
Model Mini-Test - Network Science 2021/2022 (CC4063 & CC4095)

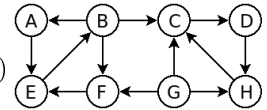
Duration: 1h30m+30m

DCC/FCUP

Answer to only **five** of the following six groups

Essential Concepts (20%)

1 - **Directed Networks.** Consider the right figure, representing a simple directed network:



- What is the **distance** from node C to node E?
- Identify all **strongly connected components** (SCCs) (and their correspondent nodes)
- Redraw the network as a **DAG** (directed acyclic graph) of SCCs.

2 - **Bipartite Networks.**

- Give an example of a **real world bipartite network**, identifying what are the nodes and the edges.
- What is a **bipartite network projection**? Give an example projection of the bipartite network you chose.

Graph Models (20%)

3 - **Erdős-Rényi $G_{(n,p)}$ model (ER).** Imagine you generate an ER graph with $n = 1000$ and $p = 0.01$.

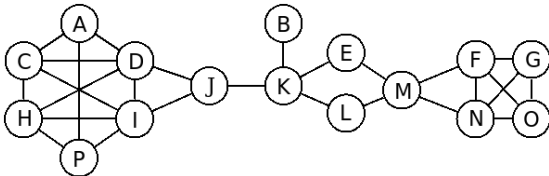
- What is the **average degree** of its nodes?
- Draw a plot showing what you expect to be its **degree distribution**.
- How many **connected components** do you expect the graph to have? Why?
- If you increase the amount of nodes to 1 million (keeping the same average degree), what happens to the **average path length**? Does it stay the same? decreases? increases? by how much?

4 - **Small-World Networks.** Imagine you generate a small world network using the Watts-Strogatz (WS) model.

- Which model generates a network with higher **clustering coefficient**: ER or WS? Why?
- A purely regular network can have an high **average path length**. Explain how the WS model lowers this value.

Node Centrality (20%)

5 - **Centrality Metrics.** Consider the following network, and indicate:



- a node with high degree but low betweenness centrality
- a node with high closeness centrality but low betweenness centrality
- a node with high betweenness centrality, high closeness centrality and high degree

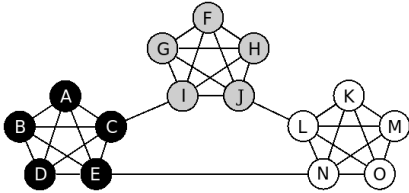
6 - **The PageRank Algorithm.**

- Problems.** Explain what a **spider trap** is, why it might be a problem for the PageRank algorithm and how we can avoid that problem.
- Customization.** Explain what is a **Personalized PageRank**, what is different from the normal PageRank and in what kind of situation it should be used.

Community Structure (20%)

7 - **Girvan-Newman and Louvain algorithms.** Consider these two algorithms for community discovery.

- Classify both algorithms in terms of being **divisive or agglomerative**.
- Which one is faster in terms of **execution time**?
- Consider the following network with three very natural and intuitive communities. Explain how one of the two algorithms would work on this network, by explaining a couple of its initial **greedy iterations**.



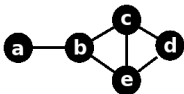
8 - **Modularity.**

- Give a brief informal description of the **concept of modularity**, and what can we use it for.

Subgraph Patterns (20%)

9 - **Subgraph census.**

- Calculate the **frequency of all possible induced subgraphs of size 3** on graph of the following figure (draw the topology of each possible subgraph and indicate all occurrences as sets of nodes).



- Explain the relationship between the concepts of **network motifs** and **subgraph frequency**.

10 - **G-Tries.**

- Explain what **G-Trie** is. Explain what does it store, draw an example, indicate in what kind of task it might be useful and give a brief explanation of the intuition on why it is efficient.

Diffusion and Cascading Behavior (20%)

11 - **Spreading in Random Trees.** Consider a simple epidemic model based on random trees: a patient meets d people and infects each one with probability q .

- Suppose you start with a **single infected patient** and that $d = 200$ and $p = 1\%$. What is the expected number of infected persons at depth level 1 and 2? Will the epidemic run forever or will it stop?
- For a general case, what is the **reproductive number**? How can we use it to determine if a disease will spread or die out? Does it matter the amount of people that initially have the disease?

12 - **Spreading models of viruses.**

- Describe the **SIS model**: indicate states, transition probabilities and dynamics on a complete graph.
- Give an **example of a disease** that fits the SIS model.