#### More on Diff-In-Diffs Methods

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#### Outline

- Propensity Score Matching-DID
- Semi-parametric DID
- Synthetic DID
- Non-linear DID
- Staggered Difference-In-Differences

## Propensity Score Matching - DID

#### Potential Outcome Framework

#### General Idea of Matching

Think of whether taking Chen's RMEB class as a binary decision,  $D_i$ ; and income in the future  $Y_i$  is the interested outcome:

$$Potential\ Outcome = \begin{cases} Y_i^1, & \textit{if}\ D_i = 1 \\ Y_i^0, & \textit{if}\ D_i = 0 \end{cases}$$

We aim to estimate

$$ATT = \mathbf{E}(Y_i^1 - Y_i^0 | D_i = 1, \mathbf{X}_i), \quad D_i \in \{0, 1\}$$

The problem is only  $Y_i = D_i Y_i^1 + (1 - D_i) Y_i^0$  is observed. In lack of ideal experiment, we use nonparticipants (control group) to approximate participants:

$$\underbrace{E(Y_i|D_i=1,\mathbf{X}_i) - E(Y_i|D_i=0,\mathbf{X}_i)}_{observed \ income \ difference} = \underbrace{E(Y_i^1 - Y_i^0|D_i=1,\mathbf{X}_i)}_{\delta(\mathbf{X}_i) = ATT|\mathbf{X}_i} + \underbrace{E(Y_i^0|D_i=1,\mathbf{X}_i) - E(Y_i^0|D_i=0,\mathbf{X}_i)}_{Selection \ Bias}$$

## Conditional Independence Assumption

General Idea of Matching

#### Selection Bias:

$$Bias(\mathbf{X}_i) = \mathbf{E}(Y_i^0|D_i = 1, \mathbf{X}_i) - \mathbf{E}(Y_i^0|D_i = 0, \mathbf{X}_i)$$

- Conditional Independence Assumption (CIA):  $\{Y_i^0, Y_i^1\} \perp D_i | \mathbf{X}_i$ 
  - Under CIA,  $Bias(\mathbf{X}_i) = 0$
  - CIA essentially states that  $D_i$  is randomly assigned conditioning on observeable characteristics  $\mathbf{X}_i$  (see Angrist (1998) on voluntary military service)
  - CIA is likely to fail when unobserved characteristics determine select-into-treatment

"The idea of matching between treated and untreated units assumes that  $Bias(\mathbf{X}_i) = 0$  so that conditioning on  $\mathbf{X}_i$  eliminates the bias."

## Regression vs. Matching

#### General Idea of Matching

- Statisticians more often use matching methods (Cochrane and Rubin, 1973); the idea is quite similar to conditioning on observeables in the regression analysis
- "Regression can be motivated as a particular sort of weighted matching estimator" (Angrist and Pischke: *Mostly Harmless Econometrics*).
  - **Matching estimator** puts the most weight on covariate cells containing units who are most likely to be treated: high  $Pr(D_i = 1 | \mathbf{X}_i)$
  - **Q** Regression puts the most weight on covariate cells with the largest conditional variance of treatment status: when  $Pr(D_i = 1|\mathbf{X}_i) \times [1 Pr(D_i = 1|\mathbf{X}_i)]$ ) is large
  - **Ommon Support:** No weights assigned to covariate cells containing no treated and control units:  $0 < \Pr(D_i = 1 | \mathbf{X}_i) < 1$

## **Propensity Score Matching**

#### A Structural Interpretation by Heckman, Ichimura, and Todd (1998)

- The dimension of  $X_i$  can be high, adding to difficulty of matching
- The well-known PSM is to match over the propensity of selection into the treatment (Rosenbaum and Rubin, 1983)
- Exclusion Restrictions: Partition X into (T, Z) such that:

$$Y_i^0 = g_0(\mathbf{T}_i) + U_i^0, (1)$$

$$Y_i^1 = g_1(\mathbf{T}_i) + U_i^1, (2)$$

$$Pr(D_i = 1 | \mathbf{X}_i) = Pr(D_i = 1 | \mathbf{Z}_i) = P(\mathbf{Z}_i)$$
 (propensity score) (3)

T and Z are not necessarily mutually exclusive

• To identify the ATT, it's enough to assume

$$U_i^0 \perp D_i | P(\mathbf{Z}_i)$$
 (PSM CIA)

## **Propensity Score Matching**

A Structural Framework by Heckman, Ichimura, and Todd, (1998)

PSM method can be put in the context of classical econometric selection models:

$$Y_i^0 = g_0(\mathbf{T}_i) + U_i^0 \tag{4}$$

$$D_{i} = \begin{cases} 1 & if \quad \lambda(\mathbf{Z}_{i}) - v \ge 0 \\ 0 & otherwise \end{cases}$$
 (5)

• If  $\mathbf{Z}_i$  and v are independent, then  $P(\mathbf{Z}_i) = \Pr\{v \leq \lambda(\mathbf{Z}_i)\} = F_v[\lambda(\mathbf{Z}_i)];$  the CIA implies that

$$E\{U_i^0|D_i=1, F_v[\lambda(\mathbf{Z}_i)]\} = E\{U_i^0|D_i=0, F_v[\lambda(\mathbf{Z}_i)]\}$$

• If also  $\lambda(\mathbf{Z}_i) \perp (U_i^0, v)$  and  $E(U_i^0) = 0$ , then for any s:

$$E(U_i^0|v=s) = 0$$
 (No Selection on Observables)

## Propensity Score Matching based DID

Heckman et al. (1998, ReSTud)

Recall the OLS estimator the two-by-two DID model

$$Y_{it} = \delta D_{it} + \underbrace{\lambda t + \eta_i + \nu_{it}}_{U_{it}(D_{it})}$$

is simply:

$$\delta^{OLS} = E[Y_{i1} - Y_{i0}|D_{i1} = 1] - E[Y_{i1} - Y_{i0}|D_{i1} = 0]$$

A sufficient condition for the identification is

$$Pr(D_{i1} = 1 | v_{it}) = Pr(D_{i1} = 1)$$

• If  $v_{it}$  is correlated in time, the model cannot be identified: What are possible solutions?

• The traditional solution is to add covariates  $X_i$  into the model:

$$Y_{it} = \mu + \tau D_{it} + \mathbf{X}_i' \alpha(t) + \lambda t + \varepsilon_{it}$$

$$\implies Y_{i1} - Y_{i0} = \tau(D_{i1} - D_{i0}) + \mathbf{X}_i' \alpha + \lambda + \varepsilon_{i1} - \varepsilon_{i0}$$

- $\bullet$   $X_i$  usually represents pre-treatment characteristics
- Ideally, covariates  $X_i$  should be treated non-parametrically as  $H(X_i)$
- The PSM-DID method is a semi-parametric way of obtaining ATT:

$$\delta^{PSM-DID} = E(Y_{i1} - Y_{i0} | \mathbf{X}_i, D_{i1} = 1) - E(Y_{i1} - Y_{i0} | \mathbf{X}_i, D_{i1} = 0)$$

$$Parallel\ Trend \Rightarrow \delta^{PSM-DID} = E(U_{i1} - U_{i0} | P(\mathbf{Z}_i), D_{i1} = 1)$$

$$-E(U_{i1} - U_{i0} | P(\mathbf{Z}_i), D_{i1} = 0)$$

- The PSM-DID estimator permits:
  - Selection to be dependent on potential outcomes
  - Selection on unobservables

#### How to Implement PSM-DID for Panel Data?

#### • Estimation:

- Calculate propensity score  $P(\mathbf{Z}_i)$  using probit or logit models
- Matching by cohort-year over the estimated propensity score
- **③** Generate differenced outcome variables  $\Delta Y_{it} = Y_{it} Y_{i0}$  and calculate treatment effects for oberservation it; Using appropriate weights to obtain  $ATT_{gt}$  at a desired aggregation level

These steps apply to almost any matching-DID estimators (NN matching...)

- **Inference**: Analytical standard errors proposed by Abadie and Imbens (2006).
- Stata command: teffects with a nice intro PDF

# Semi-parametric DID

Under the conditional common trend assumption:

$$E[Y_{i1}^0 - Y_{i0}^0 | \mathbf{X}_i, D_{i1} = 1] = E[Y_{i1}^0 - Y_{i0}^0 | \mathbf{X}_i, D_{i1} = 0]$$

Abadie (2005) shows that ATT can be estimated using simple weighting schemes:

$$E[Y_{i1}^{1} - Y_{i1}^{0} | \mathbf{X}_{i}, D_{i1} = 1] = E[\rho_{0}(Y_{i1} - Y_{i0}) | \mathbf{X}_{i}]$$
where 
$$\rho_{0} = \frac{D_{i1} - P(D_{i1} = 1 | \mathbf{X}_{i})}{P(D_{i1} = 1 | \mathbf{X}_{i})[1 - P(D_{i1} = 1 | \mathbf{X}_{i})]}$$

$$ATT = E[Y_{i1}^{1} - Y_{i1}^{0} | D_{i1} = 1] = E\left[\frac{Y_{i1} - Y_{i0}}{P(D_{i1} = 1)} \cdot \frac{D_{i1} - P(D_{i1} = 1 | \mathbf{X}_{i})}{1 - P(D_{i1} = 1 | \mathbf{X}_{i})}\right]$$
(6)

## Semi-parametric DID

The dis-aggregated ATT can be obtained by approximating  $E[Y_{i1}^1 - Y_{ii}^0 | D_{i1} = 1, \mathbf{X}_i^{sub}]$  as:

$$\delta(\mathbf{X}_{i}^{sub})^{semi-DID} = argmin_{\theta} E\left\{P(D_{i1} = 1 | \mathbf{X}_{i}) \cdot \left[\rho_{0}(Y_{i1} - Y_{i0}) - g(\mathbf{X}_{i}^{sub}; \theta)\right]^{2}\right\}$$

 $\mathbf{X}_{i}^{sub}$  is a function of  $\mathbf{X}_{i}$ ;  $\mathbf{X}^{sub}$  may contain a subset of variables in  $\mathbf{X}_{i}$ 

- Estimation Strategy:
  - Estimate propensity score  $P(D_{i1} = 1 | \mathbf{X}_i)$
  - Plug the fitted values into the sample analogue of the above equation
- Stata implementation: absdid [Link to Stata Manual]

# Synthetic DID

## Synthetic DID

#### Arkhangelsky et al. (2021, AER)

- *DID methods:* A substantial number of treated units; researchers invoke "parallel trend" to control for slection effects
- Synthetic Control (SC) methods: A single (or small number) of units exposed, seek to compensate for the lack of parallel trends by re-weighting units to match their pre-exposure trends
  - Synthetic control estimator by Abadie et al. (2010, JASA):

$$\hat{\delta}_{t}^{SC} = Y_{t}^{1} - \sum_{i=2}^{I+1} \omega_{i}^{*} Y_{it}^{*}$$
 (7)

where  $\omega^* = (\omega_2^*, \cdots, \omega_{I+1}^*)$  is chosen to minimize  $||X_1 - X_0 \omega||$ 

- Synthetic DID have features of both SC and DID:
  - Like SC, it re-weights and matches pre-exposure trends to weaken the reliance on parallel trend type assumptions
  - ② Like DID, it is invariant to additive unit-level shifts, and allows for valid large-panel inference.

- A balanced panel with N units and T periods, binary treatment  $D_{it} \in \{0,1\}$ :
  - first  $N_0$  units untreated, last  $N_1 = N N_0$  units treated
  - Units exposed to treatment after  $T_0$
- Basic procedures:
  - Like SC, find weights  $\{\hat{\omega}_i^{sc}\}$  such that  $\sum_{i=1}^{N_0} \hat{\omega}_i^{sc} \approx \frac{1}{N_1} \sum_{i=N_0+1}^{N} Y_{it}$  for all  $t=1,\cdots,T_0$
  - Then use these weights in a basic panel data DID regression (two-way fixed effects) to estimate the ATT:

$$(\hat{\boldsymbol{\delta}}^{sdid}, \hat{\mu}, \hat{\alpha}_i, \hat{\beta}_t) = \operatorname{argmin}_{\delta, \mu, \alpha_i, \beta_t} \left\{ \sum_{i=1}^{N} \sum_{t=1}^{T} (Y_{it} - \mu - \alpha_i - \beta_t - \delta D_{it})^2 \hat{\omega}_i^{sdid} \right\}$$
(8)

 DID can be thought of a special case of Synthetic DID without unit weights

## Synthetic DID

#### Benefits of Synthetic DID:

- Using similar units and similar periods makes the estimator more robust [*Intuition*: emphasize units that are more similar to treated units]
  - Example: Effect of anti-smoking legislation on California (Abadie et al., 2010)
- Use of the weights may (Not always) improve the estimator's precision by removing systematic (predictable) parts of the outcome

Software Implementation: [Stata]: sdid; [R]: synthdid

"The synthetic control DID may gain its popularity in the future, especially in the area of public policy evaluation"

# Nonlinear Diff-In-Diffs: Change-In-Changes

## Nonlinear DID (Change-In-Changes)

#### Athey and Imbens (2006, Econometrica)

 The classical DID model assumes potential outcomes are separably additive:

$$Y_{gt}^{I} = g_{I}(D_{gt}) + U_{gt}^{I}, \text{ for } I \in \{0, 1\}$$

• Athey and Imbens (2006) generalizes the function of potential outcomes:

$$Y_{gt}^0 = h(D_{gt}, U_{gt}^0)$$

Now the goal is to back out their distribution. Under very similar conditions as in DID, the distribution of  $Y_{11}^0$  can be identified as

$$F_{11}^{0}(y) = F_{10}[F_{00}^{-1}(F_{01}(y))]$$

Linear DID: 
$$E(Y_{11}^0|D_{11}=1) = E(Y_{10}^0|D_{11}=1) + [E(Y_{01}^0|D_{01}=0) - E(Y_{00}^0|D_{01}=0)]$$

- Can be extended to count data models (patents), quantile treatment effects...
- Stata implementation: cic

# Staggered Diff-In-Diffs

## Why Staggered Diff-In-Diffs

- Canonical DID framework assumes that policy happen at one time
- But in many empirical settings, policy interventions enact in a staggered manner, generating variations in treatment timing: China's pilot programs, export/import decision, digitalization...
- The two-way fixed effects (TWFE) model were widely adopted by researchers:

$$Y_{it} = \delta D_{it} + \lambda_i + \lambda_t + \varepsilon_{it}$$

- It turns out our understanding of the TWFE estimator is pretty limited, and the interpretation of  $\delta$  is unclear
- Several recent methodological papers start to deal with this problem

#### **Understanding TWFE**

3 groups and 3 periods (Goodman-Bacon, 2020)

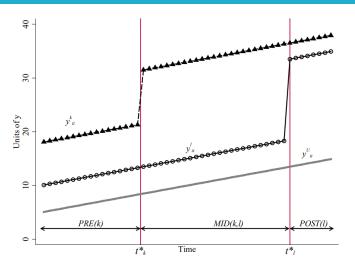


Figure 1: Staggered Difference-in-Differences: 3-by-3 Case

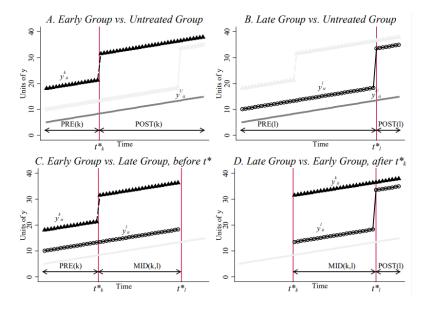


Figure 2: All four simple (2-by-2) DID Estimates

**Bacon's Decomposition Theorem** (Goodman-Bacon, 2020): Assume that the data contain  $k = 1, \dots, K$  groups ordered by the treatment timing. The TWFE estimate is a weighted average of all possible two-by-two DID estimators:

$$\hat{\delta}^{TWFE} = \sum_{k \neq U} \underbrace{w_{kU} \hat{\delta}_{kU}^{2 \times 2}}_{\text{treat vs. never treated}} + \sum_{k \neq U} \sum_{\ell > k} w_{k\ell} \underbrace{\left[ \underbrace{\mu_{k\ell} \hat{\delta}_{k\ell}^{2 \times 2, k}}_{\text{early vs. late}} + \underbrace{\left( 1 - \mu_{k\ell} \right) \delta_{k\ell}^{2 \times 2, \ell}}_{\text{late vs. early}} \right]}_{\text{late vs. early}}$$
(9)

where  $\sum_{k\neq U} w_{kU} + \sum_{k\neq U} \sum_{\ell>k} w_{k\ell} = 1$  and the weights are

$$\begin{split} w_{kU} &= \frac{N_k N_U \bar{D}_k (1 - \bar{D}_k)}{v \hat{a} r(\tilde{D}_{it})} \\ w_{k\ell} &= \frac{N_k N_\ell (\bar{D}_k - \bar{D}_\ell) [1 - (\bar{D}_k - \bar{D}_\ell)]}{v \hat{a} r(\tilde{D}_{it})} \\ \mu_{k\ell} &= \frac{1 - \bar{D}_k}{1 - (\bar{D}_k - \bar{D}_\ell)} \end{split}$$

where 
$$\tilde{D}_{it} = (D_{it} - \bar{\bar{D}}) - (\bar{D}_i - \bar{\bar{D}}) - (\bar{D}_t - \bar{\bar{D}})$$
 with  $\bar{\bar{D}} = \frac{1}{NT} \sum_i \sum_t D_{it}$ ,  $\bar{D}_i = \sum_t D_{it}$  and  $\bar{D}_t = \sum_i D_{it}$ 

#### Causal Interpretation of the TWFE Estimator

Goodman-Bacon shows the coefficient  $\delta^{TWFE}$  can be interpreted as

$$\delta^{TWFE} = \underbrace{VWATT}_{\text{variance-weighted ATT}} + \underbrace{VWCT}_{\text{variance-weighted CT}} + \underbrace{\Delta ATT}_{\text{Bias}}$$
 (10)

- VWCT = 0 if parallel trends assumption hold
- $\Delta ATT = \sum_{k \neq U} \sum_{\ell > k} \sigma_{k\ell} (1 \mu_{k\ell}) [ATT_k^{post(\ell)} ATT_k^{mid(k,\ell)}]$  is the bias term: **Negative weights** occur when already-treated units act as controls, *changes in their treatment effects over time* get subtracted
- Interpretating the TWFE estimator:
  - If treatment effects only vary across units:  $\Delta ATT = 0$ , but the weights may be far away from sample weights
  - ② If treatment effects only vary across time:  $\Delta ATT \neq 0$ , yielding implausible estimates

## Application

Impact of unilateral divorce law on female suicide rates (Stevenson and Wolfers, 2006)

- Unilateral (or no-fault) divorce allowed either spouse to end a marriage, redistributing property rights and bargaining power relative to fault-based divorce regimes
- Stevenson and Wolfers exploit "the natural variation resulting from the different timing of the adoption of unilateral divorce laws" in 37 states from 1969-1985
- Bacon replicates their study and provide new insights

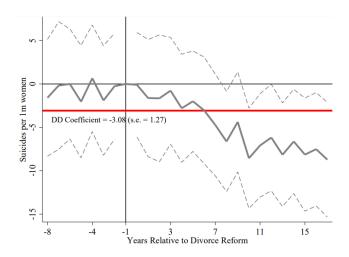


Figure 3: Event-Study and Difference-in-Differences Estimates

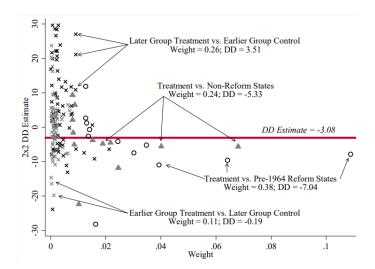


Figure 4: Bacon's Difference-in-Differences Decomposition for Unilateral Divorce and Female Suicide

Sensible Estimators for Staggered DID Designs

#### Sun and Abraham (2020, JOE)

- Consider group g units start to receive treatment at time  $g = 2, \dots, T$ .  $G_g \in \{0,1\}$  and  $G_{ig} \in \{0,1\}$  are group indicator and unit-group indicator
- **Parallel Trend Assumption**: For all  $t = 2, \dots, T$ , all  $g = 2, \dots, T$ ,

$$E[Y_t^0 - Y_{t-1}^0 | G_g = 1] = E[Y_t^0 - Y_{t-1}^0 | C = 1]$$
$$= E[Y_t^0 - Y_{t-1}^0]$$

 S&A proposes the following interaction-weighted estimator for ATT(g,t):

$$Y_{it} = \sum_{g=2}^{T-1} \sum_{e \neq 0} \delta_{ge} G_{ig} \mathbf{I}(t - G_i + 1 = e) + \lambda_i + \lambda_t + \varepsilon_{it}$$

- $\hat{\delta}_{ge}$  is consistent for ATT(g,t), t-g+1=e
- If no never treated group, dropping the last time period is unecessary
- Stata command: eventstudyinteract

#### de Chaisematin and D'Haultfoeuille (2020, AER)

- Parallel Trend Assumption as Sun and Abraham (2020)
- dC&D is interested in estimating instantaneous treatment effect:

$$\delta^{dc\&D} = E\left[\frac{\sum_{i}^{N} \sum_{t=2}^{T} G_{ig}(Y_{it}^{1} - Y_{it}^{0})}{\sum_{i}^{N} \sum_{t=2}^{T} G_{it}}\right]$$

$$\hat{\delta}^{dc\&D} = \sum_{g=2}^{T} \hat{P}(G_{g} = 1 | \text{Treated for period} \ge 1) \cdot \widehat{ATT}(g, g)$$

$$\widehat{ATT}^{ny}(g, t) = \underbrace{\sum_{i}^{T} G_{ig}(Y_{it} - Y_{ig-1})}_{\sum_{i}^{T} G_{it}} - \underbrace{\sum_{i}^{T} (1 - D_{it})(1 - G_{ig})(Y_{it} - Y_{ig-1})}_{\text{controls}}$$

- One needs to consider alternative causal parameters if interested in treatment effects dynamics
- Stata command: did\_multiplegt

## Callaway and Sant'Anna (2020, JOE)

- **Parallel Trend Assumption**: For all  $g, s, t = 2, \dots, T$  such that  $t \ge g$ , s > t:
  - (1) Never treated units:  $E[Y_t^0 Y_{t-1}^0 | G_g = 1] = E[Y_t^0 Y_{t-1}^0 | C = 1]$
  - (2) Not-yet-treated units:  $E[Y_t^0 Y_{t-1}^0 | G_g = 1] = E[Y_t^0 Y_{t-1}^0 | D_s = 1]$
- C&S consider two groups of ATT:

$$\begin{split} ATT^{never}(g,t) &= E[Y_t - Y_{g-1}|G_g = 1] - E[Y_t - Y_{g-1}|C = 1] \\ ATT^{ny}(g,t) &= E[Y_t - Y_{g-1}|G_g = 1] - E[Y_t - Y_{g-1}|D_t = 0, G_g = 0] \\ \widehat{ATT}^{ny}(g,t) &= \frac{\sum_i G_{ig}(Y_{it} - Y_{ig-1})}{\sum_i G_{ig}} - \frac{\sum_i C_i (1 - G_{ig})(Y_{it} - Y_{ig-1})}{\sum_i C_i} \end{split}$$

• Stata Command: csdid

## More on Staggered DID

- Borusyak, Jaravel, and Spiess (2021): Imputation based estimator
  - Stata command: did\_imputation
- Athey and Imbens (2022, JOE): standard DID estimator still works for randomly assigned adoption date and more .....
- Looking forward:
  - Level of treatment heterogeneity is somewhat limited: by cohort-time
  - Non-absorbing treatment: e.g., high-tech firms certification in China
  - Researchers may see benefits by combining different identification methods