TSCS Data Analysis

Ye Wang

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- ▶ We then introduce several approaches to fix the problems, such as counterfactual estimation.
- ▶ We next review available methods under sequential ignorability.
- We conclude by discussing the idea of manifold learning and how to deal with interference in TSCS data (hopefully!).

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- ▶ In the cross-sectional setting, we need ignorability: $D_i \perp Y_i(D_i) | \mathbf{X}_i$, $0 < P(D_i = 1 | \mathbf{X}_i) < 1$.
- Now we can relax the assumption along two possible directions (based on your belief).

▶ Strict exogeneity: $D_{it} \perp Y_{is}(D_{is})|\mathbf{X}_{i}^{1:t}, \mathbf{U}_{i}^{1:t}$, where $\mathbf{X}_{i}^{1:t}$ the history of observable confounders and $\mathbf{U}_{i}^{1:t}$ the history of unobservable confounders.

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- We always assume that SUTVA holds and there is no reversal causality (future treatments won't affect past outcomes).

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- ► The dataset available to the analyst only includes each unit's observable attributes, health status, treatment status, and the vaccination date.

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- You will be vaccinated on Feb. 15 with probability 0.72 if you are an old Asian male who were not infected by Covid last week.
- ► The dataset includes the observable attributes, the history of outcomes and treatment assignments of each unit, and the vaccination date.

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- ► All the classic methods (regression, weighting, matching, etc.) still apply with some modifications.
- ▶ Under strict exogeneity, we have the problem of omitted variables as part of the confounders $(\mathbf{U}_{i}^{1:T})$ is also unknown.
- ➤ To eliminate the influence of the unobservables, we need stronger assumptions on the DGP, such that the values of U_i^{1:T} can be learned from other variables.

► Note that the strict exogeneity assumption is satisfied by the following outcome model:

$$\begin{aligned} Y_{it} &= g_{it}(D_{it}, \mathbf{X}_{i}^{1:t}, \mathbf{U}_{i}^{1:t}) + \varepsilon_{it} \\ \mathbb{E}[\varepsilon_{is}|D_{it}, \mathbf{X}_{it}, \mathbf{U}_{it}] &= 0, \end{aligned}$$

which is too general for identification.

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▶ For example, we can assume that $h_{it}(\mathbf{U}_{it}) = \mu + \alpha_i + \xi_t$, then

$$Y_{it} = \mu + \delta_{it}D_{it} + \mathbf{X}_{it}\beta + \alpha_i + \xi_t + \varepsilon_{it}.$$

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Now, the assumption of strict exogeneity becomes: $\mathbb{E}[\varepsilon_{is}|D_{it},\mathbf{X}_{it},\alpha_i,\xi_t]=0$ for any s.

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- ▶ We no longer have consistent estimates if treatment effects are heterogeneous.
- ▶ There are several solutions to the problem.
- ▶ It is easier to fix the problem under the DID setting (Strezhnev, 2017).
- Otherwise, we need to modify the estimand or impose stronger assumptions.

 Chaisemartin and D'Haultfœuille (2020) advocate that we should focus only on the instant effects generated by the treatment and estimate

$$\frac{1}{N^*} \sum_{t \geq 2, D_{it} \neq D_{i,t-1}} \delta_{it}$$

where
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- ▶ This estimand is well-defined and can be consistently estimated.
- Yet it throws away a ton of information as well thus is not very efficient.
- ► There are more efficient approaches which require stronger assumptions on model specification.

➤ Xu (2017) and Liu, Wang, and Xu (2020) combine factor models with the Neyman-Rubin framework

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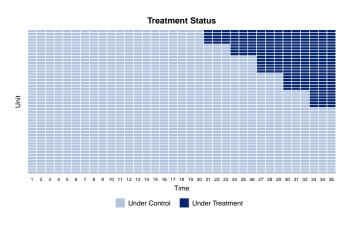
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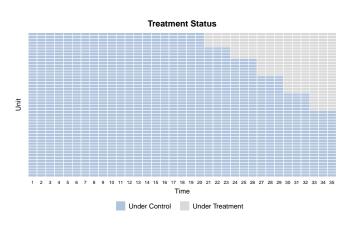
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- lacktriangle Clearly, $\hat{\delta}_{it}=Y_{it}-\hat{Y}_{it}(0)$ and

$$\widehat{ATT} = \frac{1}{|\mathcal{M}|} \sum_{(i,t) \in \mathcal{M}} \hat{\delta}_{it}$$

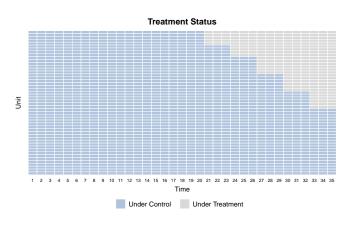
▶ In a TSCS setting, treat Y(1) as missing data



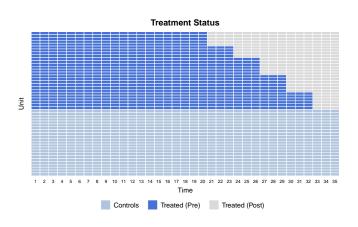
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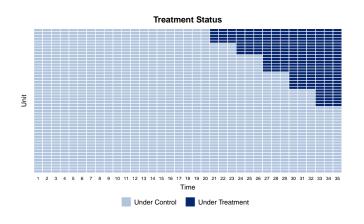
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- ▶ (Use pre-treatment data for model selection)
- ▶ Estimate ATT by averaging differences between Y(1) and $\hat{Y}(0)$



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- The model specification has to be correct:
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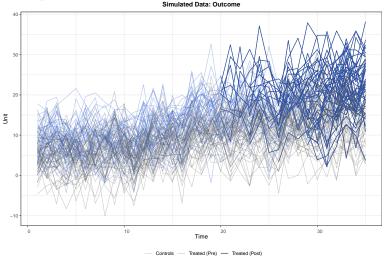
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- ▶ It also requires strict exogeneity, thus excludes temporal interference.

One advantage of the framework is that testing the identification assumption becomes straightforward.

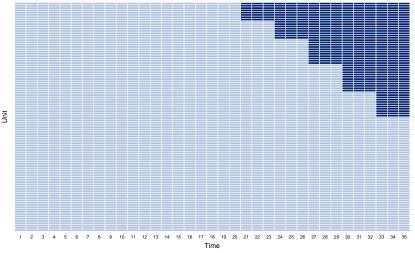
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- ► A plot of dynamic effects.
- ► A placebo test: estimate treatment effects before the treatment's onset and test their significance.
- An equivalence test: test whether all the pre-treatment ATTs are equal to zero using TOST.

Let's consider a DGP with two factors (hence FEct will be biased).

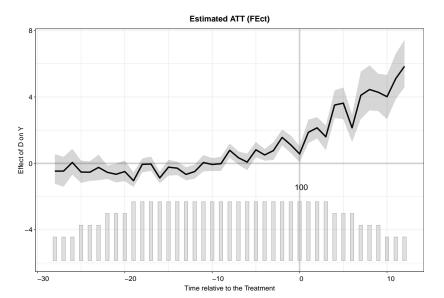


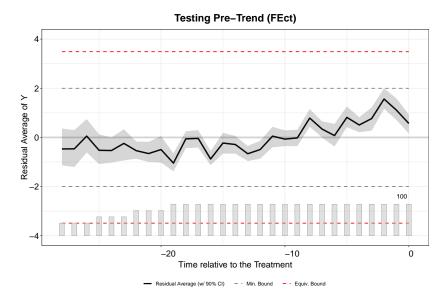
Simulated Data: Treatment Status



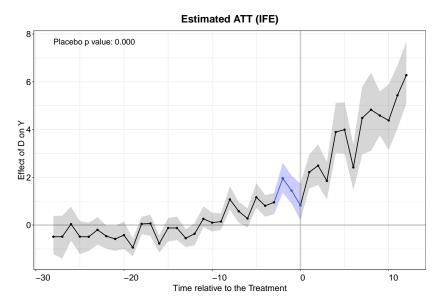
Under Control Under Treatment

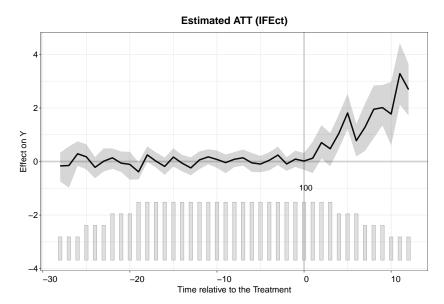
Bootstrapping for uncertainties100 runs

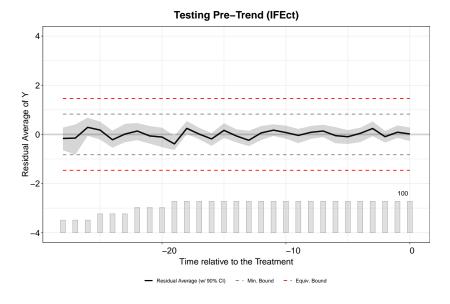




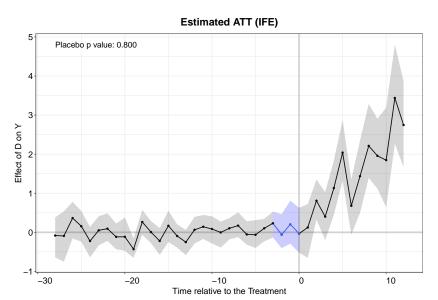
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- Modern approach: IPTW estimators (MSMs) and doubly robust estimators.
- Notice that the propensity score for unit i in period t has the form of $P(D_{it}|\mathbf{Y}_i^{1:(t-1)},\mathbf{X}_i^{1:t})$.

► Suppose there is no interference and we are interested in the contemporary effect generated by the treatment:

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$$\hat{\tau}_t = \frac{1}{N} \sum_{i=1}^{N} \frac{D_{it} Y_{it}}{e_{it}} - \frac{1}{N} \sum_{i=1}^{N} \frac{(1 - D_{it}) Y_{it}}{1 - e_{it}}.$$

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It is just another Horvitz-Thompson estimator.

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- Hazlett and Xu (2018) suggest that we may instead balance kernels of the history.
- The implicit assumption is that the distribution of the outcome history can be well approximated by its kernels.

▶ We search for a group of positive weights $\{w_i\}$ which sum to one such that

$$\frac{1}{|\mathcal{T}|} \sum_{i \in \mathcal{T}} \mathbf{K}_i = \sum_{i \in \mathcal{C}} w_i \mathbf{K}_i$$

where $\mathbf{K}_i = (k(\mathbf{Y}_i, \mathbf{Y}_1), k(\mathbf{Y}_i, \mathbf{Y}_2), \dots, k(\mathbf{Y}_i, \mathbf{Y}_N)), \mathbf{Y}_i$ is the pre-treatment outcome history of unit i, and

$$k(\mathbf{Y}_i, \mathbf{Y}_i) = e^{-||\mathbf{Y}_i - \mathbf{Y}_j||^2/h}.$$

Next, we estimate the treatment effect on the treated in each period

$$\hat{ au}_t = \frac{1}{|\mathcal{T}|} \sum_{i \in \mathcal{T}} Y_{it} - \sum_{i \in \mathcal{C}} w_i Y_{it}$$

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- ► Kim, Imai, and Wang (2019) show that we can do the same via matching.

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- ▶ They show that now $\mathbf{D}_i^T \perp \mathbf{Y}_{jt}(\mathbf{D}^T)|\frac{1}{T}\sum_{t=1}^T D_{it}$.
- ▶ $S_i = \frac{1}{T} \sum_{t=1}^{T} D_{it}$ is a sufficient statistic for α_i .

Arkhangelsky and Imbens (2019) further illustrate that under this assumption, all we need to do is to find a group of weights (again!) such that:

$$\begin{split} \{\hat{w}_{it}\} &= \arg\min\frac{1}{NT} \sum_{i=1}^{N} \sum_{t=1}^{T} \hat{w}_{it}^{2}, \\ \text{s.t. } \frac{1}{NT} \sum_{i=1}^{N} \sum_{t=1}^{T} \hat{w}_{it} D_{it} = 1, \frac{1}{N} \sum_{i=1}^{N} \hat{w}_{it} = 0, \frac{1}{T} \sum_{t=1}^{T} \hat{w}_{it} = 0, \\ \frac{1}{NT} \sum_{i=1}^{N} \sum_{t=1}^{T} \hat{w}_{it} \psi_{it}(S_{i}) &= 0, \hat{w}_{it} D_{it} \geq 0. \end{split}$$

▶ Then, the estimate of the ATE,

$$\hat{\tau} = \frac{1}{NT} \sum_{i=1}^{N} \sum_{t=1}^{I} \hat{w}_{it} Y_{it}$$

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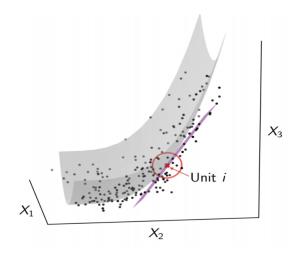
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- ▶ As long as we can learn the value of λ_i from Y_{it} , all the classic techniques can be applied.
- ▶ To estimate λ_i , we first match each unit i to K nearest neighbors.
- ▶ Next, we estimate λ_i via PCA on the K+1 outcome histories.

Figure 2: Local Tangent Space Approximation



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- Treatment assignments are usually temporally dependent in panel data.
- ▶ Now, past treatments become confounders.
- ▶ We have to control their influence under either assumption.

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- Otherwise, there is one more reason for the invalid second difference to occur.
- ▶ We can still estimate the quantity defined by Chaisemartin and D'Haultfœuille (2020), but nothing else.
- Sequential ignorability is the more suitable assumption if we have temporal interference

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- We can obtain valid estimate for the history's aggregated effect after weighting observations properly.
- Now,

$$P(\mathbf{D}_{i}^{s:t} = \mathbf{d}^{s:t}) = \prod_{s'=s}^{t} P(D_{is'}|\mathbf{D}_{i}^{1:(s'-1)}, \mathbf{Y}_{i}^{1:(s'-1)}, \mathbf{X}_{i}^{1:s'}).$$

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- Wang (2020) shows that the limit of the DID estimator has no substantive interpretation under spatial interference.
- ▶ Intuitively, the control group is contaminated and we cannot eliminate the bias via outcome adjustment.
- Yet under sequential ignorability, we can combine the idea in Aronow, Samii, and Wang (2020) and MSMs to generate consistent estimates for both the direct and the indirect effects.

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