Medical Data Science, WS 2023/2024

Prof. Dr. Nico Pfeifer Chair for Methods in Medical Informatics University of Tuebingen



2023-12-04

Assignment 4

Deadline: Tuesday, December 19, 6:00 p.m.

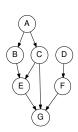
This problem set is worth 50 points. You can submit in groups of two people. Submit your solutions digitally by uploading to the ILIAS page. Just upload a zipped folder containing all necessary files and name the folder by your last names. The folder should be named according to the following scheme:

[MDS] [Assignment4]_lastname1_lastname2

Problem 1 (T, 11 Points)

Graphical models and Causality.

- (a) (3P) What makes a model (such as a Bayesian Network) generative? How can dependencies be taken into account in such networks?
- (b) (3P) True or false? Explain briefly or give counter examples:
 - a) If there are many paths between two nodes we always have to test every single path to say whether the two nodes are d-separated.
 - b) If A is d-separated from B, B is d-separated from A.
 - c) If A is d-separated from B and B is d-separated from C, A is d-separated from C.
- (c) (4P) Consider the following DAG G.



Can you show conditional independence with the help of d-separation for the following examples? Keep in mind the results from (c) and write down how many paths you need to test to show independence.

- \bullet $a \perp \!\!\! \perp B$
- $A \perp \!\!\!\perp G \mid C, E, B$
- $D \perp \!\!\! \perp C \mid E, G$
- $G \perp \!\!\! \perp B \mid A, E$
- (d) (1P) What's the difference between P(A|B) and P(A|do(B))?

Problem 2 (P, 18 Points)

Evaluate the performance of different graph kernels using GraKeL (in case you have problems installing GraKel with python 3.8 or larger, please try python 3.7): link to website and the MUTAG data set from here: website with benchmark data sets.

• Compute the graphlet kernel using sampling for the graphlets of size 3 (1000 samples). Perform a 10-fold cross-validation for the binary classification problem using the kernel with an SVM. What is the accuracy for the best *C* ∈ {0.0001, 0.001, 0.01, 0.1, 1, 10}? You can expect sth. larger than 0.8. Evaluating the accuracy for the best C value is sufficient, you do not have to evaluate the test accuracy separately.

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How many samples would you need such that the deviation from the real distribution is less than 0.05 with probability larger than 0.9?

- Compute the Weisfeiler-Lehman subtree kernel for 4 iterations. Perform a 10-fold cross-validation for the binary classification problem using the kernel with an SVM. What is the accuracy for the best *C* ∈ {0.0001, 0.001, 0.01, 0.1, 1, 10}? You can expect sth. larger than 0.85. Evaluating the accuracy for the best C value is sufficient, you do not have to evaluate the test accuracy separately.
- Compute the Weisfeiler-Lehman edge kernel (WLedge.m) for 1 iteration. Perform a 10-fold cross-validation for the binary classification problem using the kernel with an SVM. What is the accuracy for the best $C \in \{0.0001, 0.001, 0.01, 0.1, 1, 10, 100\}$?

Problem 3 (P/T, 21 Points)

Consider the Similarity Network Fusion (SNF) method with number of neighbors k = 2 (remember that the first neighbor of a node is the node itself). Note: if you provide runnable code for a) and b), you can use c) to solve a) and b).

(a) (7P) Given matrices

$$\mathbf{W}^{(1)} = \begin{pmatrix} 1.00 & 0.50 & 0.30 & 0.10 & 0.10 \\ 0.50 & 1.00 & 0.40 & 0.10 & 0.10 \\ 0.30 & 0.40 & 1.00 & 0.30 & 0.30 \\ 0.10 & 0.10 & 0.30 & 1.00 & 0.50 \\ 0.10 & 0.10 & 0.30 & 0.50 & 1.00 \end{pmatrix}$$

and

$$\mathbf{W}^{(2)} = \begin{pmatrix} 1.00 & 0.20 & 0.50 & 0.10 & 0.10 \\ 0.20 & 1.00 & 0.30 & 0.10 & 0.10 \\ 0.50 & 0.30 & 1.00 & 0.30 & 0.30 \\ 0.10 & 0.10 & 0.30 & 1.00 & 0.50 \\ 0.10 & 0.10 & 0.30 & 0.50 & 1.00 \end{pmatrix}$$

provide $P^{(1)}$, $P^{(2)}$, $S^{(1)}$, and $S^{(2)}$.

- (b) (7P)Perform two steps of the similarity network fusion method (i.e., compute $\mathbf{P}_1^{(1)}$, $\mathbf{P}_1^{(2)}$, $\mathbf{P}_2^{(1)}$, and $\mathbf{P}_2^{(2)}$ as well as the corresponding $\mathbf{P}^{(c)}$ s).
- (c) (7P) Implement the SNF starting from the similarity matrices $\mathbf{W}^{(i)}$ in Python with the convergence criterion ϵ as described in the supplement of the paper (Uni Tuebingen VPN necessary to access the full paper, supplement is freely available) and check whether the graph structure of the $\mathbf{P}^{(c)}$ s changes for t > 2 for the above described data.