ARS: Analisi Reti Sociali

01/06/2017

Name	
Student ID	

Note: Whenever an exercise requires the application of a formula/algorithm both the methodology and the solution must be *reported* and *discussed*.

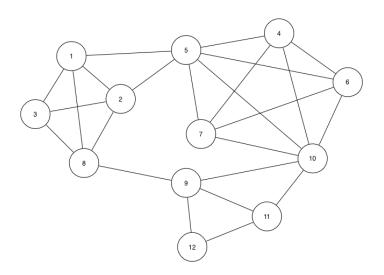


Figure 1:

Exercise 1: Community Evaluation [6 points]

Given the Graph $\mathcal G$ shown in Figure 1 compare the following partitions:

$$P1 = [(1, 2, 3, 5), (4, 6, 7, 10), (8, 9, 11, 12)]$$

$$\mathbf{P2} = [(2, 3, 8), (1, 5, 4, 7), (6, 9, 10, 11, 12)]$$

$$\mathbf{P3} = [(1, 2, 3, 8), (4, 5, 6, 7, 10), (9, 11, 12)]$$

Which partition is the best with respect to the *modularity* score? and for *conductance*?

Solution.

Modularity:

P1:
$$wP1 = min(mod(C1), mod(C2), mod(C3)) = mod(C3)$$

- $mod(C1) = (\frac{4}{24} - \frac{17}{2*24})^2 \simeq 0.035$

$$- \operatorname{mod}(C2) = \left(\frac{4}{24} - \frac{18}{2*24}\right)^2 \simeq 0.043$$

$$- \operatorname{mod}(C3) = (\frac{4}{24} - \frac{13}{2*24})^2 \simeq \mathbf{0.011}$$

P2: wP2 = min(mod(C1), mod(C2), mod(C3)) = mod(C1)

$$- \mod(C1) = (\frac{3}{24} - \frac{11}{2*24})^2 \simeq \mathbf{0.011}$$

$$- \operatorname{mod}(C2) = \left(\frac{4}{24} - \frac{18}{2*24}\right)^2 \simeq 0.043$$

$$- \operatorname{mod}(C3) = \left(\frac{5}{24} - \frac{19}{2*24}\right)^2 \simeq 0.035$$

P3: wP3 = min(mod(C1), mod(C2), mod(C3)) = mod(C3)

$$- \operatorname{mod}(C1) = (\frac{4}{24} - \frac{15}{2*24})^2 \simeq 0.021$$

$$- \operatorname{mod}(C1) = \left(\frac{4}{24} - \frac{15}{2*24}\right)^2 \simeq 0,021$$
$$- \operatorname{mod}(C2) = \left(\frac{5}{24} - \frac{24}{2*24}\right)^2 \simeq 0,085$$

$$- \operatorname{mod}(C3) = (\frac{3}{24} - \frac{9}{2*24})^2 \simeq \mathbf{0,003}$$

Best partitions w.r.t. modularity: P1, P2 max(wP1, wP2, wP3) = wP1 = wP2

Conductance:

P1: wP1 = max(cond(C1), cond(C2), cond(C3)) = cond(C1)

$$- \text{ cond(C1)} = \frac{2*7}{2*5+7} \simeq \mathbf{0.823}$$

$$- \operatorname{cond}(C2) = \frac{2*6}{2*8+6} \simeq 0.545$$

$$- \text{ cond(C3)} = \frac{2*5}{2*4+5} \simeq 0.769$$

P2: wP2 = max(cond(C1), cond(C2), cond(C3)) = cond(C2)

$$- \operatorname{cond}(C1) = \frac{2*5}{2*3+5} \simeq 0,909$$

$$- \operatorname{cond}(C2) = \frac{2*10}{2*4+10} \simeq 1,111$$

$$- \text{ cond(C3)} = \frac{2*7}{2*6+7} \simeq 0,736$$

P3: wP3 = max(cond(C1), cond(C2), cond(C3)) = cond(C3)

$$- \text{ cond(C1)} = \frac{2*3}{2*6+3} \simeq 0.4$$

$$- \operatorname{cond}(C2) = \frac{2*4}{2*10+4} \simeq 0.333$$

$$- \text{ cond(C3)} = \frac{2*3}{2*3+3} \simeq 0,666$$

Best partition w.r.t. conductance: P3 min(wP1, wP2, wP3) = wP3

Exercise 2: Community Discovery [6 points]

Given the Graph $\mathcal G$ shown in Figure 1 compute its communities applying k-clique for k=3,4,5.

Solution. The communities identified by k-clique are:

 $k=3: \{1, 2, 3, 5, 8\}, \{4, 5, 6, 7, 10\}, \{9, 10, 11, 12\}$

 $\mathbf{k=4}: \{1, 2, 3, 8\}, \{4, 5, 6, 7, 10\}$

 $k=5: \{4, 5, 6, 7, 10\}$

Exercise 3: Threshold Model [8 points]

Given the graph $\mathcal G$ shown in Figure 1 apply the Threshold model considering the following two scenarios:

- S1 Set of initial infected nodes: $I = \{5\}$;
 - Node threshold τ : 1/4 for even nodes, 2/3 for odd nodes.
- S2 Set of initial infected nodes: $I = \{6, 9\}$;
 - Node threshold τ : 1/4
 - Set of blocked nodes (i.e. nodes that are not allowed to change their status): $B = \{10\}$

Consider a node infected at time t iff at least $\tau\%$ of its neighbors were already infected at time t-1.

Solution. The nodes infected in the two scenarios during each iteration are:

S1

 $I_0: \{5\}$

 $I_1:\{2,4,6\}$

 $I_2: \{7, 8, 10\}$

 $I_3:\{1,3\}$

S2

 $I_0: \{6, 9\}$

 $I_1: \{4, 7, 8, 11, 12\}$

 $I_2:\{1,2,3,5\}$

Exercise 4: Resilience [6 points]

Given the graph \mathcal{G} shown in Figure 1 remove 6 nodes, one at a time, applying the following selection criteria:

- random selection;
- decreasing degree centrality.

Compare how each removal impacts on the graph structure in terms of number of connected components and average component size. Specify the node removed - and obtained partition - at each iteration: if multiple nodes generate the same score solve the tie removing the one with lowest identifier.

Solution.

Decreasing Degree Centrality

```
it_1: node 5, degree 6, components 1, avg. size 11 it_2: node 10, degree 5, components 2, avg. size 5 it_3: node 8, degree 4, components 3, avg. size 3.3 it_4: node 1, degree 2, components 3, avg. size 2.6 it_5: node 4, degree 2, components 3, avg. size 2.3 it_6: node 9, degree 2, components 3, avg. size 2
```

Random

```
\begin{array}{l} it_1 : \text{node } 9, \text{ degree } 4, \text{ components } 1, \text{ avg. size } 11 \\ it_2 : \text{node } 12, \text{ degree } 1, \text{ components } 1, \text{ avg. size } 10 \\ it_3 : \text{node } 6, \text{ degree } 4, \text{ components } 1, \text{ avg. size } 9 \\ it_4 : \text{node } 4, \text{ degree } 3, \text{ components } 1, \text{ avg. size } 8 \\ it_5 : \text{node } 1, \text{ degree } 4, \text{ components } 1, \text{ avg. size } 7 \\ it_6 : \text{node } 5, \text{ degree } 3, \text{ components } 2, \text{ avg. size } 2.5 \\ \end{array}
```

Exercise 5: Multiple-choice Questions [4 points]

Identify the correct answer(s), among the ones proposed, for to the following questions:

- **Q1:** Which of the following Community Discovery methods produce non-overlapping communities?
 - (A) Girvan-Newman
 - (B) DEMON
 - (C) k-Clique
 - (D) Greedy Modularity/Louvain
- **Q2:** In which of the following network topologies we define a vanishing epidemic threshold?
 - (A) Random network
 - (B) Scale-free network
 - (C) Regular grid network
 - (D) Small-world network
- **Q3:** Is a scale-free network more resilient to *random* node failures than a random graph?
 - (A) Yes
 - (B) No
- **Q4:** Is a scale-free network more resilient to *targeted* node removal than a random graph?
 - (A) Yes
 - **(B)** No

Exercise 6: Open Question [2 points]

Consider the SIS epidemic model with infection rate β and recovery rate μ : which are the possible equilibrium the model can reach? in which condition the epidemic dies out?

Solution. The epidemic dies out for $\lambda = \frac{\beta}{\mu} < 1$. λ is the reproductive number: it identifies the average number of infectious individuals generated by a single infected node in a fully susceptible population.

The equilibrium states are then:

- Die out: $\lambda < 1$, as described before.
- Outbreak: $\lambda > 1$. In this scenario the recovery process is slower than disease spreading.

Once reached the stationary state the fraction of infected nodes is $1 - \frac{\mu}{\beta}$