

Applied Microeconometrics I

Lecture 10: Regression discontinuity design

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October 5, 2023
Lecture Slides

What did we do last time?

- **Difference-in-differences (DID):**
 - RCT is seldom feasible, CIA is unlikely to hold, and good instruments are hard to find
 - Often there is a *before* and *after* a treatment is assigned (e.g. policy change) and some units are treated while others are not.
 - We need *panel data*
- **Idea:** we use the evolution of Y in the control group to build the counterfactual of “what would have happened” in the treatment group in the absence of the treatment.
- **Identifying assumption:** in the absence of the treatment, the outcomes of both groups would follow *parallel trends*.
- We allow for time-invariant (level) differences between the two groups, but rule out time-variant differences (trends).

What did we do last time?

- Regression form (FE notation):

$$Y_{i,t} = \alpha_i + \lambda_t + \rho D_{i,t} + \varepsilon_{i,t}$$

- $D_{i,t} = 1$ if death penalty is abolished and zero otherwise
- $E(\varepsilon_{i,t} | \alpha_i, \lambda_t, D_{i,t}) = 0$
- Simplest case: 2 time periods, 2 cross-sectional units (States)
 - Comparison of the two States across time periods gives ρ

- An equivalent way to write the 2x2 DID equation:

$$Y_{ist} = \alpha + \gamma TREAT_s + \lambda AFTER_t + \rho(AFTER_t * TREAT_s) + \varepsilon_{ist}$$

- Threats to validity:
 - Lack of parallel trends
 - Anticipatory behavior (e.g. leading to GE effects)
- ρ interpretation: shocks/policies implemented at same time?

What did we do last time?

- Several model extensions:
 - Add controls X (to achieve identification)
 - Multiple time periods (before and after)
 - inspect pre-trends
 - placebo policy changes
 - dynamic effects
 - Multiple treated groups, treated at different times
 - Different types of controls (never-treated; treated later)
 - i and t can be *anything!*
 - Jimeno and Boeri (2005): “workers” and “firms”

Today: Regression Discontinuity Design

- Exploit institutional rules that create RCT's
 - a known (to us) threshold rule assigns people to treatments
 - close to the threshold, the treatment assignment is randomized
 - if nothing else discontinuously jumps at the threshold (other than the treatment status), we can causally relate outcome change to the treatment change at the threshold
- Randomization in *observational data*
- Identification: Imperfect manipulation, continuity assumption
- Types of RDD: sharp vs. fuzzy
- Sharp RDD:
 - model specification, estimation, and testing validity
 - sharp RDD as a local RCT

Regression discontinuity design (RDD)

Institutional rules create experiments

- Institutional rules often assign individuals to “treatments”; this can be exploited for estimating causal effects
- Typical case: **threshold rule** that is based on ex-ante variable
 - *Score* in entry exams for college admission
 - *Income* for subsidy eligibility
 - *Project quality score* for public R&D subsidies eligibility
 - *Age limit* for legal alcohol consumption
- The ex-ante variable X is called the **running variable** (W refers to pre-determined observable characteristics)
- A pre-determined **threshold** c of X assigns individuals to the binary **treatment status** $D \in \{0, 1\}$
 - $D = 1$ if treated and $D = 0$ if not treated
 - In principle, D can be continuous

Regression discontinuity design

Institutional rules create experiments

- RDD exploits the **randomness** of the treatment assignment **around the threshold**
- Provided that D is as good as randomly assigned at c , we can compare the Y below (control) and above (treated) c .
- **Main assumption:** *people cannot perfectly manipulate X*
 - X manipulation means being able to set own D_i value
 - some manipulation is allowed
 - but no one can perfectly decide to be just below/above c
 - imperfect self-selection around c cannot be fully tested

Regression discontinuity design

Imperfect manipulation and identification

- *Imperfect manipulation:*
 - It **implies** that X is continuous around c
 - lack of continuity is consistent with precise sorting around c
 - The continuity of X implies *local randomization*
 - Randomization is key to identify the causal estimand of interest
- **Why does RDD work?**
 1. Aim: estimate effect of D on Y for people at the threshold
 2. Institutional rules assign D based on X (above/below c)
 3. At the threshold, we want to causally relate the change in Y with the discontinuity in D induced by the running variable
 4. For this to work, *no other variable can jump at c*
 5. In other words, any other discontinuities at c (in either X or observable/unobservable characteristics) will bias the estimates
 6. Continuity implies randomization: people close to c are similar

Regression discontinuity design

Continuity assumption

- Why do we require continuity (and not randomization)?
- In RDD:
 - All pre-determined observable and unobservable characteristics (W and U) are **allowed to be correlated with D** ! (and Y)
 - we only require that they evolve smoothly around c .
 - X is also **allowed to be correlated with W and U**
 - we require that, conditional on (W, U) , X passes smoothly across c
- Assuming that D is as good as randomly assigned conditional on (W, X) would mean relying on CIA (and we don't want to!)
- Remember, randomization *follows* from X continuity at c .
 - X continuity ensures that at c the effect of the discontinuous jump of D on Y is not confounded by a jump in X
 - X continuity implies that, conditional on X , on average the potential outcomes evolve smoothly across c

Regression discontinuity design

Consequences of the local random assignment

- Local random assignment means that treated and comparison units are not systematically different in any dimension at c
 - It is “pure luck” whether i is just above vs. below c
 - This follows from the inability of precisely manipulating X to be on one side vs. the other
- Three implications local random assignment:
 1. **Identification:** It is a necessary condition (and sufficient, in sharp RDD) to identify causal effect of D on Y at c
 2. **Validity testing:** test whether pre-determined W jump at c (similar to randomization test in RCTs)
 3. **Irrelevance of W :** their inclusion/exclusion should not matter for consistency (only for precision)

Types of Regression discontinuity designs

- Overall, RDD is compelling design (high internal validity):
 - local RCT
 - if X and c are known to the researcher, one can exploit local randomization *in observational data*
 - local randomization does not need to be assumed, it *follows* from imperfect manipulation of X .

- **Sharp RDD:**

1. Treatment status D is a deterministic function of X
2. D is a discontinuous function of X

That is, D jumps from 0 to 1 at c (*everyone* with $X_i \geq c$ receives $D_i = 1$).

- **Fuzzy RDD** allows for a smaller jump at c .

Types of Regression discontinuity designs

- We can often redefine D to have a sharp vs. fuzzy design:

X	D	Y	RDD
Test score	college admission	Income	sharp
Test score	college enrollment	Income	fuzzy
Income	welfare benefits eligibility	Employment	sharp
Income	welfare benefits take-up	Employment	fuzzy
Age	legal alcohol consumption	Death	sharp
Age	actual alcohol consumption	Death	fuzzy

TABLE 1 – Example of RDD applications

- Typically:
 - Sharp RDD if X fully/deterministically captures the eligibility
 - Fuzzy RDD if D is participation/take-up
 - Fuzzy RDD if D assignment follows more rules than just X

Effect of Minimum Legal Drinking Age on death rates

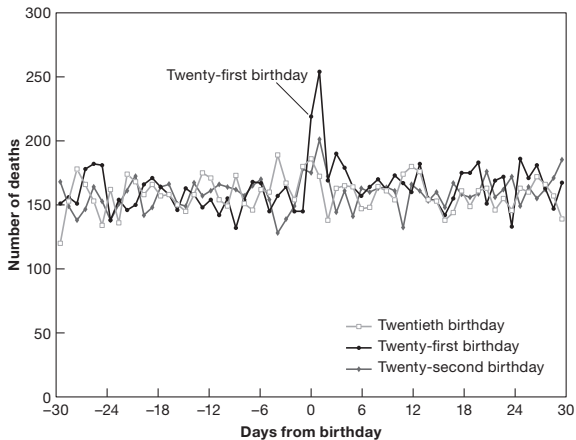
Carpenter and Dobkin (2009)

- We are interested in the effect of the minimum legal drinking age (in the US it's 21) on mortality.
- What type of RDD is this?
- Setting:
 - outcome variable Y_i : death rate
 - treatment status D_i : legal drinking status
 - running variable X_i : age
 - cutoff $c = 21$: MLDA rule transforms 21-year-olds from underage minors to legal alcohol consumers.

Effect of Minimum Legal Drinking Age on death rates

Carpenter and Dobkin (2009)

FIGURE 4.1
Birthdays and funerals

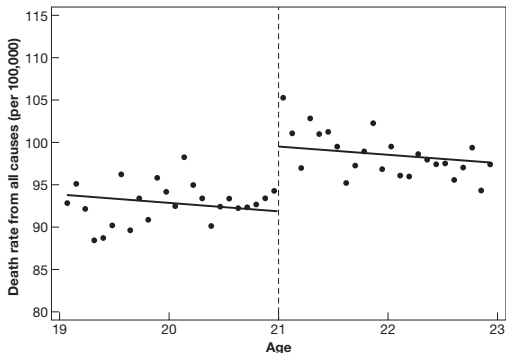


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Effect of Minimum Legal Drinking Age on death rates

Carpenter and Dobkin (2009)

FIGURE 4.2
A sharp RD estimate of MLDA mortality effects



Notes: This figure plots death rates from all causes against age in months. The lines in the figure show fitted values from a regression of death rates on an over-21 dummy and age in months (the vertical dashed line indicates the minimum legal drinking age (MLDA) cutoff).

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Sharp Regression Discontinuity Design

- Suppose that D_i is a deterministic and discontinuous function of the running variable X_i
- D jumps from 0 to 1 at c . For each i :

$$D_i = \begin{cases} 1 & \text{if } X_i \geq c \\ 0 & \text{if } X_i < c \end{cases}$$

Or, more compactly, $D_i = 1[X_i \geq c]$.

- **All people** to the right of the cut off are exposed to the treatment and **all those** to the left are denied the treatment

Sharp Regression Discontinuity Design

Linear case

- What is the effect of *college admission* (D) on *income* (Y)?
- We have information on the college admission rules:
 - admission is entirely based on applicant's *entry test score* (X).
 - students scoring at least c are admitted: $D = 1[X \geq c]$.
- We can write the model for Y_i as:

$$Y_i = \alpha + D_i\tau + X_i\beta + u_i$$

- Note that:
 - Y_i varies discontinuously with D_i (if $\tau \neq 0$)
 - D_i is discontinuous function of X_i generating treat. effect τ
 - here, *by assumption* the relationship between Y_i and X_i is linear (and the same above/below c)
- Side note: can we use X as an instrument for D ?

Sharp Regression Discontinuity Design

Linear case

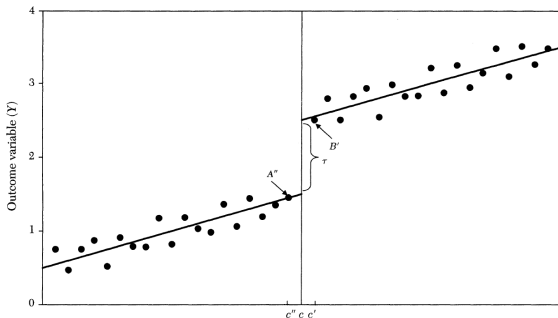


FIGURE 1 – Linear RDD

- The causal effect for a person whose $X_i = c$ requires us to *extrapolate* beyond c
- Suppose that all factors (other than college admission) affecting Y_i evolve smoothly with respect to X_i
- Then B' (A'') is a reasonable guess for Y_i when $D_i = 1$ ($D_i = 0$)
- And $B' - A''$ would be the impact of treatment on Y_i

Sharp Regression Discontinuity Design

Specifying the functional form

- **Tradeoff:**
 - Estimates are more accurate, the closer we are to the threshold
 - The closer we are to the threshold, the less data we have
- We *always* need to use data away from the threshold
 - In the example, no individual had $X_i = c$!
 - extrapolation means that we must *always* control for X_i
- As a result we need to **assume a functional form** for the relationship between Y and X
 - In the example, a linear relationship seems appropriate
 - If the true population model is linear, OLS using all observations provides the best estimate for τ
- Since we need data away from threshold, the τ **estimate depends on chosen functional form**
 - In the example, if we restricted $\beta = 0$, OLS would be biased

Sharp Regression Discontinuity Design

Specifying the functional form

- We saw the linear example
- But in general the relationship can be any $f(X_i)$:

$$Y_i = \alpha + \tau D_i + f(X_i) + u_i$$

- For example, ρ :th order polynomial:

$$f(X_i) = \beta_1 X_i + \beta_2 X_i^2 + \beta_3 X_i^3 + \dots + \beta_\rho X_i^\rho$$

- $f(X_i)$ can also be different at each side of the cutoff point
- Relies on the assumption that $f(X_i)$ is an adequate description of the relationship between Y and X
- The further away from c we are, the bolder this assumption is
- Therefore, it makes sense to use estimation methods for $f(X_i)$ well-suited *locally*

Sharp Regression Discontinuity Design

Nonlinear case 

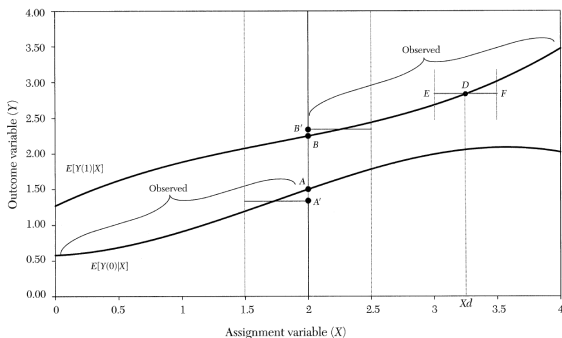


FIGURE 2 – Non-linear RDD

- $Y_i(0)$, $Y_i(1)$: potential outcomes. The i -level treatment effect is never observed.
- $E[Y_i(0)|X]$, $E[Y_i(1)|X]$: underlying relationships between the average outcomes and X (non-linear; different above/below c).
- We *never* observe $E[Y_i(1)|X]$ below $c = 2$ (the opposite holds for $E[Y_i(0)|X]$).

Sharp Regression Discontinuity Design

Nonlinear case

- With what is observable, we can estimate:

$$B - A = \lim_{\epsilon \rightarrow 0} E[Y_i | X_i = c + \epsilon] - \lim_{\epsilon \rightarrow 0} E[Y_i | X_i = c - \epsilon]$$

- Assume that X evolves continuously across c . This implies that also the underlying $E[Y_i(0)|X]$ and $E[Y_i(1)|X]$ do so.
- Then $B - A$ at the limit is equal to:

$$\tau = E[Y_i(D_i = 1) - Y_i(D_i = 0) | X_i = c]$$

- This is the treatment effect at the threshold c
- Continuity enables us to use the average outcomes of those right below c and valid counterfactuals for those right above
- We can *estimate* smooth $E[Y|X]$ relationships before/after c , but must *assume* relationship is smooth in the counterfactuals

Sharp Regression Discontinuity Design

Estimation within a bandwidth

- How to estimate $E[Y_i|X_i = c + \epsilon]$ and $E[Y_i|X_i = c - \epsilon]$?
 - $f(X_i)$ approximates well relationship between X and Y ?
 - The further away from c we are, the bolder this assumption is
 - We need estimation methods for $f(X_i)$ well-suited *locally*
- **Non-parametric methods** flexibly estimate $E[Y|X]$ below and above c within a bandwidth (window) of width h .
- E.g., *local constant estimator* (Figure 2):
 - choose interval h around $c = 2$, e.g. $(-1.5, 1.5)$
 - use data in $(-1.5, 2)$ to compute A' ; repeat above c to get B'
 - $B' - A'$ is the estimate for τ
- Alternatively, *local linear/polynomial regression*, and we can give more weight to observations closest to c
- How to choose h ? **Precision vs. bias tradeoff**
 - Literature on optimal bandwidths

Model specification and estimation

Sharp RDD

- Standard approach is to choose a window of width h and use *local linear regression* with rectangular kernel.
- This is equivalent to estimating two linear regressions over window h on both sides of c :

$$\begin{aligned} Y_i &= \alpha_l + \beta_l \tilde{X}_i + u_i & \text{if } \tilde{X}_i \in [-h, 0) \\ Y_i &= \alpha_r + \beta_r \tilde{X}_i + u_i & \text{if } \tilde{X}_i \in [0, h] \end{aligned}$$

- with $\tilde{X}_i = X - c$, hence $D_i = 1[\tilde{X}_i \geq 0]$
 - this way, α_l α_r are the value of the regression functions at c .
 - then, $\tau = \alpha_r - \alpha_l$
- Equivalently, **do OLS** once to estimate *pooled regression*:

$$Y_i = \alpha_l + \tau D_i + \beta_l \tilde{X}_i + (\beta_r - \beta_l) D_i \tilde{X}_i + u_i, \quad \tilde{X}_i \in [-h, h]$$

- again, $\tau = \alpha_r - \alpha_l$, but now we directly get $\hat{\sigma}(\hat{\tau})$

Model specification and estimation

Sharp RDD

- Note that in general we can rewrite the pooled regression as:

$$Y_i = \alpha_l + \tau D_i + f(\tilde{X}_i) + u_i, \quad \tilde{X}_i \in [-h, h]$$

with $f(\tilde{X}_i) = f_l(\tilde{X}_i) + D[f_r(\tilde{X}_i) - f_l(\tilde{X}_i)]$

- for local linear regression: $f_l(\tilde{X}_i) = \beta_l \tilde{X}_i$ and $f_r(\tilde{X}_i) = \beta_r \tilde{X}_i$
- we can alternatively use local polynomial regression for $f(\tilde{X}_i)$
- alternative non-parametric methods exist, but local linear regression with rectangular kernel (same weight to all observations in h) is a good start
- optimal bandwidth algorithms and manual inspection help deciding h value (and robustness of results to it)

Testing for sharp/fuzzy RDD validity

- We cannot test whether individuals influence the running variable precisely. But we can check:
 1. **institutional rules defining X_i :**
 - is it likely/feasible to precisely sort on one side vs. the other?
 - do people even know about c ?
 2. **local random assignment:**
 - do pre-determined W jump at c ? (similar to balance test in RCTs)
 - does W inclusion change estimates? (should only affect precision)
 3. **continuity (McRary test):**
 - Is the density of X continuous across c ?
 4. **relationship between Y and X**
 - are results robust to the use of different h ? (bias-variance trade-off)
 - what is the optimal bandwidth?
 - are estimates sensitive to different functional forms for $f(X_i)$?
- **Transparency:** plot Y vs. X to see variation that identifies τ