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:

install.packages("bpbounds")

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

Methods

- Balke and Pearl (1997) showed it is possible to derive bounds for the ACE.
- Bounds have the interpretation of:

There is some distribution of the unobserved confounders (of the exposure-outcome association) that yields a true ACE as small as the lower bound, while another choice produces an ACE as large as the upper bound.

- There are at least two ways to implement the Balke-Pearl bounds:
- i. conditional probabilities 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 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 phenotype and outcomes.
- 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).
- 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 \leftarrow array(mt3, dim = c(2, 2, 3),
           \underline{\text{dimnames}} = \mathbf{list}(\underline{x} = \mathbf{c}(0, 1),
                               y = 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
                                         13.33333
                    CRR
                                0.25
## Monotonicity inequality: FALSE
```



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

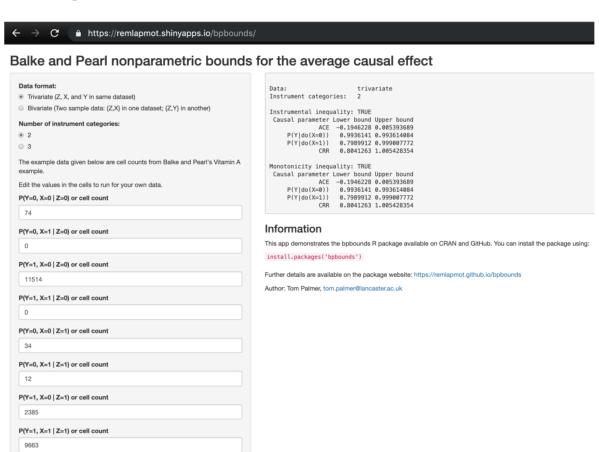


Figure 2: Screenshot of our Shiny app



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



