

Introduction to Network Theory

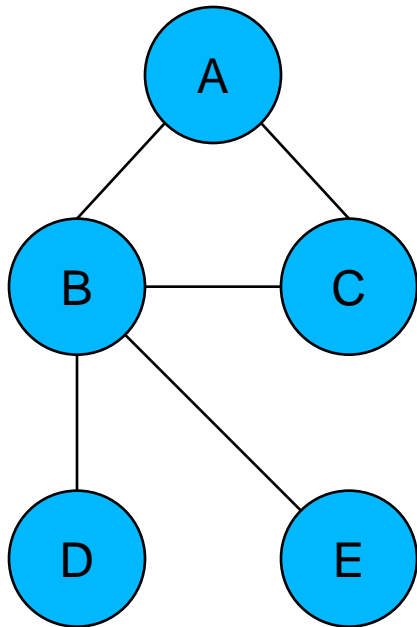
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Local metrics

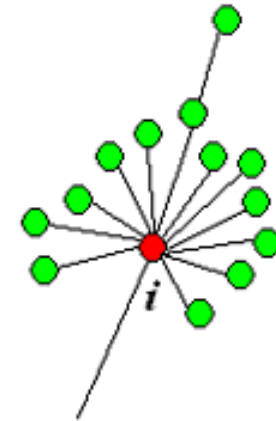
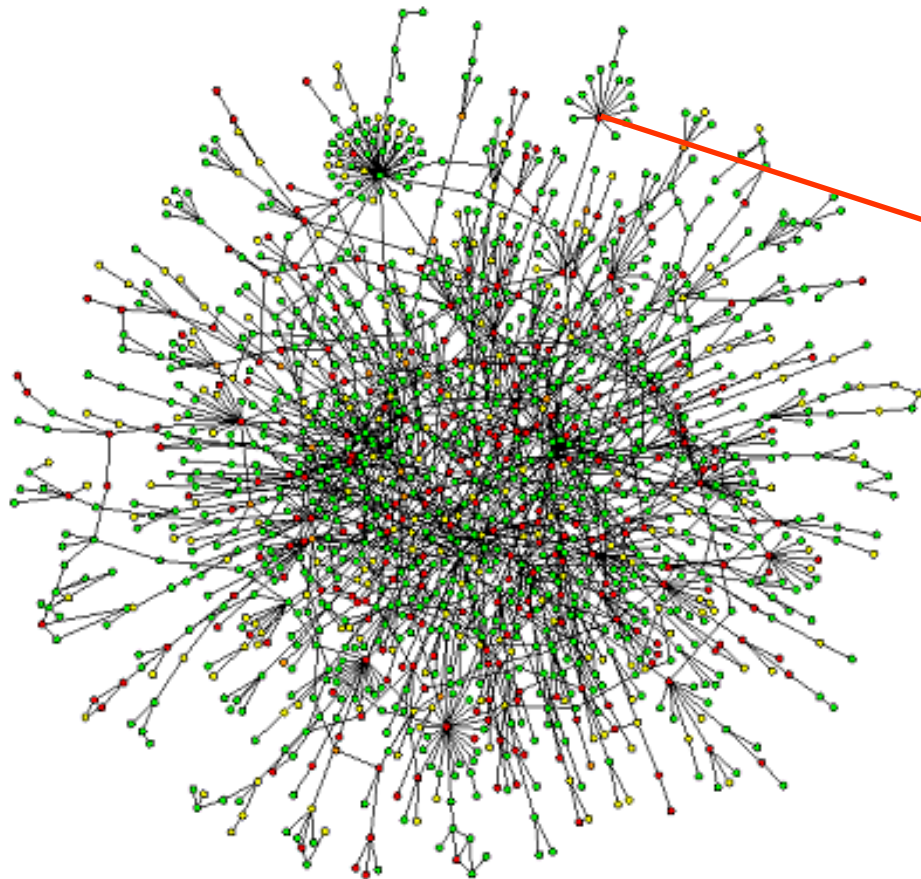
- Local metrics provide a measurement of a structural property of a single node
- Designed to characterise
 - Functional role – what part does this node play in system dynamics?
 - Structural importance – how important is this node to the structural characteristics of the system?

Degree Centrality



	<i>A</i>	<i>B</i>	<i>C</i>	<i>D</i>	<i>E</i>	degree
<i>A</i>	0	1	1	0	0	2
<i>B</i>	1	0	1	1	1	4
<i>C</i>	1	1	0	0	0	2
<i>D</i>	0	1	0	0	0	1
<i>E</i>	0	1	0	0	0	1

DEGREE DISTRIBUTION

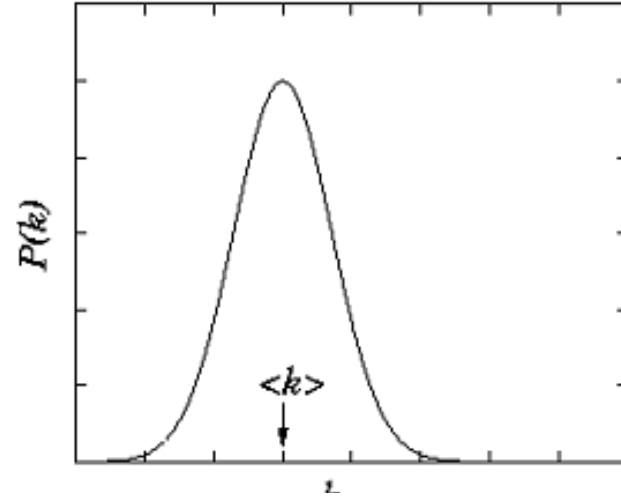


$$k_i = 13$$

protein-protein interaction network (PIN)

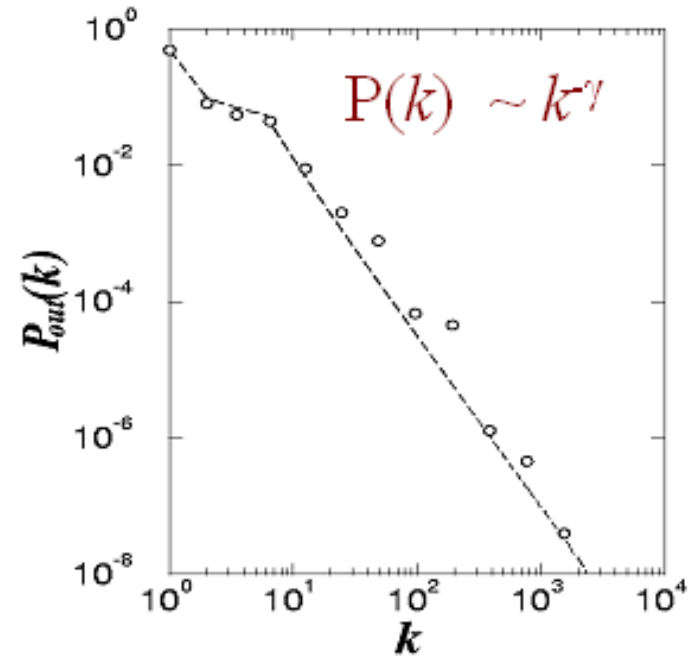
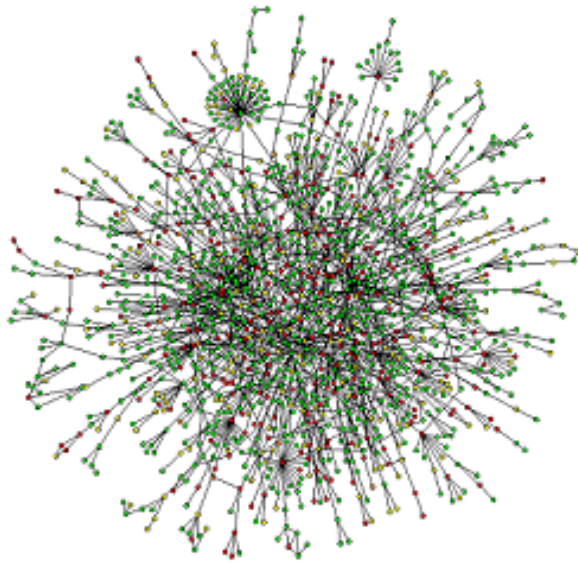
DEGREE DISTRIBUTION

Random Network



Expected

Real-World Network



Found

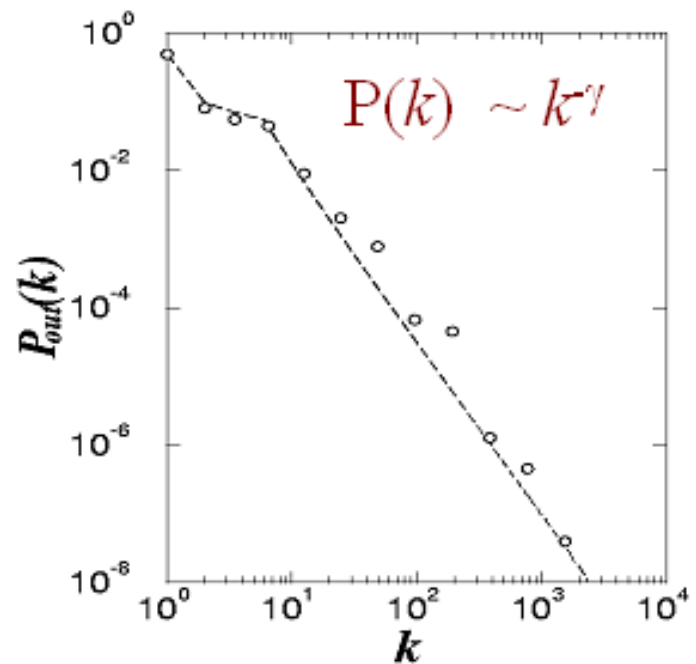
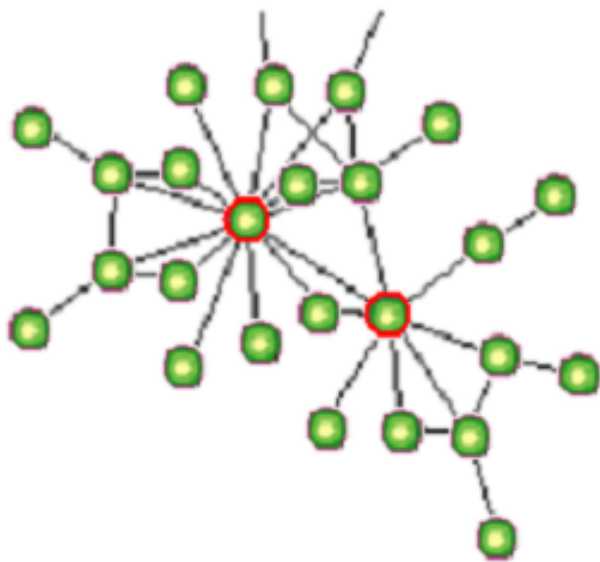
Scale-Freeness

Power-law: $f(x) = ax^\gamma$ $\log f(x) = \gamma \log x + \log a$

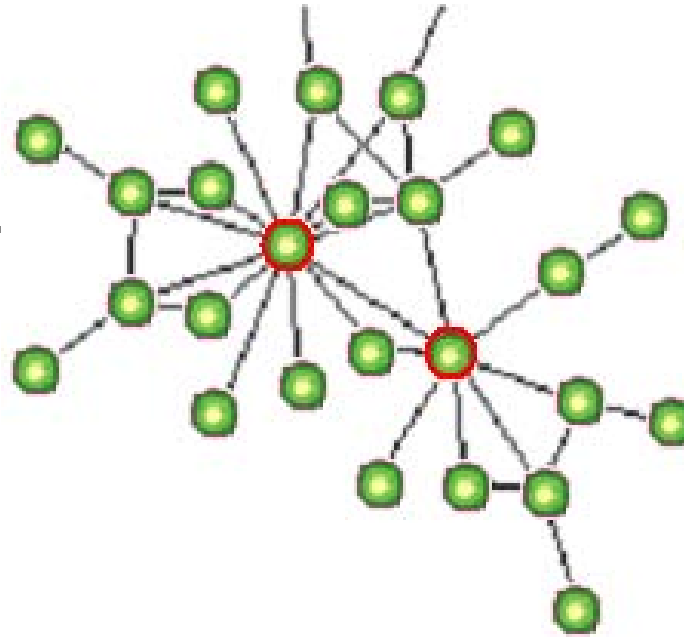
Scale invariance: $\log f(cx) = \gamma \log x + \log ca$

$f(cx) \propto f(x)$

Scale-free Network

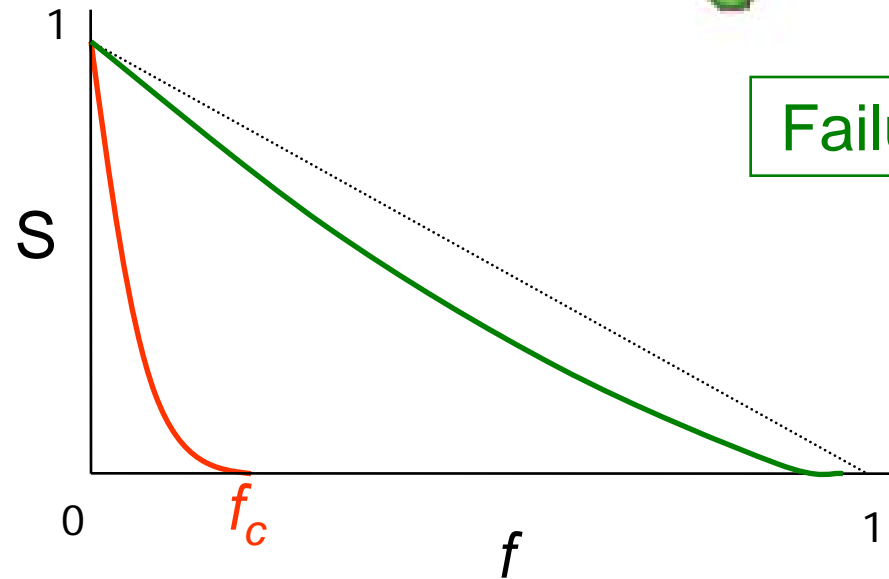


Consequences

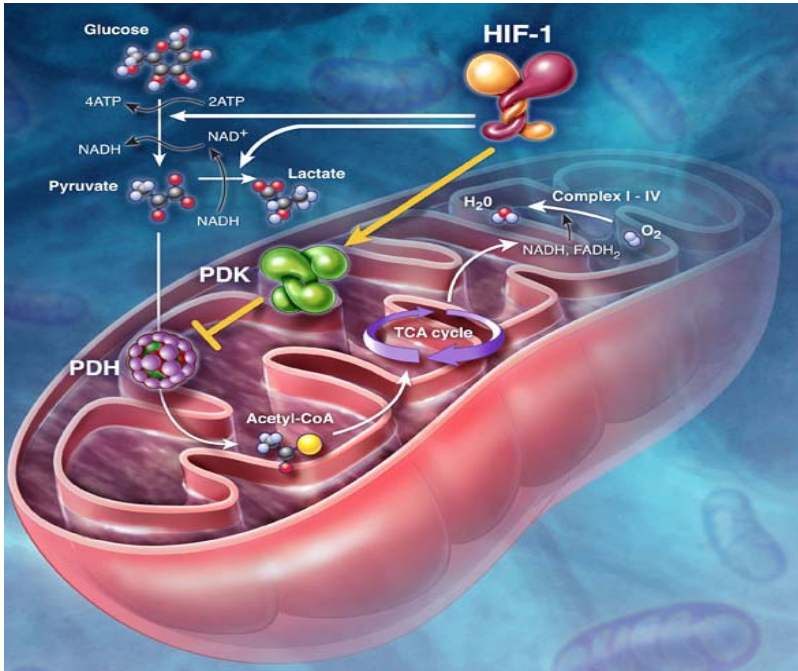
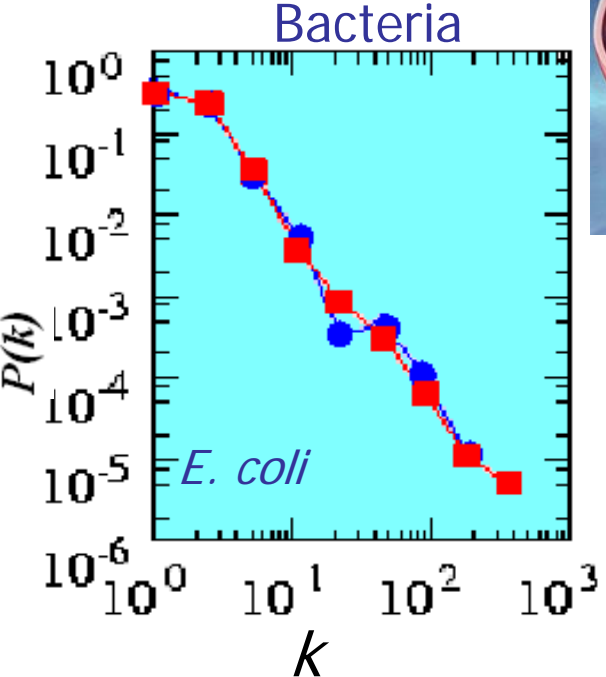
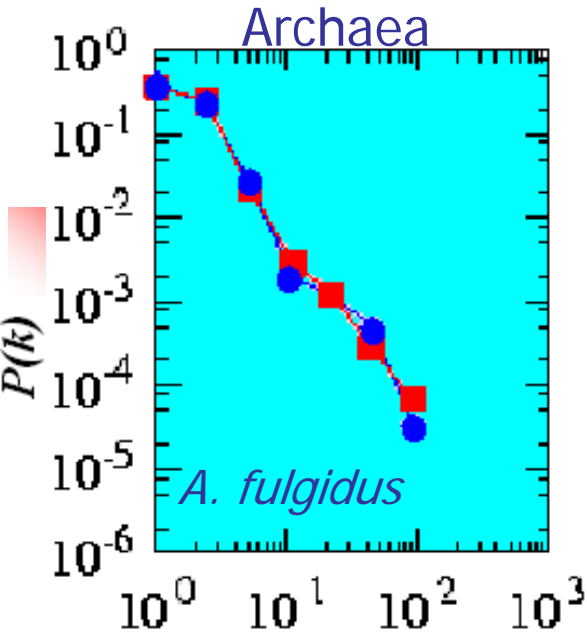


Attacks

Failures

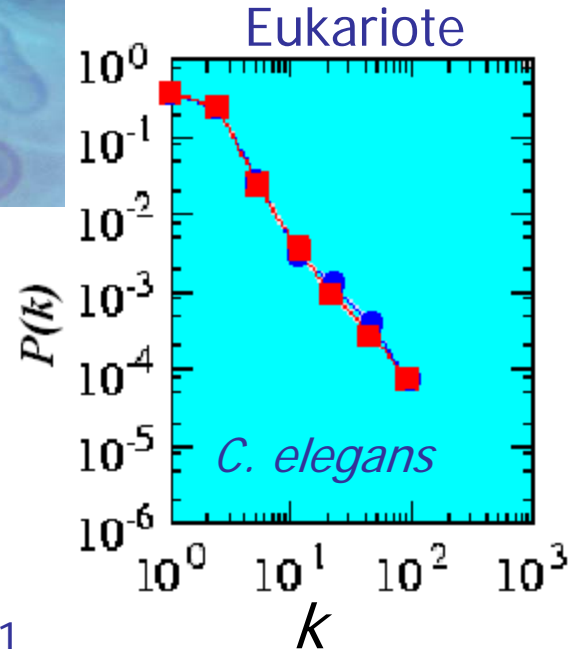


Metabolic Networks



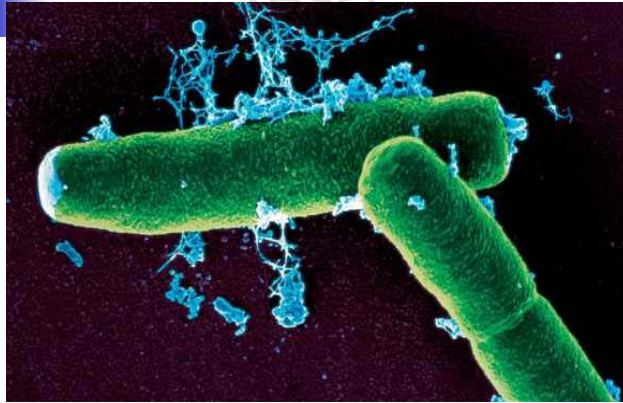
IN ●

OUT ■

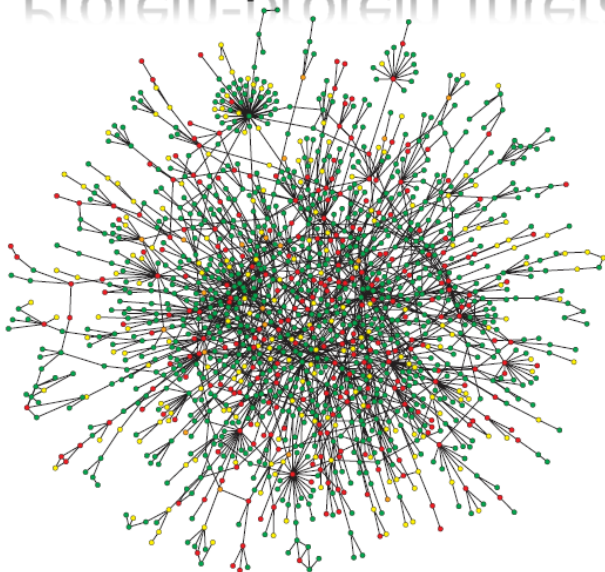


Protein Essentiality

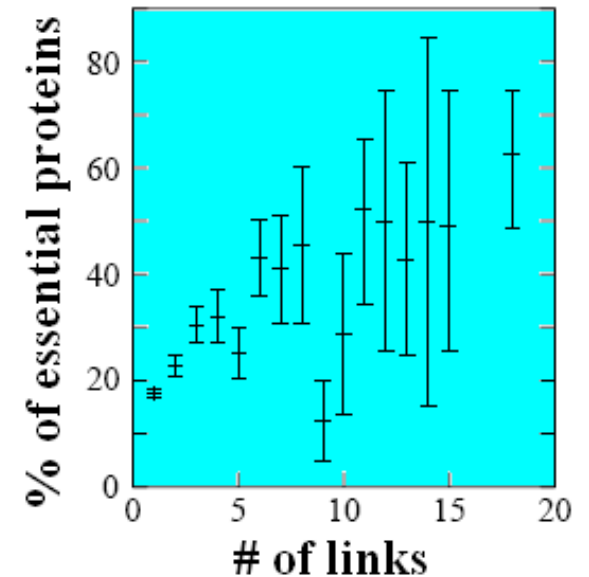
Yeast



Protein-protein Interactions



% Essential proteins vs. Protein Degree



H. Jeong, et al., Nature 411, 41 (2001)

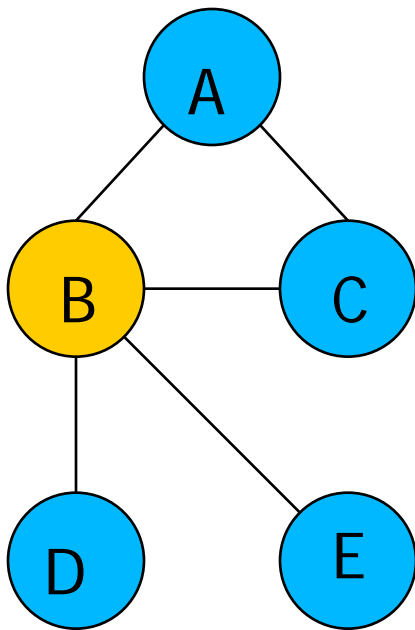


Betweenness centrality

- The number of shortest paths in the graph that pass through the node divided by the total number of shortest paths.

$$BC(k) = \sum_i \sum_j \frac{\rho(i, k, j)}{\rho(i, j)}, \quad i \neq j \neq k$$

Betweenness centrality



- Shortest paths are:

- AB, AC, ABD, ABE, BC, BD, BE, CBD, CBE, DBE

$$\rho(A, B, D) = 1; \quad \rho(A, D) = 1$$

$$\rho(A, B, E) = 1; \quad \rho(A, E) = 1$$

$$\rho(C, B, D) = 1; \quad \rho(B, D) = 1$$

$$\rho(C, B, E) = 1; \quad \rho(C, E) = 1$$

$$\rho(D, B, E) = 1; \quad \rho(D, E) = 1$$

- B has a BC of 5



Betweenness centrality

- Nodes with a high betweenness centrality are interesting because they
 - control information flow in a network
 - may be required to carry more information
- And therefore, such nodes
 - may be the subject of targeted attack

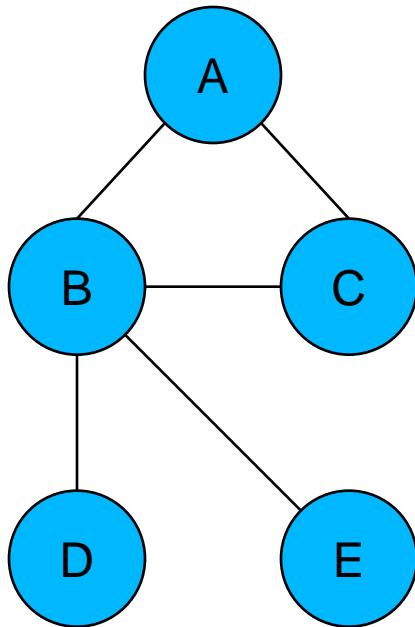


Closeness centrality

- The normalised inverse of the sum of topological distances in the network.

$$CC(i) = \frac{N - 1}{\sum_j d(i, j)}$$

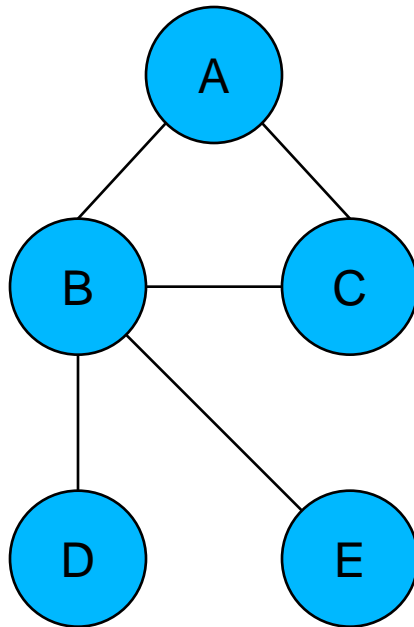
Closeness centrality



	<i>A</i>	<i>B</i>	<i>C</i>	<i>D</i>	<i>E</i>	$\sum_{j=1}^n d(i, j)$
<i>A</i>	0	1	1	2	2	6
<i>B</i>	1	0	1	1	1	4
<i>C</i>	1	1	0	2	2	6
<i>D</i>	2	1	2	0	2	7
<i>E</i>	2	1	2	2	0	7



Closeness centrality



Closeness

0.67

1.00

0.67

0.57

0.57

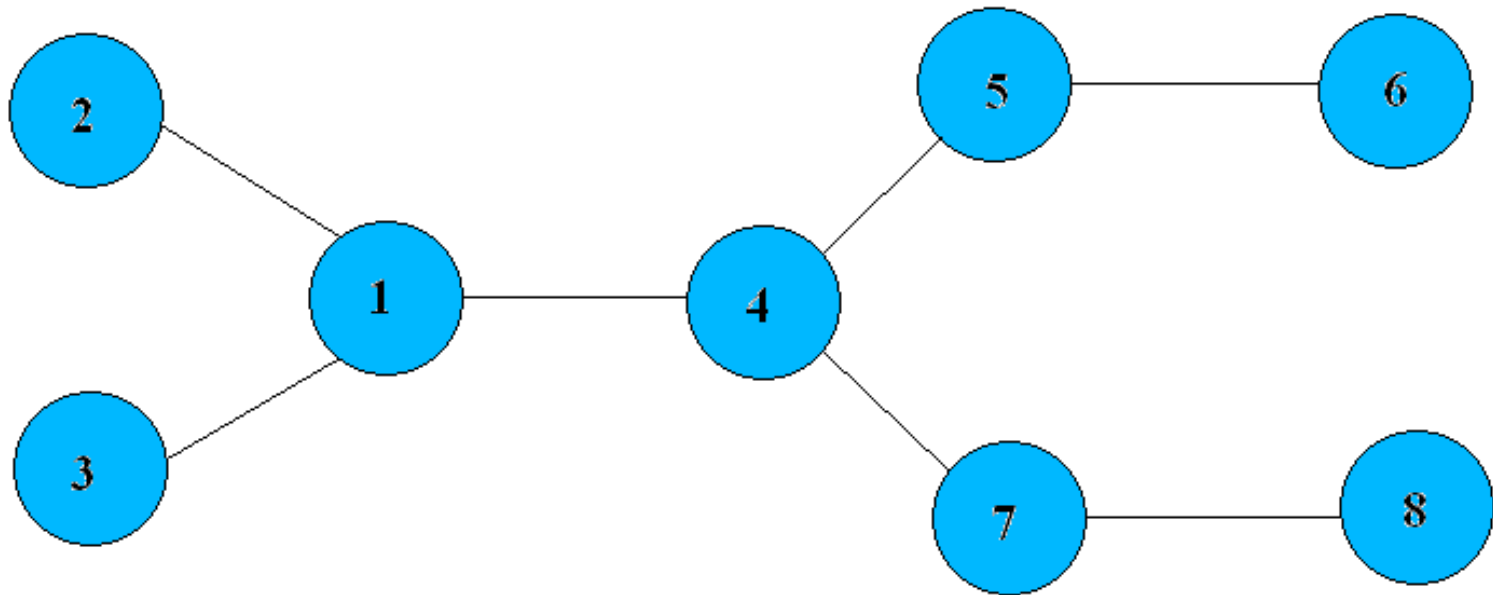


Closeness centrality

- Node B is the most central one in spreading information from it to the other nodes in the network.



Degree: Difficulties





Extending the Concept of Degree

Make x_i proportional to the average of the centralities of its i 's network neighbors

$$x_i = \frac{1}{\lambda} \sum_{j=1}^n A_{ij} x_j$$

where λ is a constant. In matrix-vector notation we can write

$$\mathbf{x} = \frac{1}{\lambda} \mathbf{A}\mathbf{x}$$

In order to make the centralities non-negative we select the *eigenvector* corresponding to the *principal eigenvalue* (Perron-Frobenius theorem).



Eigenvalues and Eigenvectors

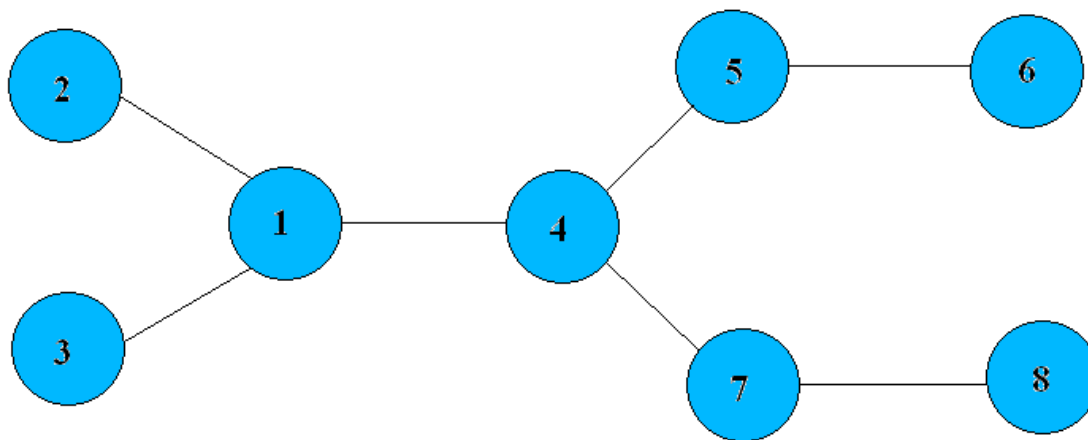
- The value λ is an **eigenvalue** of matrix A if there exists a non-zero vector x , such that $Ax = \lambda x$. Vector x is an **eigenvector** of matrix A
 - The largest eigenvalue is called the **principal eigenvalue**
 - The corresponding eigenvector is the **principal eigenvector**
 - Corresponds to the direction of maximum change



Eigenvector Centrality

- The corresponding entry of the principal eigenvector of the adjacency matrix of the network.
- It assigns relative scores to all nodes in the network based on the principle that connections to high-scoring nodes contribute more.

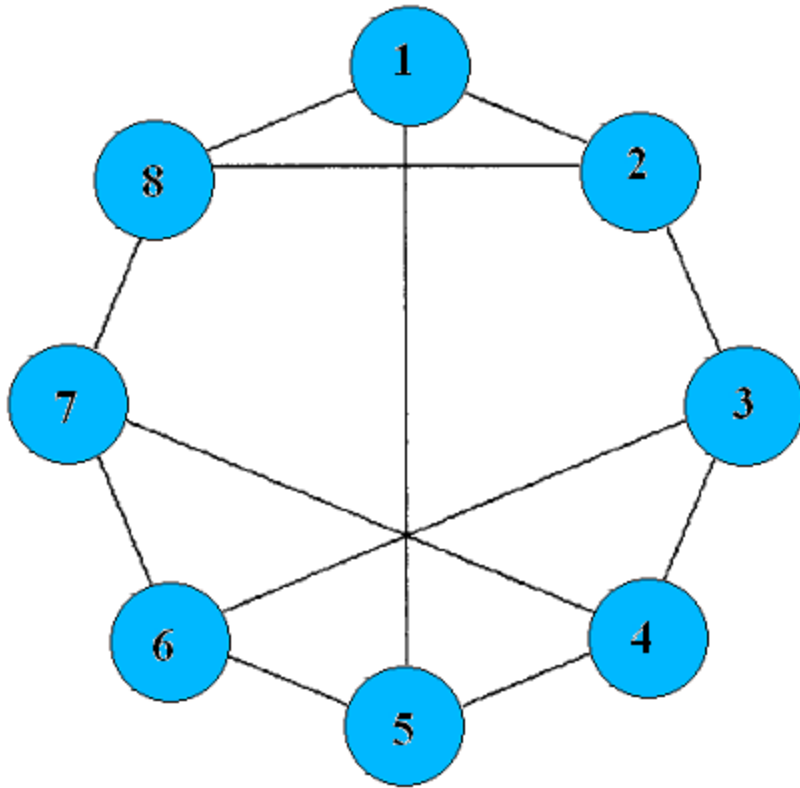
Eigenvector Centrality



Node EC

1	0.500
2	0.238
3	0.238
4	0.575
5	0.354
6	0.354
7	0.168
8	0.168

Eigenvector Centrality: Difficulties



In *regular networks* all nodes have exactly the same value of the *eigenvector centrality*, which is equal to $\frac{1}{\sqrt{n}}$

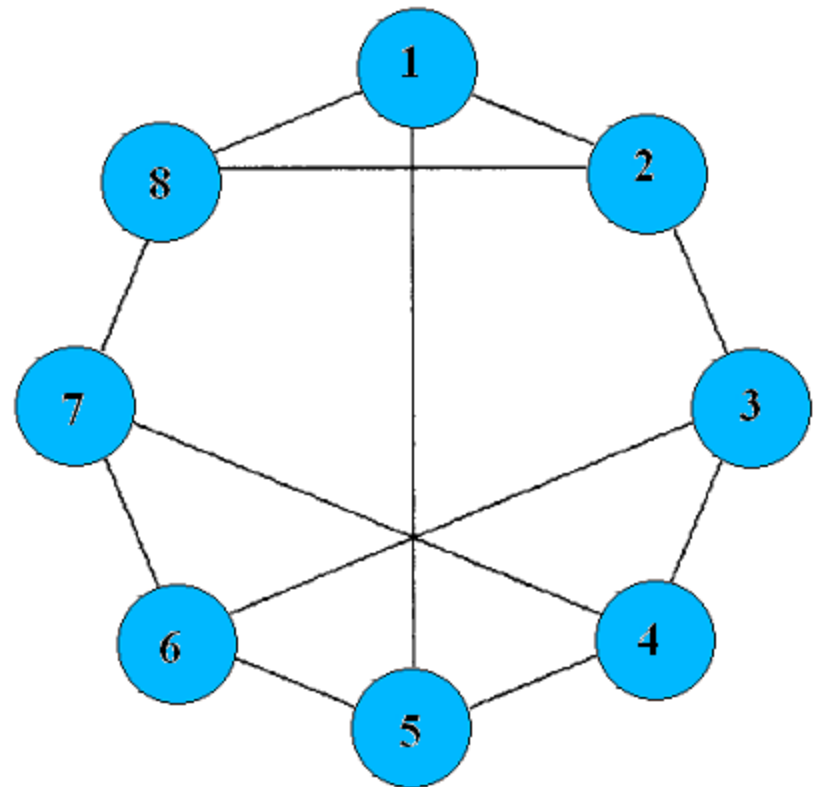
Subgraph Centrality

A *closed walk of length* k in a graph is a succession of k (not necessarily different) edges starting and ending at the same node, e.g.

1,2,8,1 (length 3)

4,5,6,7,4 (length 4)

2,8,7,6,3,2 (length 5)





Subgraph Centrality

The number of *closed walk of length k* starting at the same node i is given by the ii -entry of the k th power of the adjacency matrix

$$\mu_k(i) = (\mathbf{A}^k)_{ii}$$

$$\mathbf{A} = \begin{pmatrix} 0 & 1 & \cdots \\ 1 & 0 & \cdots \\ \vdots & \vdots & \ddots \end{pmatrix}$$

$$\mathbf{A}^2 = \begin{pmatrix} \mu_2(1) & & \\ & \mu_2(2) & \\ & & \ddots \end{pmatrix}$$

$$\mathbf{A}^3 = \begin{pmatrix} \mu_3(1) & & \\ & \mu_3(2) & \\ & & \ddots \end{pmatrix}$$



Subgraph Centrality

- We are interested in giving weights in decreasing order of the length of the *closed walks*. Then, visiting the closest neighbors receive more weight than visiting very distant ones.
- The subgraph centrality is then defined as the following weighted sum

$$\begin{aligned} EE(i) &= c_0 (A^0)_{ii} + c_1 (A)_{ii} + c_2 (A^2)_{ii} + c_3 (A^3)_{ii} + c_4 (A^4)_{ii} + \dots \\ &= \sum_{l=0}^{\infty} c_l \mu_l(i) \end{aligned}$$



Subgraph Centrality

- By selecting $c_l = 1/l!$ we obtain

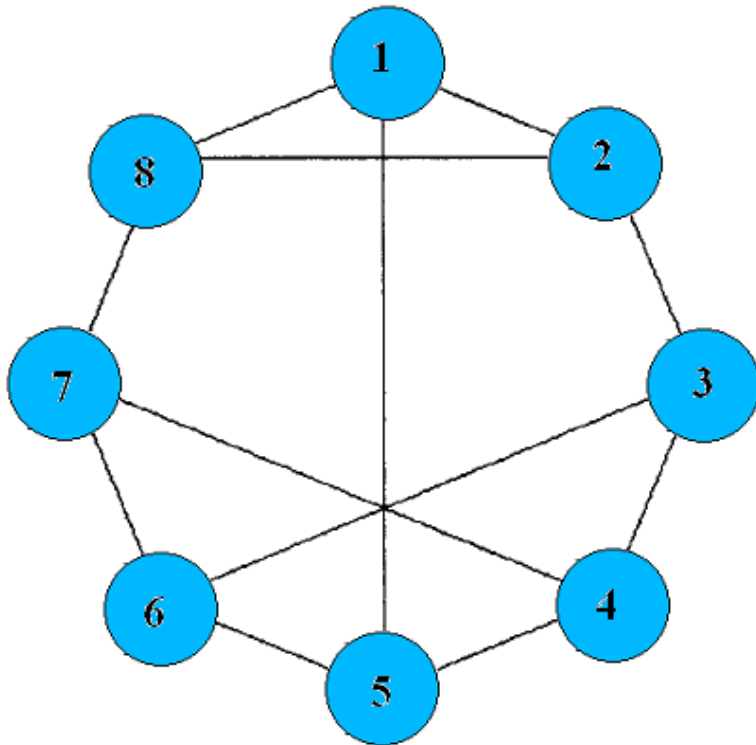
$$EE(i) = \sum_{l=0}^{\infty} \frac{\mu_l(i)}{l!} \quad EE(i) = \left(e^{\mathbf{A}} \right)_{ii}$$

where $e^{\mathbf{A}}$ is the exponential of the adjacency matrix.

- For simple graphs we have

$$EE(i) = \sum_{j=1}^n [x_j(i)]^2 e^{\lambda_j}$$

Subgraph Centrality



$$DC(i) = DC(j), \forall i \in V, j \in V$$

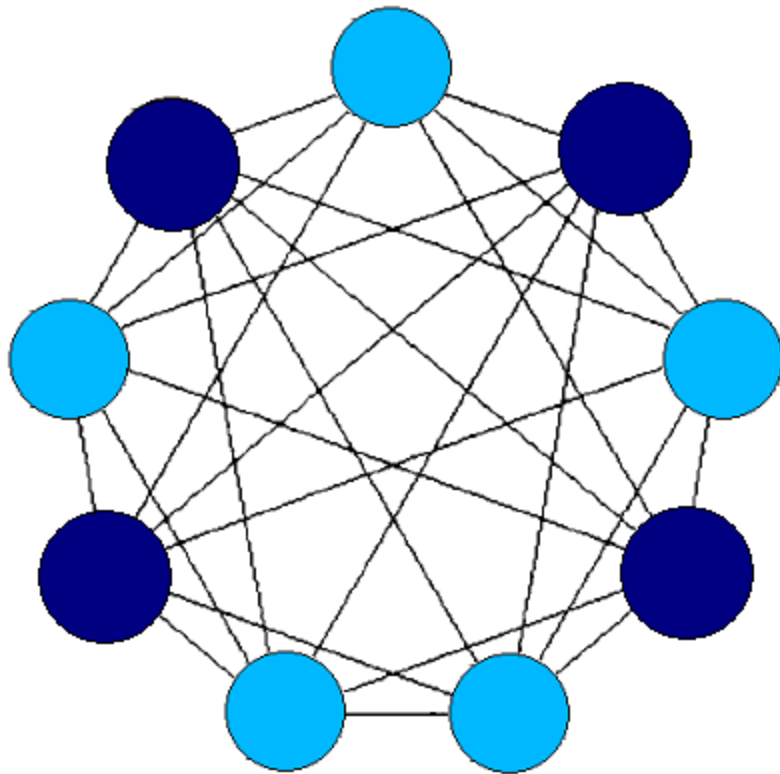
$$CC(i) = CC(j), \forall i \in V, j \in V$$

$$EC(i) = EC(j), \forall i \in V, j \in V$$

<u>Nodes</u>	<u>BC(i)</u>
1,2,8	9.528
4,6	7.143
3,5,7	11.111

<u>Nodes</u>	<u>EE(i)</u>
1,2,8	3.902
4,6	3.705
3,5,7	3.638

Subgraph Centrality





$$DC(i) = DC(j), \forall i \in V, j \in V$$

$$CC(i) = CC(j), \forall i \in V, j \in V$$

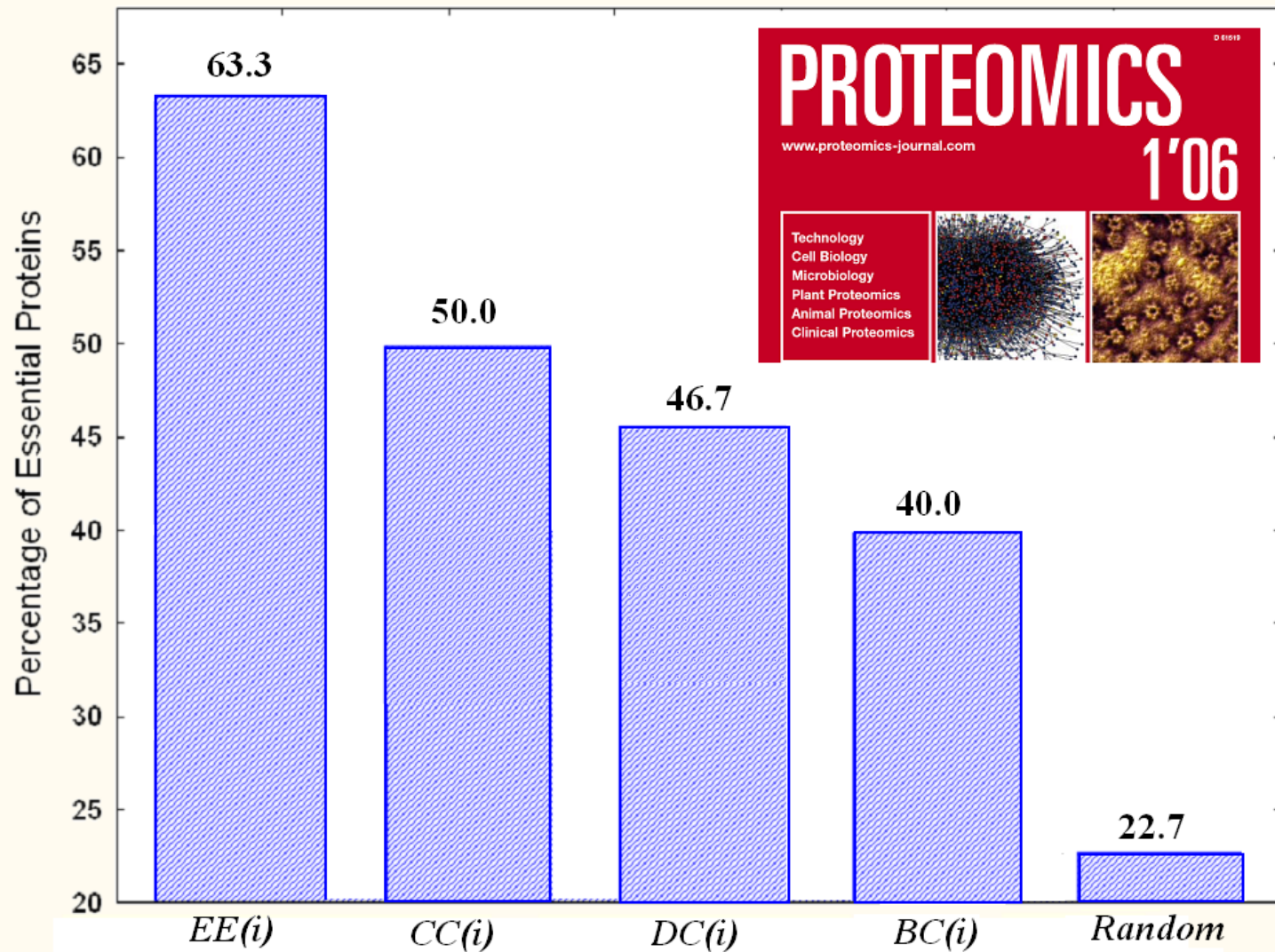
$$BC(i) = BC(j), \forall i \in V, j \in V$$

$$EC(i) = EC(j), \forall i \in V, j \in V$$

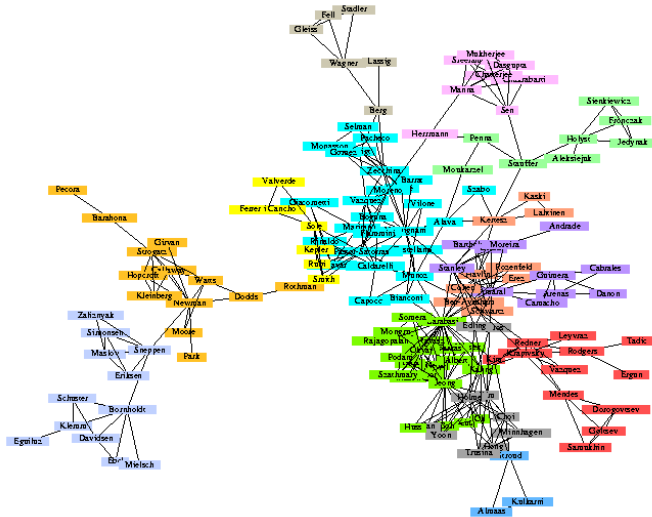
<u>Nodes</u>	<u>EE(i)</u>
	45.696
	45.651

Centrality and Protein Essentiality

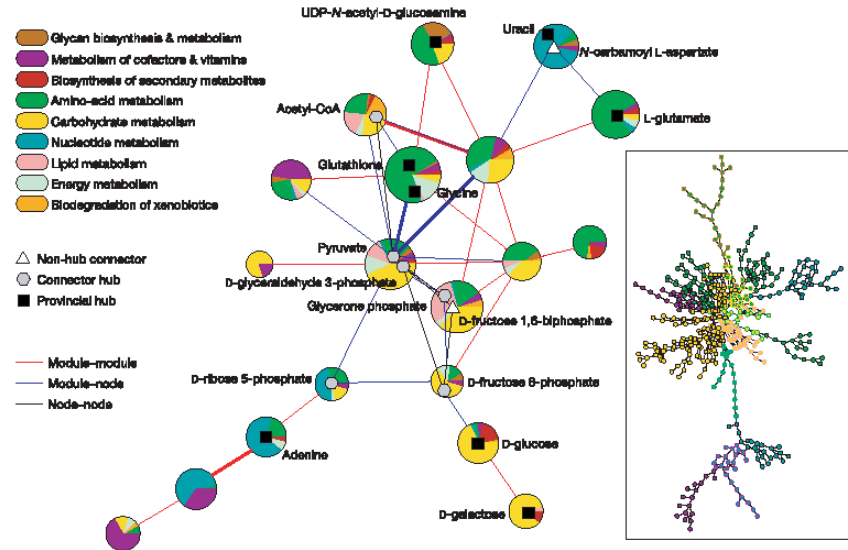
Estrada, E., *Proteomics* 6, 35 (2006)



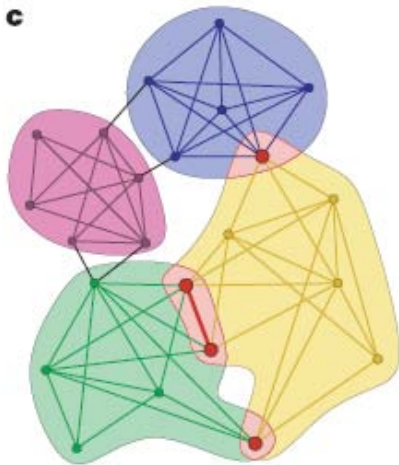
Communities



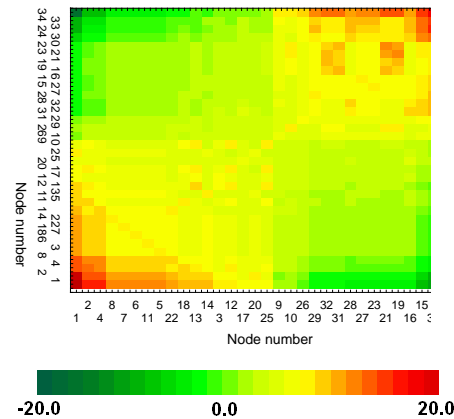
Girvan & Newman, *PNAS-USA* 99, 2002, 7821



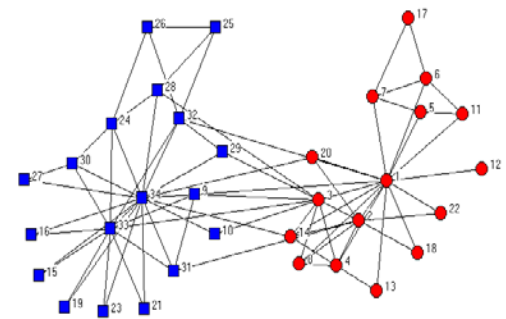
Guimerá & Amaral, *Nature* 433, 2005, 895



Palla, et al., *Nature* 435, 2005, 814



Estrada & Hatano, *Phys. Rev. E* 77, 2008, 036111





References

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- Wasserman & Faust, *Social Network Analysis*, Cambridge University Press, 2008.
- Estrada & Rodríguez-Velázquez, *Phys. Rev. E* **2005**, 71, 056103.
- Estrada & Hatano, *Phys. Rev. E* **2008**, 77, 036111.

Exercise 1

Identify the most central node according to the following criteria:

- (a) the largest chance of receiving information from closest neighbors;
- (b) spreading information to the rest of nodes in the network;
- (c) passing information from some nodes to others.

