



LUND
UNIVERSITY

Matching and synthetic controls

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2021 ClimBEco course



Causal Inference from observational data

Introduction

Matching

Synthetic Controls

References

Synopsis: Today, we will be looking into methods that help us find (aka *match*) or simulate (aka *synthesize*) a control group for inferring causal effects from observational data, and its recent developments

In particular, we will develop an understanding of



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- matching approaches
- synthetic controls
- machine-based learning methods



Intuition

Consider a situation where the untreated are very different from the treated:

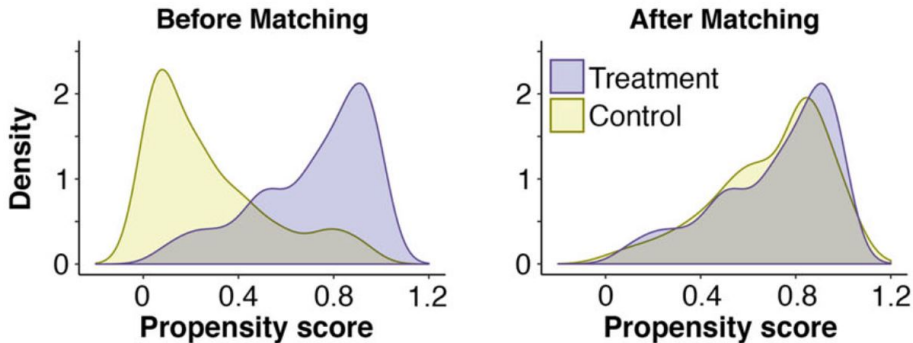


Image source: Schleicher et al. 2020



basic conditions

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The classical overarching conditions for robust causal inference:

- stable unit treatment value assumption (SUTVA)
 - treating one individual unit does not affect another's (potential) outcome
 - treatment is comparable [no (strong) variation in treatment]



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- unconfoundedness (strong ignorability)
 - $(Y(1), Y(0)) \perp T$: treatment assignment is independent of the outcomes
 - i.e. no omitted variable bias (recall the storch example)
 - or, at least, conditional unconfoundedness $(Y(1), Y(0)) \perp T | X$



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→ $\pi(X_i) = Pr(D_i = 1 | X_i)$ or *propensity score* can be used for matching



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- $\pi(X_i) = Pr(D_i = 1 | X_i)$ or *propensity score* can be used for matching
- but should maybe not (King and Nielsen 2019), we will see alternatives



Overview

Here is a general overview of possible matching methods

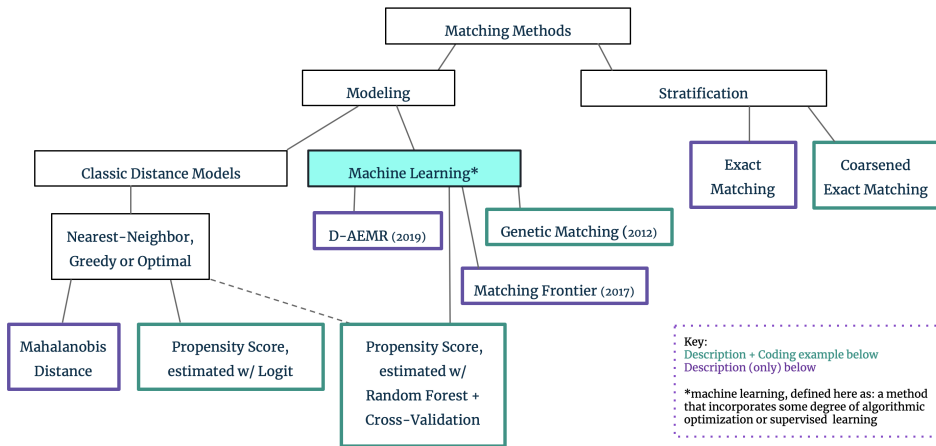


Image source: [Sizemore and Alkurdi 2019](#)



References I

- King, Gary and Richard Nielsen (2019). 'Why Propensity Scores Should Not Be Used for Matching'. In: *Political Analysis* 27.4, pp. 435–454. ISSN: 14764989. DOI: 10.1017/pan.2019.11.
- Schleicher, Judith et al. (2020). 'Statistical matching for conservation science'. In: *Conservation Biology* 34.3, pp. 538–549. ISSN: 15231739. DOI: 10.1111/cobi.13448.

