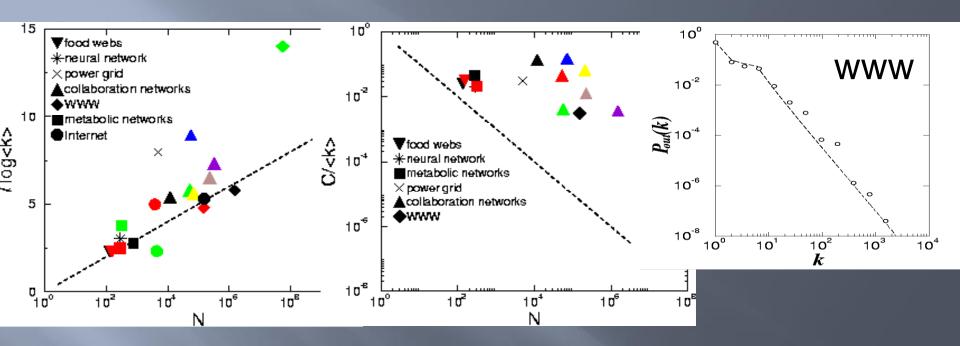
INTRODUCTION TO NETWORK SCIENCE

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6. NETWORK GROWTH MODELS

Complex networks: observations



$$\langle d \rangle \sim \log N$$

Small World:

Average distance scales logarithmically with the network size

$$C/\langle k \rangle = const$$

Clustered:

Clustering coefficient is large, it does not depend on network size.

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

Scale-free:

The degree distribution is broad, with a power law tail.

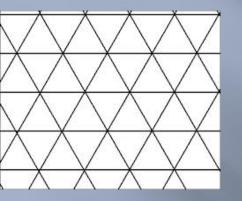
Complex networks: static models

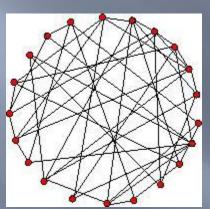
Lattice

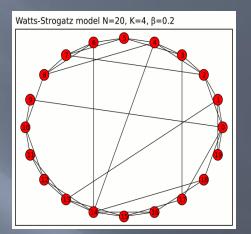
ER graph

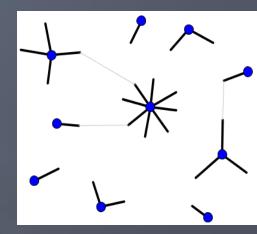
WS graph

Configuration









Tayloring vs Modeling

With configuration type models all requested features can be described – no model in the deeper sense

Amazing universal behavior across a large number of networks: small world, high clustering, broad degree distribution (qualitative universality).

Universality: common mechanism for emergence?

What is common in collaboration, internet, genetic etc. networks?!

Modeling: matter of culture

For a statistician: Modeling is to find a function, which fits data best.

The more we learn about our system the more sophisticated models of this kind are needed.

Such models can be of practical use: You don't need to memorize all the details, you can take the model.

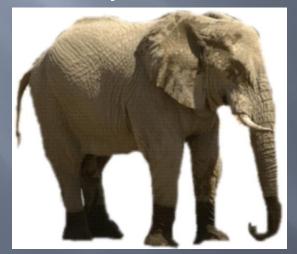
It is like a map of a country. However, if you want more details the resolution of your map has to be better (the fitting functions will have more parameters).

Folklore in physics:

With 2 parameters

Straight line

With 4 parameters



With 3 parameters



Parabola

With parameters > 4



- "What do you consider the largest map that would be really useful?"
- "About six inches to the mile."
- "Only six inches!" exclaimed Mein Herr. "We very soon got six yards to the mile. Then we tried a hundred yards to the mile. And then came the grandest idea of all! We actually made a map of the country, on the scale of a mile to the mile!"
- "Have you used it much?" I enquired.
- "It has never been spread out, yet," said Mein Herr: "The farmers objected: they said it would cover the whole country, and shut out the sunlight! So now we use the country itself, as its own map, and I assure you it does nearly as well."

 Lewis Carroll's Sylvie and Bruno Concluded

Without any theoretical support, such models (fitting functions, maps) do not help much in understanding the phenomenon.

We require from a model more!
It should give insight into the basic mechanisms.

Kepler's 3 laws: Description

Newton's laws + gravity law: Cause → consequence.

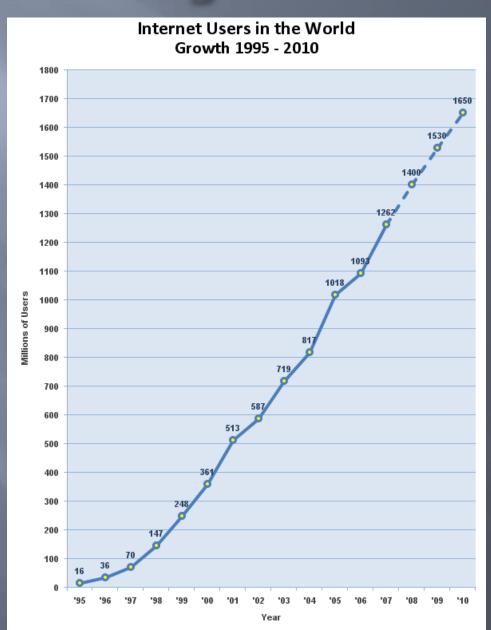
From Newton-type laws there is hope to draw conclusions beyond the observations, predictions e.g., discovery of new planets.

We need Newton's laws for networks.

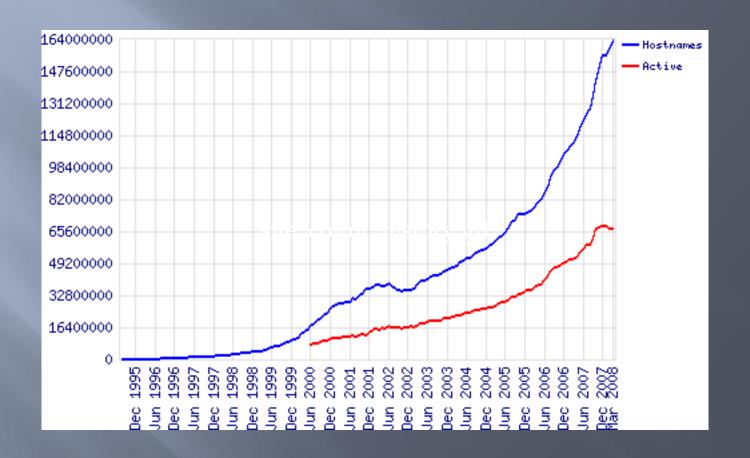
It is not enough to study complex networks as they are given. We have to study how they emerge!

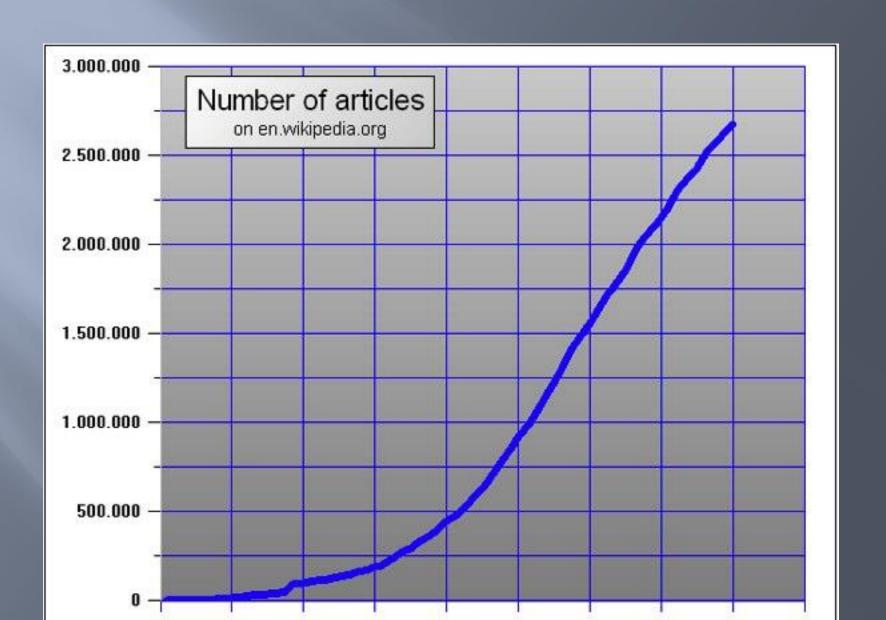
Most networks result from a growth process.

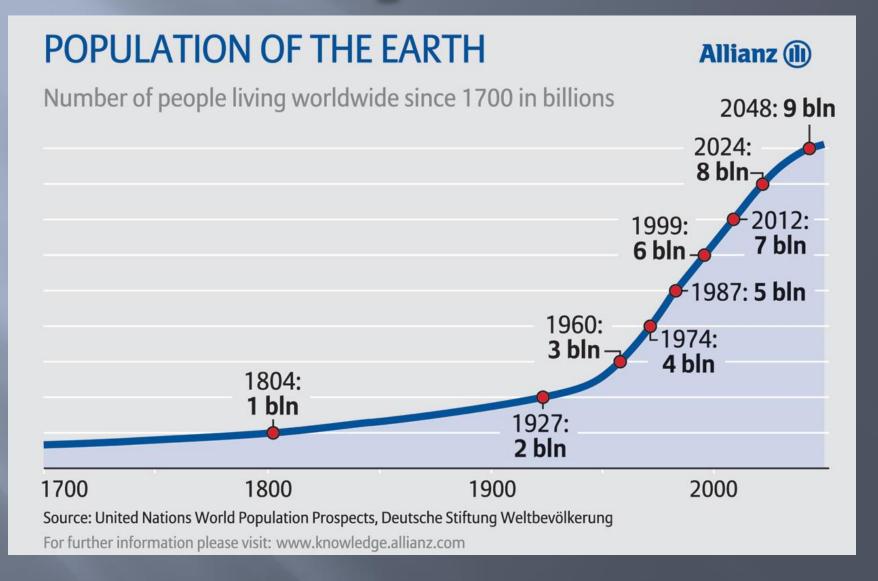
Random networks have constant number of nodes – they are unable to capture the growth aspect.

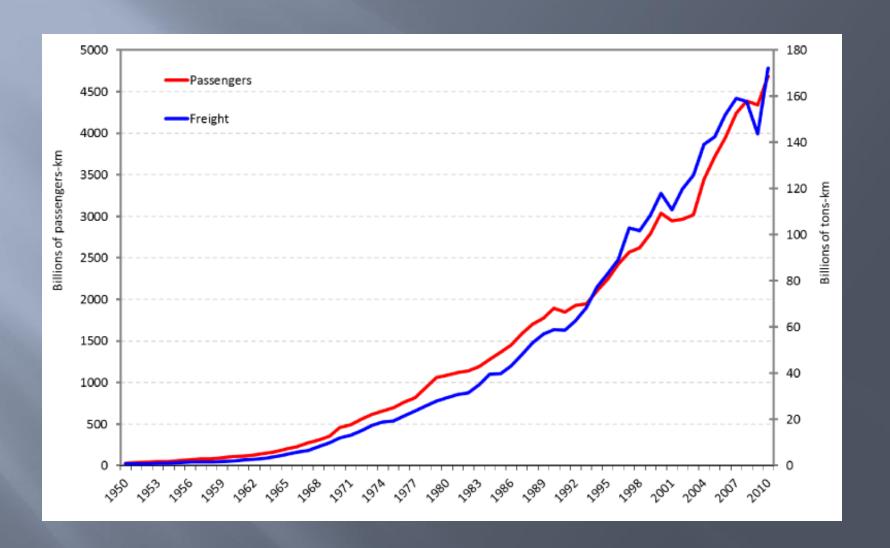


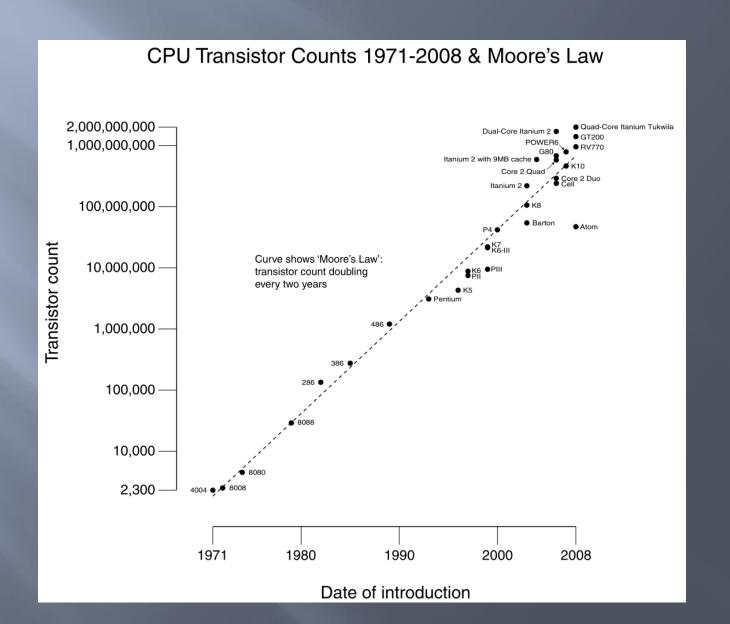
Source: www.internetworldstats.com - January, 2008 Copyright © 2008 Miniwatts Marketing Group

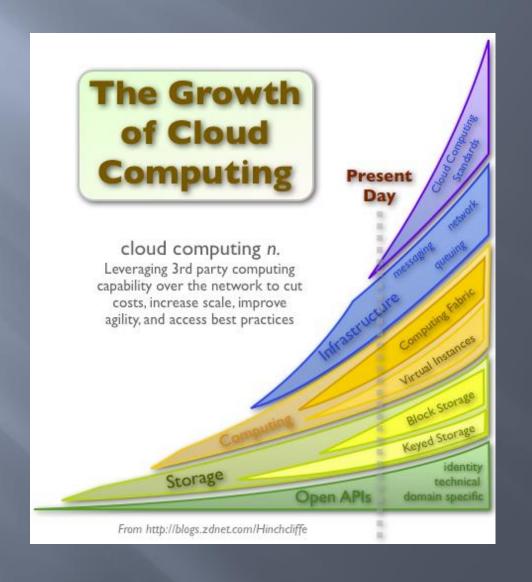








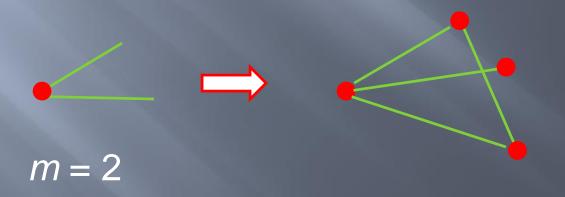




Growth

The simplest case is if we add nodes one by one and assume that all of them bring in *m* links.

The process needs a "seed", i.e., an initial small set of nodes linked together.



Next question: how to attach?

Preferential attachment

The spirit of ER or WS: attach entirely randomly!

Wrong! It does not lead to the required broad degree distribution.

Attachment is not purely random. There are hubs indicating that the Matthew effect is in play

Combination of

- Growth
- Preferential attachment

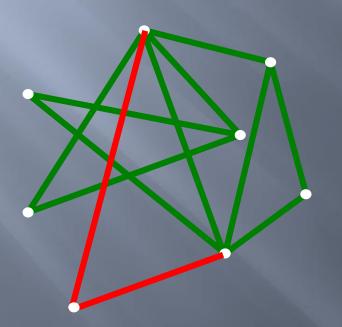
New nodes prefer to link to highly connected nodes.







Réka Albert



PREFERENTIAL ATTACHMENT:

The probability that a node connects to a node with *k* links is proportional to *k*.

$$\Pi(i) = \frac{k_i}{\sum_j k_j}$$

Normalization

(1) Networks continuously expand by the addition of new nodes

WWW: addition of new documents **GROWTH**

(2) New nodes prefer to link to highly connected nodes.

WWW: linking to well known sites

PREFERENTIAL ATTACHMENT

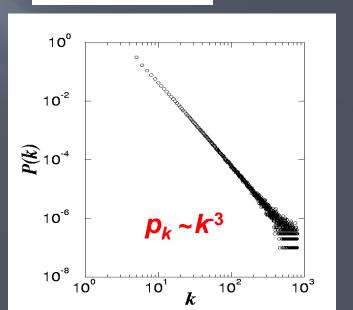
What comes out?

Most interesting: Degree distribution

Power law distribution of degrees

y=3 independent of m

$$\Pi(k_i) = \frac{k_i}{\sum_j k_j}$$



Power law out of preferential attachment is not that new:

- Gyorgy Polya (1887-1985) 1923: *Polya process* in the mathematics literature George Udmy Yule (1871-1951) in 1925: the number of species per genus of flowering plants; *Yule process* in statistics
- Robert Gibrat (1904-1980), 1931: rule of proportional growth independent of system size. *Gibrat process* in economics
- George Kinsley Zipf (1902-1950), 1949: the distribution of wealth in the society. Herbert Alexander Simon (1916-2001), 1955, the distribution of city sizes and
- other phenomena
- Derek de Solla Price (1922-1983), 1976, used it to explain the citation statistics of scientific publications, "cumulative advantage"
- Robert Merton (1910-2003), 1968: Matthew effect,

B A: simple network model

+ the ubiquity of preferential attachment for networks resulted in a radically new approach to modelling them

Derivation [N(t) = t]; one node per time step]:

$$< N(k,t) > = tp_k(t)$$

 $< N(k,t) >= tp_{\nu}(t)$ Number of nodes with degree k at time t.

$$\Pi(k) = \frac{k}{\sum_{j} k_{j}} = \frac{k}{2mt}$$

 $\Pi(k) = \frac{k}{\sum_{i} k_{i}} = \frac{k}{2mt}$ 2m: each node adds m links, but each link contributs to the degree of 2 nodes

Number of links added to degree *k* nodes after the arrival of a new node: $\frac{k}{2mt} \times tp_k(t) \times m = \frac{k}{2} p_k(t)$

of degree k-1 nodes that acquire a new link, becoming

$$\frac{k-1}{2} p_{k-1}(t)$$

degree k $\frac{k-1}{2}p_{k-1}(t)$ # of degree k nodes that acquire a new link, becoming degree k+1 k

Preferential

attachment

$$(t+1)p_k(t+1) = tp_k(t) + \frac{k-1}{2}p_{k-1}(t) - \frac{k}{2}p_k(t)$$

New node adds m new links

Gain of k-

nodes via $k-1 \rightarrow k$

nodes via

k-nodes at time t+1

k-nodes at time t

Loss of k $k \rightarrow k+1$

Total number of

k-nodes

$$(t+1)p_k(t+1) = tp_k(t) + \frac{k-1}{2}p_{k-1}(t) - \frac{k}{2}p_k(t)$$

No nodes with degree < m. We need a separate eq. for that case:

$$(t+1)p_m(t+1) = tp_m(t) + 1 - \frac{m}{2}p_m(t)$$
The just arriving new node

We are interested in the long time, stationary solution:

$$\lim_{t\to\infty}p_k(t)=p_k$$

$$(t+1)p_k(t+1) = tp_k(t) + \frac{k-1}{2}p_{k-1}(t) - \frac{k}{2}p_k(t) \quad k > m$$

$$(t+1)p_m(t+1) = tp_m(t) + 1 - \frac{m}{2}p_m(t)$$

Stationary equations with

$$\lim_{t\to\infty}p_k(t)=p_k$$

$$p_k = \frac{k-1}{2} p_{k-1} - \frac{k}{2} p_k \quad k > m$$

$$p_m = 1 - \frac{m}{2} p_m$$

$$p_k = \frac{k-1}{k+2} p_{k-1} \quad k > m$$

$$p_m = \frac{2}{m+2}$$

$$p_k = \frac{k-1}{k+2} \, p_{k-1}$$

$$p_k = \frac{k-1}{k+2} p_{k-1}$$
 \Rightarrow $p_{k+1} = \frac{k}{k+3} p_k$

$$p_m = \frac{2}{m+2}$$

"Initial condition"

$$p_{m+1} = \frac{m}{m+3} p_m = \frac{2m}{(m+2)(m+3)} = \frac{2m(m+1)}{(m+1)(m+2)(m+3)}$$

$$p_{m+2} = \frac{m+1}{m+4} p_{m+1} = \frac{2m(m+1)}{(m+2)(m+3)(m+4)}$$

$$p_{m+3} = \frac{m+2}{m+5} p_{m+2} = \frac{2m(m+1)}{(m+3)(m+4)(m+5)}$$

$$p_k = \frac{2m(m+1)}{k(k+1)(k+2)}$$

For large *k*:

 $P(k) \sim k^{-3}$ Power law tail

A simple route

Start from eq.
$$P(k) = \frac{k-1}{2}P(k-1) - \frac{k}{2}P(k)$$

$$2P(k) = (k-1)P(k-1) - kP(k) = -P(k-1) - k[P(k) - P(k-1)]$$

Let's take a continuum limit

$$2P(k) = -P(k-1) - k\frac{P(k) - P(k-1)}{k - (k-1)} = -P(k-1) - k\frac{dP(k)}{dk}$$

$$P(k) = -\frac{1}{2} \frac{d[kP(k)]}{dk}$$

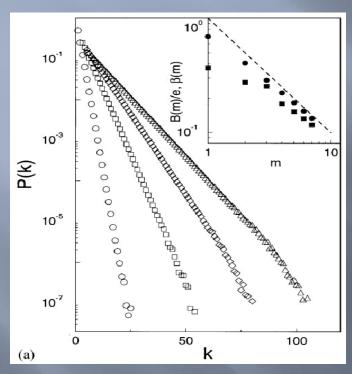
$$P(k) = -\frac{1}{2}P(k) - \frac{1}{2}k\frac{dP(k)}{dk} \qquad \frac{3}{2}P(k) = -\frac{1}{2}k\frac{dP(k)}{dk}$$

$$P(k) = Ak^{-3}$$

because

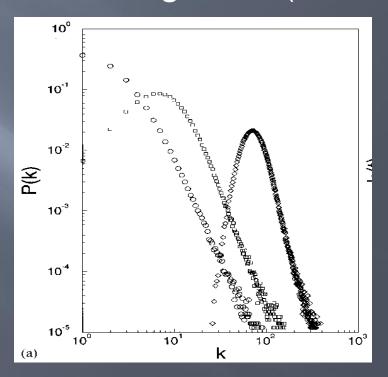
$$y = ax \frac{dy}{dx} \Longrightarrow y = Cx^{1/a}$$

Growth without P.A.



Exponential p_k

P.A. without growth (*N* fixed)



Power law only at the beginning, then sharply peaked p_k finally complete graph. No stationary sol'n

	Lattice	ER	WS	BA
<d></d>	~ L	~ In N	~ In N	
С	const	<k>/N</k>	const	
p_k	δ(k,k ₀)	Poisson	shifted Poisson ≅੍ਹ	~ <i>k</i> -Y

Many questions:

Average distance <*d*>?

Clustering C?

What about exponents other than 3?

Origin of preferential attachment (global rule)?

	const.	$\gamma = 2$
Ultra Small World	ln ln N	$2 < \gamma < 3$
$\langle d \rangle$ ~ <	$\ln(\gamma - 1)$ $\ln N$	2 < y < 3
$\langle \alpha \rangle$	ln N	$\gamma = 3$
	ln ln N	$\gamma = 3$
Small World	ln <i>N</i>	$\gamma > 3$

Size of the biggest hub O(N)

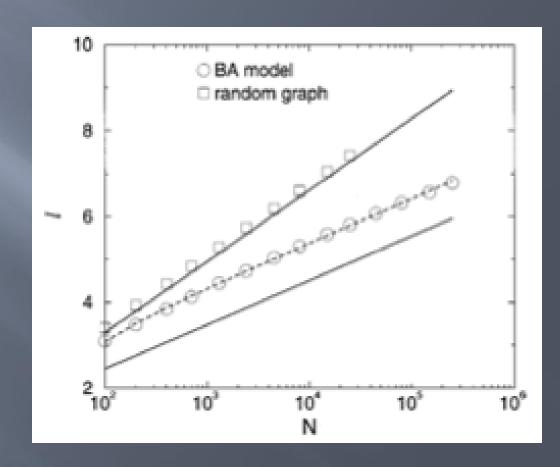
Av. distance increases slower than log

BA: Boarderline case: slightly slower than log

Finite second moment, result like in ER.

Config. model results but broader validity

$$\langle d \rangle \approx \frac{\ln N}{\ln \ln N}$$

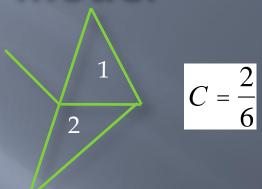


	Lattice	ER	WS	BA
<d>></d>	~ L	~ In <i>N</i>	~ In N	~ InN/ InInN
С	const	<k>/N</k>	const	
p_k	$\delta(k,k_c)$	Poisson	shifted Poisson ≒	~ <i>k</i> - ^v

BA model is a small world. The mechanism is through the hubs! (C.f. Milgram experiment!)

Clustering Coefficient in the BA model

$$C_i = \frac{N_i^{\Delta}}{k_i(k_i - 1)/2}$$



Denote the probability to have a link between node i and j with P(i,j) The probability that three nodes i,j,l form a triangle is P(i,j)P(i,l)P(j,l). The expected number of triangles, in which a node l with degree k_l participates is thus:

$$N_l^{\Delta} = \int_{i=1}^N di \int_{j=1}^N dj P(i,j) P(i,l) P(j,l)$$

We need to calculate P(i,j).

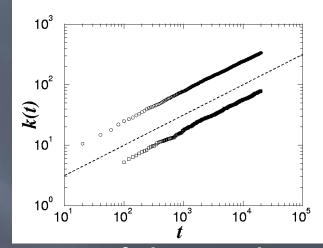
Time evolution of k

Let us describe the time evolution of degrees

$$k_i(t+1) = k_i(t) + m\frac{k_i}{\sum_j k_j} \Rightarrow \frac{dk_i}{dt} = m\frac{k_i}{\sum_j k_j} = \frac{k_i}{2t}$$

$$\frac{dk_i}{dt} = \frac{k_i}{2t}$$

$$k_i(t) = m\sqrt{\frac{t}{t_i}} \sim t^{\beta} \quad \text{with } \beta = 1/2$$



This tells us that what matters is the age of the nodes. Old nodes will always have advantage!

Calculate P(i,j)

Node *j* arrives at time $t_i = j$ and the probability that it will link to node i with degree k_i already in the network is determined by preferential attachment:

time
$$j$$

$$P(i, j) = m\Pi(k_i(j)) = m\frac{k_i(j)}{\sum_{l=1}^{j} k_l} = m\frac{k_i(j)}{2mj}$$

$$k_i(t) = m \left(\frac{t}{t_i}\right)^{1/2} = m \left(\frac{j}{i}\right)^{1/2}$$

 $P(i,j) = \frac{m}{2} (ij)^{-\frac{1}{2}}$ Where we used that the arrival time of node j is t=jand the arrival time of node is $t_i = i$

$$N_{i}^{\Delta} = \int_{i=1}^{N} di \int_{j=1}^{N} dj P(i,j) P(i,l) P(j,l) = \frac{m^{3}}{8} \int_{i=1}^{N} di \int_{j=1}^{N} dj (ij)^{-\frac{1}{2}} (il)^{-\frac{1}{2}} (jl)^{-\frac{1}{2}} = \frac{m^{3}}{8l} \int_{i=1}^{N} \frac{di}{i} \int_{j=1}^{N} \frac{dj}{j} = \frac{m^{3}}{8l} (\ln N)^{2}$$

$$C = \frac{\frac{m^3}{8l} (\ln N)^2}{k_l (k_l - 1)/2}$$

$$C = \frac{m}{8} \frac{(\ln N)^2}{N}$$

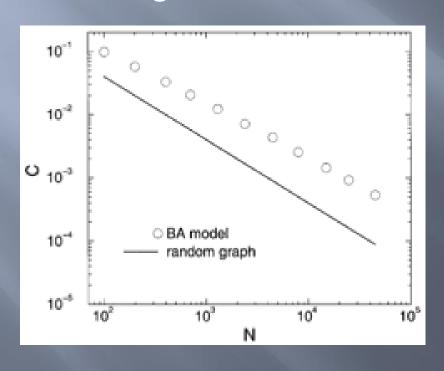
 $C = \frac{\frac{m^3}{8l} (\ln N)^2}{k_l(k_l - 1)/2}$ $k_l(t) = m \left(\frac{N}{l}\right)^{1/2}$ of node *i* at current time, at time *t=N* Which is the degree

Let us approximate:

$$k_l(k_l-1) \approx k_l^2 = m^2 \frac{N}{l}$$

Clustering coefficient:

For ER we have $C = p = \langle k \rangle / N$ Decreasing with size. BA:



Bad news!

$$C = \frac{m}{8} \frac{(\ln N)^2}{N}$$

	Lattice	ER	WS	BA
<d></d>	~ L	~ In <i>N</i>	~ In N	~ In/V /InIn/V
С	const	<k>/N</k>	const	~(ln/V) ² //V
p_k	$\delta(k,k_c)$	Poisson	shifted Poisson	~ <i>k</i> -Y

Shall we throw BA away? No!

It is an important aspect to capture the mechanism by which complex networks emerge. A new approach, a starting point.

Barabási-Albert (BA) model

Summary of the BA model:

• Nr. of links:
$$L = m t$$

•Average degree:
$$\langle k \rangle = \frac{2L}{N} \rightarrow 2m$$

• Degree dynamics
$$k_i(t) = m \left(\frac{t}{t_i}\right)^{\beta}$$
 $\beta = \frac{1}{2}$

•Degree distribution:
$$P(k) \sim k^{-g}$$
 $g = 3$

•Average Path Length:
$$l \approx \frac{\ln N}{\ln \ln N}$$

•Clustering Coefficient:
$$C \sim \frac{(\ln N)^2}{N}$$

β: dynamical exponent

γ: degree exponent

The network grows, but the degree distribution becomes stationary.

N = t

How to cure the clustering problem?

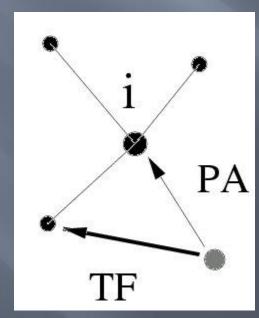
Something is missing from the attachment mechanism

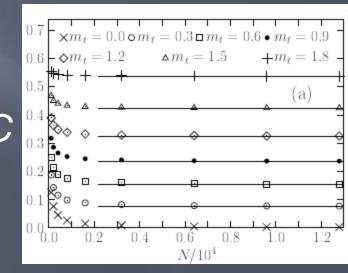
Besides preferential attachment (hunting for popular nodes), there is an additional process: Friends of friends get easily friends. This should be incorporated.

PA: Preferencial attachment

TF: triad formation

First link: PA then TF with prob. *p* PA with 1-p



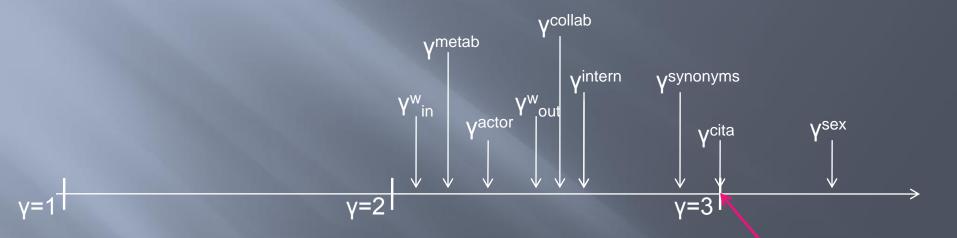


This was easy! (Much easier than the effort with the configuration model!)

	Lattice	ER	WS	BA+
<d>></d>	~ L	~ In N	~ In N	~ In/V /InIn/V
С	const	<k>/N</k>	const	const.
p_k	$\delta(k,k_c)$	Poisson	shifted Poisson	~k-Y

It is worth concentrating on the mechanisms!

There is no strict universality: empirical exponents are system dependent and not 3!



Can we produce different exponents with slight modification of BA, keeping the concept?

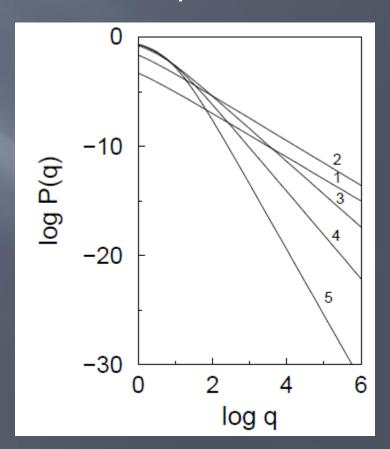
BA

Yes!

Initial attractiveness : $\Pi(k) \sim A + (k-m) = A + q$

 \rightarrow P(k) \sim k^{- γ} where γ =2 + A/m

Tunable exponent between 2 and ∞



A/m=0.001; 0.05; 1; 2; 4

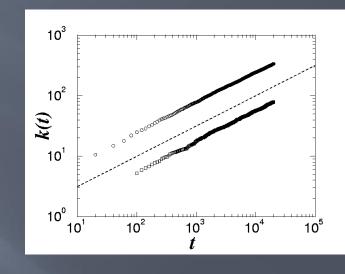
We saw k(t)

Let us describe the time evolution of degrees

$$\frac{\partial k_i}{\partial t} \propto \Pi(k_i) = A \frac{k_i}{\sum_j k_j} = \frac{k_i}{2t}$$

During a unit time (time step): $\Delta k = m \rightarrow A = m$

$$k_i(t) = m\sqrt{\frac{t}{t_i}} \sim t^{\beta}$$
 with $\beta = 1/2$

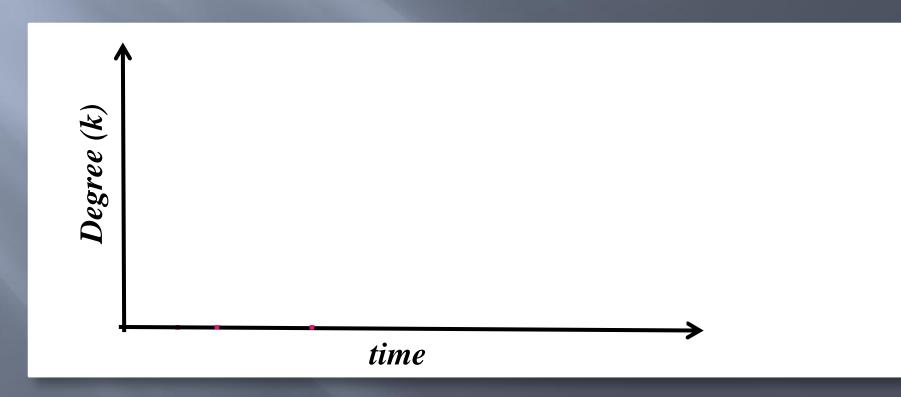


This tells us that what matters is the age of the nodes. Old nodes will always have advantage!

Further question: Can new ones make it? (Google!)

Fitness Model: Can Latecomers Make It?

BA model: $k(t) \sim t^{1/2}$ (first mover advantage)



Fitness model: fitness (η)

$$\Pi(k_i) = \frac{\eta_i k_i}{\sum_j \eta_j k_j}$$

$$k(\eta,t)\sim t^{\beta(\eta)}$$

$$\beta(\eta) = \eta/C$$



A key element in the BA model (and related ones) is that the new nodes attach to the old ones via preferential attachment.

We have seen that this is quite common assumption – but it is strange!

Imagine, you are making a new www site and put appropriate links into it. Do you have to search through the 10¹⁰ web sites and make a statistics to calculate the probability of linking to one of them?!

No way!

In reality we cannot test a huge network. The decision is local – but what comes out *is* preferential attachment. (Self organization – invisible hand...)

There could be other mechanisms. It is a legitimate question to ask: Does a system obey PA?

We can measure this!

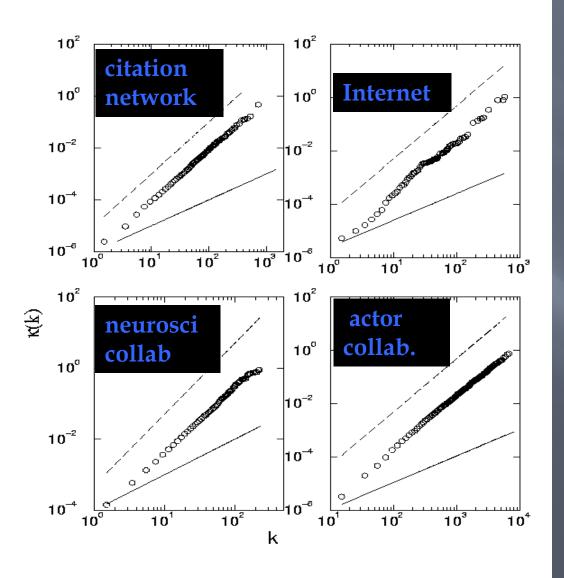
$$\frac{\partial k_i}{\partial t} \propto \Pi(k_i) \sim \frac{\Delta k_i}{\Delta t}$$

Plot the change in the degree Δk during a fixed time Δt for nodes with degree k, and you get $\sim \Pi(k)$

To reduce noise, plot the sum of $\Pi(k)$ over k:

$$\Pi_{<}(k) = \sum_{k' < k} \Pi(k')$$

One has to measure the cumulative increment of the nodes with degree smaller than *k*



Plots of the cumulative

$$\Pi_{<}(k) = \sum_{k' < k} \Pi(k')$$

No pref. attach:
$$\kappa \sim k$$

Linear pref. attach:
$$\kappa \sim k^2$$

What could be the local mechanism leading the PA?

- Copying mechanism
 directed network
 select a node and an edge of this node
 attach to the endpoint of this edge
- 2. Walking on a network directed network the new node connects to a node, then to every first, second, ... neighbor of this node
- 3. Attaching to edges select an edge attach to both endpoints of this edge (clustering!)
- Node duplication
 duplicate a node with all its edges
 randomly prune edges of new node

Consider the citation network.

Nodes: Papers

Directed: Out degrees - references cited

In degrees – citing papers

In an ideal case those papers are cited, which have been read by the authors and have impact on the results.

Authors are often sloppy: They simply copy the reference list of other papers. (This is evidenced by propagation of typos.)

Similar mechanism may work in other networks too.

To make it more realistic: Only a fraction of an old paper's refs is copied and the remaining is filled in with others selected at random. (For simplicity we assume that all bibliographies have the same size, c.)

Old prob.
$$\gamma$$
1) aaa $\xrightarrow{\text{prob. } \gamma}$ 1) aaa (copied)
2) bbb $\xrightarrow{\text{prob. } 1-\gamma}$ 2) bla (randomly chosen)

c)
$$zzz \longrightarrow c) zzz$$

As a result, we will have on the average $c\gamma$ copied items and $c(1 - \gamma)$ ones selected at random.

Starting set: E.g., n_0 vertices interconnected randomly such that everybody has c out links (no mulitlinks allowed).

Q: What is the in degree distribution?

Node *i* can get a link in two ways: a) part of a list and copied b) randomly selected. The number of nodes at time *t* is *n*.

- a) Node *i* has indegree k^{in}_{i} . The prob. of choosing *i* is thus $\gamma k^{in}_{i} / n$. (Case a).
- b) $c(1-\gamma)/n$.

$$\frac{\gamma k_i^{in}}{n} + \frac{(1-\gamma)c}{n} = \frac{\gamma k_i^{in} + (1-\gamma)c}{n}$$

The expected number of nodes with in degree *k* receiving a new link is

$$np_k \frac{\gamma k + (1 - \gamma)c}{n} =$$

$$p_k [\gamma k + (1 - \gamma)c] = \frac{c(k + a)}{c + a} p_k$$
with $a = c(\frac{1}{\gamma} - 1)$

which can be rewritten as

resulting in a rate equation:

$$np_k(n) + \frac{c(k-1+a)}{c+a} p_{k-1}(n) - \frac{c(k+a)}{c+a} p_k(n)$$

$$np_k(n) + \frac{c(k-1+a)}{c+a}p_{k-1}(n) - \frac{c(k+a)}{c+a}p_k(n)$$

$$(n+1)p_0(n+1) = np_0(n) + 1 - \frac{ca}{c+a}p_0(n)$$

k=0

For $n \rightarrow \infty$ oo stationary solution:

$$p_k = \frac{c}{c+a} [(k-1+a)p_{k-1}(n) - (k+a)p_k(n)]$$

$$p_0 = 1 - \frac{ca}{c+a} p_0$$

k=0

$$p_0 = \frac{1 + a/c}{a + 1 + a/c}$$

For every k iterative sol'n. The asymptotics is: $p_k \sim k^{-\alpha}$

th $\alpha = 2 + a/c = 1 + 1/\gamma$

Tunable exponent

For more details see Newman's book

Summary of models

Linear growth, linear pref. attachment

Nonlinear preferential attachment $\Pi(k_i) \sim k_i^{\alpha}$

Asymptotically linear pref. attachment

$$\Pi(k_i) \sim a_{\infty} k_i$$
 as $k_i \rightarrow \infty$

Initial attractiveness

$$\Pi(k_i)\sim A+k_i$$

Accelerating growth $\langle k \rangle \sim t^{\theta}$ constant initial attractiveness

Internal edges with probab. p

Rewiring of edges with probab. q

c internal edges or removal of c edges

Gradual aging $\Pi(k_i) \sim k_i (t - t_i)^{-\nu}$

Multiplicative node fitness

$$\Pi_i \sim \eta_i k_i$$

Edge inheritance

Copying with probab. p

Redirection with probab. r

Walking with probab. p

Attaching to edges

p directed internal edges $\Pi(k_i, k_i) \propto (k_i^{in} + \lambda)(k_i^{out} + \mu)$ $\gamma = 3$

no scaling for $\alpha \neq 1$

$$\gamma \rightarrow 2$$
 if $a_{\infty} \rightarrow \infty$

$$\gamma \rightarrow \infty$$
 if $a_\infty \rightarrow 0$

$$\gamma = 2$$
 if $A = 0$

$$\gamma \rightarrow \infty$$
 if $A \rightarrow \infty$

$$\gamma = 1.5 \text{ if } \theta \rightarrow 1$$

$$\gamma \rightarrow 2 \text{ if } \theta \rightarrow 0$$

$$\gamma = 2 \text{ if}$$

$$q = \frac{1 - p + m}{1 + 2m}$$

$$\gamma \rightarrow \infty$$
 if $p,q,m \rightarrow 0$

$$\gamma \rightarrow 2$$
 if $c \rightarrow \infty$

$$\gamma \rightarrow \infty$$
 if $c \rightarrow -1$

$$\gamma \rightarrow 2$$
 if $\nu \rightarrow -\infty$

$$\gamma \rightarrow \infty$$
 if $\nu \rightarrow 1$

$$P(k) \sim \frac{k^{-1-C}}{\ln(k)}$$

$$P(k_{in}) = \frac{d}{k_{in}^{\sqrt{2}}} \ln(ak_{in})$$

$$\gamma = (2-p)/(1-p)$$

$$y = 1 + 1/r$$

$$\gamma \simeq 2$$
 for $p > p_c$

$$\gamma = 3$$

$$\gamma_{in} = 2 + p\lambda$$

 $\gamma_{out} = 1 + (1 - p)^{-1} + \mu p/(1 - p)$

Barabási and Albert, 1999

Krapivsky, Redner, and Leyvraz, 2000

Krapivsky, Redner, and Leyvraz, 2000

Dorogovtsev, Mendes, and Samukhin, 2000a, 2000b

Dorogovtsev and Mendes, 2001a

Albert and Barabási, 2000

Dorogovtsev and Mendes, 2000c

Dorogovtsev and Mendes, 2000b

Bianconi and Barabási, 2001a Dorogovtsev, Mendes, and Samukhin, 2000c

Kumar et al., 2000a, 2000b

Krapivsky and Redner, 2001

Vázquez, 2000

Dorogovtsev, Mendes, and Samukhin, 2001a

Krapivsky, Rodgers, and Redner, 2001

Homework

Generate graphs with nonlinear preferential attachment:

$$\Pi(i) = \frac{k_i^{\beta}}{\sum_j k_j^{\beta}}$$

Calculate the degree distribution and try to find a characteristic degree as a function of β . (Chose β values not far from 1.)