the only link which is negative.]



(b) How many distinct configurations will have 3 links positive and 3 links negative?

(c) Find how many of the total number of configurations with distinct assignment of link signs that you calculated in (a) are balanced. Note that a network is balanced if every closed loop or cycle is balanced, i.e., product of the signs of the links in the loop is + ve. However, instead of having to look at all 4-cycles as well as 3-cycles (triads), you can use the Cartwright-Harary theorem, according to which a fully connected network is balanced if each of the triads (ABC, ABD, etc.) are each individually balanced.

[Hint: find how many distinct triads are there in the network. If for a given configuration, even one of these triads is <u>not</u> balanced (i.e., has an odd number of negative links) the configuration will be <u>not</u> balanced.]