

Networks and hypernetworks 1

Topology, dynamics and applications

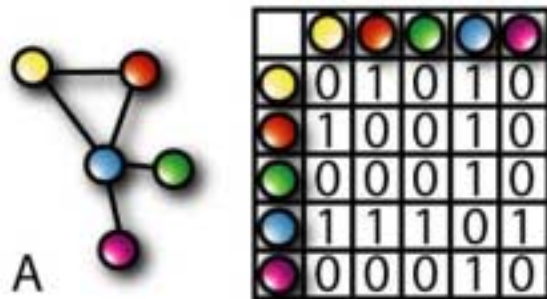
Rui Vilela Mendes

<http://label2.ist.utl.pt/vilela/>

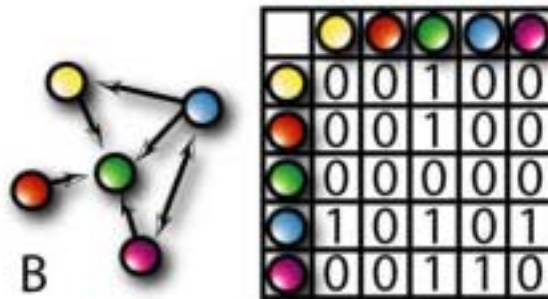
Network basic elements

- Nodes
- Links (directed, undirected, weighted, non-weighted)
- Simple or bipartite
- Adjacency matrix
- Networks are graphs

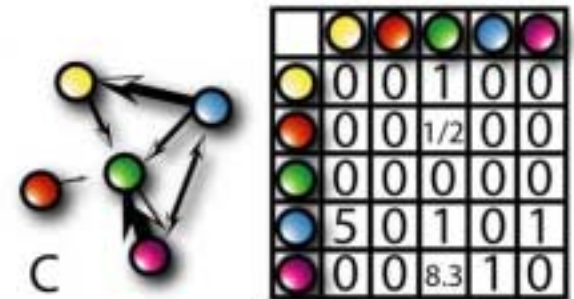
- Simple graph
Symmetric



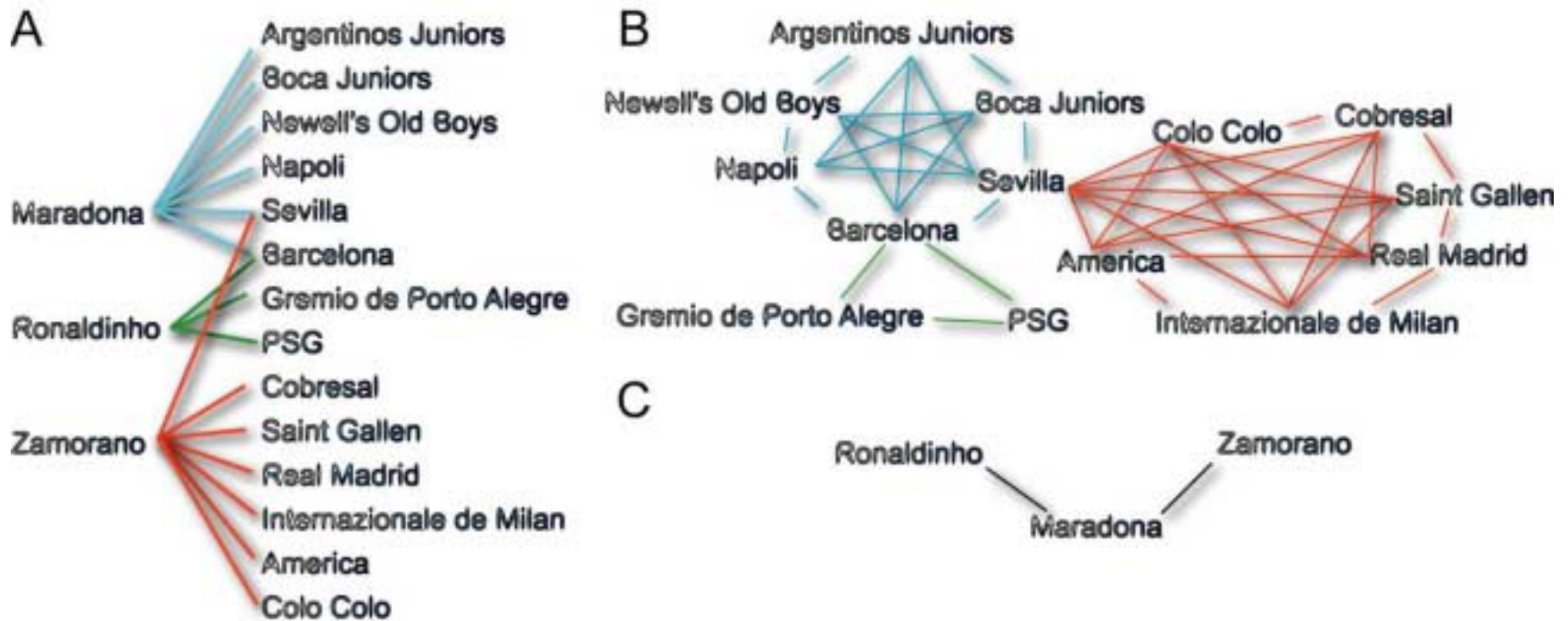
Directed



Directed
Weighted



A bi-partite graph and their derived simple graphs



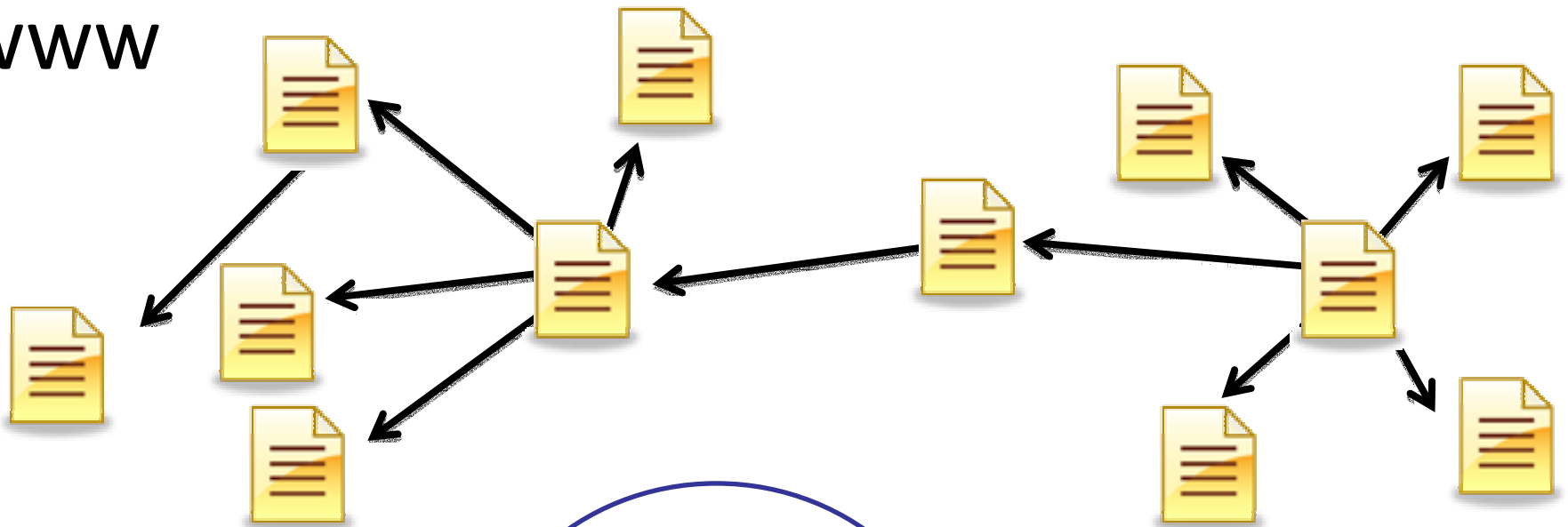
Where does one find networks?



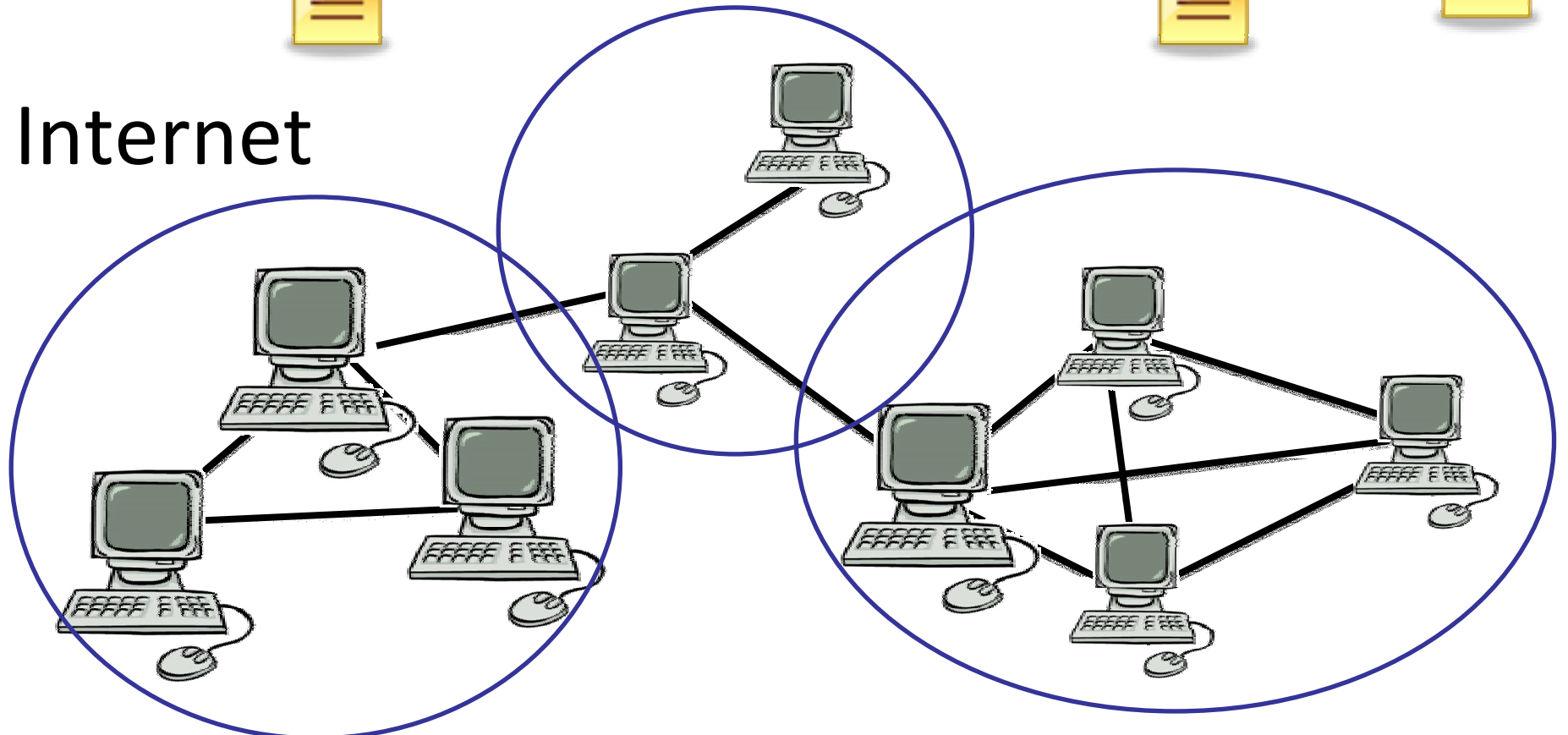
Where does one find networks?

- EVERYWHERE

WWW



Internet

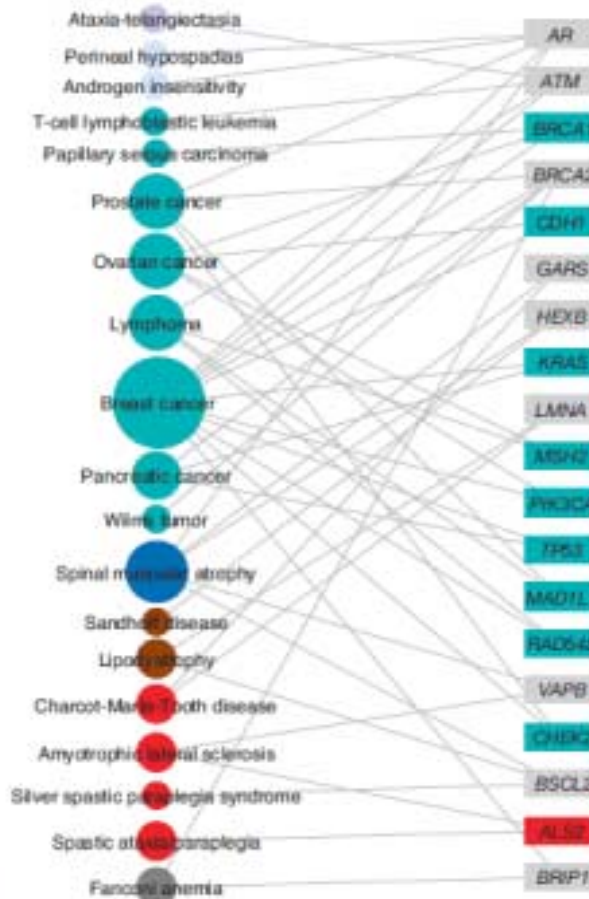


The disease network

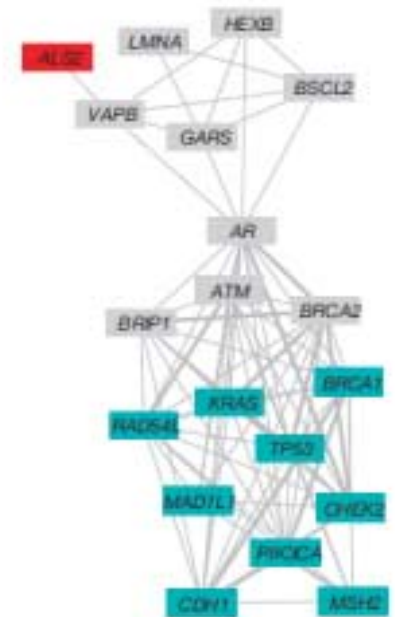
DISEASOME

disease phenotype

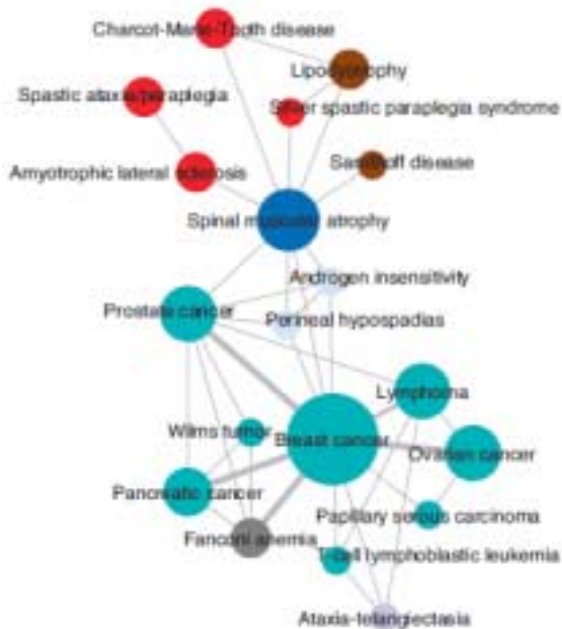
disease genome

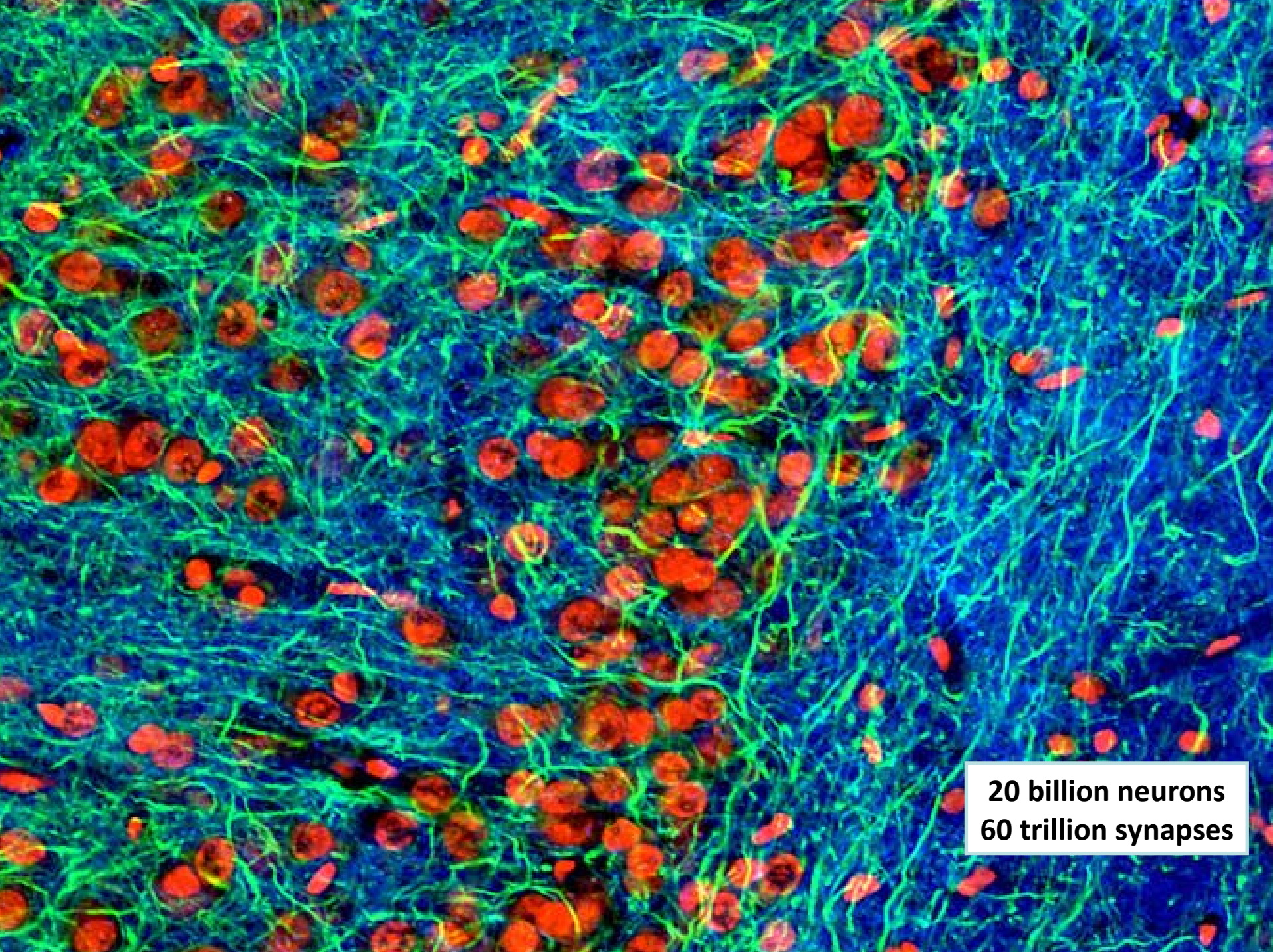


Disease Gene Network (DGN)

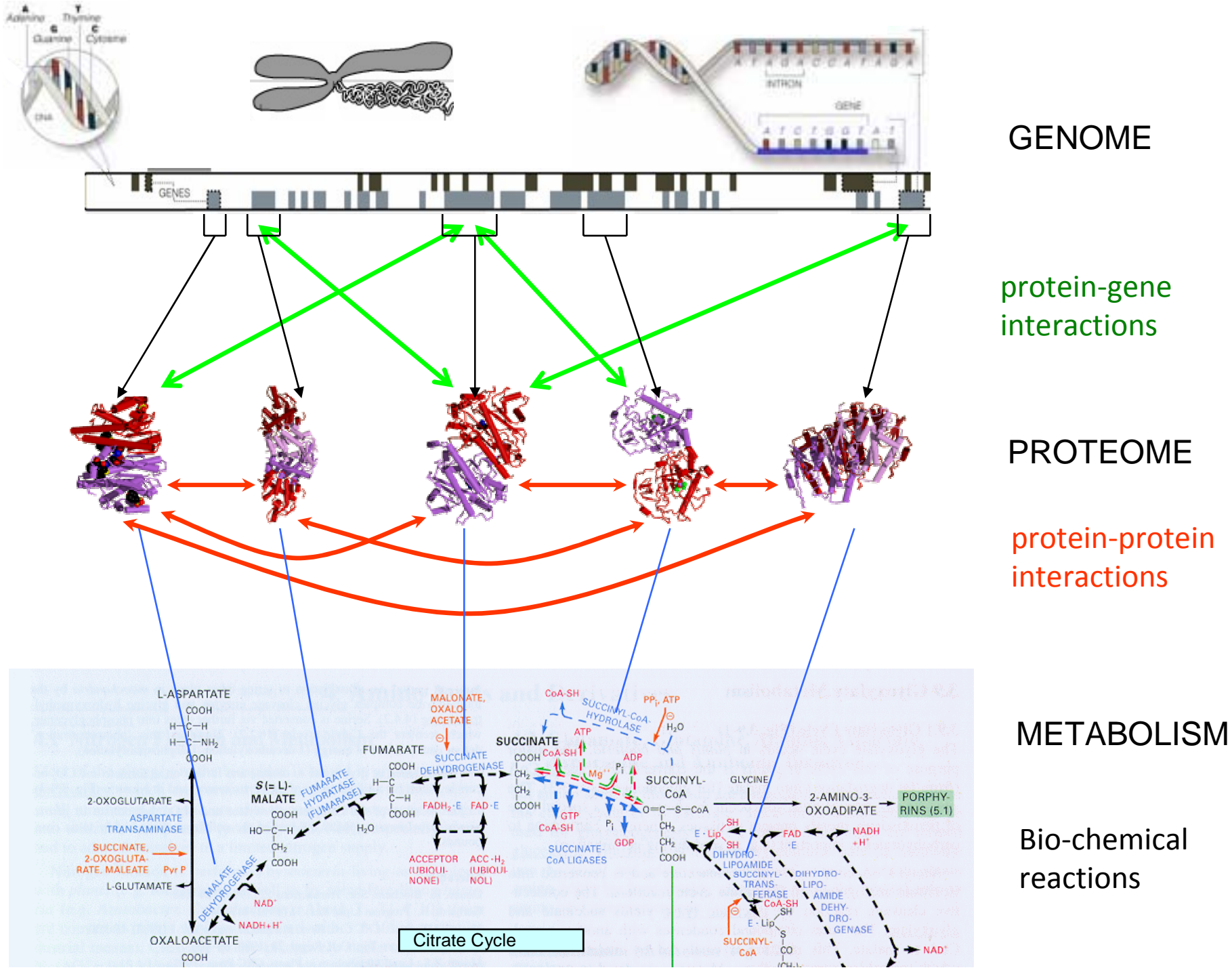


Human Disease Network (HDN)





20 billion neurons
60 trillion synapses



Biochemical Pathways

1

2

3

4

5

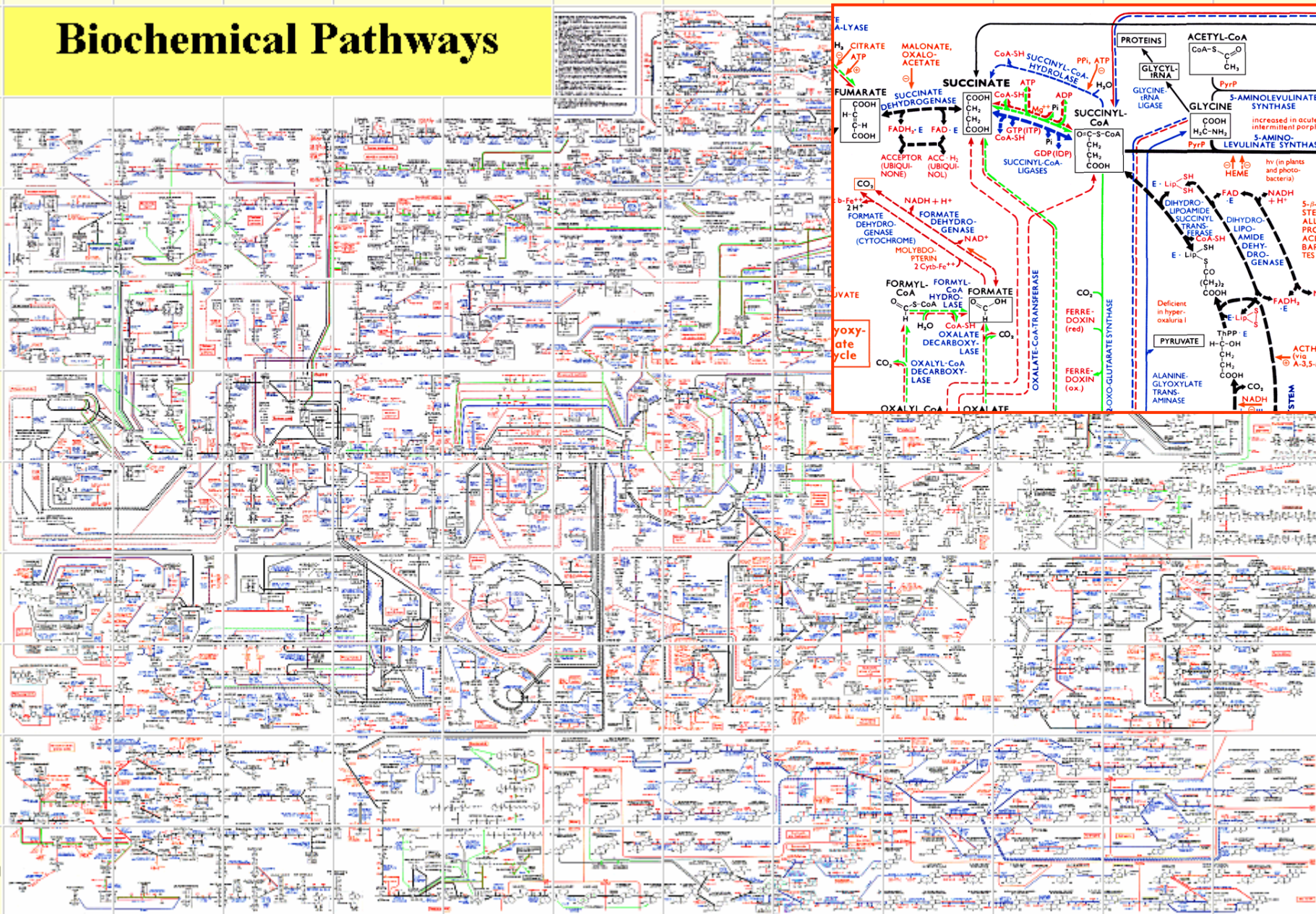
6

7

8

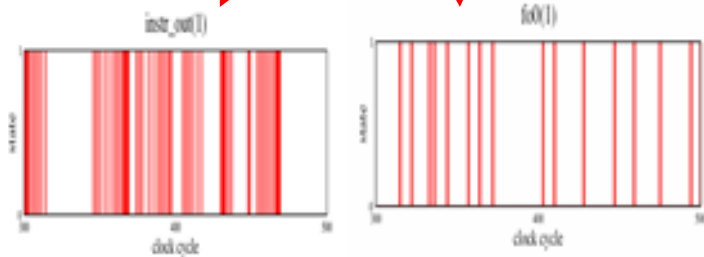
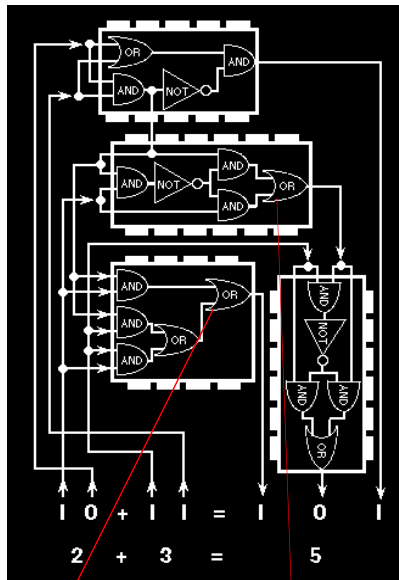
9

10



Computer chip

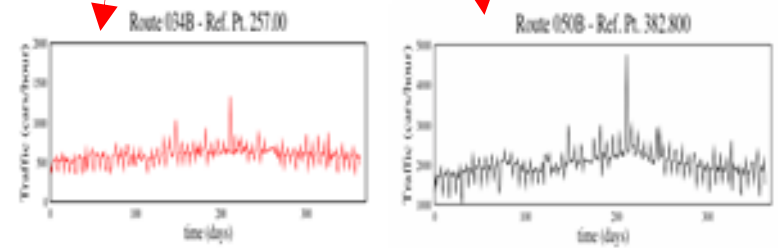
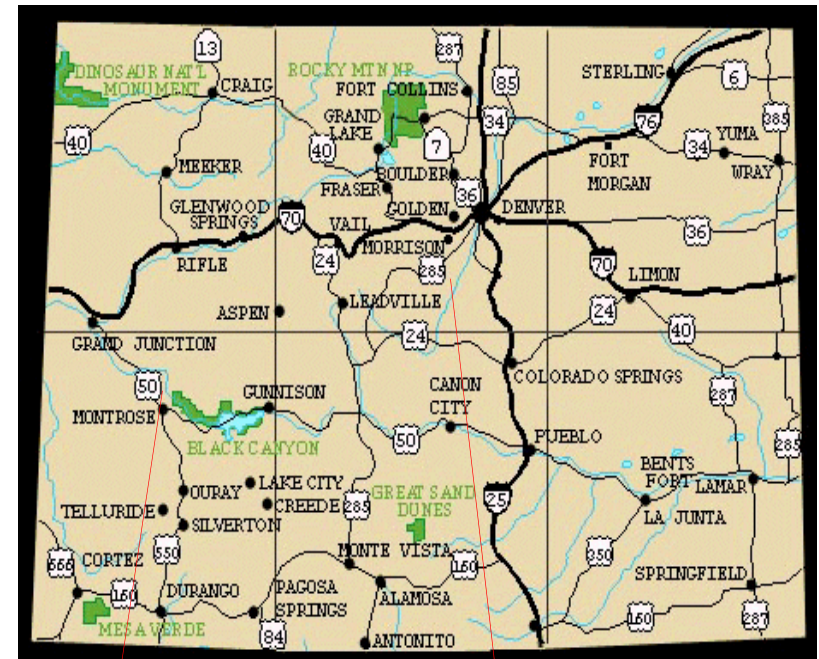
• $f_i(t)$ = state of a given logic component i at clock cycle t .



- 462 signal carriers
- 8,862 clock cycles.

Highways

• $f_i(t)$ = traffic at a given point of a road i at day t .

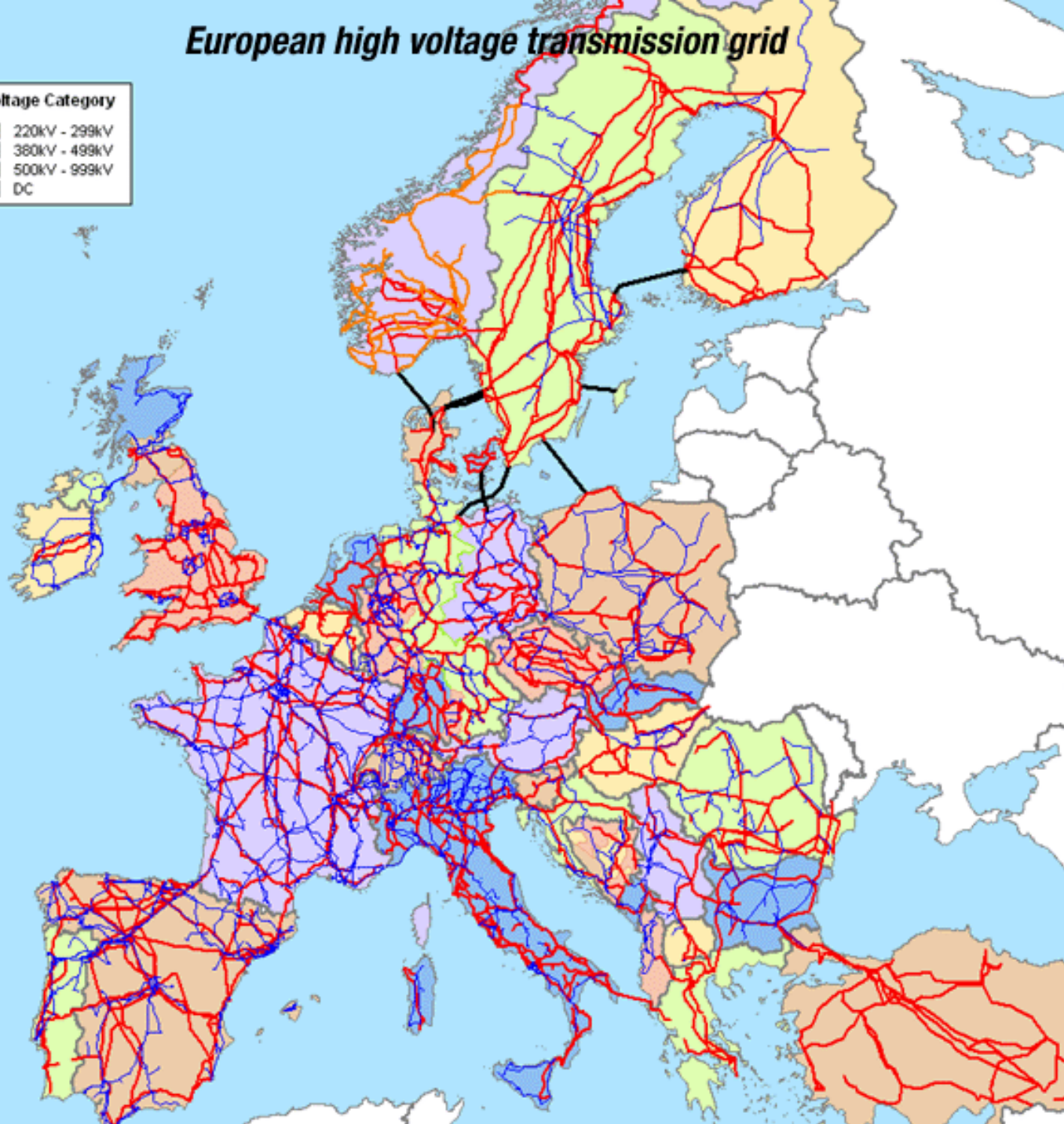


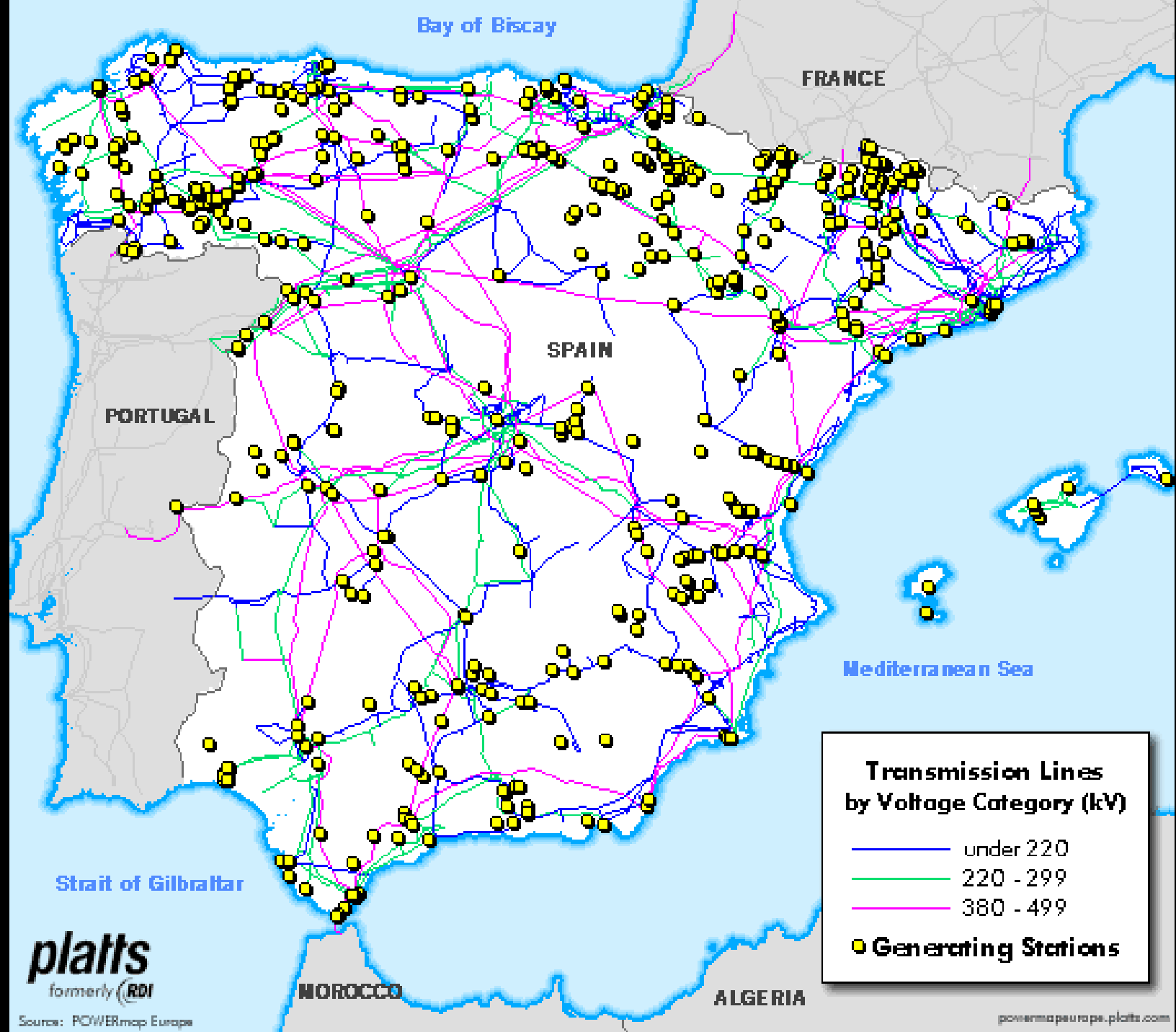
- Daily traffic on 127 Colorado roads from 1998 to 2001.

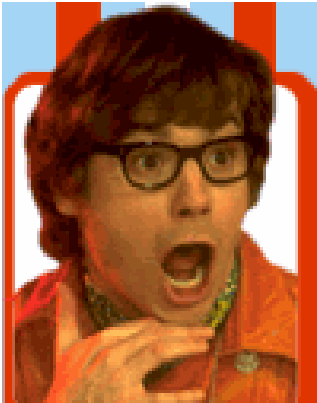
European high voltage transmission grid

Voltage Category

- 220kV - 299kV
- 380kV - 499kV
- 500kV - 999kV
- DC





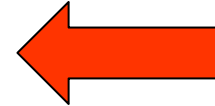


Austin Powers:
The spy who
shagged me

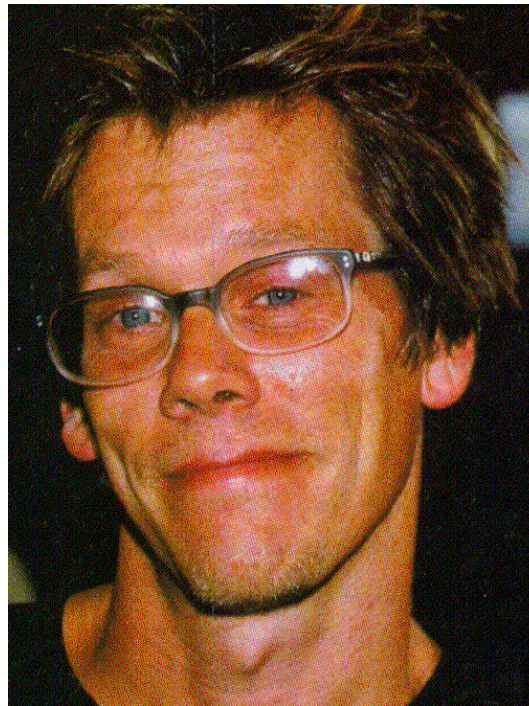


Robert Wagner

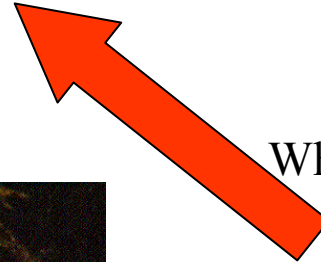
Let's make
it legal



Wild Things



What Price Glory



Barry Norton



Monsieur
Verdoux



A Few
Good Men





Characterizing networks: Topological parameters

- Degree
- Path length
- Clustering
- Centrality
- Betweenness
- Eigenvalue centrality
- Motifs

Local and global measures

Local Measures

Centrality measures (degree, closeness, betweenness, eigenvector, page-rank)

Clustering measures (Clustering, Topological Overlap or Mutual Clustering)

Motifs

Global Measures

Degree Correlations, Correlation Profile.

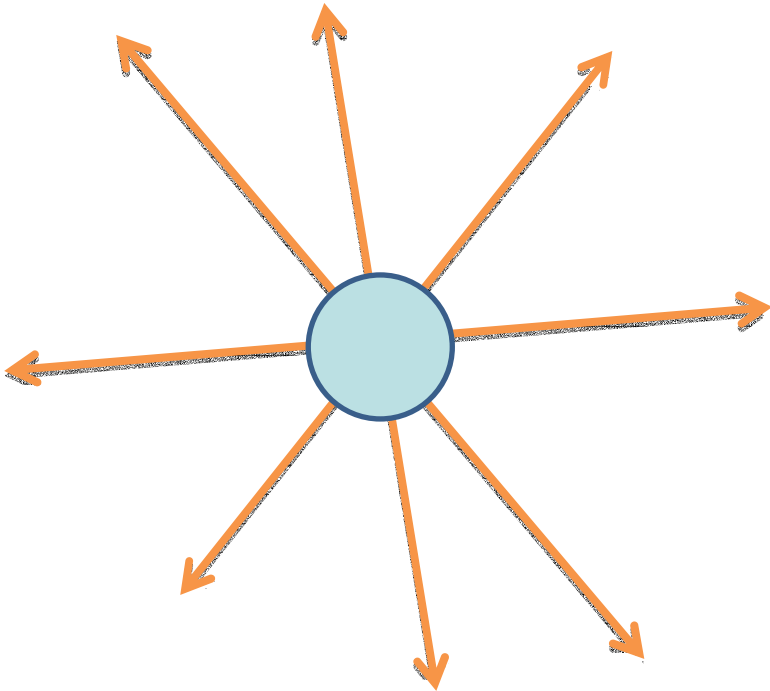
Hierarchical Structure

Fractal Structure

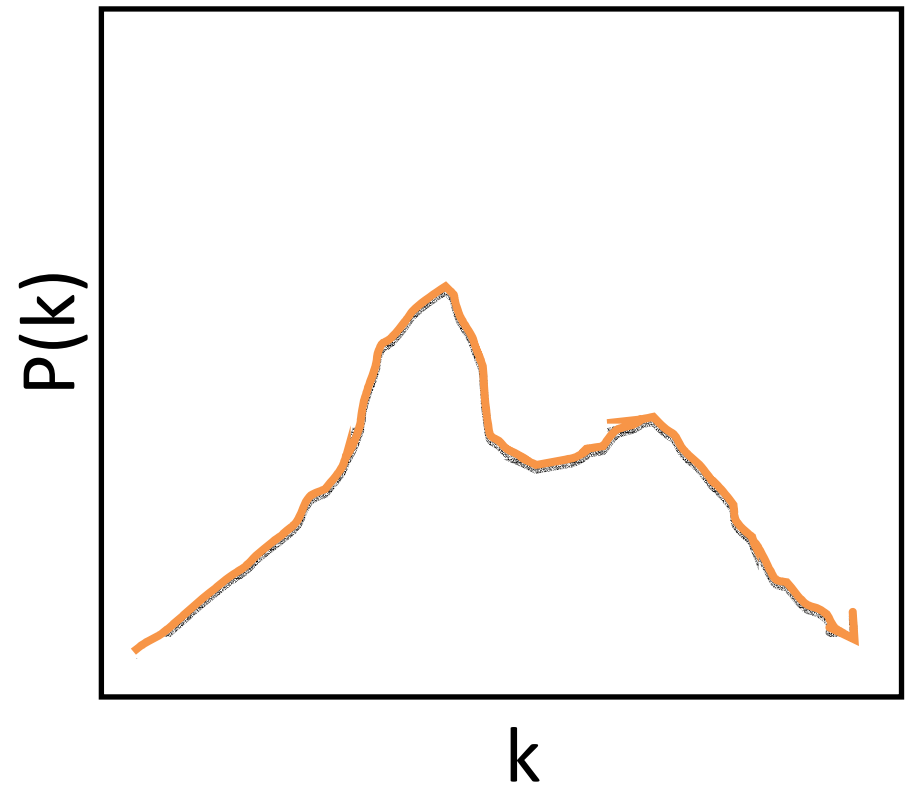
Communities

Modularity

Degree (k)

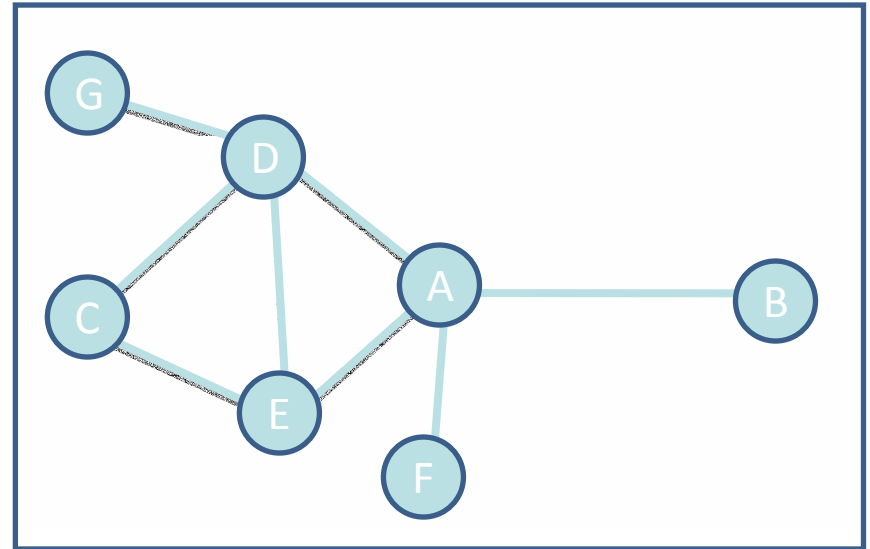


Degree Distribution



Average Path Length

	A	B	C	D	E	F	G
A		1	2	1	1	1	2
B			3	2	2	2	3
C				1	1	3	2
D					1	2	1
E						2	2
F							3
G							



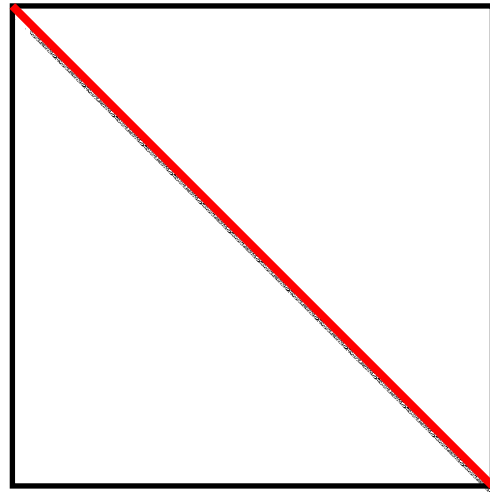
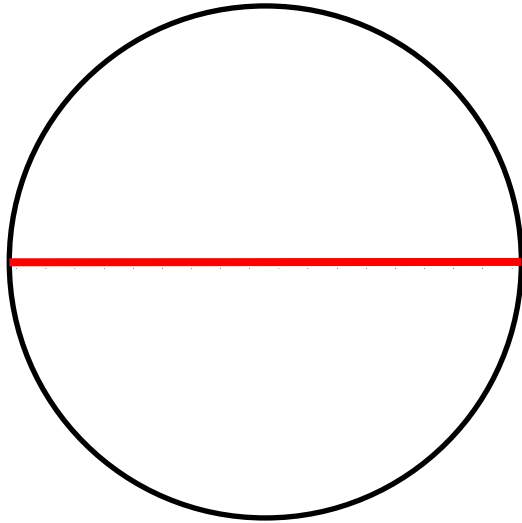
$$D(1)=8$$

$$D(2)=9$$

$$D(3)=4$$

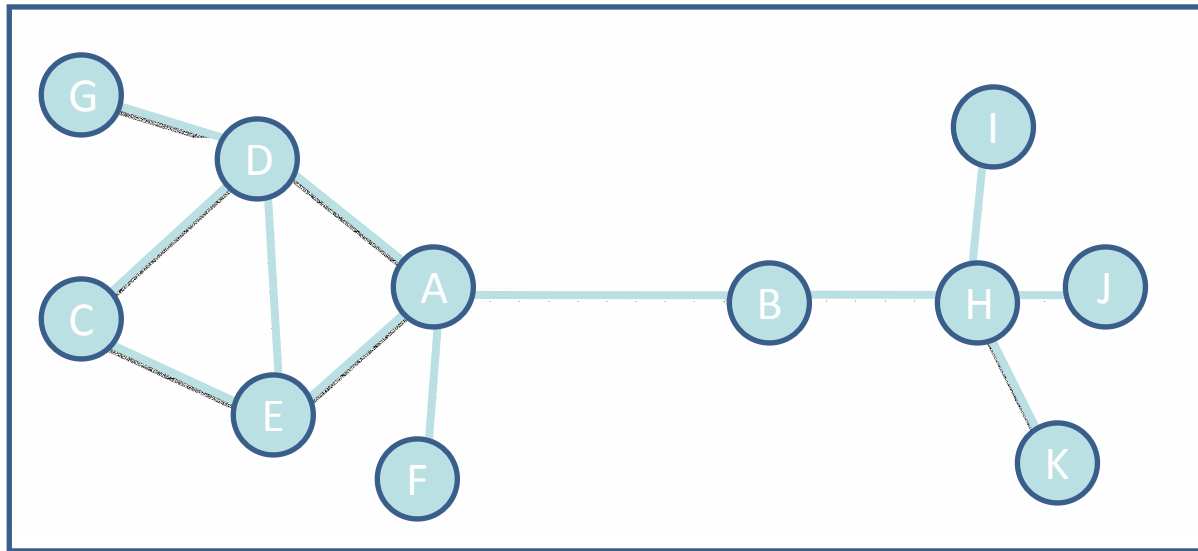
$$L=(8+2 \times 9+3 \times 4)/(8+9+4) \quad L=1.8$$

Diameter



Diameter=Maximum Distance Between Elements in a Set

Diameter= $D(G,J)=D(C,J)=D(G,I)=\dots=5$



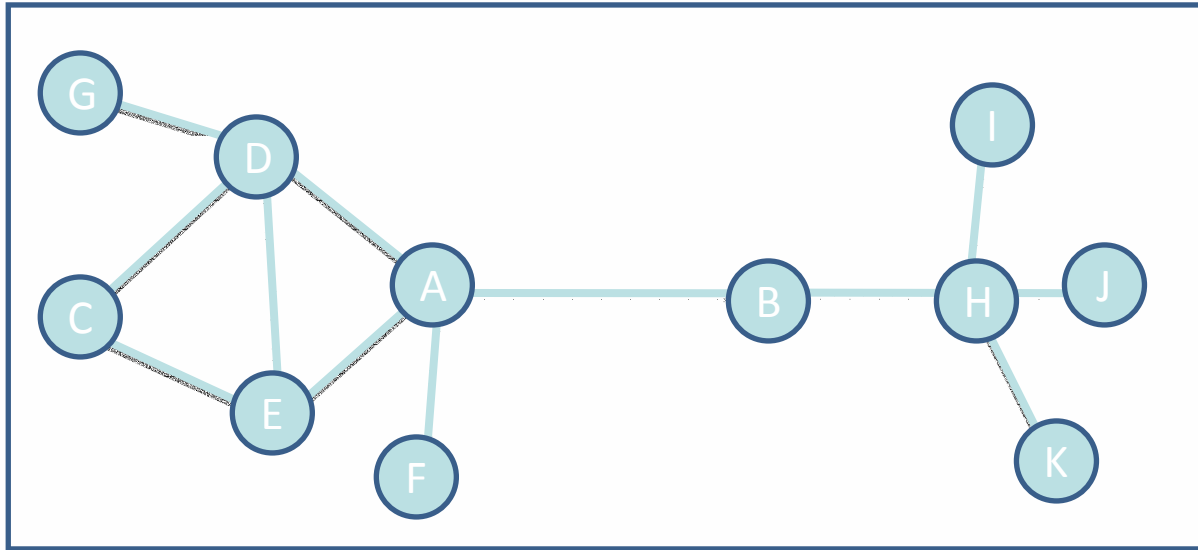
Clustering Coefficient

$$C_i = 2\Delta / k(k-1)$$

$$C_A = 2/12 = 1/6$$

$$C_C = 2/2 = 1$$

$$C_E = 4/6 = 2/3$$



Degree centrality $C(v)$

- N =number of nodes
- *For a node:*
$$C(v_i) = \text{degree}(v_i) / (N-1)$$
- *For a graph:*
$$C(G) = \sum_i (C(v^*) - C(v_i)) / (N-2)$$
- v^* is the node with maximum $C(v)$
- $N-2$ is the maximum value of the sum (a star graph)

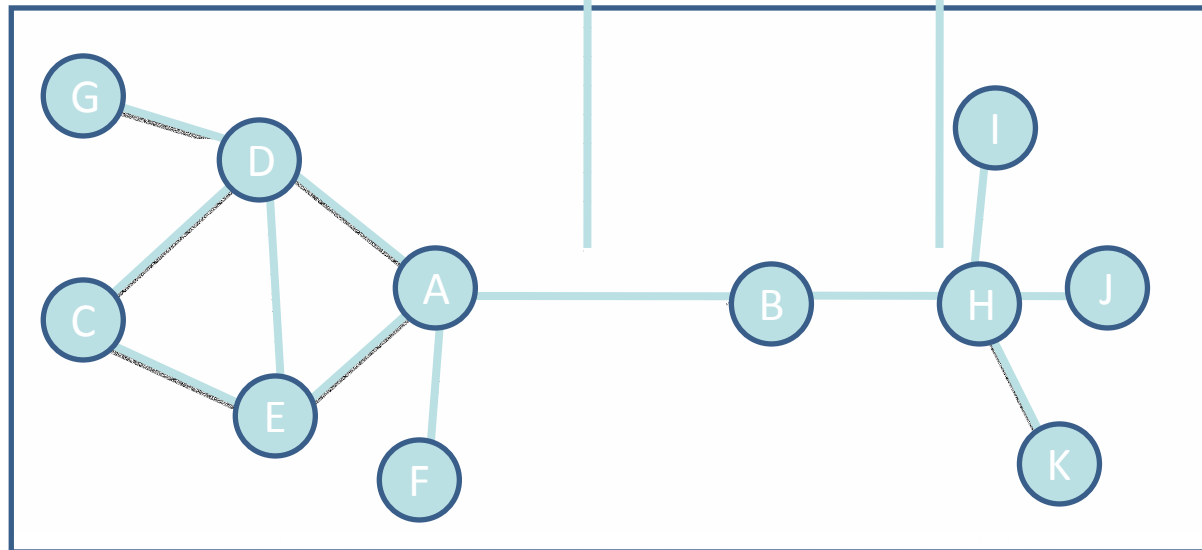
BETWEENNESS CENTRALITY

BC= number of shortest Paths that go through a node.

$$BC(G)=0$$

$$BC(A)=5*5+4=29$$

$$BC(B)=4*6=24$$



N=11

$$C(G) = 1/10(1 + 2 \cdot 3 + 2 \cdot 3 + 4 + 3 \cdot 5)$$

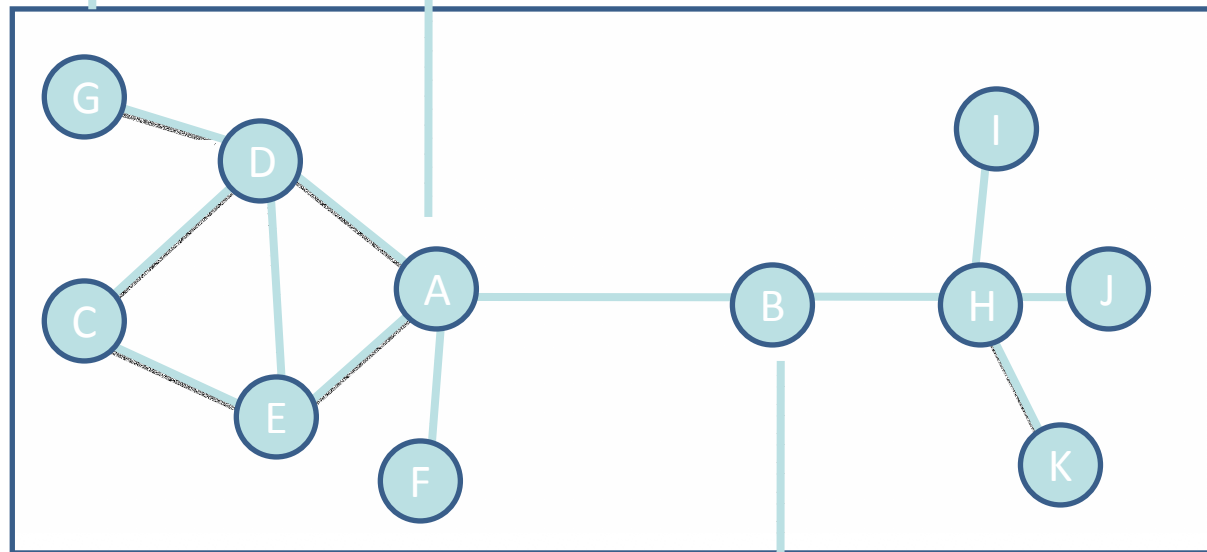
$$C(G) = 3.2$$

$$C(A) = 1/10(4 + 2 \cdot 3 + 3 \cdot 3)$$

$$C(A) = 1.9$$

CLOSENESS CENTRALITY

C = Average Distance to neighbors



$$C(B) = 1/10(2 + 2 \cdot 6 + 2 \cdot 3)$$

$$C(B) = 2$$

N=11

Consider the Adjacency Matrix $A_{ij} = 1$ if node i is connected to node j and 0 otherwise.

Consider the eigenvalue problem:

$$A\mathbf{x} = \lambda\mathbf{x}$$

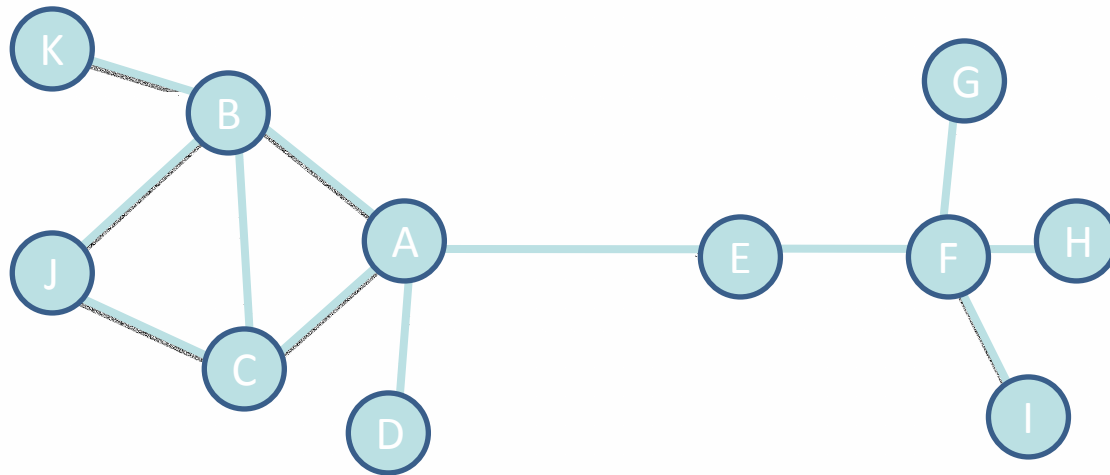
Then the eigenvector centrality of a node is defined as:

$$x_i = \frac{1}{\lambda} \sum_{j \in M(i)} x_j = \frac{1}{\lambda} \sum_{j=1}^N A_{i,j} x_j$$

where λ is the largest eigenvalue associated with A .

PAGE RANK

PR=Probability that a random walker with interspersed Jumps would visit that node.
PR=Each page votes for its neighbors.



$$PR(A) = PR(B)/4 + PR(C)/3 + PR(D) + PR(E)/2$$

A random surfer eventually stops clicking

$$PR(X) = (1-d)/N + d(\sum PR(y)/k(y))$$

PAGE RANK

PR=Probability that a random Walker would visit that node.
PR=Each page votes for its neighbors.

$$\mathbf{R} = \begin{bmatrix} PR(p_1) \\ PR(p_2) \\ \vdots \\ PR(p_N) \end{bmatrix}$$

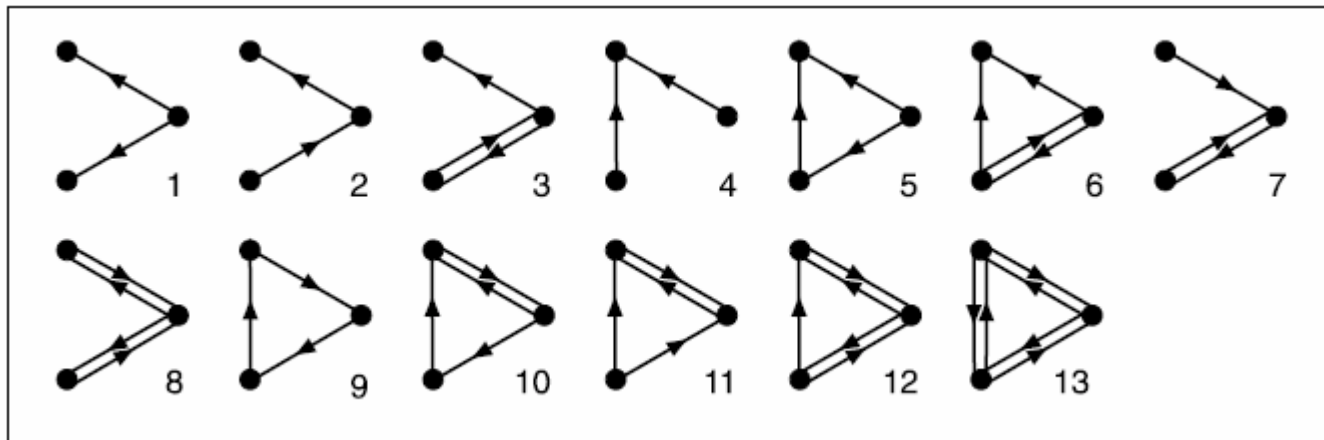
$$\mathbf{R} = \begin{bmatrix} (1-d)/N \\ (1-d)/N \\ \vdots \\ (1-d)/N \end{bmatrix} + d \begin{bmatrix} \ell(p_1, p_1) & \ell(p_1, p_2) & \cdots & \ell(p_1, p_N) \\ \ell(p_2, p_1) & \ddots & & \vdots \\ \vdots & & \ell(p_i, p_j) & \\ \ell(p_N, p_1) & \cdots & & \ell(p_N, p_N) \end{bmatrix} \mathbf{R}$$

$$\sum_{i=1}^N \ell(p_i, p_j) = 1,$$

Network Motifs: Simple Building Blocks of Complex Networks

R. Milo,¹ S. Shen-Orr,¹ S. Itzkovitz,¹ N. Kashtan,¹ D. Chklovskii,²
U. Alon^{1*}

Motifs



Networks are graphs

- The first well-studied graphs:
- Random graphs

Random Graph Theory

Original Formulation:

N nodes, n links chosen randomly from the $N(N-1)/2$ possible links.

Alternative Formulation:

N nodes. Each pair is connected with probability p .
Average number of links $= p(N(N-1))/2$;

Random Graph Theory Works on the limit $N \rightarrow \infty$ and studies when properties on a graph emerge as a function of p .

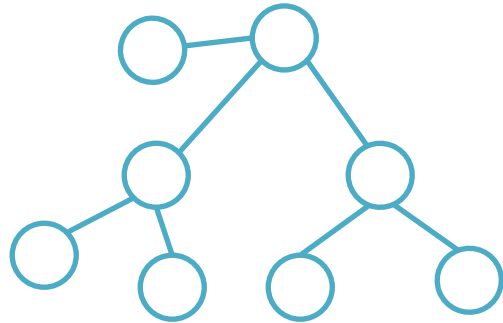


Alfred Renyi

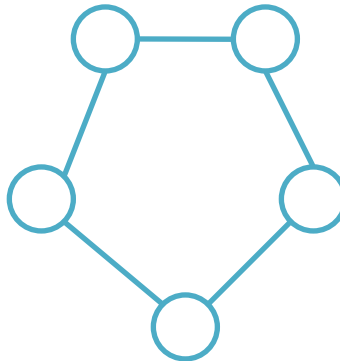
Random Graph Theory: Erdos-Renyi (1959)

Subgraphs

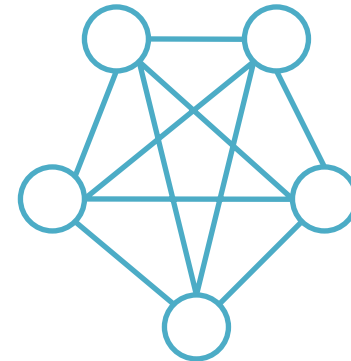
Trees



Cycles



Cliques



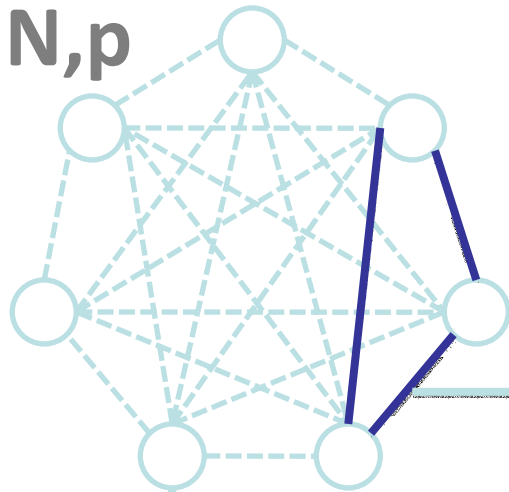
Nodes: k
Links: $k-1$

k
 k

k
 $k(k-1)/2$

Among N nodes choose k

$G_{N,p}$



$F(k,l)$

$$E = C_k^N p^l \frac{k!}{a} \cong N^k p^l / a$$

Each link occurs with
Probability p

Which in the large N
goes like

We can permute the nodes we choose
in $k!$ ways, but have to remember not to double
count isomorphisms (a)

$$E \cong N^k p^l / a$$

One expects the number of subgraphs to be finite if:

$$p(N) \sim cN^{-k/l}$$

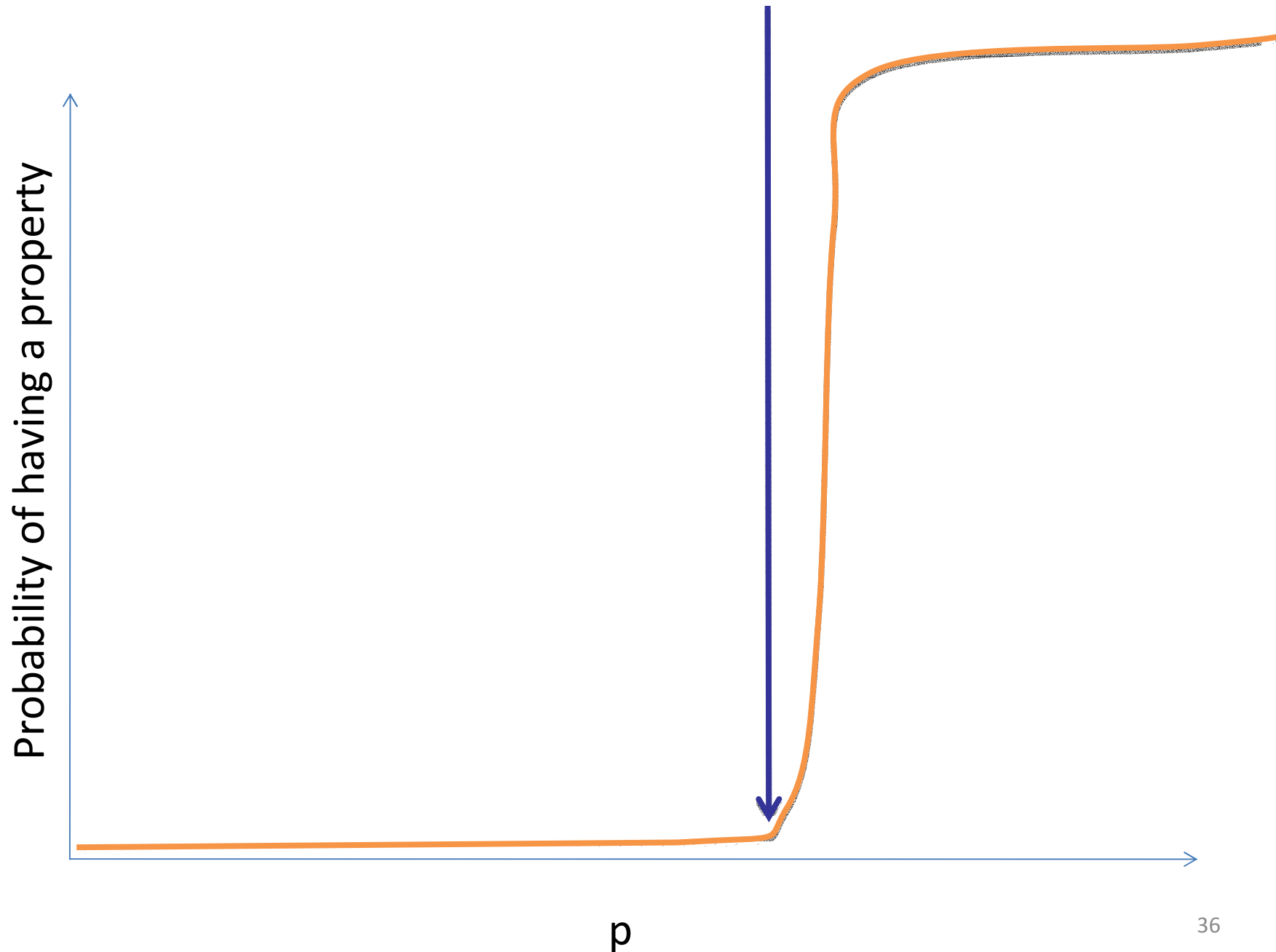


Which implies a number of subgraphs:

$$E = c^l / a = \lambda$$

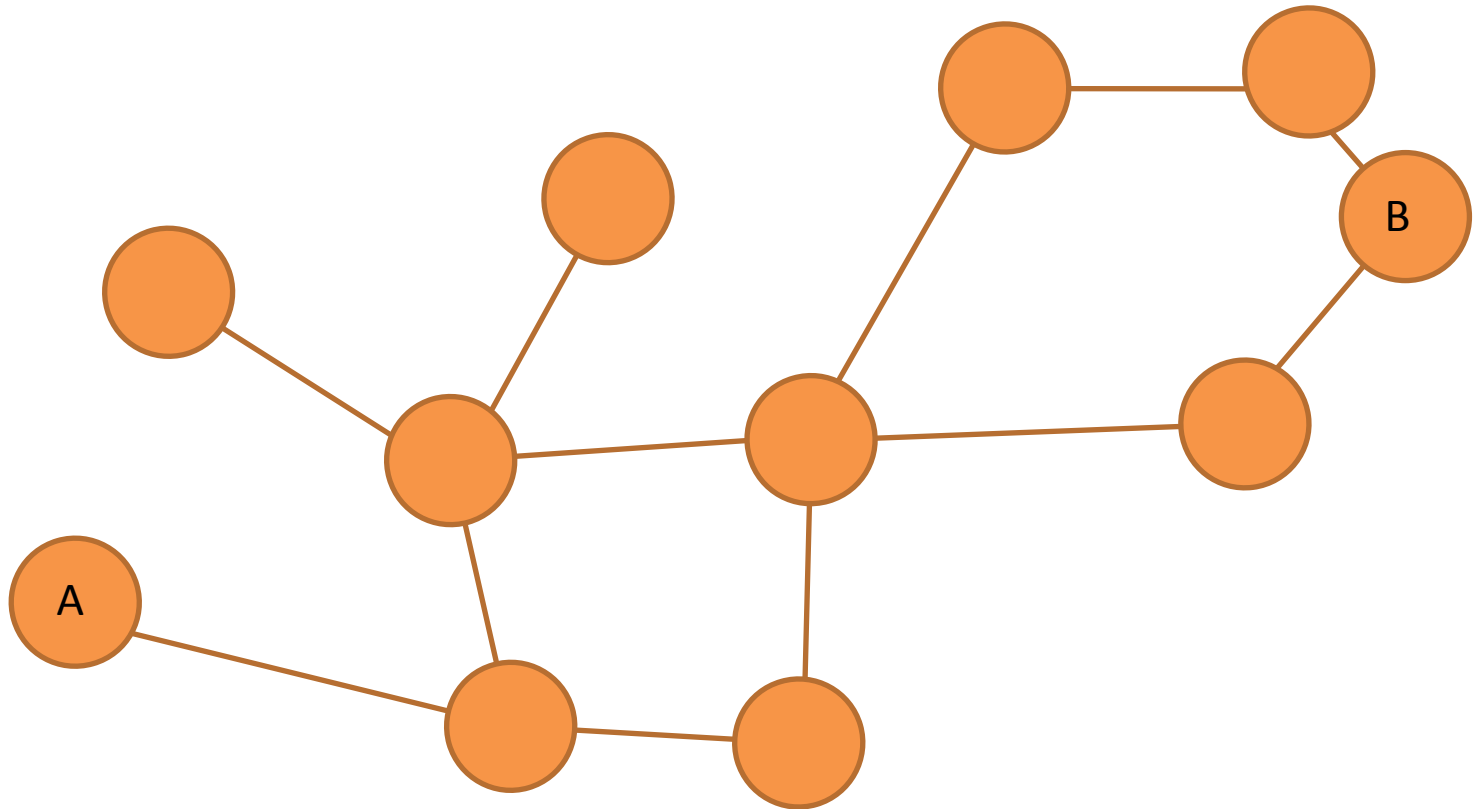
- (a) The critical probability of having a tree of order k is $p_c(N) = cN^{-k/(k-1)}$;
- (b) The critical probability of having a cycle of order k is $p_c(N) = cN^{-1}$;
- (c) The critical probability of having a complete subgraph of order k is $p_c(N) = cN^{-2/(k-1)}$.

Subgraphs appear suddenly (percolation threshold)

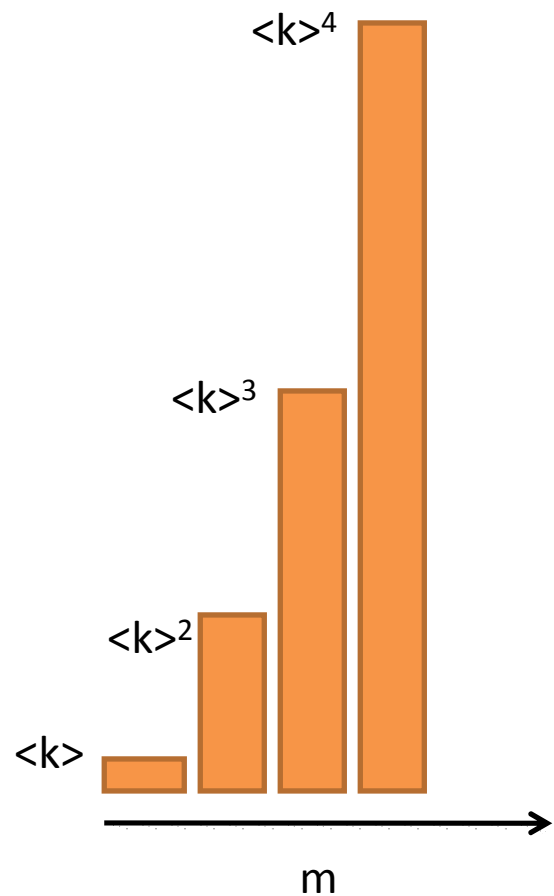


- Average degree $\langle k \rangle = p(N-1) \sim pN$
- Let $p(N) \sim N^z$
- For $z < -3/2$ almost all graphs contain only isolated nodes and edges
- When z passes $-3/2$, trees of order 3 appear
- When z reaches $-4/3$, trees of order 4 appear
- As z approaches -1 , trees of larger and larger order
- But as long as $z < -1$, $\langle k \rangle \sim pN \sim 0$ for large N , the graph is still a union of disjoint trees.
- At $z = -1$ a giant component appears
A connected graph although $\langle k \rangle \sim 1$

Distance Between A and B?



Number of nodes at distance
 m from a randomly chosen node



Hence the average path length is

$$N \sim \langle k \rangle^{\langle l \rangle}$$

$$\log(N) / \log(\langle k \rangle) \sim \langle l \rangle$$

- **Degree distribution**

$$P(k_i = k) = C_k^{N-1} p^k (1-p)^{N-1-k}$$

- X_k = Number of nodes with degree k

$$E(X_k) = NP(k_i = k) = NC_k^{N-1} p^k (1-p)^{N-1-k}$$

- Approaches a Poisson distribution

$$P(X_k) = e^{-\lambda} \frac{\lambda^k}{k!}$$

- **Clustering coefficient**

$$C_{rand} = p = \frac{\langle k \rangle}{N}$$

Are most naturally occurring networks random graphs?

- NO!
- Small worlds
- Clustered
- Scale free

Six Degrees (Stanley Milgram)

160 people

1 person

Stanley Milgram

Map illustrating the concept of Six Degrees of Separation, showing a path from Chicago to Boston, with labels indicating the number of people at each step: 160 people and 1 person.

Stanley Milgram

Six Degrees (Stanley Milgram)

160 people

1 person

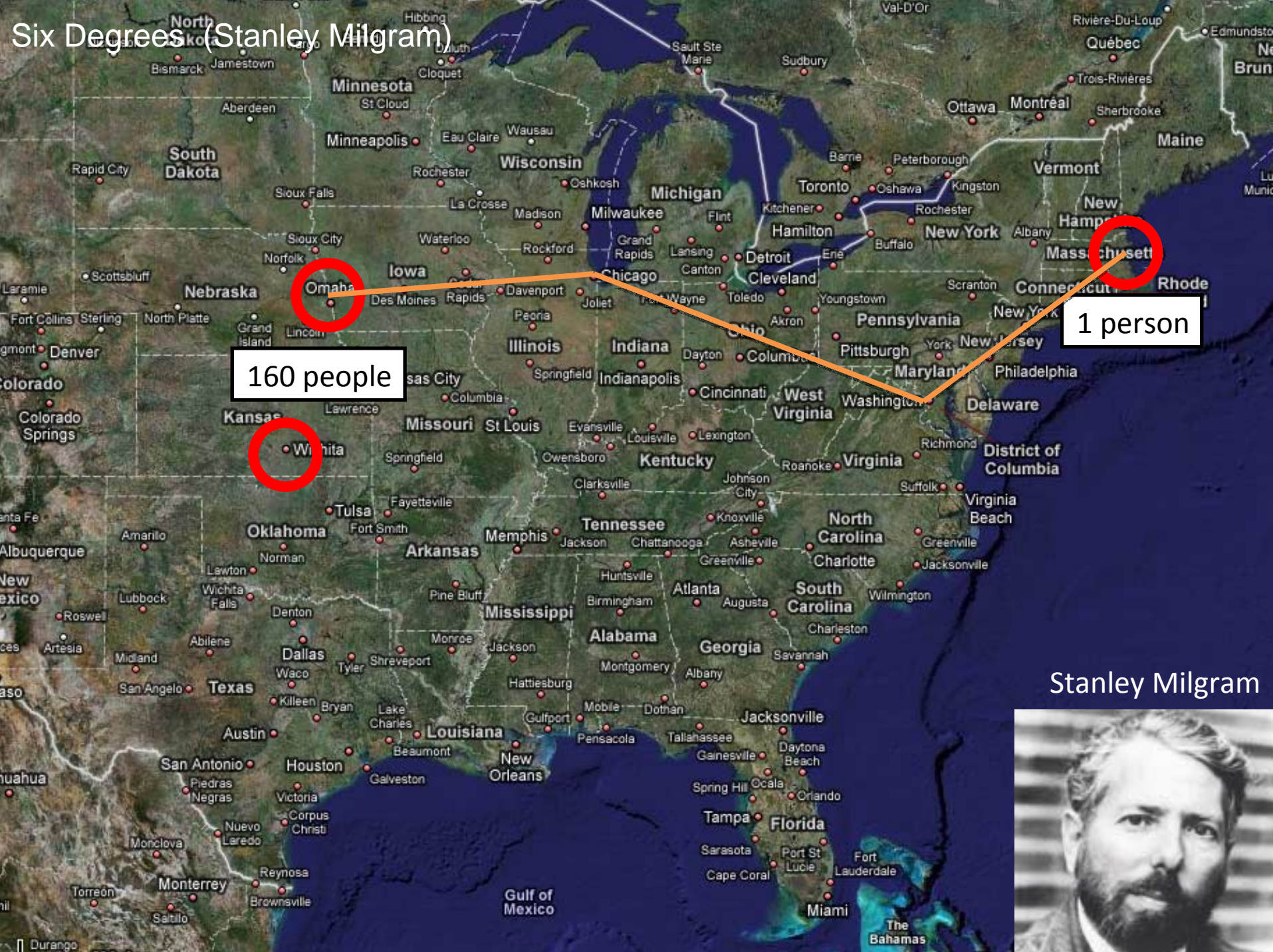
Stanley Milgram

Six Degrees (Stanley Milgram)

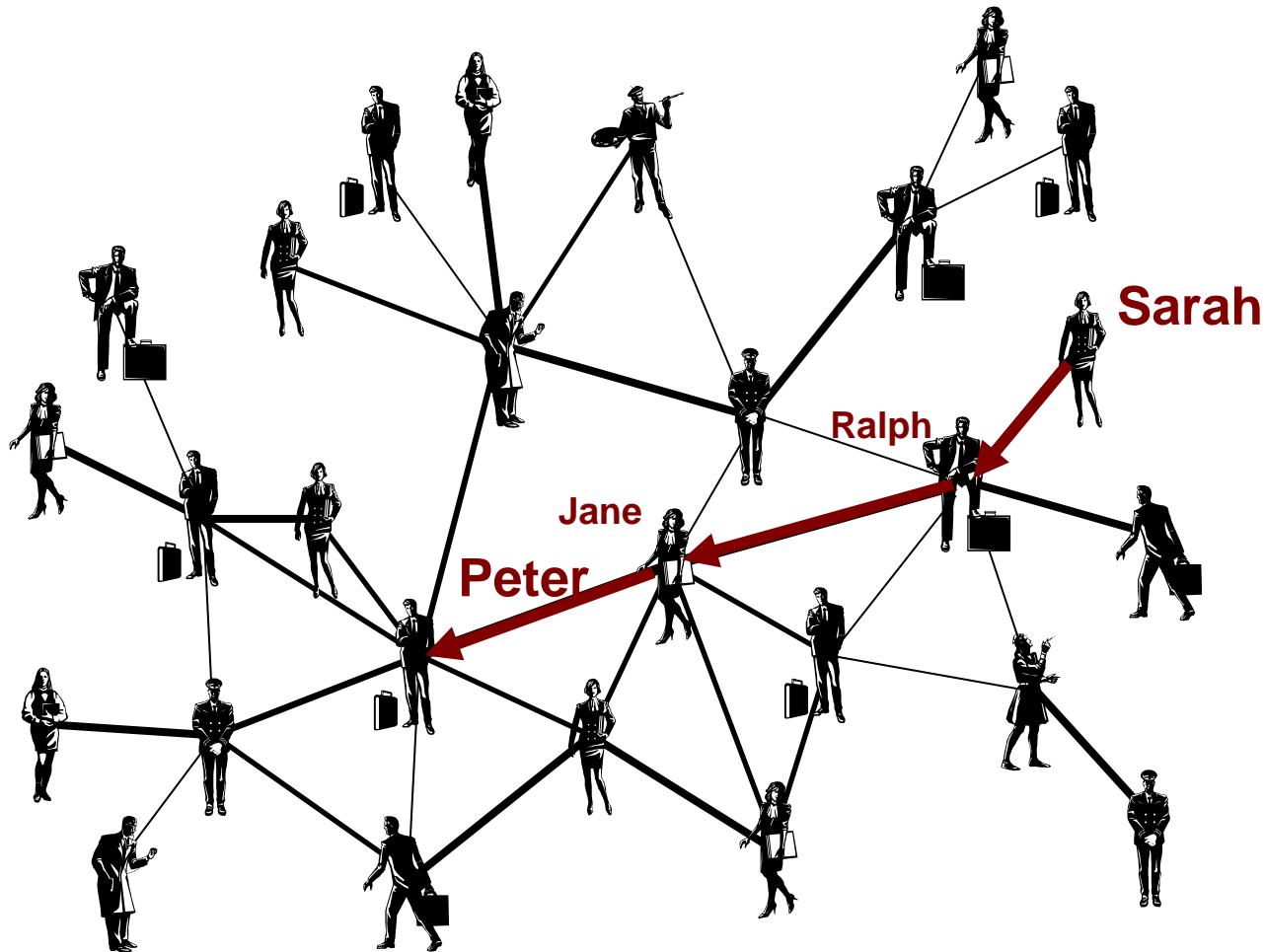
160 people

1 person

Stanley Milgram



Small worlds



Society:

Six degrees

S. Milgram 1967

F. Karinthy 1929

WWW:

19 degrees

Albert *et al.* 1999

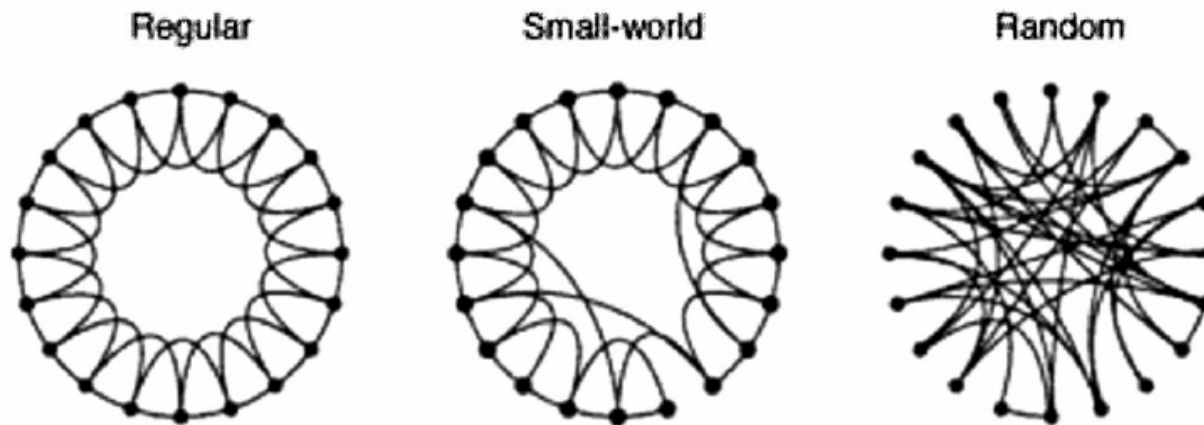
Small Worlds

- Stanley Milgram found that the average length of the chain connecting the sender and receiver was of length 5.5
(The origin of the mythical “Six degrees of separation”)
- 19 degrees in the Web
- Random networks have short path length and small clustering
- Many natural networks have short path length, but high clustering
- The Watts-Strogatz model. Is it the model of natural networks?

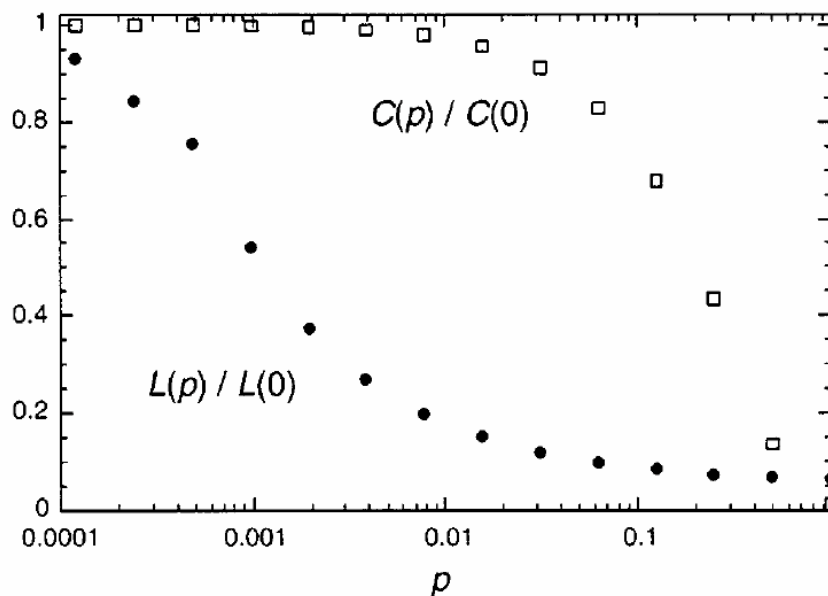
Collective dynamics of 'small-world' networks

letters to nature

Duncan J. Watts* & Steven H. Strogatz



$p = 0$ $\xrightarrow{\text{Increasing randomness}}$ $p = 1$



Duncan Watts



Steve Strogatz

Collective dynamics of ‘small-world’ networks

Duncan J. Watts* & Steven H. Strogatz

Department of Theoretical and Applied Mechanics, Kimball Hall,
Cornell University, Ithaca, New York 14853, USA

Table 1 Empirical examples of small-world networks

	L_{actual}	L_{random}	C_{actual}	C_{random}
Film actors	3.65	2.99	0.79	0.00027
Power grid	18.7	12.4	0.080	0.005
<i>C. elegans</i>	2.65	2.25	0.28	0.05

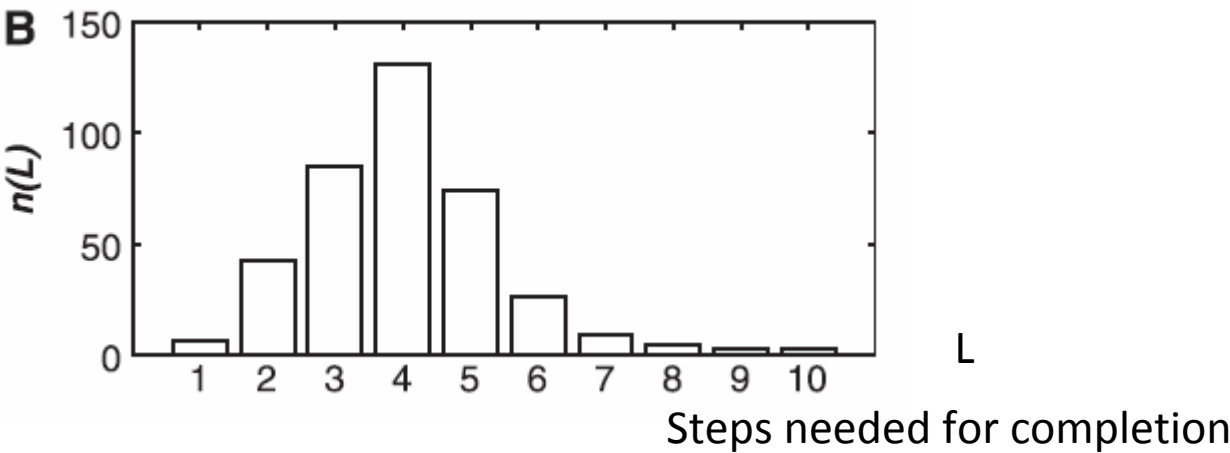
Other small worlds

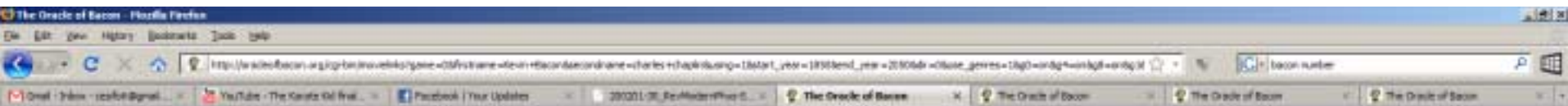


An Experimental Study of Search in Global Social Networks

Peter Sheridan Dodds,¹ Roby Muhamad,² Duncan J. Watts^{1,2*}

We report on a global social-search experiment in which more than 60,000 e-mail users attempted to reach one of 18 target persons in 13 countries by forwarding messages to acquaintances. We find that successful social search is





Help
Credits
How it Works
Contact Us
Other games »

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Heurich. All rights reserved.



Kevin Bacon to charles chaplin Find link More options »



Total number of linkable actors: 953840

Average Kevin Bacon number: 2.946

<u>Kevin Bacon</u> Number	# of People
<u>0</u>	1
<u>1</u>	2108
2	204188
3	601747
4	136178
5	8656
6	839
7	111
8	12

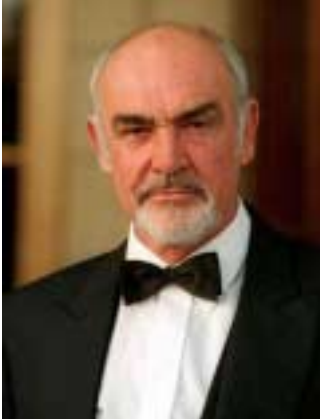
Kevin Bacon



Average Connery number: 2.731

Connery Number	# of people
0	1
1	2272
2	218560
3	380721
4	40263
5	3537
6	535
7	66
8	2

Sean Connery

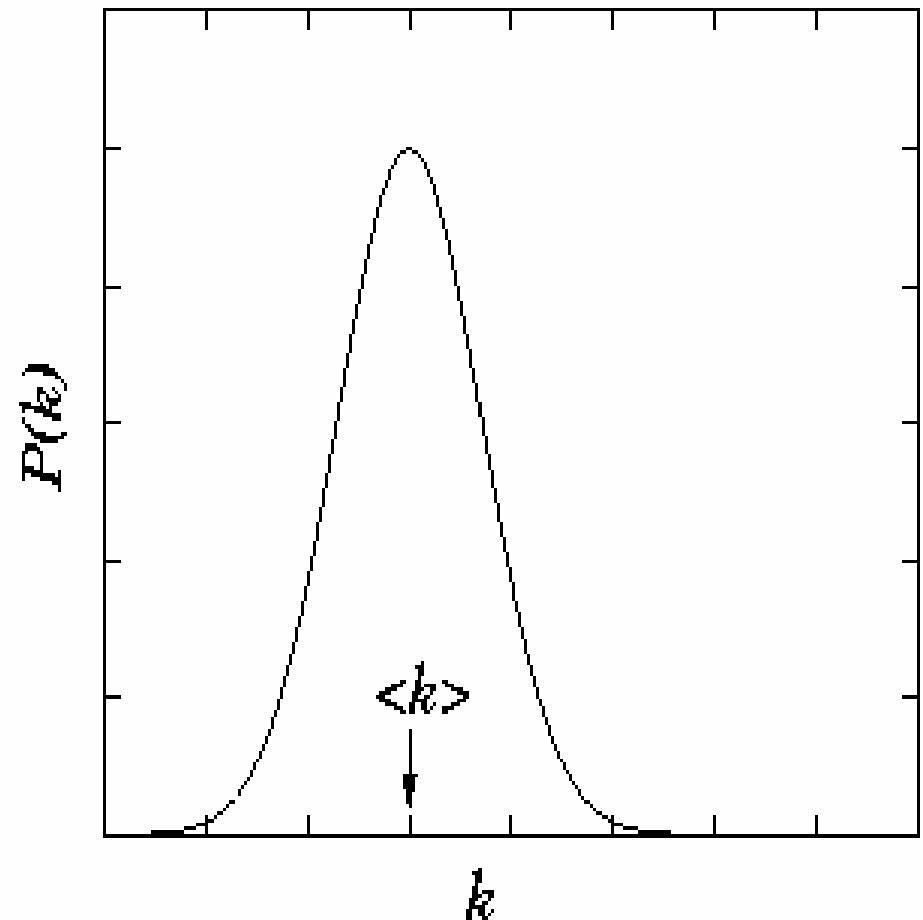
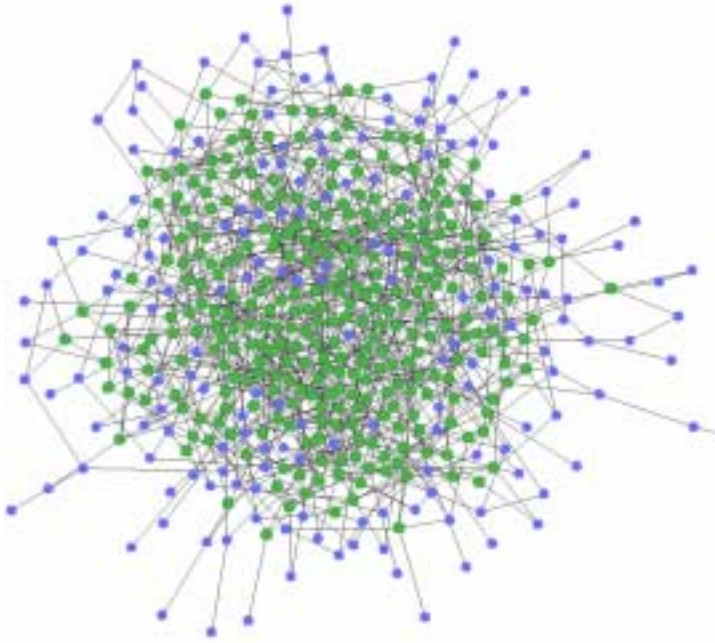


Is Watts-Strogatz the model of most natural occurring networks?

- No!
- Most naturally occurring networks have scale-free degree distribution
- They are small worlds but neither random, nor Watts and Strogatz
- Examples of scale-free networks

Erdős-Rényi model (1960)

Degree distribution (Poisson)



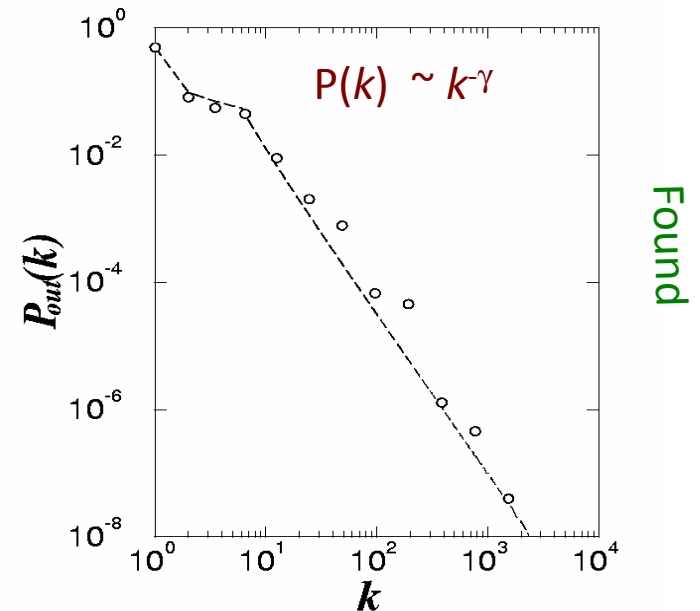
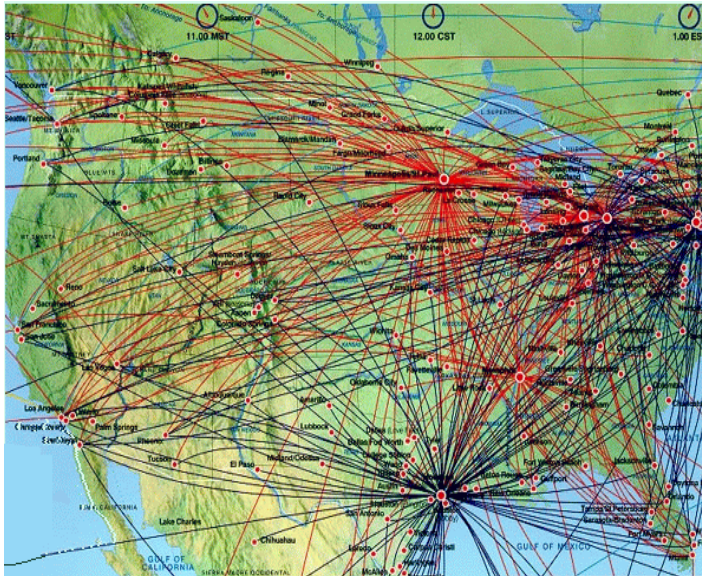
WorldWideWeb

Nodes: WWW documents Links:
URL links

Over 3 billion documents

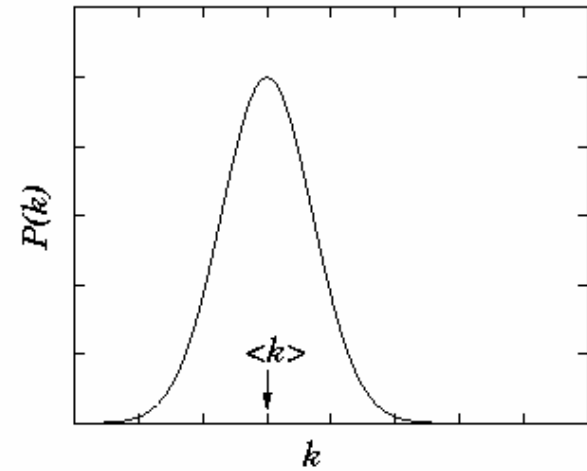
ROBOT: collects all URL's found
in a document and follows them
recursively

Scale-free Network

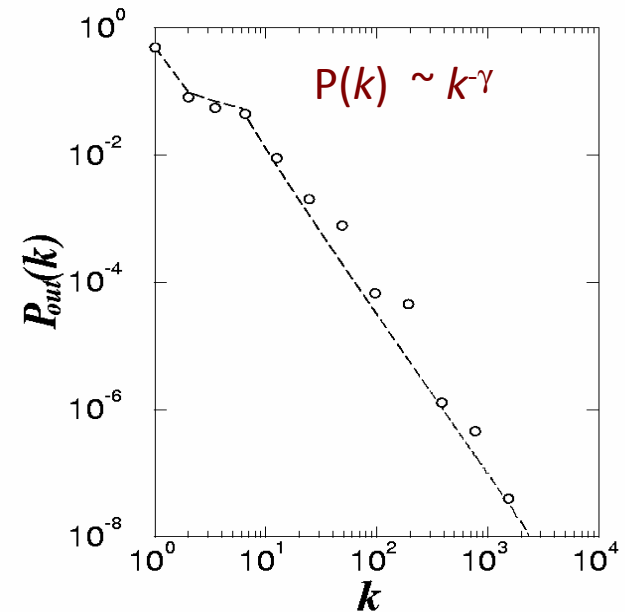
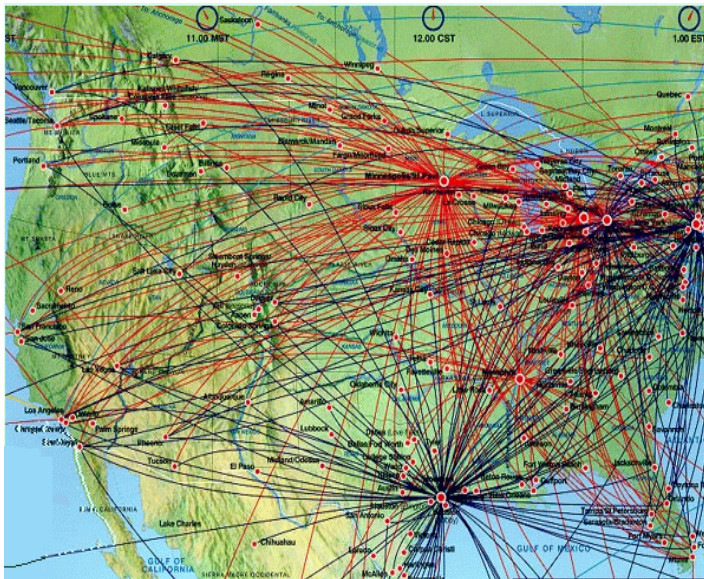


R. Albert, H. Jeong, A-L Barabasi, *Nature*, 401 130 (1999).

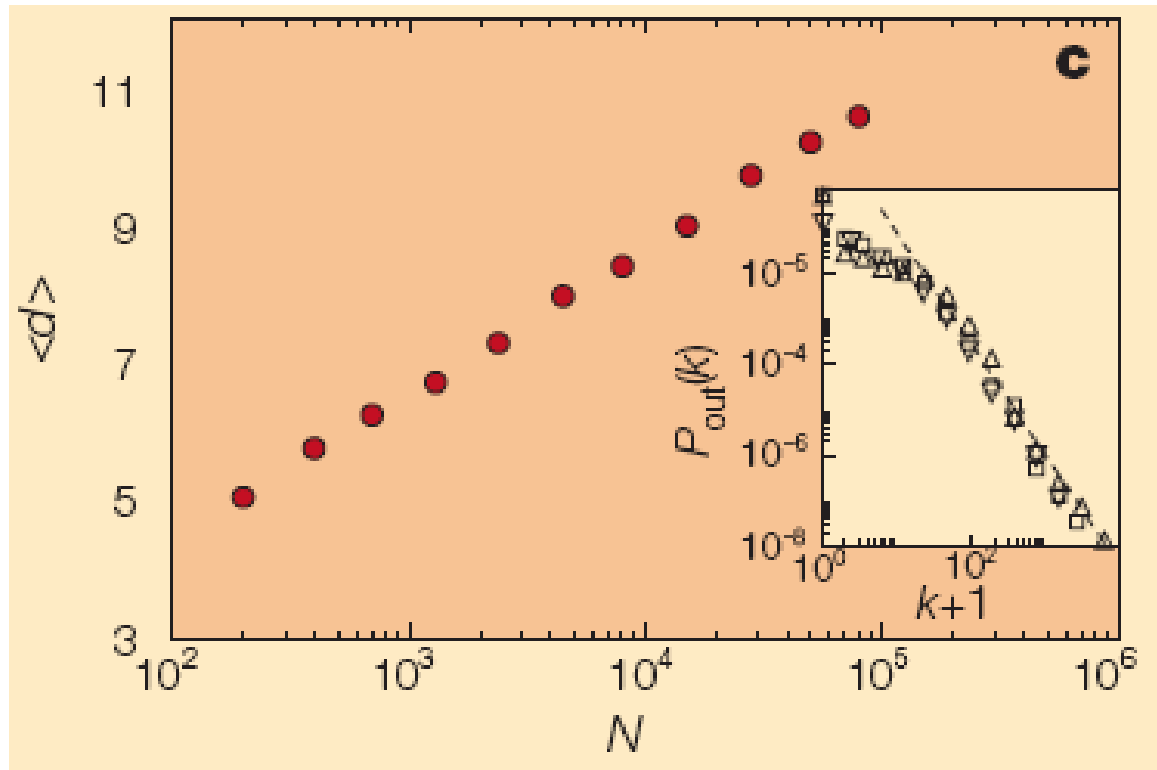
Exponential Network



Scale-free Network



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



Réka Albert, Hawoong Jeong,
Albert-László Barabási

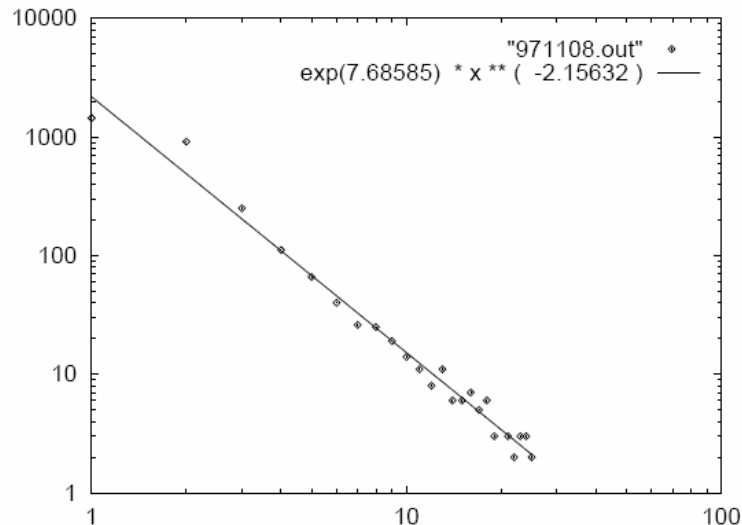
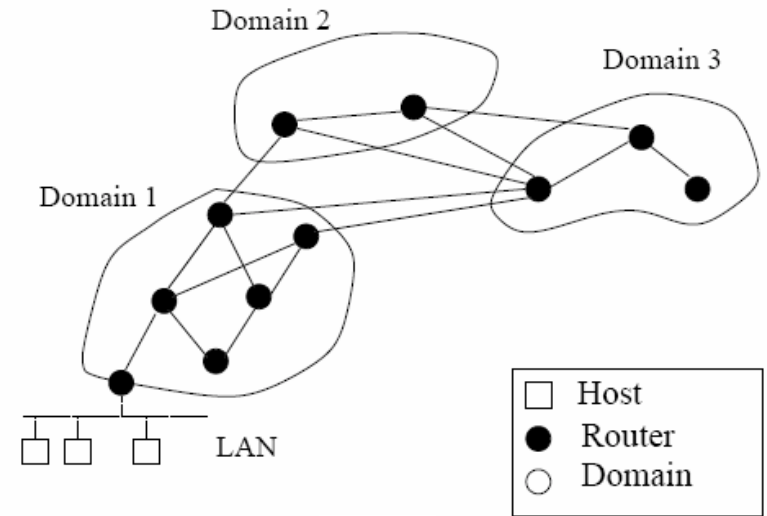
Diameter of the World-Wide Web

"On Power-Law Relationships of the Internet Topology", Michalis Faloutsos, Petros Faloutsos, Christos Faloutsos, ACM SIGCOMM'99, Cambridge, Massachussets, pp 251-262, 1999

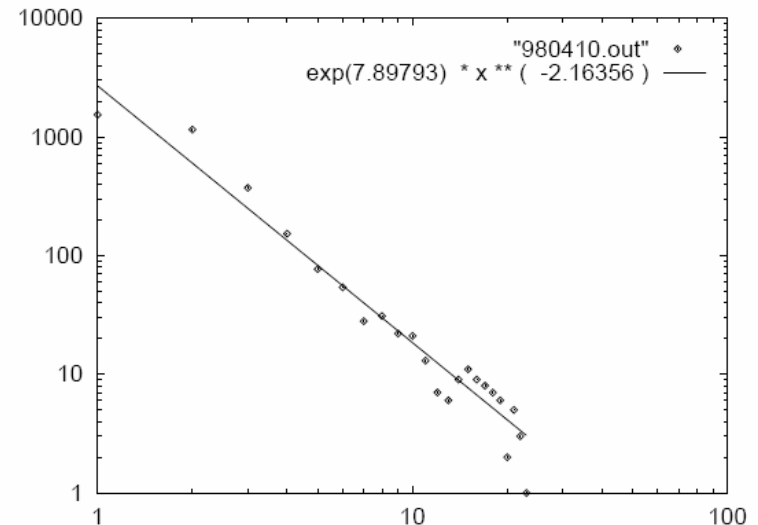
Abstract

Despite the apparent randomness of the Internet, we discover some surprisingly simple power-laws of the Internet topology. These power-laws hold for three snapshots of the Internet, between November 1997 and December 1998, despite a 45% growth of its size during that period. We show that our power-laws fit the real data very well resulting in correlation coefficients of 96% or higher.

Our observations provide a novel perspective of the structure of the Internet. The power-laws describe concisely



(a) Int-11-97



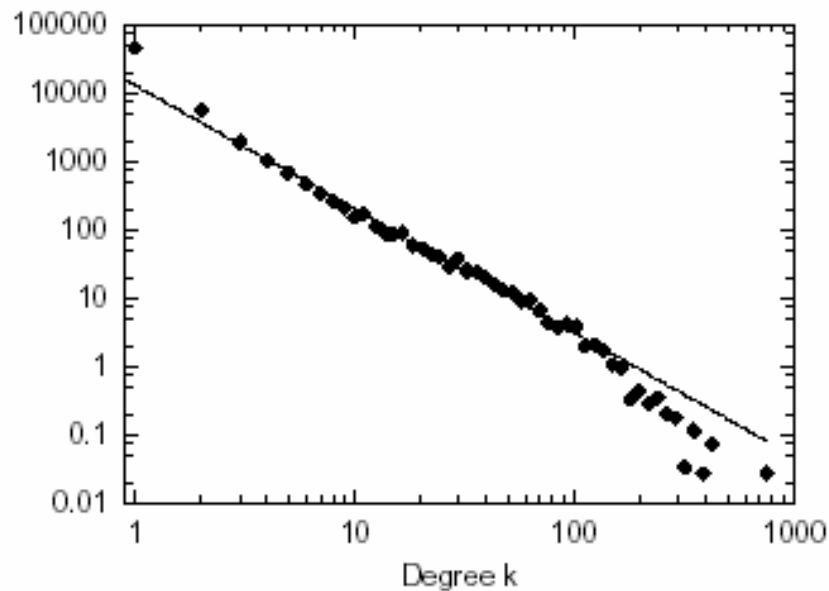
(b) Int-04-98

Online communities

Nodes: online user

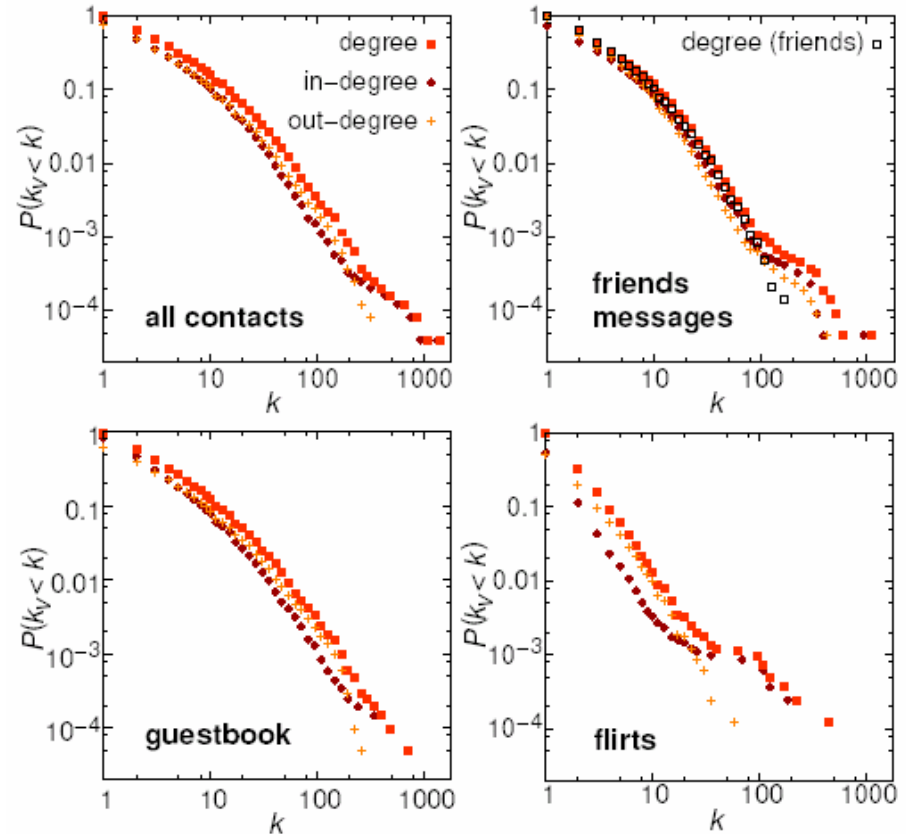
Links: email contact

Kiel University log files
112 days, $N=59,912$ nodes



Ebel, Mielsch, Bornholdtz, PRE 2002.

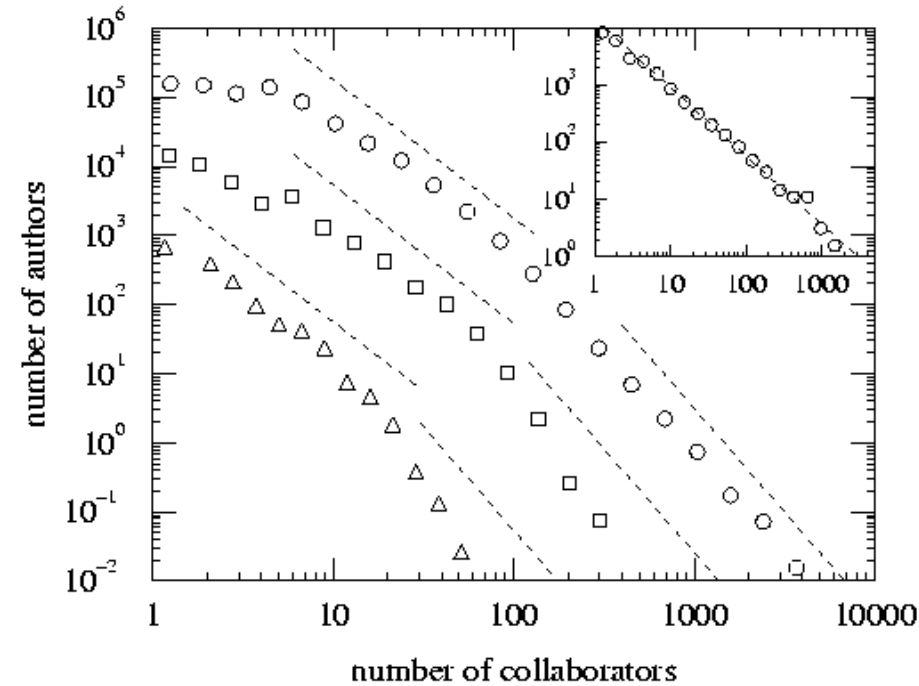
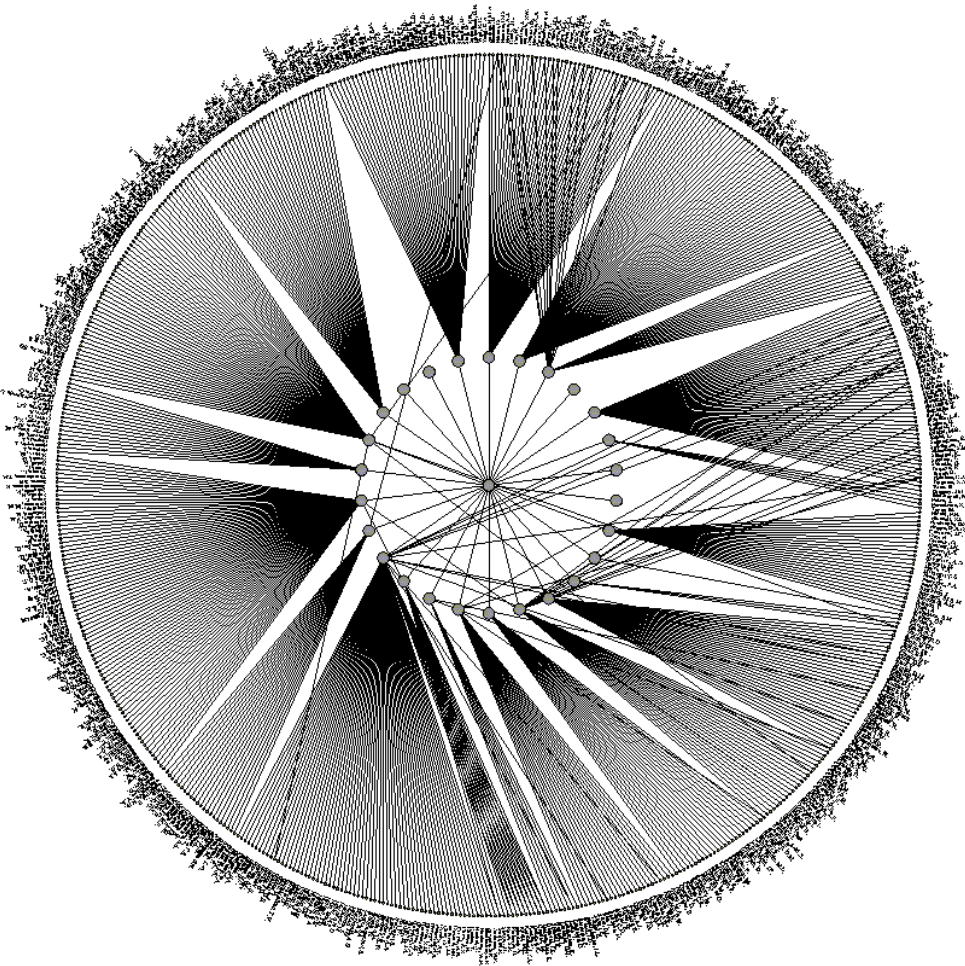
Pussokram.com online
community; 512 days,
25,000 users.



Holme, Edling, Liljeros, 2002.

SCIENCE COAUTHORSHIP

Nodes: scientist (authors) Links: write paper together



(Newman, 2000, A.-L. B. *et al* 2001)

SCIENCE CITATION INDEX

1,000 Most Cited Physicists, 1981-June 1997
Out of over 500,000 Examined
(see <http://www.sst.nrel.gov>)

Author name		Institute	Country	Field	avg. cites	total art.	total cites	rank by total cit.
Witten	E	Princeton (U)	USA, NJ	High-energy (T)	168	138	23235	1
Gossard	AC	UCSB (U)	USA, CA	Semicon				2
Cava	RJ	Bell Labs (I)	USA, NJ	Supercon				3
Ballogg	RJ	Bell Labs (I)	USA, NJ	Supercon				4
Ploog	K	Max Planck (NL)	Germany	Semicon				5
Ellis	J	Euro Nuclear Cent.	Switzerland	Astroph				6
Fisk	Z	Florida State (U)	USA, FL	Solid S				7
Cardona	M	Max Planck (NL)	Germany	Semicon				8
Nanopoulos	DV	Texas A&M (U)	USA, TX	High-e				9
Heeger	CA	Stanford (U)	USA, CA	Polym				10
Lee*								11
Suzuki*								12
Anderson	PW	Princeton (U)	USA, NJ	Solid S				13
Suzuki*	M							14
Freeman				Solid S				15
Tanaka								16
Muller			nd	Supercon				17
Schnee				Supercon				18
Chen				Optics (E)	60	162	9668	19
Morko				Semiconductors (E)	20	477	9668	19
Miller				Semiconductors (E)	67	144	9652	21
Chu				Superconductivity (E)	44	213	9453	22
Bednorz			nd	Superconductivity (E)	110	85	9311	23
Cohen				Solid State (T)	33	284	9311	23
Metg				Superconductivity (E)	86	108	9300	25
Waszc				Superconductivity (E)	57	112	9170	26
Shirane				Superconductivity (E)	23	169	8841	27
Wieg				Semiconductors (E)	85	104	8822	28
Vando				Magnetism (E)	67	129	8686	29
Uchida					28	301	8520	30
Hor	PH	Houston Univ. (U)	USA, TX	Superconductivity (E)	72	119	8512	31
Murphy	DW			Astronomy (E)	111	76	8439	32
Birgeneau	RJ	MIT (U)	USA, MA	Superconductivity (E)	41	286	8375	33
Jorgensen	JD	Argonne (NL)	USA, IL	Superconductivity (E)	167	107	8298	34
Hinks	DG	Argonne (NL)	USA, IL	Superconductivity (E)	57	225	8253	35

Nodes: papers

Links: citations

1736 PRL papers (1988)

H.E. Stanley,...

1 2 ... 25

1234... 1078...

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

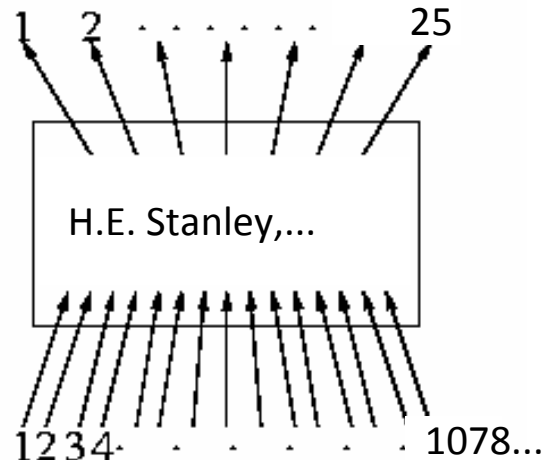
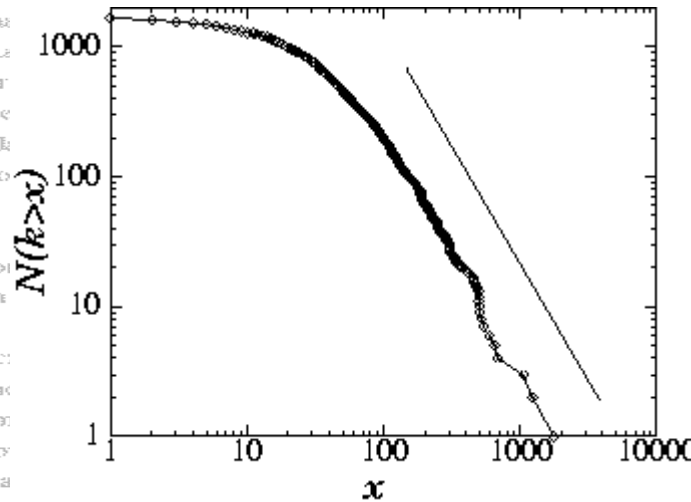
$(\gamma = 3)$

(S. Redner, 1998)

Nodes: papers

Links: citations

1736 PRL papers (1988)



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

$$(\gamma = 3)$$

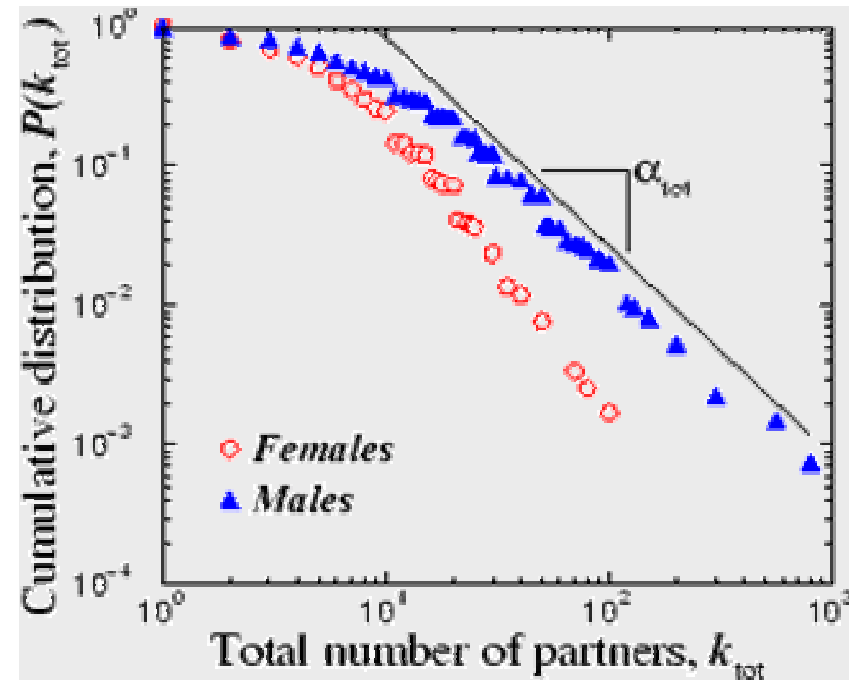
(S. Redner, 1998)

* citation total may be skewed because of multiple authors with the same name

Swedish sex-web

Nodes: people (Females; Males)

Links: sexual relationships



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

Liljeros et al. *Nature* 2001

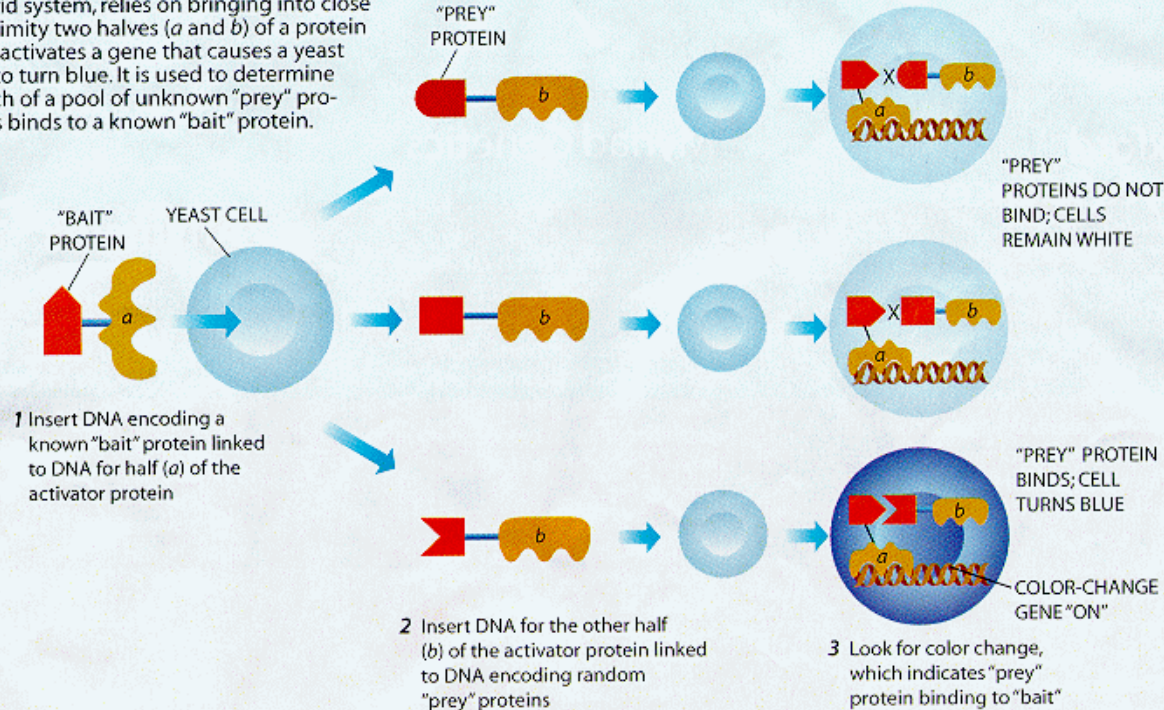
Yeast protein network

Nodes: proteins

Links: physical interactions (binding)

Finding Proteins That Interact

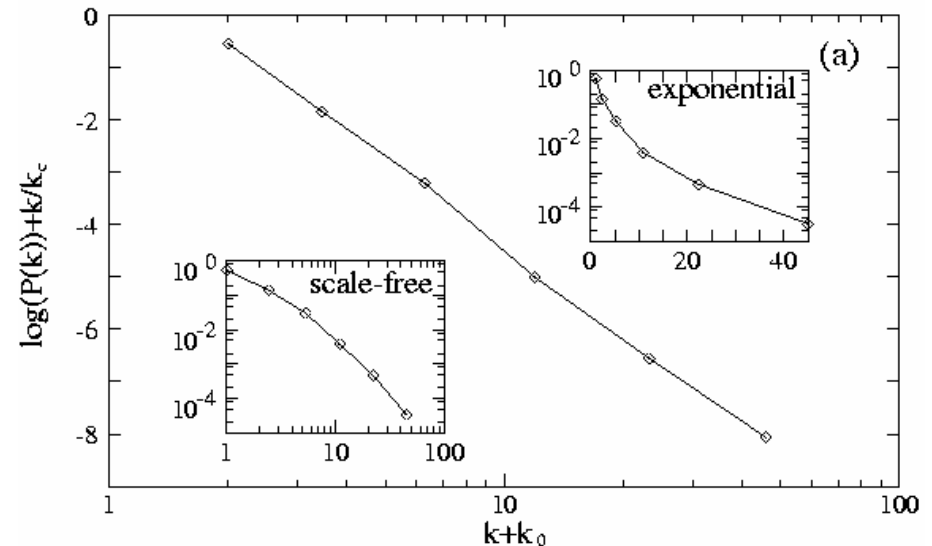
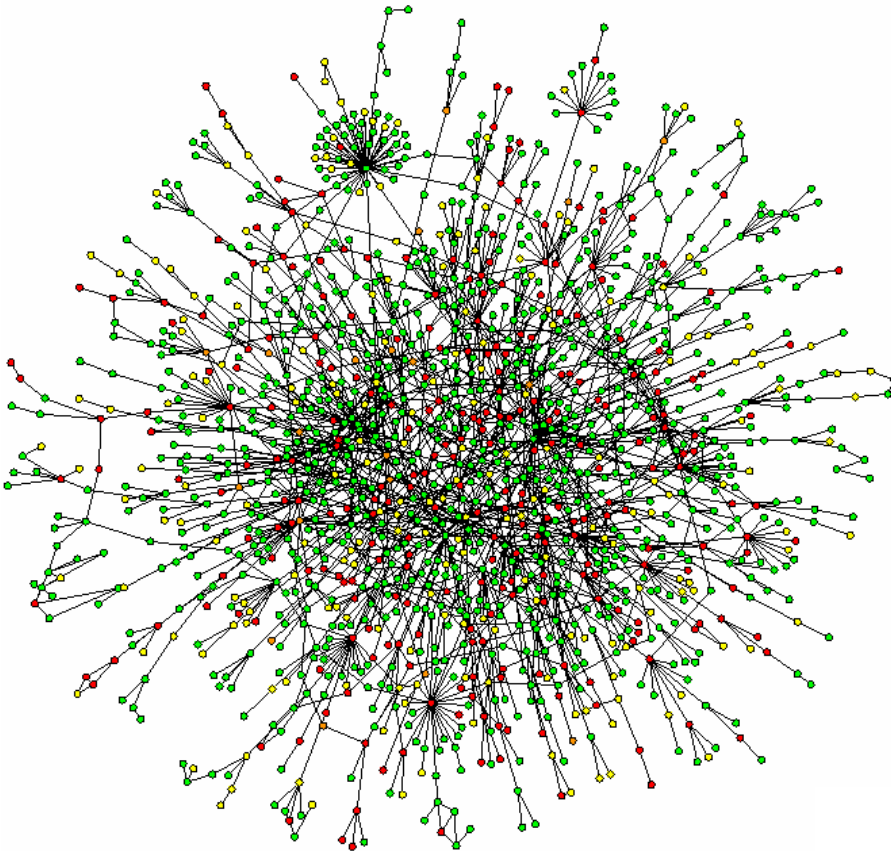
One technique, called the yeast two-hybrid system, relies on bringing into close proximity two halves (*a* and *b*) of a protein that activates a gene that causes a yeast cell to turn blue. It is used to determine which of a pool of unknown "prey" proteins binds to a known "bait" protein.



Protein interaction network

Nodes: proteins

Links: physical interactions (binding)

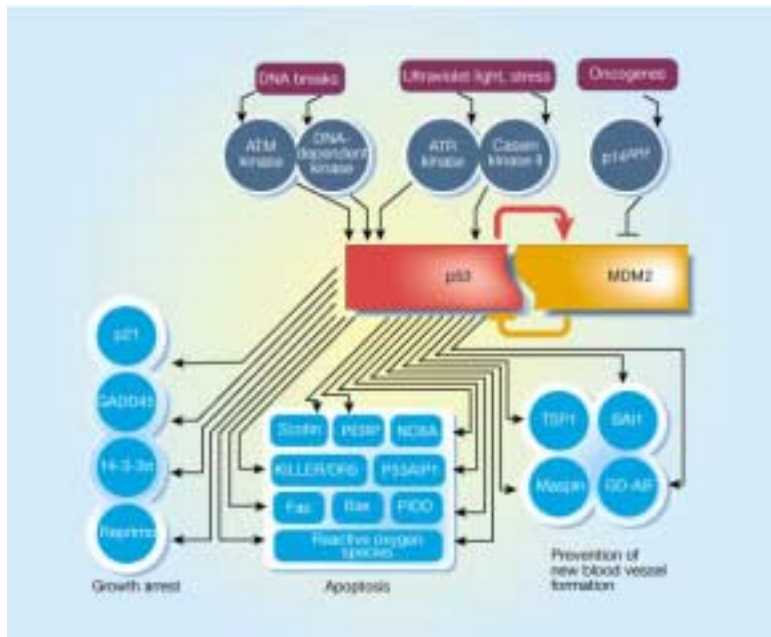


$$P(k) \sim (k + k_0)^{-\gamma} \exp\left(-\frac{k + k_0}{k_\tau}\right)$$

Surfing the p53 network

Bert Vogelstein, David Lane and Arnold J. Levine

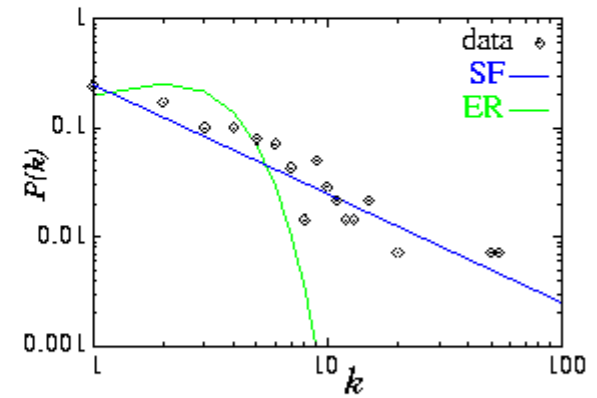
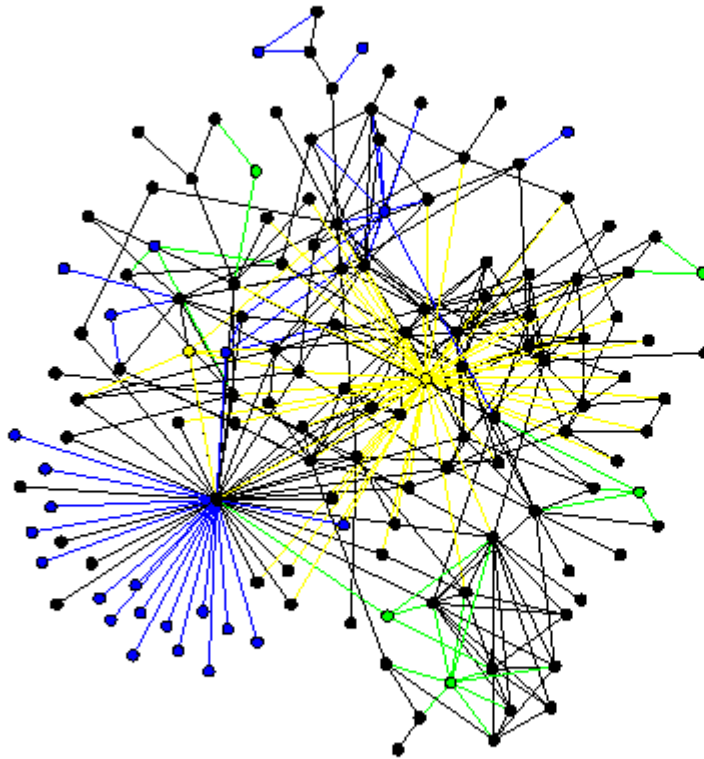
The p53 tumour-suppressor gene integrates numerous signals that control cell life and death. As when a highly connected node in the Internet breaks down, the disruption of p53 has severe consequences.



...

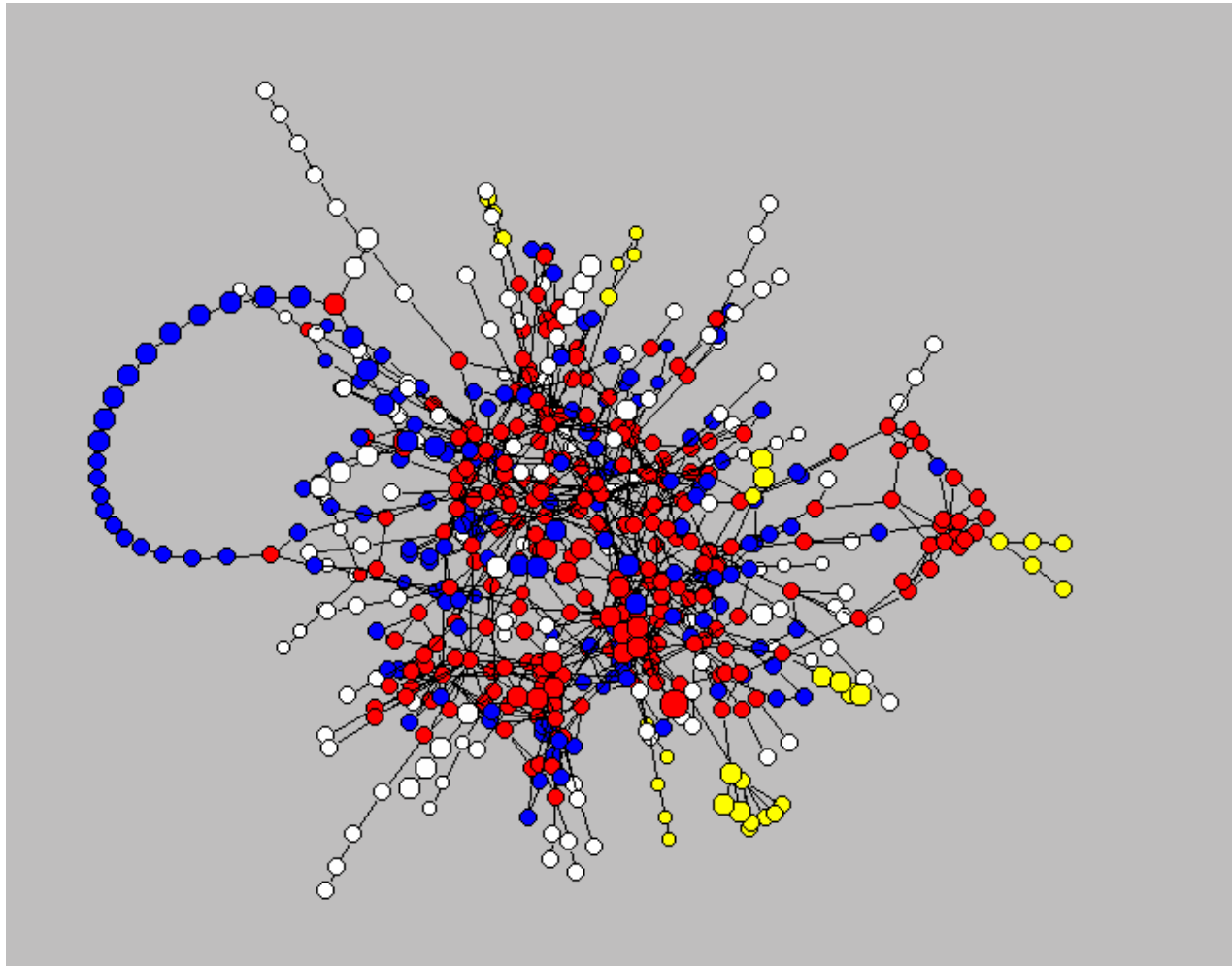
“One way to understand the p53 network is to compare it to the Internet. The cell, like the Internet, appears to be a ‘**scale-free network**’.”

p53 network (mammals)

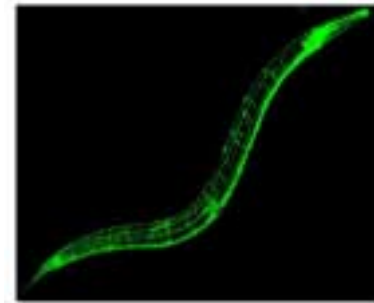
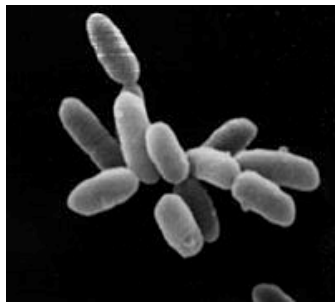
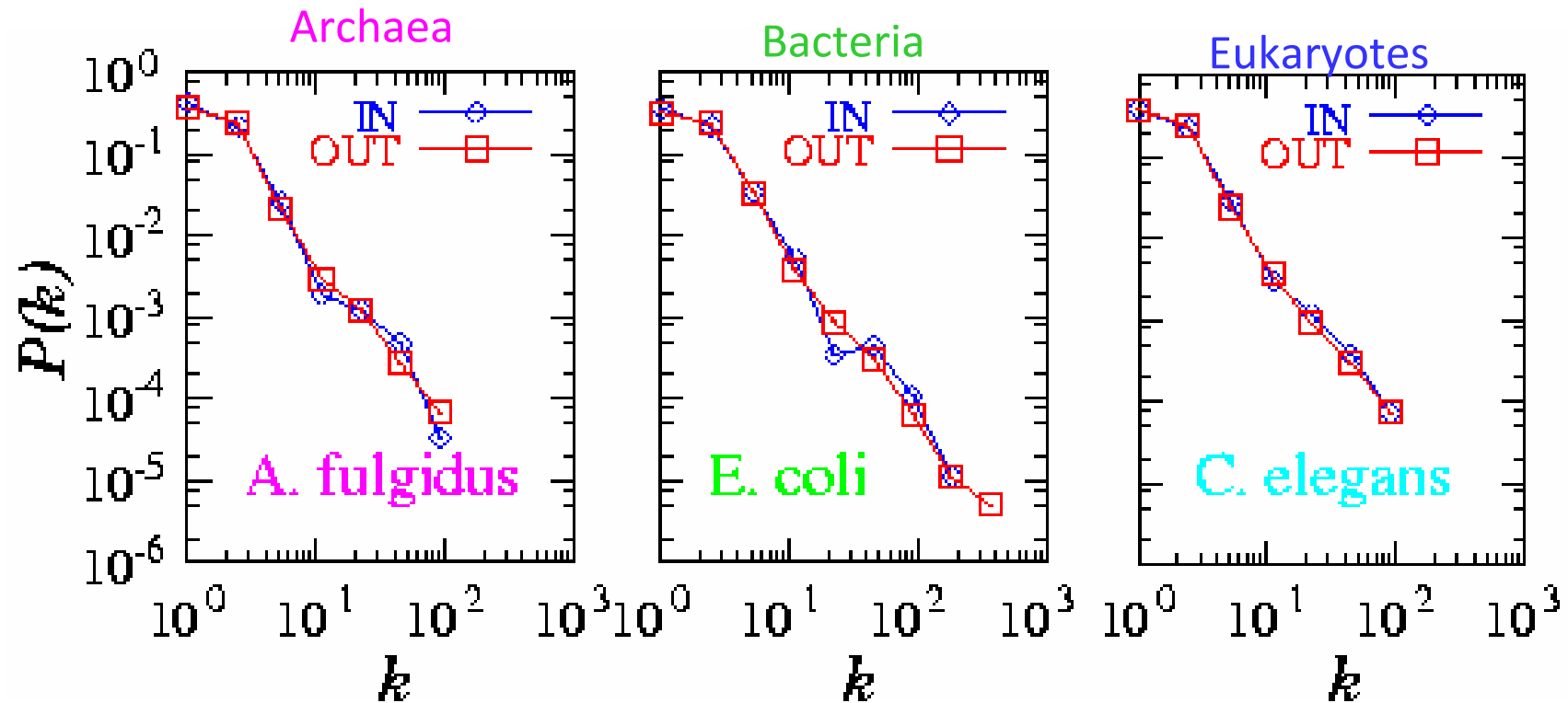


Metabolic Network

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



Metabolic network



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

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

Small World Features of Scale free Networks

Random Networks:

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

(small world effect)

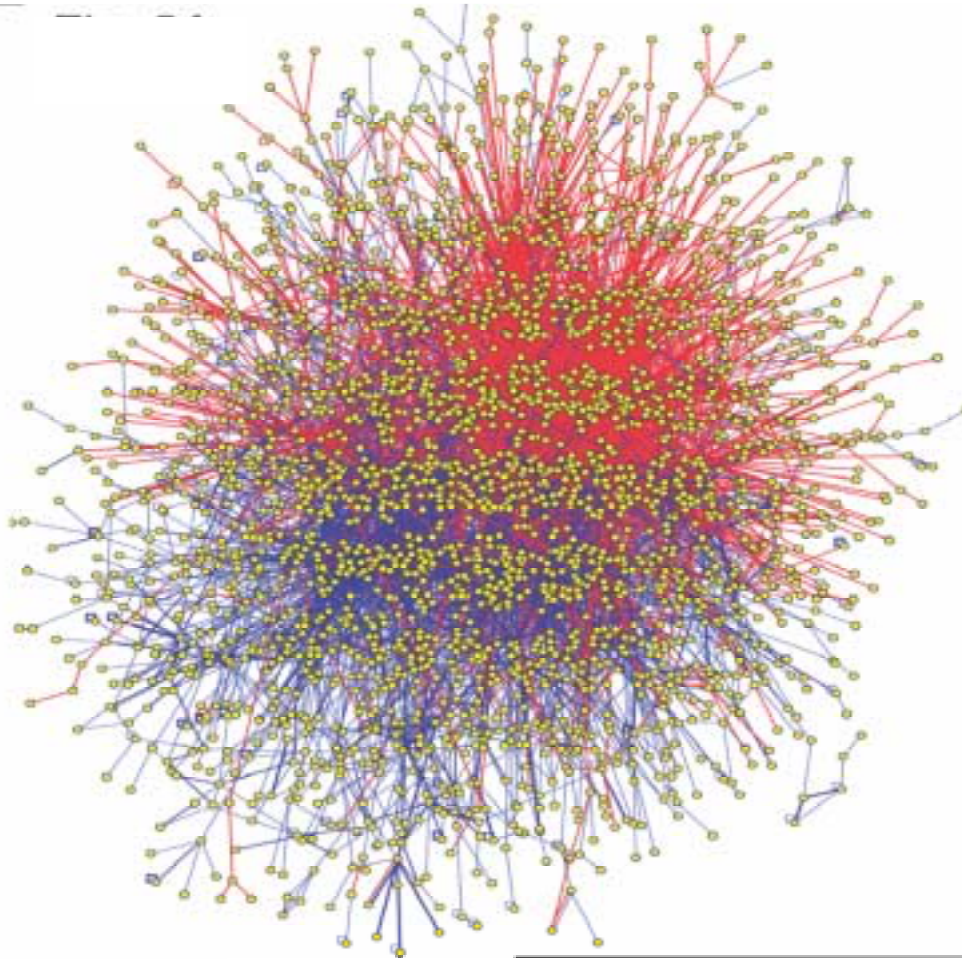
Scale-Free Networks: $P(k) \sim k^{-\gamma}$

$$D = \begin{cases} \log N & \gamma > 3 \\ \log \log N & 2 < \gamma < 3 \\ \text{const} & \gamma = 2 \end{cases}$$

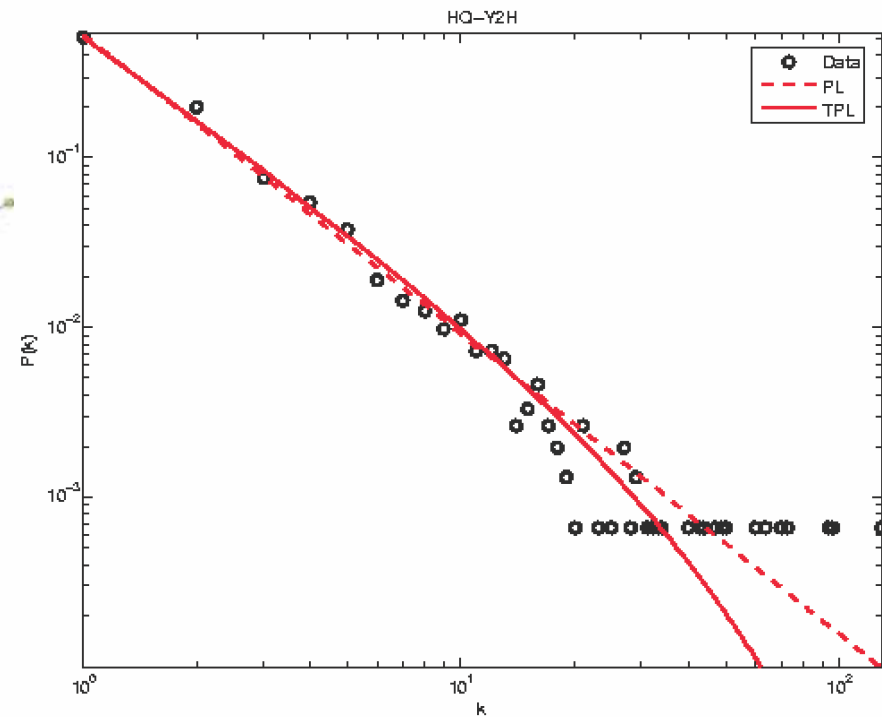
(ultra small world)

Cohen, Havlin, PRL '03

Human Interaction Network

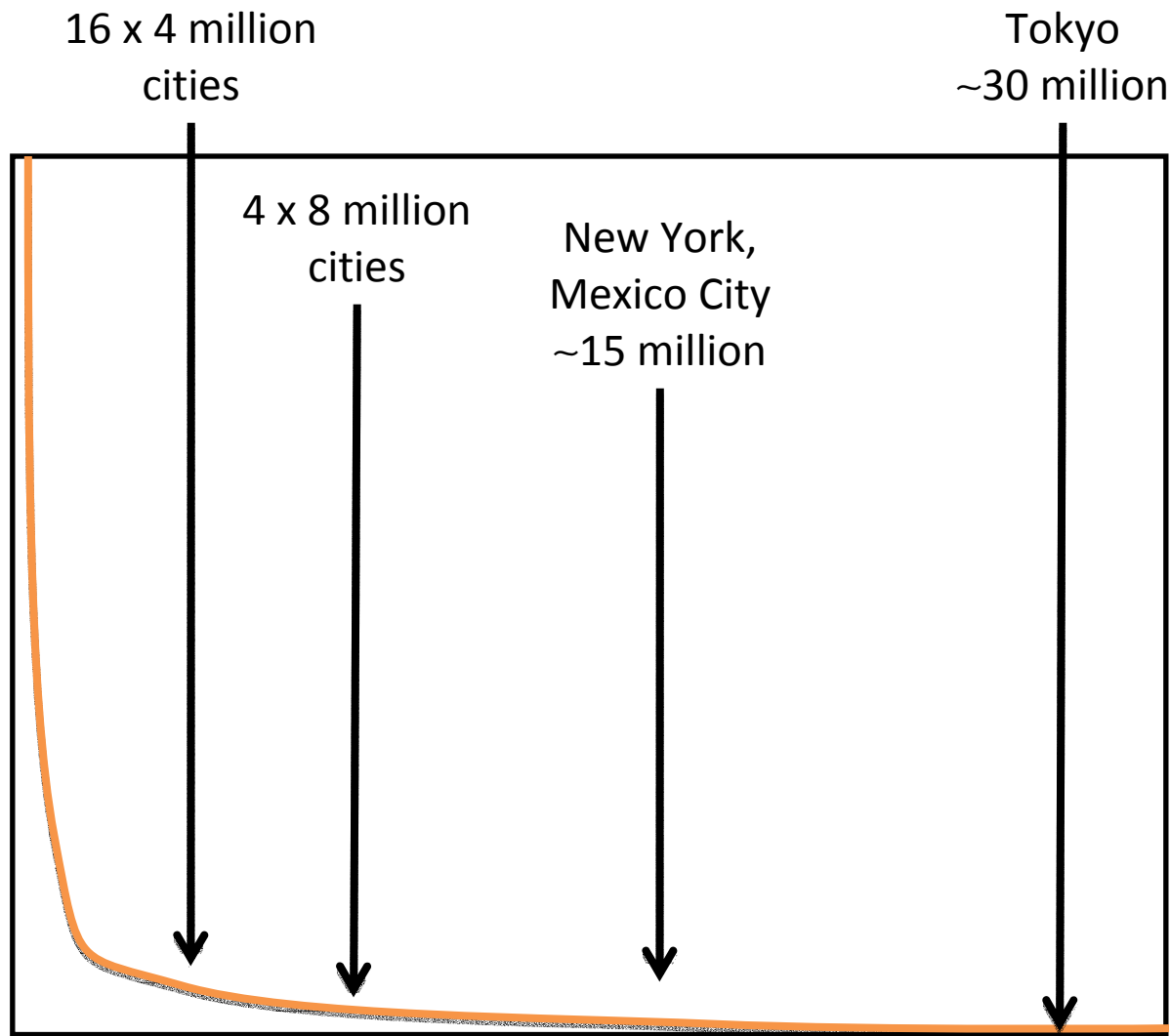


2,800 Y2H interactions
4,100 binary LC interactions
(HPRD, MINT, BIND, DIP, MIPS)



Rual *et al.* Nature 2005; Stelze *et al.* Cell 2005

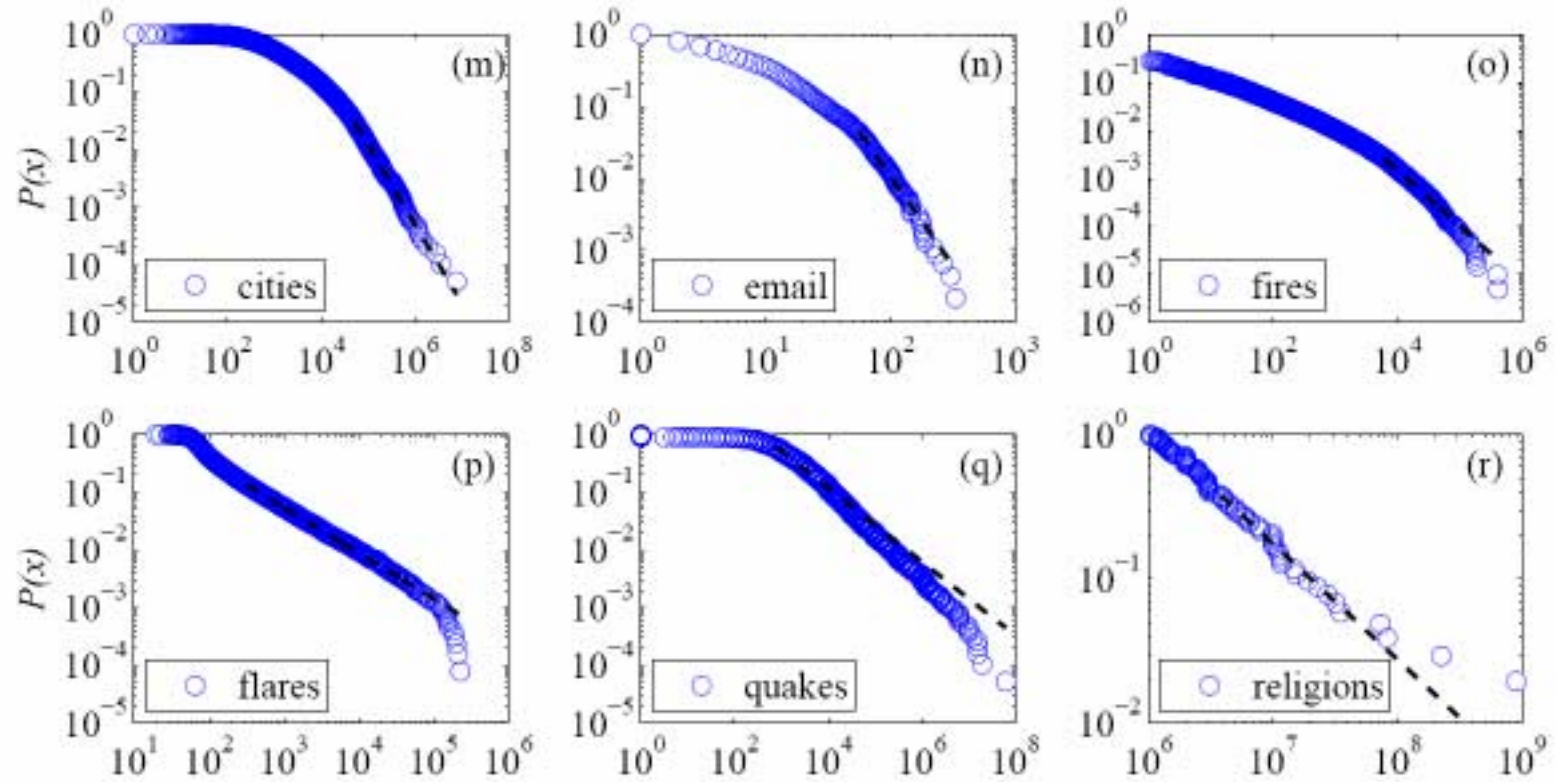
Number of Cities



Size of Cities

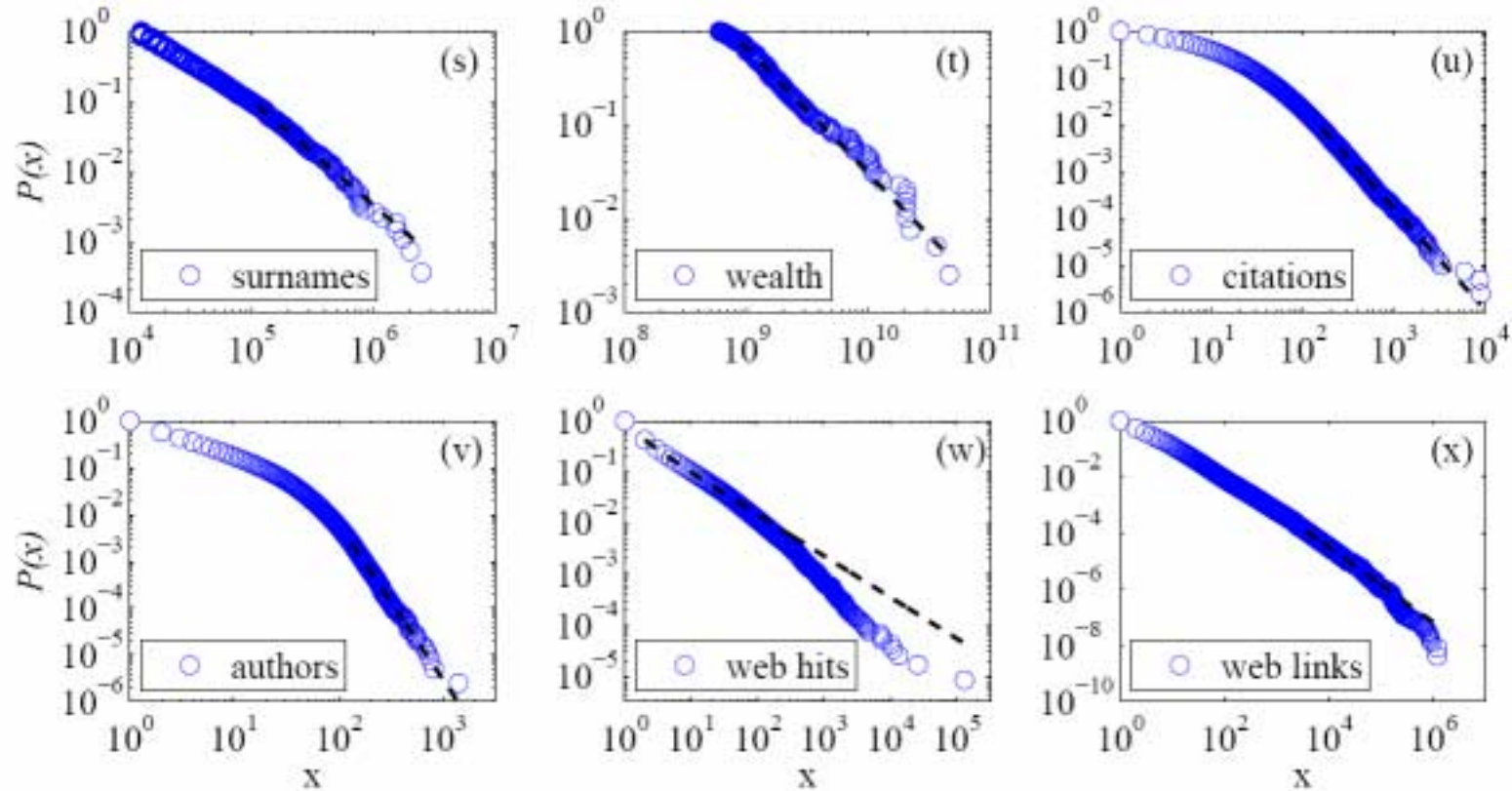
$$P \sim 1/x$$

Power laws everywhere



[Power-law distributions in empirical data](#), Aaron Clauset, Cosma Rohilla Shalizi, and M. E. J. Newman, submitted to *SIAM Review*.

Power laws everywhere



[Power-law distributions in empirical data](#), Aaron Clauset, Cosma Rohilla Shalizi, and M. E. J. Newman, submitted to *SIAM Review*.

ACTOR CONNECTIVITIES

Nodes: actors

Links: cast jointly



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

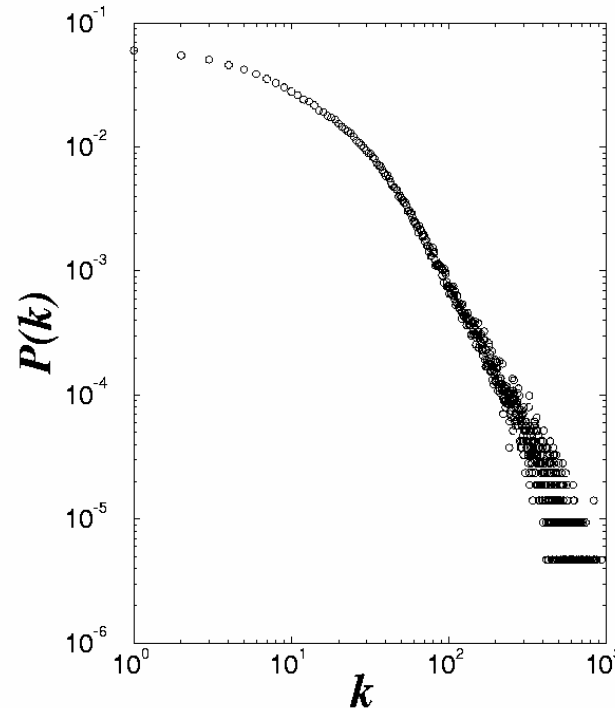


$N = 212,250$ actors

$\langle k \rangle = 28.78$

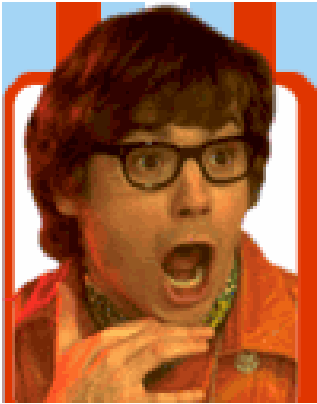
$P(k) \sim k^{-\gamma}$

$\gamma = 2.3$



Scale free networks

- In scale free networks most nodes have few connections, whereas a few ones have very many nodes.
- They are called the HUBS
- Despite their clustered structure, scale free networks are small worlds because the hubs provide the long-range connections
- Is Kevin Bacon an Hub ?

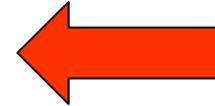


Austin Powers:
The spy who
shagged me

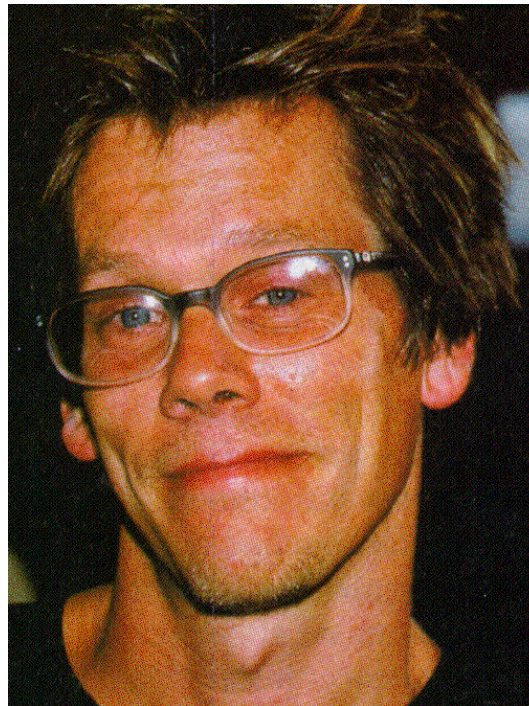


Robert Wagner

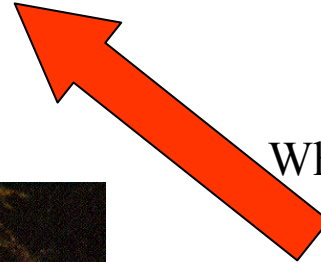
Let's make
it legal



Wild Things



What Price Glory



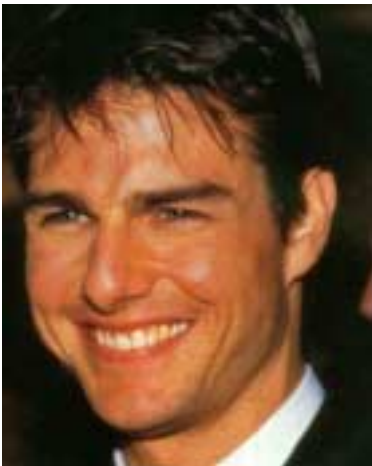
Barry Norton



Monsieur
Verdoux



A Few
Good Men



Nothing special about Kevin Bacon

Kevin Bacon

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

Average separation: 2.79

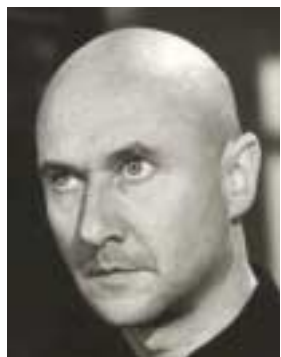
*Is Kevin Bacon
the most
connected actor?*

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
...				
876	Kevin Bacon	2.786981	46	1811
...				



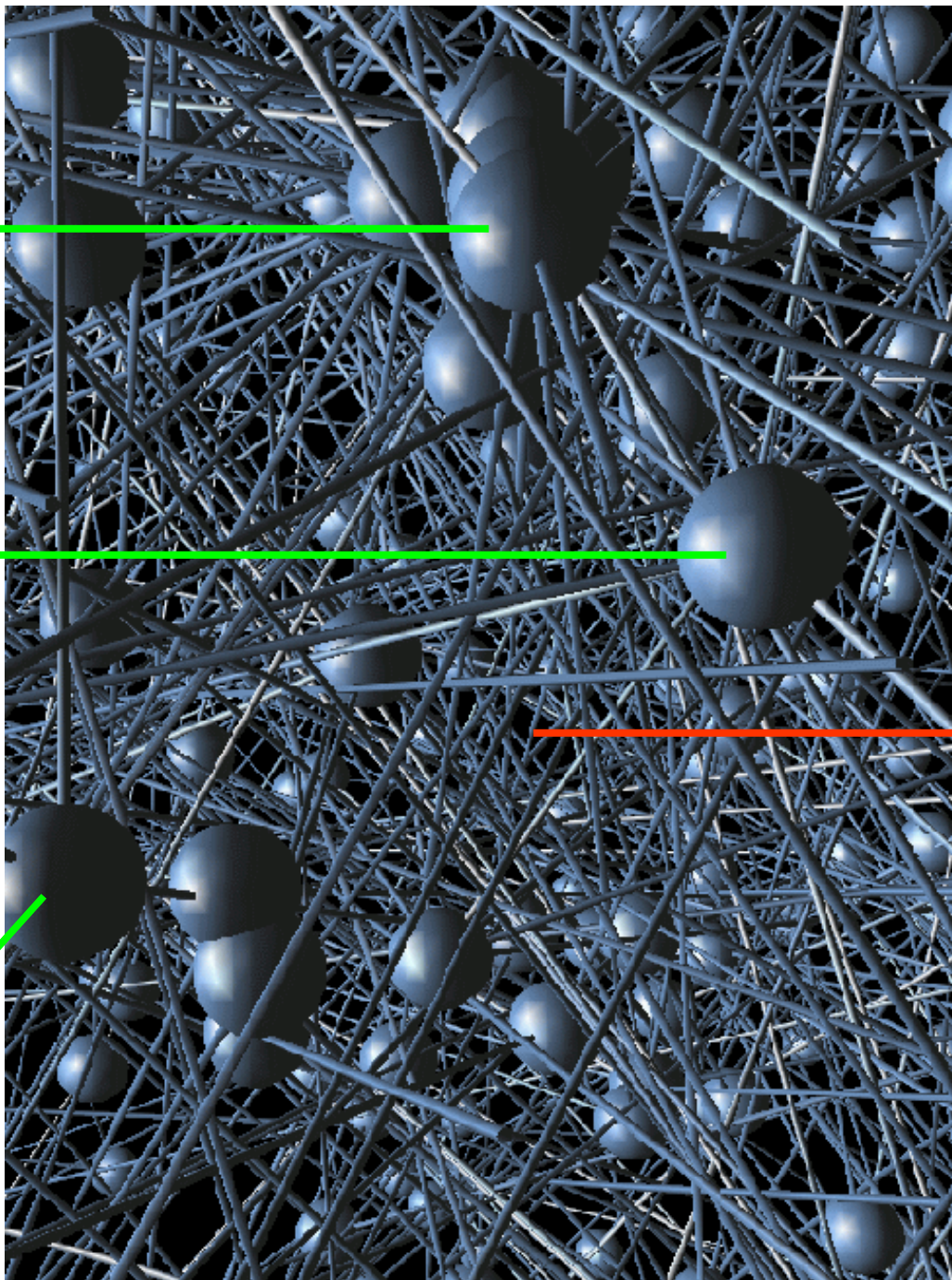
#1 Rod Steiger



#2 Donald
Pleasence



#3 Martin Sheen



#876
Kevin Bacon



Why are networks scale-free?

- **DYNAMICS**

Dynamics on networks

Dynamics of networks

- **Dynamics of networks**

Preferential attachment

Fitness model

Gene duplication

Growth versus preferential attachment

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

WWW : addition of new documents

Citation : publication of new papers

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

WWW : linking to well known sites

Citation : citing again highly cited papers

GROWTH:

add a new node with m links

PREFERENTIAL ATTACHMENT: the probability that a node connects to a node with k links is proportional to k .

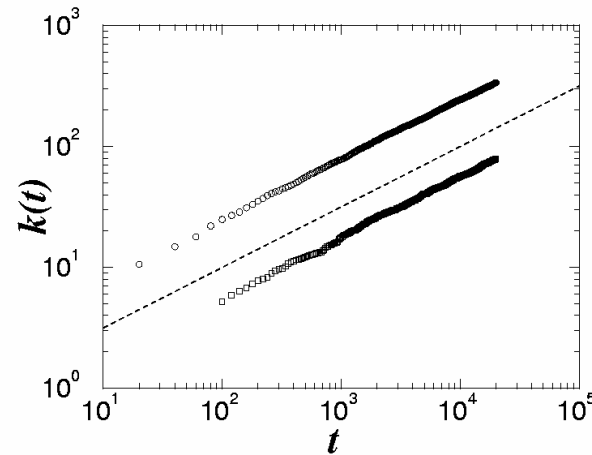
$$\Pi(k_i) = \frac{k_i}{\sum_j k_j}$$

Mean Field Theory (Preferential attachment)

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

, with initial condition $k_i(t_i) = m_0$

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



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

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

$$\gamma = 3$$

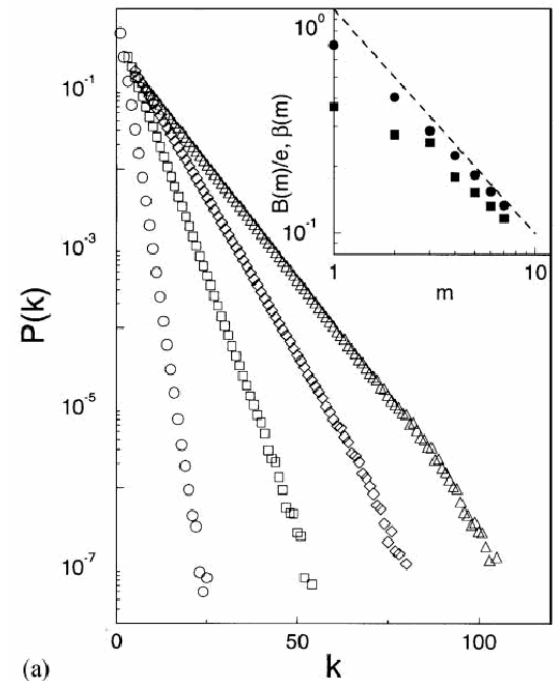
Growth model

$\Pi(k_i)$: uniform

$$\frac{\partial k_i}{\partial t} = A\Pi(k_i) = \frac{m}{m_0 + t - 1}$$

$$k_i(t) = m \left(\ln\left(\frac{m_0 + t - 1}{m + t_i - 1}\right) + 1 \right)$$

$$P(k) = \frac{e}{m} \exp\left(-\frac{k}{m}\right) \sim e^{-k}$$



Can Latecomers Make It? Fitness Model

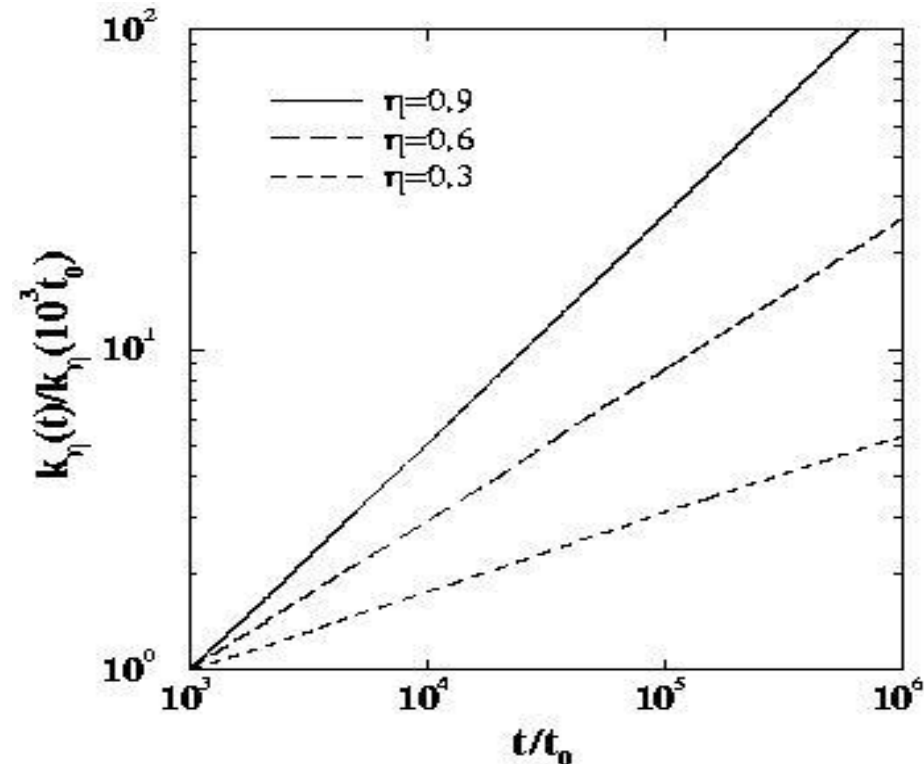
SF model: $k(t) \sim t^{1/2}$ (first mover advantage)

Real systems: nodes compete for links

Fitness Model: fitness (η)

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

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

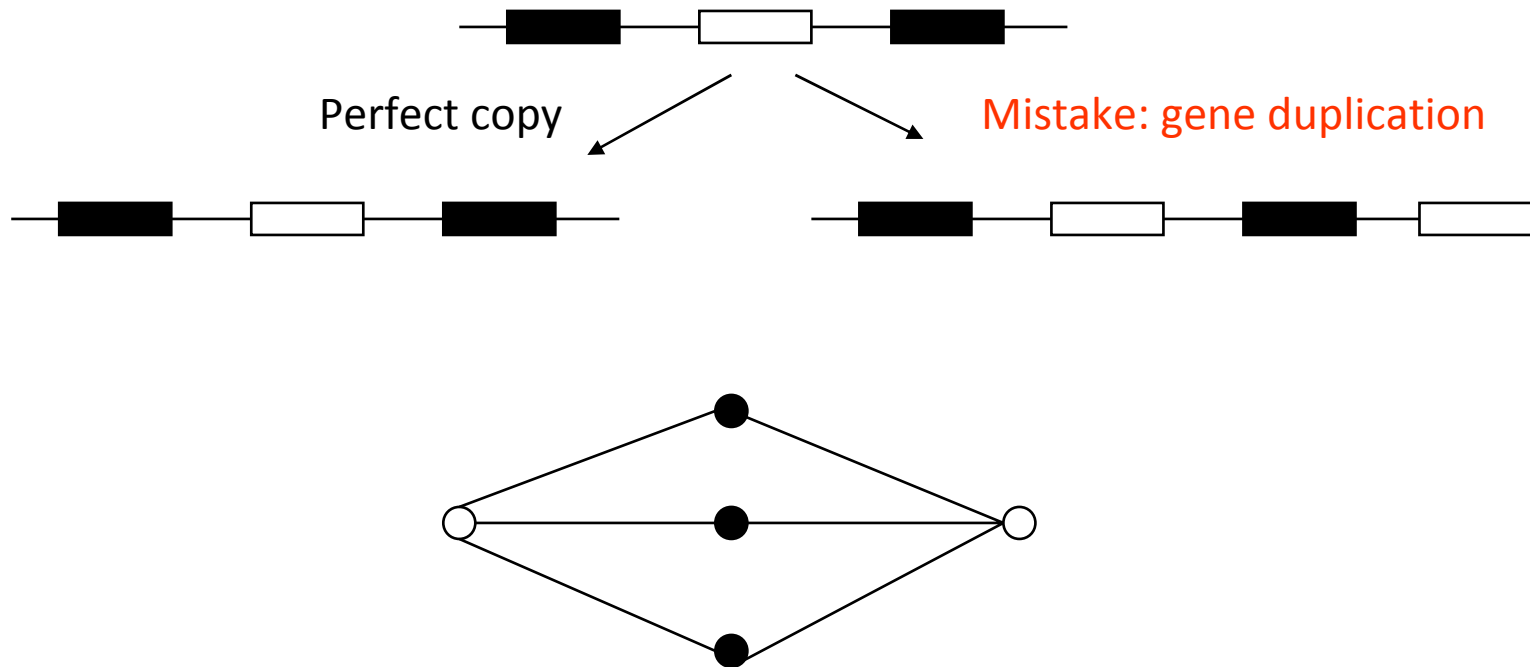


G. Bianconi and A.-L. Barabási, *Europhysics Letters*. **54**, 436 (2001).

Other Models

New concept or mechanism	Limits of γ	Reference
Linear growth, linear pref. attachment	$\gamma = 3$	Barabási and Albert, 1999
Nonlinear preferential attachment $\Pi(k_i) \sim k_i^\alpha$	no scaling for $\alpha \neq 1$	Krapivsky, Redner, and Leyvraz, 2000
Asymptotically linear pref. attachment $\Pi(k_i) \sim a_\infty k_i$ as $k_i \rightarrow \infty$	$\gamma \rightarrow 2$ if $a_\infty \rightarrow \infty$ $\gamma \rightarrow \infty$ if $a_\infty \rightarrow 0$	Krapivsky, Redner, and Leyvraz, 2000
Initial attractiveness $\Pi(k_i) \sim A + k_i$	$\gamma = 2$ if $A = 0$ $\gamma \rightarrow \infty$ if $A \rightarrow \infty$	Dorogovtsev, Mendes, and Samukhin, 2000a, 2000b
Accelerating growth $\langle k \rangle \sim t^\theta$ constant initial attractiveness	$\gamma = 1.5$ if $\theta \rightarrow 1$ $\gamma \rightarrow 2$ if $\theta \rightarrow 0$	Dorogovtsev and Mendes, 2001a
Accelerating growth $\langle k \rangle = at + 2b$	$\gamma = 1.5$ for $k \ll k_c(t)$ $\gamma = 3$ for $k \gg k_c(t)$	Barabási <i>et al.</i> , 2001 Dorogovtsev and Mendes, 2001c
Internal edges with probab. p	$\gamma = 2$ if $q = \frac{1-p+m}{1+2m}$	
Rewiring of edges with probab. q	$\gamma \rightarrow \infty$ if $p, q, m \rightarrow 0$	Albert and Barabási, 2000
c internal edges or removal of c edges	$\gamma \rightarrow 2$ if $c \rightarrow \infty$ $\gamma \rightarrow \infty$ if $c \rightarrow -1$	Dorogovtsev and Mendes, 2000c
Gradual ageing $\Pi(k_i) \sim k_i(t-t_i)^{-\nu}$	$\gamma \rightarrow 2$ if $\nu \rightarrow -\infty$ $\gamma \rightarrow \infty$ if $\nu \rightarrow 1$	Dorogovtsev and Mendes, 2000b
Multiplicative node fitness $\Pi_i \sim \eta_i k_i$	$P(k) \sim \frac{k^{-1-C}}{\ln(k)}$	Bianconi and Barabási, 2001a
Additive-multiplicative fitness $\Pi_i \sim \eta_i(k_i - 1) + \xi_i$	$P(k) \sim \frac{k^{-1-m}}{\ln(k)}$ $1 \leq m \leq 2$	Ergün and Rodgers, 2001
Edge inheritance	$P(k_{in}) = \frac{d}{k_{in}^2} \ln(ak_{in})$	Dorogovtsev, Mendes, and Samukhin, 2000c
Copying with probab. p	$\gamma = (2-p)/(1-p)$	Kumar <i>et al.</i> , 2000a, 2000b
Redirection with probab. r	$\gamma = 1 + 1/r$	Krapivsky and Redner, 2001
Walking with probab. p	$\gamma \rightarrow 2$ for $p > p_c$	Vázquez, 2000
Attaching to edges	$\gamma = 3$	Dorogovtsev, Mendes, and Samukhin, 2001a
p directed internal edges $\Pi(k_i, k_j) \propto (k_i^{in} + \lambda)(k_j^{out} + \mu)$	$\gamma_{in} = 2 + p\lambda$ $\gamma_{out} = 1 + (1-p)^{-1} + \mu p/(1-p)$	Krapivsky, Rodgers, and Redner, 2001
$1-p$ directed internal edges Shifted linear pref. activity	$\gamma_{in} = 2 + p$ $\gamma_{out} = 2 + 3p$	Tadić, 2001a

Another origin of the scale-free topology: Gene Duplication



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

$$\Pi(k) \sim k$$

(preferential attachment).

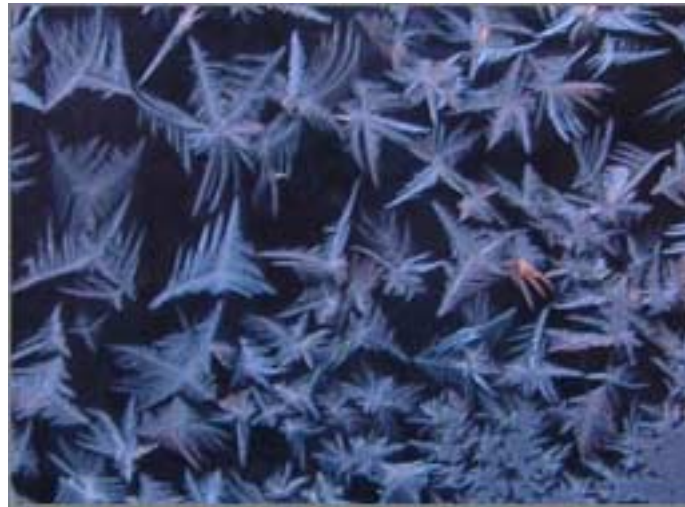
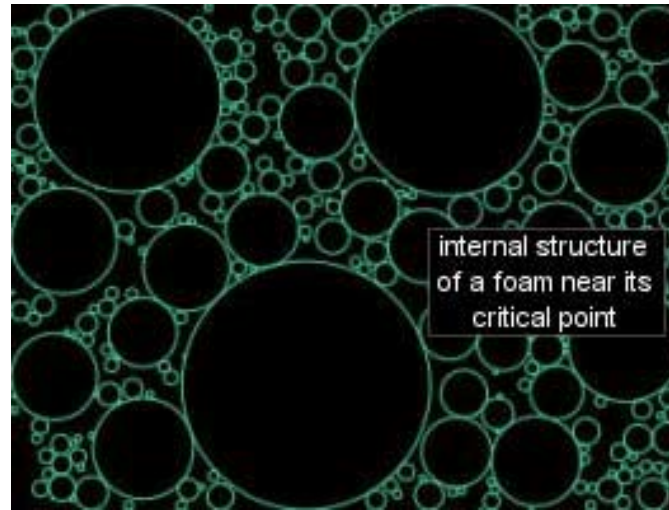
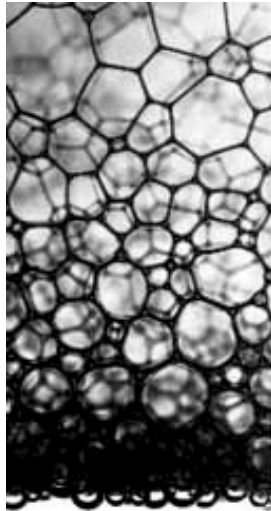
Wagner (2001); Vazquez *et al.* 2003; Sole *et al.* 2001; Rzhetsky & Gomez (2001); Qian *et al.* (2001); Bhan *et al.* (2002).

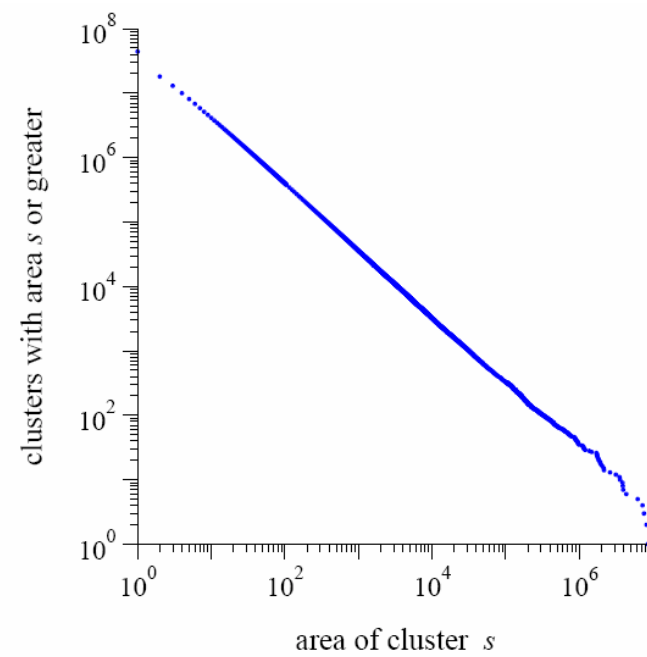
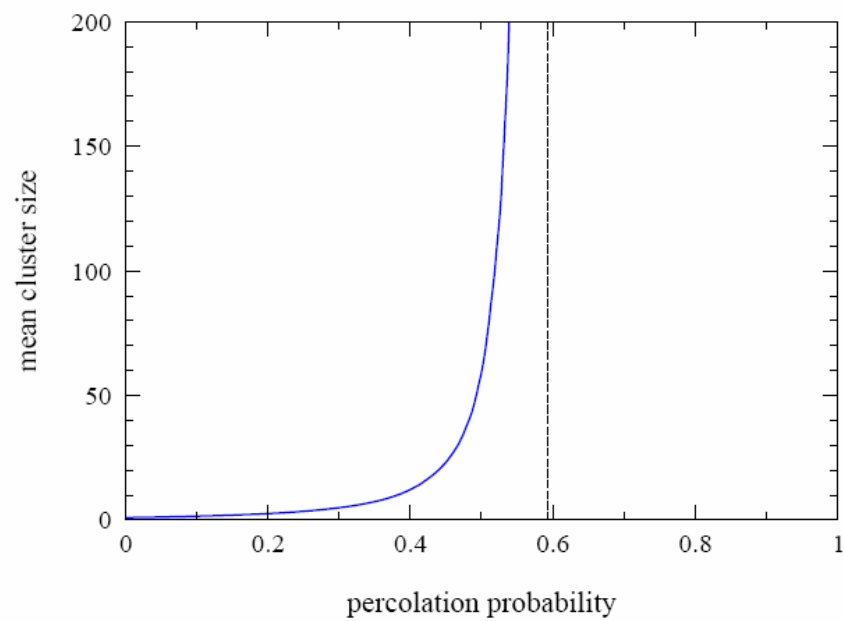
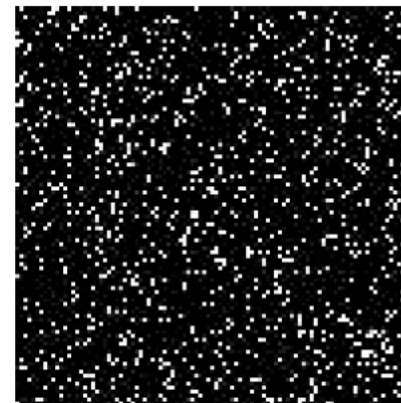
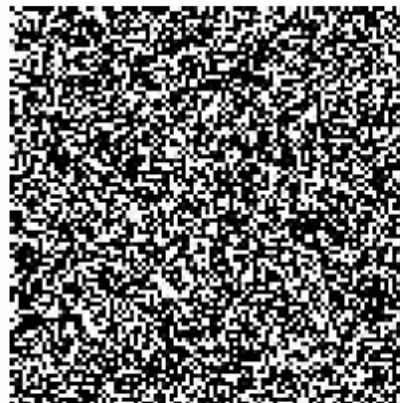
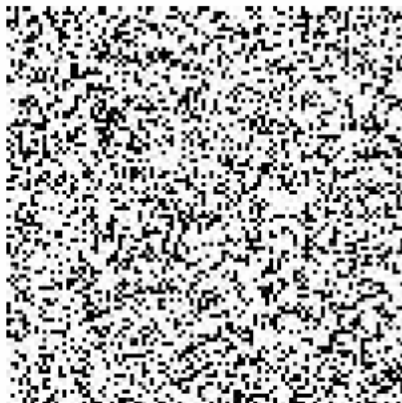
Power laws elsewhere

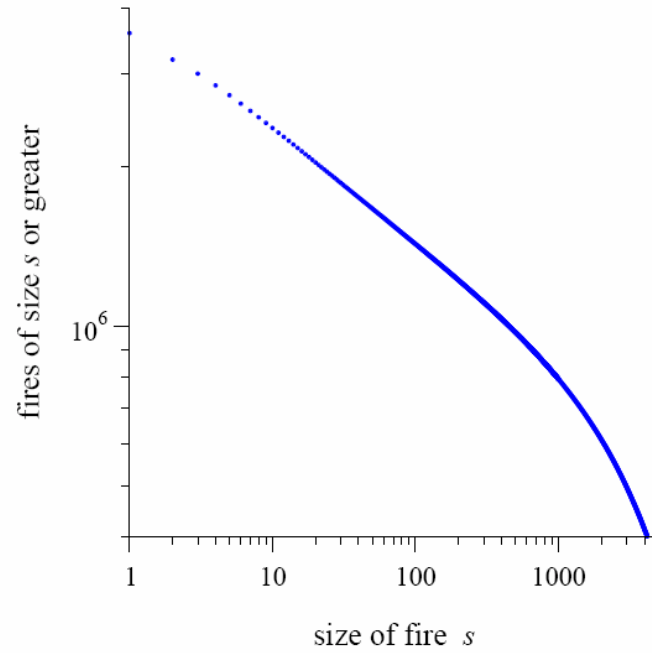
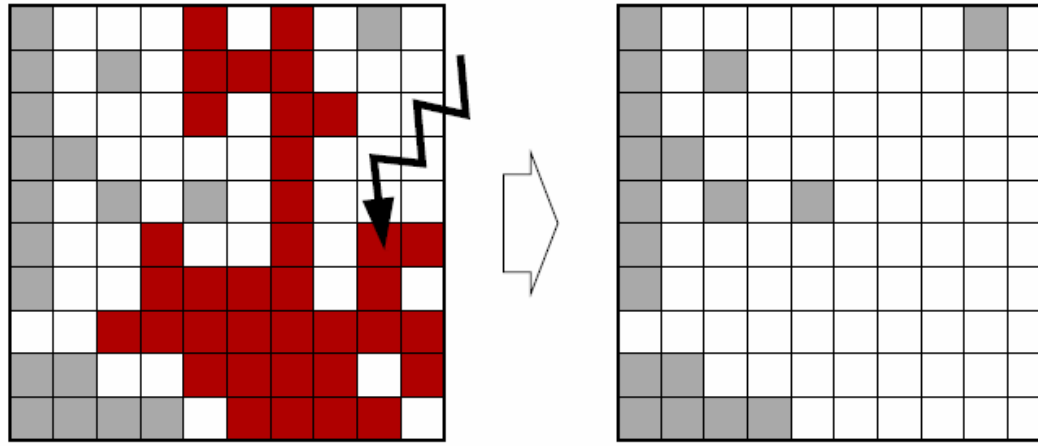
Is there a connection?

- Phase transitions
- Self-organized criticality
- Absence of a characteristic scale

Phase transitions







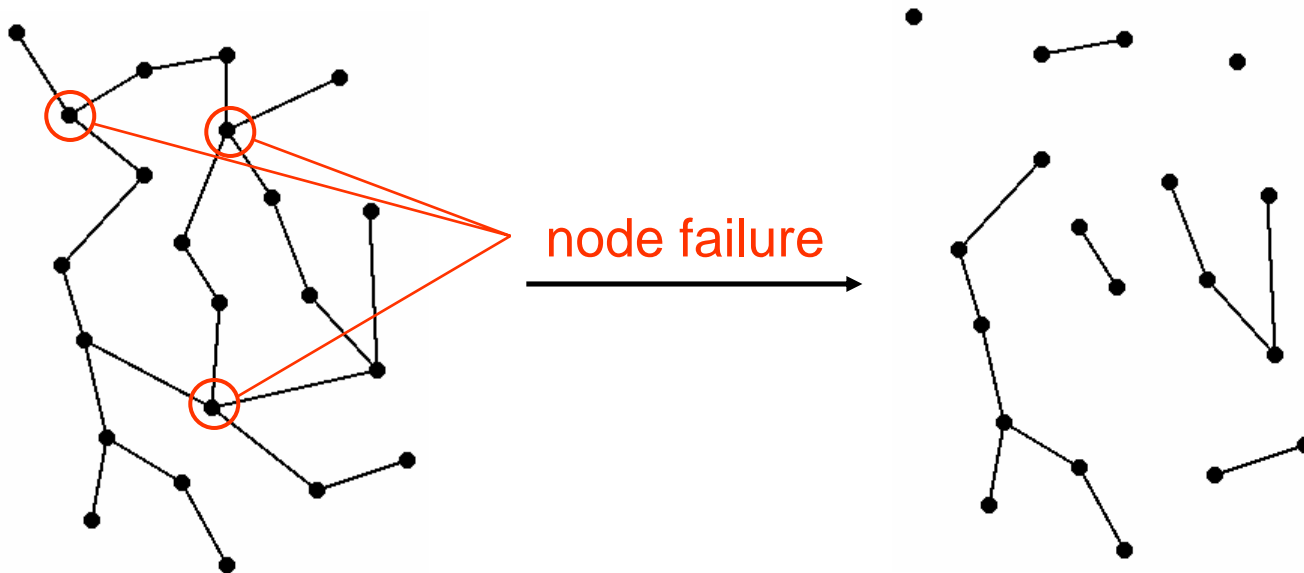
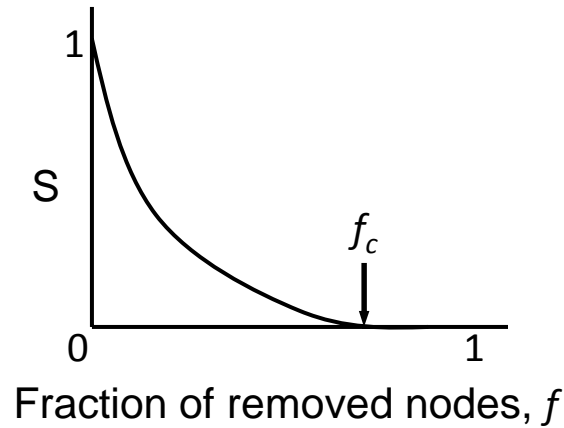
Self-Organized Criticality

Error and attack tolerance

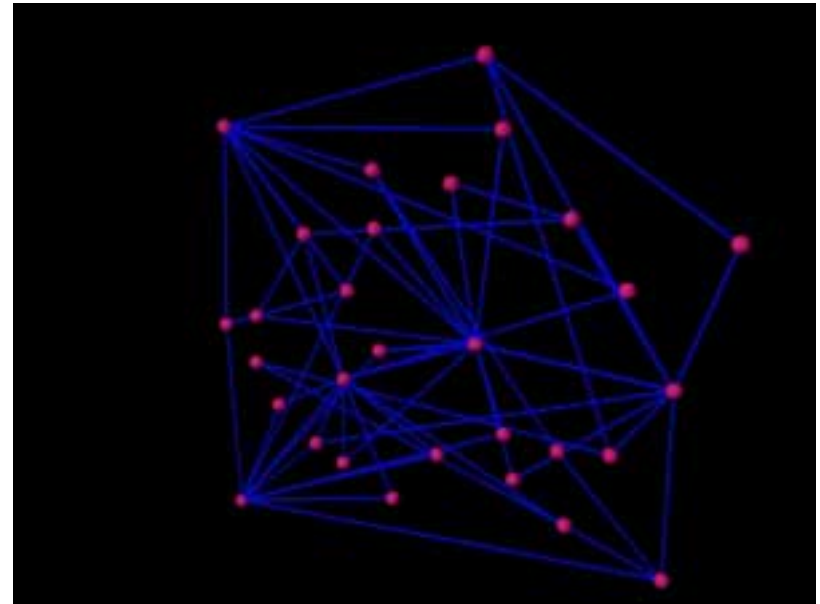
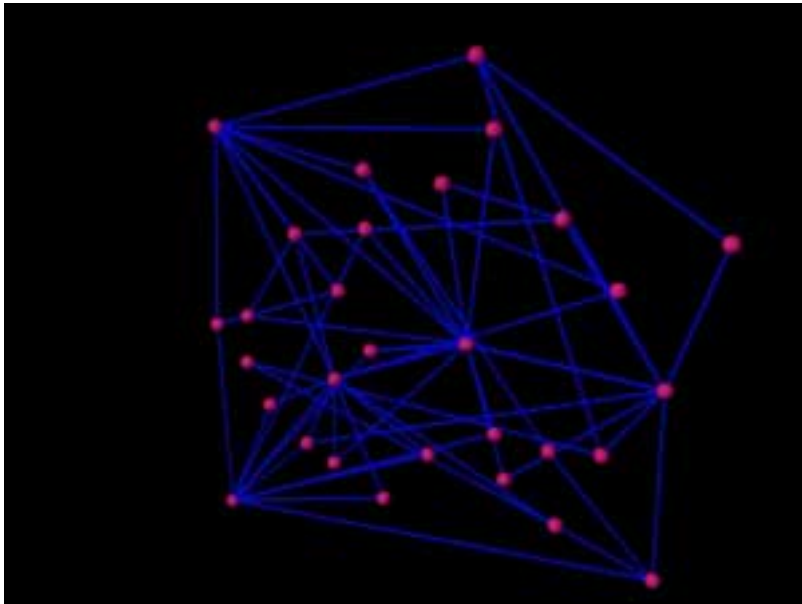


Robustness

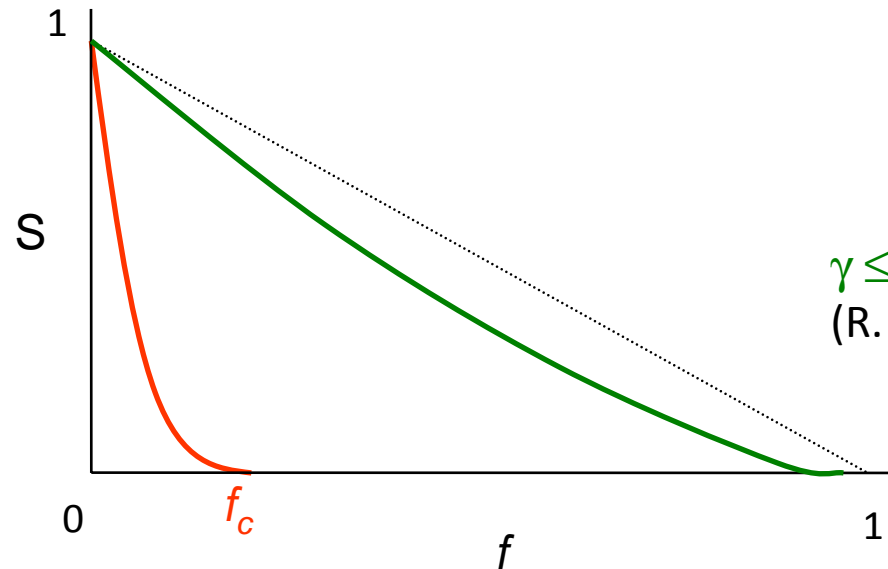
Complex systems maintain their basic functions even under errors and failures
(cell \rightarrow mutations; Internet \rightarrow router breakdowns)



Robustness of scale-free networks



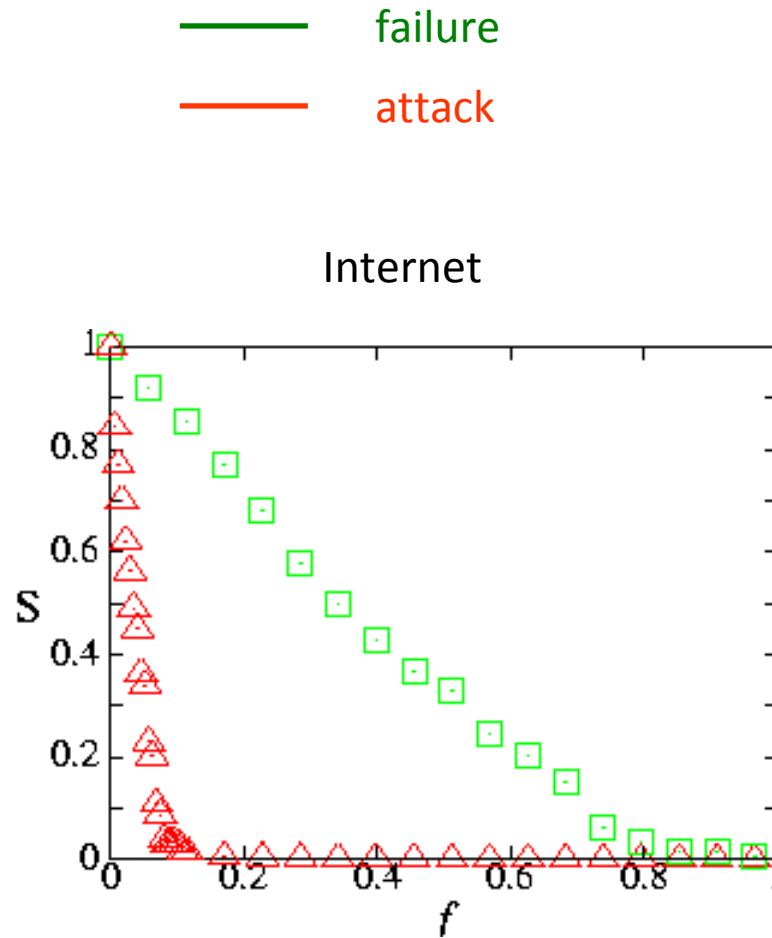
Attacks



Failures

$\gamma \leq 3 : f_c = 1$
(R. Cohen et al PRL, 2000)

Achilles' Heel of complex networks



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

Scale free networks

- Natural networks are not random graphs
- Many networks in nature are Scale-Free (SF), meaning that just a few nodes have a disproportionately large number of connections.
- Power-law distributions are ubiquitous in nature.
- While power-laws are associated with critical points in nature, systems can self-organize to this critical state.
- Important dynamical implications of the Scale-Free topology:
 - SF Networks are robust to failures, yet vulnerable to targeted attacks
 - The SF structure has important implications for the dynamics of the agents on the networks (epidemics, etc.)