# Regression Discontinuity

Drew Dimmery drewd@nyu.edu

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#### Structure

- RDD interpretation
- RDD estimation
- Placebo tests
- Sorting
- Other stuff

## Interpretation

- It's a LATE!
- A different kind of LATE!
- It can be interpreted as a weighted average over all units (Lee & Lemieux 2010)
- $\bullet$  (W, U) are observed and unobserved factors which explain all heterogeneity.
- X=c is the cutpoint on the running variable, Y is the outcome  $\lim_{\epsilon\downarrow 0} E[Y|X=c+\epsilon] \lim_{\epsilon\uparrow 0} E[Y|X=c+\epsilon]$   $= \sum_{w,u} \tau(w,u) p(W=w,U=u|X=c)$   $= \sum_{w,u} \tau(w,u) \frac{f(c|W=w,U=u)}{f(c)} p(W=w,U=u)$
- What does this mean?
- It's a weight of individual treatment effects weighted by the likelihood that a unit will lie near the threshhold on the running variable.
- Keep this in mind as you interpret results.

#### Estimation

- If only someone wrote a package to do this...
- http://github.com/ddimmery/rdd

- The current best practices is to use local polynomial regression.
- Typically linear
- There are also some interesting methods using randomization inference, though. (Cattaneo et al n.d.)

## Replication

- I'll be replicating the recent Meyersson paper that's been making noise.
- Replication materials
- The paper shows a (local) result that when Islamic parties won elections in Turkey, this resulted in better outcomes for women.
- Running variable: vote margin (but not exclusively 2 party system as in Lee)
- Outcome that we'll look at: high school education

. . .

```
require(foreign, quietly = TRUE)
d <- read.dta("regdata0.dta")
summary(d$iwm94)

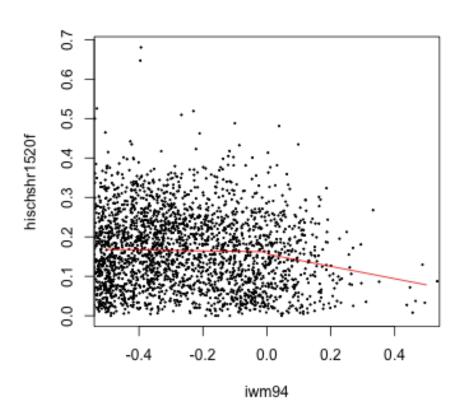
## Min. 1st Qu. Median Mean 3rd Qu. Max. NA's
## -1.0 -0.5 -0.3 -0.3 -0.1 1.0 544</pre>
```

## Explore data

• Plot the raw data.

. . .

```
with(d, plot(iwm94, hischshr1520f, pch = 19, cex = 0.2, xlim = c(-0.5, 0.5)))
left.lm <- lm(hischshr1520f ~ iwm94, d, subset = iwm94 < 0)
right.lm <- lm(hischshr1520f ~ iwm94, d, subset = iwm94 >= 0)
left.x <- seq(-0.5, 0, 0.01)
right.x <- -left.x
lines(left.x, predict(left.lm, newd = data.frame(iwm94 = left.x)), col = "red")
lines(right.x, predict(right.lm, newd = data.frame(iwm94 = right.x)), col = "red")</pre>
```



#### Estimation

- So the basic estimation would just take the difference of the intercepts from left.lm and right.lm.
- And there's an equivalency to just running a single regression as Cyrus showed in class.
- But I'm just going to use rdd

```
rd.out <- RDestimate(hischshr1520f ~ iwm94, d)
## Error: could not find function "RDestimate"
rd.out
## Error: object 'rd.out' not found

Full Results
summary(rd.out)
## Error: object 'rd.out' not found

Plot it
plot(rd.out, range = c(-0.4, 0.4))
## Error: object 'rd.out' not found

title(xlab = "Islamic Party Vote Margin", ylab = "Female High School Education Share")
## Error: plot.new has not been called yet</pre>
```

#### Placebo tests

- Do place bo tests on other covariates and other outcomes.
- They're "placebo" because there "shouldn't" be an effect on them (except occasionally by chance)

```
# Age 19+
RDestimate(ageshr19 ~ iwm94, d)[c("est", "se")]
## Error: could not find function "RDestimate"
# Log Population
RDestimate(lpop1994 ~ iwm94, d)[c("est", "se")]
## Error: could not find function "RDestimate"
# Household Size
RDestimate(shhs ~ iwm94, d)[c("est", "se")]
## Error: could not find function "RDestimate"
More Placebos
# Men in 2000
RDestimate(hischshr1520m ~ iwm94, d)[c("est", "se")]
## Error: could not find function "RDestimate"
# Women in 1990 (pre-treatment)
RDestimate(c90hischshr1520f ~ iwm94, d)[c("est", "se")]
## Error: could not find function "RDestimate"
# Men in 1990 (pre-treatment)
RDestimate(c90hischshr1520m ~ iwm94, d)[c("est", "se")]
```

## Error: could not find function "RDestimate"

# Sorting

• As Cyrus discussed, density tests are also a good way to examine the possibility of sorting.

DCdensity(d\$iwm94, verbose = TRUE, plot = FALSE)
## Error: could not find function "DCdensity"

## **Density Plot**

```
DCdensity(d$iwm94)
## Error: could not find function "DCdensity"
```

## Fuzzy designs

- I don't have an example for this, but it's quite easy.
- Do it the same way as before, but with RDestimate(Y~runvar+treatment)

#### Overall

- Some big things for RDD:
  - Lots of plots
  - Think about locality in interpretation
  - Use your covariates for robustness/placebo tests
  - Everything should be robust to different bandwidths, etc
  - If effects start disappearing as bw goes down, that's a bad sign.
  - Your bandwidth is probably to wide.
- If there's still more time, maybe I'll go through some high points of the rdd code.