

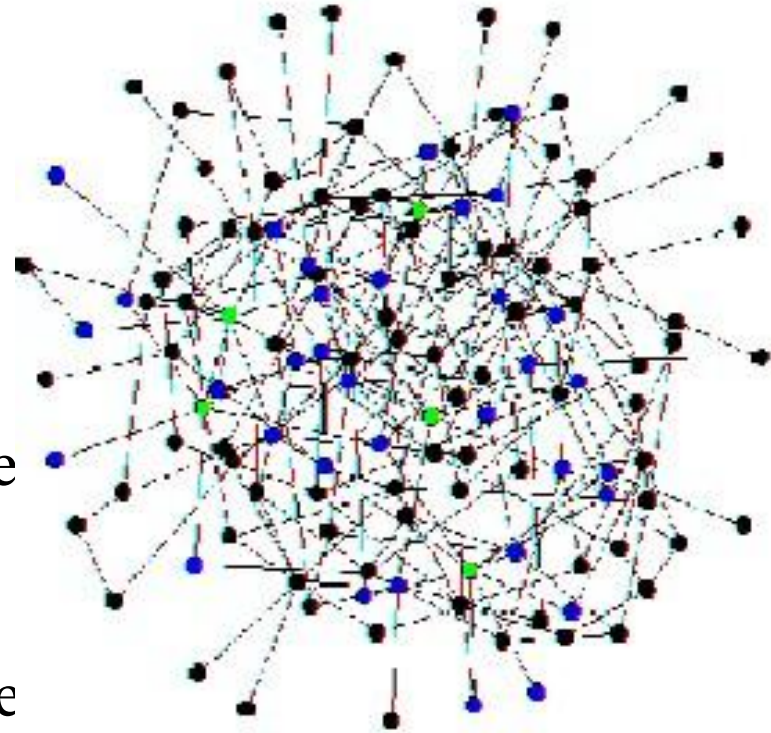
# Complex Networks

- **Network** is a structure of  $N$  **nodes** and  $2M$  **links** (or  $M$  **edges**)
- Called also **graph** – in Mathematics
- Many examples of networks

**Internet:** nodes represent computers  
links the connecting cables

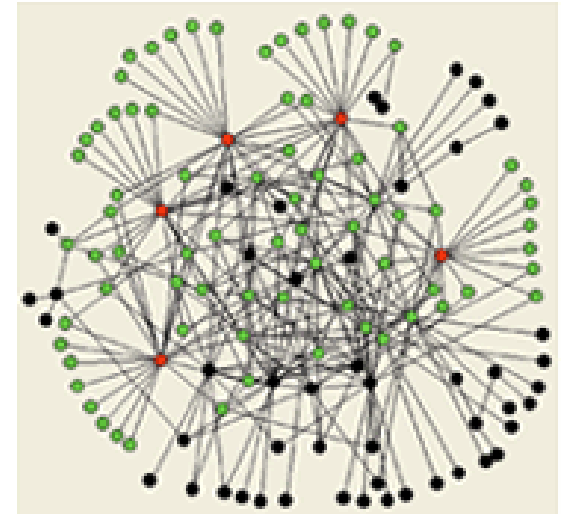
**Social network:** nodes represent people  
links their relations

**Cellular network:** nodes represent mole  
links their interactions

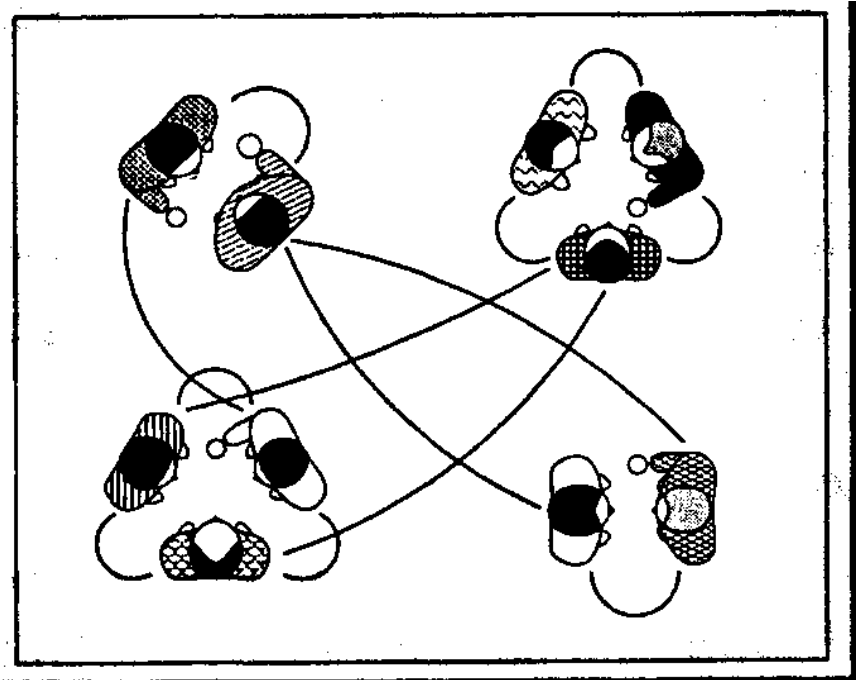
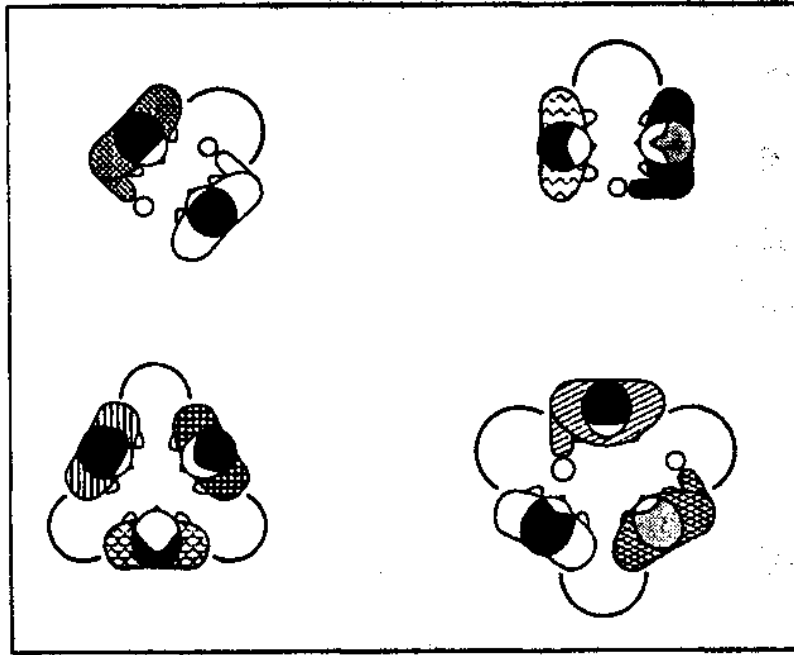


# Networks as a tool for Complex systems

- Complex systems are usually composed of many **interacting entities**.
- Can be well represented by **networks**: nodes represent the **entities** and links their **interactions**.
- Examples: biological systems, social systems, climate, earthquakes, epidemics, transport, economic, etc.
- **Weighted** networks each link (node) has a weight determining the strength or cost of the link (node).



# Social Networks- Stanley Milgram (1967)

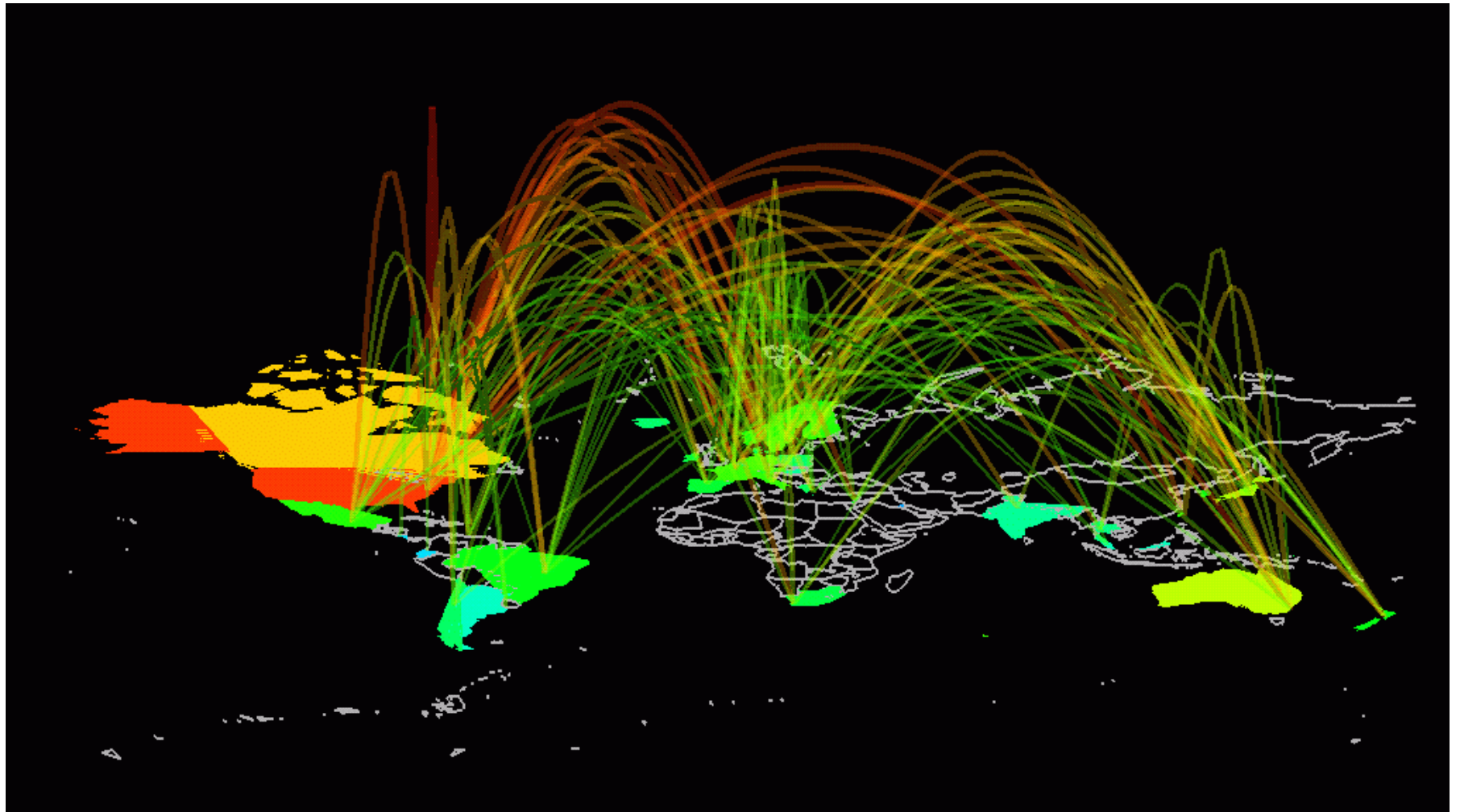


**Nodes**: individuals

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

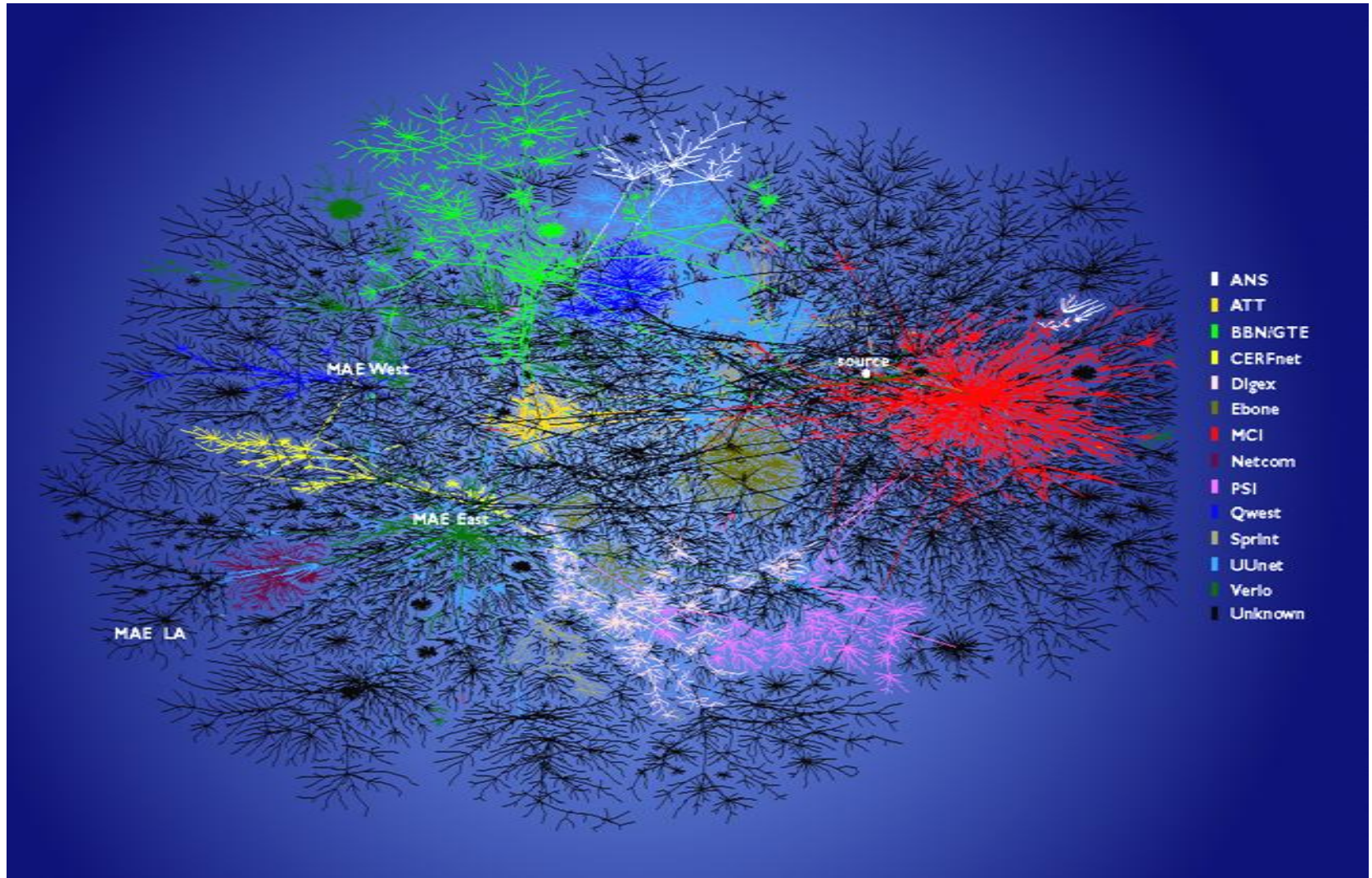
John Guare  
(1992)

**Six Degrees of Separation**

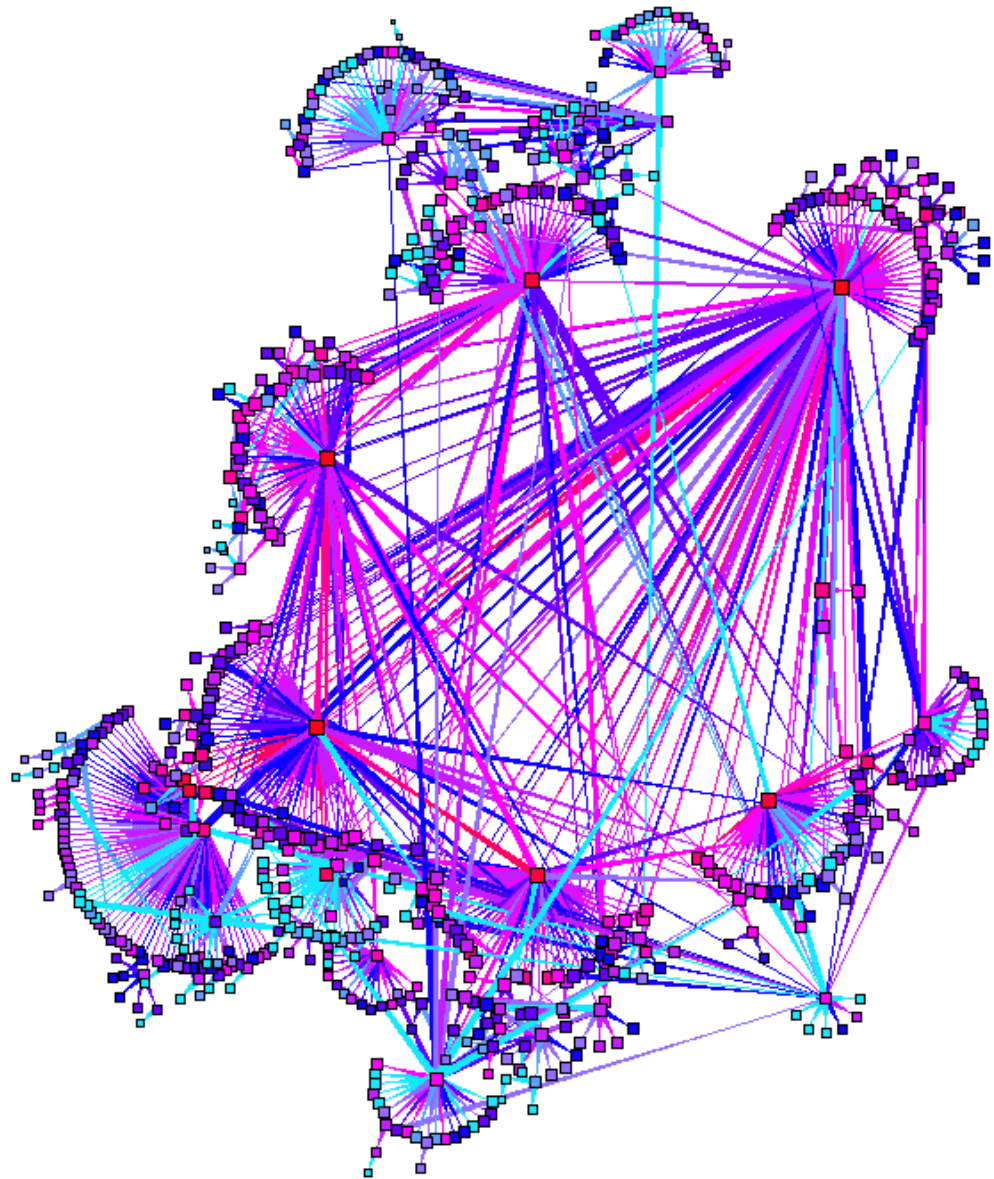
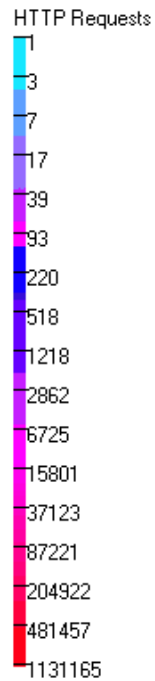


Map showing the world-wide internet traffic





A snapshot of Internet connectivity.



Hierarchical topology of the international web cache



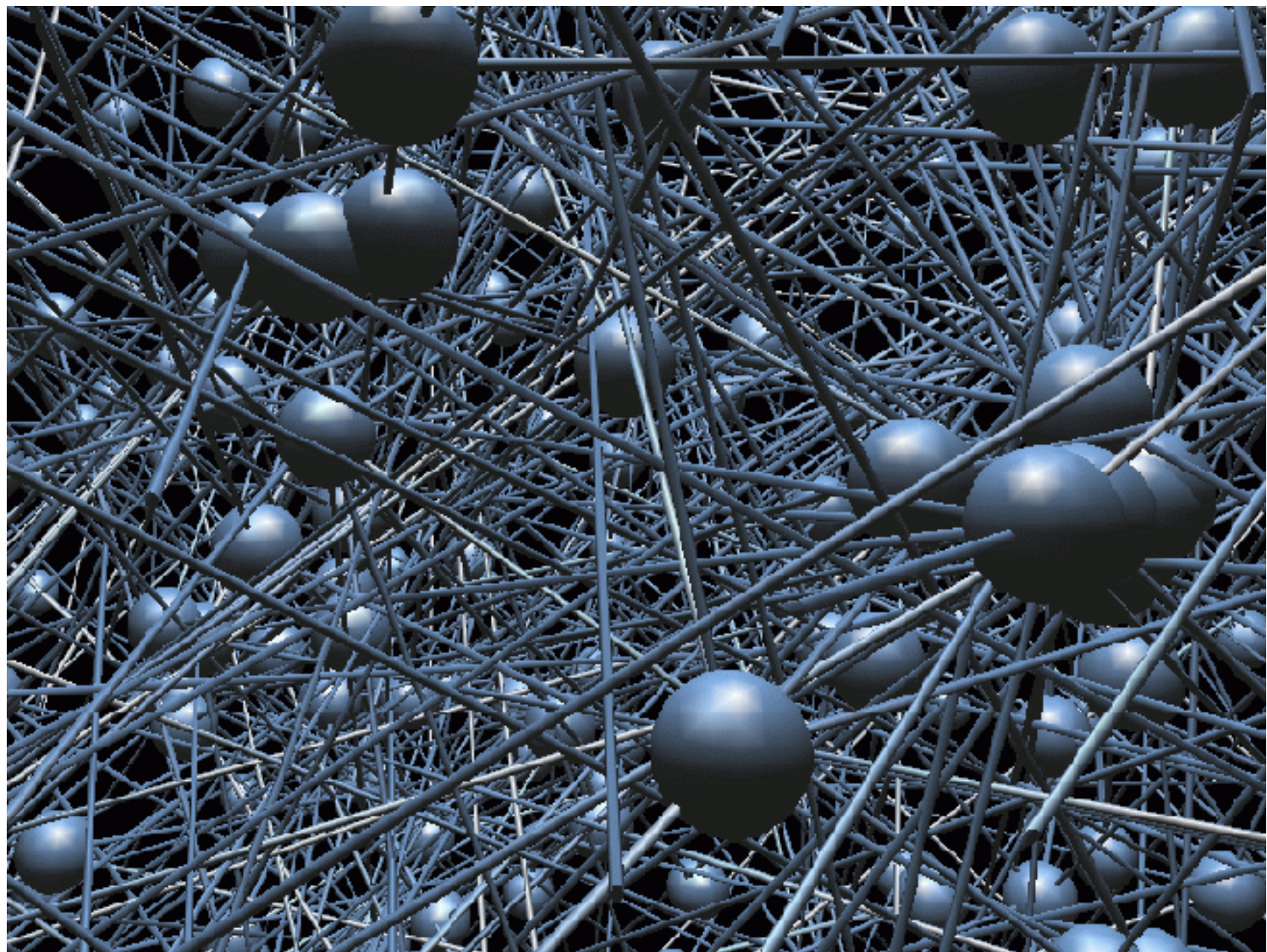
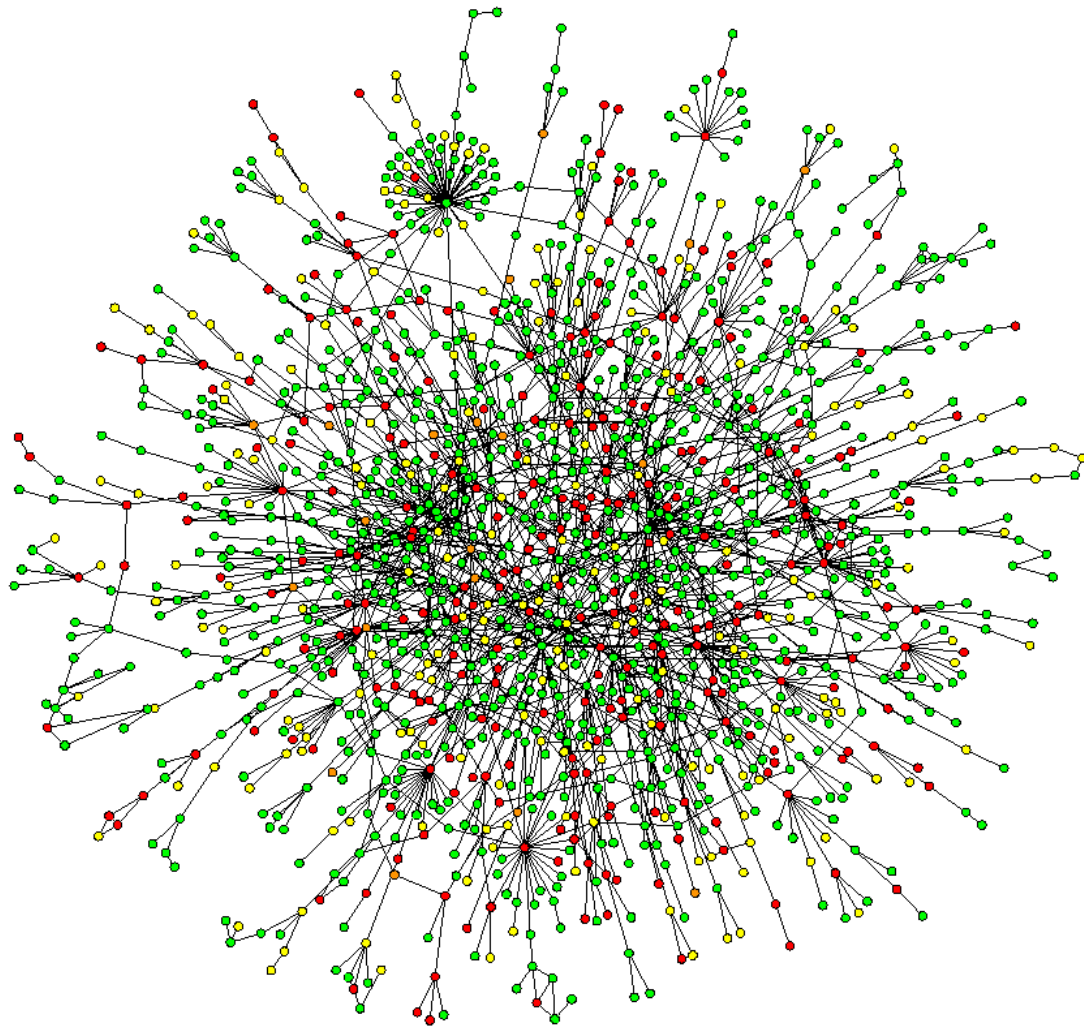


Image of Social links in Canberra, Australia



Network of protein-protein interactions. The color of a node signifies the phenotypic effect of removing the corresponding protein (red, lethal; green, non-lethal; orange, slow growth; yellow, unknown).



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

DROSOPHILA MELANOGASTER  
(Fruit fly)

HOMO SAPIENS

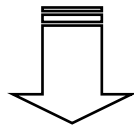
In this example the fly has 40 genes, and the human

▲ 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-

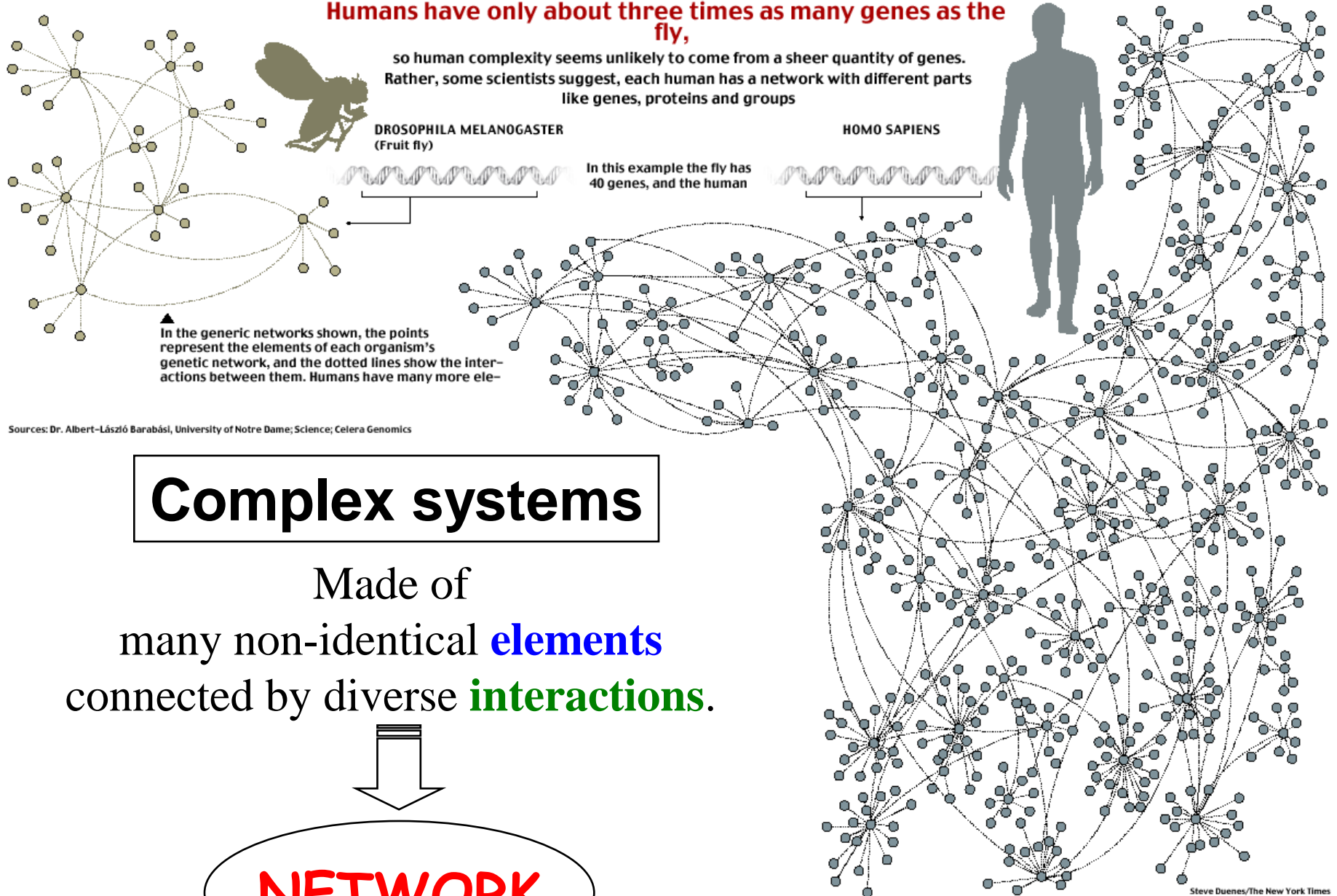
Sources: Dr. Albert-László Barabási, University of Notre Dame; Science; Celera Genomics

## Complex systems

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



**NETWORK**



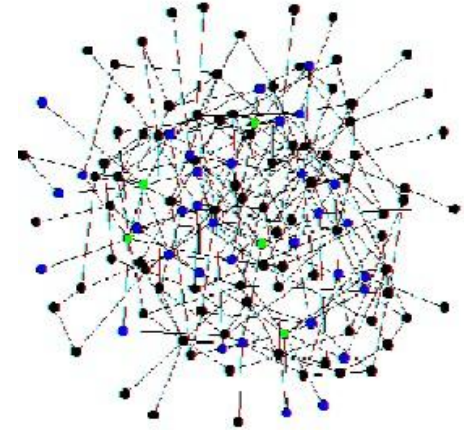
Steve Duenes/The New York Times

# Network Properties

- ❖ **Degree** distribution  $P(k)$  --  $k$ - degree of a node
- ❖ **Diameter** or **distance** -- Average distance between nodes-- $d$
- ❖ **Clustering Coefficient**  $c(k) = \frac{\text{no. of links between } k \text{ neighbors}}{k(k-1)/2}$   
How many of my friends are also friends?
- ❖ **Centrality** or **Betweenness** --  $b$

Number of times a bond or a node is relatively used for the shortest path

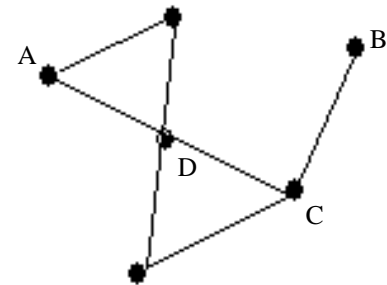
- ❖ **Critical Threshold:** The concentration of nodes that are removed and the network collapses



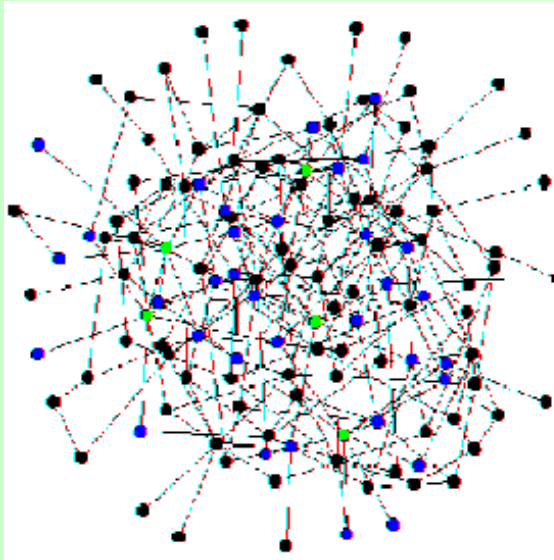
$$d(AB) = 3$$

$$c(D) = \frac{2}{6} = \frac{1}{3}$$

$$b(BC) = \frac{N-1}{N-1} = 1$$



# Random Graph Theory



- **Developed in the 1960's by Erdos and Renyi.** (Publications of the Mathematical Institute of the Hungarian Academy of Sciences, 1960).
- **Discusses the ensemble of graphs with N vertices and M edges (2M links).**
- **Distribution of connectivity per vertex is Poissonian (exponential),** where  $k$  is the number of links :

$$P(k) = e^{-c} \frac{c^k}{k!}, \quad c = \langle k \rangle = \frac{2M}{N}$$

- **Distance  $d = \log N$  -- SMALL WORLD**



# More Results

- **Phase transition at average connectivity,  $\langle k \rangle = 1$ :**

$\langle k \rangle < 1$  No spanning cluster (giant component) of order  $\log N$

$\langle k \rangle > 1$  A spanning cluster exists (unique) of order  $N$

$\langle k \rangle = 1$  The largest cluster is of order  $N^{2/3}$

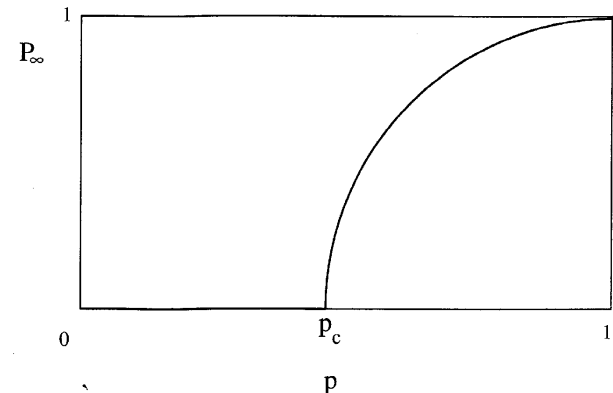
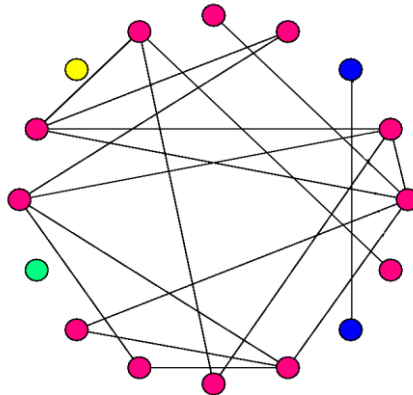
- **Behavior of the spanning cluster size near the transition is linear:**

$P_\infty \propto (p - p_c)^\beta$ ,  $\beta = 1$ , where  $p$  is the probability of a site to exist,

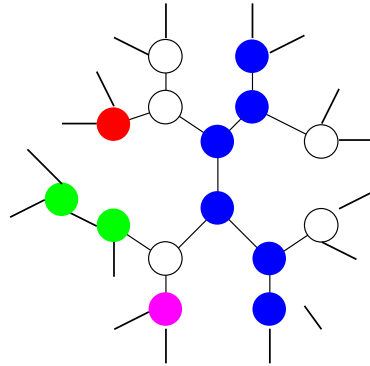
$$p_c = 1 / \langle k \rangle$$

- **Size of the spanning cluster is determined by the self-consistent equation:**

$$P_\infty = 1 - e^{-\langle k \rangle P_\infty}$$



# Percolation on a Cayley Tree



- Contains no loops
- Connectivity of each node is fixed ( $z$  connections)

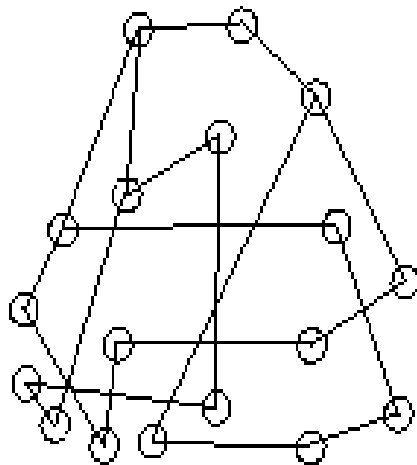
- Critical threshold: 
$$p_c = \frac{1}{z-1}$$

- Behavior of the spanning cluster size near the transition is linear:

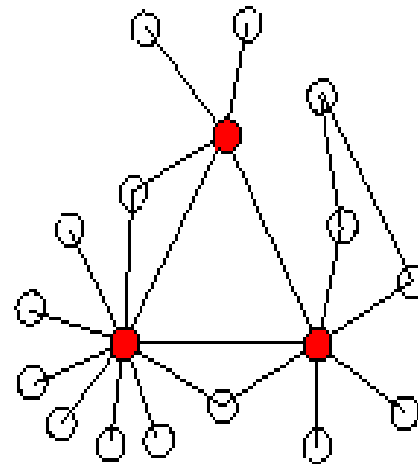
$$P_\infty \propto (p_c - p)^\beta, \quad \beta = 1$$

# In Real World - Many Networks are non-Poissonian

**Exponential**



**Scale-free**



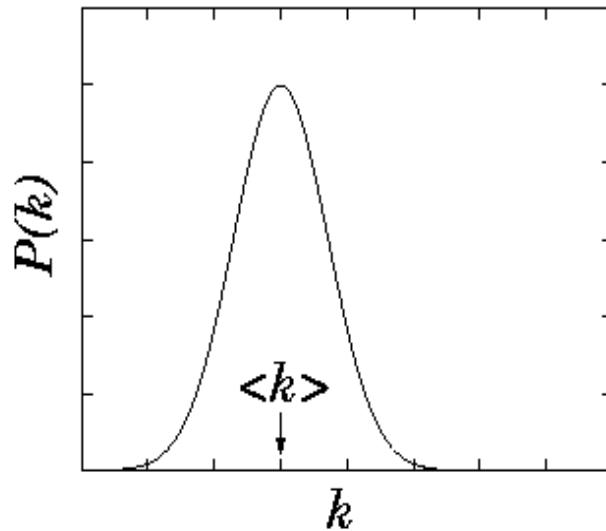
$$P(k) = e^{-\langle k \rangle} \frac{\langle k \rangle^k}{k!}$$

$$P(k) = \begin{cases} ck^{-\lambda} & m \leq k \leq K \\ 0 & \text{otherwise} \end{cases}$$

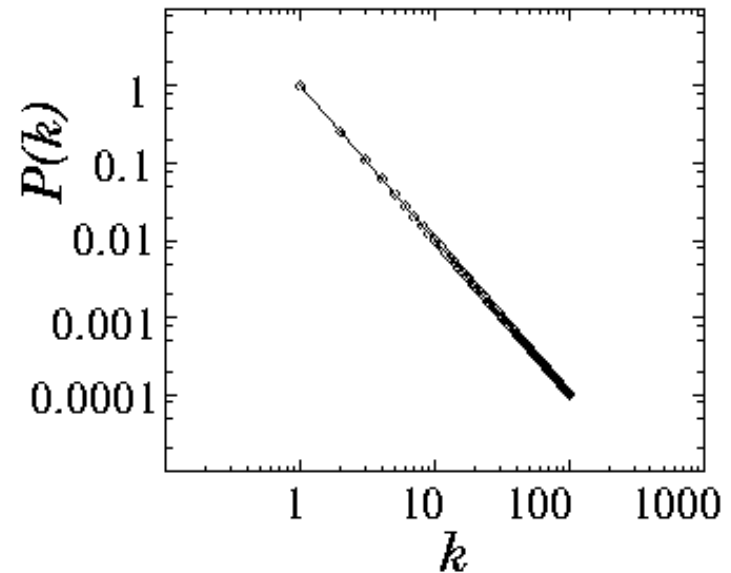


# New Type of Networks

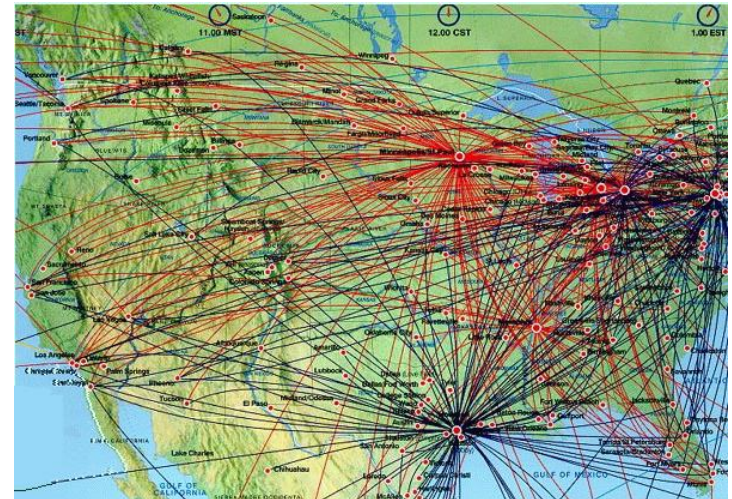
Poisson distribution



Power-law distribution

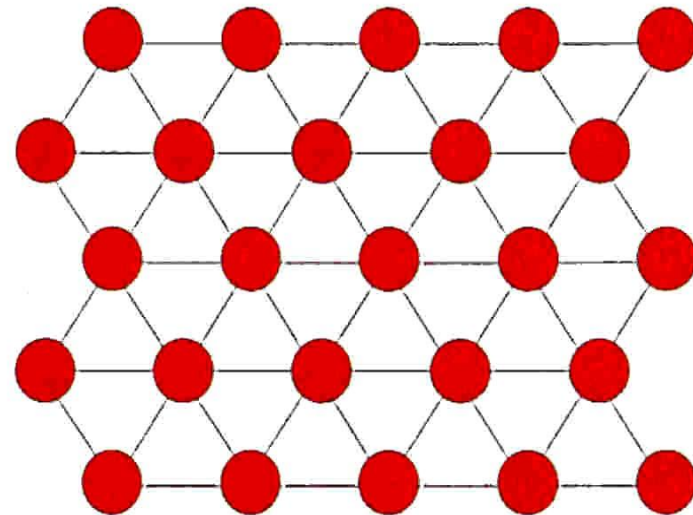
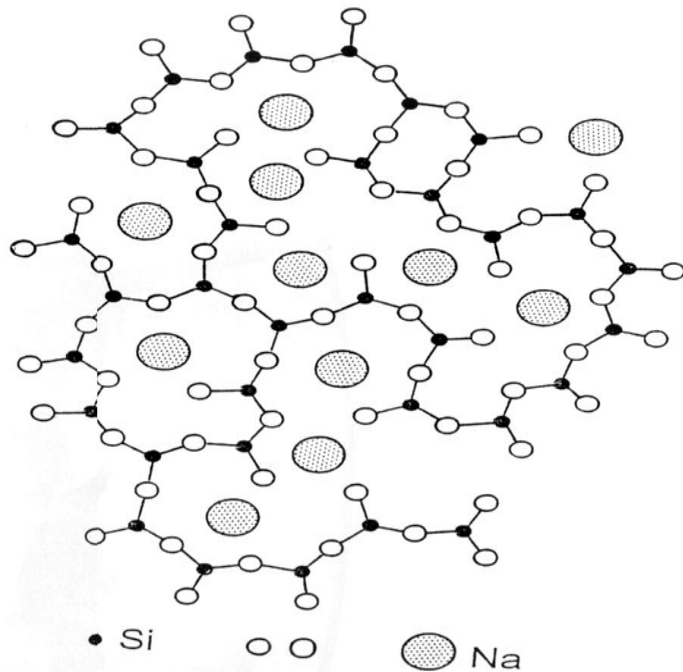
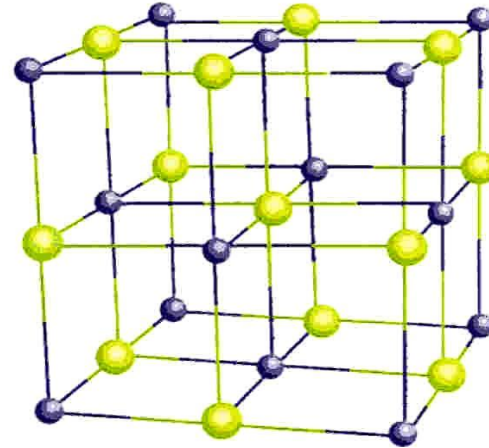
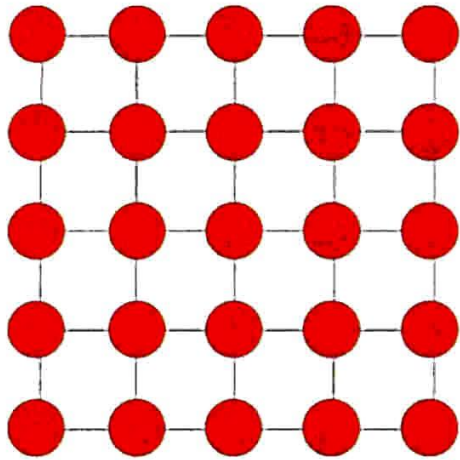


**Exponential Network**



**Scale-free Network**

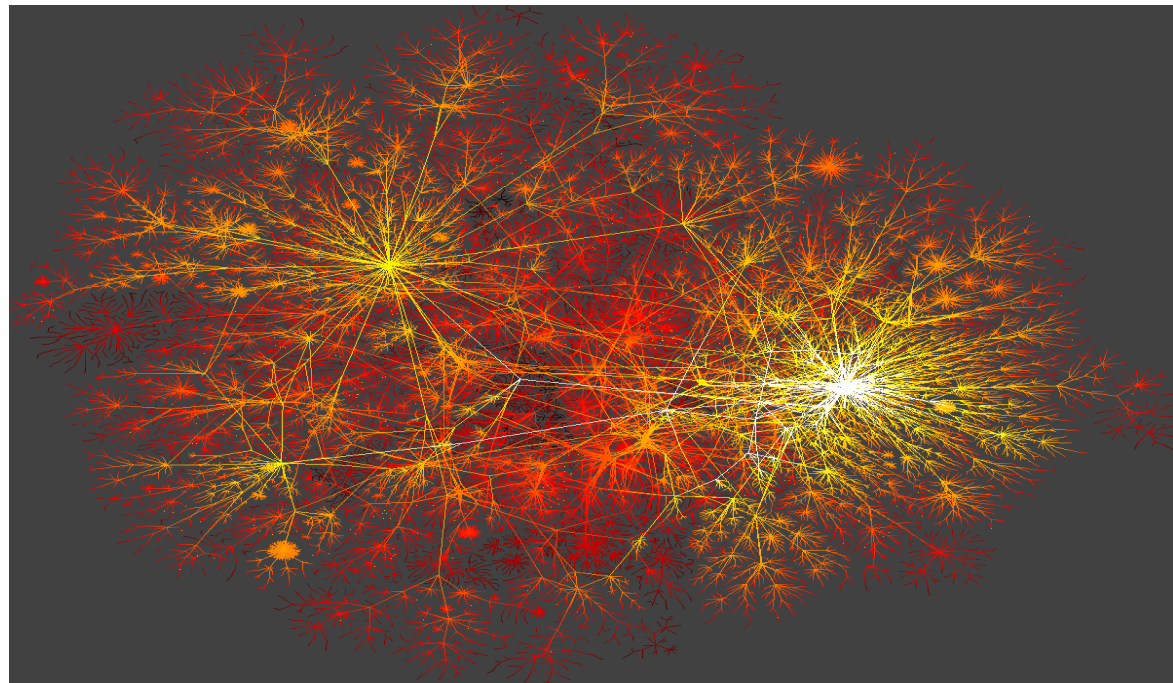
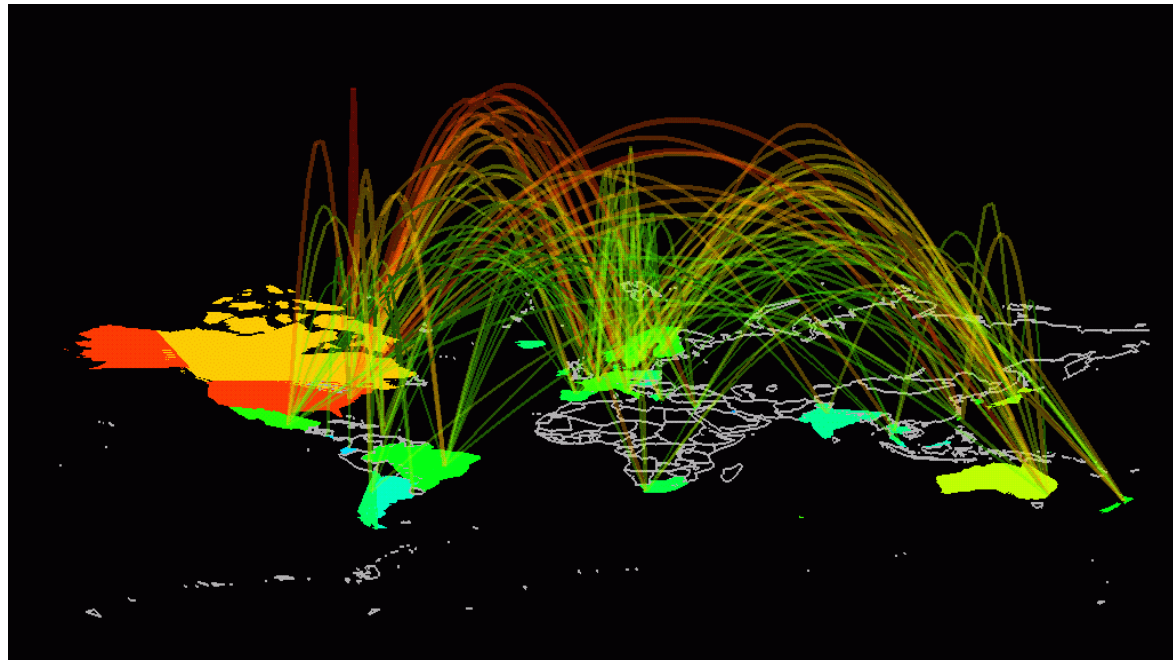
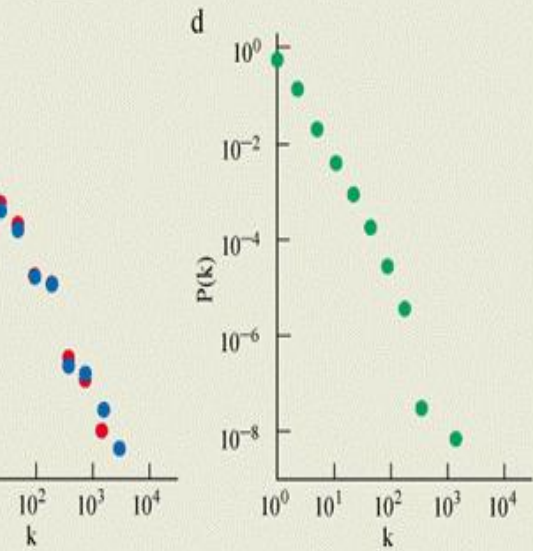
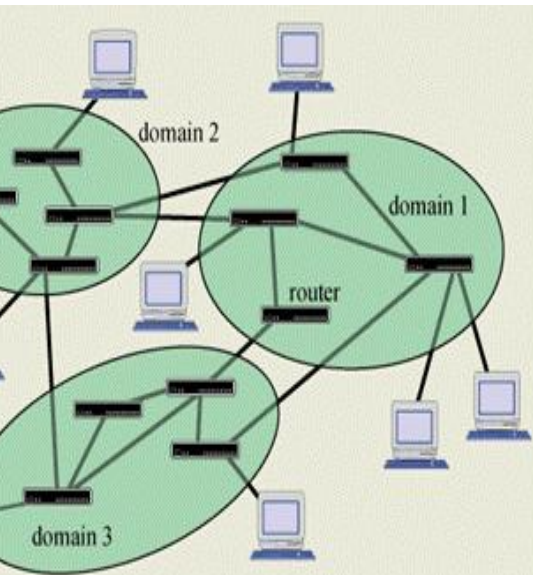
# Networks in Physics





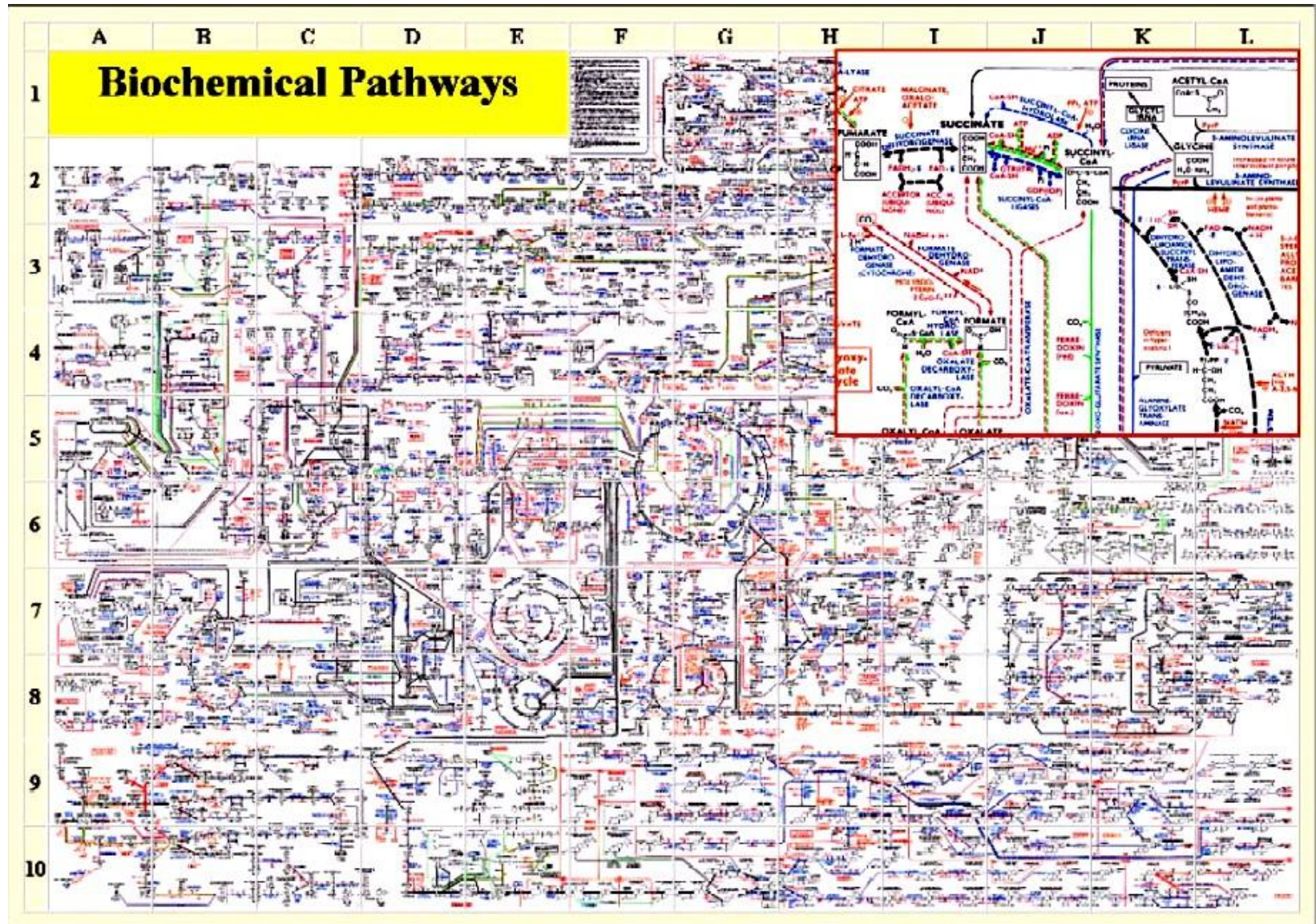
# Internet Network

Faloutsos et. al., SIGCOMM '99





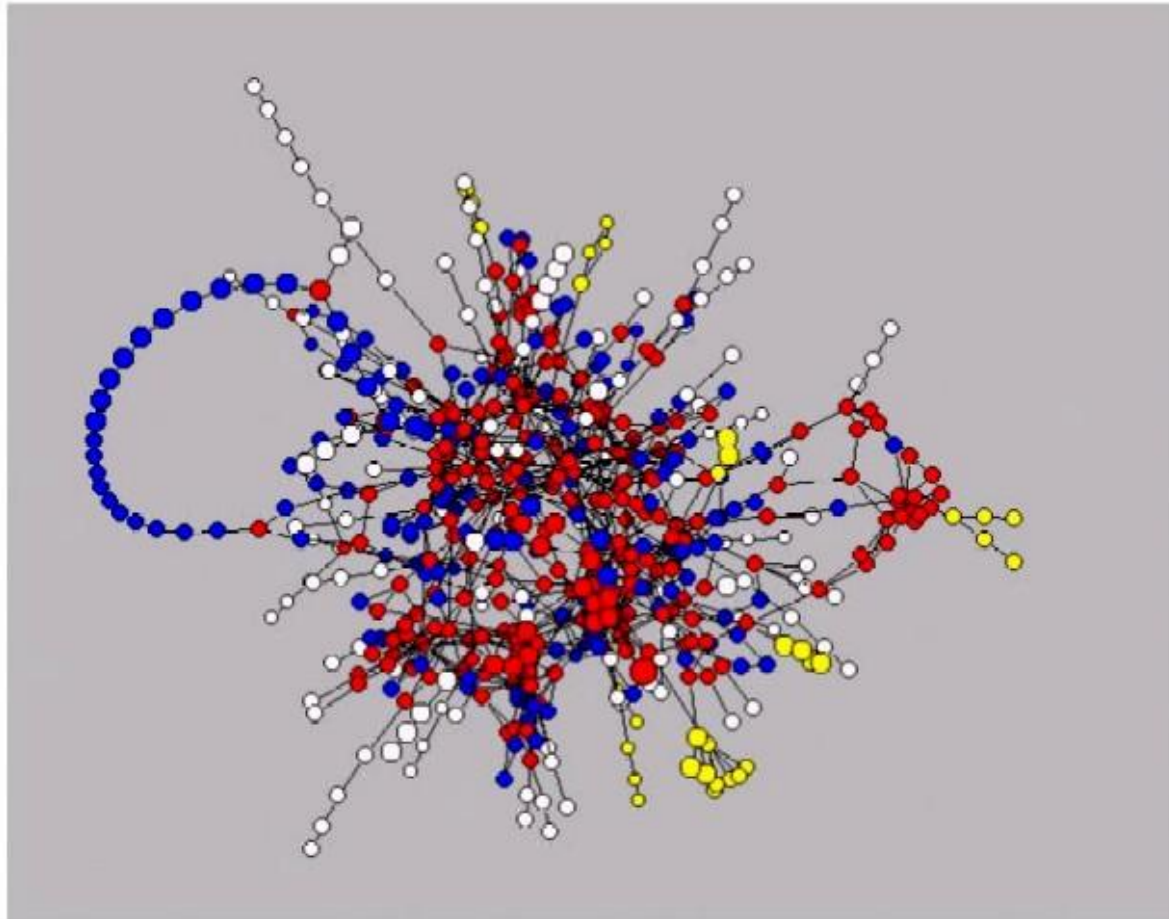
# Metabolic network



# Metabolic Network

Nodes: chemicals (substrates)

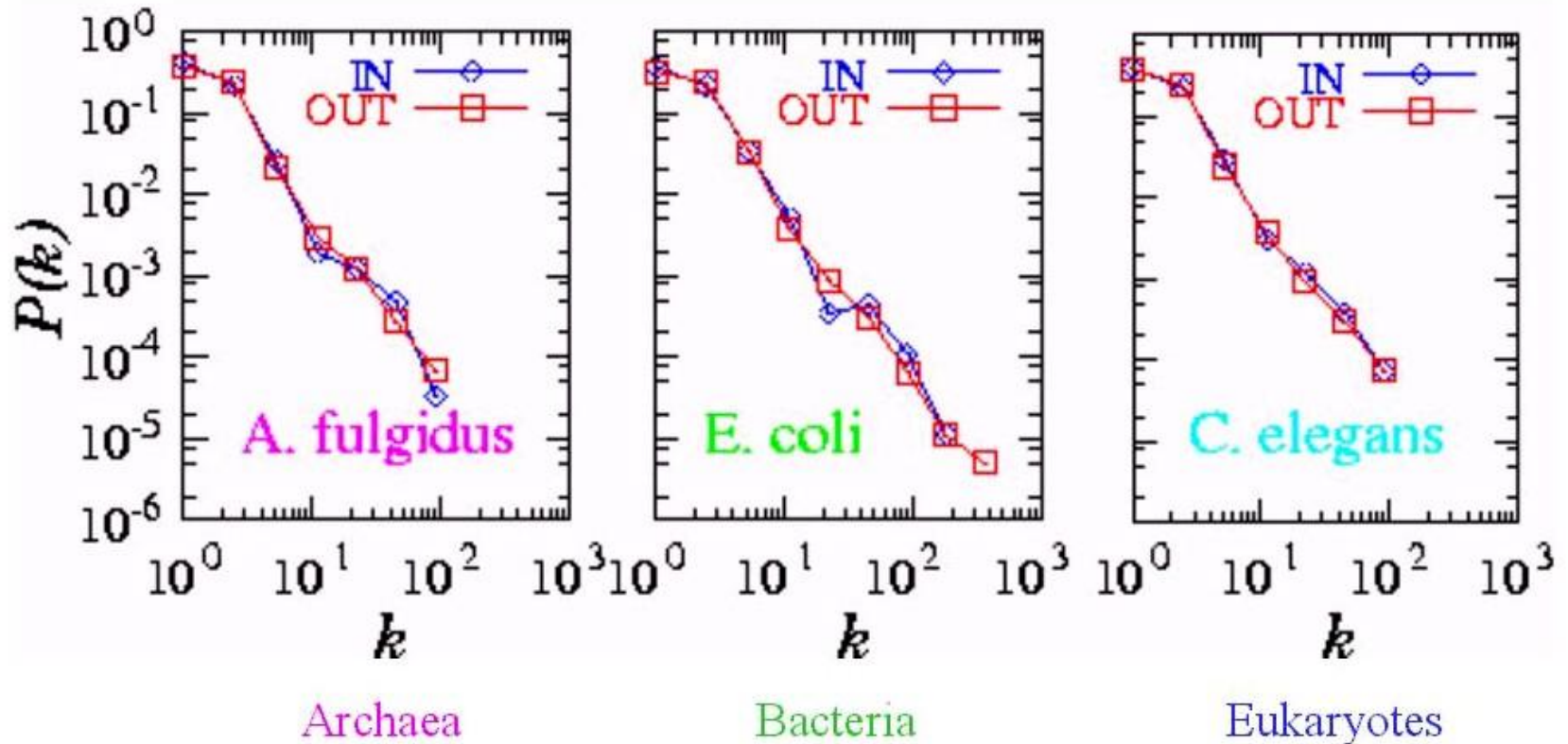
Links: bio-chemical reactions



Jeong et al Nature 2000



# Metabolic network

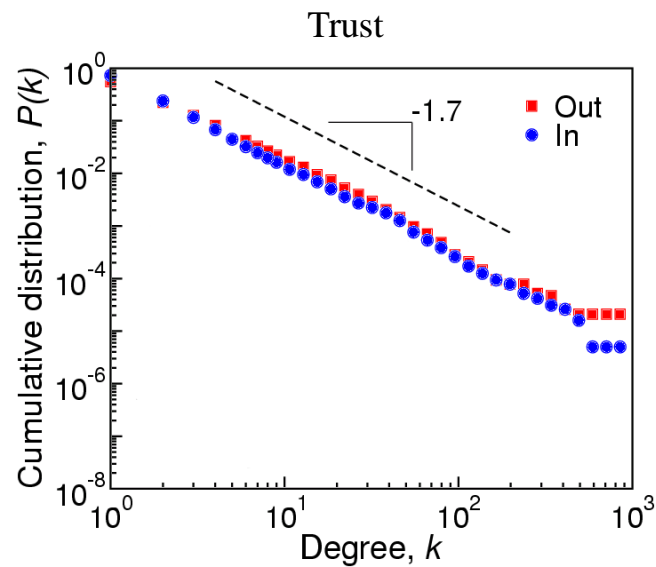
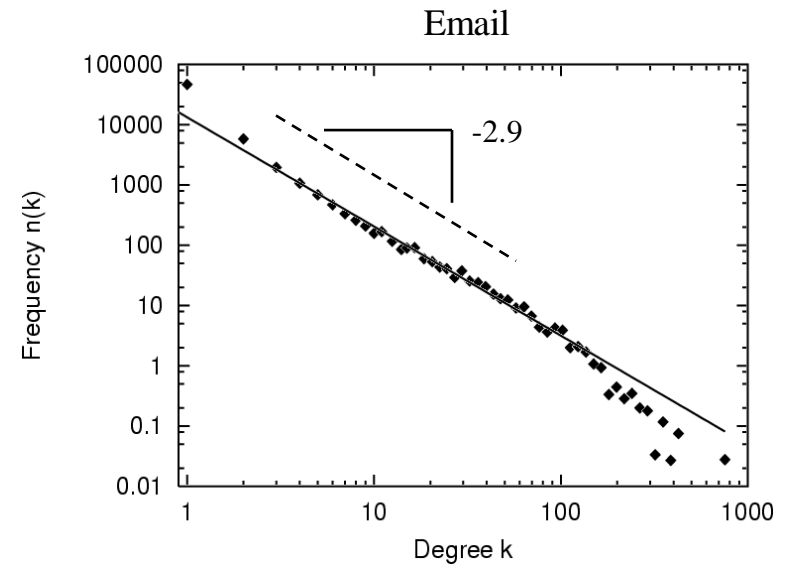
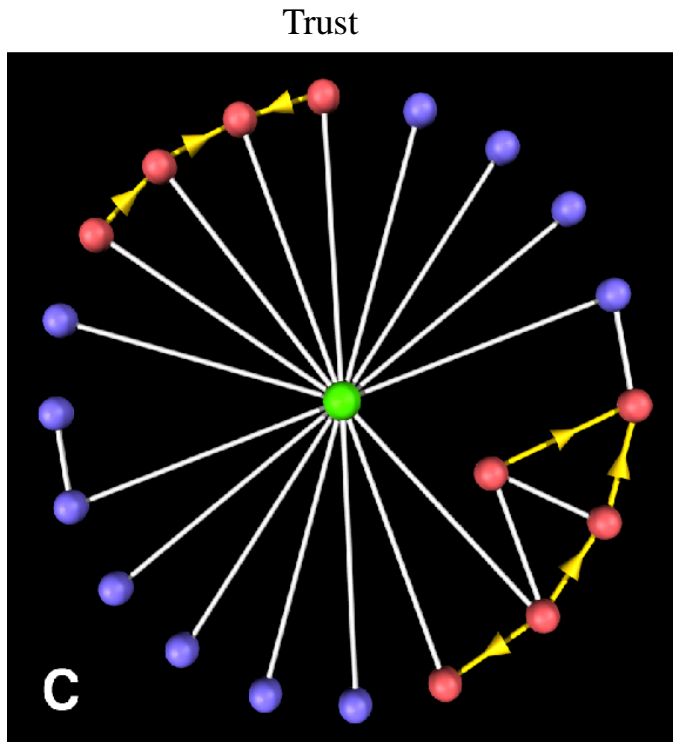


Organisms from all three domains of life are  
**scale-free** networks!

Jeong et al, Nature (2000)

# More Examples

- Trust networks: Guardiola et al (2002)
- Email networks: Ebel et al PRE (2002)





# Erdős Theory is Not Valid

## Stability and Immunization

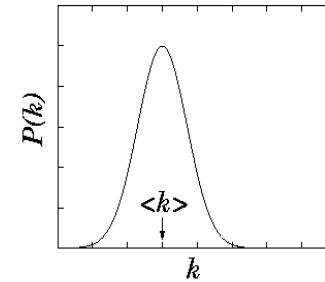
$$q_c = 1 - p_c = 1 - \frac{1}{\langle k \rangle}$$

Critical concentration 30-50%

## Distance

$$d \sim \log N$$

## Distribution

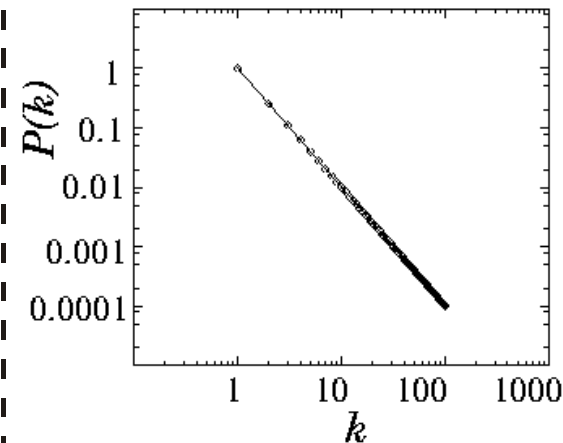


**T**  
**h**  
**e**  
**o**  
**r**  
**y**

### Infectious disease Critical concentration

Malaria	99%
Measles	90-95%
Whooping cough	90-95%
Fifths disease	90-95%
Chicken pox	85-90%
Mumps	85-90%
Rubella	82-87%
Poliomyelitis	82-87%
Diphtheria	82-87%
Scarlet fever	82-87%
Smallpox	70-80%
INTERNET	99%

Almost constant  
(Metabolic Networks,  
Jeong et. al.  
(Nature, 2000))



**E**  
**x**  
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**t**

**Generalization** of Erdős Theory:

Cohen, Erez, ben-Avraham, Havlin, PRL **85**, 4626 (2000)

**Epidemiology Theory:** Vespignani, Pastor-Satoral,  
PRL (2001), PRE (2001)

Cohen, Havlin,

Phys. Rev. Lett. 90, 58701(2003)

**Modelling:** Albert, Jeong, Barabasi (Nature 2000)

# Experimental Data: Virus survival

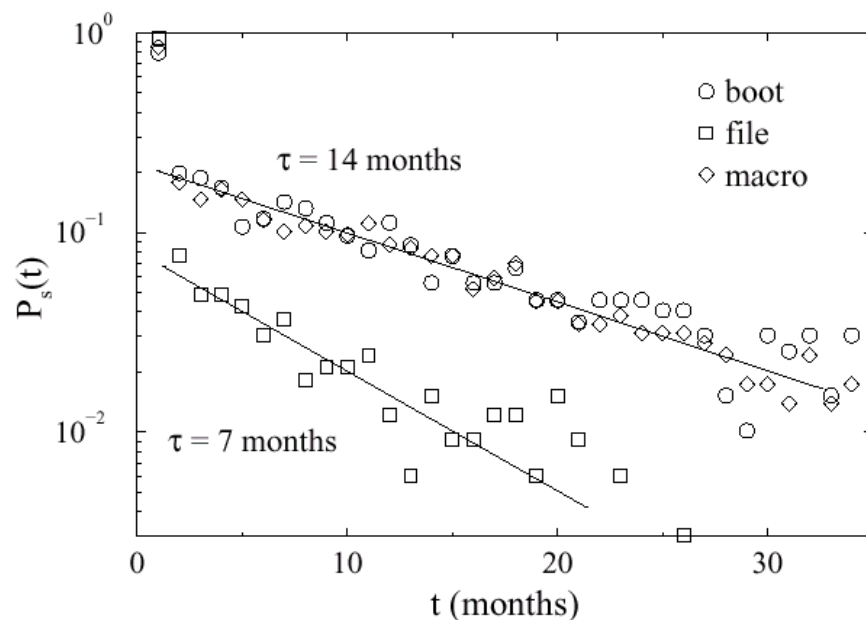
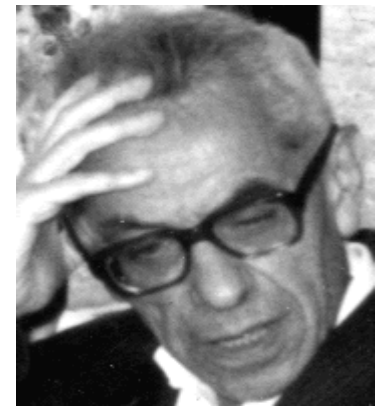


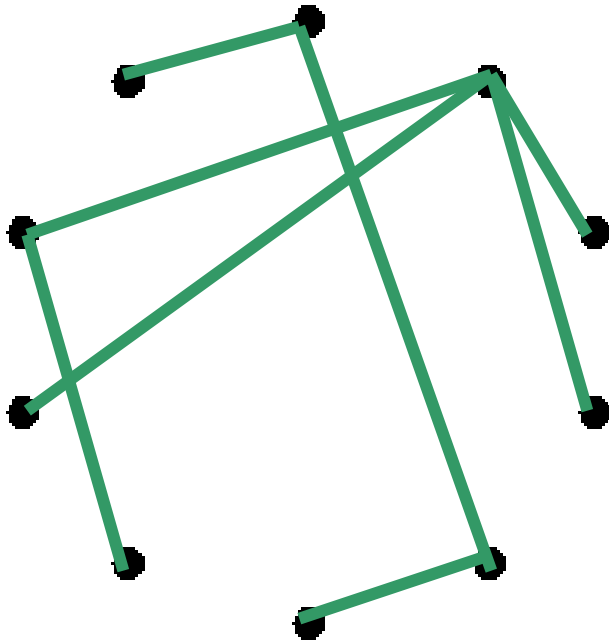
FIG. 1. Surviving probability for viruses in the wild. The 814 different viruses analyzed have been grouped in three main strains [9]: file viruses infect a computer when running an infected application; boot viruses also spread via infected applications, but copy themselves into the boot sector of the hard-drive and are thus immune to a computer reboot; macro viruses infect data files and are thus platform-independent. It is evident in the plot the presence of an exponential decay, with characteristic time  $\tau \simeq 14$  months for macro and boot viruses and  $\tau \simeq 7$  months for file viruses.

**(Pastor-Satorras and Vespignani, Phys. Rev Lett. 86, 3200 (2001))**

# Erdős-Rényi model (1960)



**Pál Erdős  
(1913-1996)**



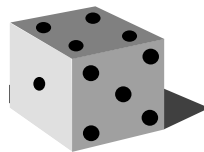
Connect with  
probability  $p$

$$p=1/6$$

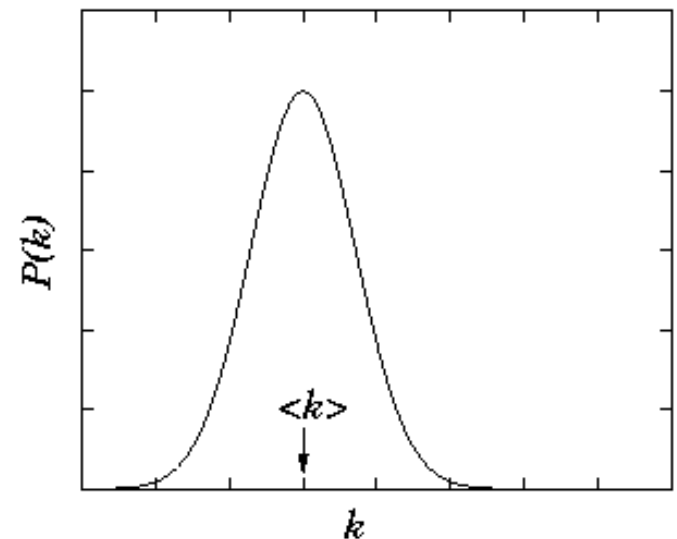
$$N=10$$

$$\langle k \rangle \sim 1.5$$

Poisson distribution



- Democratic
- Random



# Scale-free model

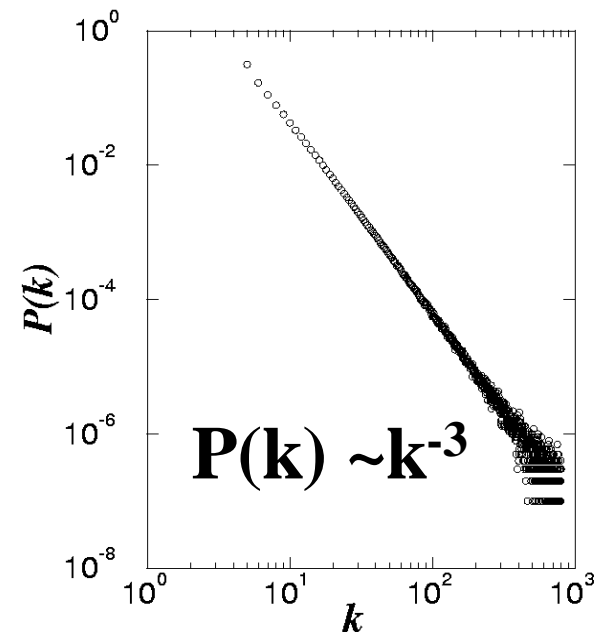
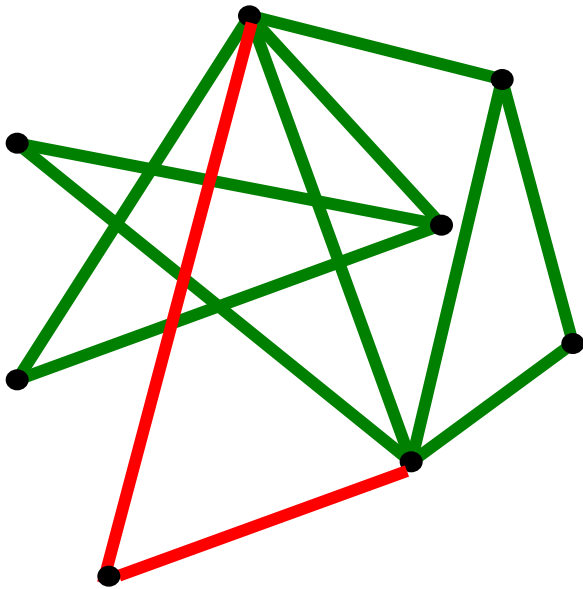
## (1) GROWTH :

At every time step 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

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



# Shortest Paths in Scale Free Networks

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

$$d = \text{const.}$$

$$\lambda = 2$$

Ultra  
Small  
World

$$d = \log \log N$$

$$2 < \lambda < 3$$

$$d = \frac{\log N}{\log \log N}$$

$$\lambda = 3$$

(Bollobas, Riordan, 2002)

Small World

$$d = \log N$$

$$\lambda > 3$$

(Bollobas, 1985)

(Newman, 2001)

Cohen, Havlin Phys. Rev. Lett. 90, 58701(2003)

Cohen, Havlin and ben-Avraham, in Handbook of Graphs and Networks

eds. Bornholdt and Shuster (Wiley-VCH, NY, 2002) chap.4

Confirmed also by: Dorogovtsev et al (2002), Chung and Lu (2002)



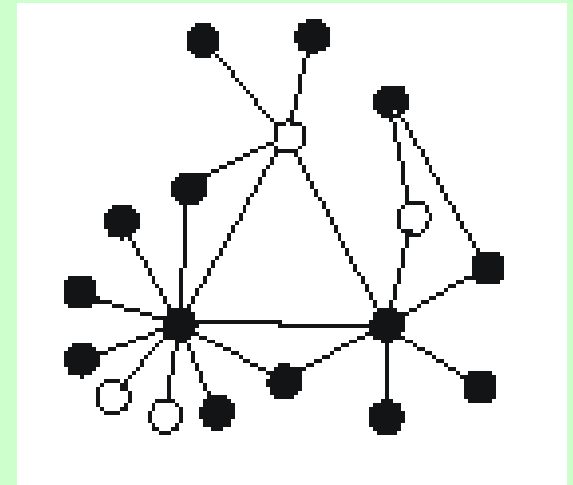
# Model of Stability

## Random Breakdown (Immunization)

The Internet is believed to be almost randomly connected scale-free network, where

$$P(k) \propto k^{-\lambda}, \lambda \approx 2.5$$

Nodes are randomly removed (or immune)  
with probability  $q=1-p$

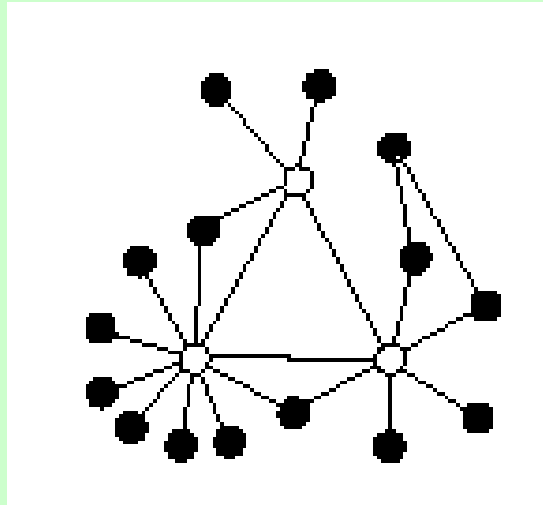


Where does the phase transition occur ?

# Model for Stability

## Targeted Attack (Immunization)

The fraction,  $q$ , of nodes with the highest degree are removed (or immunized).



Is this fundamentally different from random breakdown?

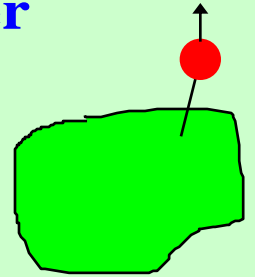
We find that not only **critical thresholds** but also **critical exponents** are different !

**THE UNIVERSALITY CLASS DEPENDS ON THE WAY CRITICALITY REACHED**

# THEORY FOR ANY DEGREE DISTRIBUTION

## Condition for the Existence of a Spanning Cluster

If we start moving on the cluster from a single site, in order that the cluster does not die out, we need that each site reached will have, on average, at least 2 links (one “in” and one “out”).



This means:  $\langle k_i | i \leftrightarrow j \rangle = \sum_{k_i} k_i P(k_i | i \leftrightarrow j) \geq 2$ , where  $i \leftrightarrow j$  means that site  $i$  is connected to site  $j$ .

But, by Bayes rule:  $P(k_i | i \leftrightarrow j) = \frac{P(i \leftrightarrow j | k_i) P(k_i)}{P(i \leftrightarrow j)}$

We know that  $P(i \leftrightarrow j | k_i) = \frac{k_i}{N-1}$  and  $P(i \leftrightarrow j) = \frac{\langle k \rangle}{N-1}$

Combining all this together:  $\kappa \equiv \frac{\langle k^2 \rangle}{\langle k \rangle} = 2$   
(for every distribution) at the critical point.

**Exponential graph:**

$$\frac{\langle k^2 \rangle}{\langle k \rangle} = \frac{\langle k \rangle^2 + \langle k \rangle}{\langle k \rangle} = 2$$

$$\Rightarrow \langle k \rangle = 1$$

**Cayley Tree:**

$$p_c = \frac{1}{z-1}$$

# Percolation for Random Breakdown

If percolation is considered the connectivity **distribution** changes according

to the law:  $\bar{P}(k) = \sum_{k' > k} P(k') \binom{k'}{k} p^{k'-k} (1-p)^k$

Calculating the change in  $\kappa$  gives the percolation threshold:

$p_c = 1 - q_c = \frac{1}{\kappa_0 - 1}$ , where  $\kappa_0 = \frac{\langle k_0^2 \rangle}{\langle k_0 \rangle}$ . **compared to**  $p_c = 1 / \langle k_0 \rangle$  for Erdos Renyi

For **scale-free** distribution with lower cutoff  $m$ , and upper cutoff  $K$ , gives

$$\kappa_0 = \left( \frac{2 - \lambda}{3 - \lambda} \right) \frac{K^{3-\lambda} - m^{3-\lambda}}{K^{2-\lambda} - m^{2-\lambda}}, \quad K \gg N^{\frac{1}{\lambda-1}}.$$

For scale-free graphs where the second moment diverges.

No critical threshold!

Network is stable (or not immunized) even for  $q \rightarrow 1$

# Critical Threshold Scale Free

General result:

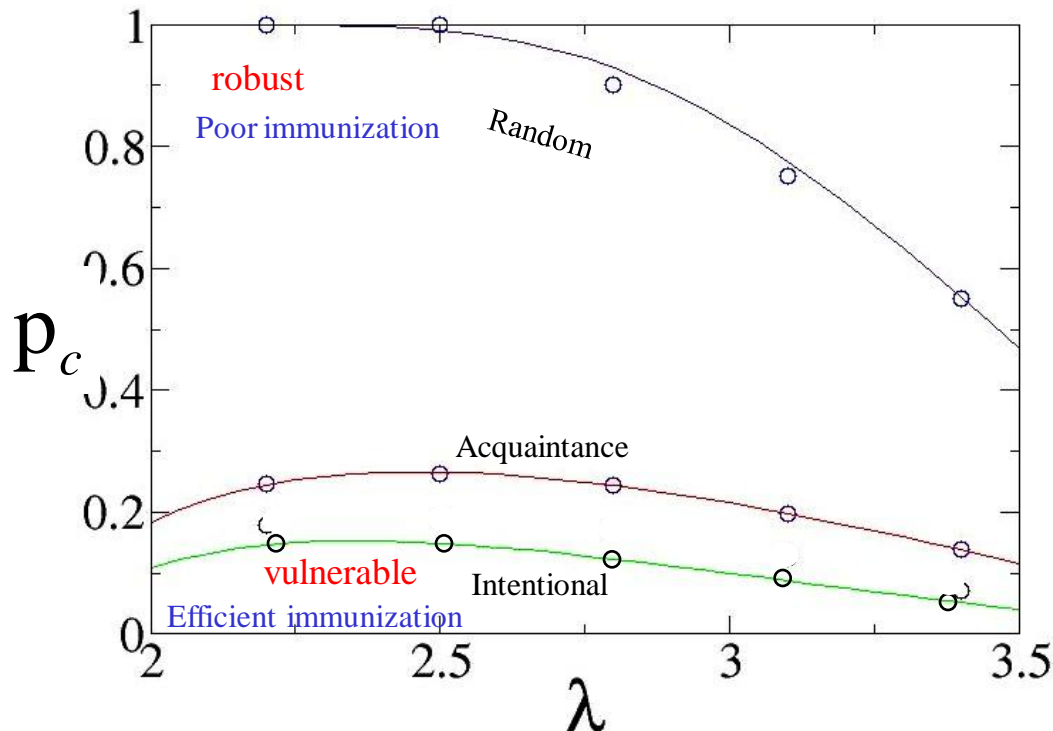
$$p_c = \frac{1}{K_0 - 1}$$

$$K_0 \equiv \frac{\langle k^2 \rangle}{\langle k \rangle}$$

For Poisson:

$$K_0 = \frac{\langle k^2 \rangle}{\langle k \rangle} = \frac{\langle k \rangle^2 + \langle k \rangle}{\langle k \rangle}$$

$$p_c = \frac{1}{\langle k \rangle}$$



Cohen et al. Phys. Rev. Lett. 91, 168701 (2003)

Efficient Immunization  
Strategies:

Acquaintance Immunization



# Percolation for Targeted Attack (Immunization)

Attack has two kinds of influence on the connectivity distribution:

- Change in the upper cutoff

Can be calculated by  $\sum_{k=\bar{K}}^K P(k) = p$ ,

or approximately:  $\bar{K} = mp^{1/(1-\lambda)}$ .

- Change in the connectivity of all other sites due to possibility of a broken link (which is different than in random breakdown). The probability of a link to be removed can be calculated by:

$$\bar{p} = \frac{1}{\langle k_0 \rangle} \sum_{k=\bar{K}}^K kP(k),$$

or approximately:  $\bar{p} = p^{(2-\lambda)/(1-\lambda)}$ .

Substituting this into:  $1 - \bar{p}_c = \frac{1}{\bar{k} - 1}$ ,

where  $\bar{k} = \left( \frac{2-\alpha}{3-\alpha} \right) \frac{\bar{K}^{3-\lambda} - m^{3-\lambda}}{\bar{K}^{2-\lambda} - m^{2-\lambda}}$ , gives the critical threshold.

There exists a finite percolation threshold even for networks robust to random removal

# Critical Threshold Scale Free

General result:

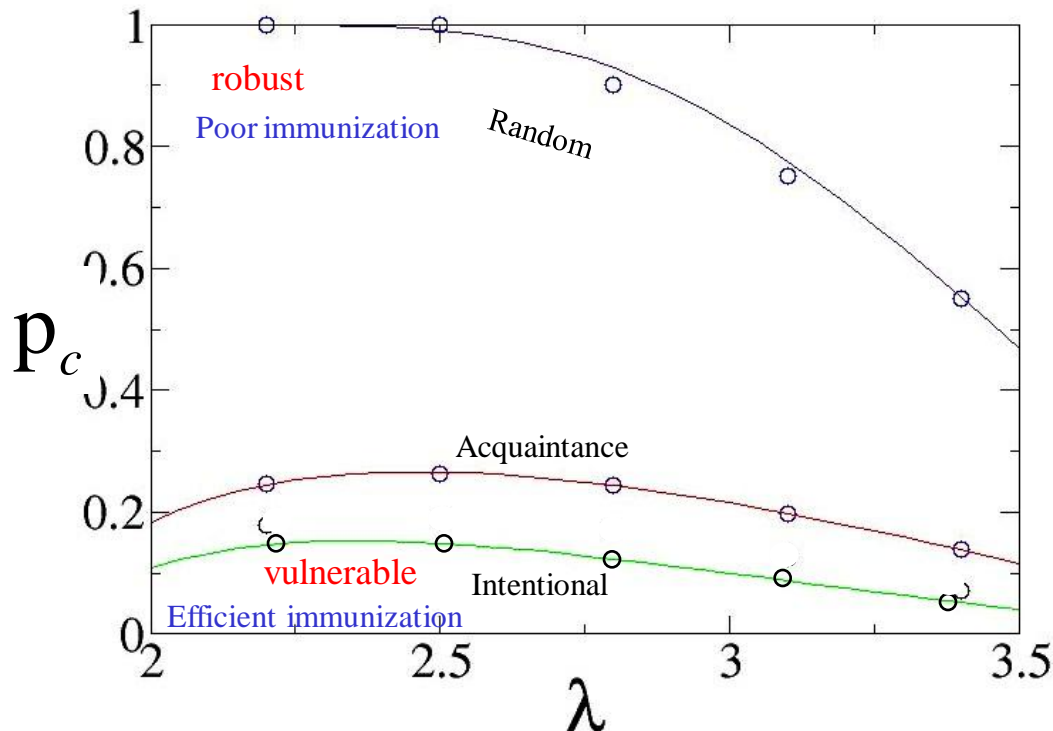
$$p_c = \frac{1}{K_0 - 1}$$

$$K_0 \equiv \frac{\langle k^2 \rangle}{\langle k \rangle}$$

For Poisson:

$$K_0 = \frac{\langle k^2 \rangle}{\langle k \rangle} = \frac{\langle k \rangle^2 + \langle k \rangle}{\langle k \rangle}$$

$$p_c = \frac{1}{\langle k \rangle}$$



Cohen et al. Phys. Rev. Lett. 91, 168701 (2003)

Efficient Immunization  
Strategies:

Acquaintance Immunization

# Critical Exponents

Using the properties of power series (generating functions) near a singular point (Abelian methods), the behavior near the critical point can be studied.

(Diff. Eq. Melloy & Reed (1998) Gen. Func. Newman Callaway PRL(2000), PRE(2001))

**For random breakdown the behavior near criticality in scale-free networks is different than for random graphs or from mean field percolation. For intentional attack-same as mean-field.**

Even for networks with  $3 < \lambda < 4$ , where  $\langle k \rangle$  and  $\langle k^2 \rangle$  are finite, the critical exponents change from the known mean-field result  $\beta = 1$ . The order of the phase transition and the exponents are determined by  $\langle k^3 \rangle$ .

**Size of the infinite cluster:**

$$P_{\infty} \sim (p - p_c)^{\beta} \quad \beta = \begin{cases} \frac{1}{3-\lambda} & 2 < \lambda < 3 \\ \frac{1}{\lambda-3} & 3 < \lambda < 4 \\ 1 & \lambda > 4 \end{cases} \quad (\text{known mean field})$$

**Distribution of finite clusters at criticality:**

$$n_s \sim s^{-\tau} \quad \tau = \begin{cases} \frac{2\lambda-3}{\lambda-2} & \lambda < 4 \\ 2.5 & \lambda \geq 4 \end{cases} \quad (\text{known mean field})$$

# Fractal Dimensions

From the behavior of the critical exponents the fractal dimension of scale-free graphs can be deduced.

Far from the critical point - the dimension is infinite - the mass grows exponentially with the distance.

At criticality - the dimension is finite for  $\lambda > 3$ .

Random Graphs – Erdos Renyi(1960)

Largest cluster at criticality

$$S \propto N^{\frac{2}{3}}$$

Scale Free networks

$$S \propto R^{d_f} \propto N^{\frac{d_f}{d_c}} \propto N^{\frac{\lambda-2}{\lambda-1}} \quad \lambda \leq 4$$

$$S \propto N^{\frac{2}{3}} \quad \lambda \geq 4$$

**Chemical dimension:**

$$d_l = \begin{cases} \frac{\lambda-2}{\lambda-3} & \lambda < 4 \\ 2 & \lambda \geq 4 \end{cases}$$

$$S \propto \ell^{d_l}$$

**Fractal dimension:**

$$d_f = \begin{cases} 2 \frac{\lambda-2}{\lambda-3} & \lambda < 4 \\ 4 & \lambda \geq 4 \end{cases}$$

$$S \propto R^{d_f}$$

**Embedding dimension:**  
(upper critical dimension)

$$d_c = \begin{cases} 2 \frac{\lambda-1}{\lambda-3} & \lambda < 4 \\ 6 & \lambda \geq 4 \end{cases}$$

The dimensionality of the graphs depends on the distribution!