What range could your causal effect lie between if the instrumental variable assumptions held?

## Find out with our bpbounds R package and Shiny app!

# bpbounds: R package and web app

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## Introduction

- We present our bpbounds R package and Shiny web app for the nonparametric bounds for the average causal effect (ACE) due to Balke and Pearl (Palmer et al. 2018).
- This is an R implementation of our Stata programs (Palmer et al. 2011).
- The package can be installed from CRAN as follows:

install.packages("bpbounds")

• Code development is on the GitHub repository: https://github.com/remlapmot/bpbounds

## Methods

- Under the instrumental variable assumptions alone, without additional parametric model assumptions, the ACE is not identified.
- Balke and Pearl (1997) showed it is possible to derive bounds for the ACE.
- The bounds have the following interpretation:

There is some joint distribution of the unobserved confounders and the observed variables that yields a true ACE as small as the lower bound, while another choice produces an ACE as large as the upper bounds (the bounds are tight).

- There are at least two ways to implement the Balke-Pearl bounds:
  - i. using conditional probabilities calculated from
- contingency tables; ii. the polytope method due to **Dawid (2003)**.
- We implemented the polytope method since it is generalisable for identified IV models with

exposures, outcomes, and instruments with more than 2 categories.

• Currently, we allow for a binary or 3 category instrument, and binary exposure and outcome.

## Example Mendelian randomization analysis

- We extract an example from Meleady et al. (2003).
- We have a 3 category instrument and binary exposure and outcome.
- We use the 677CT polymorphism (rs1801133) in the MTHFR gene, involved in folate metabolism, as an instrumental variable to investigate the causal effect of homocysteine on the risk of cardiovascular disease.
- The code is shown on the right.
- The ACE lies between a risk difference of -9% to 74% increase in absolute risk.
- Additionally, we see that the monotonicity inequality is not satisfied.

## Conclusion

- Use of bounds in instrumental variable analyses is regaining interest (Swanson et al. 2018; Labrecque and Swanson 2018).
- The empirical experience that the bounds are often wide is not a bad property of the method, it is a property of the typical data: Mendelian randomization data simply often are uninformative in that sense due to weak instrumental variables.
- We recommend using the bounds when the variables are genuinely discrete, but not when the exposure is genuinely continuous (Sheehan and Didelez 2019).
- Our R package and app provide a convenient interface to the bounds.

#### References

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Dawid, A. P. 2003. "Causal Inference Using Influence Diagrams: The Problem of Partial Compliance (with Discusssion)." In Highly Structured Stochastic Systems, edited by P. J. Green, N. L. Hjort, and S. Richardson, 45-65. New York: Oxford University Press.

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Sheehan, Nuala A, and Vanessa Didelez. 2019. "Epidemiology, genetic epidemiology and Mendelian randomisation: more need than ever to attend to detail." Human Genetics, 1-16. https://doi.org/10.1007/s00439-019-02027-3.

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## Extra Figures & Tables

```
library (bpbounds)
mt3 < -c(.83, .05, .11, .01,
          .88, .06, .05, .01,
          .72, .05, .20, .03)
p3 \leftarrow array(mt3, \underline{dim} = c(2, 2, 3),
            \underline{\text{dimnames}} = \mathbf{list}(\underline{x} = \mathbf{c}(0, 1),
                                \underline{y} = \mathbf{c}(0, 1),
                                z = c(0, 1, 2))
bpres3 <- bpbounds(as.table(p3))</pre>
summary (bpres3)
## Data:
                                  trivariate
## Instrument categories:
## Instrumental inequality: TRUE
## Causal parameter Lower bound Upper bound
                                             0.74000
                                 -0.09
                                             0.12000
         P(Y|do(X=0))
                                  0.06
                                             0.80000
         P(Y|do(X=1))
                                  0.03
                                  0.25
                                           13.33333
                    CRR
## Monotonicity inequality: FALSE
```



Figure 1: Shiny app https://remlapmot.shinyapps.io/bpbounds

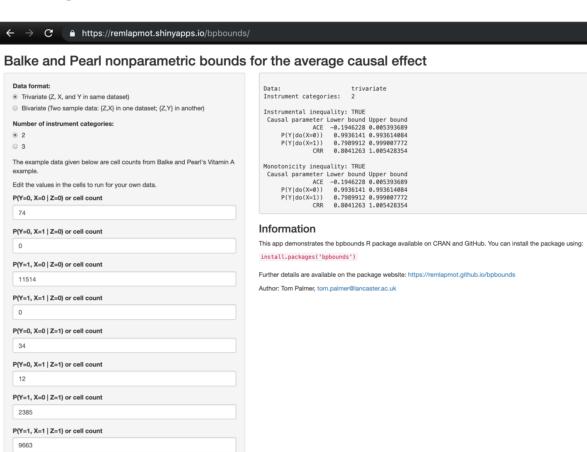


Figure 2: Screenshot of our Shiny app



Figure 3: Package website https://remlapmot.github.io/bpbounds/

