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AT URBANA-CHAMPAIGN

LEARNING & LABOR

Matching Estimators



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Outline

- Why matching and what is needed?
- Types of matching
- How to choose X 's and run specification tests
- Matching and Difference-in-Difference
- Bounding Bias
- Practical Issues



Why Match and What is Needed?



Why Match?

- Randomization is not always feasible
 - ...and doesn't always work
- Selection bias: program bias, administrative bias
- E.g. protected area established based on criteria on elevation, slope and species habitat
- E.g.2 Irrigation intervention targeted at vulnerable households in semi-arid areas



The Problem

- We want to estimate the effect of a policy that targets a specific area



Matching



- Or, more likely...
- Find a set of ‘matched’ control (untreated) parcels with similar characteristics to the treated parcels
- If we do a good job finding control parcels that look like the treated parcels, we can replicate what would have happened without protection*

*Assumes unobservable characteristics are distributed in the same way as observables: i.e. Matching over observables = matching over unobservables



Why Not Just Run a Regression?

- Can still control for covariates – but some characteristics may have a non-linear effect
- Often is a matter of throwing away ‘bad’ controls



Assumptions Needed

- Conditional Independence Assumption (CIA)
 - Once you've controlled for the observable X 's, you've controlled for selection (either X 's explain everything, or U 's are distributed in the same way as the X 's)
- Common Support Assumption
 - Enough control observations have characteristics in the same range as the treated observations
 - E.g. if irrigation scheme targets all high elevation land in an eco-region, one might not be able to find good controls
 - E.g.2 if new sustainable intensification schemes target all farm households with less than 1 ha, might have difficulties finding controls



Types of Matching



Types of Matches

- Characteristic matching (nn matching; mahalanobis matching)
- Propensity Score Matching
- Exact matching (stratified matching)
- Caliper matching
- Kernel matching



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i	T	Edu	Income
1	0	2	60
2	0	3	80
3	0	5	90
4	0	12	200
5	1	5	100
6	1	3	80
7	1	4	90
8	1	2	70

From Heinrich et al 2010



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i	T	Edu	Income	Match	Yi	Yo	Diff
1	0	2	60				
2	0	3	80				
3	0	5	90				
4	0	12	200				
5	1	5	100	[3]	100	90	10
6	1	3	80	[2]	80	80	0
7	1	4	90	[2,3]	90	85	5
8	1	2	70	[1]	70	60	10

ATT = 6.25



Dimensionality Issue

- Can easily match over only one variable (dimension)
- Once one has multiple variables, one needs to weight them and create an index
- What is the appropriate weighting scheme?



Characteristic Matching

- Based on X 's, where they are weighted by inverse variance

X_1 slope, $\sigma^2 = 5$

X_2 wealth, $\sigma^2 = 700$



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- Mahalanobis: Proximity to the center of the mass of the treated



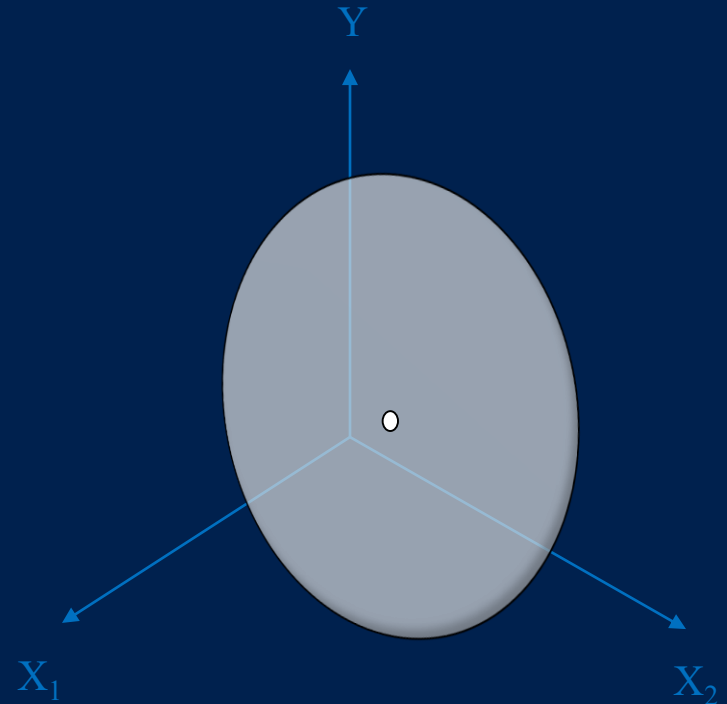
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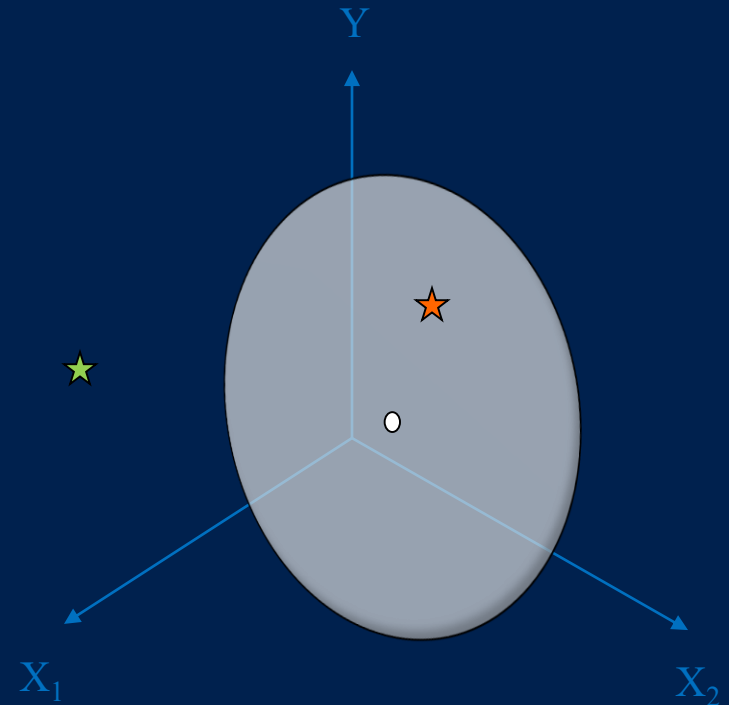
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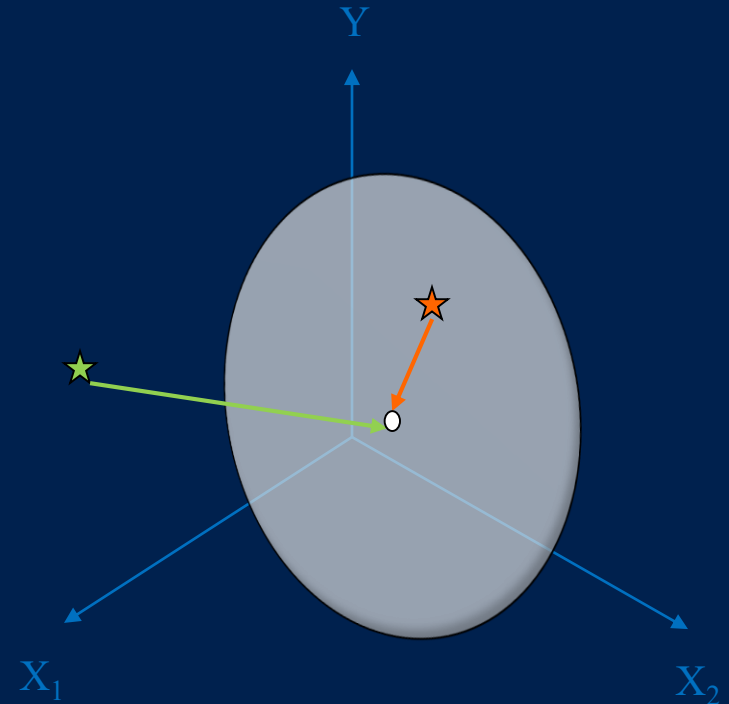
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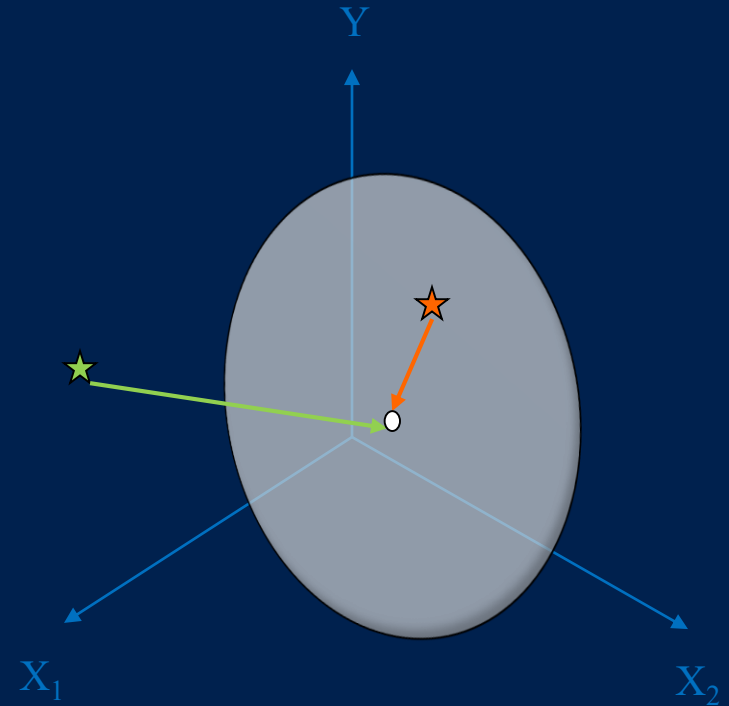
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X_1 slope, $\sigma^2 = 5$

X_2 wealth, $\sigma^2 = 700$

- Mahalanobis: Proximity to the center of the mass of the treated
- Different ways of measuring 'distance'



Propensity Score Matching

- Measure ‘distance’ by probability of treatment
- First estimate the probability of treatment given observable characteristics using a probit or logit
- Then predict that probability for all treated and control observations
- Use these predicted probabilities as ‘coordinates’ to allocate which controls are near to which treatment observations



Common support



Exact Matching

- Forcing the control observations to be exactly the same over some dimension
 - E.g. sex; ecoregion



How to define a 'nearby' control?



- 1) Pick 'n' closest observations in terms of propensity score, weighted equally: $\frac{1}{n}$
- 2) Pick a certain distance and include all observations within that distance
- 3) Weight observations using a kernel

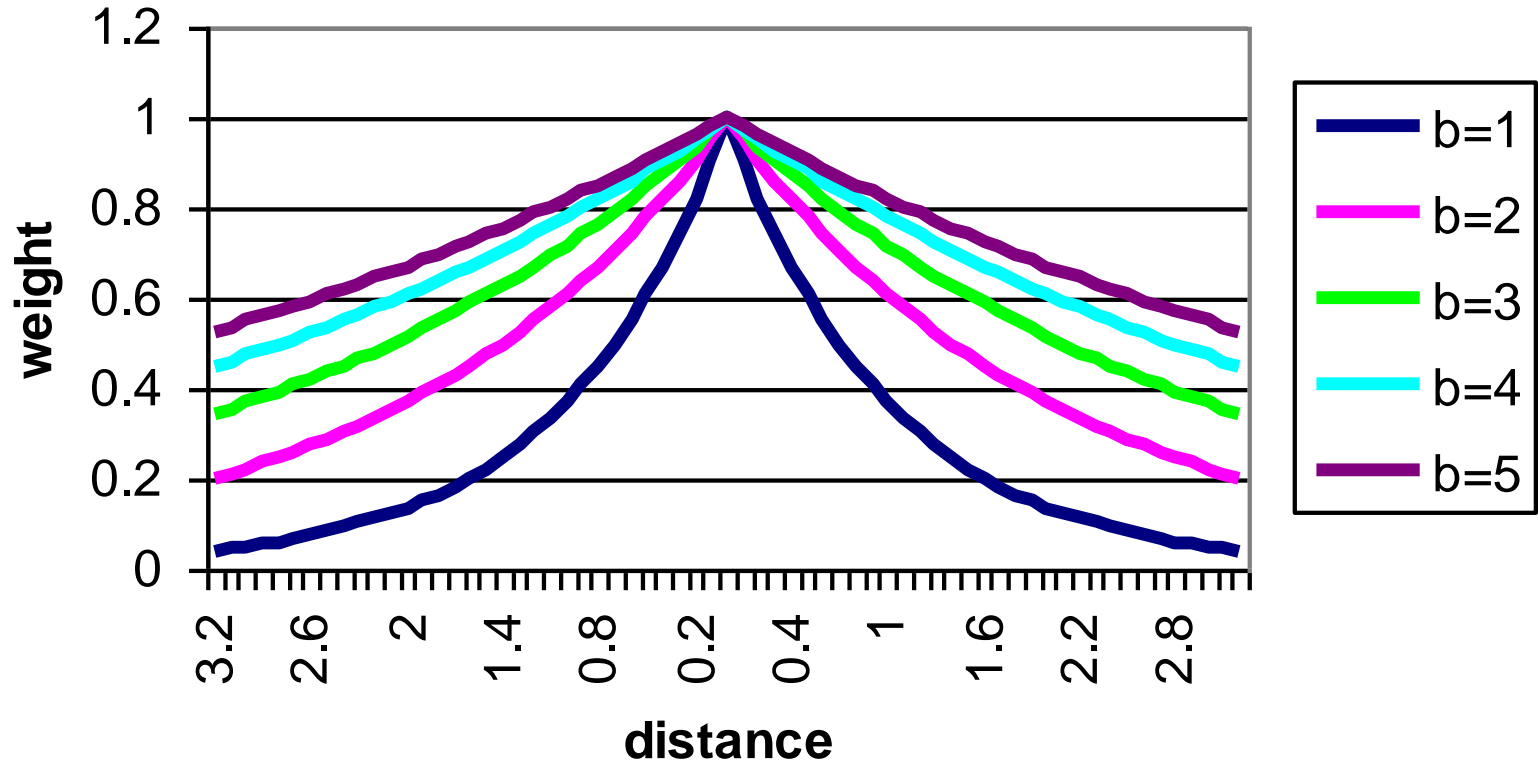


Kernel matching

- weight each observation based on its ‘distance’
- $\omega_{ij} = \left[1 - \left(\frac{\delta_{ij}}{d_i} \right)^3 \right]^3 I(\delta_{ij} < d_i)$
- Where δ_{ij} is the distance between i and j
- d_i is the distance to the q^{th} nearest neighbor
- q is often referred to as the ‘window size’
- Alternatively, $\omega_{ij} = \Phi \left[\frac{\delta_{ij}}{s_i b} \right]$
- where s is the std. dev. of distances between i and all others, and b is the bandwidth



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Effect of bandwidth on distance weights



Matching Options

- With or without replacement
- Weighted
- Efficiency vs bias trade-off



How to Choose X 's and Run Specification Tests



How to Choose Covariates?

1. Things that affect selection into treatment
2. Things that affect outcome
 - If don't condition on them in the estimation
- Regressions on treatment and on outcome
- Do not use characteristics that might be affected by treatment
 - Consider spillovers



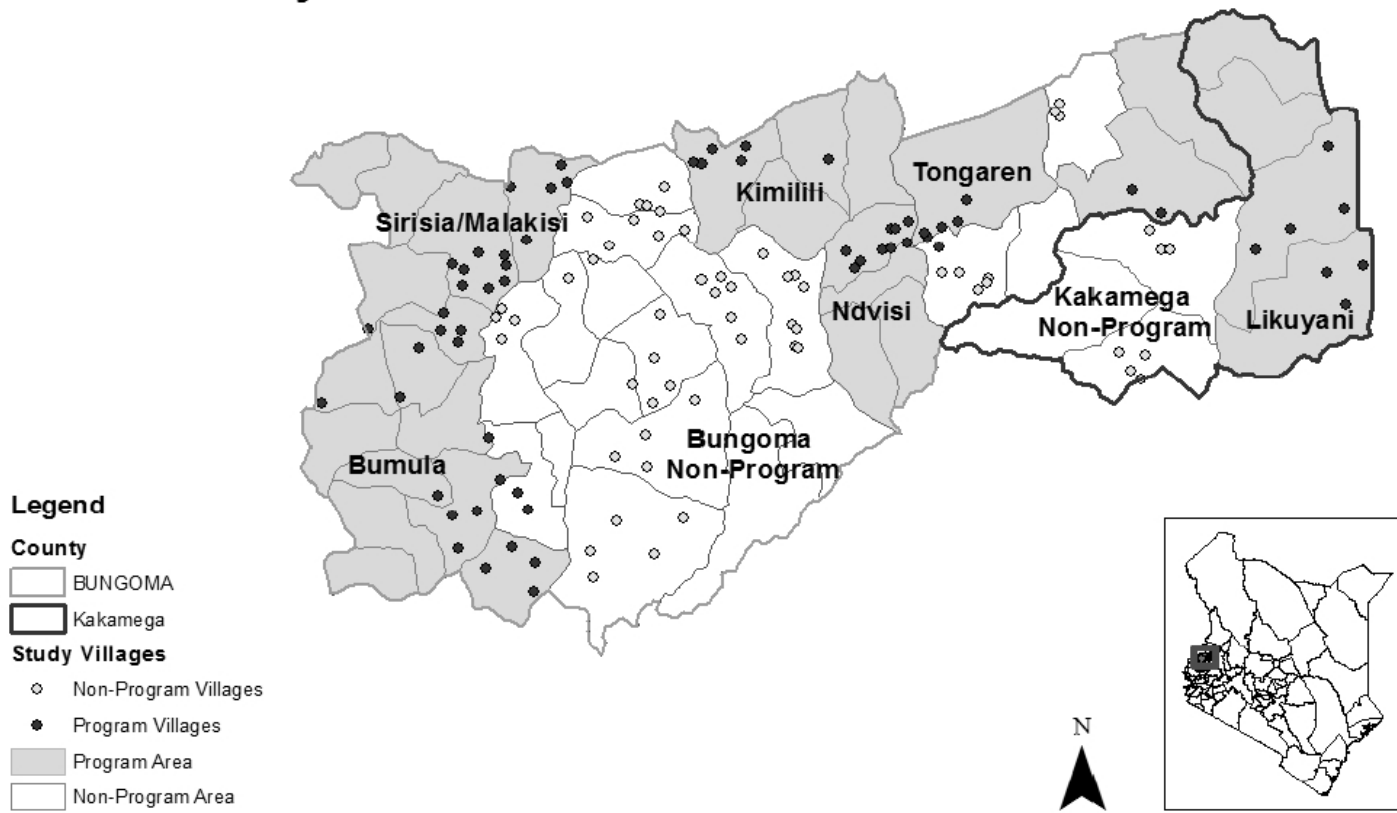
Things to Keep in Mind...

- Unlike a regression, adding in a covariate that doesn't affect selection and/or outcome can matter
- Ensure data for controls and treatment are from the same source (same frequency/granularity; same probability of missing etc)
- Heterogeneity analysis – may want to match over subsets (exact matching)
 - E.g. gender in PSM



Pre-Survey Matching

Study Area
Bungoma & Kakamega Counties
Western Kenya



Administrative boundaries compiled by Daniel K. Thompson, Emory University, available at mapeastafrica.com.
Project boundaries provided by Vi Agroforestry
Prepared by Seth Morgan, the University of Illinois at Urbana-Champaign

0 5 10 20 Kilometers

What if we don't want to throw away
data?

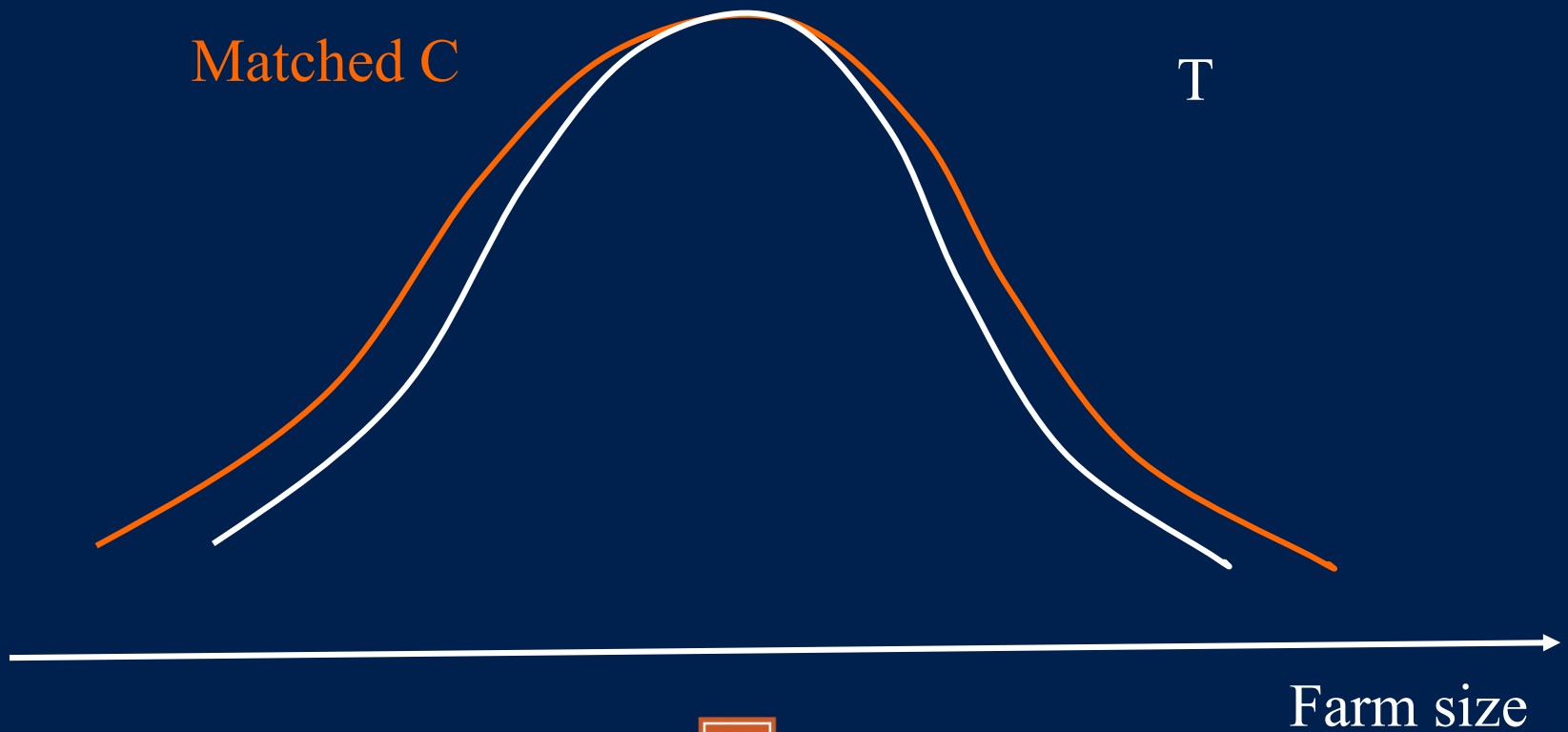


Specification Tests

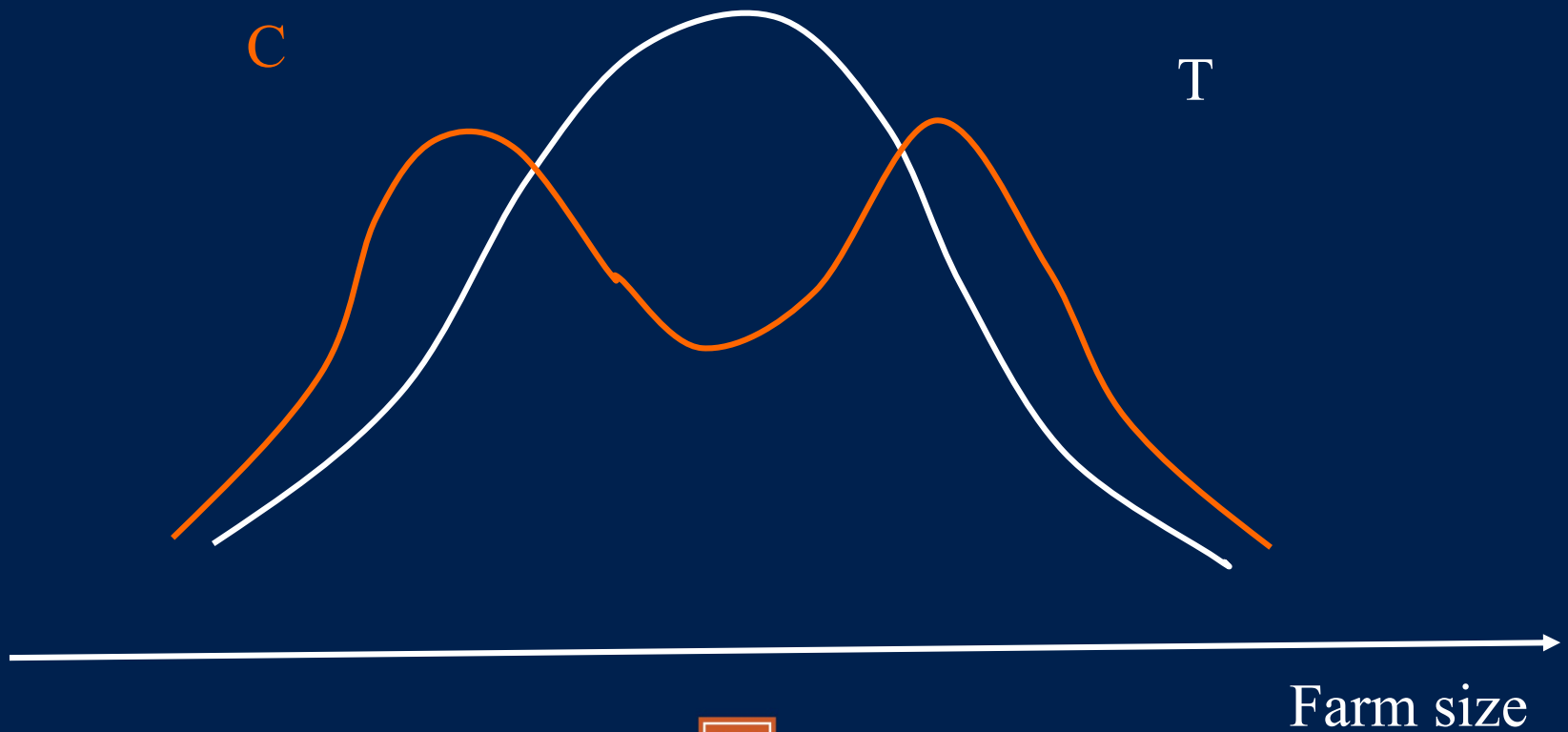
- Covariate balance test
 - Including distribution
- Multiple algorithms for robustness
 - Test to examine if results are sensitive to a few bad matches
- Visual inspection of propensity scores (before and after)
- Mapping



Checking X's



But maybe...



Matching and Difference-in-Difference



DiD Matching

- Matching can be used in a DiD framework
- Control for time-invariant unobservables
- (Heckman Ichimura and Todd 1997; Heckman et al 1998)



Other options

- Matching with Continuous Treatment (Imbens 2000)
- Matching with a roll-out design



Measuring the degree of potential bias



Rosenbaum Bounds

- Rosenbaum (2002): Identify “hidden bias” from unobservable covariates
- Ask how much unobservables might affect results (make the ATT insignificant)
- Specifically, estimates an odds ratio of how much could an unobserved variable bias outcome by affecting selection

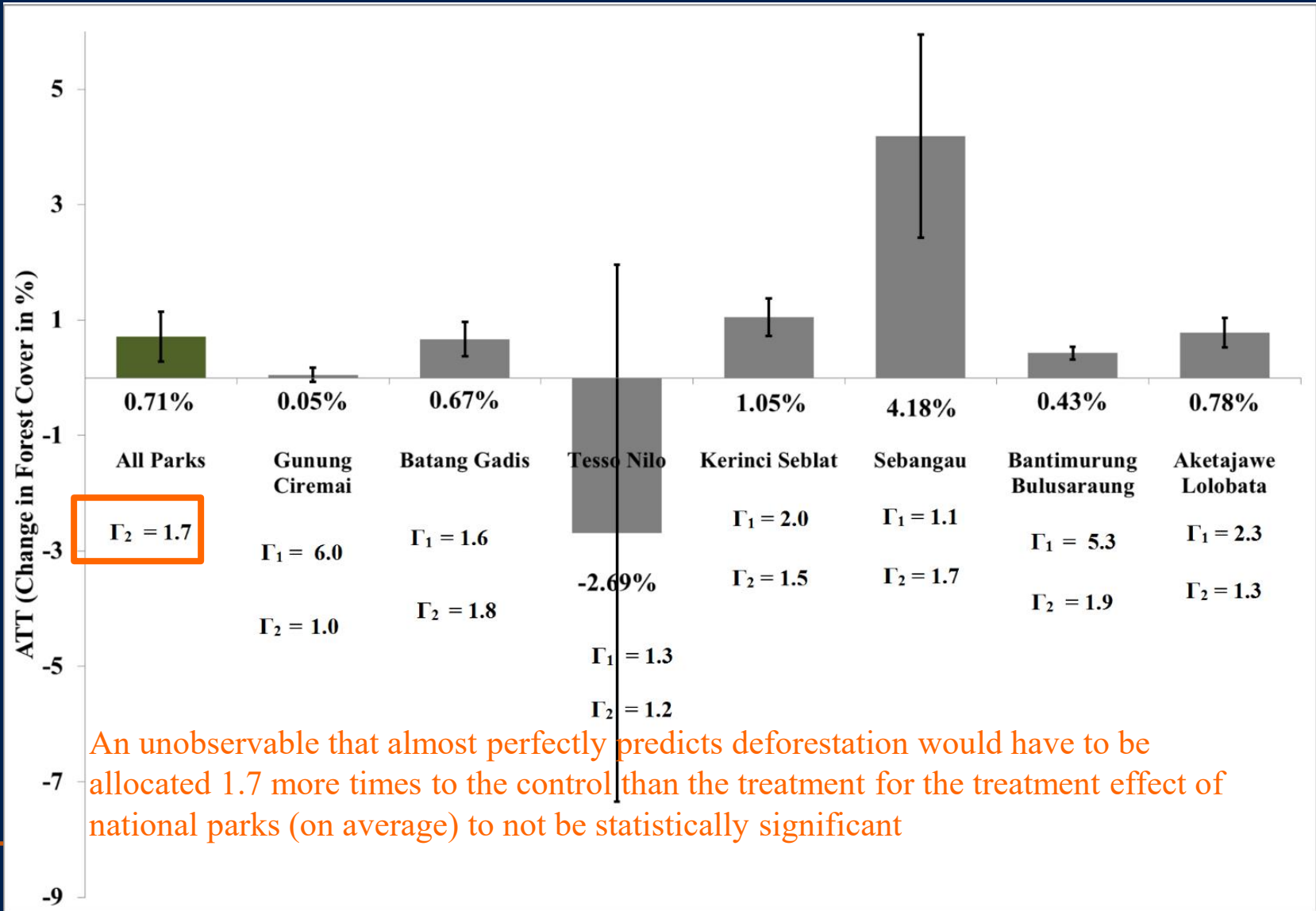


e.g. Shah and Baylis

- Comparing effect of parks across Indonesia
 - Do unobservable covariates affect whether individual park ATT is different from the national ATT (Γ_1)
 - Do unobservable covariates affect the park level ATT estimates (Γ_2)



Test for Hidden Bias



Interpretation of the Γ

- Lalonde (1985): effect of job training on wages
- match on age, education, race, marital status, high school degree, earnings for the two years before the training program and unemployment before the training program.

ATT: 1767.7 (830.85); p.stat = 0.033; $\Gamma = 1.05$



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```
> #Sensitivity Tests
> psens(mgen1, Gamma=1.5, GammaInc=.1)
Rosenbaum Sensitivity Test for Wilcoxon Signed Rank P-Value
      Gamma L. Bound P-Value U. Bound P-Value
[1,]    1.0      0.0346      0.0346
[2,]    1.1      0.0062      0.1271 ←
[3,]    1.2      0.0009      0.3000
[4,]    1.3      0.0001      0.5164
[5,]    1.4      0.0000      0.7139
[6,]    1.5      0.0000      0.8539
```

If the odds of a person being in the training program are only 1.1 times higher b/c of an unobservable that affects income, the p-value as high as 0.127



Can also bound the estimated treatment effect

Rosenbaum Sensitivity Test for Hodges-Lehmann Point Estimate

	Gamma	L. Bound	HL Est.	U. Bound	HL Est.
[1,]	1.0	1194.000000		1194.0	
[2,]	1.1	560.780000		1231.2	
[3,]	1.2	274.080000		1598.4	
[4,]	1.3	-0.015006		1944.9	
[5,]	1.4	-113.220000		2218.3	
[6,]	1.5	-333.220000		2424.8	

← Median effect size if
no difference in
unobservables

← If people are 1.5 times
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Practical note: most socio-economic studies have $\Gamma < 2$.



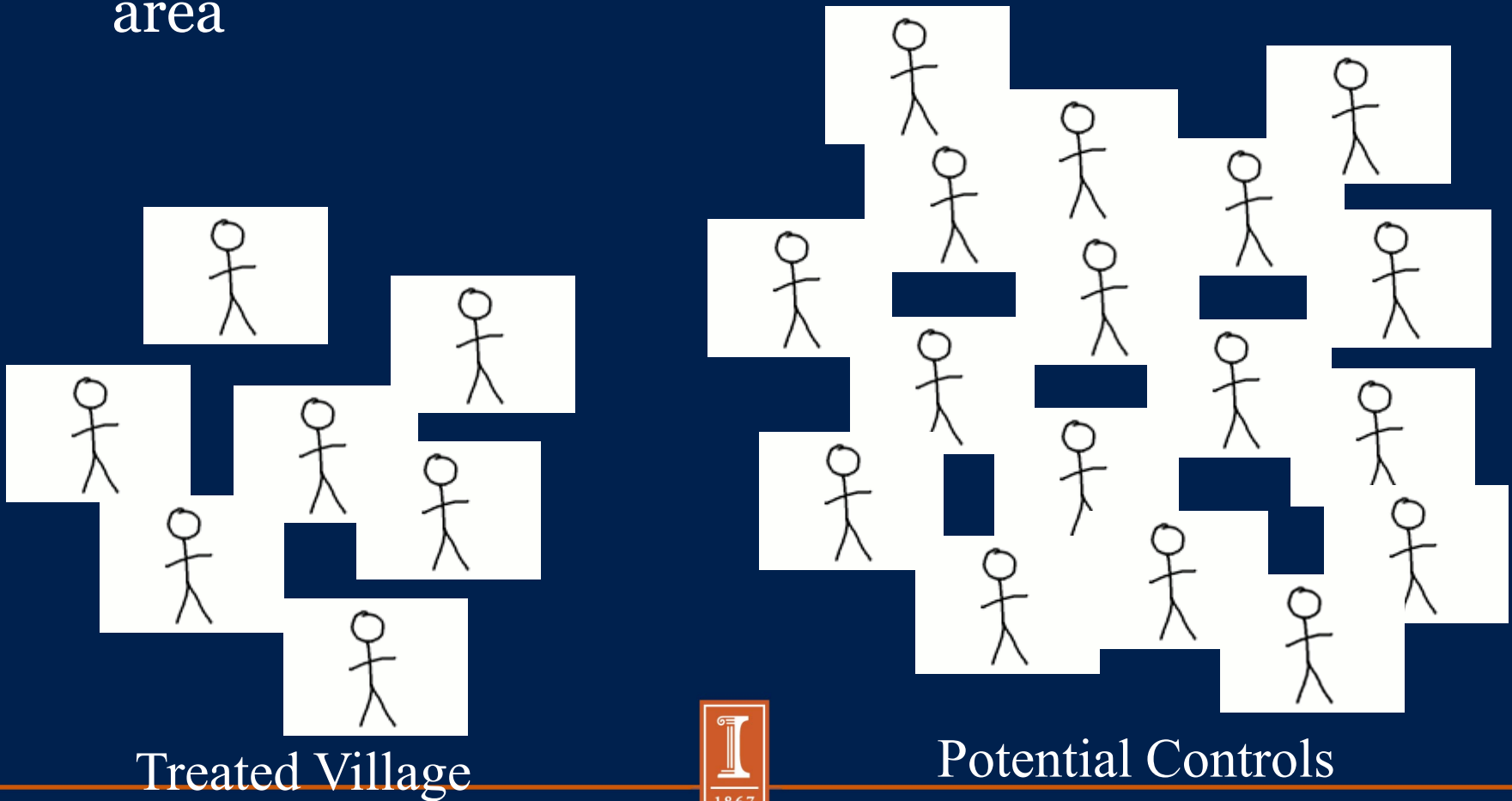
Practical Issues

- Unit of observation
- Number of treated vs control
 - Can we afford to throw out controls?
- Unobservables
- Spillovers (contaminated controls)
- Different data sources (what happens when you observe treated obs at greater granularity than controls?)



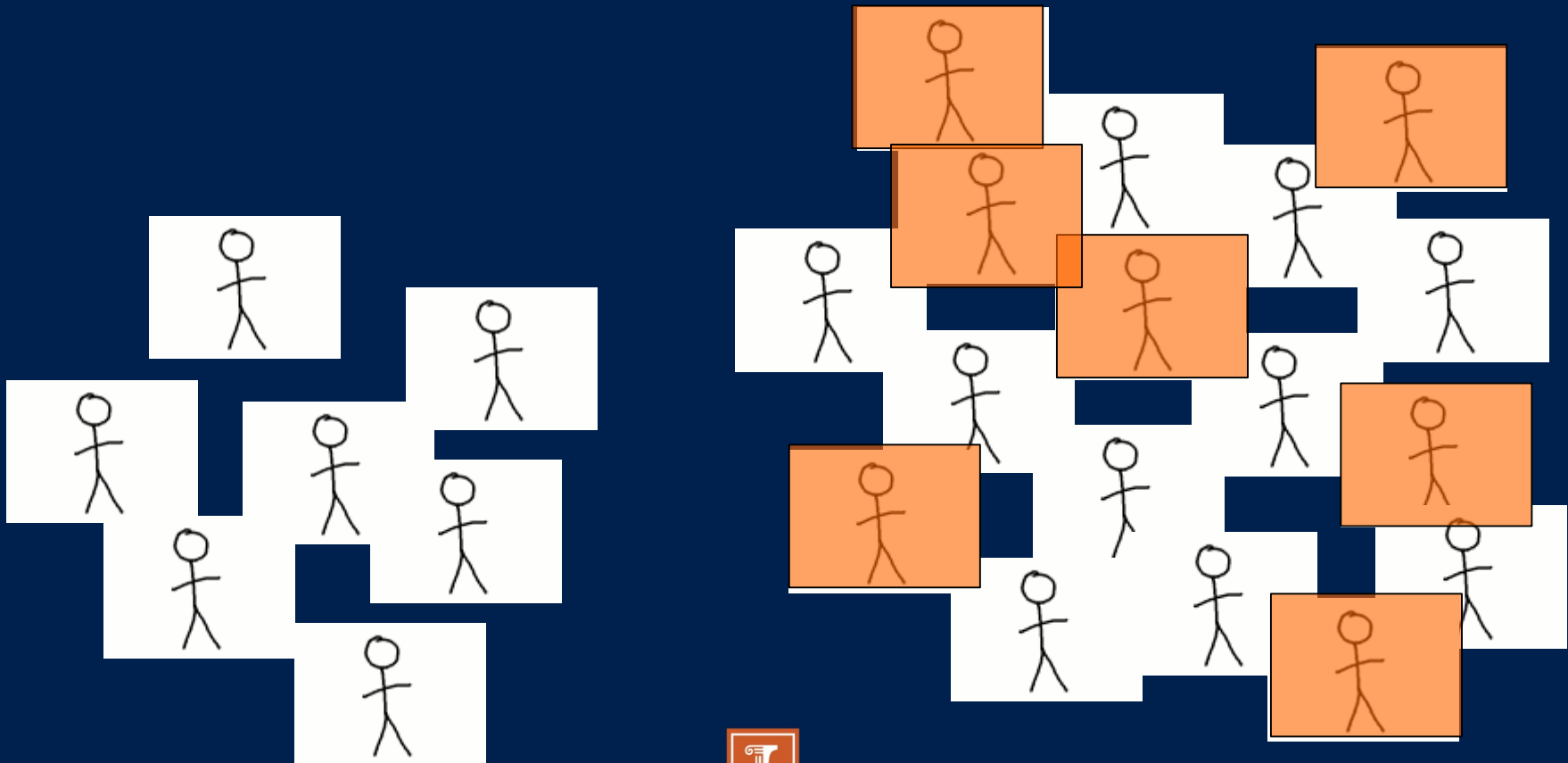
example

- High-yielding seed variety introduced to a limited area



example

- 1 to 1 matching



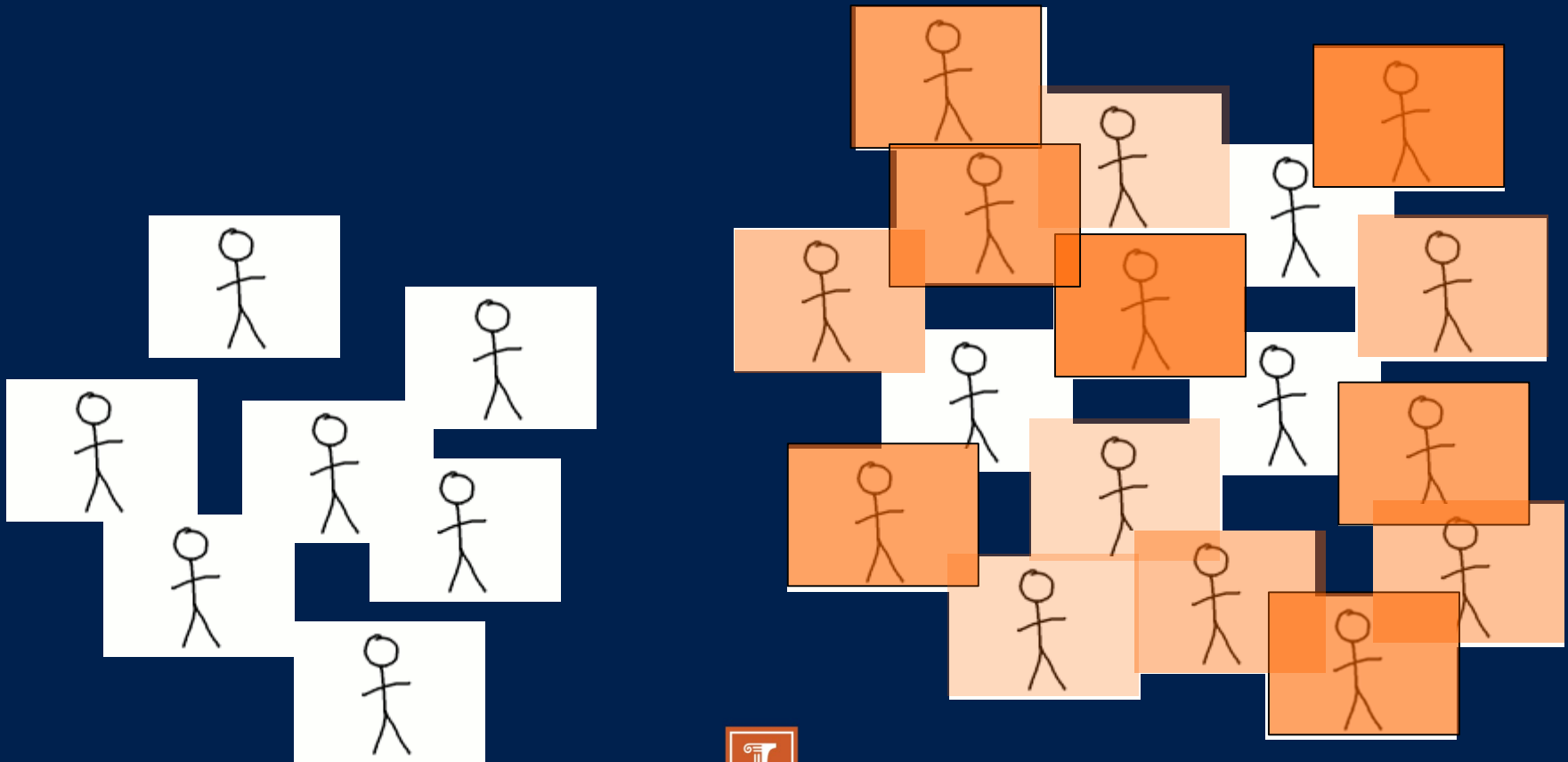
Treated Village



Matched Controls

example

- Kernel matching



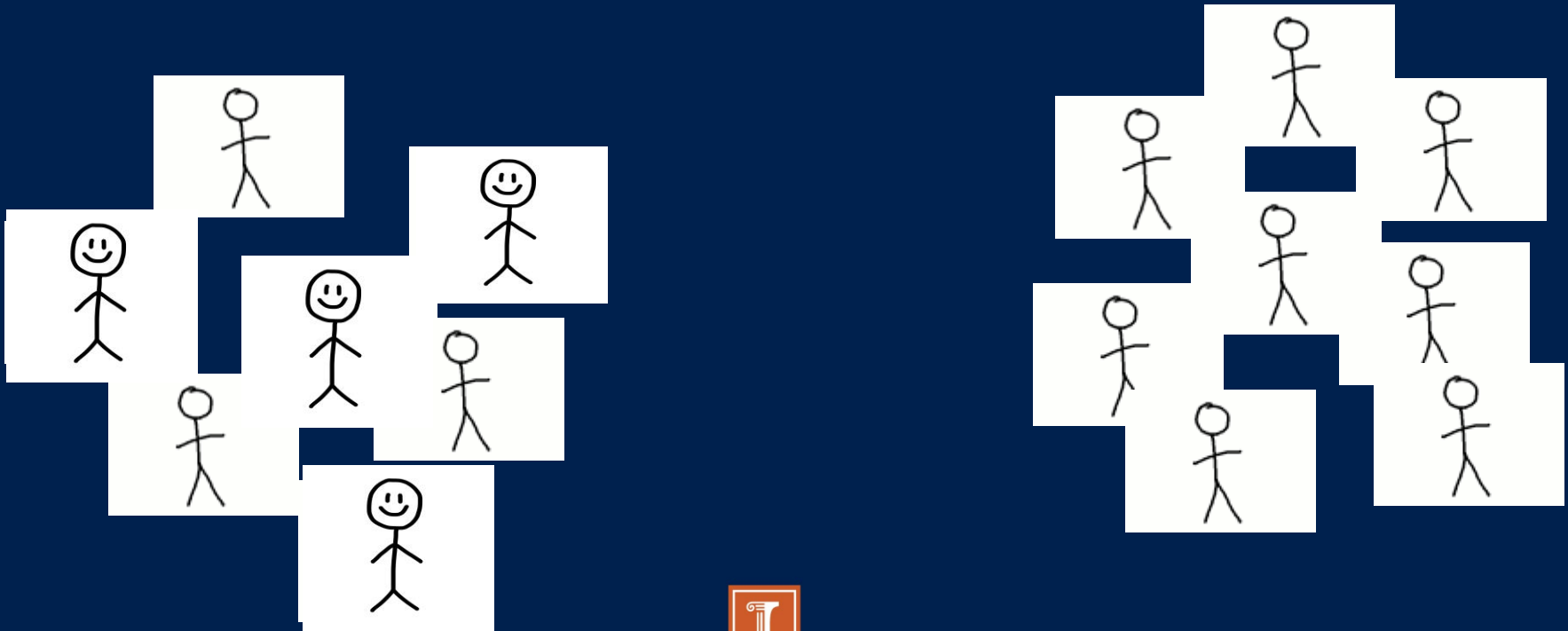
Treated Village



Weighted Matched
Controls

example

- Now assume only some people adopt



example

- PSM

