

# causaloptim

# Introduction to the package

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### Crash course in causal inference

The causal roadmap:

- 1. Translate the scientific question of interest into a formal causal parameter optional: define the ideal study
- 2. Specify a model for the generating mechanism of the observed data, with the causal parameter in mind i.e., a directed acyclic graph (DAG) and some other assumptions
- 3. State the identifying assumptions, and, derive the statistical parameter
- 4. Estimate it and do inference

Point 3: under what conditions can the observed data narrow down the causal parameter to a single value?

Sometimes, under the most reasonable set of assumptions, not even an infinite amount of data can narrow down the causal parameter to a single point.

#### A causal parameter

Suppose we have a new treatment that we think will prolong life. How do we measure its efficacy?

I would say let X represent the treatment indicator, 0 for the standard of care, and 1 for the new treatment.

Let Y represent a bad outcome, 0 for alive at 3 years, 1 if dead within 3 years. Y(X = 1) denotes the outcome if X were intervened upon to have value 1, called a potential outcome. Y(X = 0) also exists.

One way to measure the efficacy of X on Y is with the causal risk difference/average treatment effect:

$$p\{Y(X=1)=1\} - p\{Y(X=0)=1\} := \theta.$$

#### DAGs and assumptions - random treatment assignment



This is a functional causal model  $\{F_V : pa(V) \rightarrow V \mid V \in V\}$  for the measured variables  $\mathcal{V} = \{C, X, Y\}$ 

Here we assume that no variable influences X, so the observation Y|X = x coincides with the potential outcome Y(X = x) for  $X \in \{0, 1\}$ . Thus we can estimate  $\theta$  easily using the observed data, i.e., it is *identifiable*.

# DAGs and assumptions - confounding



Here,  $Y = F_Y(X, C)$  and  $X = F_X(C)$ , there is confounding. We can still estimate  $\theta$  easily using the observed data, it is just a little more complicated, we have to adjust for C.

# DAGs and assumptions - unmeasured confounding



Here,  $Y = F_Y(X, U)$  and  $X = F_X(U)$ , there is confounding, but we are in trouble because U is not measured (as indicated by the dashed circle). Not even knowing the true probabilities  $\mathbf{p} = \{p(Y = y, X = x) \text{ for } y, x \in \{0, 1\}\}$  suffices

to point identify  $\theta$ . Instead, we aim to get an upper and lower bound for  $\theta$ :

$$L(\boldsymbol{p}) \leq \theta \leq U(\boldsymbol{p}).$$

This is called partial identification, or nonparametric causal bounding.

### What does causaloptim do?

- Provides a framework for deriving  $L(\mathbf{p}), U(\mathbf{p})$  in specific scenarios.
- Users specify a DAG (and/or other assumptions), and a parameter of interest
- They get out  $L(\mathbf{p}), U(\mathbf{p})$  for that parameter under that DAG.
- These are *symbolic* nonparametric bounds: expressions in terms of estimable probabilities, not specific numbers.
- The method only works under certain constraints on the DAG and the parameter of interest.



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# How does it work?

- Graph is split into left and right sides
- Observed variables are categorical, unobserved can be anything
- Complete confounding within each side but paths connecting sides are unconfounded
- Assume that we observe p{right vars|left vars} for all levels of the variables



# How does it work? (2)



The graph is translated to the equivalent response function variable formulation. Since the observable variables are discrete, all possible response patterns can be enumerated.

# How does it work? (3)

Example response functions in the graph  $Z \rightarrow X; Z \leftarrow U \rightarrow X$ :

<b>Response Pattern</b>	Function	Interpretation
r = 0	$f_X(z,0)=0$	never takes $X$ , regardless of $Z$
r = 1	$f_X(z,1) = z$	full compliance with assignment $Z$
r = 2	$f_X(z,2) = 1 - z$	total defiance of assignment $Z$
r = 3	$f_X(z,3) = 1$	always takes $X$ , regardless of $Z$

We define a vector of unobservable parameters  $\boldsymbol{q}$  each element of which represents the probability of a particular response pattern.

# How does it work? (4)

for p the observable probabilities and q the unobservable response function probabilities, we have a linear system of equations

 $\boldsymbol{p} = R\boldsymbol{q}$  for some matrix  $R \in \{0, 1\}^{m \times n}$ .

Algorithm 1 of Sachs et al. [2023] gives a way to find the matrix R. That paper also shows that these linear constraints are complete for all graphs that meet our criteria.

# How does it work? (5)

Further, we can express potential outcomes of Y under intervention on X in terms of the parameters q by the adjustment formula;

$$P(Y(X = x) = y) = \sum_{r_X, r_Y} P(y|x, r_Y) q_{(r_X, r_Y)}$$

Hence, we have

$$\nu = P(Y(X = 1) = 1) - P(Y(X = 0) = 1)$$
  
=  $c^T q$  for some vector  $c \in \{0, 1, -1\}^n$ 

Algorithm 2 of Sachs et al. 2022 gives a way to find the c vector and describes conditions under which it is linear.

# Linear Programming

Now we have a set of linear constraints as well as our effect of interest in terms of q and we are ready to optimize. The following LP gives a tight lower bound on  $\nu$ :

$$\begin{array}{ll} \min & c^{T}q\\ st & \Sigma q = 1\\ \& & Rq = p\\ \& & q \ge 0 \end{array}$$

This "primal" problem is stated in terms of the q s, which are not estimable. We want a solution that gives us an expression in terms of the observable p s, thus we convert to the dual problem.

#### Linear Programming continued

- By *the Strong Duality Theorem* of convex optimization, the optimal value of this primal problem equals that of its dual.
- Furthermore, its constraint space is a convex polytope and by *the Fundamental Theorem of Linear Programming*, this optimum is attained at one of its vertices.
- Thus, we enumerate the vertices of the dual problem which are in terms of the *p* s. We use the *Double description method* as implemented in cddlib (very fast!)

The vertices correspond to the expressions given in the min/max output of the bounds.

# About the package development

- Came out of a close collaboration with Arvid Sjölander and Erin Gabriel since 2017 when we were all at Karolinska in Sweden
- Needed help getting Balke's C++ code from 1996 running again
- First released on Github in 2019 and on CRAN in 2020
- We've used it ourselves and "hacked" it to do some additional things
- Some of these hacks are now features in an unreleased version that will be close to a 1.0.0 release (stable API).

Some examples of what we've used it for: [Gabriel et al., 2021, 2022, 2023, 2024b,a]

It also includes Balke's original code from his thesis, which was previously unreleased

Some of the new hacks/features:

- Causal model object creation and tools
- $\rightarrow$  E.g., sampling from model, deriving observable constraints, testing linearity, testing observable constraints
  - Flexibility in specifying causal model and observables
  - Testing linearity of causal effect

# Observable constraints

- $\boldsymbol{p} = R \boldsymbol{q}$  relates the observables to the response pattern probabilities.
  - Geometrically, it also describes a convex polytope by the locations of its vertices (the V-representation)
  - These vertices are related to the bounds.

Another view of this same polytope gives us *observable constraints* that can be used to falsify the causal model. These are called the instrumental inequalities in the IV model.

- The other view is the H-representation, which describes the polytope in terms of its faces.
- These faces are described by inequalities in terms of the observable probabilities.

The was described by Bonet [2001] and we have implemented it in the causal model object creation.



# New features demo

# The future of causaloptim

- Plans to enhance flexibility and applicability
- Extend to non-linear cases tricks for symbolic optimization, and numeric optimization in other cases
- Inference on the bounds (confidence intervals)
- Integrate into a complete causal pipeline

This is thanks to the creation of the new **Pioneer Center for SMART Biomed**, locally headed by Erin Gabriel. It specifically includes funding for the development of high-quality software for research on common complex diseases.

There are lots of opportunities to get involved. See more: https://smartbiomed.dk/



# What have you used causaloptim for? Feature wishes? Would you use it now if you haven't before? What's the best wine from your region?



#### References I

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### References II

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