## RDD and diff-in-diff

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Regression discontinuity design (RDD)

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## Basic assumption

RDD assumes a running variable (x) with a cut point (c) beyond which treatment is assigned (D).

$$D_i = \begin{cases} 1 & \text{if} \quad x_i \geqslant c \\ 0 & \text{if} \quad x_i < c \end{cases} \tag{1}$$

## Distinction

# It has a flavor of logit or propensity scores, but there are some differences:

- logit: x is not latent and we know the cutpoint: Both are observed and included as a predictors.
- ▶ **matching**: we have no control/treatment group. However, we assume that units on either side of the treatment are increasingly similar as their *x* is similar.

## Examples

**Administrative data are perfect:** You have some rule that kicks in at a specific threshold for otherwise almost identical observations.

- school test scores on school admission, restrictions on class size
- legal drinking age on alcohol related deaths
- election of candidates in close races.

▶ Individuals close to the threshold are interchangeable

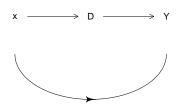
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X is a confounder ...so we only control for X

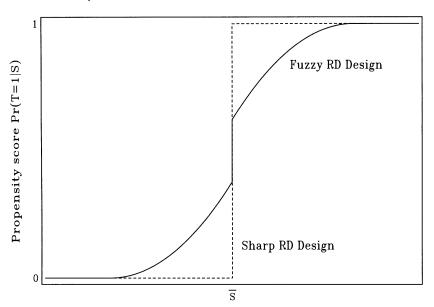


## Two designs

# We distinguish between two designs depending on how probable the treatment is:

- **sharp** RD: assignment is deterministic
- ▶ fuzzy RD: assignment is probabilistic

## A visual representation



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## Do it in R

# Check out alcohol related deaths (y) as a function of legal drinking age (D)

## Is that all?

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- 1. **the continuity assumption**: x must have a continuous effect on y
- 2. **no omited variable bias**:x must capture all influence on D.

The continuity assumption

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... that's not a regression discontinuity.

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- ightharpoonup recode the  $x \to \text{compare}$  with the recoding of y in GLMs.
- consider a sufficiently small window

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## We can create a curvilinear effect of x using polynomials (e.g.:)

$$y_i = \alpha + \rho D_i + \gamma_1 x_i + \gamma_2 x_i^2 \tag{4}$$

 $\Rightarrow$  Here, x has a symmetrical effect on both sides of the treatment.

## Recode the x: assymetric effect

We can assume x has different effects on each side of the treatment

$$y_i = \alpha + \rho D_i + \gamma x_i + \delta x_i D_i \tag{5}$$

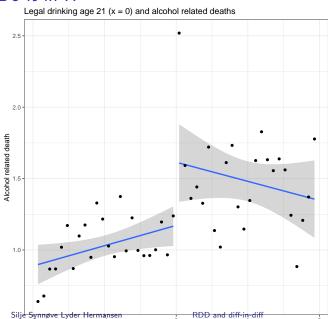
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$$y_i = \alpha + \rho D_i + \gamma x_i + \delta x_i D_i \tag{5}$$

 $\Rightarrow$  we center the x on the cutpoint  $(x_i - c) \rightarrow \rho$  still reports the change at the cutpoint.

## Do it in R



## Extrapolation

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• but we can also extrapolate y beyond the cutpoint with x:  $\rho + \delta(x - c)$ 

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- subsetting the data to tweak the window around the cutpoint is a non-parametric approach.

## Bandwidth: the idea

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► There's a tradeoff between linearity and statistical power (we need sufficient N).

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⇒ When you narrow down, do you get a weaker or stronger effect?

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- We can do it by hand
- ...or we can make an algorithm do it:
  - run a local weighted regression line
    - bandwidth is estimated accordingly
- ⇒ the point is to show robustness, not p-hack!

#### Omited variable bias

Fuzzy RD

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# Fuzzy RD

### Often the D increases the probability of a treatment, but we don't know!

⇒ This is a Instrumental Variable approach (more on Thursday)

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$$\mathbf{v} = \alpha + \beta_1 \bar{\mathbf{x}} + \beta_2 \mathbf{x} + \mathbf{e}$$

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• step 1: 
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- step 1:  $\bar{x} = \alpha_1 + \phi D + \beta_1 x + e_1$
- $\Rightarrow \gamma$  is the causal effect of D in a fuzzy design.

Differences-in-differences

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Definition: Comparing two differences

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- ⇒ Treatment effect is that difference

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- ▶ the difference between the treatment and control groups
- ⇒ based on panel data (units are observed several times).

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- Treatment: District 6 provided money to banks, while district 8 did not.
- After a year district 6 had 121 banks, while district 8 had 132
- ⇒ What was the treatment effect?

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# Example: States' monetary policy and number of banks

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within-unit difference: Number of banks before and after the crisis in each district.

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 $\Rightarrow$  Basically a 2-by-2 table

How to do it?

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- $\triangleright$   $\beta_3$  is the causal effect

### Data

### **Data requirements**

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- ightharpoonup Requires panel data ightharpoonup which means correcting the standard errors.
- Common panel types: state-year/administrative unit-time period; people over time . . .

> we want to know the trend before and after the break

Another example: drinking age and death

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# Does the legal drinking age has an effect on death rates among the young?

- y is number of deaths per 100 000
- P is post-treatment dummy
- T is dummies for states
- trend is year dummies

### Another example: step $1 \rightarrow$ calculate differences

The authors have two tricks:

- ► Hardcode the interaction effect (dummy before/after treatment)
- ▶ They remove the intercept to retain all dummies

```
load("df2.Rda")
##with intercept
mod <- lm(mrate ~ legal +
            state +
            year_fct,
          df)
##without intercept; with all dummies
mod <- lm(mrate ~ 0 +
            legal +
            state +
            year fct,
          df)
```

# Another example: step $2 \rightarrow$ calculate errors

#### Calculate robust standard errors:

library(clubSandwich)

# Another example: step $3 \rightarrow$ interpretation

### Display the results and interpret:

# Another example: step $3 \rightarrow$ interpretation

Table 1: Death rates among young as a function of legal drinking age

	Dependent variable:
	mrate
Legal drinking age (causal effect)	10.804**
	(4.479)
Observations	714
$R^2$	0.986
Adjusted R <sup>2</sup>	0.985
Residual Std. Error	17.339 (df = 649)
F Statistic	$726.005^{***} (df = 65; 649)$
Note:	*p<0.1; **p<0.05; ***p<0.01

 $\Rightarrow$  What did we find?

The parallel trends assumption

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- we can use weights
- ⇒ There's a tradeoff: treatment is at the unit level, statistical power at the subunit level.