CPTS 591 Elements of Network Science

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# Assignment 2

## **Problem 1**

The fourth graph I chose is called "Zachary's karate club". It is asocial network of friendships between 34 members of a karate club at a US university in the 1970s.

URL: http://www-personal.umich.edu/~mejn/netdata/

### Load 4 real world graphs:

```
"``{r load 4 graphs}
political_graph <- read_graph('Political_blogs.gml',format = 'gml')
neural_graph <- read_graph('Neural_network.gml', format = 'gml')
internet_graph <- read_graph('Internet.gml',format = 'gml')
karate_graph <- read_graph('karate.gml',format = 'gml')</pre>
```

## 1. Political Blogs Graph:

Highest score nodes in (i) Degree:



## (vi) PageRank:



# 2. Neural Network Graph:

### Highest score nodes in (i) Degree:

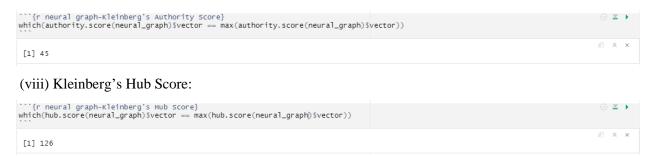


### (v) Katz index:

### (vi) PageRank:

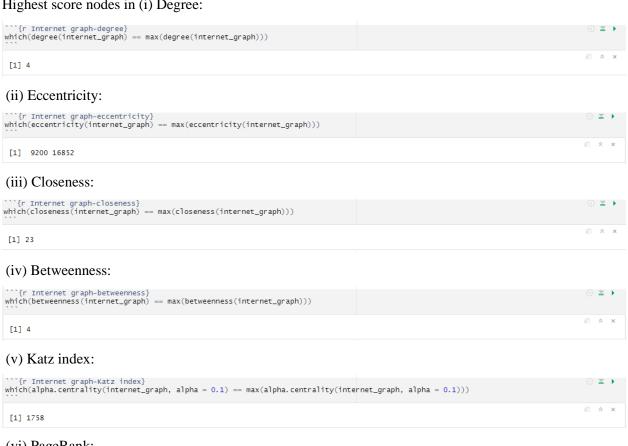


### (vii) Kleinberg's Authority Score:



## 3. Internet Graph:

### Highest score nodes in (i) Degree:



## (vi) PageRank:



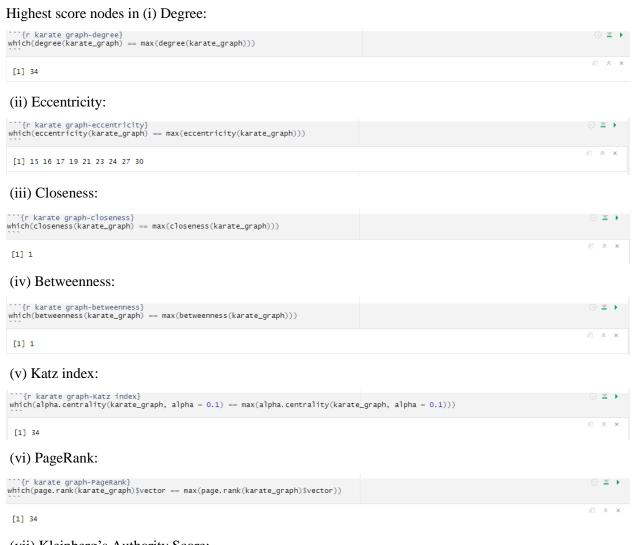
## (vii) Kleinberg's Authority Score:

```{r Internet graph-Kleinberg's Authority Score} which(authority.score(internet_graph)\$vector))	⊕ ≚ ▶
[1] 4	<i>□</i>

## (viii) Kleinberg's Hub Score:

<pre>```{r Internet graph-Kleinberg's Hub Score} which(hub.score(internet_graph)\$vector == max(hub.score(internet_graph)\$vector)) ```</pre>		⊻ 1	•
[1] 4	-	* >	×

# 4. Zachary's Karate Club Graph:



# (vii) Kleinberg's Authority Score:

```{r karate graph-Kleinberg's Authority Score} which(authority.score(karate_graph)\$vector == max(authority.score(karate_graph)\$vector))	※ ▼ ▶
[1] 34	<i>□</i>

## (viii) Kleinberg's Hub Score:

```{r karate graph-Kleinberg's Hub Score} which(hub.score(karate_graph)\$vector == max(hub.score(karate_graph)\$vector))	⊕ <b>×</b> →
[1] 34	<i>□</i>

#### Result analysis:

From the above 4 graphs, we can find there are some cases in which the different centrality measures sometimes identify the same node(s) with the highest score in one graph, in other words, different centrality measures could identify that node(s) as the most important. For example, in neural network graph, the node with the index 45 has the highest score in degree, Katz index, PageRank, and Kleinberg's authority score. In the Internet graph, the node with the index 4 has the highest score in degree, betweenness, PageRank, Kleinberg's authority score, and Kleinberg's hub score. In those cases, we can consider those nodes are the most important. However, it will not happen all the time, for example, in political blogs graph, there is not any node(s) which has most highest scores in all centrality measures, so we cannot say there is any node(s) is the most important in this graph.

### **Problem 2**

Generate 2 random graphs with 20 nodes:

```
**Yer generate 2 random graphs with node 20}
erg1 <- erdos.renyi.game(20, .1, type = "gnp")
E(erg1)
bag1 <- barabasi.game(20, power = 1.0)
E(bag1)

**

+ 19/19 edges from 8299add:
[1] 6-- 8 3-- 9 3--11 2--12 4--12 6--12 2--13 10--13 12--13 3--14 12--16 3--17 5--17 15--17 17--18 2--19 7--19 16--19
[19] 4--20
+ 19/19 edges from 82a87ae:
[1] 2-> 1 3-> 1 4-> 3 5-> 1 6-> 1 7-> 1 8-> 1 9-> 1 10-> 5 11-> 1 12-> 1 13->10 14-> 1 15->10 16->10 17-> 1 18->16 19-> 3
[19] 20-> 4
```

### Generate 2 random graphs with 40 nodes:

I also presented all edges to check whether they have roughly the same number of edges.

Get the largest connected component for each graph:

```
"{r get the largest connected components for each graph}
is.connected(ergi)
is.connected(bag)
is.connected(bag)
is.connected(bag)
decompose.graph(ergl)
decompose.graph(ergl)
v(dergl[[1]])
v(dergl[[2]])
v(dergl[[3]])
v(dergl[[3]])
decompose.graph(erg2)
derg2 <- decompose.graph(erg2)
derg2 <- decompose.graph(erg2)
v(dergl[[3]])
```

1. Compute all eigenvalues corresponding eigenvectors of the Laplacian of the graph:

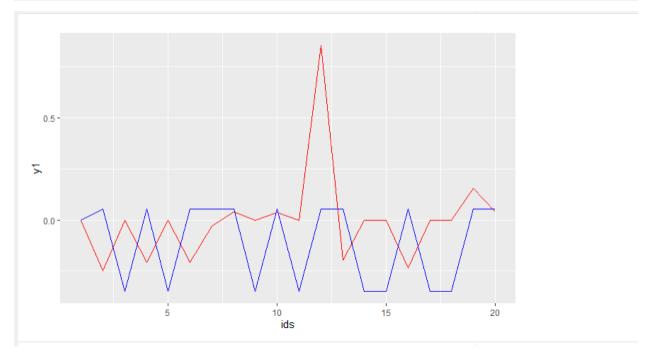
```
eigen1 <- sort(eigen(laplacian_matrix(lcderg1))$values)
eigen2 <- sort(eigen(laplacian_matrix(bag1))$values)
eigen3 <- sort(eigen(laplacian_matrix(lcderg2))$values)
eigen4 <- sort(eigen(laplacian_matrix(bag2))$values)
```

2. Compute and plot the table (For easy observation, all results are reserved with 5 valid numbers):

```
(r 2-2)
options(digits = 5)
Largest_eigenvalue
  Graph
                                 n m dmin dm...
<int> <dbl> <dbl> <dbl>
   ccg
   Second_smallest_eigenvalue
  LCC ER random graph with node 20
                                 11
                                     12
   1
   5
  2.4727
  5
   0.136364
  6.2842
  0.360467
  LCC BA random graph with node 20
  11 1.6364
  6.2842
                                 20
                                    19
   0.000000
  1.000000
  LCC ER random graph with node 40
                                 34
                                     39
  6.1836
   16
   0.048387
  0.024947
  6.2842
  LCC BA random graph with node 40
                                 40
                                     39
  2.0303
   5
   0.000000
  1.000000
  6.2842
 4 rows | 1-10 of 13 columns
```

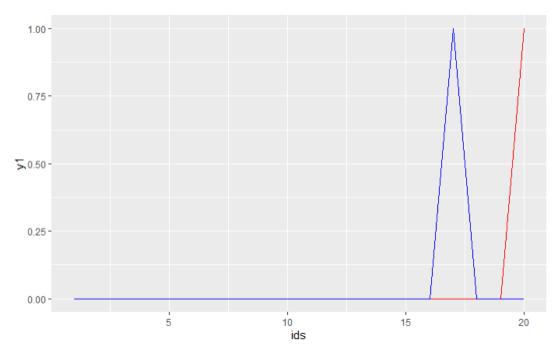
3. Plot graph for first ER model. Red line represents the eigenvectors corresponding to the largest eigenvalue, blue line represents the eigenvectors corresponding to the second smallest eigenvalue.

```
"``{r 2-3(1)}
ids <- array(1:20)
y1 <- eigen(laplacian_matrix(erg1))$vector[,1]
y2 <- eigen(laplacian_matrix(erg1))$vector[,19]
df <- data.frame(ids, y1,y2)
ggplot(df,aes(ids)) +
    geom_line(aes(y=y1),color = "red") +
    geom_line(aes(y=y2),color = "blue")</pre>
```



Plot graph for BA model. Red line represents the eigenvectors corresponding to the largest eigenvalue, blue line represents the eigenvectors corresponding to the second smallest eigenvalue.





#### Observation:

In the ER model, the eigenvectors are totally different when we choose the largest eigenvalue and the second smallest eigenvalue. However, in the BA model, most eigenvectors are 0, few eigenvectors are 1, when we correspond to different eigenvalues, the eigenvectors are almost the same.

## **Problem 3**

- 1. PageRank in biology & bioinformatics: Gene Rank, Protein Rank, Iso Rank. I personally very interested in biology. I think it is a mysterious field. I did not expect that PageRank could be applied to this field. All advanced theories or algorithms might help human-beings achieve the truth of our body closer.
- 2. PageRank in social networks: Buddy Rank, Twitter Rank. Take a simple example, such as the recommendation system on Facebook, which provides us with a lot of data to recommend relevant people. It can find many interesting connections between things through data.

3. PageRank in literature: Book Rank. Online libraries become more and more popular, so Book Rank becomes necessary in our life. It is very useful when we are learning new knowledge or finding references. I can easily find related articles and books.