## Problem Set 3 6.S091: Causality IAP 2023

Due: Friday, February 3rd at 11:59pm EST, by email (csquires@mit.edu)

- Problem sets **must** be done in LaTeX.
- You may use any programming language for your solutions, and you are not required to turn in your code

Note: while there are a total of 15 points on this pset, 10 will still be considered a perfect score. Anything above 10 will be "bonus" points and will count toward the 18 points needed to pass.

# Problem 1: Constructing Minimal I-MAPs [5 points]

You should use the partial-correlation based conditional independence test which you coded in Problem Set 2. An implementation of the necessary functions can be found at

https://github.com/csquires/6.S091-causality/blob/main/psets/pset3/utils.py

(a) Write a function

minimal\_map(samples, permutation, alpha)

which constructs the graph  $\mathcal{G}_{\pi}$  for the permutation  $\pi$ , using the partial-correlation based conditional independence test with significance level alpha.

Samples of  $(X_1, X_2, X_3, X_4, X_5)$  are in the file imap\_samples.csv at

https://github.com/csquires/6.S091-causality/blob/main/psets/pset3/imap\_samples.csv

Run minimal\_imap(imap\_samples, permutation, 0.05) for the following permutations:

$$\pi_a = [1, 2, 3, 4, 5]$$
  

$$\pi_c = [5, 4, 1, 2, 3]$$
  

$$\pi_b = [5, 4, 3, 2, 1]$$

The graph  $\mathcal{G}_{\pi_a}$  should have 4 edges. Draw the graphs for each permutation  $\pi_a$ ,  $\pi_b$ , and  $\pi_c$ .

(b) Recall that a Chickering sequence from  $\mathcal{G}$  to  $\mathcal{H}$  is a sequence of DAGs, where each consecutive pair in the sequence is related by a covered edge reversal or an edge addition. Construct a Chickering sequence from  $\mathcal{G}_{\pi_a}$  to  $\mathcal{G}_{\pi_b}$ . You can do this problem by hand, you do not need to write a function.

## Problem 2: d-separation defines a graphoid [5 points]

In this problem, you will show that two of the graphoid properties hold for d-separation. Throughout the problem, let  $\mathbf{A}, \mathbf{B}_1, \mathbf{B}_2, \mathbf{S} \subseteq \mathcal{X}$  be disjoint subsets of nodes in a DAG  $\mathcal{G}$ , and let  $\mathbf{B} = \mathbf{B}_1 \cup \mathbf{B}_2$ .

- (a) Weak Union. Suppose there is a d-connecting path  $\gamma$  from  $A \in \mathbf{A}$  to  $B \in \mathbf{B}_1$  given  $\mathbf{S} \cup \mathbf{B}_2$ . Show that there is a d-connecting path  $\gamma'$  from some  $A' \in \mathbf{A}$  to some  $B' \in \mathbf{B}$  given  $\mathbf{S}$ .
- (b) Contraction. Suppose there is a d-connecting path  $\gamma$  from  $A \in \mathbf{A}$  to  $B \in \mathbf{B}$  given S. Show that either:
  - There is a d-connecting path  $\gamma'$  from some  $A' \in \mathbf{A}$  to some  $B' \in \mathbf{B}_2$  given  $\mathbf{S}$ , or
  - There is a d-connecting path  $\gamma'$  from some  $A' \in \mathbf{A}$  to some  $B' \in \mathbf{B}_1$  given  $\mathbf{S} \cup \mathbf{B}_2$ .

### Problem 3: Searching an Equivalence Class [5 points]

The folder

https://github.com/csquires/6.S091-causality/tree/main/psets/pset3/mec\_examples contains pairs of the form

starting\_dag1.csv ending\_dag1.csv

where the starting DAG  $\mathcal{G}$  is Markov equivalent to the ending DAG  $\mathcal{H}$ . Each file represents a DAG in adjacency matrix form, i.e., each file stores a matrix A where  $A_{ij} = 1$  if and only if the associated DAG  $\mathcal{G}$  has an edge  $i \to j$ . Starter code for loading DAGs can be found at

https://github.com/csquires/6.S091-causality/blob/main/psets/pset3/search\_mec.py

#### (a) Write a function

#### covered\_edge\_neighbors(dag)

which, given a DAG  $\mathcal{G}$ , returns all DAGs  $\mathcal{G}'$  which differ from  $\mathcal{G}$  by exactly one covered edge reversal. starting\_dag1 should have 5 neighbors. How many neighbors do starting\_dag2 and starting\_dag3 have?

### (b) Write a function

### search\_mec(starting\_dag, ending\_dag)

which takes two Markov equivalent DAGs  $\mathcal{G}$  and  $\mathcal{H}$  and returns a sequence of covered edge reversals from  $\mathcal{G}$  to  $\mathcal{H}$ .

The sequence of covered edges can be found by breadth-first search using covered\_edge\_neighbors. For breadth-first search, you might need to maintain a set of already-visited states. In Python, I recommend using a frozenset containing the edges of a DAG to represent each state, which is hashable and hence can be used in a set or as keys of a dictionary.

The shortest path from starting\_dag1 to ending\_dag1 should be of length 6 (where the length of the path is the number of covered edge flips). How long is the shortest path from starting\_dag2 to ending\_dag2? How long is the shortest path from starting\_dag3 to ending\_dag3?