



the  
**abdus salam**  
international centre for theoretical physics

SMR/1423 - 19

SCHOOL ON  
"STATISTICAL PHYSICS, PROBABILITY THEORY AND  
COMPUTATIONAL COMPLEXITY"

(26 August - 4 September 2002)

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" Internet Graphs and their Generation "

presented by:

**A.L. Barabasi**  
University of Notre Dame  
U.S.A.



# The Last Banda

Gift of New Advent is called *Re: libe*  
Give it a try now. **Soon it's a book**

re:libe

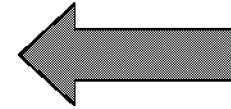


Austin Powers:  
The spy who  
shagged me

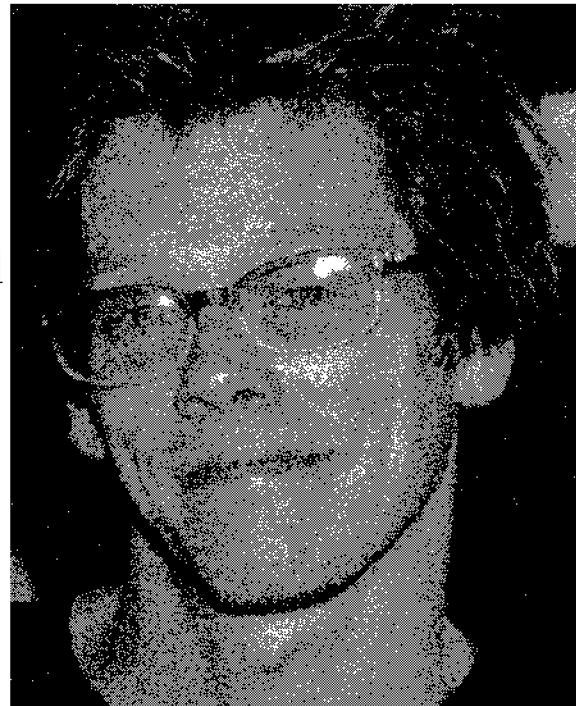
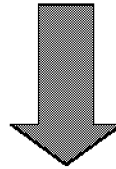


**Robert Wagner**

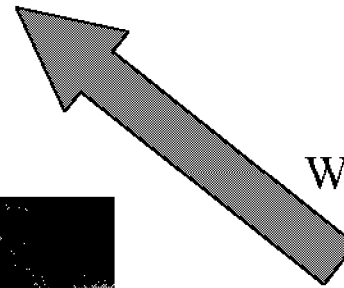
Let's make  
it legal



Wild Things

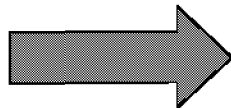


What Price Glory

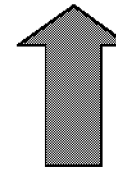


**Barry Norton**

A Few Good  
Man



Monsieur  
Verdoux



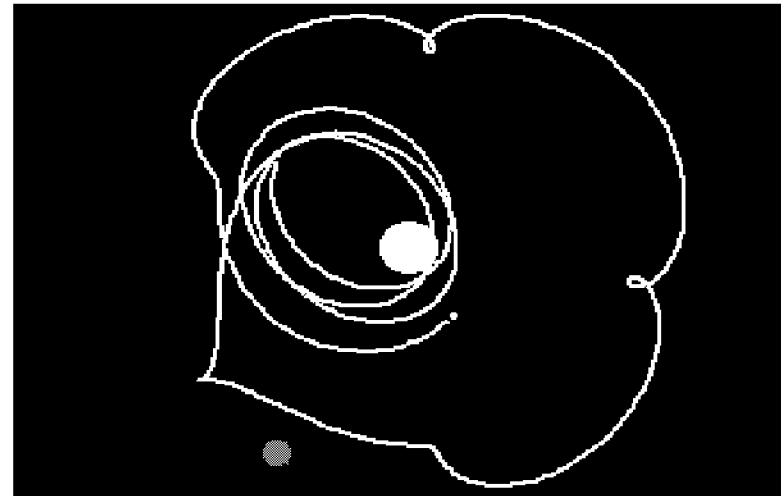
# What is Complexity?

A popular paradigm: Simple systems display complex behavior

- non-linear systems
- chaos
- fractals

## 3 Body Problem

Earth (☾) Jupiter (♃) Sun (☉)



Main Entry: <sup>1</sup>com·plex

Function: *noun*

Etymology: Late Latin *complexus* totality, from Latin, embrace, from *complecti*

Date: 1643

**1 : a whole made up of complicated or interrelated parts**

# Society

**Nodes**: individuals

**Links**: social relationship  
(family/work/friendship/etc.)



S. Milgram (1967)

John Guare

**Six Degrees of Separation**

Social networks: Many individuals with diverse social interactions between them.

# Communication networks

The Earth is developing an electronic nervous system, a network with diverse nodes and links are

← -computers

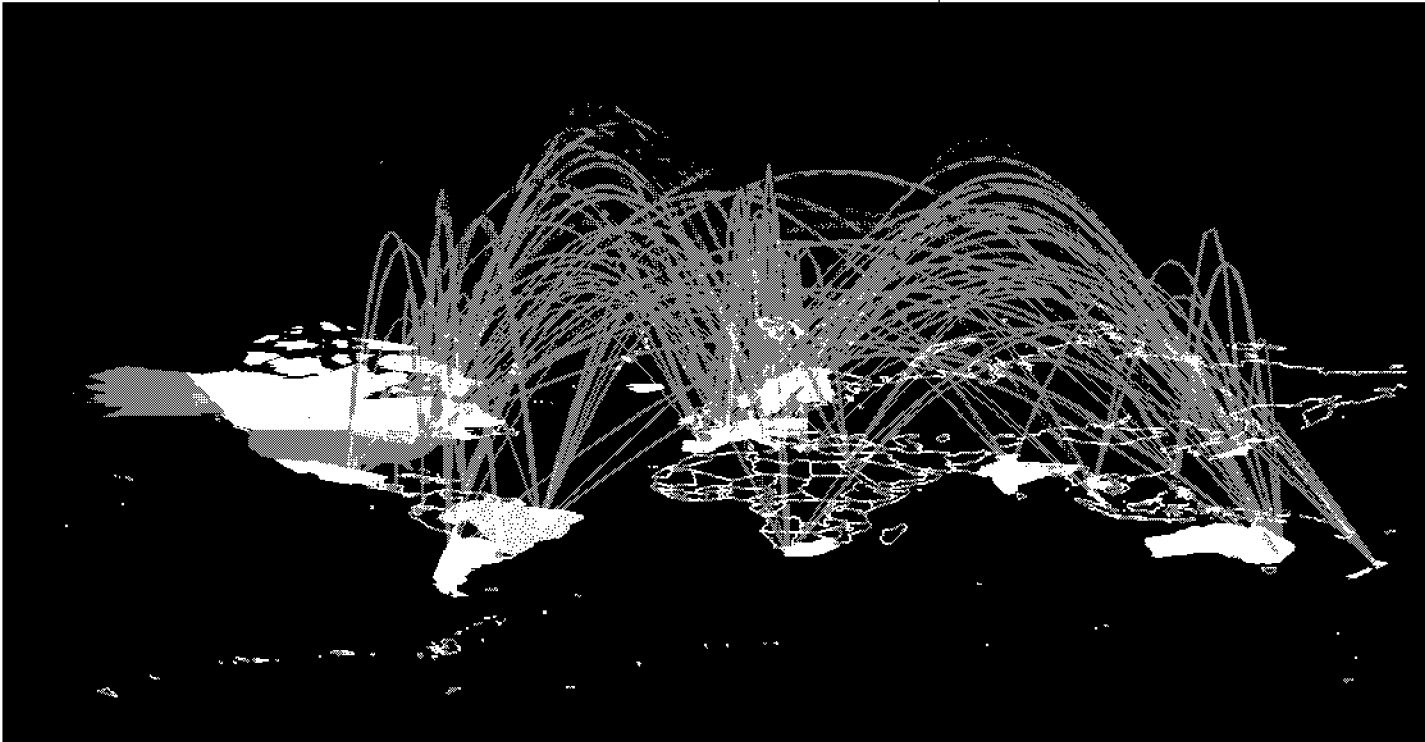
-routers

-satellites

↓ -phone lines

-TV cables

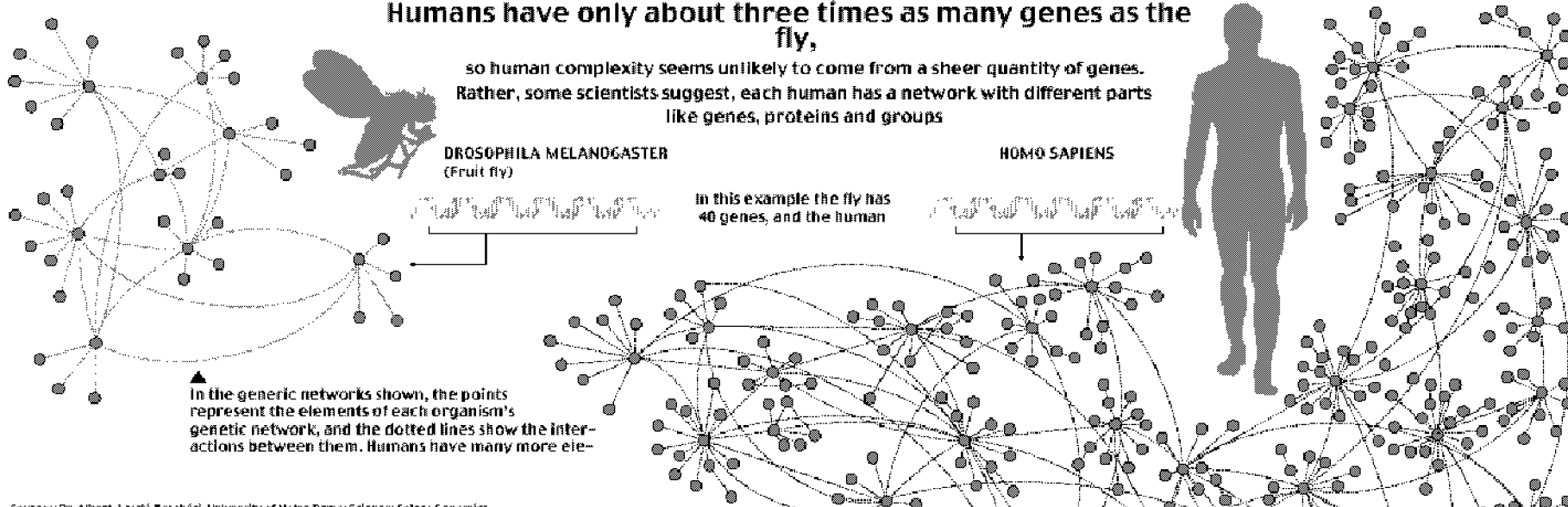
-EM waves



Communication networks: Many non-identical components with diverse connections between them.

Humans have only about three times as many genes as the fly,

so human complexity seems unlikely to come from a sheer quantity of genes. Rather, some scientists suggest, each human has a network with different parts like genes, proteins and groups



In the generic networks shown, the points represent the elements of each organism's genetic network, and the dotted lines show the interactions between them. Humans have many more ele-

Sources: Dr. Albert-László Barabási, University of Notre Dame; Science; Celera Genomics

# Complex systems

Made of many non-identical elements connected by diverse interactions.

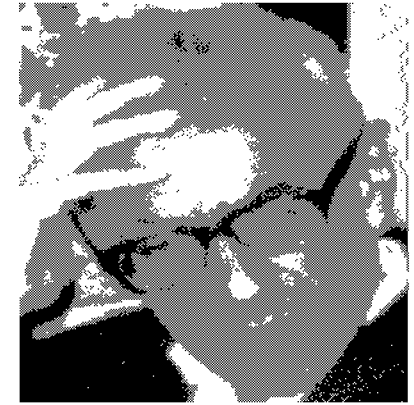


**NETWORK**

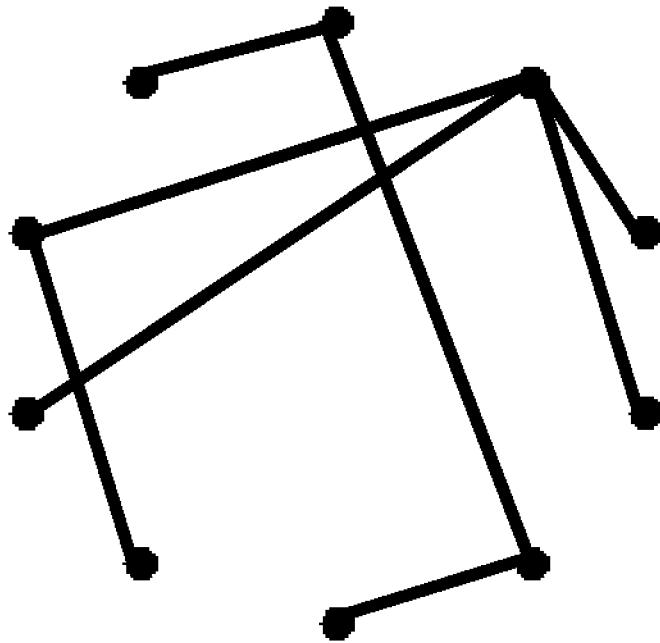
Steve Daines/The New York Times



# Erdős-Rényi model (1960)



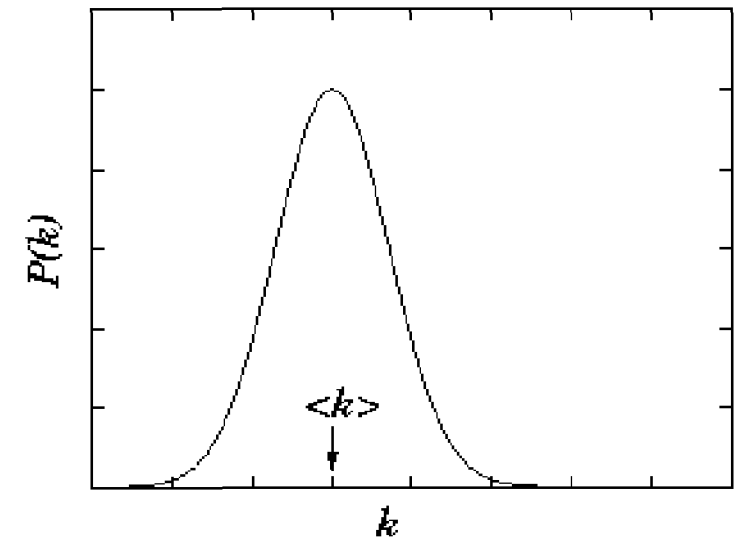
**Pál Erdős**  
**(1913-1996)**



Connect with  
probability  $p$

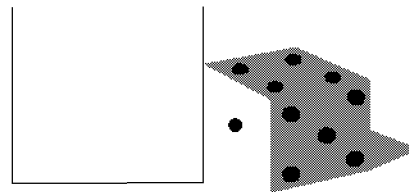
$$p=1/6$$
$$N=10 \langle k \rangle$$
$$\sim 1.5$$

Poisson distribution

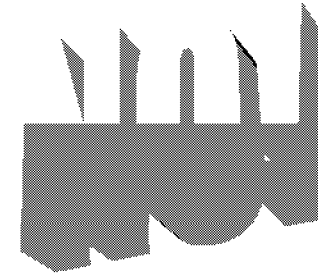


- Democratic

- Random



# ARE COMPLEX NETWORKS REALLY RANDOM?

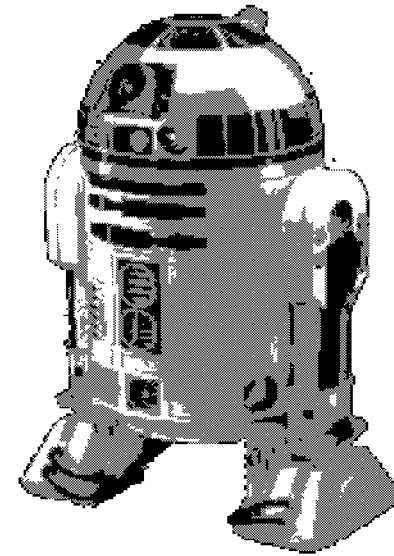
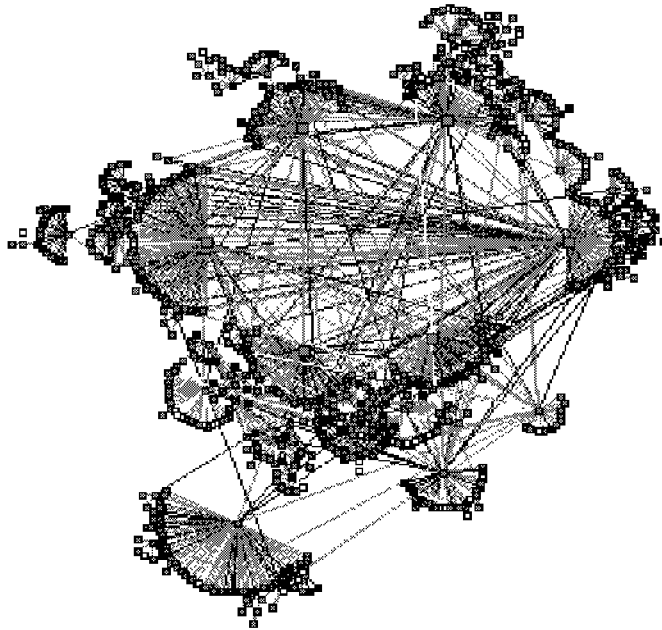


# World Wide Web

**Nodes:** WWW documents

**Links:** URL links

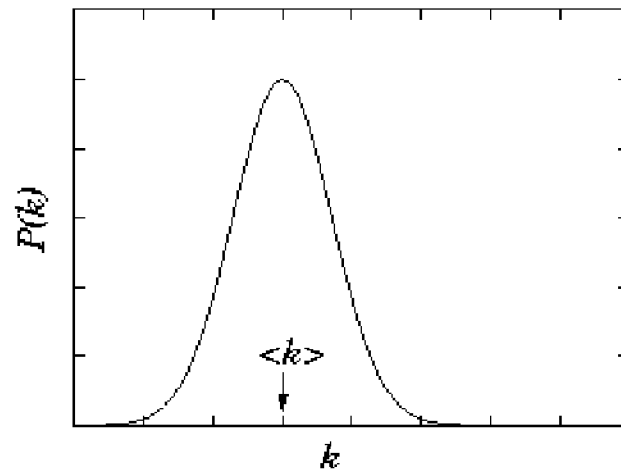
800 million documents (S. Lawrence, 1999)



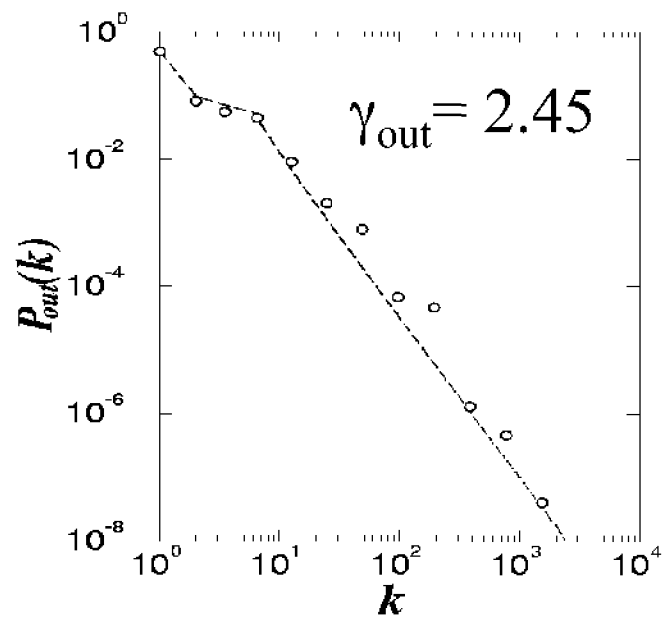
**ROBOT:** collects all URL's found in a document and follows them recursively

R. Albert, H. Jeong, A-L Barabasi, Nature, **401** 130 (1999)

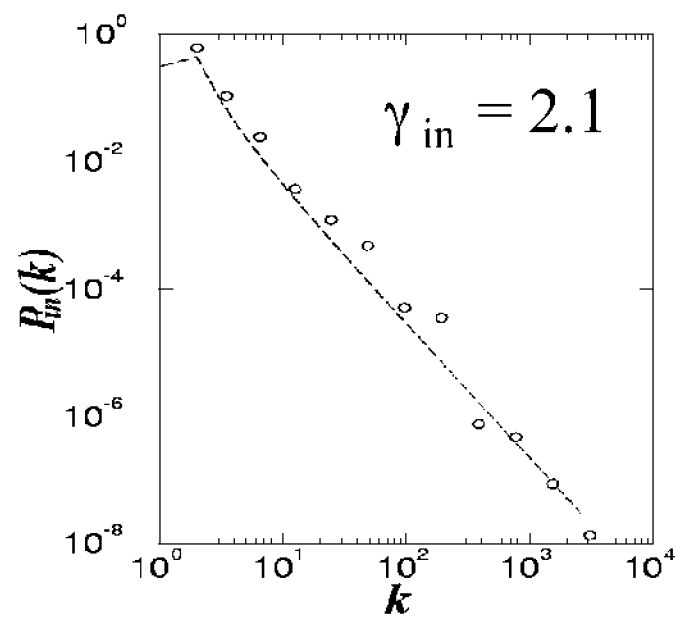
# What did we expect?



## We find:

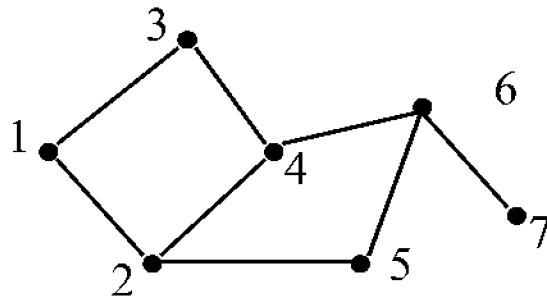


$$P_{out}(k) \sim k^{-\gamma_{out}}$$



$$P_{in}(k) \sim k^{-\gamma_{in}}$$

## 19 degrees of separation



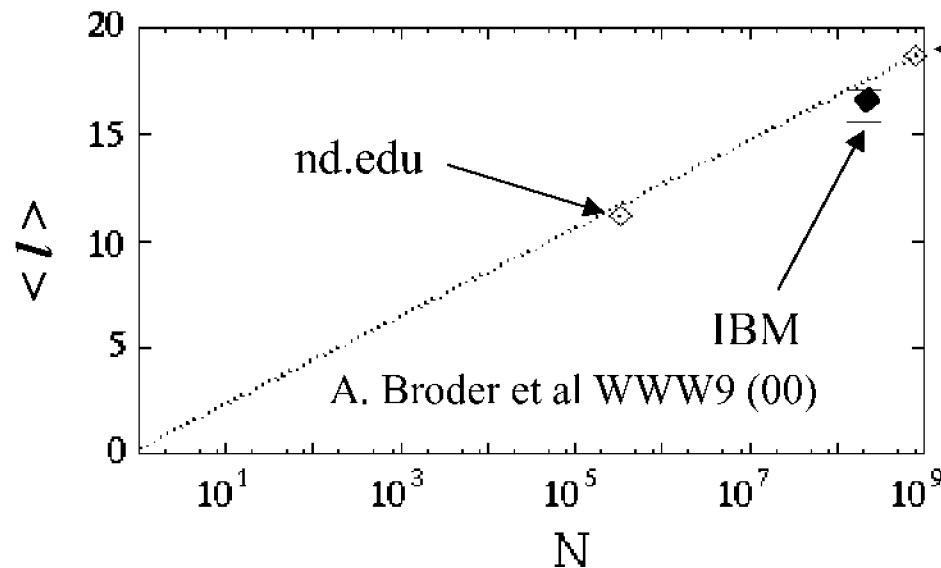
$$l_{15} = 2 [1 \rightarrow 2 \rightarrow 5]$$

$$l_{17} = 4 [1 \rightarrow 3 \rightarrow 4 \rightarrow 6 \rightarrow 7]$$

$$\dots \langle l \rangle = ??$$

- **Finite size scaling:** create a network with  $N$  nodes with  $P_{in}(k)$  and  $P_{out}(k)$

$$\langle l \rangle = 0.35 + 2.06 \log(N)$$



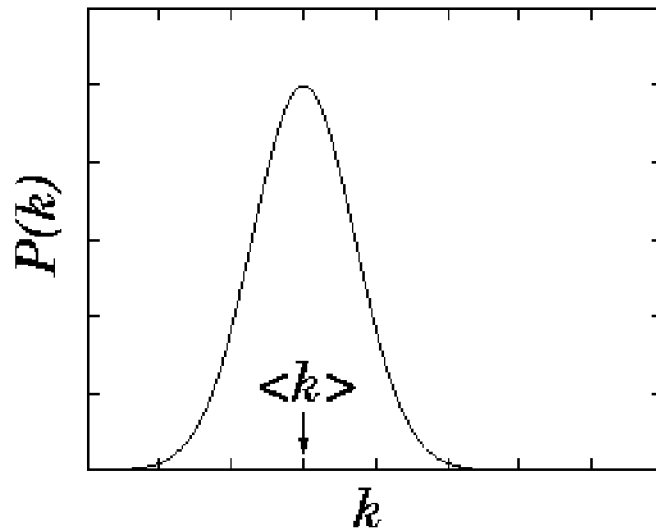
← **19 degrees of separation** R.

Albert et al Nature (99)

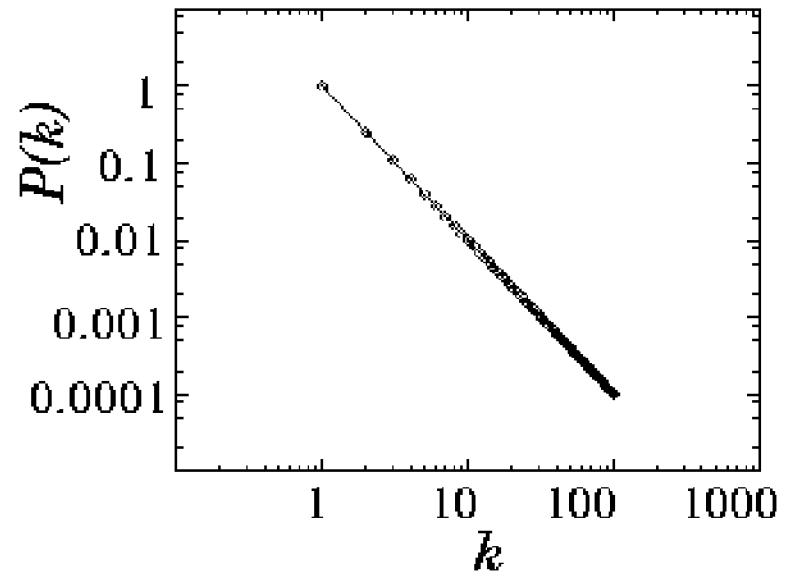
based on 800 million webpages  
[S. Lawrence et al Nature (99)]

# What does it mean?

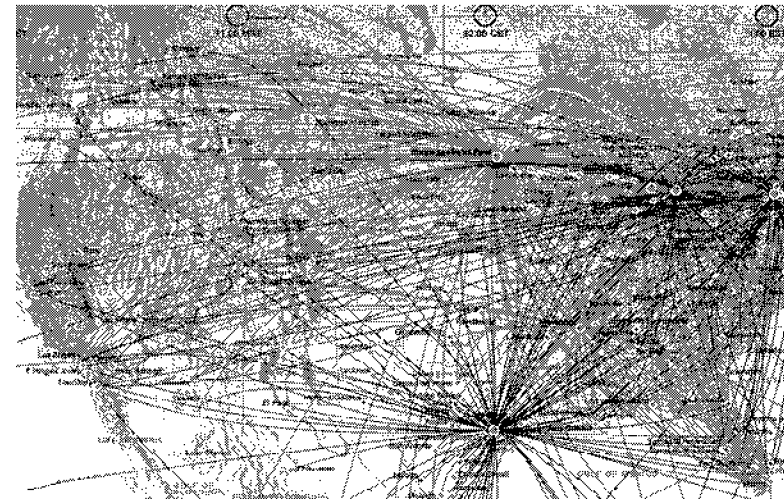
Poisson distribution



Power-law distribution



**Exponential Network**

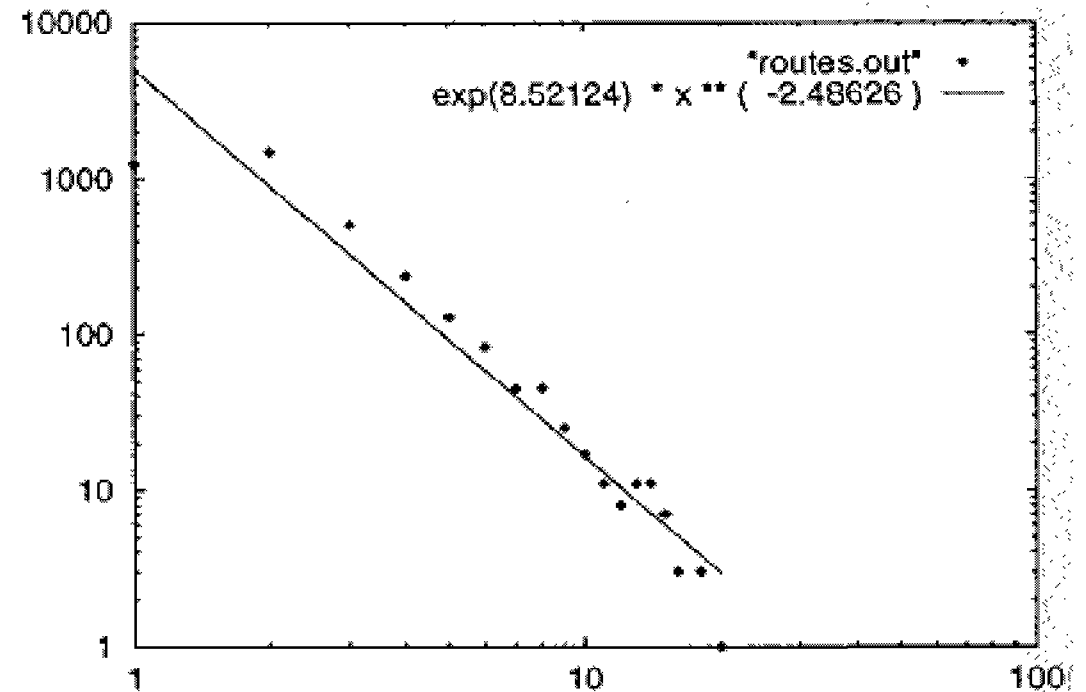
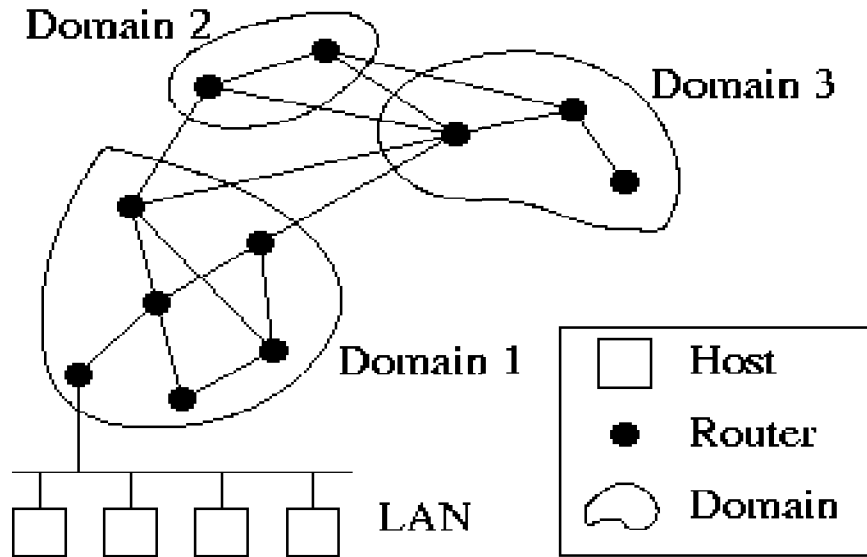


**Scale-free Network**

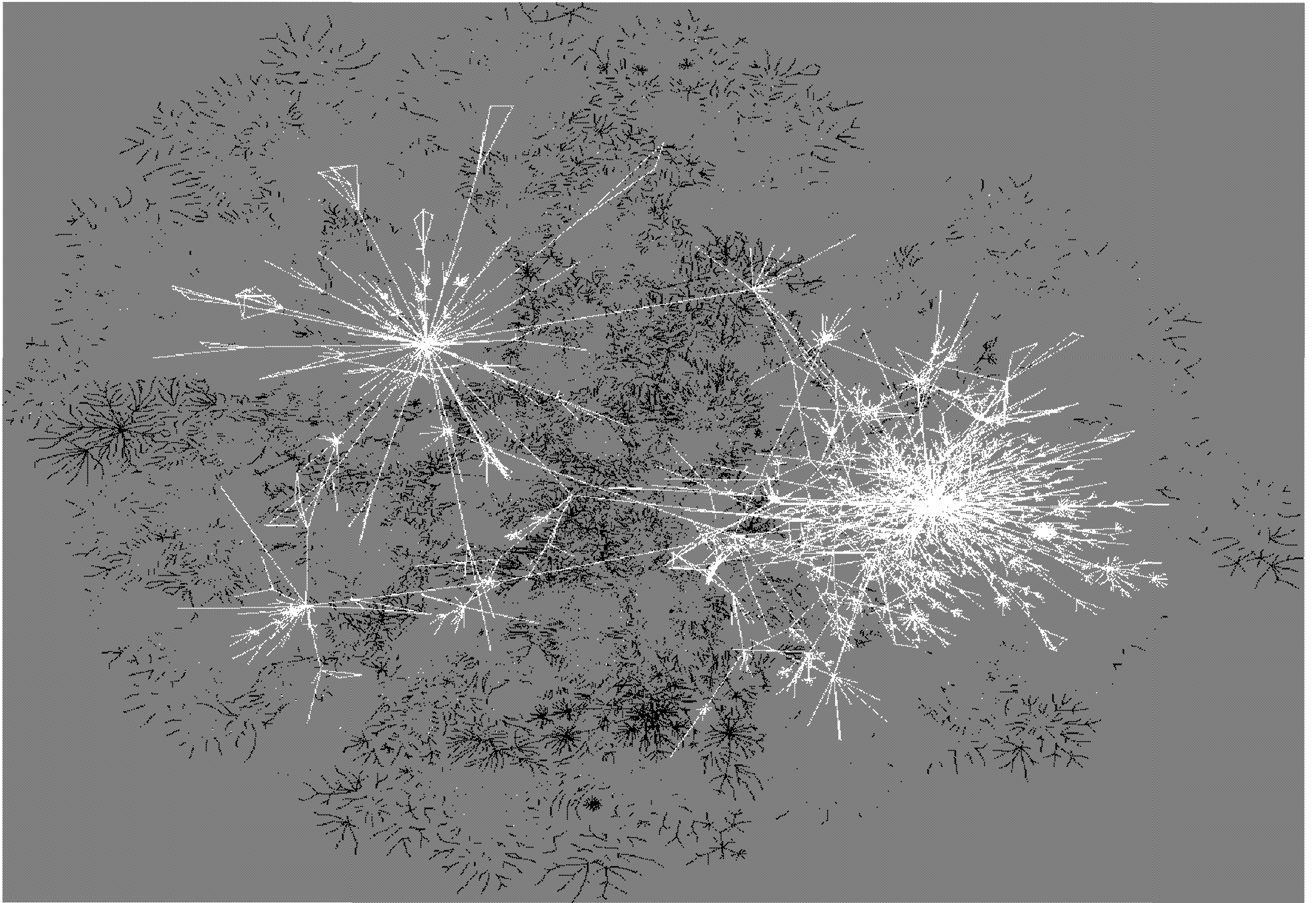
# INTERNET BACKBONE

Nodes: computers, routers

Links: physical lines



(Faloutsos, Faloutsos and Faloutsos, 1999)





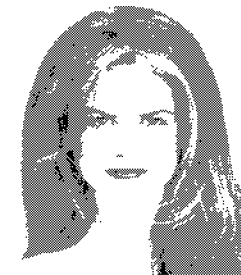
# ACTOR CONNECTIVITIES

Nodes: actors  
cast jointly

Links: **IMDb** Internet Movie Database



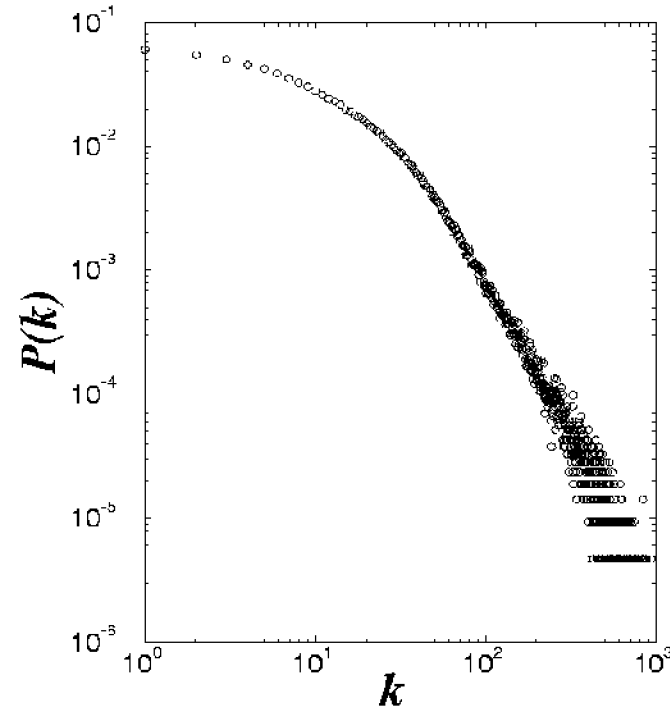
Days of Thunder (1990)  
Far and Away (1992)  
Eyes Wide Shut (1999)



$N = 212,250$  actors  
 $\langle k \rangle = 28.78$

$P(k) \sim k^{-\gamma}$

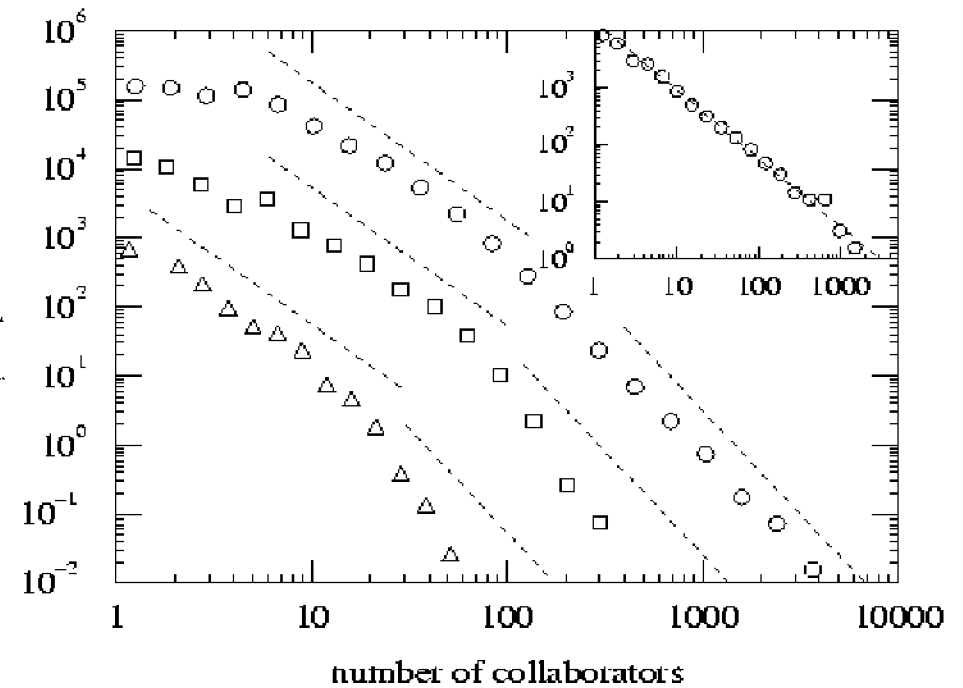
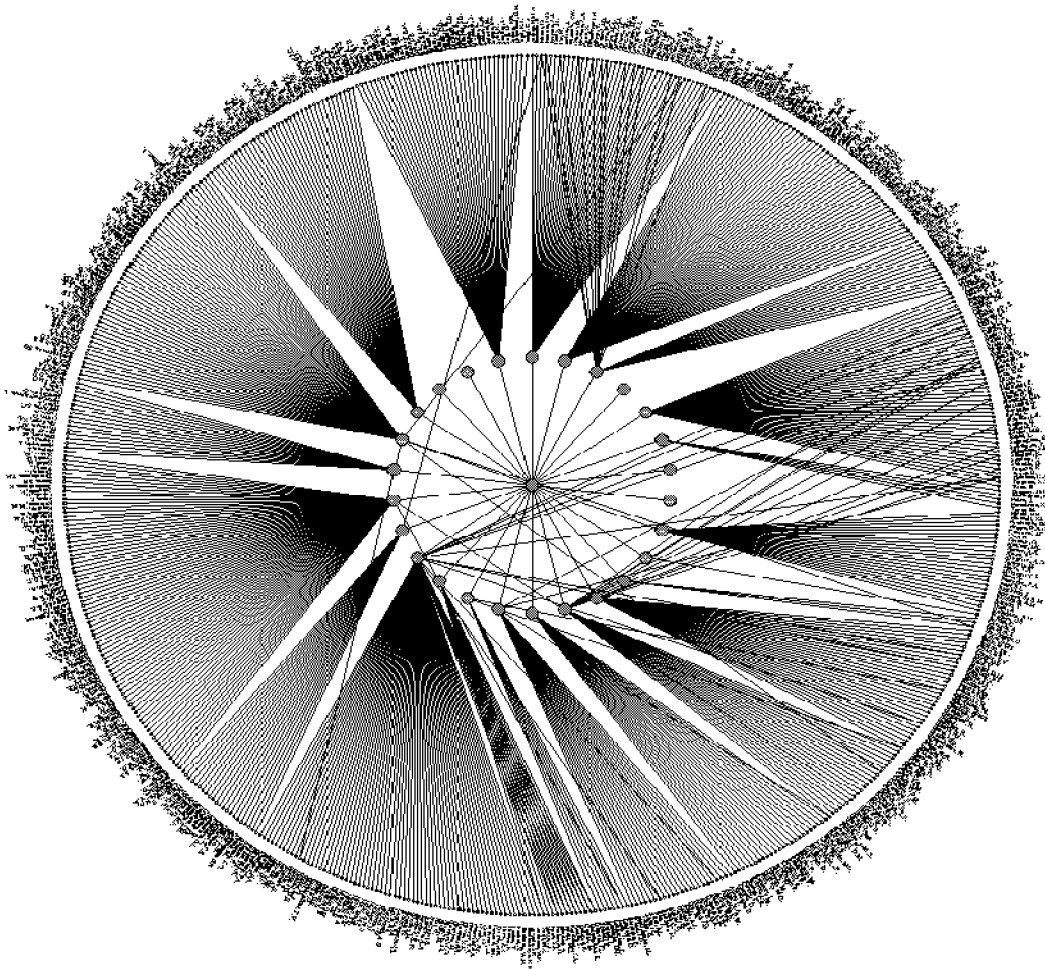
$\gamma = 2.3$



# SCIENCE COAUTHORSHIP

Nodes: scientist (authors)

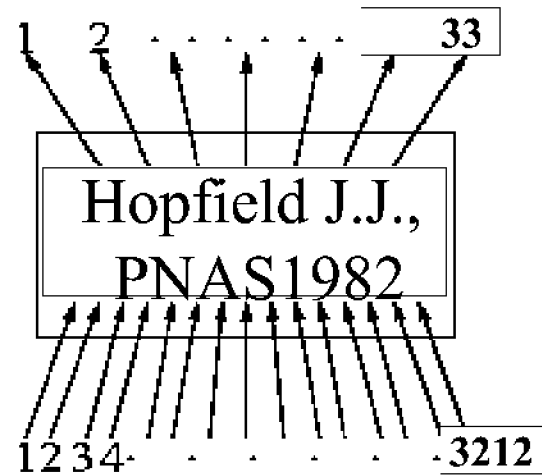
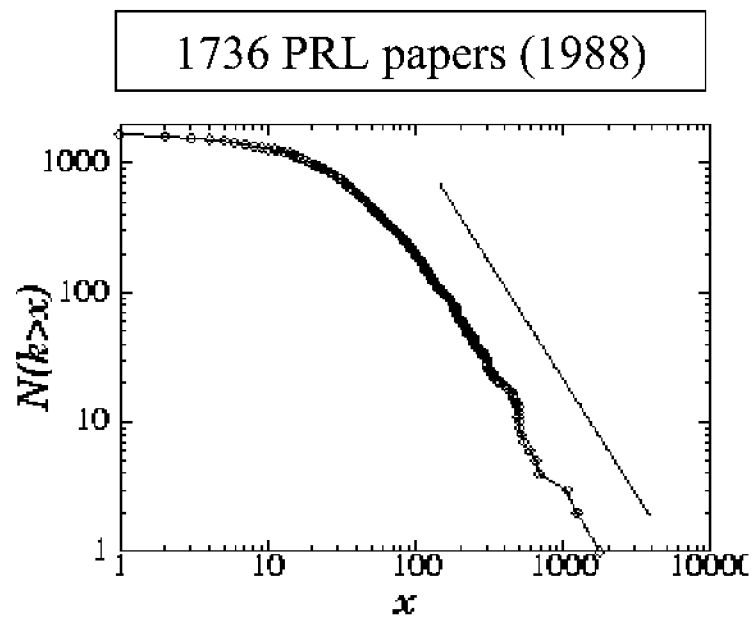
Links: write paper together



(Newman, 2000, H. Jeong et al 2001)

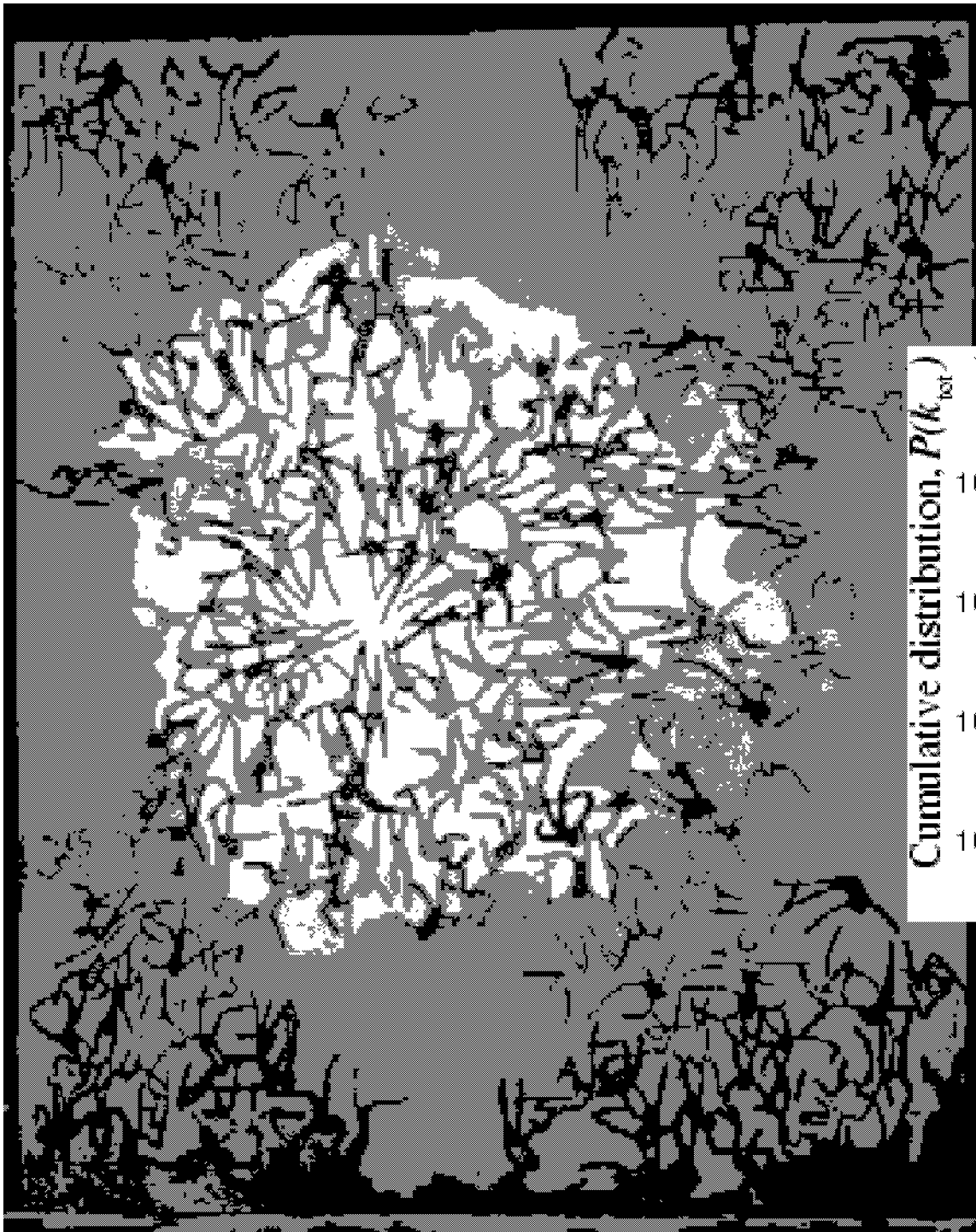
# SCIENCE CITATION INDEX

**Nodes**: papers  
**Links**: citations



$$P(k) \sim k^{-\gamma}$$
$$(\gamma = 3)$$

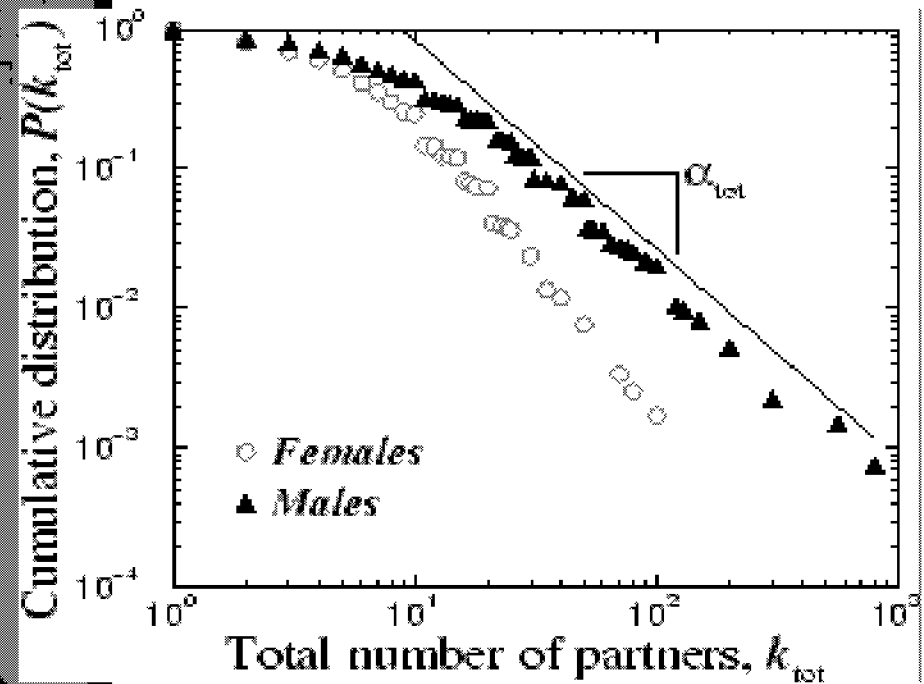
(S. Redner, 1998)



# Swedish sex-web

**Nodes:** people (Females; Males)

**Links:** sexual relationships



4781 Swedes; 18-74;  
59% response rate.

Liljeros et al. Nature 2001

**Many real world networks have  
the same internal structure:**

**Scale-free networks**

**Why?**

# SCALE-FREE NETWORKS

## **(1) The number of nodes (N) is NOT fixed.**

Networks continuously expand by the addition of new nodes

Examples:

WWW : addition of new documents

Citation : publication of new papers

## **(2) The attachment is NOT uniform.**

A node is linked with higher probability to a node that already has a large number of links.

Examples :

WWW : new documents link to well known sites (CNN,

YAHOO, NewYork Times, etc)

Citation : well cited papers are more likely to be cited again

# Scale-free model

## (1) GROWTH :

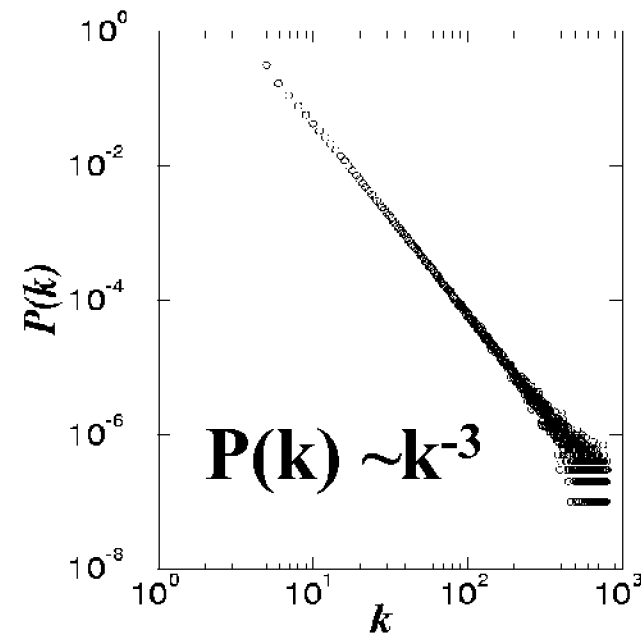
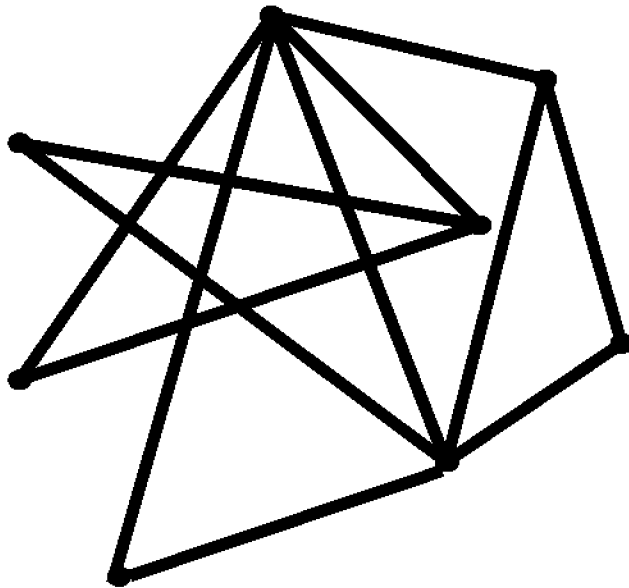
At

every timestep we add a new node with  $m$  edges (connected to the nodes already present in the system).

## (2) PREFERENTIAL ATTACHMENT :

The probability  $\Pi$  that a new node will be connected to node  $i$  depends on the connectivity  $k_i$  of that node

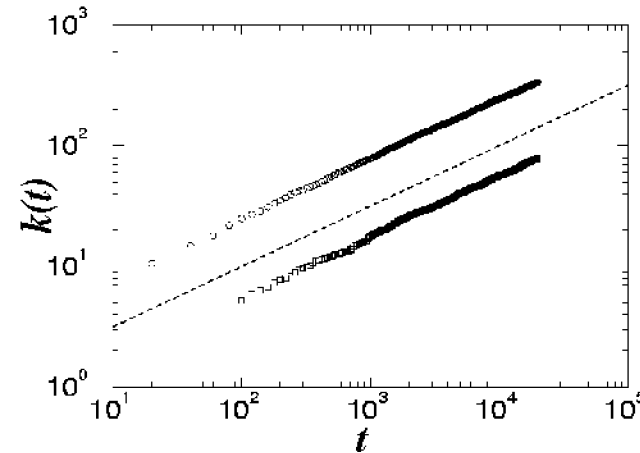
$$\Pi(k_i) = \frac{k_i}{\sum_j k_j}$$



# Mean Field Theory

$$\frac{\partial k_i}{\partial t} \propto \Pi(k_i) = A \frac{k_i}{\sum_j k_j} = \frac{k_i}{2t}, \text{ with initial condition } k_i(t_i) = m$$

$$k_i(t) = m \sqrt{\frac{t}{t_i}}$$



$$P(k_i(t) < k) = P_t(t_i > \frac{m^2 t}{k^2}) = 1 - P_t(t_i \leq \frac{m^2 t}{k^2}) = 1 - \frac{m^2 t}{k^2 (m_0 + t)}$$

$$\therefore P(k) = \frac{\partial P(k_i(t) < k)}{\partial k} = \frac{2m^2 t}{m_0 + t} \frac{1}{k^3} \sim k^{-3}$$

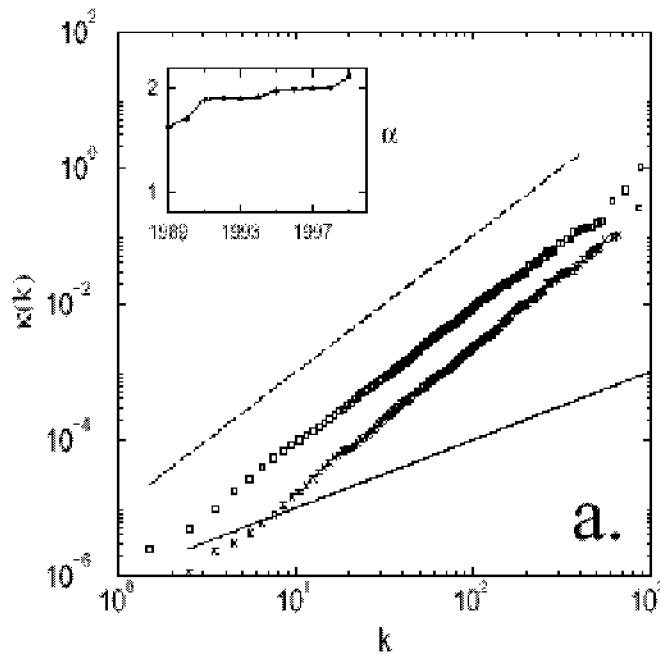
$$\underline{\quad} = 3$$



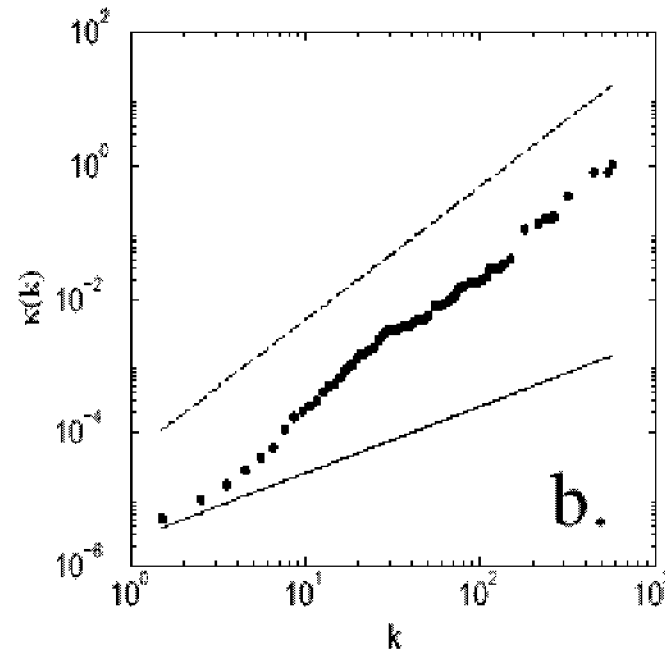
# Preferential Attachment

$$\frac{\partial k_i}{\partial t} \propto \Pi(k_i) \sim \frac{\Delta k_i}{\Delta t} \quad \text{For given } \Delta t, \therefore \Delta k \propto \Pi(k)$$

**k vs.  $\Delta k$  : increase in the No. of links in a unit time**



**Citation  
network**

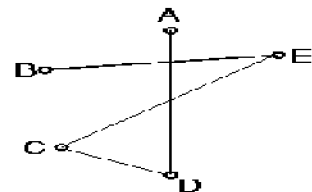


**Internet**

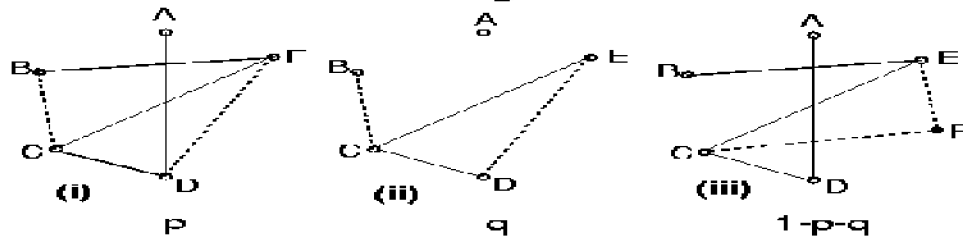
(Jeong, Neda, A.-L. B, cond-mat/0104131)

# Universality?

WWW (in)	Internet	Actor	Citation index	Sex Web	Cellular network	Phone call network	linguistics
$\gamma = 2.1$	$\gamma = 2.5$	$\gamma = 2.3$	$\gamma = 3$	$\gamma = 3.5$	$\gamma = 2.1$	$\gamma = 2.1$	$\gamma = 2.8$

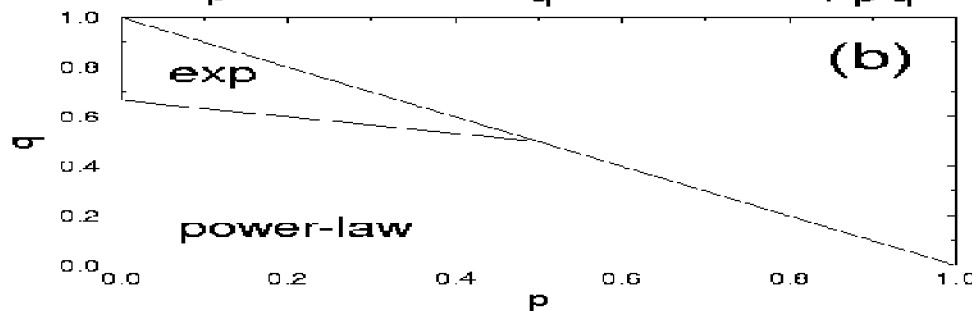


(a)



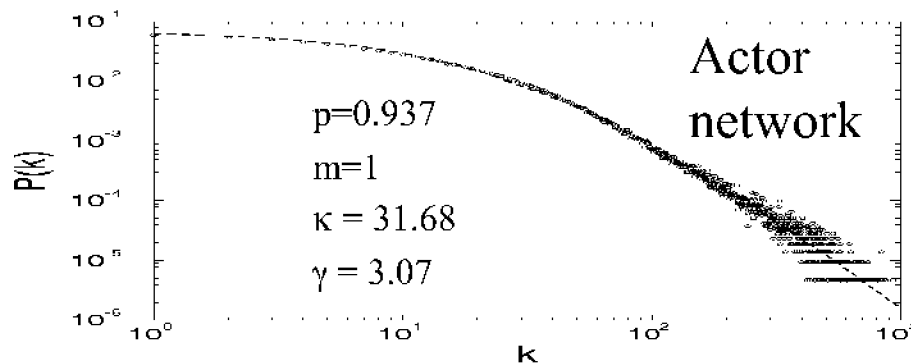
Extended Model

- prob.  $p$  : internal links
- prob.  $q$  : link deletion
- prob.  $1-p-q$  : add node



$$P(k) \sim (k + \kappa(p, q, m))^{-\gamma(p, q, m)}$$

$$\gamma \in [1, \infty)$$



- Predict the network topology from microscopic processes with parameters  $(p, q, m)$
- Scaling but no universality

## Other Models

- **Non-linear preferential attachment :**  
 $k^\alpha \rightarrow P(k) \sim$  no scaling for  $\alpha \neq 1$

$$\Pi(k) \sim$$

$\Rightarrow \alpha < 1$  : stretch-exponential

$\Rightarrow \alpha > 1$  : no-scaling ( $\alpha > 2$  : “gelation”)

(Krapivsky et al (2000).)

- **Initial attractiveness :**  $\Pi(k) \sim A + k^\alpha$

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

(Dorogovtsev et al (2000).)

- **Aging :** each node has a lifetime  
node cannot get links after retirement. (actor)

$\rightarrow$

$\Rightarrow P(k)$  : power-law with exponential cutoff

(Amaral et al (2000).)

## Other Models (continued)

- **Saturation** : each node has maximum link number.

→ node cannot get links after finite # of links

⇒  $P(k)$  : power-law with exponential cutoff

(Amaral et al (2000).)

- **Walking on the network:**

**Each new node :**

→ connects to a randomly selected node

→ with prob.  $p$  to nodes the selected node points to

(repeated recursively with the new links)

For  $p=1$ :  $P(k) \sim k^{-2}$

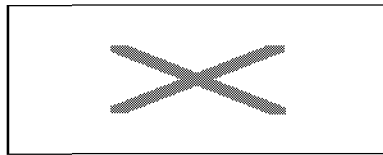
(Vazquez, 2000)

# Can Latecomers Make It? Fitness Model

SF model:  $k(t) \sim t$  ————— (first mover advantage)

Real systems: nodes compete for links -- *fitness*

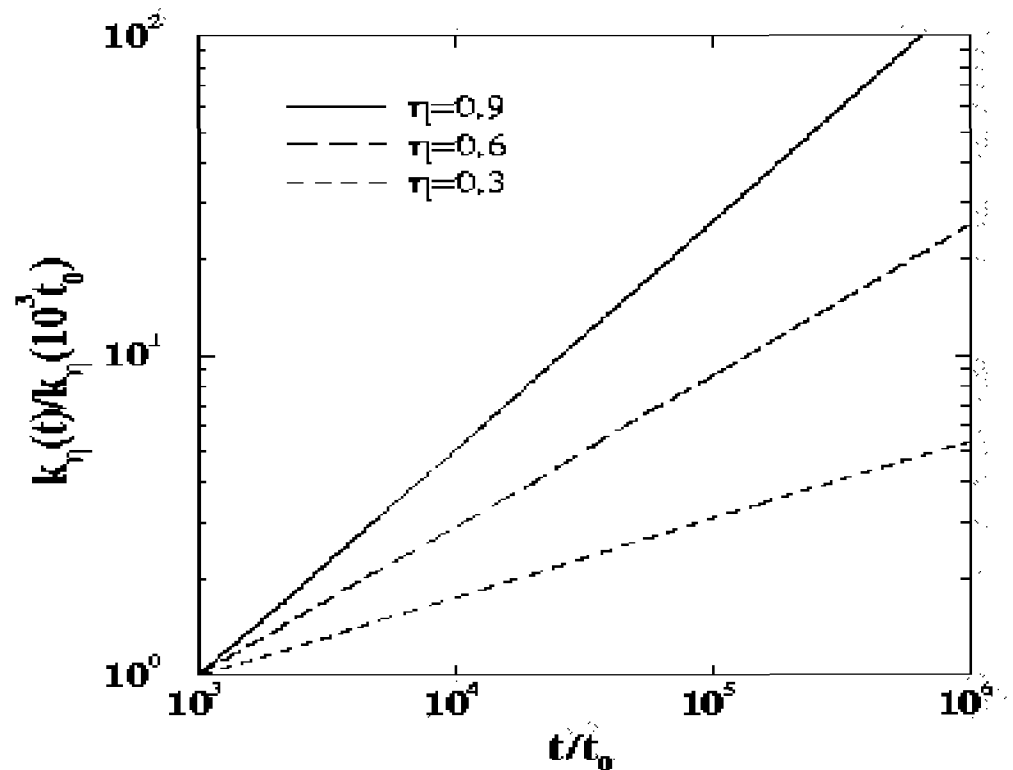
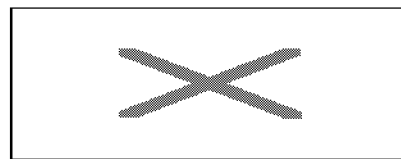
Fitness Model: fitness ( $\eta$ )



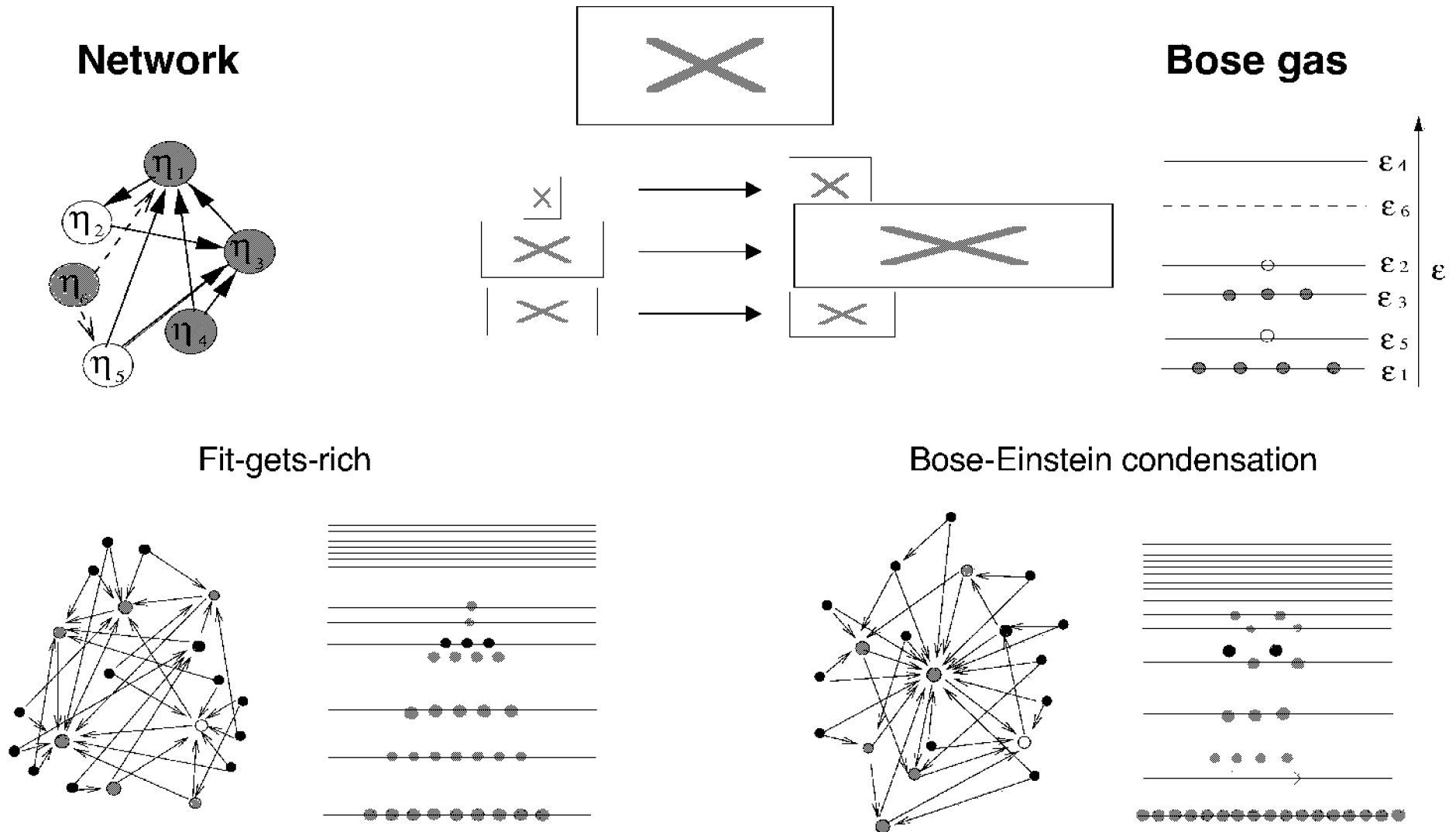
$$k(\eta, t) \sim t^{\beta(\eta)}$$

where

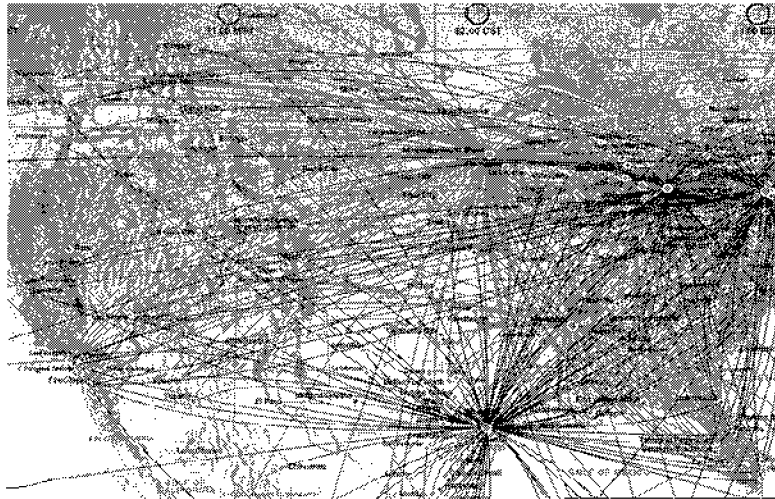
$$\beta(\eta) = \eta/C$$



# Bose-Einstein Condensation in Evolving Networks



# What is the topology of cellular networks?

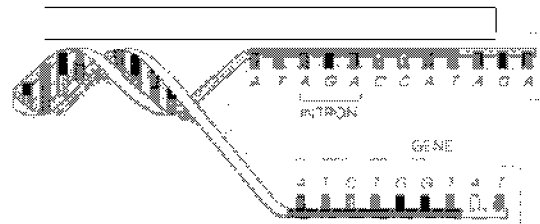
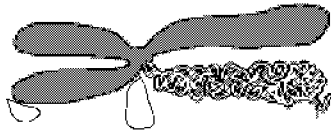
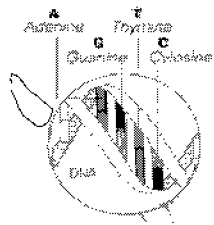


**Argument 1:**  
**Cellular networks are  
scale-free!**

**Reason:** They  
formed one node at a  
time...

**Argument 2:**  
**Cellular networks are  
exponential!**

**Reason:** They  
have been streamlined by  
evolution...



**GENOME**

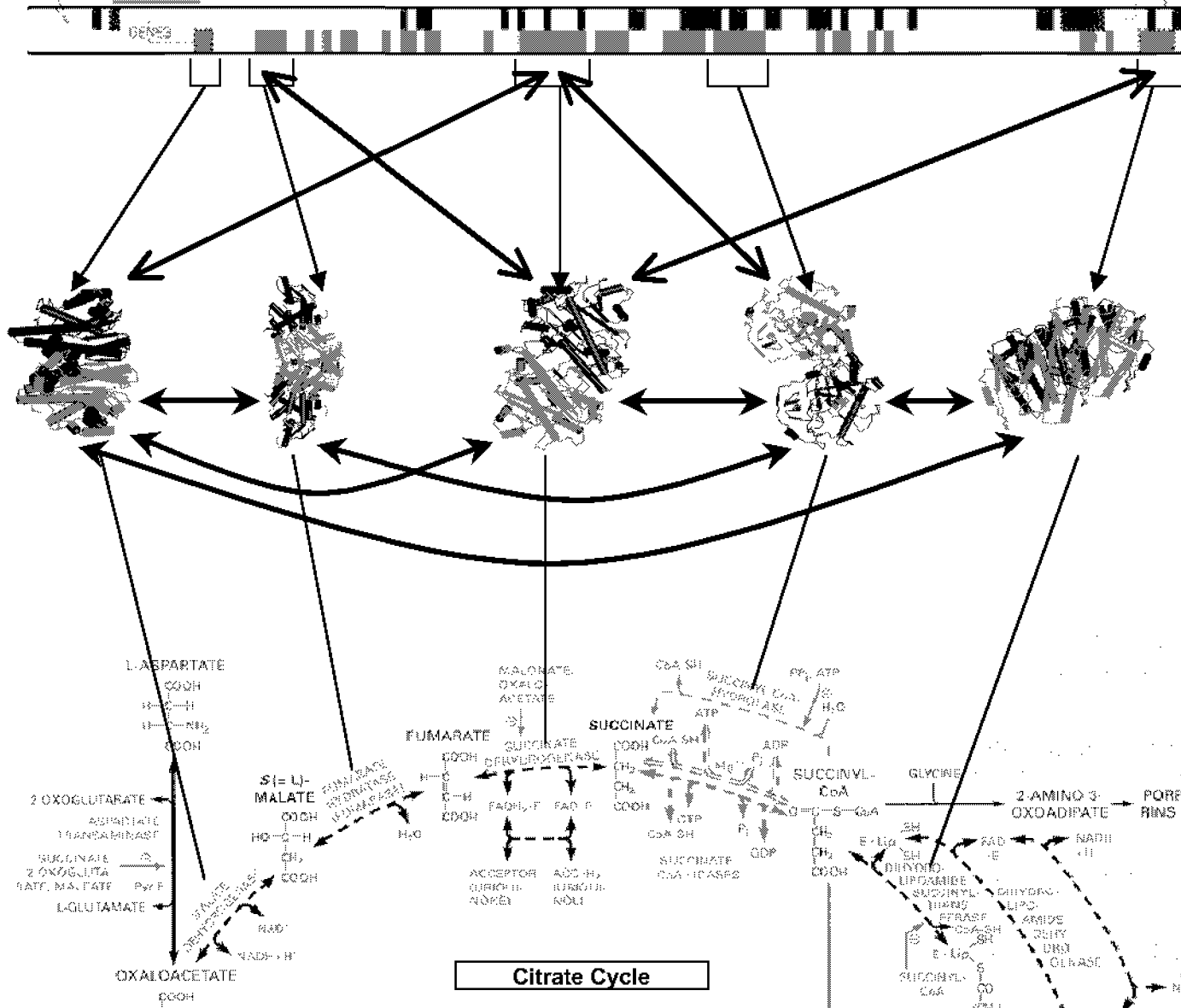
**protein-gene interactions**

**PROTEOME**

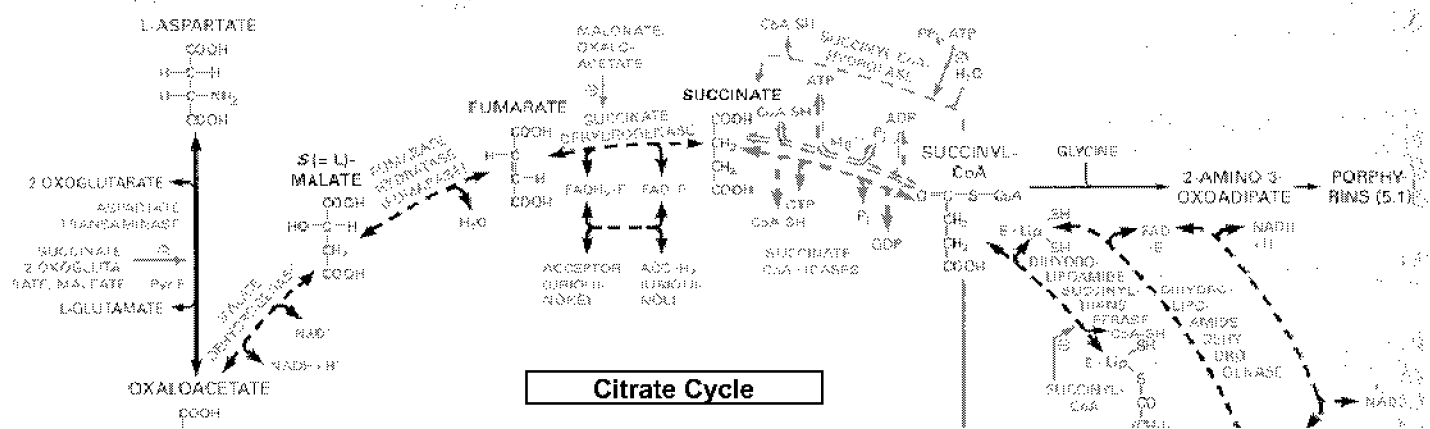
**protein-protein interactions**

**METABOLISM**

**Bio-chemical reactions**







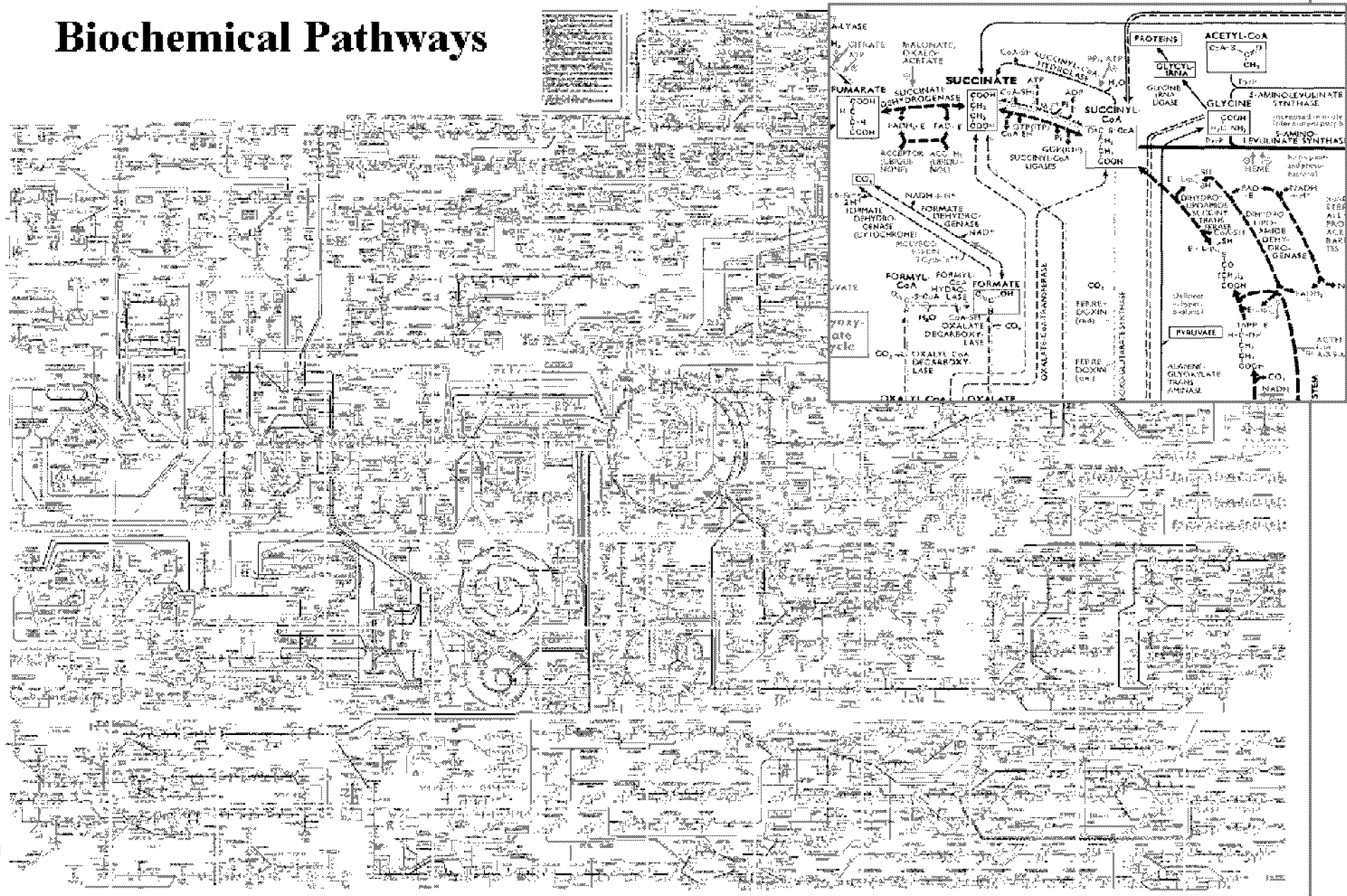
# METABOLISM

## Bio-chemical reactions

A B C D E F G H I J K L

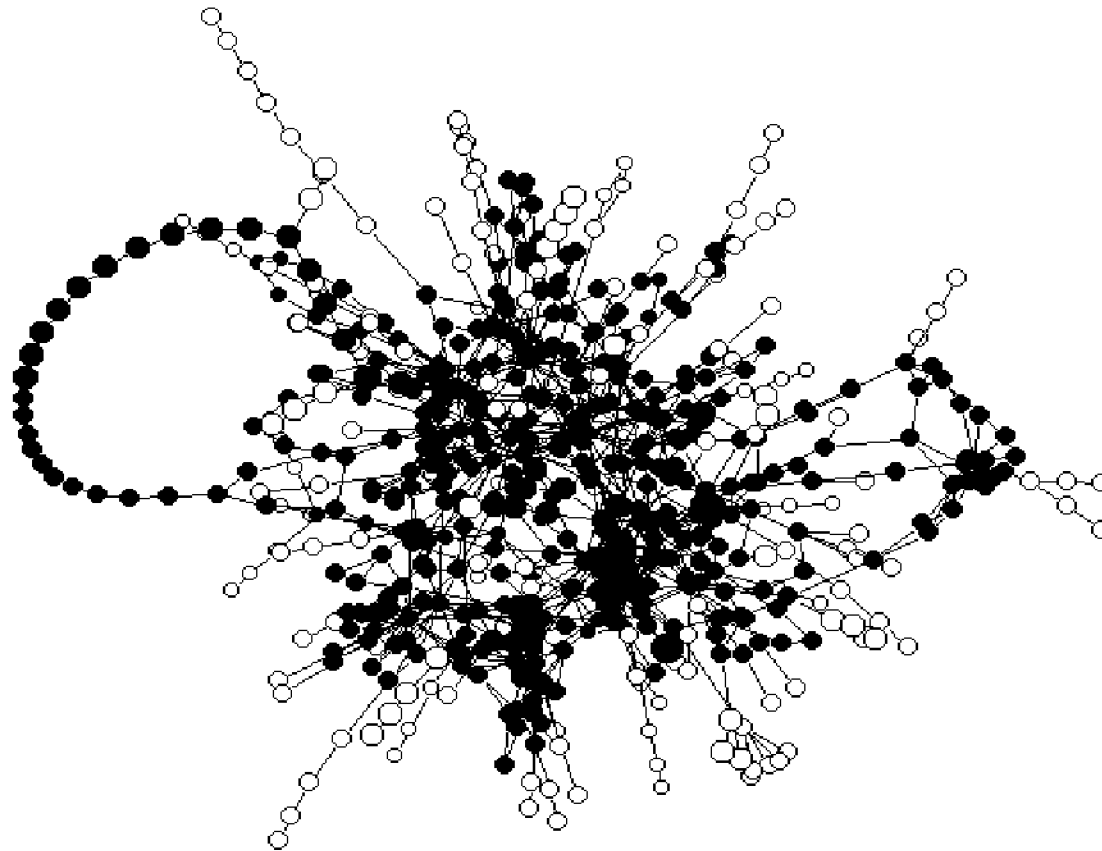
# Biochemical Pathways

1  
2  
3  
4  
5  
6  
7  
8  
9  
10

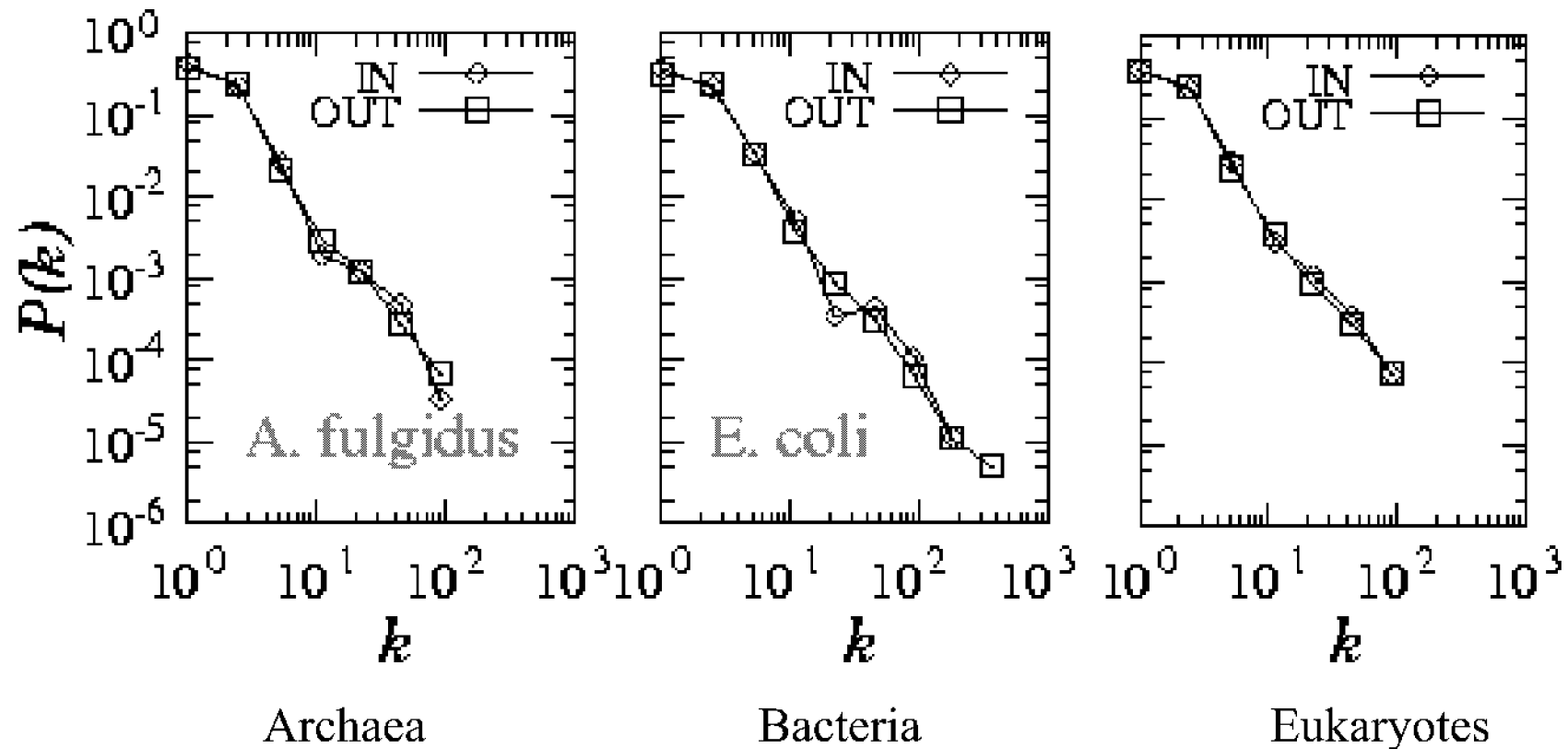


# Metabolic Network

Nodes: chemicals (substrates)  
Links: bio-chemical reactions



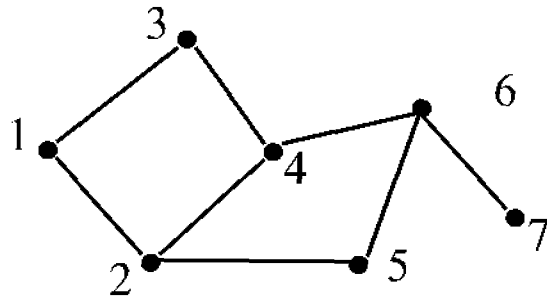
# Metabolic network



Organisms from all three domains of life are **scale-free** networks!

H. Jeong, B. Tombor, R. Albert, Z.N. Oltvai, and A.L. Barabasi, Nature, **407** 651 (2000)

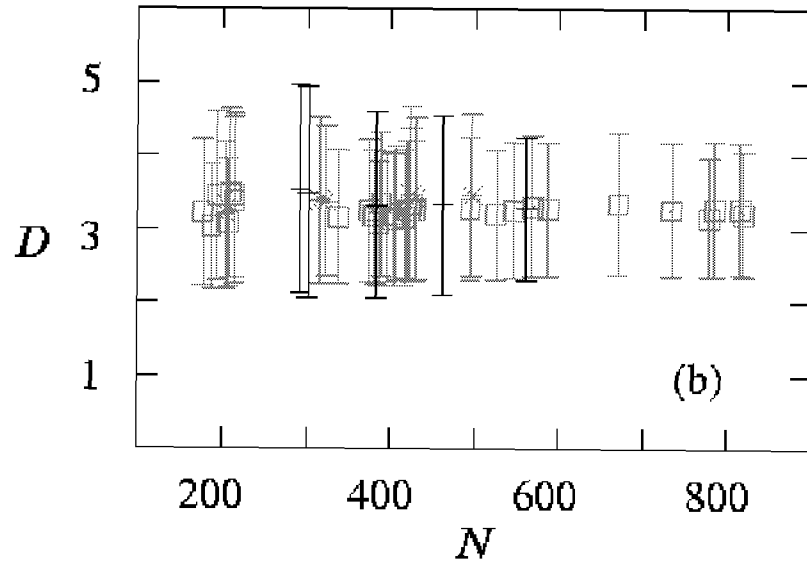
# Node-node distance in metabolic networks



$$D_{15} = 2 [1 \rightarrow 2 \rightarrow 5]$$

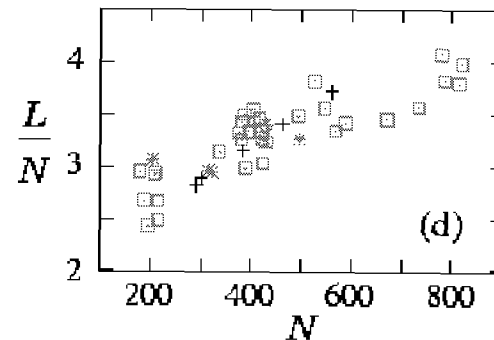
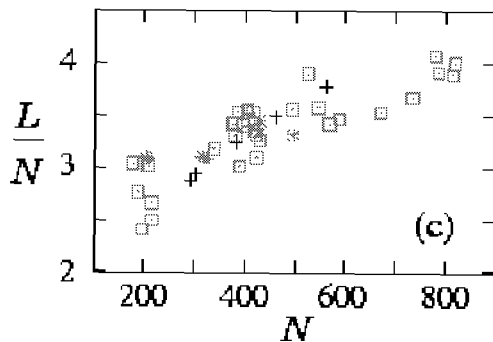
$$D_{17} = 4 [1 \rightarrow 3 \rightarrow 4 \rightarrow 6 \rightarrow 7]$$

...  $D = ??$

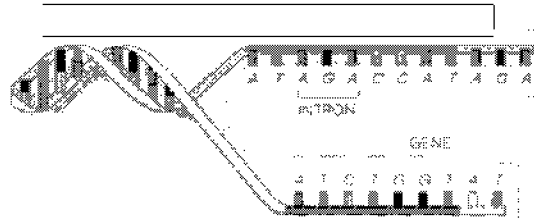
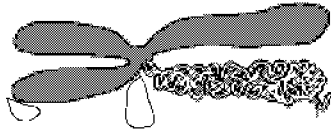
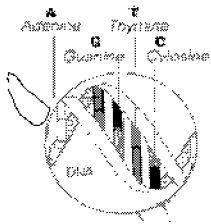


**Scale-free networks:**

$$D \sim \log(N)$$



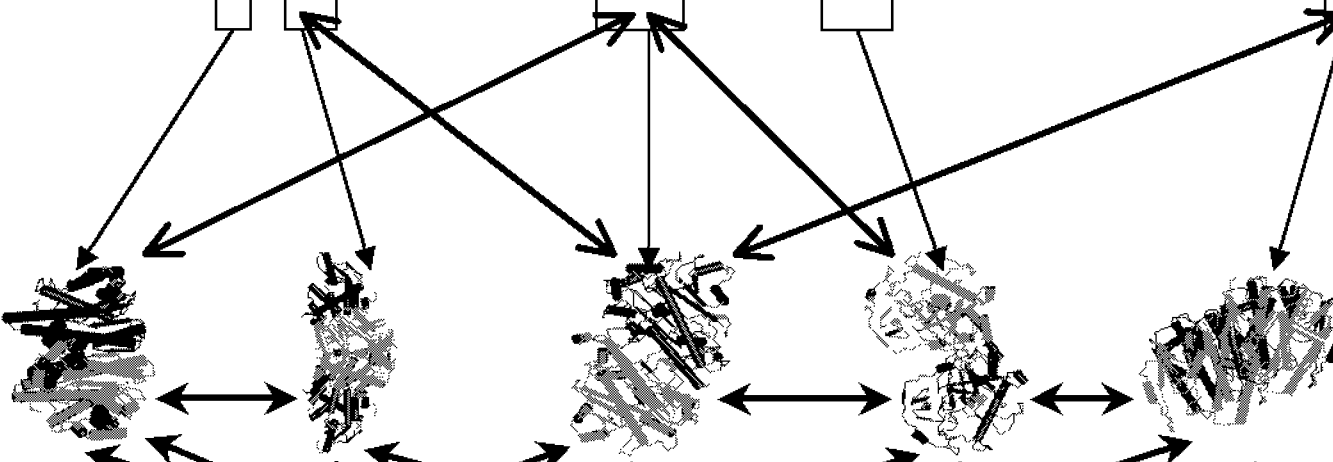
**Larger organisms are expected to have a larger diameter!**



**GENOME**



**protein-gene interactions**

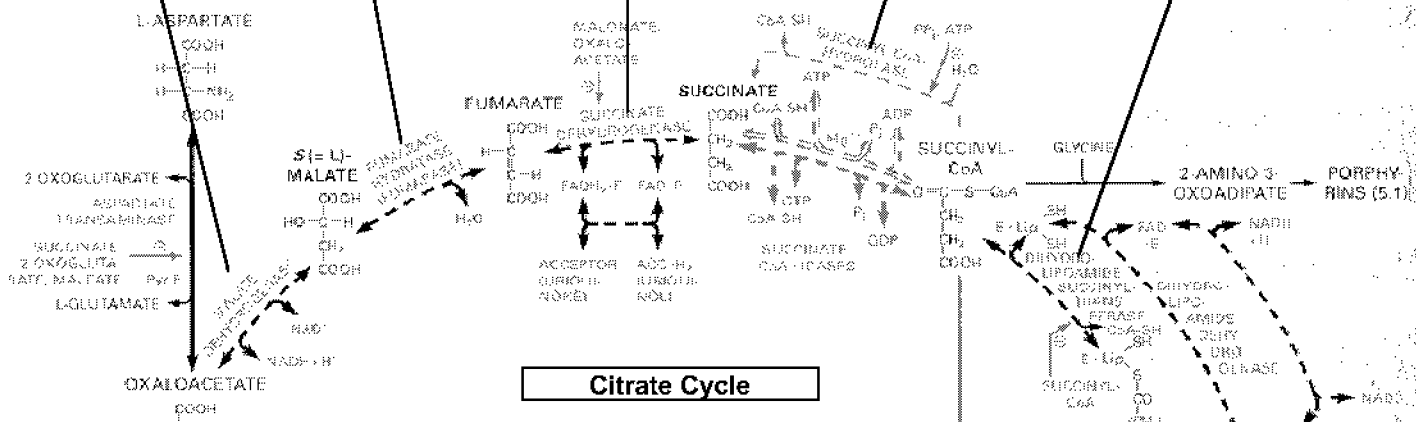


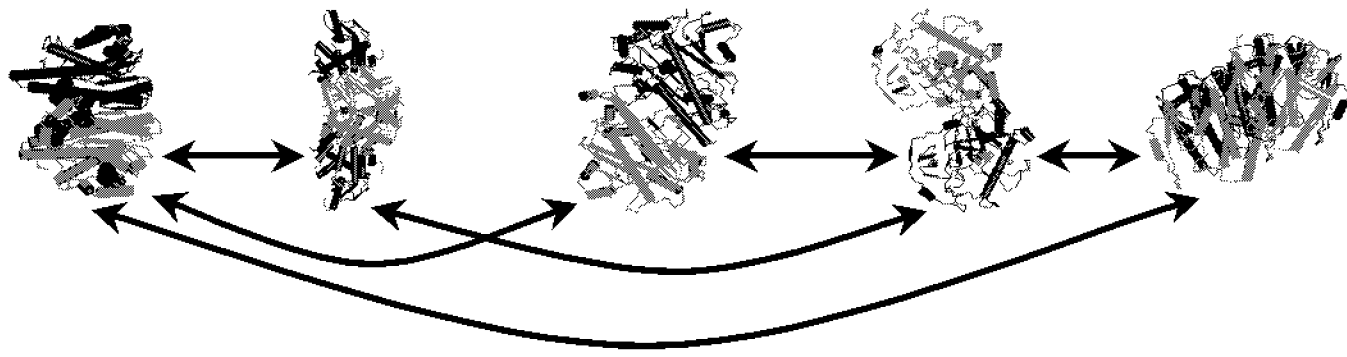
**PROTEOME**

**protein-protein interactions**

**METABOLISM**

**Bio-chemical reactions**





**PROTEOME**  
**protein-protein**  
**interactions**

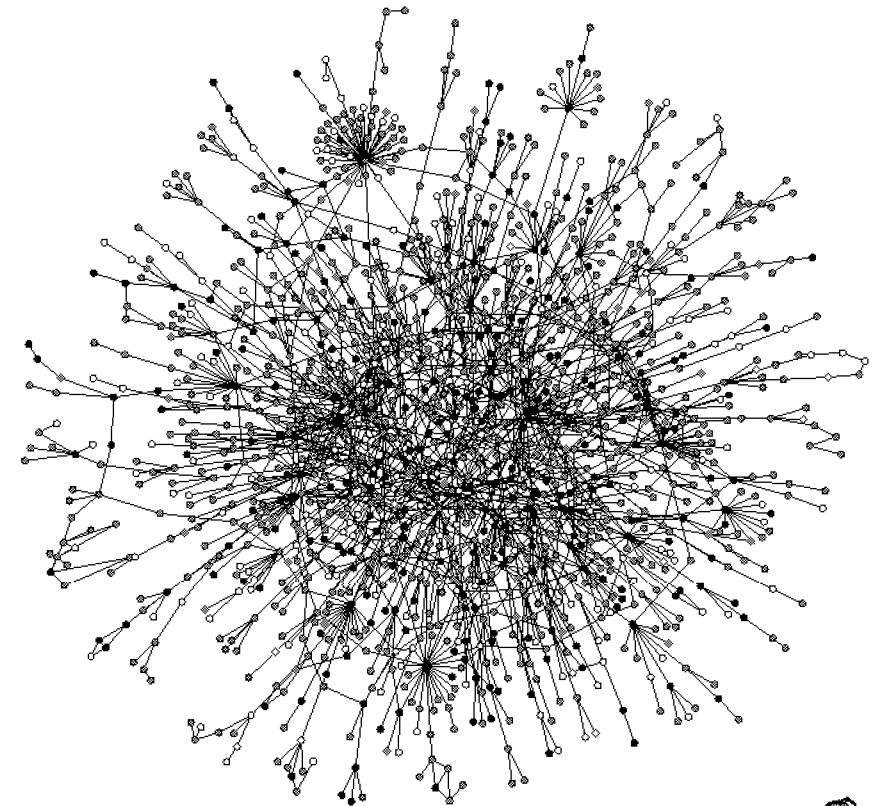
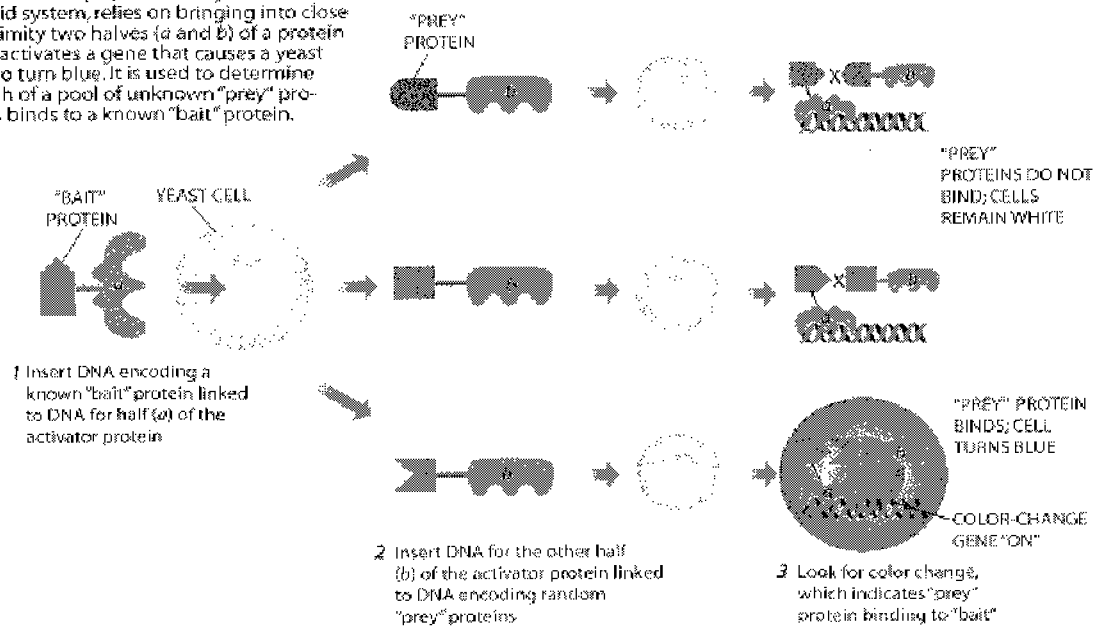
# Yeast protein network

**Nodes:** proteins

**Links:** physical interactions (binding)

## Finding Proteins That Interact

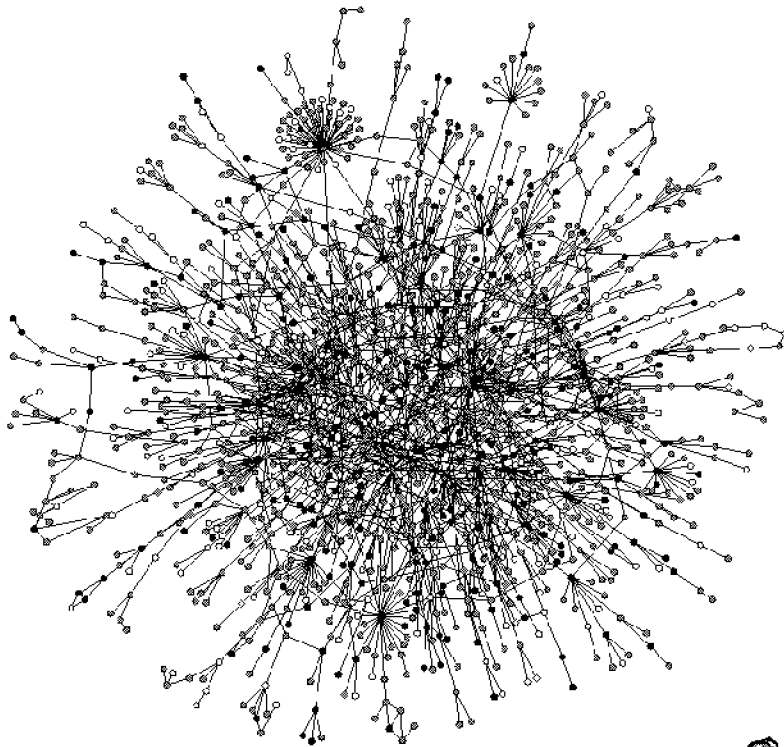
One technique, called the yeast two-hybrid system, relies on bringing into close proximity two halves (a and b) of a protein that activates a gene that causes a yeast cell to turn blue. It is used to determine which of a pool of unknown "prey" proteins binds to a known "bait" protein.



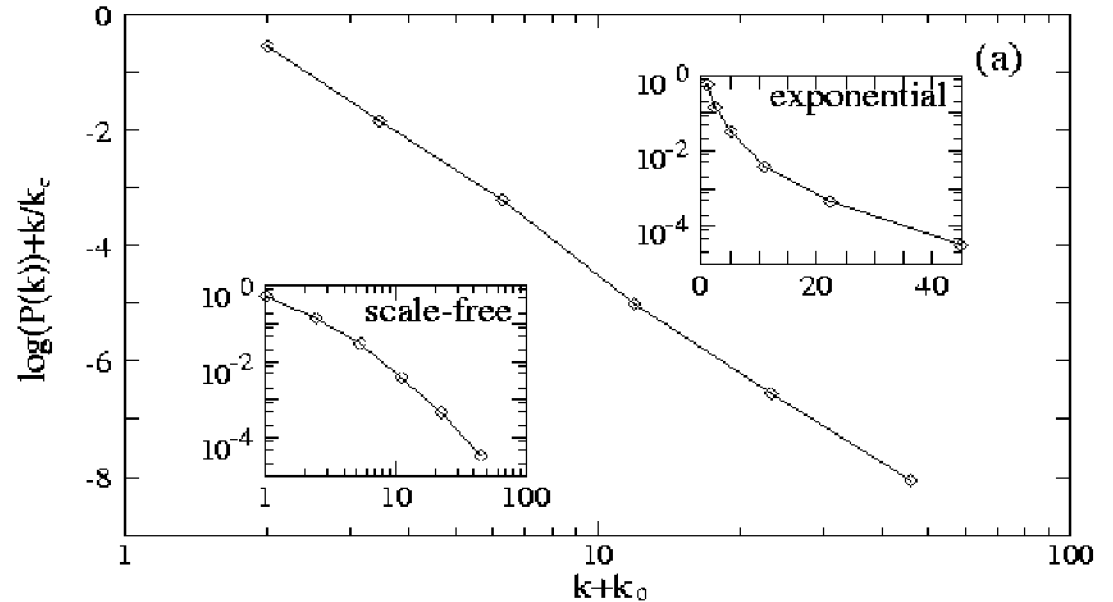
P. Uetz, *et al.* *Nature* **403**, 623-7 (2000).



# Topology of the protein network



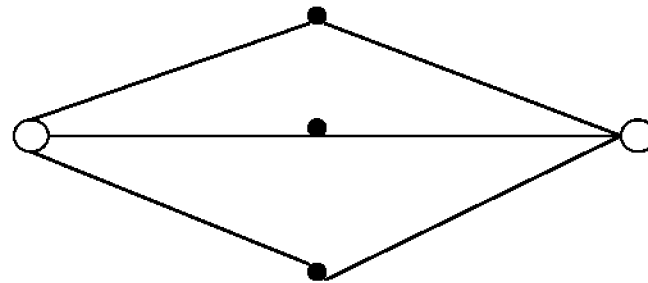
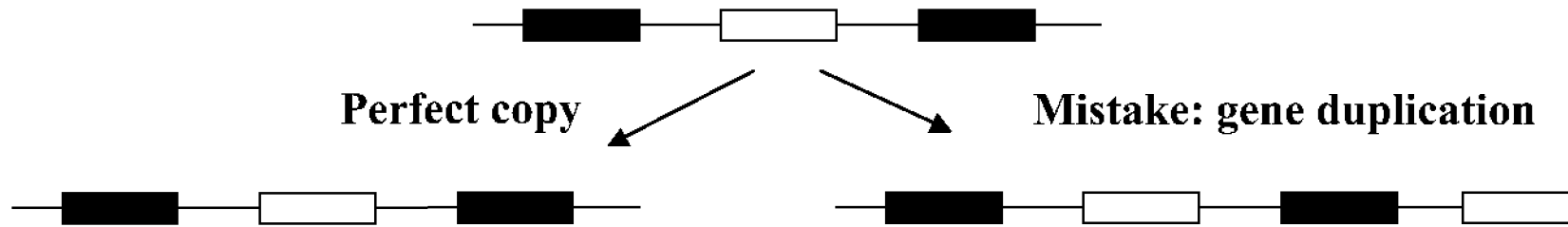
Palek



$$P(k) \sim (k + k_0)^{-\gamma} \exp\left(-\frac{k + k_0}{k_\tau}\right)$$

**H. Jeong, S.P. Mason, A.-L. Barabasi, Z.N. Oltvai, Nature 411, 41-42 (2001)**

# Origin of scaling in protein interaction networks



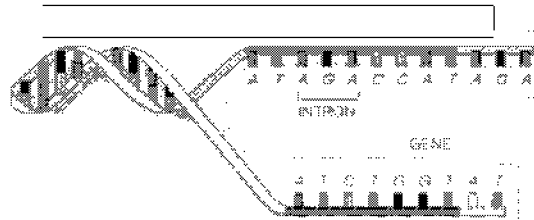
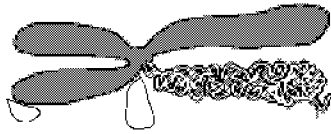
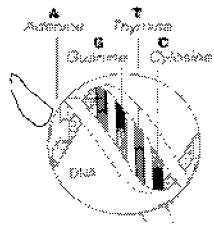
Proteins with more interactions are more likely to get a new link:

$$P(k) \sim k$$

(preferential attachment).

Vazquez, Flammini, Maritan, Vespignani cond-mat/0108043

Sole, Pastor-Satorras, Smith, Kepler, Adv. Compl. Syst. 2001



**GENOME**

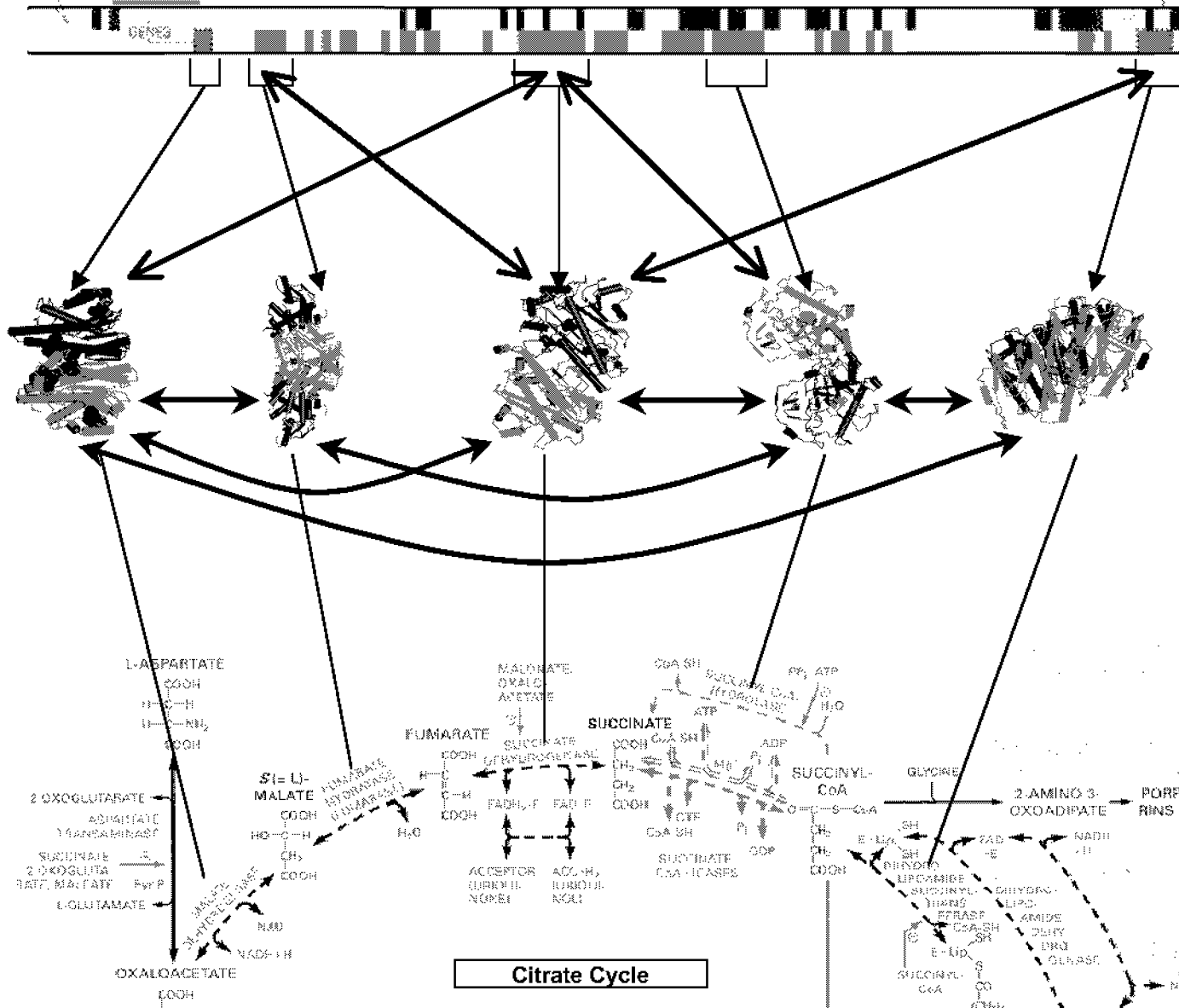
**protein-gene interactions**

**PROTEOME**

**protein-protein interactions**

**METABOLISM**

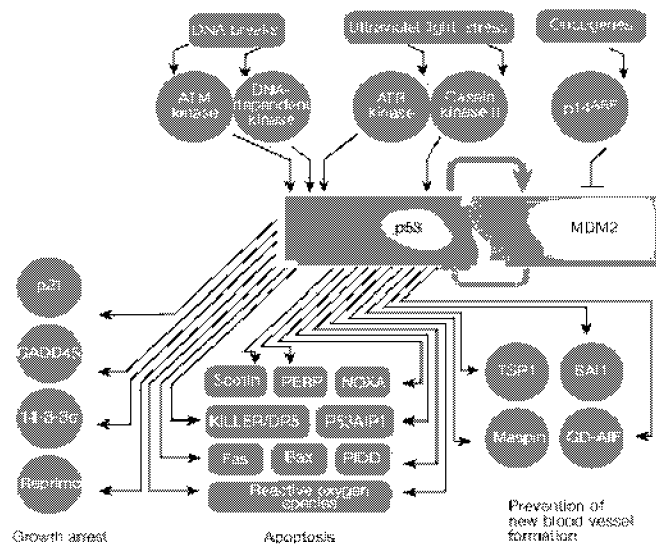
**Bio-chemical reactions**



# Surfing the p53 network

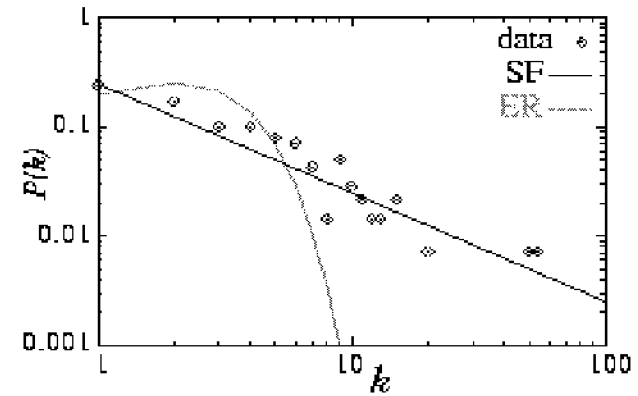
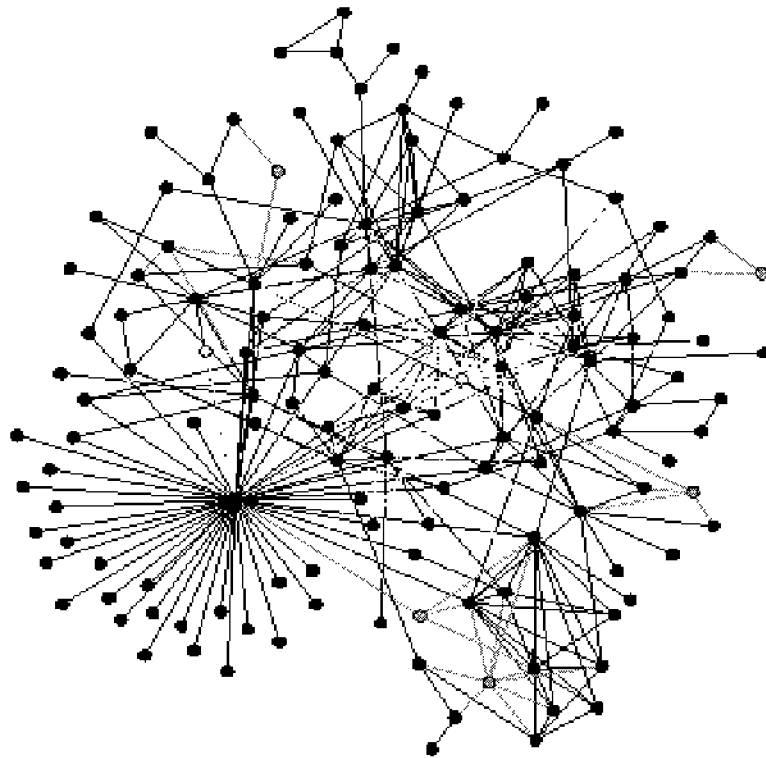
Bert Vogelstein, David Lane and Arnold J. Levine

The p53 tumour-suppressor gene integrates numerous signals that control cell life and death. As when a highly connected node in the Internet breaks down, the disruption of p53 has severe consequences.



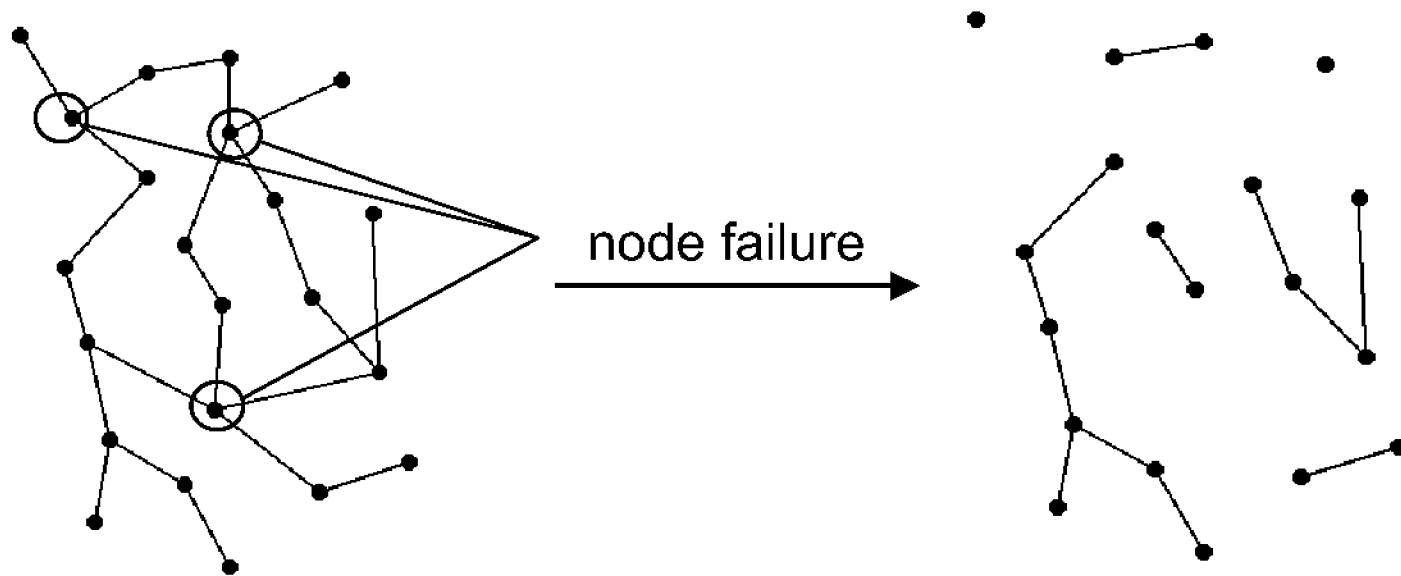
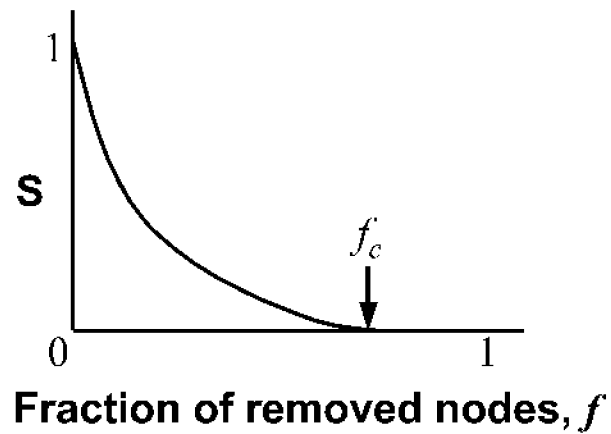
“One way to understand the p53 network is to compare it to the Internet. The cell, like the Internet, appears to be a **‘scale-free network’**.”

## p53 network (mammals)



# Robustness

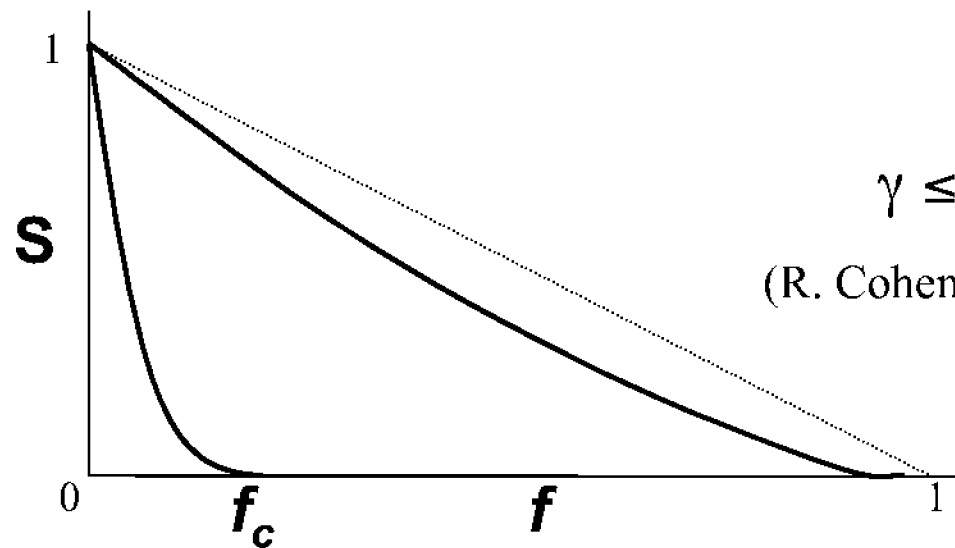
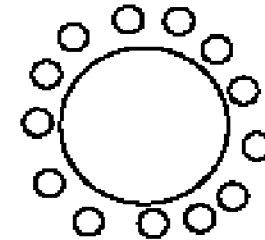
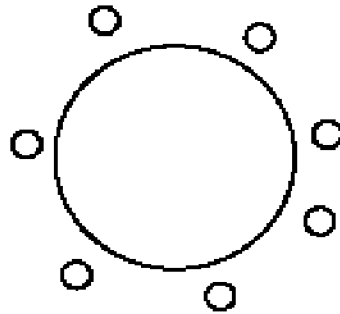
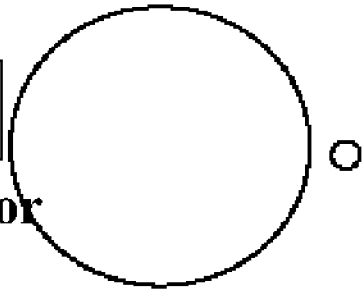
Complex systems maintain their basic functions  
even under errors and failures  
(cell  $\rightarrow$  mutations; Internet  $\rightarrow$  router breakdowns)



# Robustness of scale-free networks

Failures

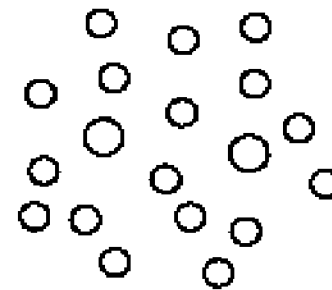
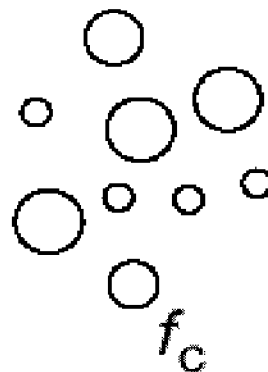
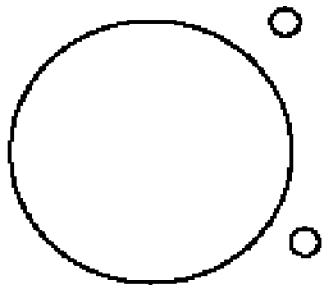
Topological error tolerance



$$\gamma \leq 3 : f_c = 1$$

(R. Cohen et al PRL, 2000)

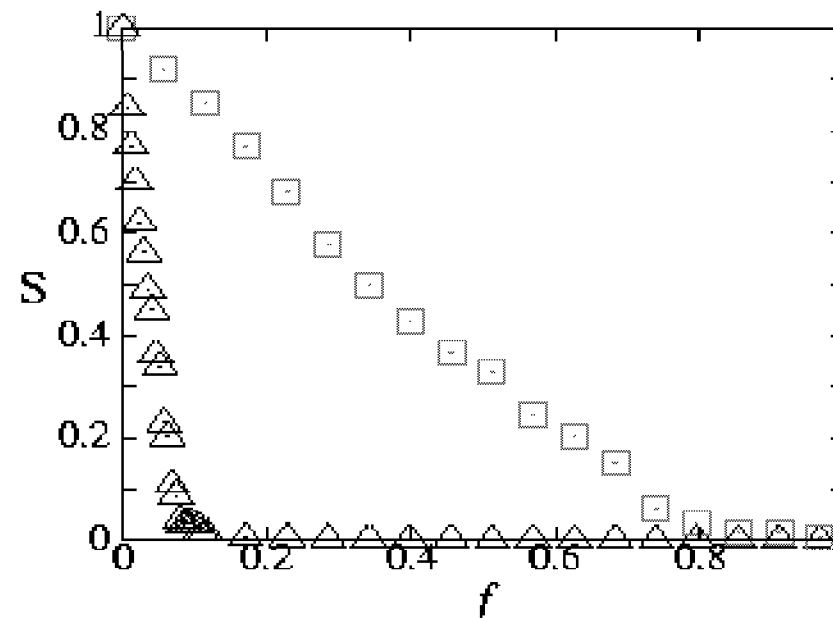
Attacks



# Achilles' Heel of complex networks

— failure  
— attack

Internet

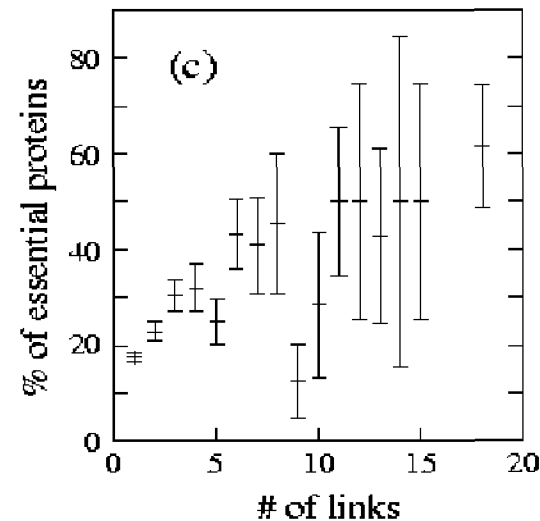
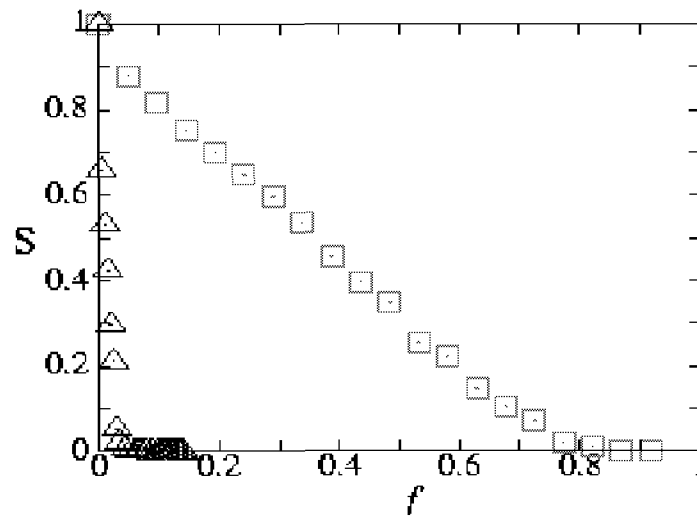


R. Albert, H. Jeong, A.L. Barabasi, Nature **406** 378 (2000)



# Yeast protein network

## - lethality and topological position -



Highly connected proteins are more **essential (lethal)**...

H. Jeong, S.P. Mason, A.-L. Barabasi, Z.N. Oltvai, Nature 411, 41-42 (2001)

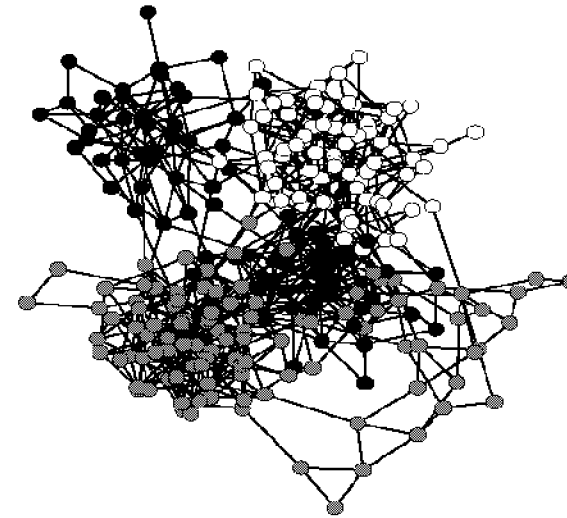
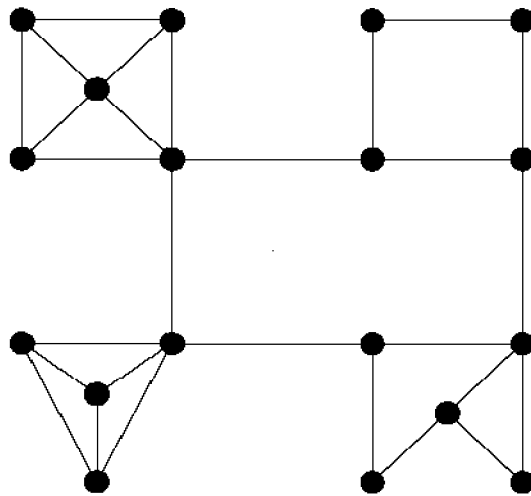
# Modularity in Cellular Networks

➤ Hypothesis:

Biological functions are carried by discrete functional modules.

❖ Hartwell, L.-H., Hopfield, J. J., Leibler, S., & Murray, A. W. (1999).

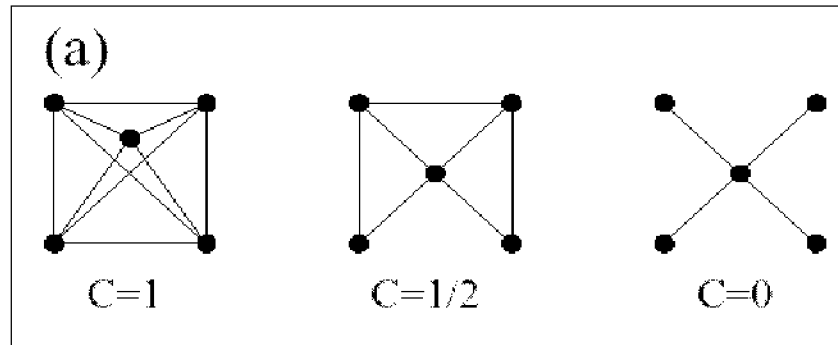
➤ Traditional view of modularity:



➤ Question: Is modularity a myth, or a structural property of biological networks?  
(are biological networks fundamentally modular?)

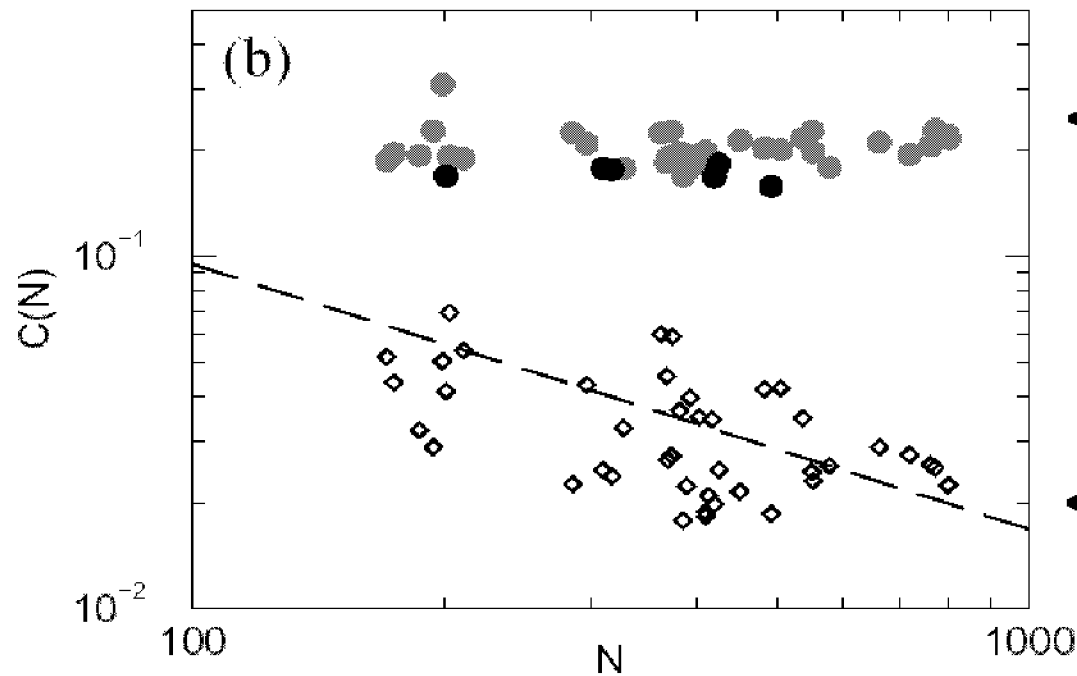
Ravasz, Somera, Mongru, Oltvai, A-L. B, *Science* 297, 1551 (2002).

# Modularity in the metabolism



**Clustering Coefficient:**

$$C(k) = \frac{\text{\# links between } k \text{ neighbors}}{k(k-1)/2}$$

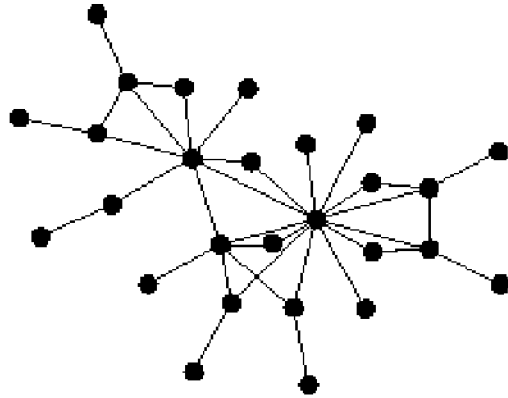


← **Metabolic network  
(43 organisms)**

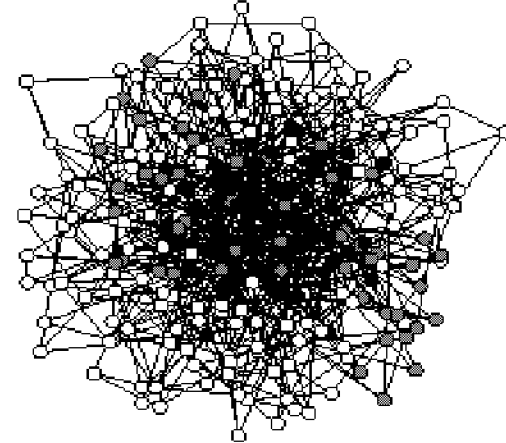
← **Scale-free model**

# Modular vs. Scale-free Topology

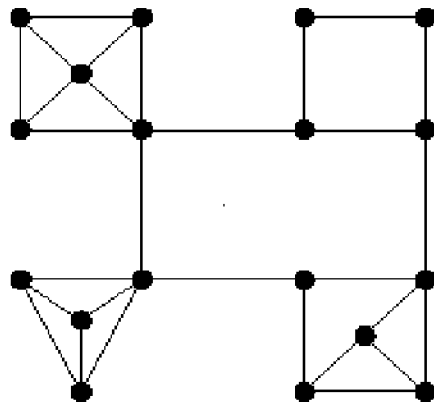
(a)



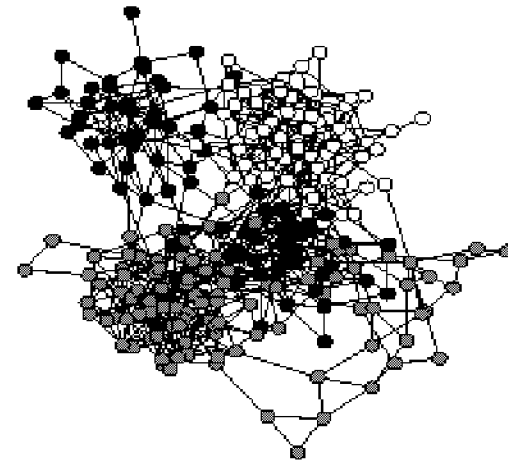
Scale-free



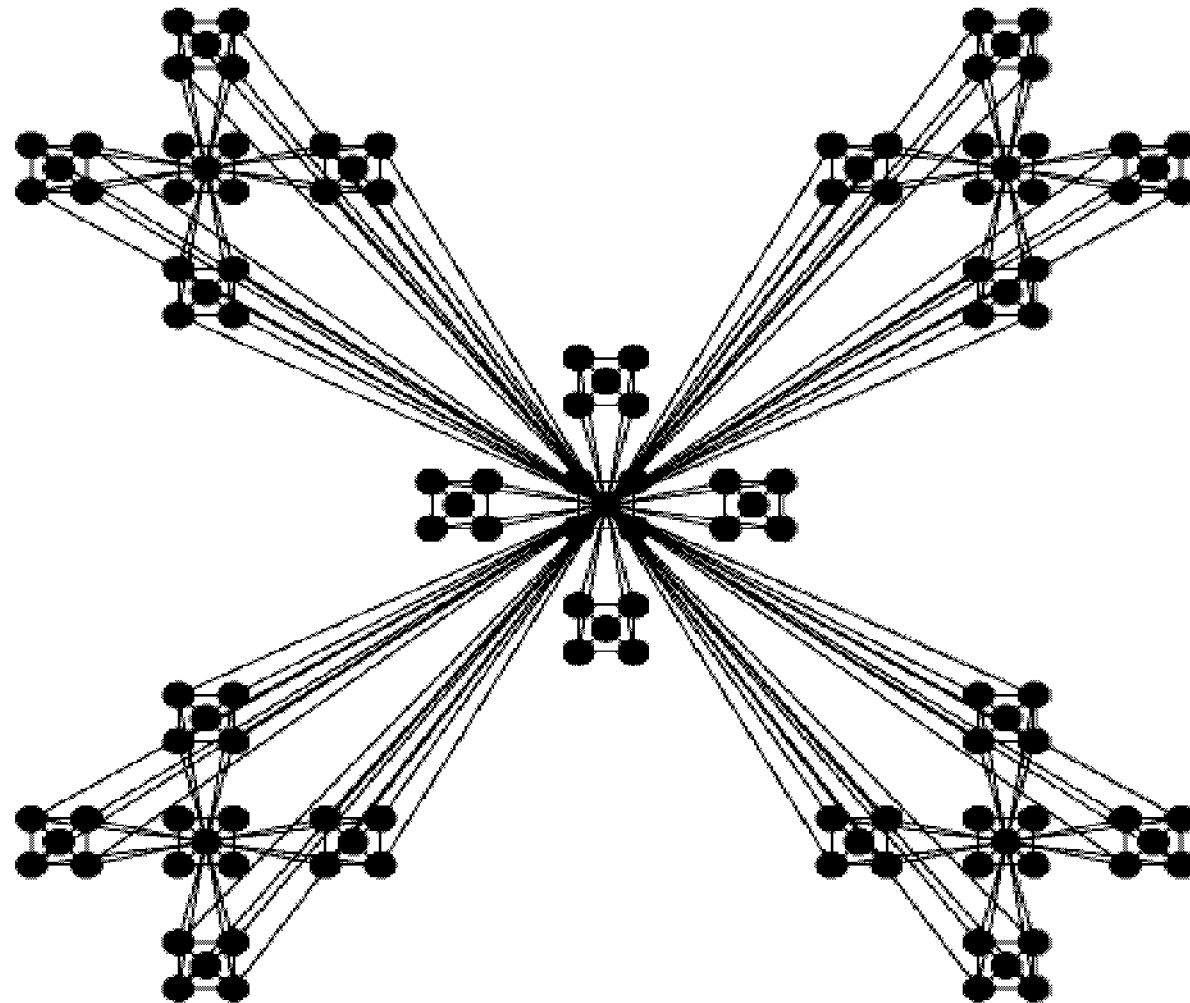
(b)



Modular



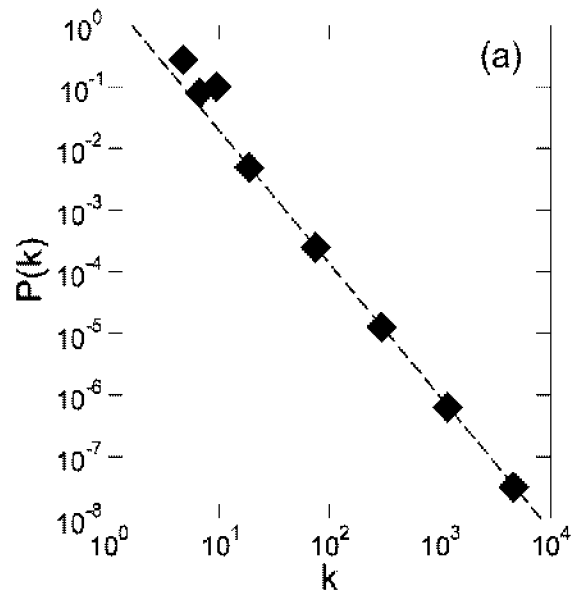
# Hierarchical Networks



# Properties of hierarchical networks

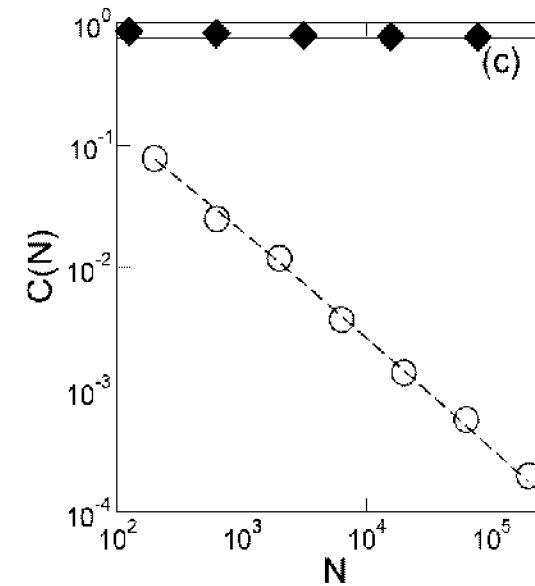
## 1. Scale-free

$$\begin{aligned}\gamma &= 1 + \frac{\ln 5}{\ln 4} \\ &= 2.161\end{aligned}$$



## 2. Clustering coefficient independent of N

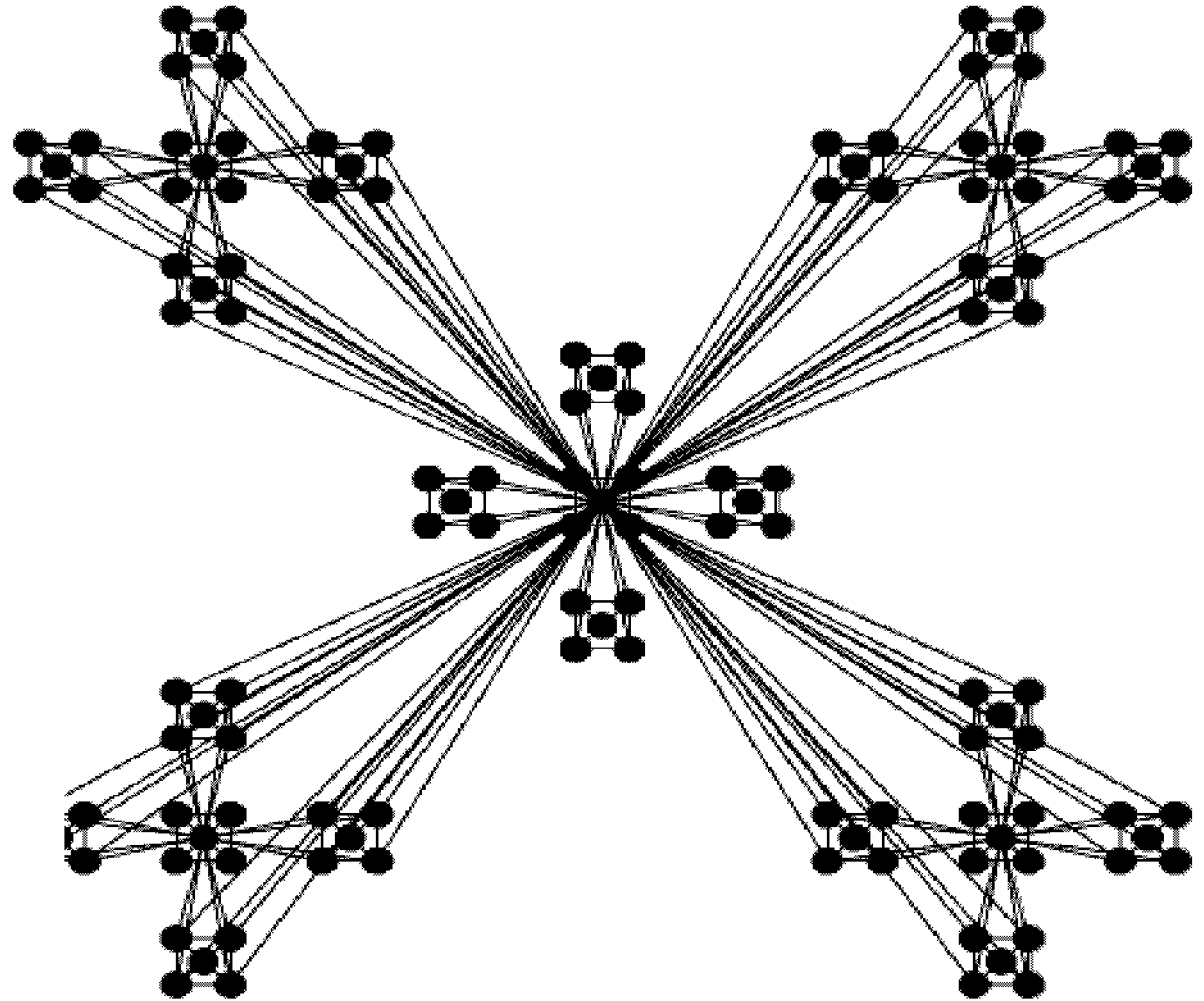
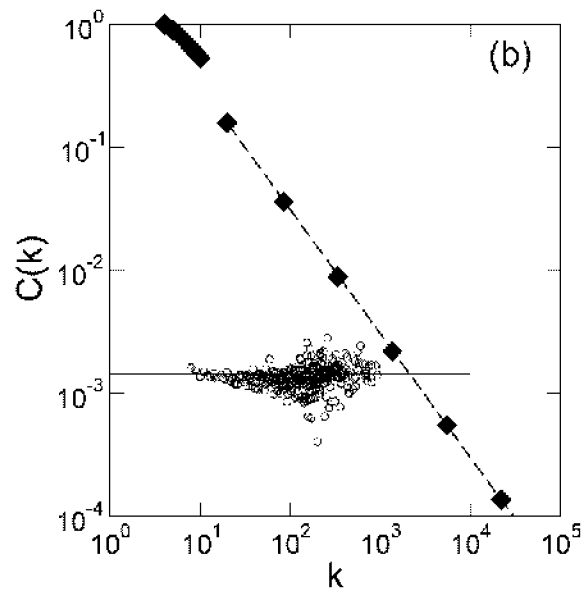
$$C(N) = \text{const.}$$



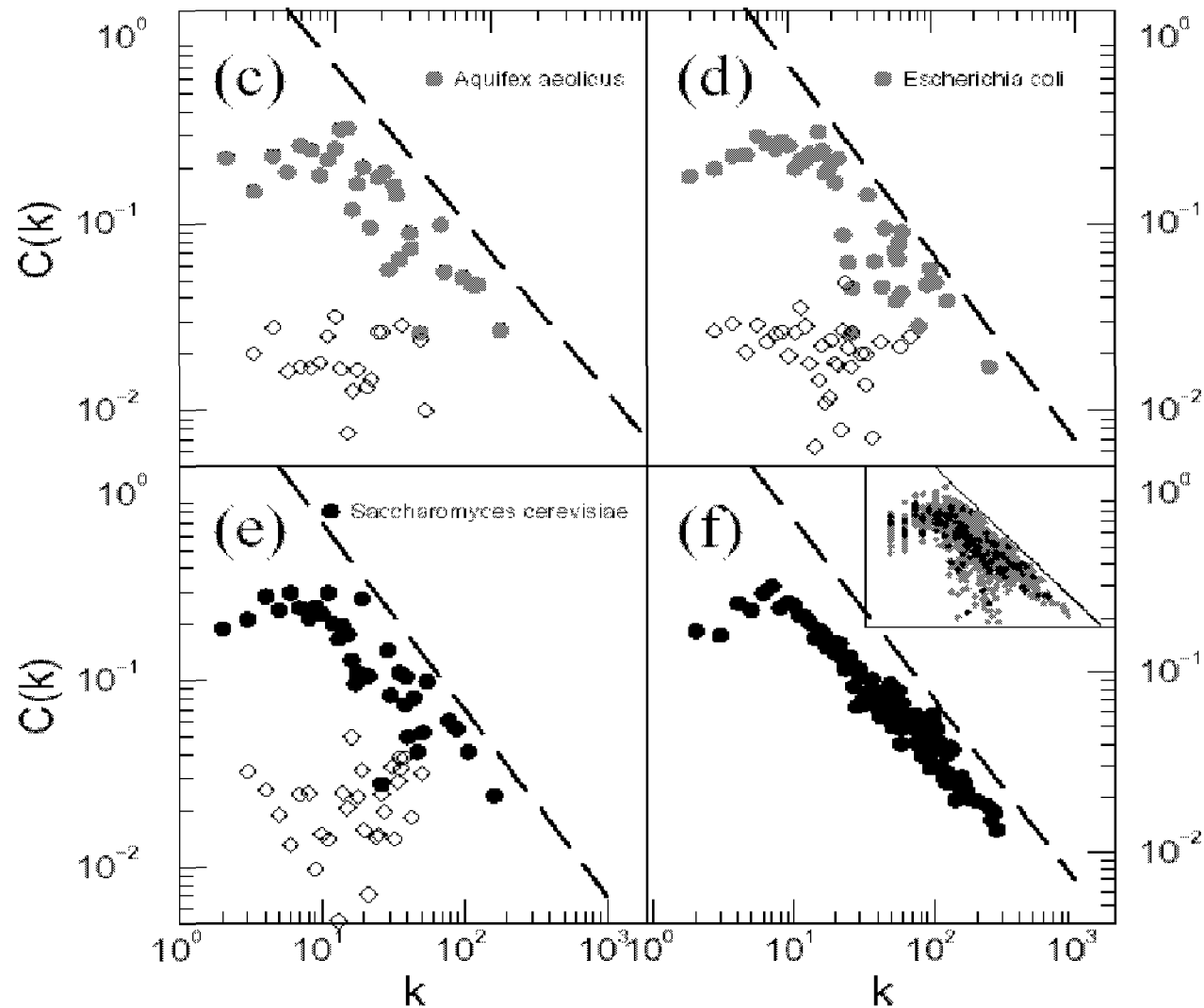
# Hierarchical Networks

## 3. Clustering coefficient depends on $k$

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



## Scaling of the clustering coefficient $C(k)$

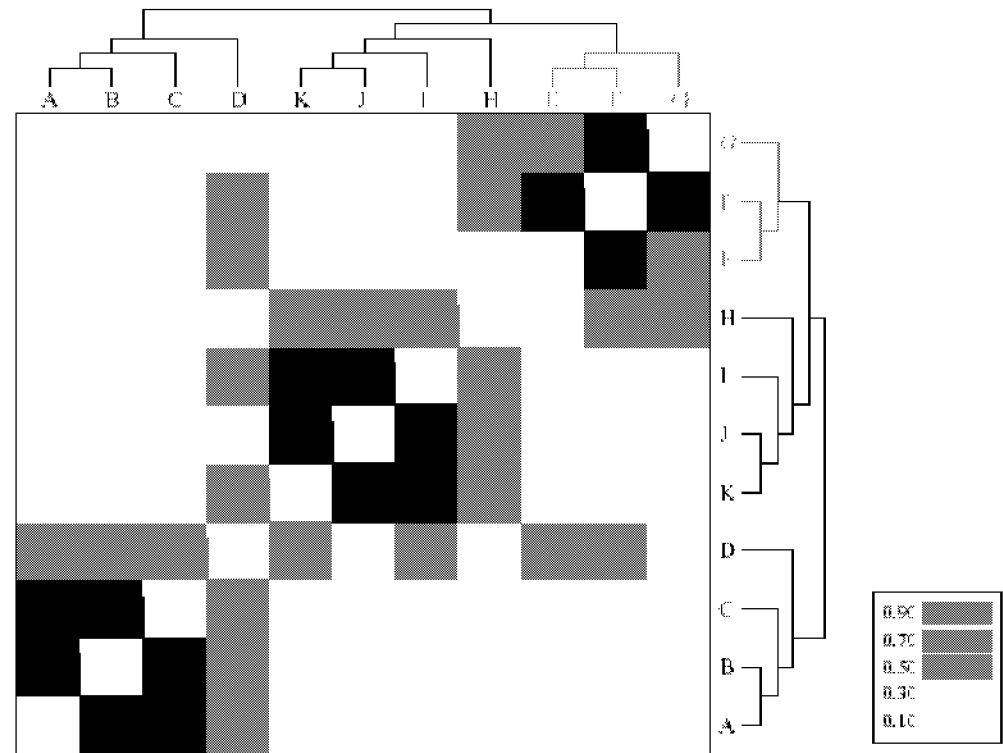
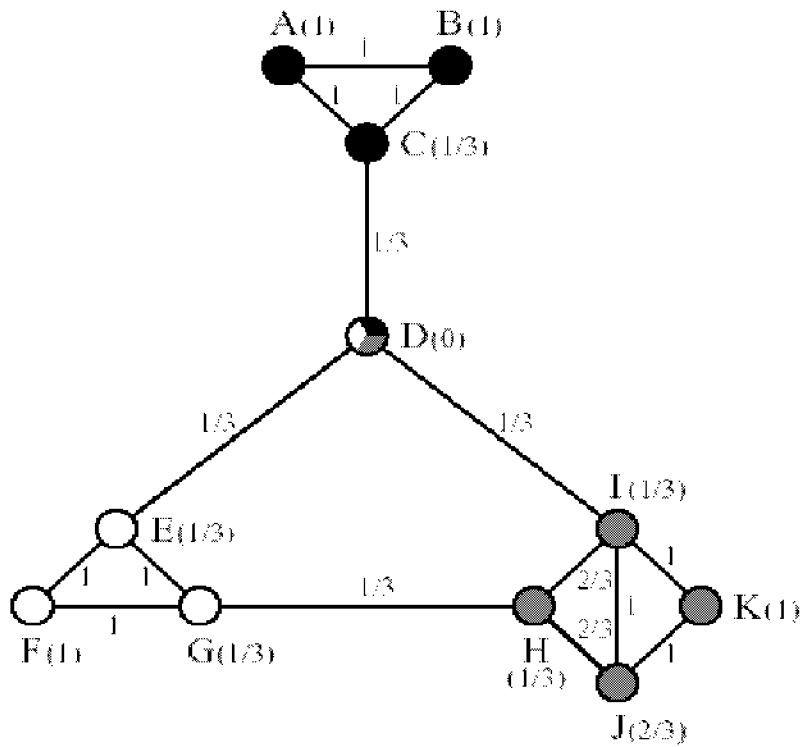


**The metabolism forms a hierarchical network.**

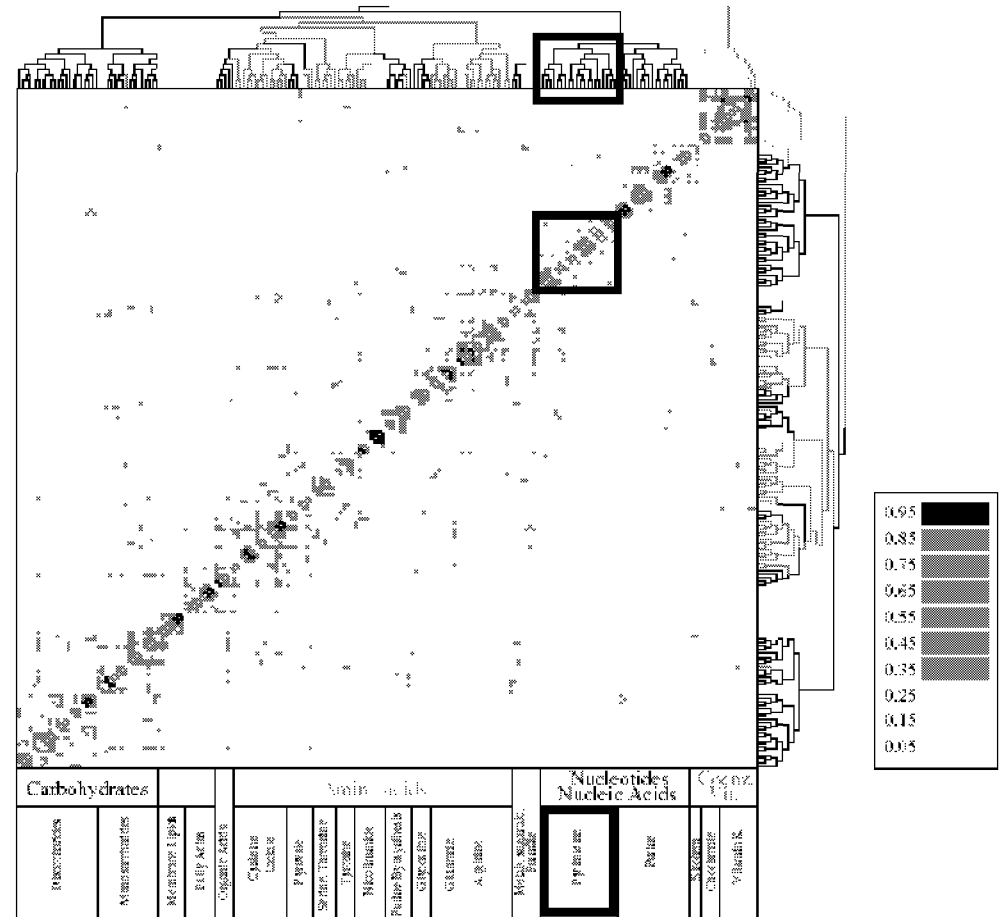
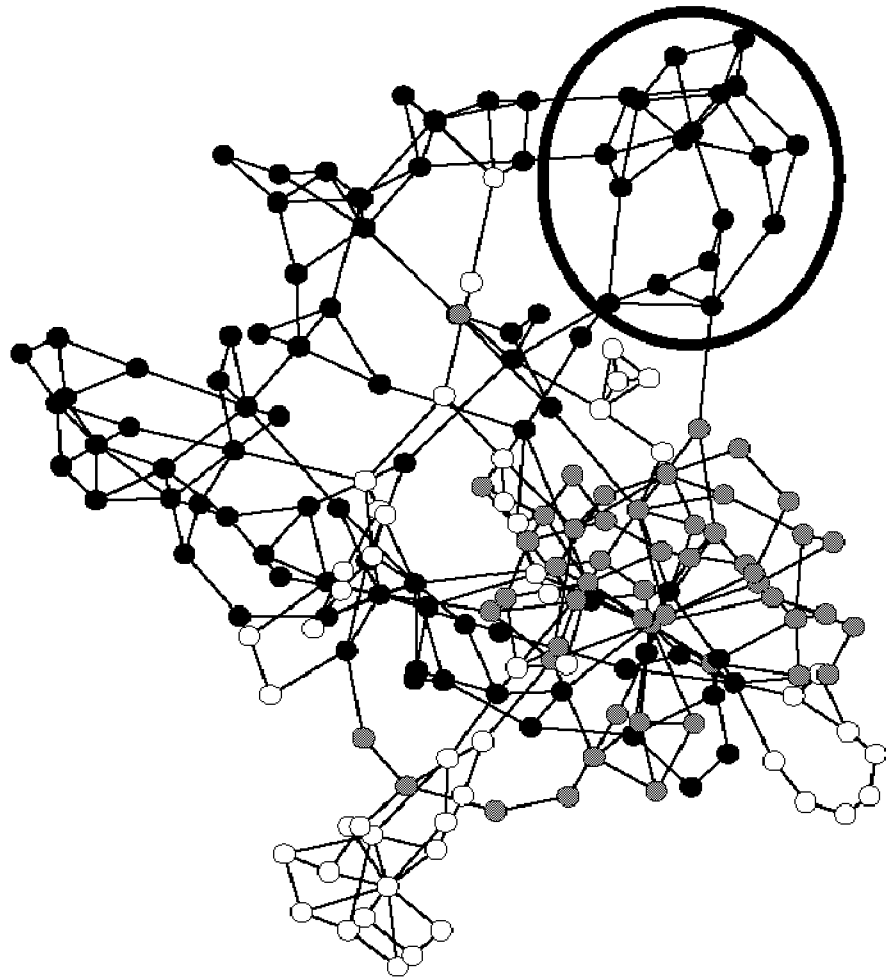
Ravasz, Somera, Mongru, Oltvai, A-L. B, *Science* **297**, 1551 (2002).



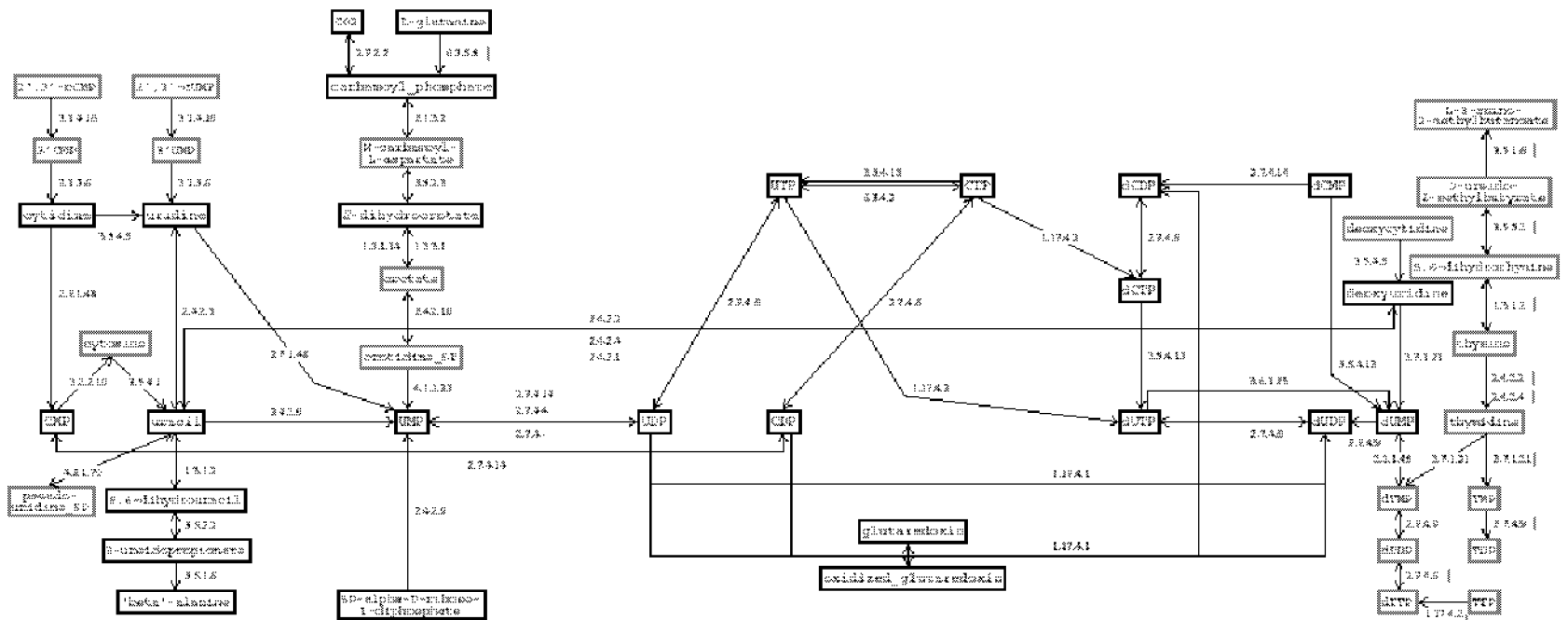
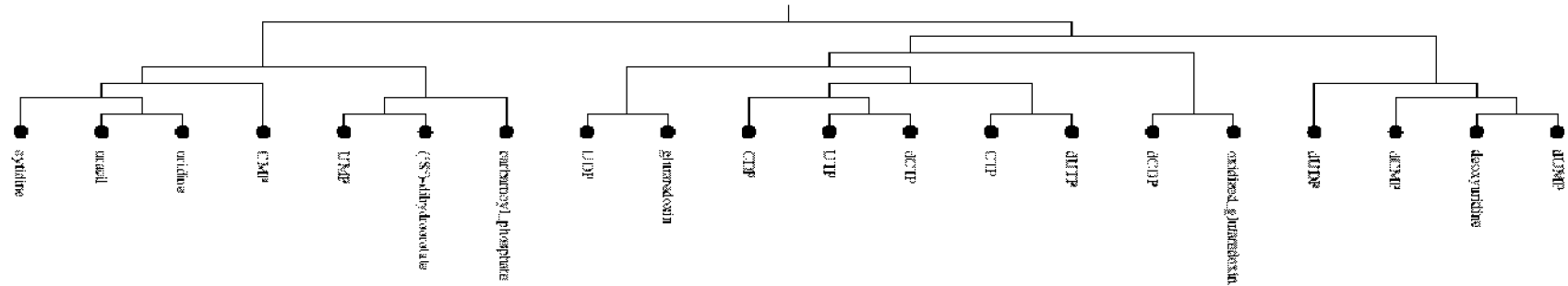
# Can we identify the modules?



# Modules in the *E. coli* metabolism



# The structure of pyrimidine metabolism

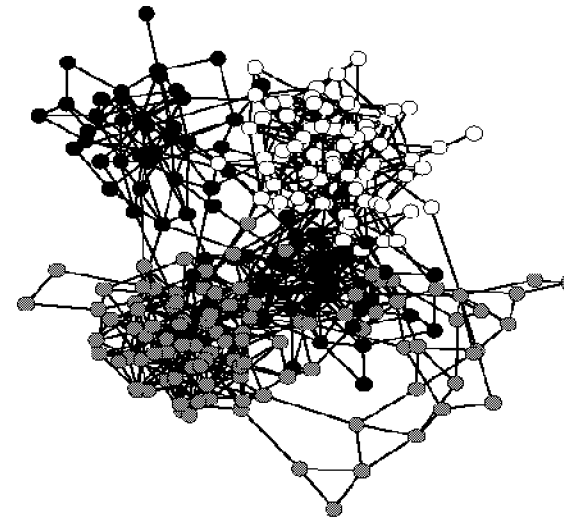
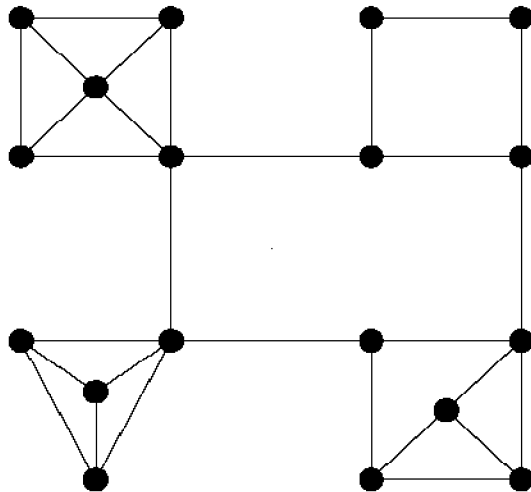


# Modularity

➤ High  $C$  → real networks are fragmented into group or modules

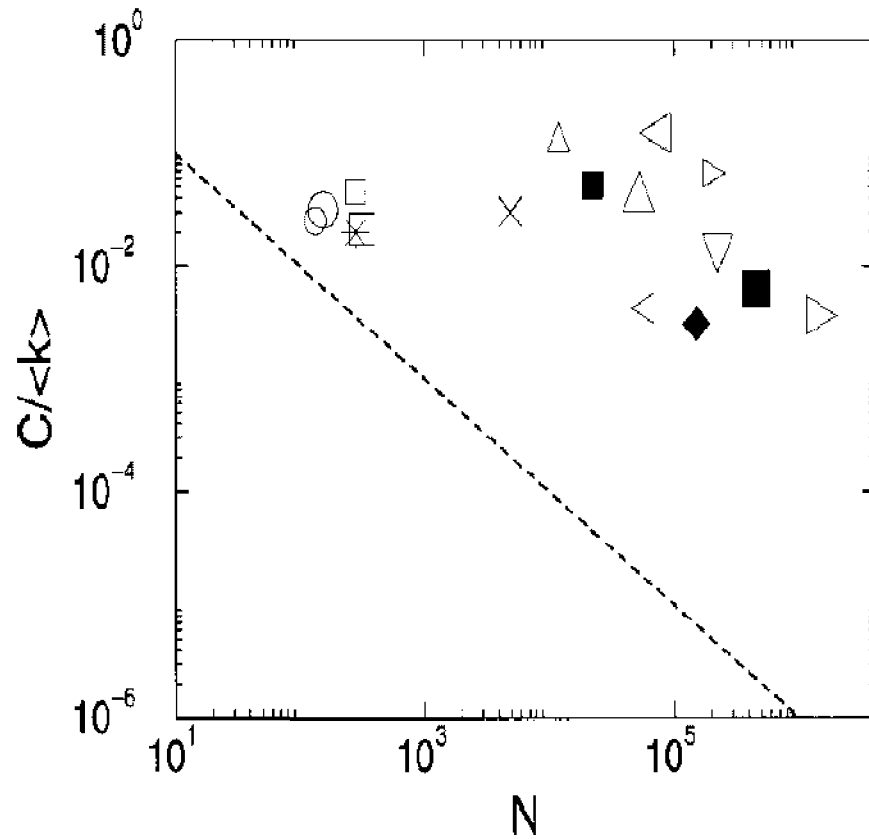
- ❖ **Society:** Granovetter, M. S. (1973) ; Girvan, M., & Newman, M.E.J. (2001); Watts, D. J., Dodds, P. S., & Newman, M. E. J. (2002).
- ❖ **WWW:** Flake, G. W., Lawrence, S., & Giles. C. L. (2000).
- ❖ **Biology:** Hartwell, L.-H., Hopfield, J. J., Leibler, S., & Murray, A. W. (1999).
- ❖ **Internet:** Vasquez, Pastor-Satorras, Vespignani(2001).

➤ Traditional view of modularity:

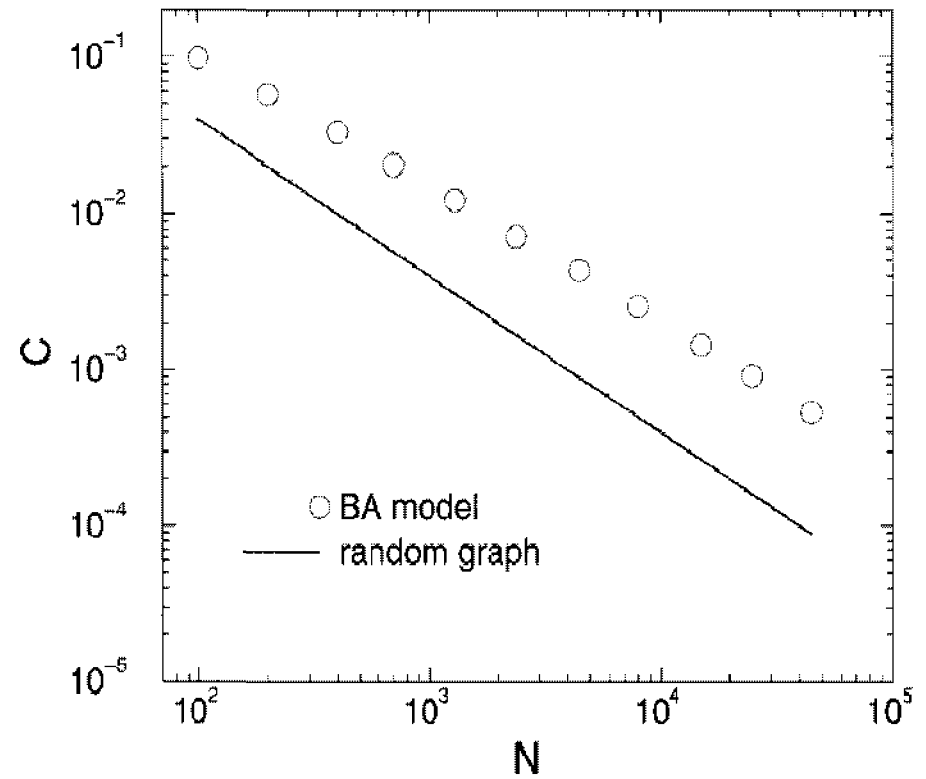


# Clustering in non-biological networks

**C is independent of N**

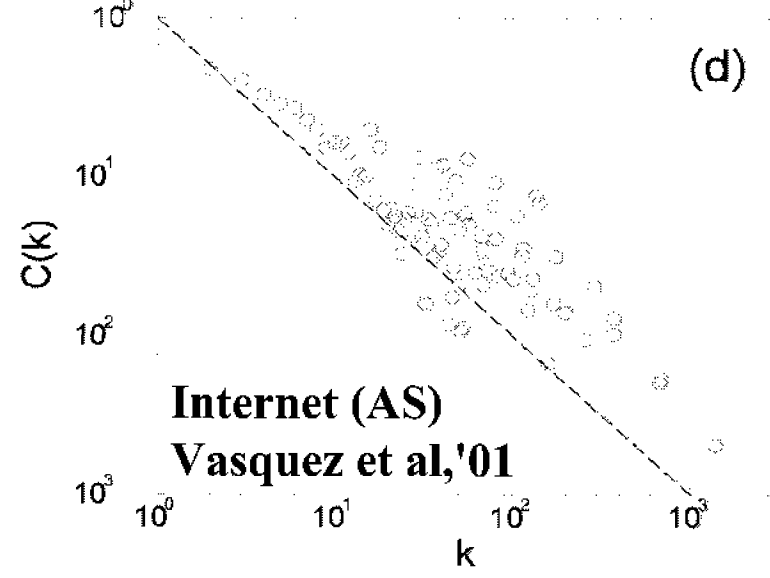
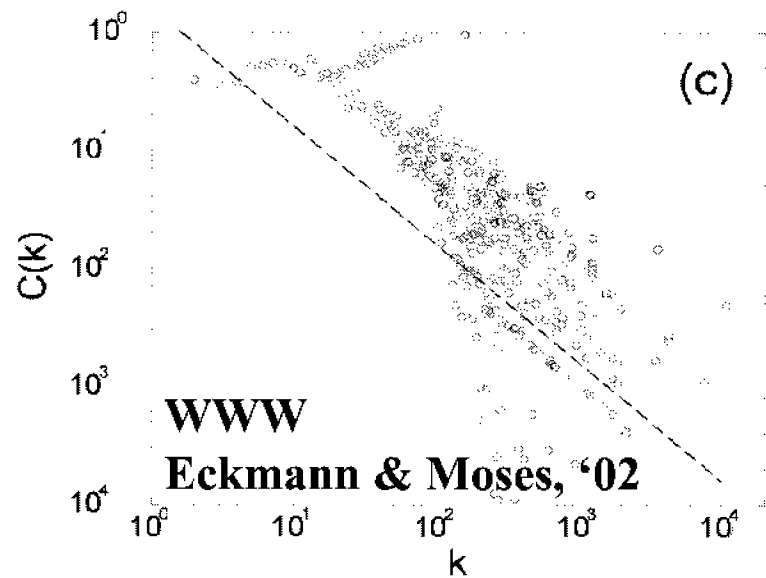
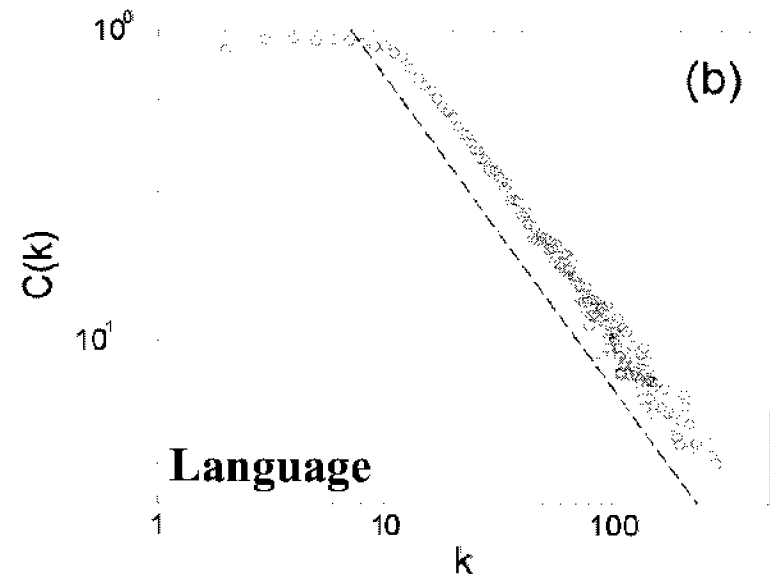
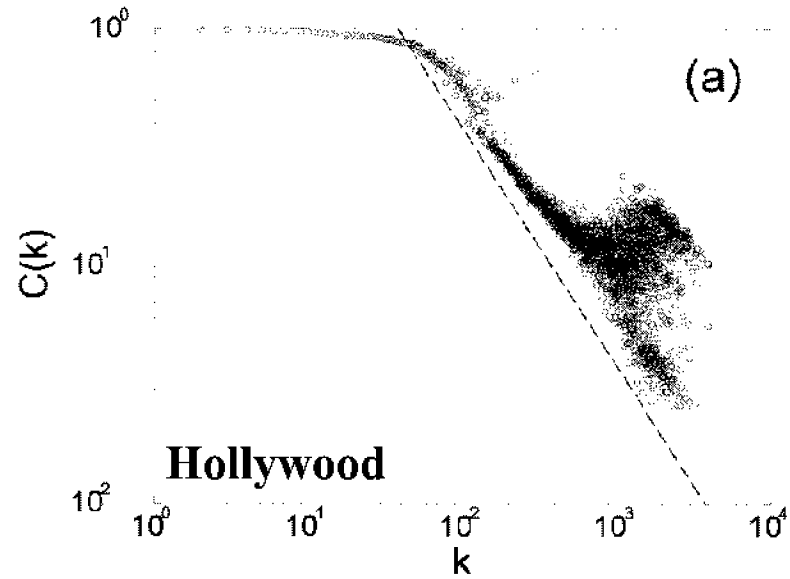


**C decreases with N**

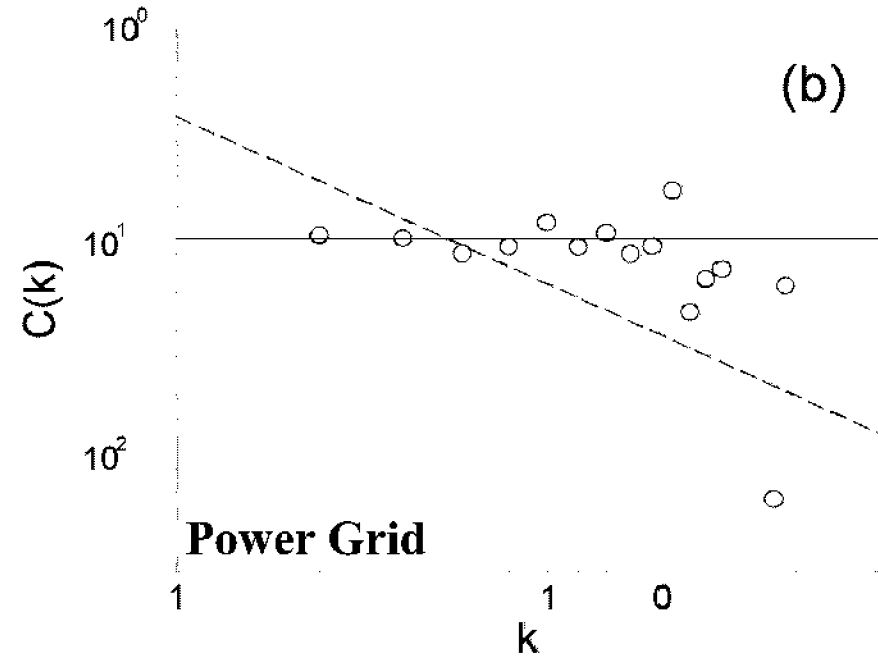
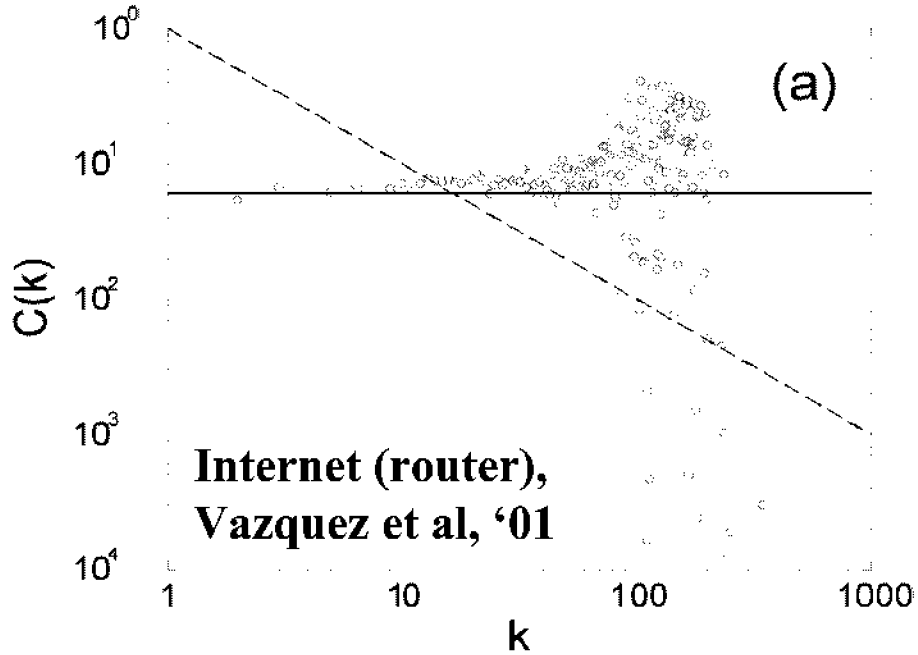


$$C_i = 2n_i/k_i(k_i-1)$$

# Real Networks



# Exceptions: Geographically Organized Networks:

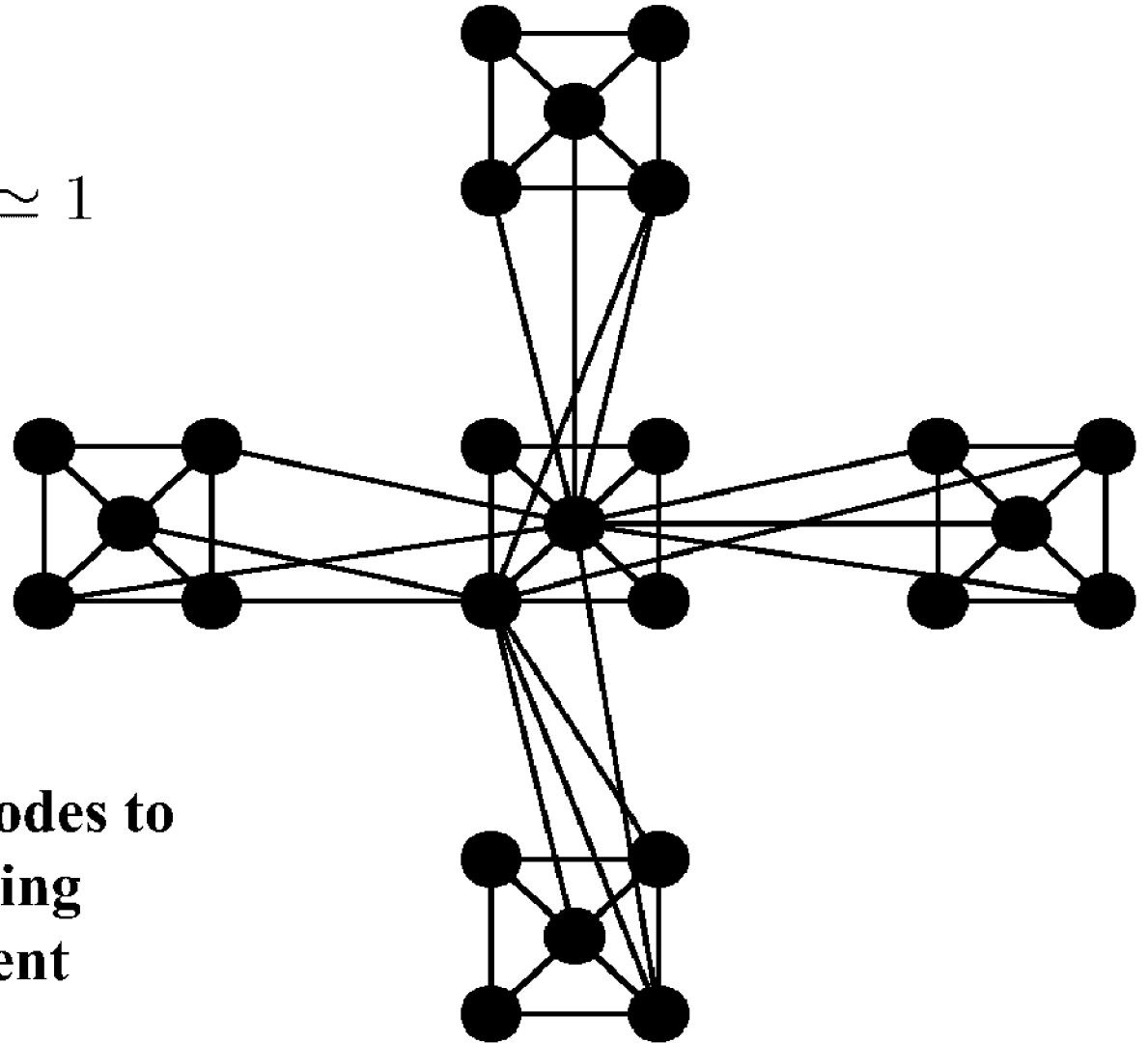


Common feature:

economic pressures towards shorter links

## Is the hierarchical exponent $\beta$ universal?

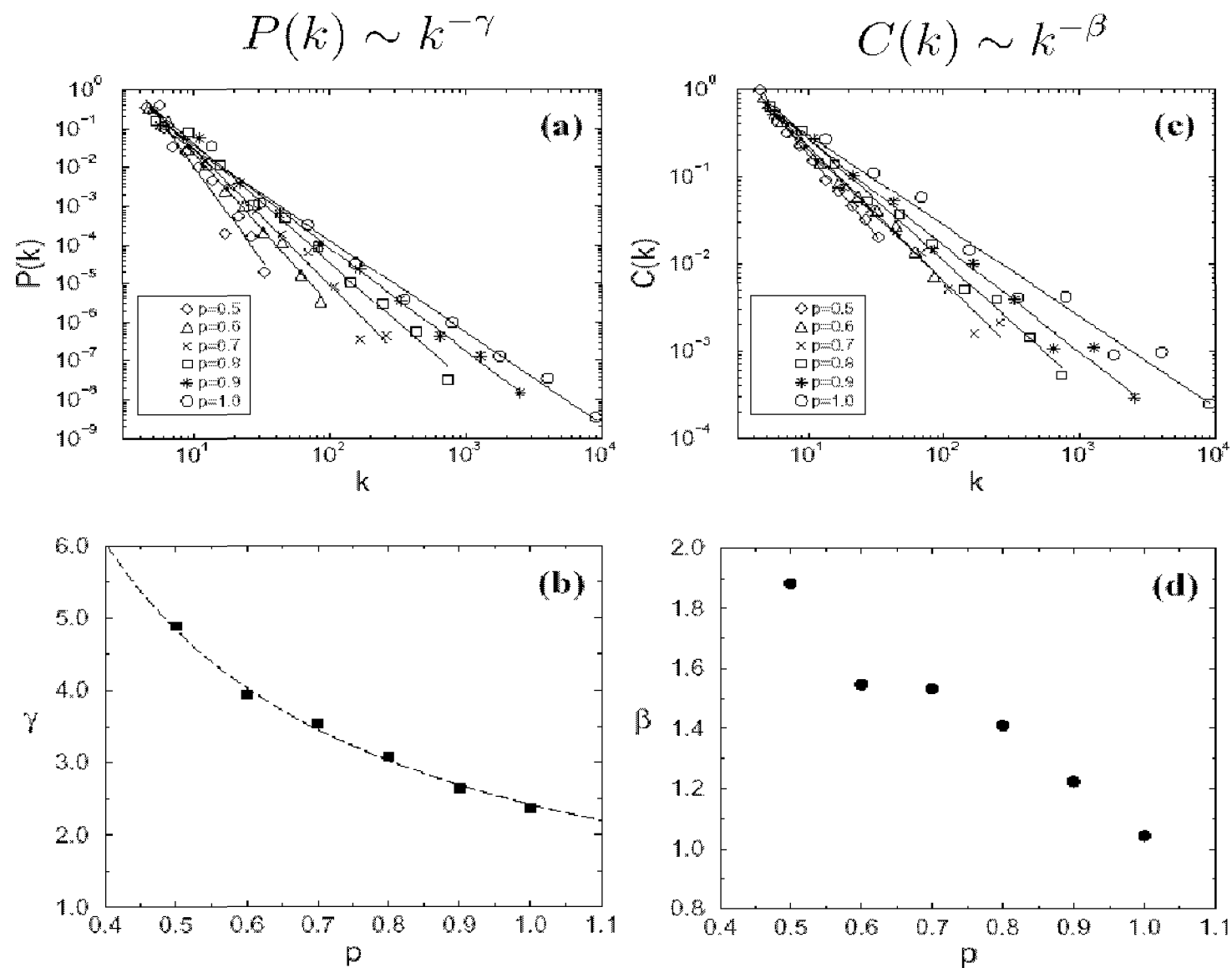
- ❖  $C(k) \sim k^{-\beta}$
- ❖ For most systems:  $\beta \simeq 1$



**Connect a  $p$  fraction of nodes to  
the central module using  
preferential attachment**



# Stochastic Hierarchical Model



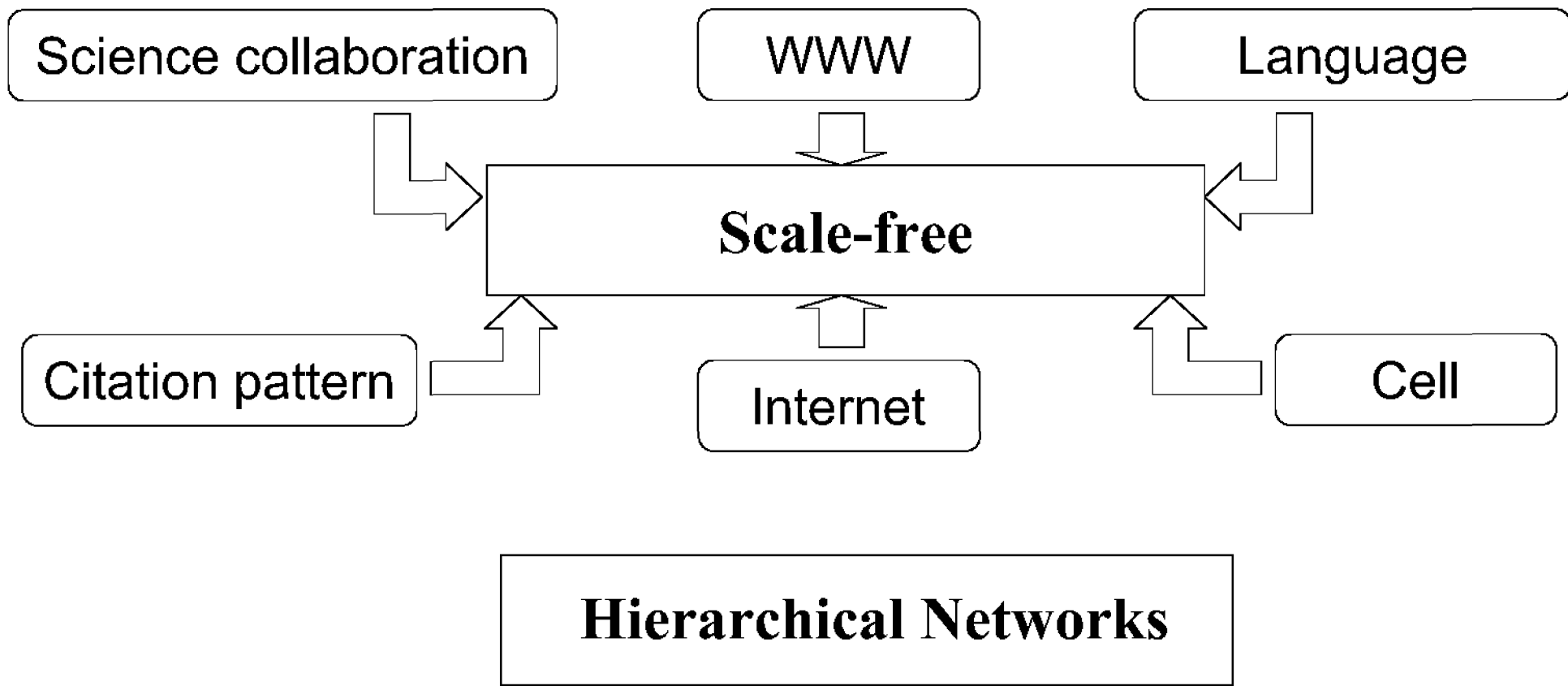
# Is hierarchy present in network models?

## **NO:**

- Scale-free model (albert & Albert, 1999)
- Erdos-Renyi model (1959)
- Watts-Strogatz (1998)

## **YES:**

- Dorogovtsev, Goltsev, Mendes, 2001 (determ.)
- Klemm and Eguiluz, 2002
- Vasquez, Pastor-Satorras, Vespignani (2001)\*  
⇒ Bianconi & albert (fitness model) (2001)



**Scale-free**

$$P(k) \sim k^{-\alpha}$$

**Modular**

$$C(N) = \text{const.}$$

**Hierarchical**

$$C(k) \sim k^{-\beta}$$

# Traditional modeling: Network as a static graph

Given a network with  $N$  nodes and  $L$  links



Create a graph with statistically identical topology

RESULT: model the static network topology

**PROBLEM: Real networks are dynamical systems!**

## Evolving networks

OBJECTIVE: capture the network dynamics

METHOD : 

- identify the processes that contribute to the network topology
- develop dynamical models that capture these processes



**BONUS: get the topology correctly.**

# Bonus: Why Kevin Bacon?

Measure the average distance between Kevin Bacon and all other actors.

Kevin Bacon

No. of movies : 46      No. of actors : 1811

Average separation: 2.79

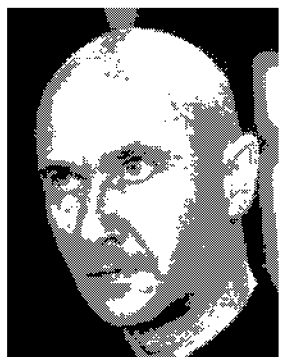
*Is Kevin Bacon the  
most connected  
actor?*

***NO!***

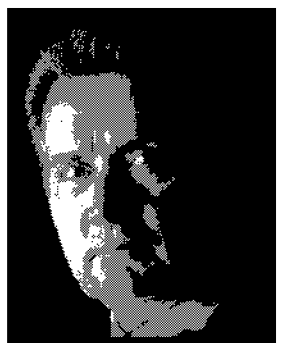
Rank	Name	Average distance	# of movies	# of links
1	Rod Steiger	2.537527	112	2562
2	Donald Pleasence	2.542376	180	2874
3	Martin Sheen	2.551210	136	3501
4	Christopher Lee	2.552497	201	2993
5	Robert Mitchum	2.557181	136	2905
6	Charlton Heston	2.566284	104	2552
7	Eddie Albert	2.567036	112	3333
8	Robert Vaughn	2.570193	126	2761
9	Donald Sutherland	2.577880	107	2865
10	John Gielgud	2.578980	122	2942
11	Anthony Quinn	2.579750	146	2978
12	James Earl Jones	2.584440	112	3787
...				
<b>876</b>	<b>Kevin Bacon</b>	<b>2.786981</b>	<b>46</b>	<b>1811</b>
...				



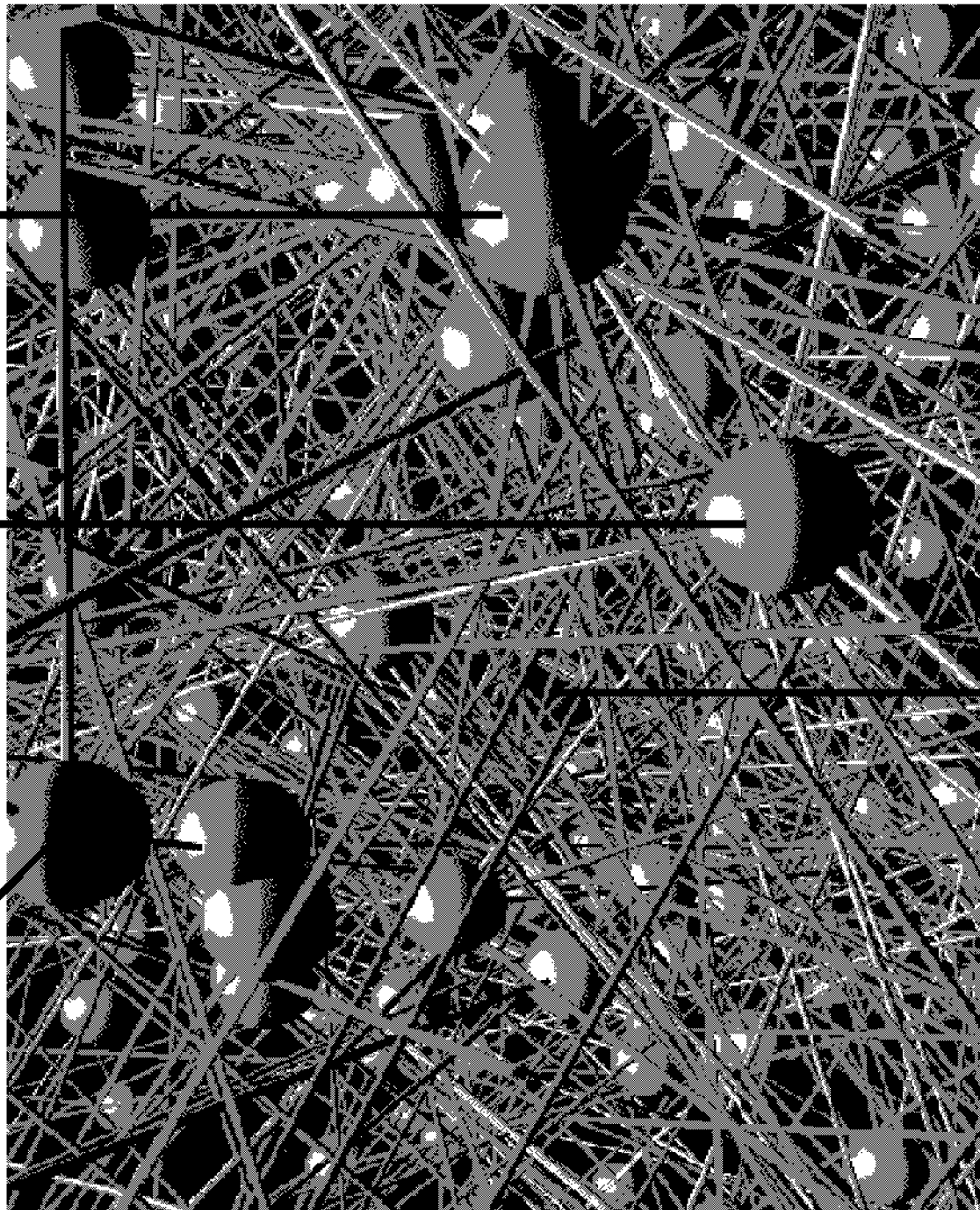
#1 Rod Steiger



#2 Donald Pleasence



#3 Martin Sheen



#876  
Kevin Bacon



**<http://www.nd.edu/~networks>**

# References

- R. Albert, H. Jeong, A.L. Barabasi, Nature **401** 130 (1999).
- R. Albert, A.L. Barabasi, Science **286** 509 (1999).
- A.L. Barabási, R. Albert and H. Jeong, Physica A **272**, 173 (1999)
- R. Albert, H. Jeong, A.L. Barabasi, Nature **406** 378 (2000).
- H.Jeong, B.Tombor, R.Albert, Z.N.Oltvai, A.L.Barabasi, Nature **407** 651 (2000).
- H. Jeong, S.P. Mason, A.L. Barabasi, Z.N. Oltvai, Nature (in press).

**URL: <http://www.nd.edu/~networks>**



<http://www.nd.edu/~networks>

**Complex Networks &  
Biological Applications**

**Thursday**

**Room 102**

**10:30—18:30**

**Mark Newman**

**Andrea Levchenko**

**Lou Pecora**

**Gabor Forgacs**

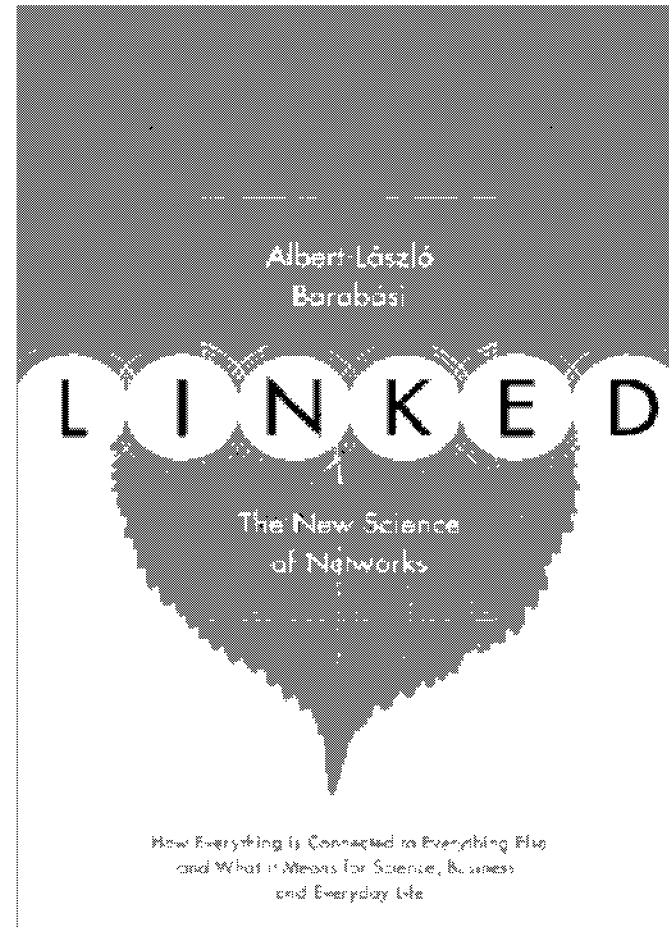
**Allesandro Vespignani**

**Reka Albert**

**Gilberto Thomas**

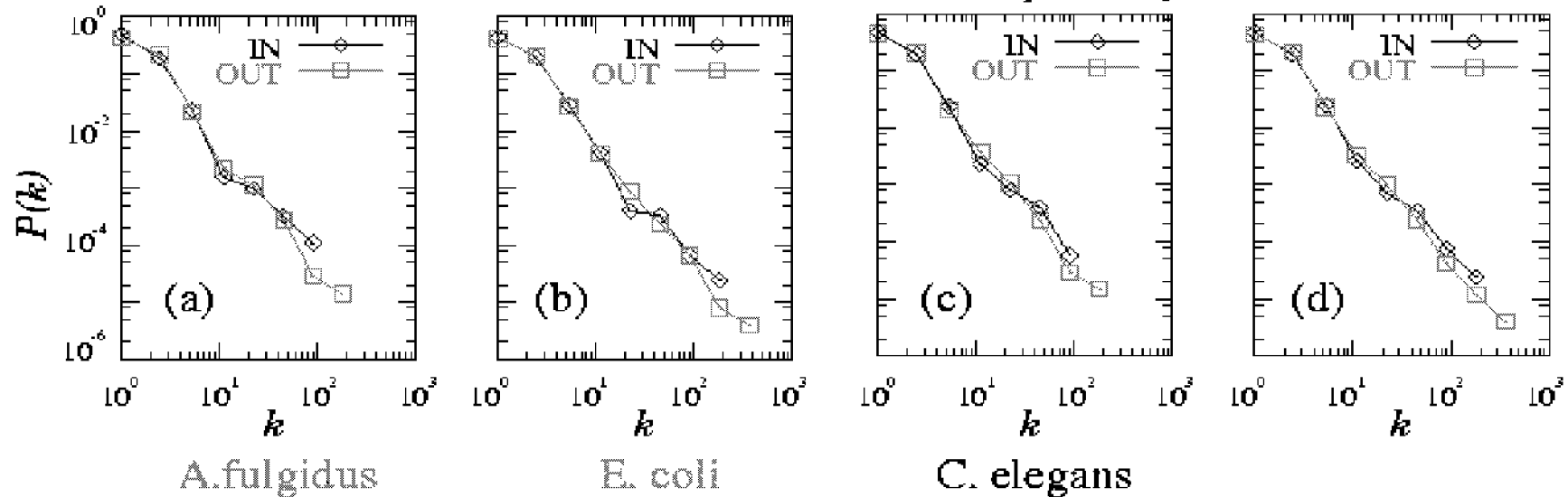
**Byungnam Kahng**

**Hawoong Jeong**



# Whole cellular network

## Metabolic and non-metabolic pathways



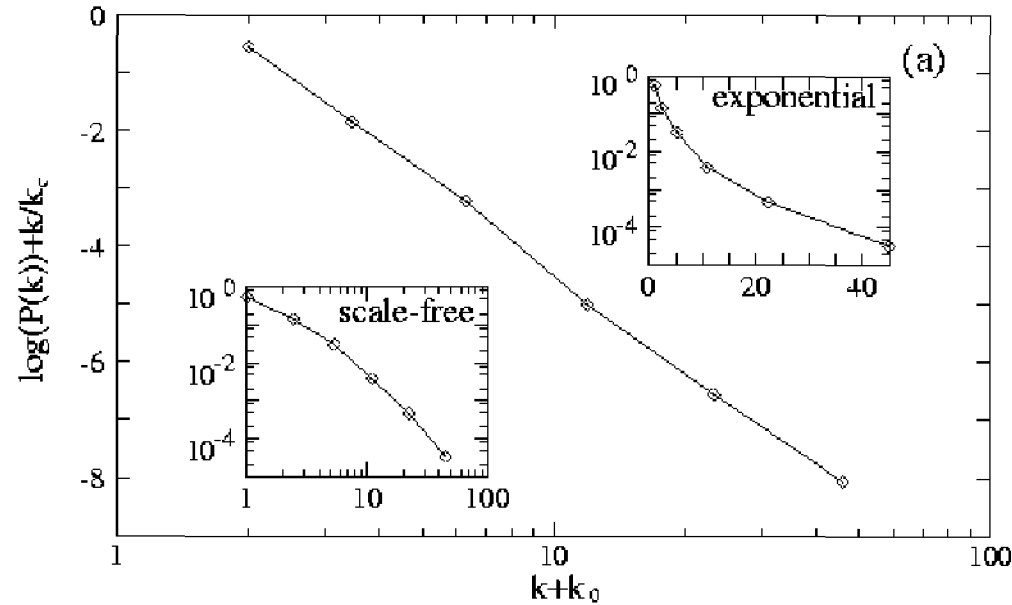
Metabolic pathways

intermediate metabolism  
bioenergetics

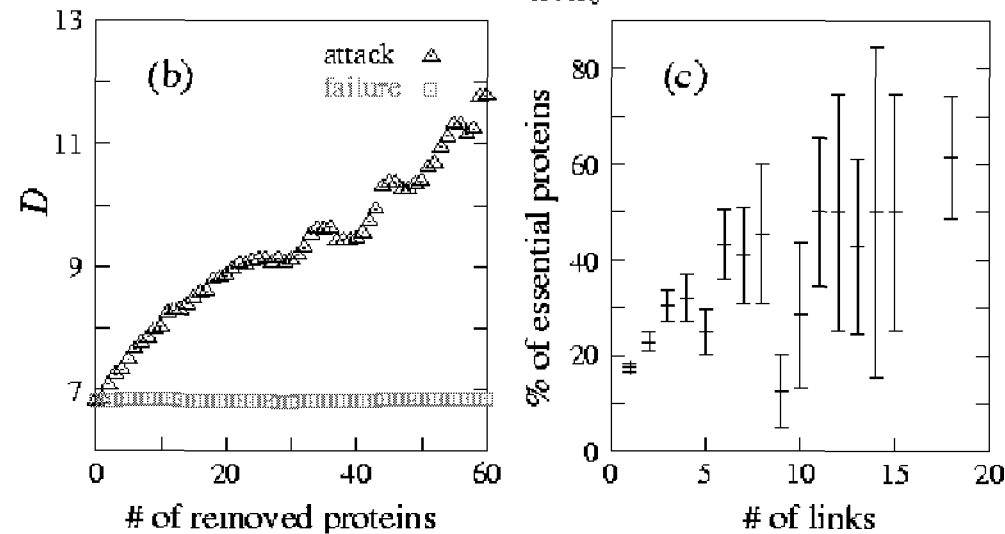
Non-metabolic pathways

information pathways  
electron, transmembrane transport  
signal transduction  
structure and function of the cell

# Properties of the protein network



$$P(k) \sim (k + k_0)^{-\gamma} \exp\left(-\frac{k + k_0}{k_\tau}\right)$$

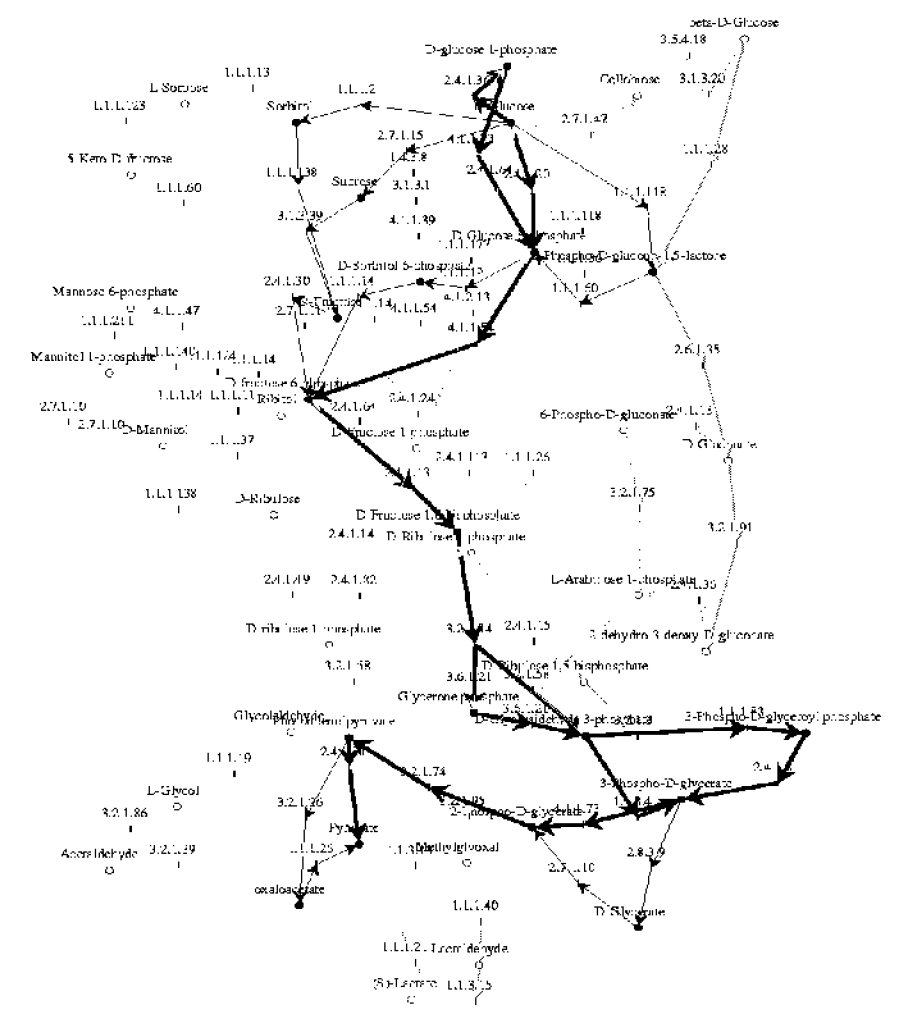
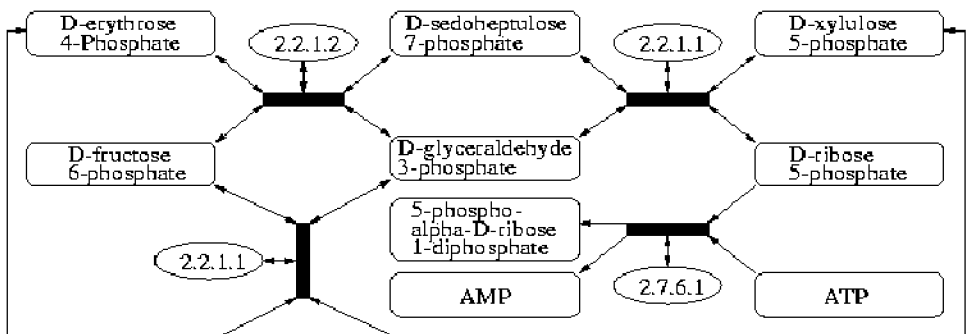
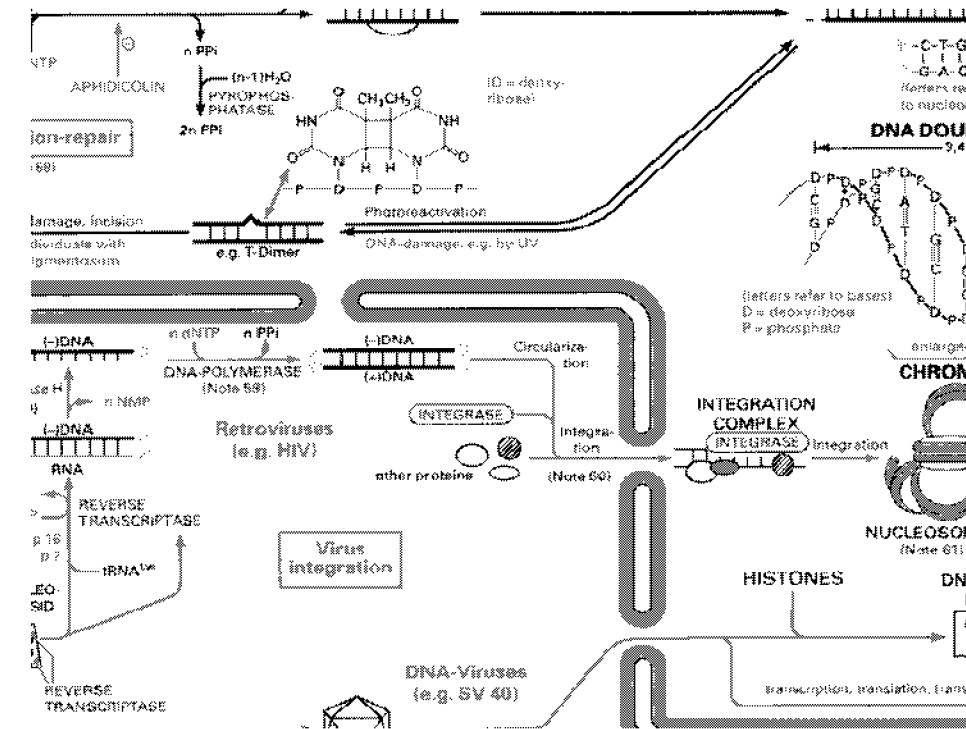


Highly connected proteins are more **essential (lethal)** than less connected proteins.

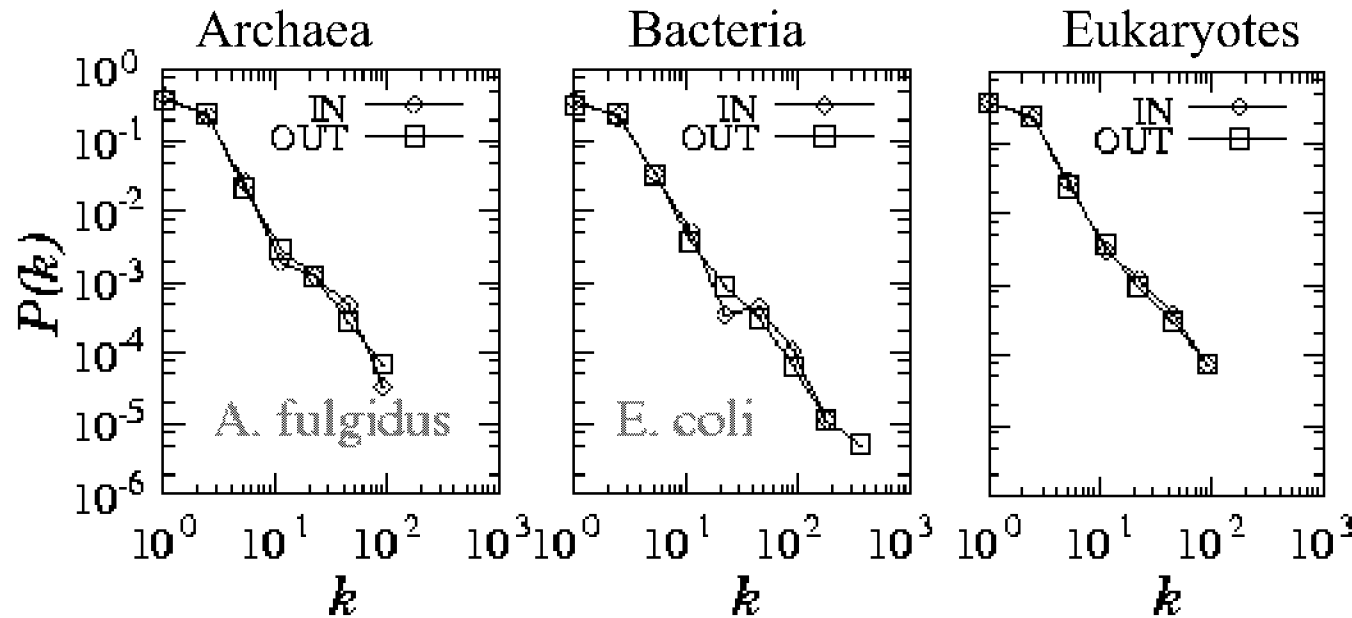
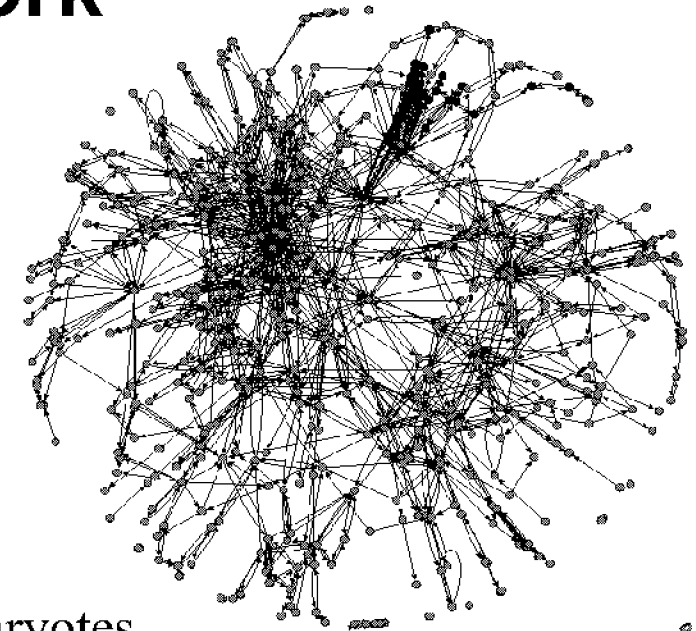
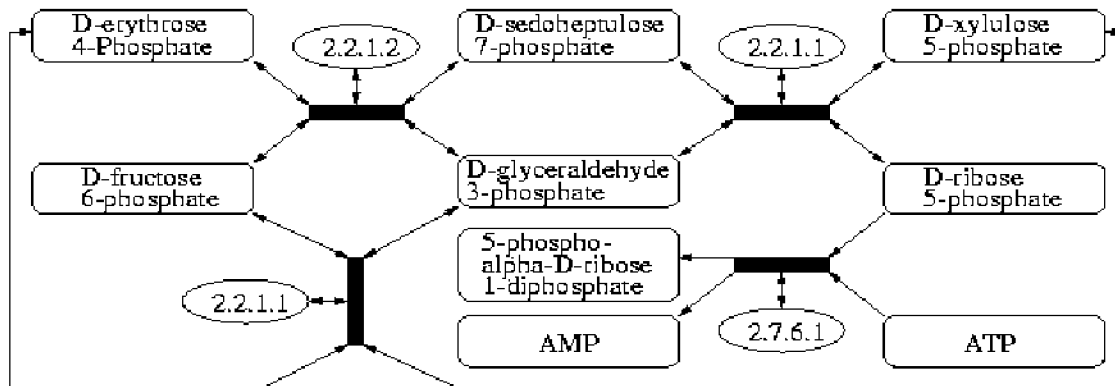
# Metabolic Network

**Nodes:** chemicals  
(substrates)

**Links:** chem. reaction

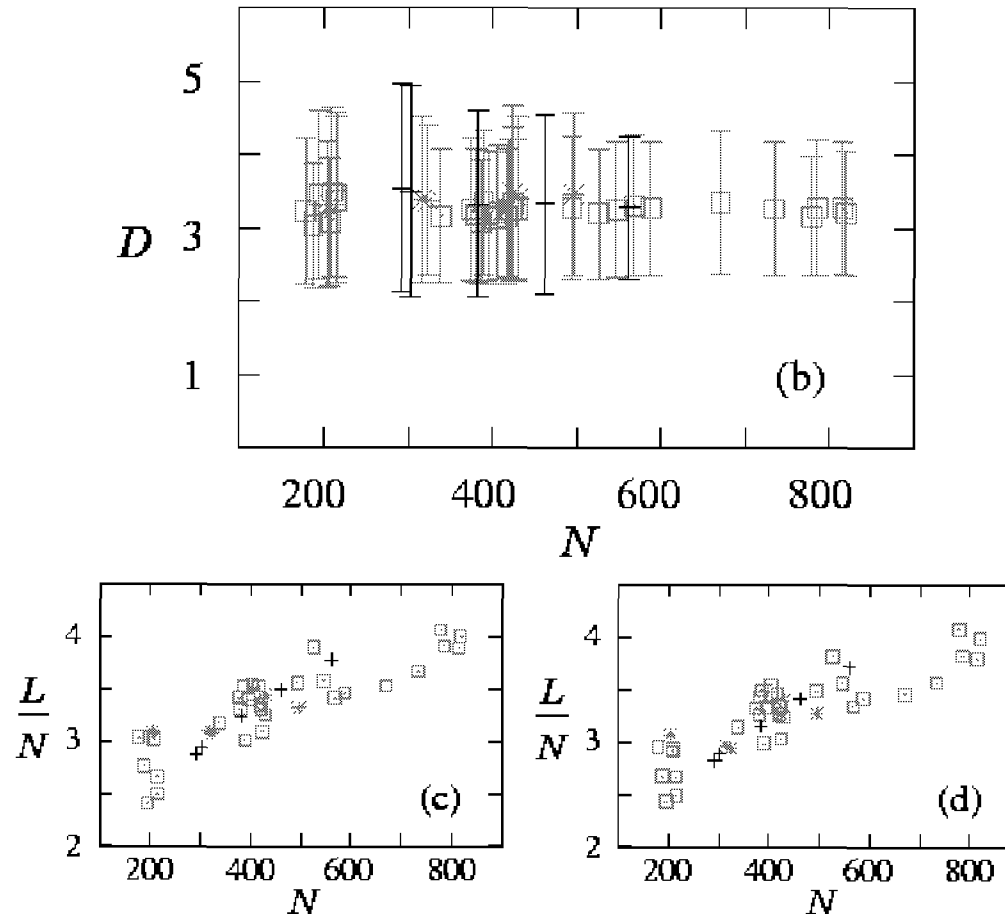


# Metabolic network



Organisms from all three domains of life are **scale-free** networks!

# Properties of metabolic networks



**Average distances are independent of organisms!**

⇐ by making more links between nodes.

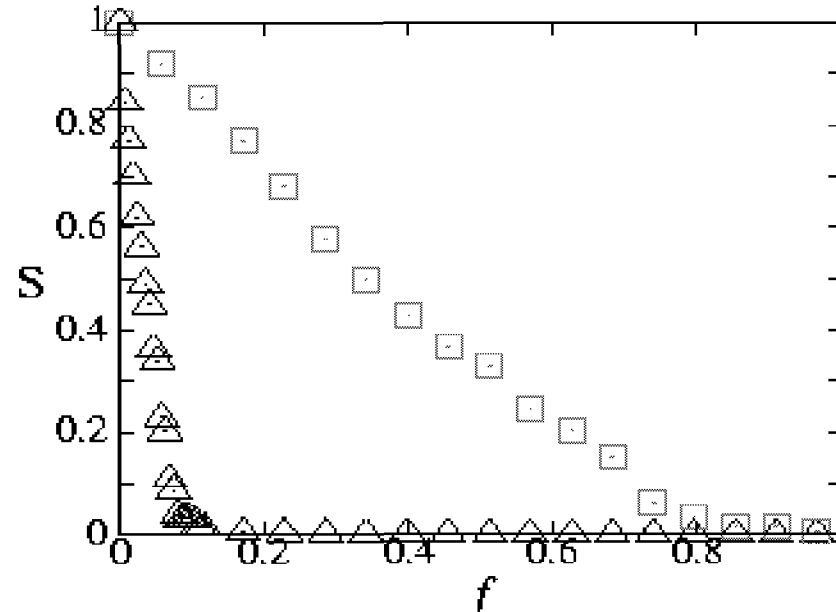
⇐ based on “**design principles**” of the cell through **evolution**.

**cf. Other scale-free network:  $D \sim \log(N)$**

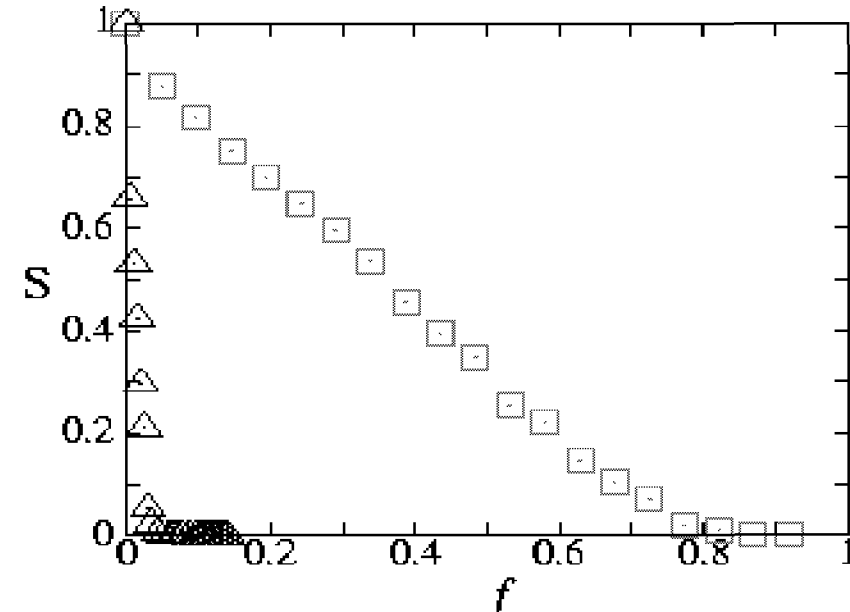
# Achilles' Heel of complex network

— failure  
— attack

## Internet

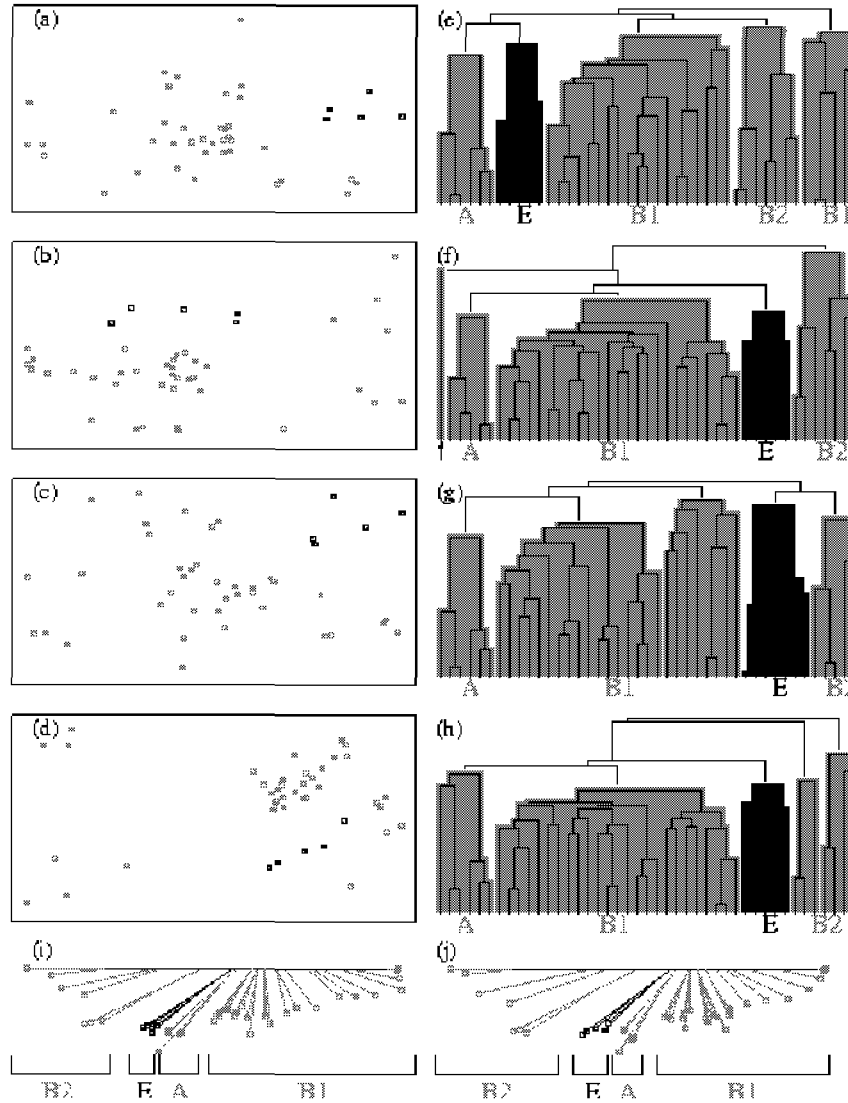


## Protein network



R. Albert, H. Jeong, A.L. Barabasi, Nature **406** 378 (2000)

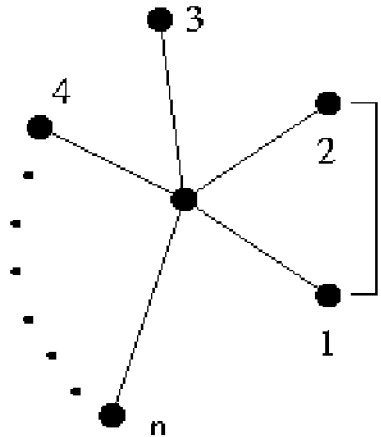
# Taxonomy using networks





# Watts-Strogatz

**Clustering:** My friends will know each other with high probability!

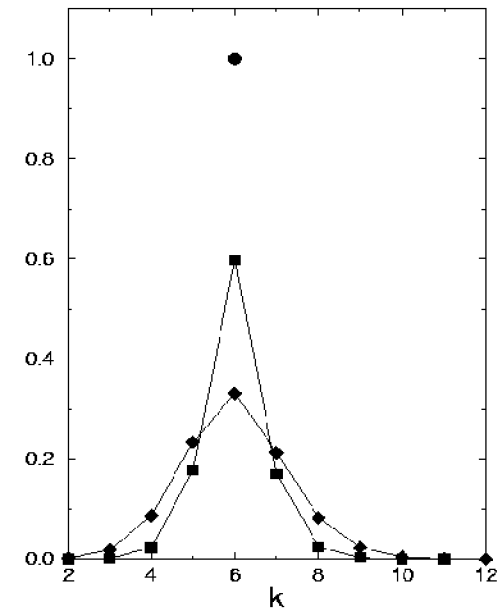
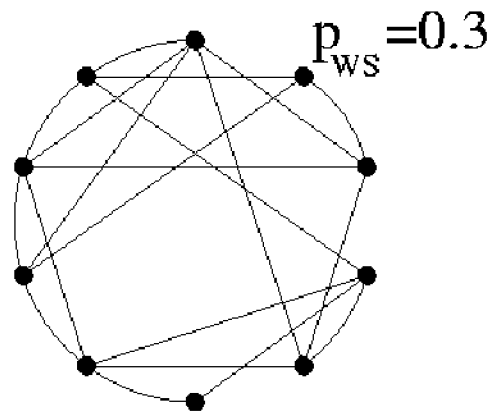
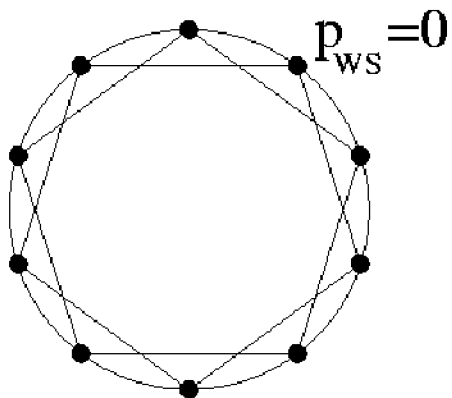


Probability to be connected  $C \gg p$

$$C = \frac{\text{\# of links between } 1, 2, \dots, n \text{ neighbors}}{n(n-1)/2}$$

**N nodes forms a regular lattice.**

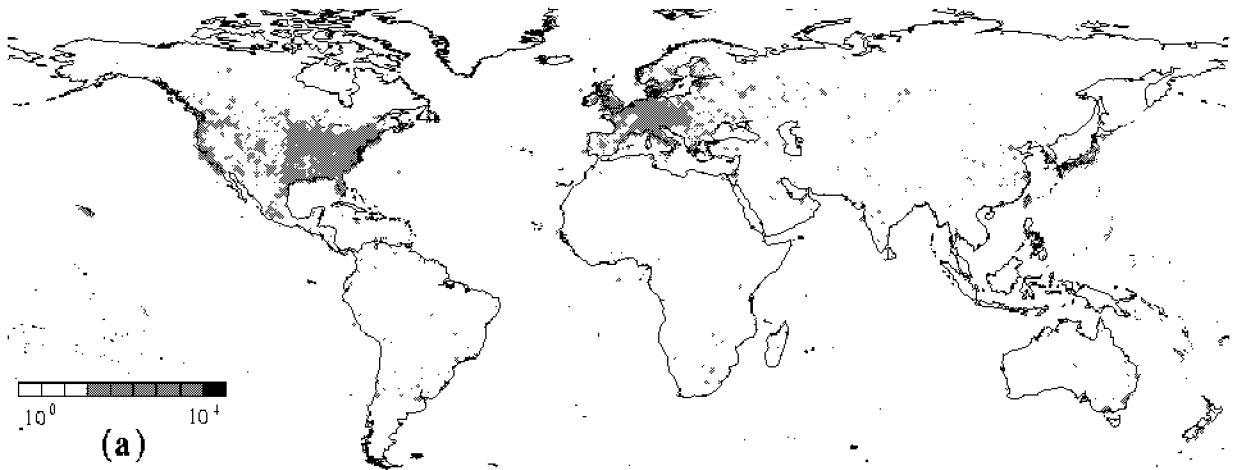
**With probability p, each edge is rewired randomly.**



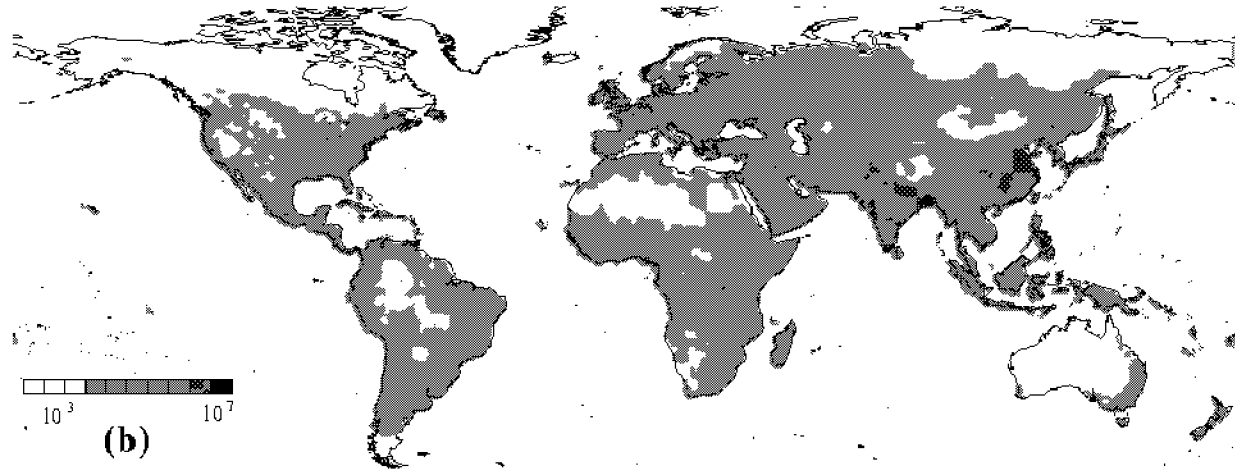
(Nature **393**, 440 (1998))

# Spatial Distributions

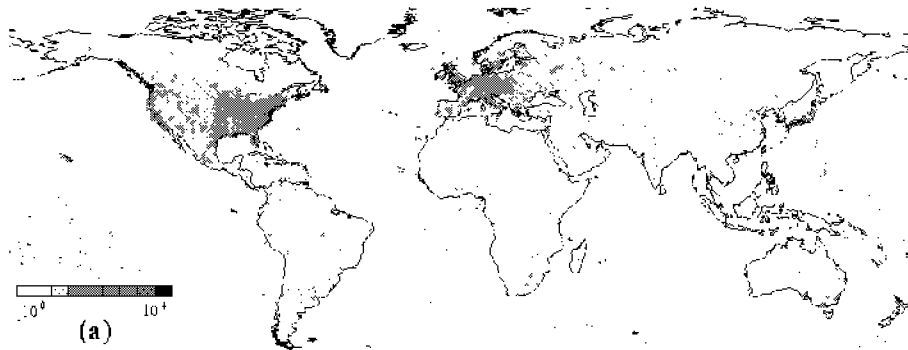
Router density



Population density



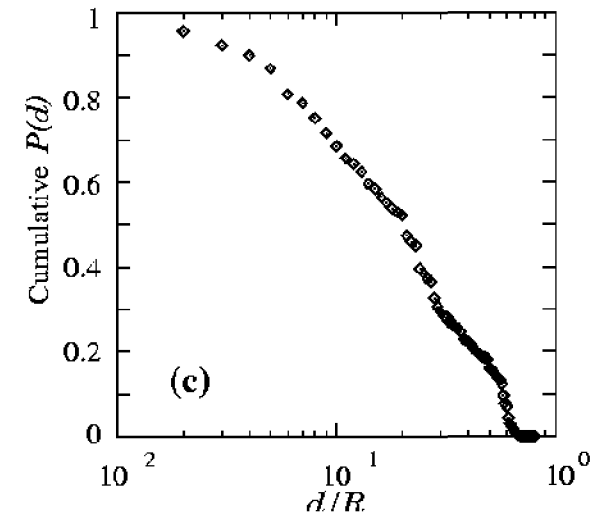
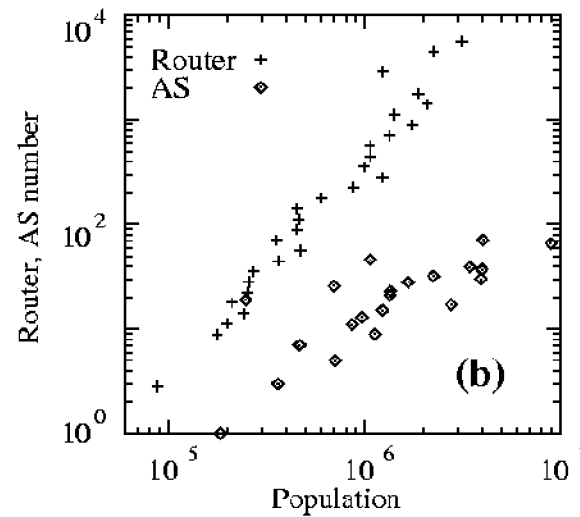
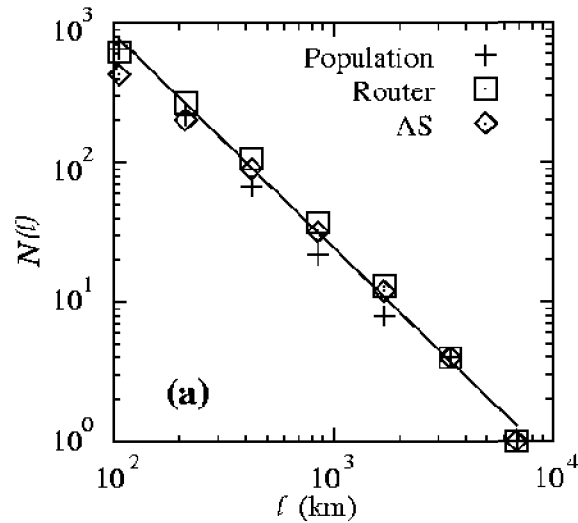
# Spatial Distribution of Routers



## Fractal set

Box counting:  $N(\ell) \equiv$  No. of boxes of size  $\ell$  that contain routers

$$N(\ell) \sim \ell^{-D_f} \quad D_f = 1.5$$

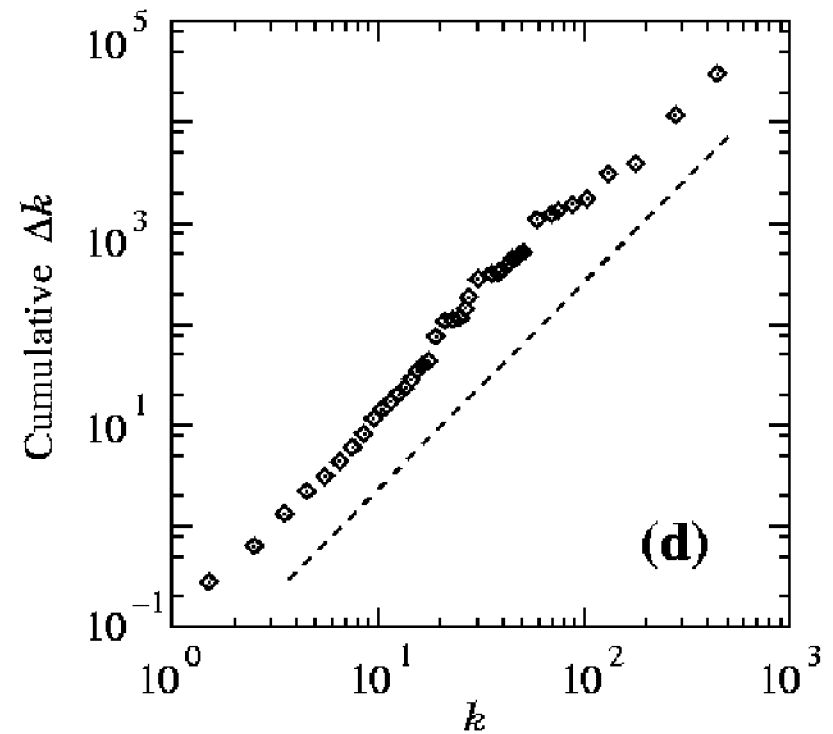


# Preferential Attachment

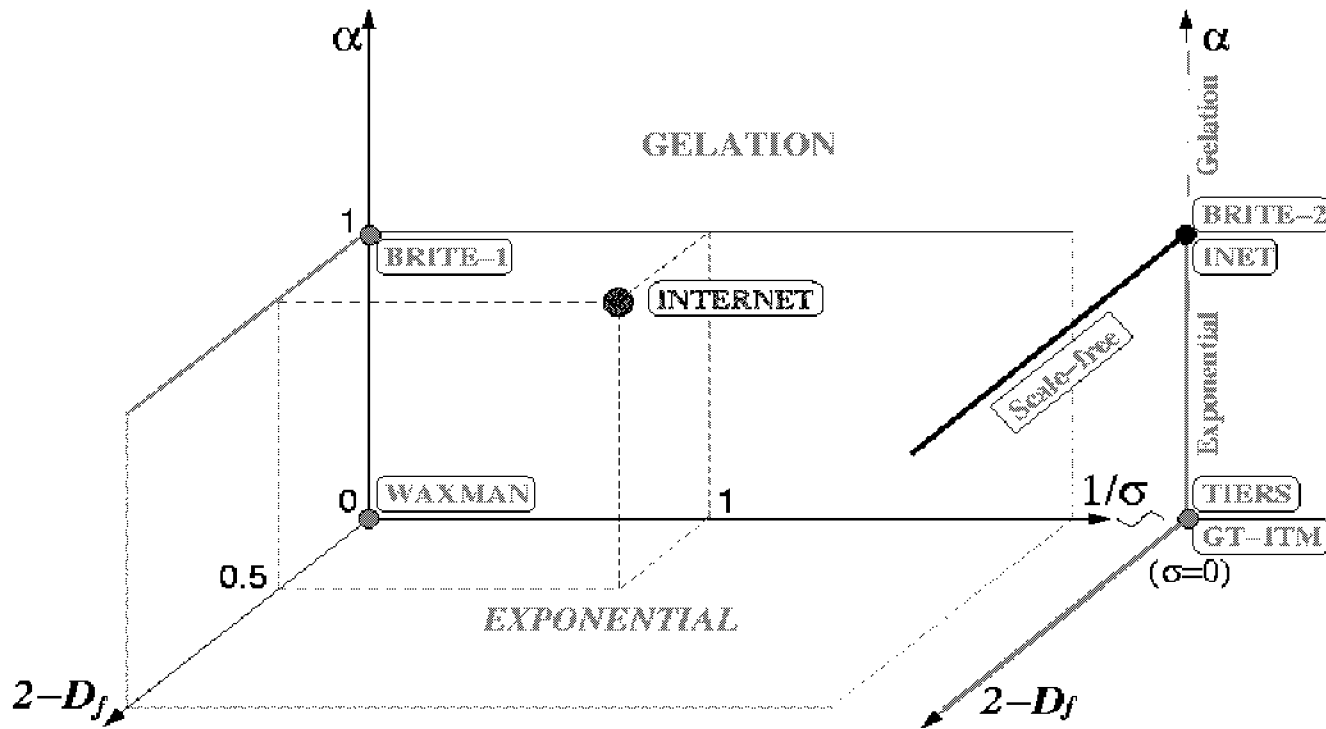
- Compare maps taken at different times ( $\Delta t = 6$  months)
- Measure  $\Delta k(k)$ , increase in No. of links for a node with  $k$  links

Preferential Attachment:

$$\Delta k(k) \sim k$$



# INTERNET



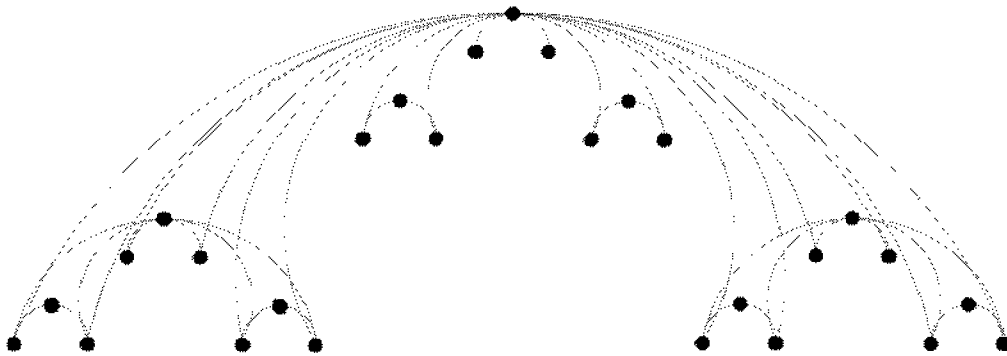
$$N(\ell) \sim \ell^{-D_f} \quad D_f=1.5$$

$$\Delta k(k) \sim k^\alpha \quad \alpha=1$$

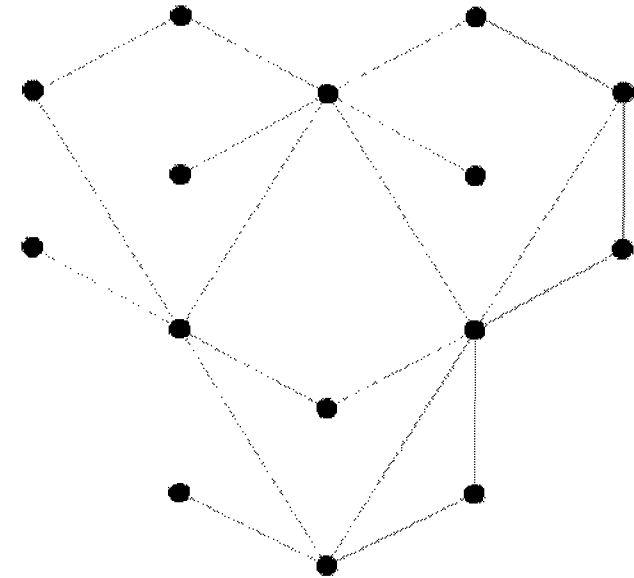
$$P(d) \sim d^{-\sigma} \quad \sigma=1$$

# Combining Modularity and the Scale-free Property

## Deterministic Scale-Free Networks

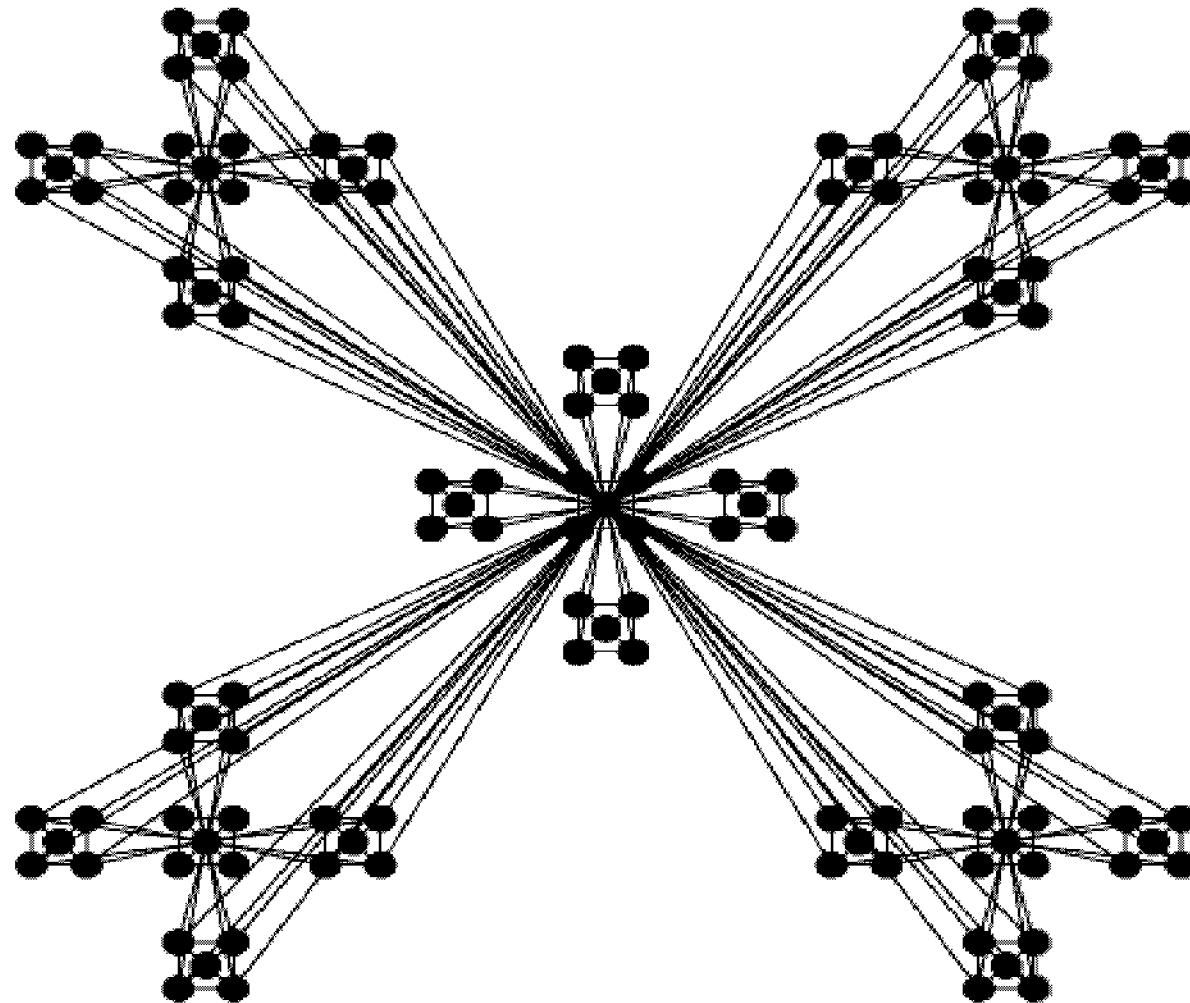


Barabási, A.-L., Ravasz, E., & Vicsek, T.  
(2001) *Physica A* **299**, 559.



Dorogovtsev, S. N., Goltsev, A. V., &  
Mendes, J. F. F. (2001) cond-mat/0112143.  
(DGM)

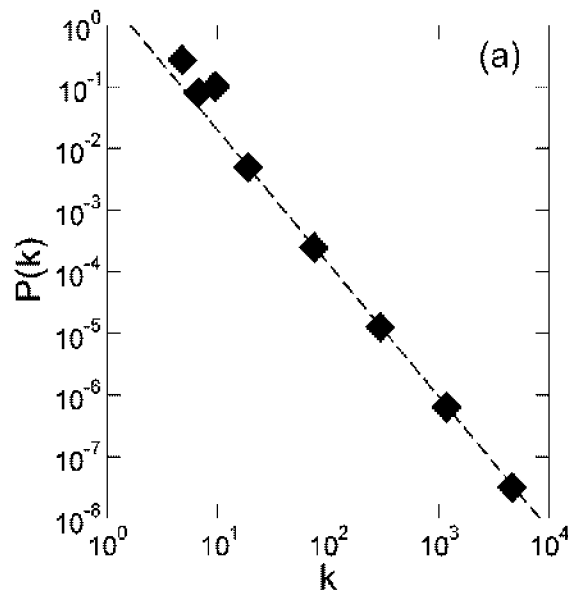
# Hierarchical Networks



# Properties of hierarchical networks

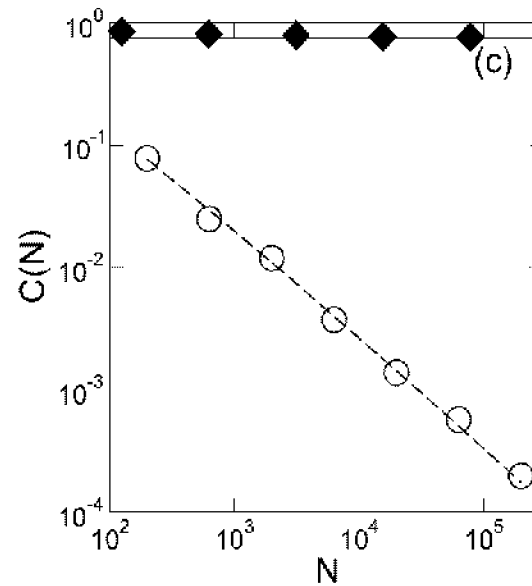
## 1. Scale-free

$$\gamma = 1 + \frac{\ln 5}{\ln 4} \\ = 2.161$$



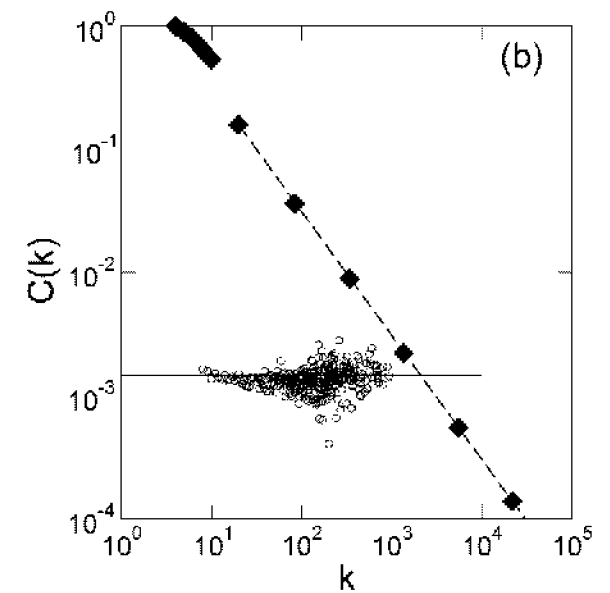
## 2. Clustering coefficient independent of N

$$C(N) = \text{const.}$$



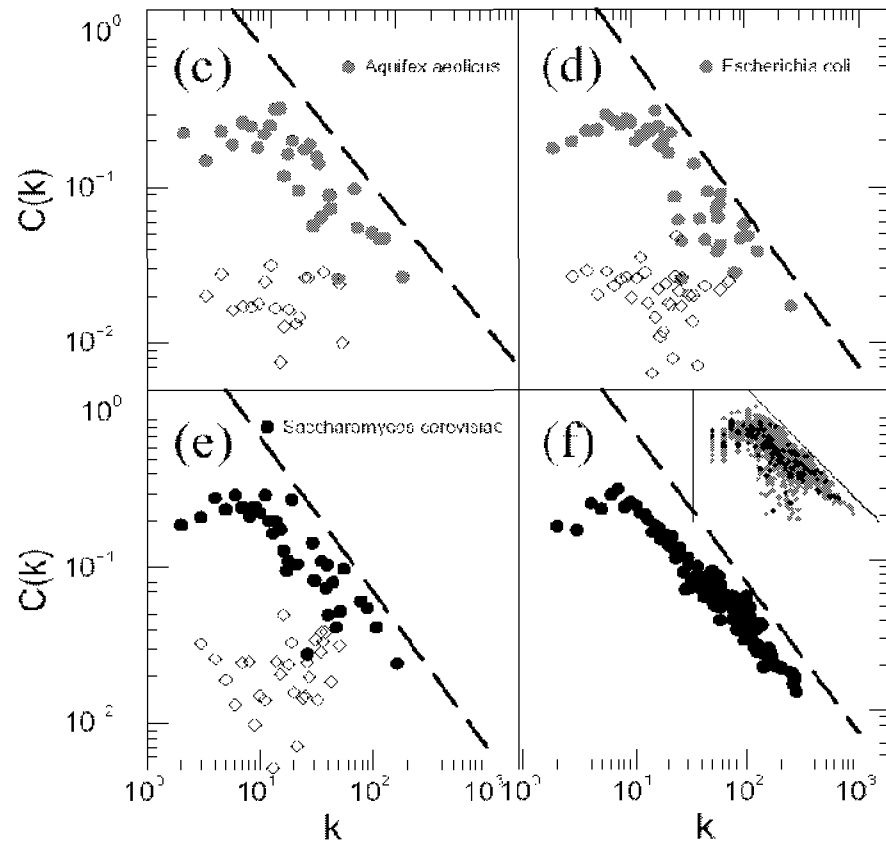
## 3. Scaling clustering coefficient (DGM)

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

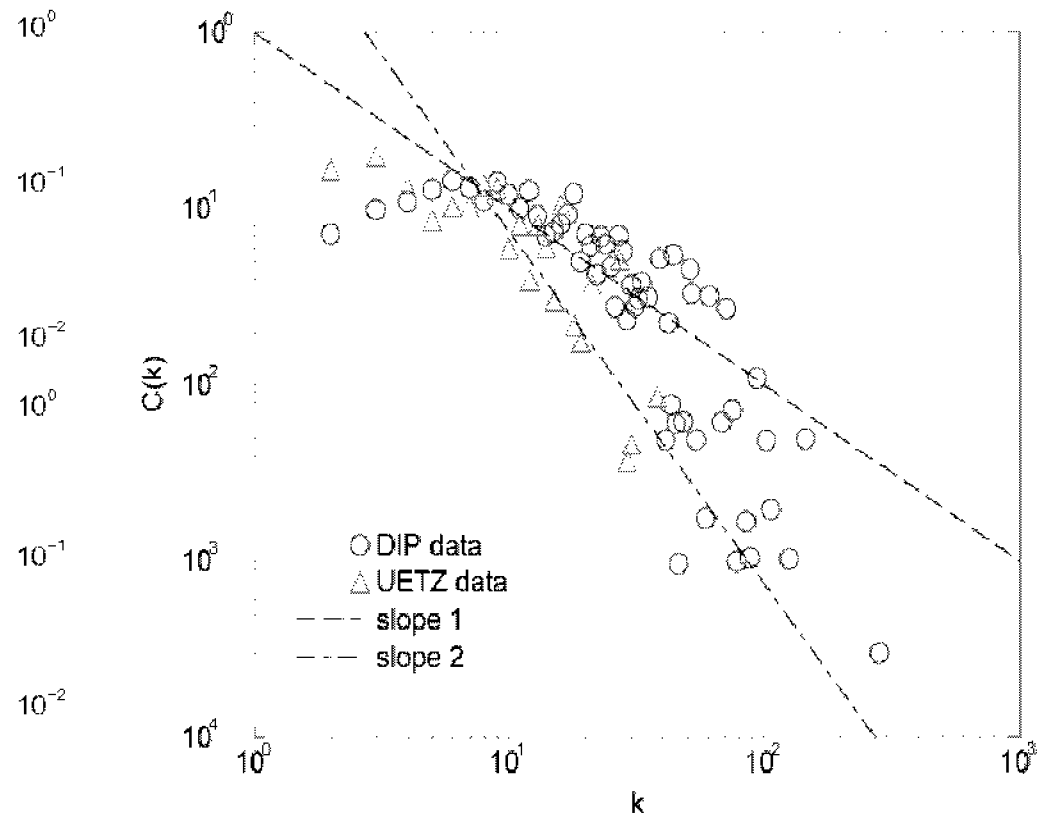




# Hierarchy in biological systems



**Metabolic networks**



**Protein networks**

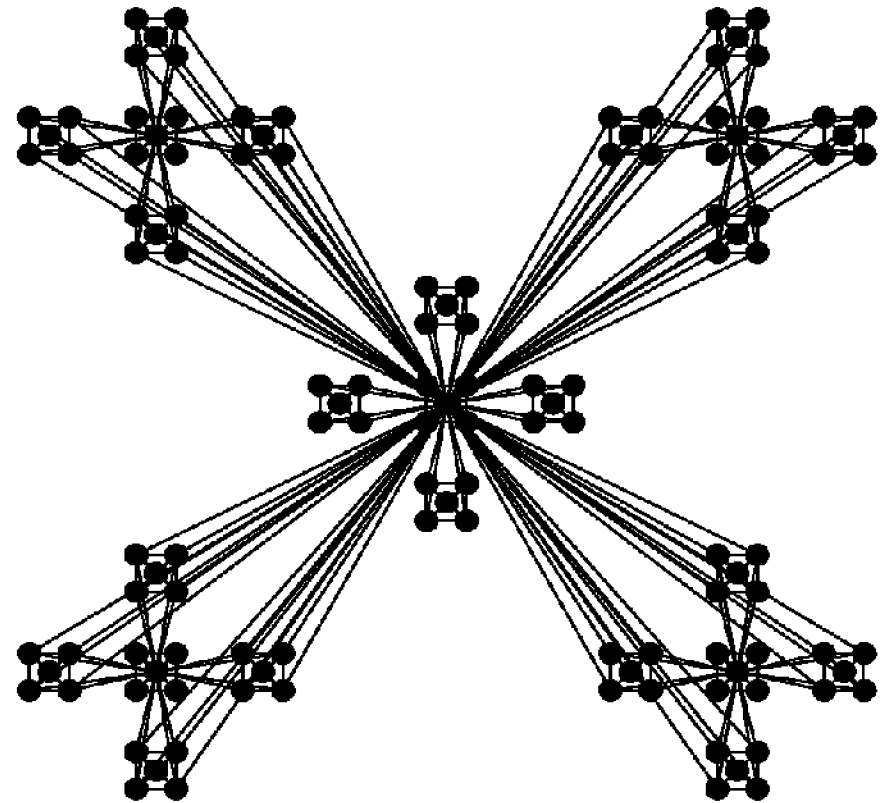
# Real Networks Have a Hierarchical Topology

What does it mean?

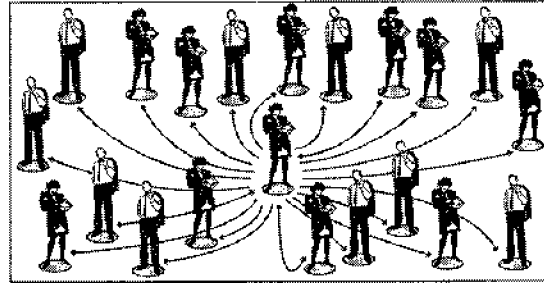
Many highly connected small clusters  
combine into  
few larger but less connected clusters  
combine into  
even larger and even less connected clusters

➤ The degree of clustering follows:

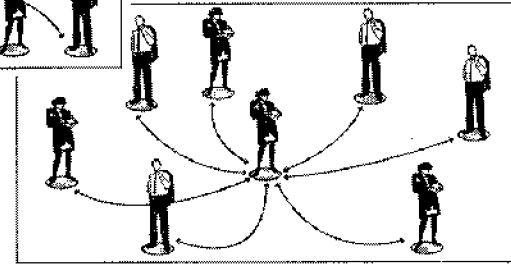
$$C(k) \sim k^{-\beta}$$



# Society



**Mega-Hub.** An MTV veejay spreads the word to thousands or millions of people through one-way links.



**Hub.** This undergraduate has spread the word to seven other people through two-way links.

# Internet

