Spectral Analysis

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Eigenvalue and Eigenvector

- Let A be an nxn matrix.
- A scalar λ is called an Eigenvalue of A if there is a nonzero vector X such that AX = λ X. Such a vector X is called an Eigenvector of A corresponding to λ .

• Example: 2
1 is an Eigenvector of A =
$$\begin{bmatrix} 3 & 2 \\ 3 & 2 \end{bmatrix}$$
 for $\lambda = 4$
 $\begin{pmatrix} 3 & 2 \\ 3 & -2 \end{bmatrix}$
 $\begin{pmatrix} 3 & 2 \\ 3 & -2 \end{bmatrix} \begin{pmatrix} 2 \\ 3 & -2 \end{bmatrix} \begin{pmatrix} 2 \\ 3 & -2 \end{bmatrix}$
 $\begin{pmatrix} 3 & 2 \\ 3 & -2 \end{bmatrix} \begin{pmatrix} 2 \\ 3 & -2 \end{bmatrix} \begin{pmatrix} 2 \\ 3 & -2 \end{bmatrix}$
 $\begin{pmatrix} 3 & 2 \\ 3 & -2 \end{bmatrix} \begin{pmatrix} 2 \\ 3 & -2 \end{pmatrix} \begin{pmatrix} 2 \\ 3 & -2 \end{pmatrix}$
 $\begin{pmatrix} 3 & 2 \\ 3 & -2 \end{pmatrix} \begin{pmatrix} 2 \\ 3 & -2 \end{pmatrix} \begin{pmatrix} 2 \\ 3 & -2 \end{pmatrix}$

Finding Eigenvalues and Eigenvectors

$$A = \begin{bmatrix} 7 & 3 \\ 3 & -1 \end{bmatrix}$$

$$(4) Solving for \lambda:
$$(\lambda - 8) (\lambda + 2) = 0$$

$$\lambda = 8 \text{ and } \lambda = -2 \text{ are the Eigen values}$$

$$(5) \text{ Consider } A - \lambda I$$

$$(7 - \lambda - 3)$$

$$(7 - \lambda - 3)$$$$

Finding Eigenvalues and Eigenvectors

For
$$\lambda = -2$$

 $\begin{bmatrix} 7 - (-2) & 3 \\ 3 & -1 - (-2) \end{bmatrix} = \begin{bmatrix} 9 & 3 \\ 3 & 1 \end{bmatrix}$

Solve B X = 0

$$\begin{bmatrix}
9 & 3 \\
3 & 1
\end{bmatrix}
\begin{bmatrix}
X1 \\
X2
\end{bmatrix} =
\begin{bmatrix}
0 \\
0
\end{bmatrix}$$

$$9X1 + 3X2 = 0 \longrightarrow X2 = -3X1$$

$$3X1 + X2 = 0 \longrightarrow X2 = -3X1$$
If X1 = 1;
$$\begin{bmatrix}
1 \\
-3
\end{bmatrix}$$
is an eigenvector for $\lambda = -2$

Verification
$$A = \begin{bmatrix} 7 & 3 \\ 3 & -1 \end{bmatrix}$$

 $AX = \lambda X$
For $\lambda = 8$ and $X = \begin{bmatrix} 3 \\ 1 \end{bmatrix}$
 $\begin{bmatrix} 7 & 3 \\ 3 & -1 \end{bmatrix} \begin{bmatrix} 3 \\ 1 \end{bmatrix} = \begin{bmatrix} 24 \\ 8 \end{bmatrix} = 8 \begin{bmatrix} 3 \\ 1 \end{bmatrix}$

An *n*x*n* matrix has *n* eigenvalues and the corresponding eigenvectors.

Spectral Analysis

- Spectral decomposition is a method of projecting the characteristics of a network graph in n-dimensions or directions (that are mutually perpendicular) where n is the number of vertices in the graph.
- The projection in each direction is represented in the form of a scalar value (called the eigenvalue) and its corresponding vector with entries for each vertex (called the eigenvector).
- The largest eigenvalue of the projection is called the principal eigenvalue (a.k.a. spectral radius) and the corresponding eigenvector is called the principal eigenvector.
- We will use the adjacency matrix of a network graph to determine its eigenvalues and eigenvectors.

Example: Eigenvalues and Eigenvectors

2

3

4

5

3

4

6

4

8

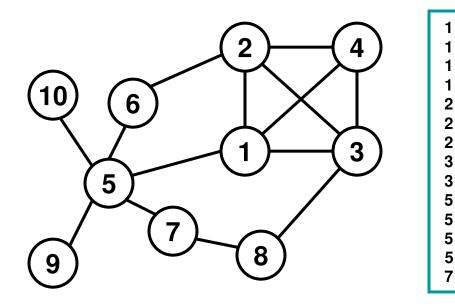
6

7

9

10

8



List of edges written in the increasing order of the node IDs from left to right and top to bottom.

Property: For a matrix with all positive entries, the Principal Eigenvalue is always positive & the entries in the Principal Eigenvector are also positive

Node I	Ds ←				Eigenva	alues —				\longrightarrow
	3.3893	1.888	1.1131	0.3197	0	-0.3741	-0.8563	-1.3098	-1.6828	-2.4871
1	0.4741	0.0641	-0.1261	0.2899	0.0000	-0.3898	0.5219	0.0776	0.3210	-0.3711
2	0.4568	0.2043	-0.2089	-0.3262	0.0000	0.0872	-0.2158	0.6614	0.1384	0.2979
3	0.4453	0.2648	0.2600	0.1054	0.0000	0.3619	0.2312	-0.0638	-0.6797	-0.0477
4	0.4060	0.2824	-0.0674	0.2161	0.0000	-0.1585	-0.6275	-0.5155	0.1309	0.0486
5	0.2986	-0.6305	-0.1241	0.0974	0.0000	-0.1448	0.1651	-0.1838	-0.1298	0.6242
6	0.2229	-0.2257	-0.2991	-0.7157	0.0000	0.1538	0.0591	-0.3646	-0.0051	-0.3708
7	0.1390	-0.3609	0.5100	-0.1521	0.0000	-0.4838	-0.3369	0.2471	-0.2518	-0.3086
8	0.1724	-0.0509	0.6917	-0.1460	0.0000	0.3258	0.1234	-0.1399	0.5536	0.1433
9	0.0881	-0.3339	-0.1114	0.3046	0.7071	0.3870	-0.1928	0.1403	0.0771	-0.2510
10	0.0881	-0.3339	-0.1114	0.3046	-0.7071	0.3870	-0.1928	0.1403	0.0771	-0.2510

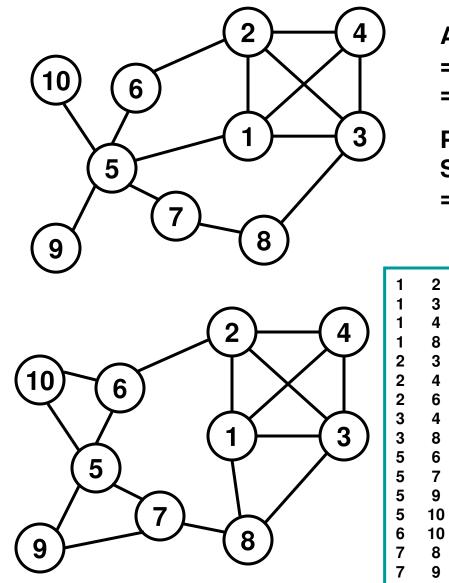
Each column corresponds to an Eigenvector with an entry for each node

Spectral Radius Ratio for Node Degree

- Spectral radius ratio for node degree (for an undirected graph) is the ratio of the principal eigenvalue of the adjacency matrix and the average degree of the vertices in the graph.
- If Kmin, Kavg and Kmax are the minimum, average and maximum values for the node degrees, then:
 - Kmin \leq Kavg \leq Principal Eigenvalue \leq Kmax
- So, the spectral radius ratio for node degree of a graph is always greater than or equal to 1.
- The spectral radius ratio for node degree is a measure of the variation in the node degrees. The farther is the value from 1, the larger the variation in node degree.
- The spectral radius ratio for node degree can be uniformly applied across networks of all size and be used to evaluate the relative variation in node degree.

N. Meghanathan, "Spectral Radius as a Measure of Variation in Node Degree for Complex Network Graphs," Proceedings of the *3rd International Conference on Digital Contents and Applications*, (DCA 2014), pp. 30-33, Hainan, China, December 20-23, 2014.

Spectral Radius Ratio for Node Degree



Avg. Degree = Sum of Node Degrees / N = (4 + 4 + 4 + 3 + 5 + 2 + 2 + 2 + 1 + 1)/10 = 2.8

Principal Eigenvalue = 3.3893 Spectral Radius Ratio for Node Degree = 3.3893 / 2.8 = 1.21

Avg. Degree = Sum of Node Degrees / N = (4 + 4 + 4 + 3 + 4 + 3 + 3 + 3 + 3 + 2 + 2)/10 = 3.2

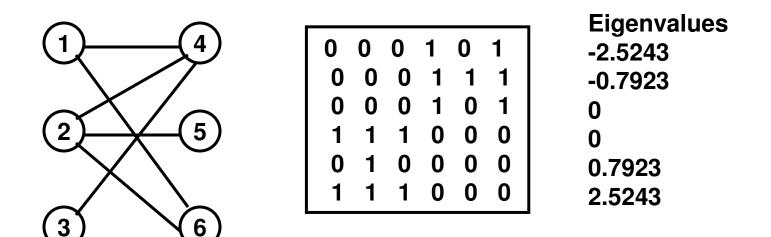
Principal Eigenvalue = 3.4735 Spectral Radius Ratio for Node Degree = 3.4735 / 3.2 = 1.09

Applications

- Bipartivity Index: Detection of bipartivity in graphs
- Estrada Index: Protein folding
- Laplacian Matrix: Determining the number of components, connectivity and number of spanning trees in a graph
- Hierarchical Community Detection
- Prediction of Graph Isomorphism

Eigen Values of a Bipartite Network

- There are even number of vertices.
- Let λ1, λ2, λ3, ..., λn be the n Eigenvalues of an nnode network.
- For any j = 1, 2, ..., n/2, if $\lambda_j = |-\lambda_{n-j+1}|$, then the network is almost bi-partite.



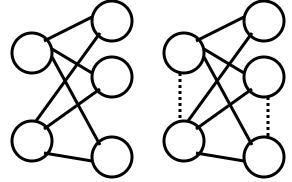
Bipartivity Check

 A bi-partite graph is the one that has two partitions V1 and V2 of its vertices such that

- V1 U V2 = V; V1 n V2 = Φ .

- No edges within V1 and within V2.
- All edges are those connecting V1 and V2.

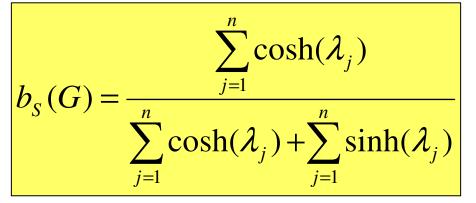
There should be NO odd length cycles in a truly bi-partite network



• Graphs can be considered close to bi-partite if there are few edges (not a significant number) called the frustrated edges that connect vertices within V1 and/or V2.

Computing the Bipartite Measure

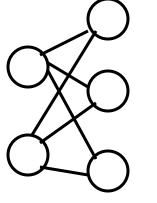
Compute the Eigenvalues (λ 1, λ 2, λ 3, ..., λ n) of the nxn adjacency matrix.

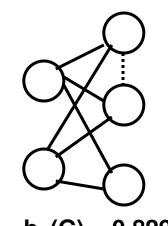


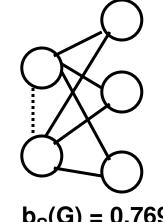
For a "truly" bi-partite graph, b_S(G) = 1; the sinh terms add to 0.

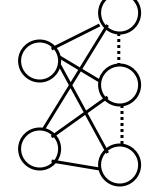
For a "close-to" bi-partite graph, $b_S(G) < 1$; the sinh terms add up to some small positive value.

Bipartivity Measure: Examples







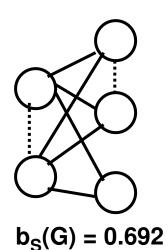


 $b_{S}(G) = 1.0$

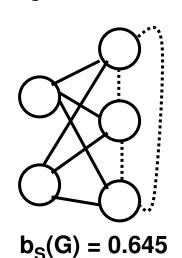
 $b_{S}(G) = 0.829$

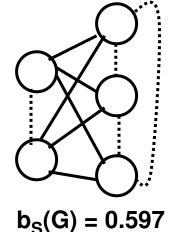
 $b_{S}(G) = 0.769$

 $b_{S}(G) = 0.731$



 $b_{S}(G) = 0.645$





For a given number of frustrated links, a larger bipartivity measure is observed if more of the frustrated links are present in the network with the Source: Estrada, E.; Rodriguez-Velazquez, J. A. Spectral measures of larger subset. bipartivity in complex networks. Physical Review E 72, 2005, 046105.

Bipartivity of Real Networks

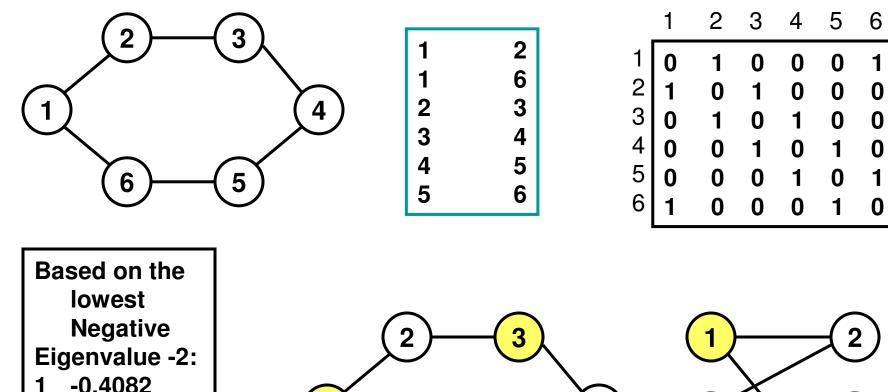
Type: Information	on	<u>Type:</u> PPIs	
Network	Bipartivity	Network	Bipartivity
	Measure		Measure
SciMet	0.500	Yeast	0.500
Roget	0.529	Human	0.576
		H. Pylori	0.711
<u>Type:</u> Social		A. Fulgidus	0.976
Network	Bipartivity		
	Measure	Type: Transcrip	
Drugs	0.500	Network	Bipartivity
Corporate Elite	0.500		Measure
Karate Club	0.597	Urchin	0.618
Saw Mill	0.749	E. Coli	0.831
		Yeast	0.960
Type: Food web	S		
Network	Bipartivity	Type: Technolo	gical
	Measure	Network	Bipartivity
Coachella	0.500		Measure
El Verde	0.500	USAir97	0.500
Grassland	0.743	Internet	0.502
Stony stream	0.815	Electronic3	0.952

Source: Estrada, E.; Rodriguez-Velazquez, J. A. Spectral measures of bipartivity in complex networks.*Physical Review E* 72, **2005**, 046105.

Identifying Bipartite Subsets using Eigenvalue and Eigenvector

- We identify the smallest Eigenvalue (most likely a negative value), hereafter called the bi-partite Eigenvalue, and its corresponding Eigenvector, hereafter called the bi-partite Eigenvector.
- The values in the bi-partite Eigenvector will be positive and negative.
 - The node IDs whose entries are of the same sign in the bi-partite Eigenvector form the two subsets.
 - The vertices that are of the same sign are more likely not to have links between them, and are more likely to have links with vertices of the other sign.
 - Each of the two subsets will have the minimum (or zero, if possible) number of frustrated links. Most of the links are likely to be between the vertices in the two subsets.

Identifying Bipartite Subsets using Eigenvalue and Eigenvector: Ex. 1 (1)



6

1

4

5

3

5

6

- -0.4082
 0.4082
- 3 -0.4082
- 4 0.4082
- 5 -0.4082
- 6 0.4082

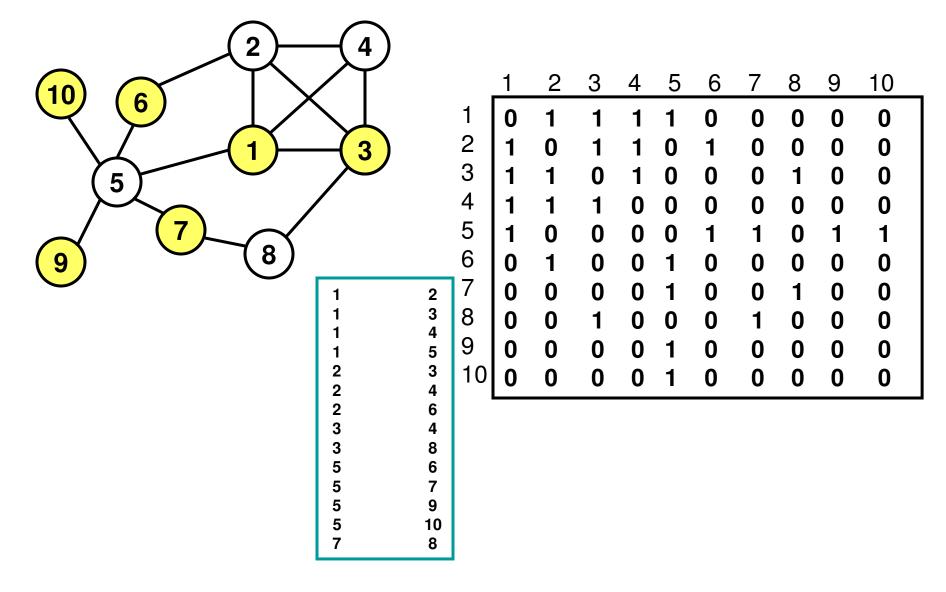
Identifying Bipartite Subsets using Eigenvalue and Eigenvector: Ex. 1 (2)

Eigenvalue, λ	cosh(λ)	sinh(λ)
-2	3.7622	-3.6269
-1	1.5431	-1.1752
-1	1.5431	-1.1752
1	1.5431	1.1752
1	1.5431	1.1752
2	3.7622	3.6269
Total	13.6968	0

$$b_{S}(G) = \frac{\sum_{j=1}^{n} \cosh(\lambda_{j})}{\sum_{j=1}^{n} \cosh(\lambda_{j}) + \sum_{j=1}^{n} \sinh(\lambda_{j})}$$

$$b_{S}(G) = \frac{13.6968}{(13.6968 + 0)}$$
$$= 1.0$$

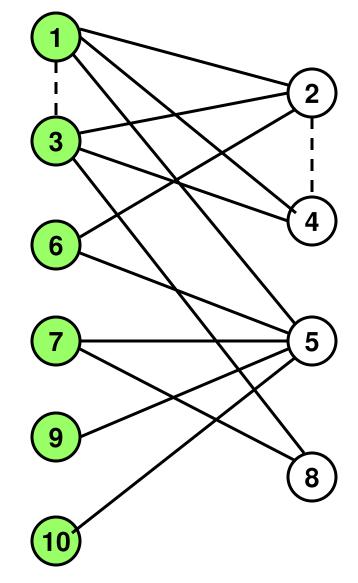
Identifying Bipartite Subsets using Eigenvalue and Eigenvector: Ex. 2 (1)



Identifying Bipartite Subsets using Eigenvalue and Eigenvector: Ex. 2 (2)

Based on the lowest Negative Eigenvalue -2.4870836366555165:

- 1 0.3711379191456978
- 2 -0.2979483534591263
- 3 0.04772178019913417
- 4 -0.048615713642947846
- 5 -0.6242087587467021
- 6 0.3707784887547619
- 7 0.3085825736313854
- 8 -0.14326191068896146
- 9 0.25098020410206323
- 10 0.250980204102063



Identifying Bipartite Subsets using Eigenvalue and Eigenvector: Ex. 2 (3)

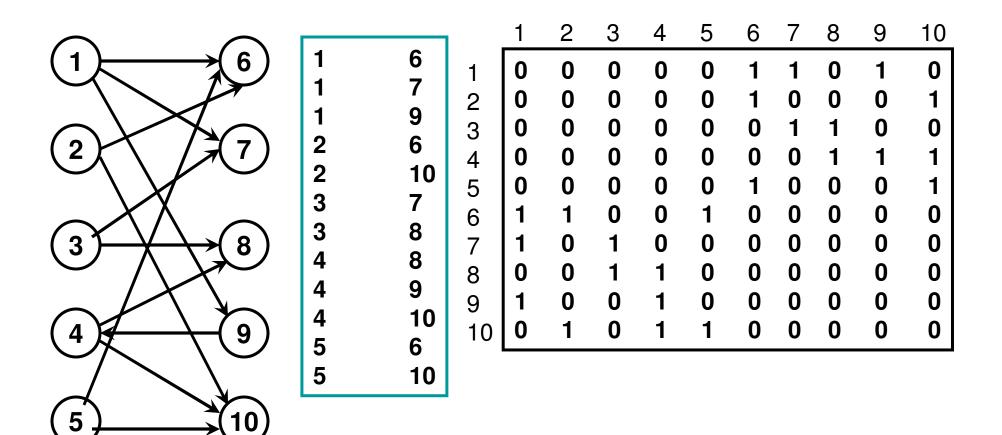
Eigenvalue, λ	cosh(λ)	sinh(λ)
-2.4871	6.0547	-5.9716
-1.6828	2.7832	-2.5974
-1.3098	1.9877	-1.7178
-0.8564	1.3897	-0.9650
-0.3741	1.0708	-0.3829
0	1.0	0
0.3197	1.0515	0.3252
1.1131	1.6862	1.3576
1.8880	3.3788	3.2274
3.3893	14.839	14.8057
Total	35.2416	8.0812

$$b_{S}(G) = \frac{\sum_{j=1}^{n} \cosh(\lambda_{j})}{\sum_{j=1}^{n} \cosh(\lambda_{j}) + \sum_{j=1}^{n} \sinh(\lambda_{j})}$$

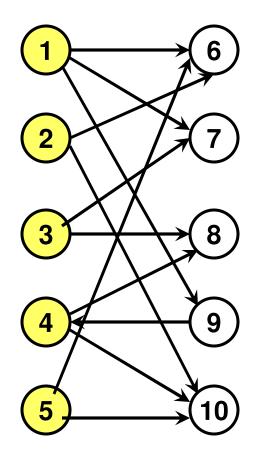
Bipartite Partition Detection: Digraph

- When confronted with a directed graph, first transform the directed graph to an undirected graph and determine the two partitions as explained previously using the Eigenvector approach.
- After identifying the partitions, restore the directions of the edges.
- In a directed graph, the edges typically point from one set of vertices to the other set of vertices.
 - There could be scenarios where the edges could point in the reverse direction; as long as we know the direction of the edges, we could restore them after determining the two partitions.

Digraph Bipartivity Detection: Example (1)



Digraph Bipartivity Detection: Example (2)



Base	d on the lowest Negative
	Eigenvalue -2.4998:
1	-0.3625951312994192
2	-0.32242994615264337
3	-0.17066751642864314
4	-0.3625951312994189
5	-0.32242994615264337
6	0.40300972011359426
7	0.21331972680920852
8	0.21331972680920852
9	0.29009605184492576
10	0.4030097201135942

Protein Folding

- Protein folding is the process by which a protein transforms from a random coil (sequence of amino acids: linear polypeptide chain) to its characteristic 3-dimensional structure that is essential to its expected function.
- The correct three-dimensional structure is essential to function, although some parts of functional proteins may remain unfolded.
- Failure to fold into native structure generally produces inactive proteins, but in some instances misfolded proteins have modified or toxic functionality.
- When modeled as a graph, the more flat (linear chain) is the graph, the less the folding and vice-versa.

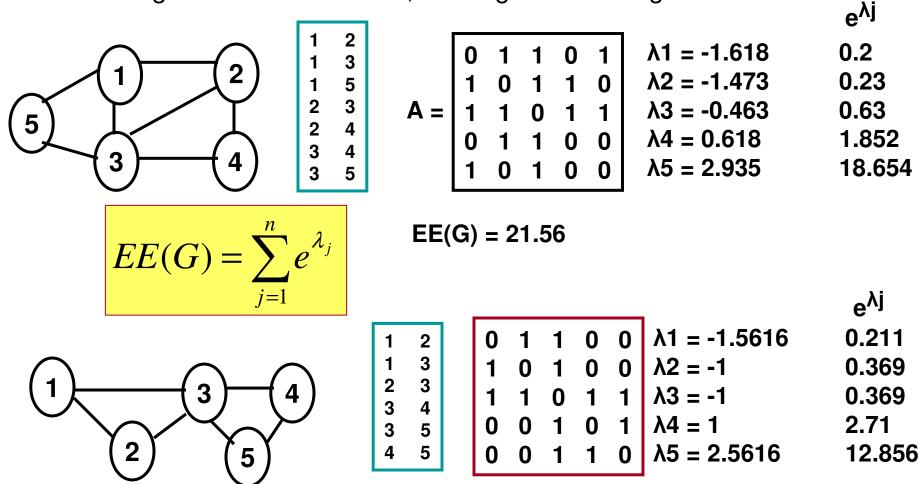
Estrada, E. (2000). "Characterization of 3D molecular structure". *Chem. Phys. Lett.* (319): 713

Source: Wikipedia

Estrada Index of Graphs

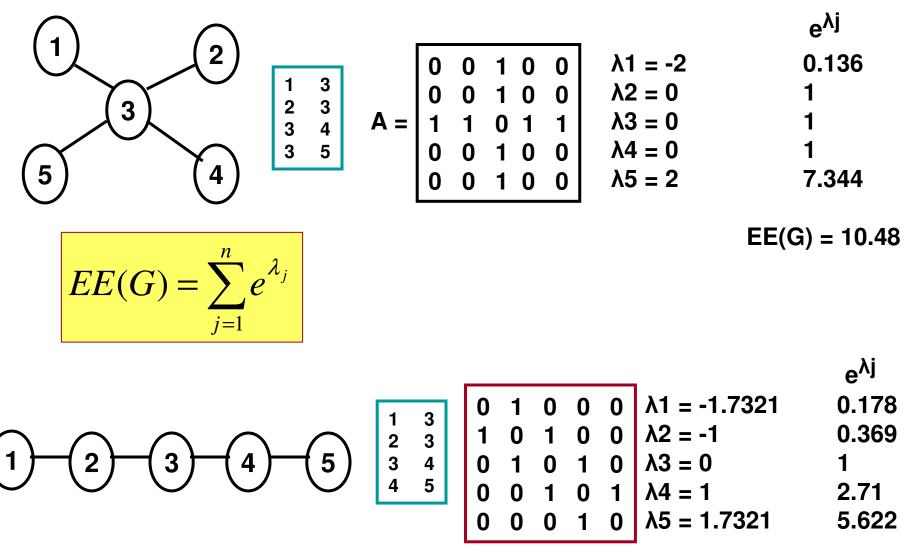
• The Estrada Index can be used to determine the degree of folding of a protein.

- Larger the Estrada Index, the larger the folding.



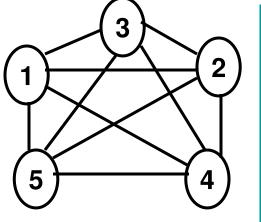
EE(G) = 16.51

Estrada Index of Star and Chain



EE(G) = 9.88

Estrada Index of Complete Graph

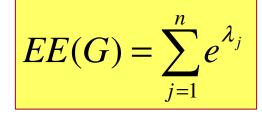


1	2	
1	3	
1	4	
1	5	
2	3	
2	4	
2	5	
3	4	
2 2 3 3 4	2 3 4 5 3 4 5 4 5 5	
4	5	

	0	1	1	1	1	λ1 = -1
	1	1 0 1 1	1	1	1	λ2 = -1
A =	1	1	0	1	1	λ3 = -1
	1	1	1	0	1	λ4 = -1
	1	1	1	1	0	λ5 = 4

e ^{λj}
0.368
0.368
0.368
0.368
54.576

EE(G) = 56.048



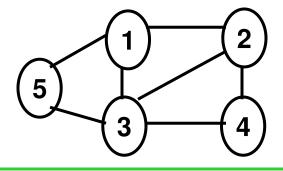
Foldina Effectiveness

Folding Effectiveness(G) = $-\frac{1}{2}$

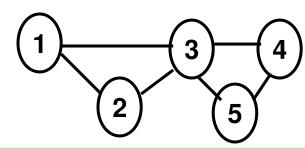
Estrada Index Complete Graph (G)

Estrada Index (G)

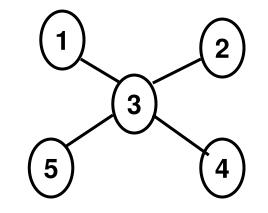
Closer is the value to 1, the more folded is the protein



Folding Effectiveness (G) = 21.56 / 56.048 = 0.385



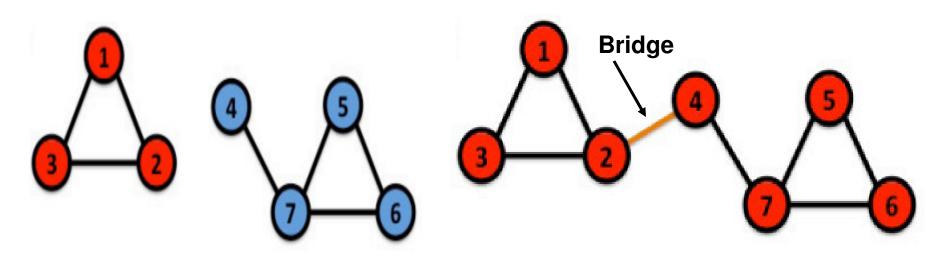




Folding Effectiveness (G) = 10.48 / 56.048 = 0.187

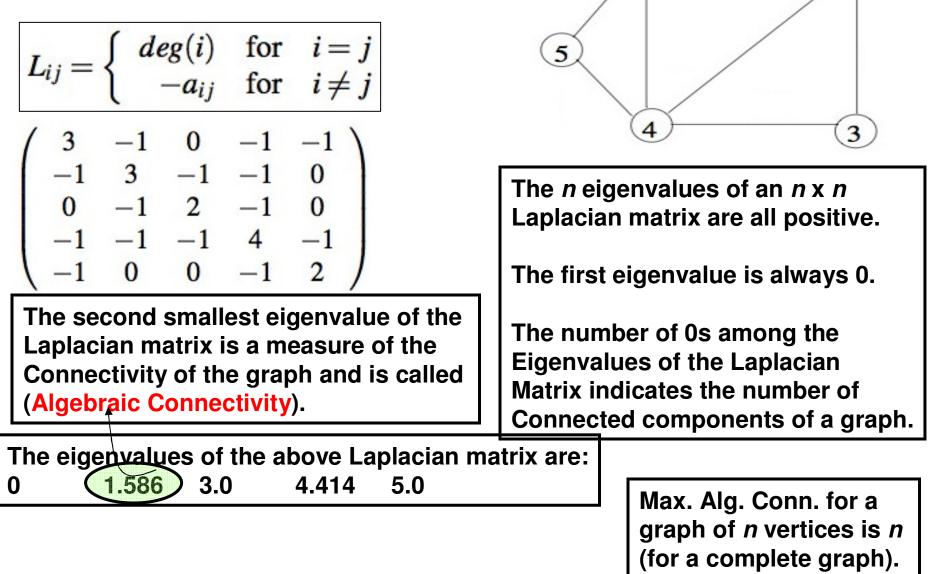
Components (Clusters)

- The vertices of a graph are said to be in a single component if there is a path between the vertices.
- A graph is said to be connected if all its vertices are in one single component; otherwise, the graph is said to be disconnected and consists of multiple components.
 - Adding one or more links (bridges) can connect the different components



Laplacian Matrix

• Laplacian Matrix L = D - A



Algebraic Connectivity

Algebraic Connectivity: 1.108

Traditional Connectivity: 2

L	ap	lac	ia	n N	lat	rix	Eigenvalues
	1	2	3	4	5	6	0
1	2	-1	0	-1	0	0	1.108
2	-1	4	-1	0	-1	-1	
3	0	-1	2	0	0	-1	2.295
4	-1	0	0	2	-1	0	3.0
5	0	-1	0	-1	3	-1	4.317
6	0	-1	-1	0	-1	3	
					Gra	iph 1	5.278

If we remove vertices 1 and 5, Graph 1 will get disconnected, but not Graph 2 Larger the algebraic connectivity of a graph, lower the chances of a network to get disconnected due to node removal.

1 - 3	Laplacian Matrix	Eigenvalues
$\Psi \rightarrow \Psi$	1 2 3 4 5 6	0
	1 2 -1 0 -1 0 0 2 -1 4 -1 -1 -1 -1	1.382
	3 0 -1 2 0 0 -1	2.382
	4 -1 -1 0 2 -1 0	3.618
Algebraic Connectivity: 1.382	5 0 -1 0 -1 3 -1 6 0 -1 -1 0 -1 3	4.618
Traditional Connectivity: 2	Graph 2	5.999

Laplacian Matrix # Components

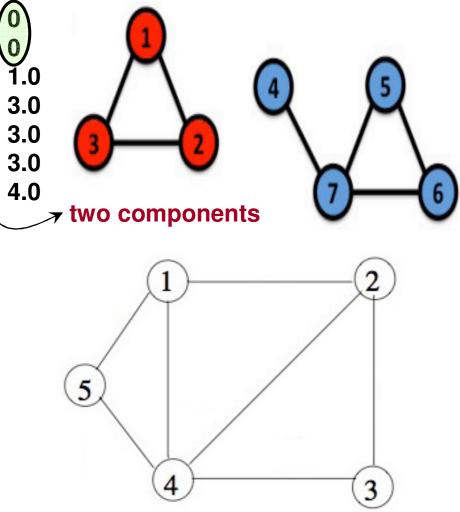
Laplacian Matrix # Spanning Trees

If $\mu_1 = 0 < \mu_2 \le \mu_3 \le ... \le \mu_n$ are the *n* Eigenvalues of the Laplacian matrix of a connected graph of *n* vertices

The # spanning trees of the graph is then

$$\frac{1}{n}\prod_{i=2}^{n}\mu_{i}$$

The 7 eigenvalues of the Laplacian Matrix are:

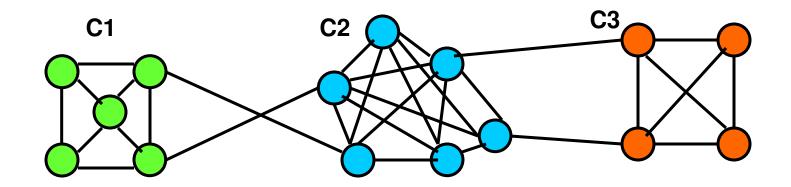


The five eigenvalues of the Laplacian matrix of the above graph are: 0, 1.586, 3.0, 4.414, 5.0

spanning trees = (1/5)(1.586*3*4.414*5) = 21.

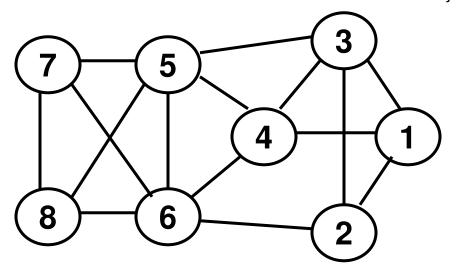
Community

- Community: It is formed by individuals such that those within a group interact with each other more frequently than with those outside the group.
- The paths among vertices within a community is most likely to involve only the other vertices within the community.
- For a highly modular community of vertices, the number of edges connecting them within the community is significantly greater than the number of edges connecting them to vertices outside the community.



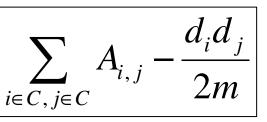
Modularity Maximization

- Modularity measures the strength of a community partition by taking into account the degree distribution.
- Given a network with *m* edges, the probability of an edge between two nodes *i* and *j* with degrees *d_i* and *d_j* respectively is *d_i***d_j* / 2m.



Probability of an edge between nodes 1 and 2 is (3)(3) / (2*15) = 0.30

Strength of a Community, C



For a network with k communities and a total of m edges

Modularity:

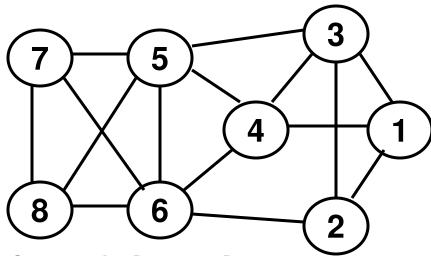
$$Q = \sum_{l=1}^{k} \sum_{i \in C_l, j \in C_l} A_{i,j} - \frac{d_i d_j}{2m}$$

A larger value for Q indicates a good community structure

Modularity Maximization

- The intuition behind the idea of modularity is that a community is a structural element of a network that has been formed in a manner far from a random process.
- If we consider the actual density of links in a community, it should be significantly larger than the density we would expect if the links in the network were formed by a random process.
 - In other words, if two nodes i and j are end up being in the same community, there should be more likely a link between them (i.e., Aij = 1, leading to an overall high value for Q).
 - If i and j end up being in a community such that the chances of having a link between them is just as the same as between any two nodes in the network (i.e., a random network), then the value of Q is more likely to be low (because there could be some Aij = 0 that will bring down the value of Q).

Evaluating Modularity (Example 1)



Community [1, 4, 5, 7] Edges with Aij = 1 Modularity $1 - (3)(4)/(2^{*}15) = 0.60$ 1 - 41 - (4)(5)/(2*15) = 0.334 – 5 5 – 7 1 - (3)(5)/(2*15) = 0.50Edges with Aij = 0 1 – 5 0 - (3)(5)/(2*15) = -0.501 - 70 - (3)(3)/(2*15) = -0.304 - 70 - (4)(3)/(2*15) = -0.40

Total Modularity Score for Community [1, 4, 5, 7]

0.23

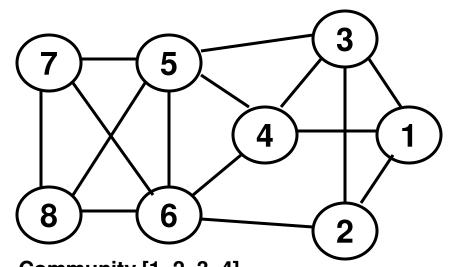
Total Modularity for the two Communities: 0.23 + 0.23 = 0.46

4 3 1
$-\underbrace{2}_{3}$
4
8]
8] Modularity
Modularity
Modularity 1 – (3)(4)/(2*15) = 0.60
Modularity 1 – (3)(4)/(2*15) = 0.60 1 – (3)(5)/(2*15) = 0.50
Modularity 1 – (3)(4)/(2*15) = 0.60 1 – (3)(5)/(2*15) = 0.50
Modularity 1 - (3)(4)/(2*15) = 0.60 1 - (3)(5)/(2*15) = 0.50 1 - (3)(5)/(2*15) = 0.50

Total Modularity Score for Community [2, 3, 6, 8]

0.23

Evaluating Modularity (Example 2)



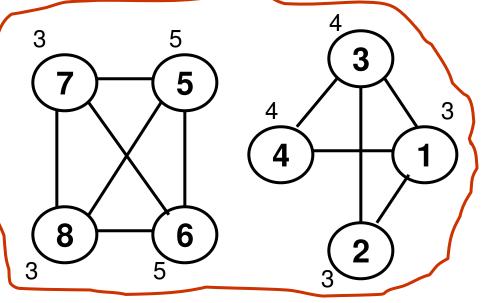
Community [1, 2, 3, 4] Edges with Aij = 1 Modularity

1 – 2	$1 - (3)(3)/(2^{*}15) = 0.70$
1 – 3	$1 - (3)(4)/(2^{*}15) = 0.60$
1 – 4	$1 - (3)(4)/(2^{*}15) = 0.60$
2 – 3	$1 - (3)(3)/(2^{*}15) = 0.70$
3 – 4	$1 - (4)(4)/(2^{15}) = 0.47$
Edges with Aij = 0	
2 – 4	$0 - (3)(4)/(2^{*}15) = -0.40$

Total Modularity Score for Community [1, 2, 3, 4]

2.67

<u>Total Modularity for the two</u> Communities: 2.67 + 2.87 = 5.54



Community [5, 6, 7, 8]		
Edges with Aij = 1	Modularity	
5 – 6	1 – (5)(5)/(2*15) = 0.17	
5 – 7	1 – (3)(5)/(2*15) = 0.50	
5 – 8	1 – (3)(5)/(2*15) = 0.50	
6 – 7	1 – (3)(5)/(2*15) = 0.50	
6 – 8	1 – (3)(5)/(2*15) = 0.50	
7 – 8	1 - (3)(3)/(2*15) = 0.70	

Total Modularity Score for Community [2, 3, 6, 8]

2.87

Using Eigenvectors to Identify Components

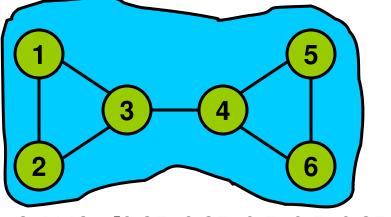
- Compute the Eigenvalues and Eigenvectors of the Adjacency matrix A
- The principal Eigenvector is the one that corresponds to the largest Eigenvalue.
- If all the entries in the "principal Eigenvector" are positive, then it implies that all the nodes are in one component.
 - Else, the vertices with the positive entries are in one component and those with the negative entries are in another component. (Note: 0 is considered positive).
- We apply the above interpretation to all the subsequent Eigenvectors (in the decreasing order of the corresponding Eigenvalues) and identify the smaller communities within the larger components.

Example 1: Eig	envectors to I	dentify
Com	<u>nmunities</u>	
1 3 4 5 A = 6	$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	2 3 3 4 5 6 6

The Eigenvalues in the decreasing order and their corresponding Eigenvectors are:

Eigenvalu	le Eigenvector	All entries are +ve; hence all vertices
2.4142	[0.35; 0.35; 0.5; 0.5; 0.35; 0.35]	are in one single component community
1.7321	[-0.44; -0.44; -0.33; 0.33; 0.44; 0.44	 Vertices 1, 2, 3 form one community Vertices 4, 5, 6 form another comm.
-0.4142	[0.35; 0.35; -0.5; -0.5; 0.35; 0.35]	Within 1-2-3; 3 is in one comm. Within 4-5-6; 4 is in one comm.
-1	[-0.71; 0.71; 0; 0; 0; 0]	
-1	[0; 0; 0; 0; -0.71; 0.71]	
-1.7321	[0.23; 0.23; -0.63; 0.63; -0.23; -0.23	3]

Example 1: Eigenvectors to Identify Communities (1)

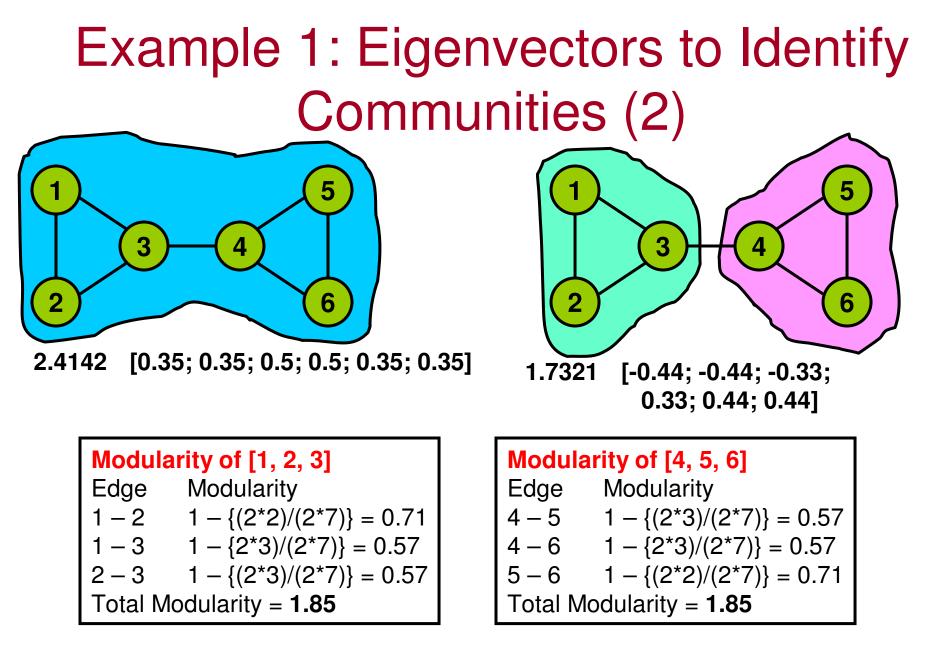


Edge	Modularity
3 – 4	$1 - (3^*3/2^*7) = 0.36$
3 – 5	0 - (2*3/2*7) = -0.43
3 – 6	$0 - (2^{*}3/2^{*}7) = -0.43$
4 – 5	$1 - (2^{*}3/2^{*}7) = 0.57$
4 – 6	1 - (2*3/2*7) = 0.57

 $2.4142 \quad [0.35; \, 0.35; \, 0.5; \, 0.5; \, 0.35; \, 0.35]$

Total Modularity = 0.47

EdgeModularity1-21-(2*2/2*7) = 0.711-31-(2*3/2*7) = 0.571-40-(2*3/2*7) = -0.431-50-(2*2/2*7) = -0.291-60-(2*2/2*7) = -0.292-31-(2*3/2*7) = 0.572-40-(2*3/2*7) = -0.432-50-(2*2/2*7) = -0.292-60-(2*2/2*7) = -0.29



Total Modularity of [1, 2, 3] and [4, 5, 6] = <u>3.7</u>

Example 1: Eigenvectors to Identify Communities (3)

Final Partition

Total Modularity

Score = 3.7

5

6

2.4142 [0.35; 0.35; 0.5; 0.5; 0.35; 0.35]

3

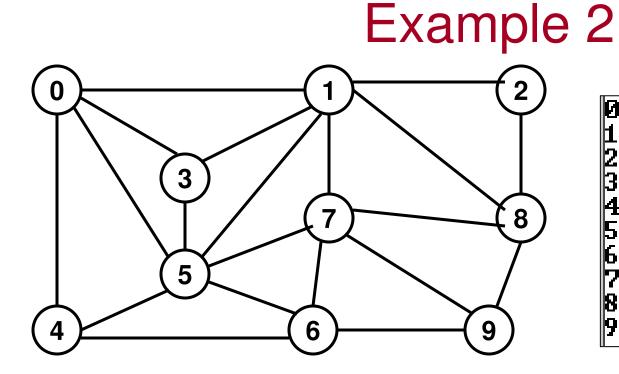
Modularity of [3] and [4] are 0 each Modularity of [1, 2] and [5, 6] are 0.71 each

5

6

Total Modularity of [1, 2] and [3] is 0.71 is less than the modularity of [1, 2, 3]. Hence, We stay with [1, 2, 3] as a community.

Total Modularity of [5, 6[and [4] is 0.71 is less than the modularity of [4, 5, 6]. Hence, We stay with [4, 5, 6] as a community.



Eigenvalue # 9 (2.1175)

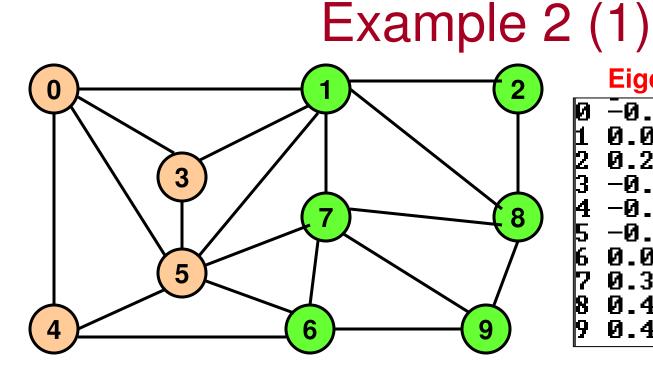
0.41478523082002594
0.0034483071344804475
0.22130808207036753
-0.32465490044157624
-0.28097915591550676
-0.276112087029933
0.09593065477768678
0.33637956450900897
0.4651662798490688
0.42384255258775577

Eigenvalue # 10 (4.3515)

0.3197102709282999 Ø 0.42642009875327136 H 2 3 0.16070352830399387 274067807962952 4 24425677852346567 5 6 0.4464825297669496 0.2966966608945022 0.3817282429845956 8 27288531134487964 0.21861536589094482 9

Eigenvalue # 8 (1.6723)

0	0.11648437679157984
<u>ا</u> ل	0.43051446084677686
2	0.40755819937708554
3	0.2421225400215074
4	-0.3357423606460691
5	-0.1420982832225889
6	-0.5358466992374123
7	-0.15516270796328716
8	0.25104348443917357
9	-0.26309093820736645



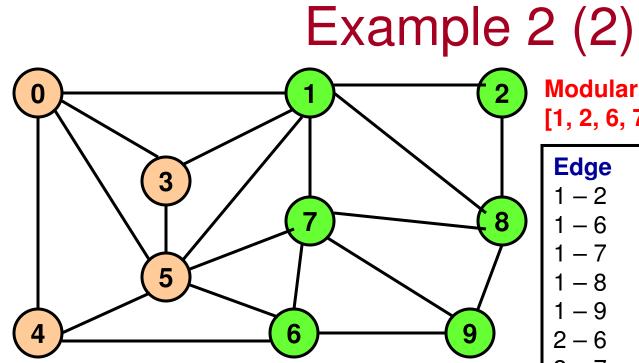
Eigenvalue # 9 (2.1175)

0 -0.41478523082002594
1 0.0034483071344804475
2 0.22130808207036753
3 -0.32465490044157624
4 -0.28097915591550676
5 -0.276112087029933
6 0.09593065477768678
7 0.33637956450900897
8 0.4651662798490688
9 0.42384255258775577

Modularity for Community [0, 3, 4, 5]		
Edge	Modularity	
0 – 3	$1 - \{(4^*3)/(2^*20)\} = 0.7$	
0 – 4	$1 - \{(4^*3)/(2^*20)\} = 0.7$	
0 – 5	$1 - \{(4^*6)/(2^*20)\} = 0.4$	
3 – 4	$0 - \{(3^*3)/(2^*20)\} = -0.23$	
3 – 5	$1 - \{(3^*6)/(2^*20)\} = 0.55$	
4 – 5	$1 - \{(3^*6)/(2^*20)\} = 0.55$	
Total N	lodularity Score for	
[0, 3, 4	, 5] = 2.67	

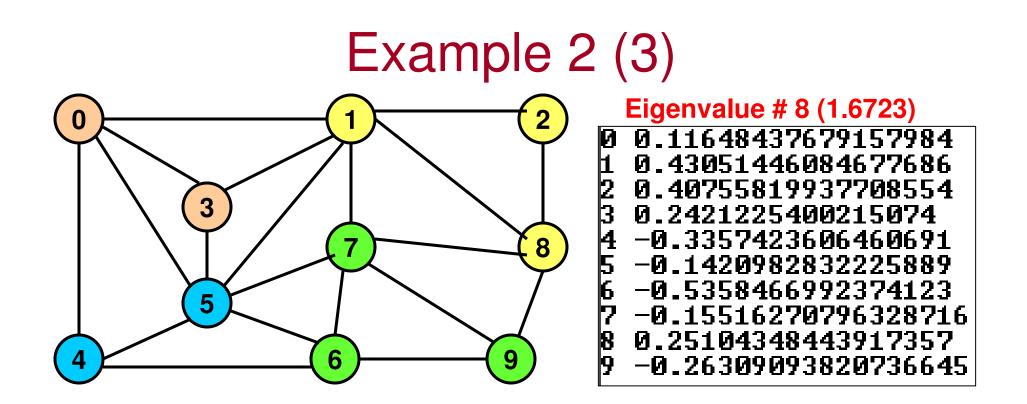
Modularity for Community [1, 2, 6, 7, 8, 9]

Edge	Modularity
1 – 2	$1 - \{(6^{*}2)/(2^{*}20)\} = 0.7$
1 – 6	$0 - \{(6^*4)/(2^*20)\} = -0.6$
1 – 7	$1 - \{(6^*5)/(2^*20)\} = 0.25$
1 – 8	$1 - \{(6^*4)/(2^*20)\} = 0.4$
1 – 9	$0 - \{(6^*3)/(2^*20)\} = -0.45$
2-6	$0 - \{(2^*4)/(2^*20)\} = -0.2$
2-7	$0 - \{(2^*5)/(2^*20)\} = -0.25$
2 – 8	$1 - \{(2^*4)/(2^*20)\} = 0.8$



Modularity for Community [1, 2, 6, 7, 8, 9]

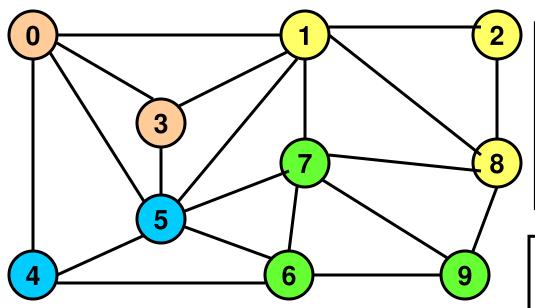
Edge	Modularity
1 – 2	$1 - \{(6^{*}2)/(2^{*}20)\} = 0.7$
1 – 6	$0 - \{(6^*4)/(2^*20)\} = -0.6$
1 – 7	$1 - \{(6^{+}5)/(2^{+}20)\} = 0.25$
1 – 8	$1 - \{(6^*4)/(2^*20)\} = 0.4$
1 – 9	$0 - \{(6^*3)/(2^*20)\} = -0.45$
2-6	$0 - \{(2^*4)/(2^*20)\} = -0.2$
2-7	$0 - \{(2^*5)/(2^*20)\} = -0.25$
2-8	$1 - \{(2^*4)/(2^*20)\} = 0.8$
2-9	$0 - \{(2^*3)/(2^*20)\} = -0.15$
6-7	$1 - \{(4^*5)/(2^*20)\} = 0.5$
6 – 8	$0 - \{(4^*4)/(2^*20)\} = -0.4$
6-9	$1 - \{(4^*3)/(2^*20)\} = 0.7$
7-8	$1 - \{(5^*4)/(2^*20)\} = 0.5$
7 – 9	$1 - \{(5^*3)/(2^*20)\} = 0.63$
8-9	$1 - \{(4^*3)/(2^*20)\} = 0.7$
Total Mc	dularity Score for
[1, 2, 6,	7, 8, 9] = 3.13



Modularity of Community [0, 3] = 0.7Modularity of Community [4, 5] = 0.55Total Modularity of [0, 3] and [4, 5] = 1.25is less than the Modularity of [0, 3, 4, 5]= 2.67 Hence, we stay with community [0, 3, 4, 5]

without further partitioning it.





Modularity of [1, 2, 8] = 1.9Modularity of [6, 7, 9] = 1.83Total Modularity of [1, 2, 8] and [6, 7, 9] = 3.73is larger than the modularity of [1, 2, 6, 7, 8, 9] = 3.13Hence, we allow the partitioning of [1, 2, 6, 7, 8, 9] into [1, 2, 8] and [6, 7, 9] Modularity of Community [1, 2, 8]EdgeModularity1-2 $1 - \{(6^*2)/(2^*20)\} = 0.7$ 1-8 $1 - \{(6^*4)/(2^*20)\} = 0.4$ 2-8 $1 - \{(2^*4)/(2^*20)\} = 0.8$ Total Modularity of [1, 2, 8] = 1.9

Modularity of Community [6, 7, 9]EdgeModularity6-7 $1 - \{(4*5)/(2*20)\} = 0.5$ 6-9 $1 - \{(4*3)/(2*20)\} = 0.7$ 7-9 $1 - \{(5*3)/(2*20)\} = 0.63$ Total Modularity of [6, 7, 9] = **1.83**

Example 2 (5)

