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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Extra Figures & Tables

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

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

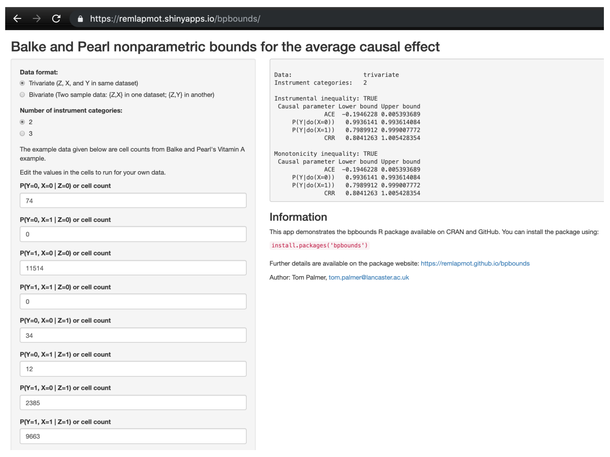

Figure 2: Screenshot of our Shiny app.

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