

# Welcome to the Sentinel Innovation and Methods Seminar Series

The webinar will begin momentarily

Please visit [www.sentinelinitiative.org](http://www.sentinelinitiative.org) for recordings of past sessions and details on upcoming webinars.

Note: closed-captioning for today's webinar will be available on the recording posted at the link above.

The Sentinel logo features the word "Sentinel" in a white serif font, with a white curved line arching over the letters "e" and "l".



# Can We Train Machine Learning Methods to Outperform the High-Dimensional Propensity Score Algorithm?



M. Ehsan. Karim; UBC

2022 Sentinel Innovation and Methods Seminar Series

May 11, 2022

# Outline

Slides at [tinyurl.com/hdps2022](https://tinyurl.com/hdps2022)

## 1. hdPS

- Basic terminology

## 2. Machine learning-based hdPS

- [Karim et al. 2018](#) Epidemiology
- Joint work with
  - Menglan Pang and Robert W Platt
  - McGill, CNODES Methods; Fund CIHR, Grant #DSE – 146021
- General idea

## 3. Related research

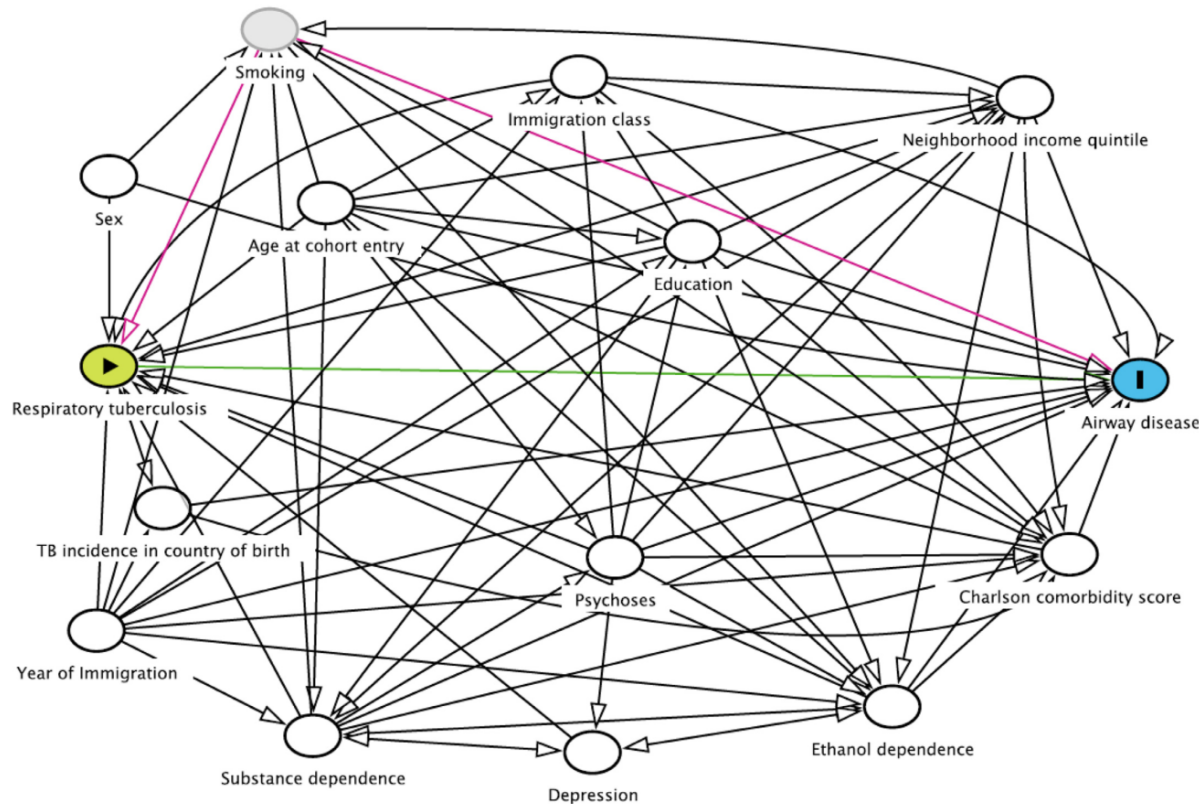
- Not exhaustive

hdPS



# Motivating Example

[Basham et al. 2021](#) EClinicalMedicine: [CC BY license](#)



Healthcare claims data for immigrants to British Columbia, Canada, 1985–2015

# Health care database: Advantages vs Disadvantages

1. Larger **sample size**;
  2. Diverse population;
  3. **Longitudinal records** /many years;
  4. **Detailed** health encounters, comorbidity, drug exposure history;
  5. possibility to **link** other databases.
1. Not specifically designed for answering a particular **research question**;
  2. **Data sparsity**: relies on visits and encounters;
  3. No control over which factors were measured.

TLDR: **May not have all confounders.**

# How to select adjustment variables?

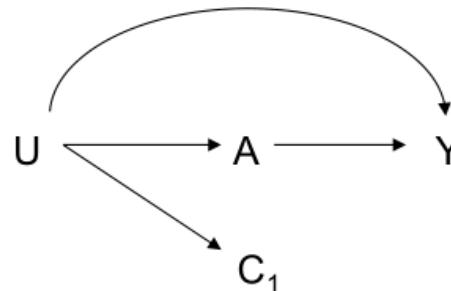
## Modified disjunctive cause criterion

Adjust for variables that are

- causes of exposure or outcome or both,
- discard: known instrument,
- including **good proxies** for unmeasured common causes

[VanderWeele et al. 2019](#) European Journal of Epidemiology: [CC BY license](#)

- $U$  =  
Smoking
- $C_1$  =  
Tobacco use

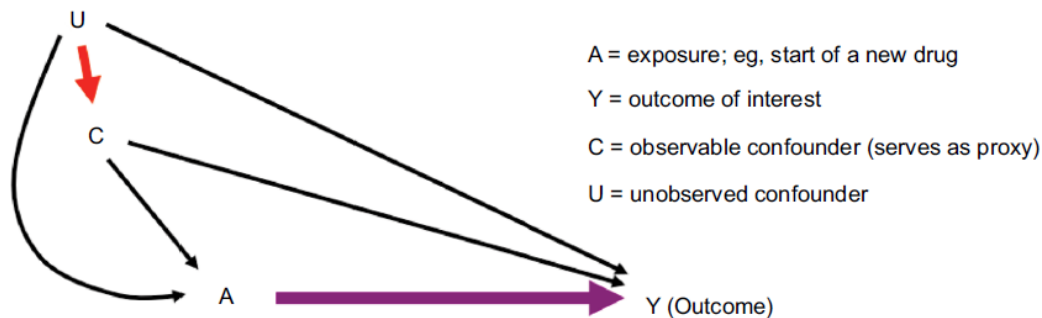


**Fig. 5** Control for a proxy confounder  $C_1$  of the true unmeasured confounder  $U$  will often, but not always, reduce confounding bias in the relationship between exposure  $A$  and outcome  $Y$

# Proxy information in Admin data

[Schneeweiss et al. 2018](#) Clinical Epidemiology: [CC BY NC license](#)

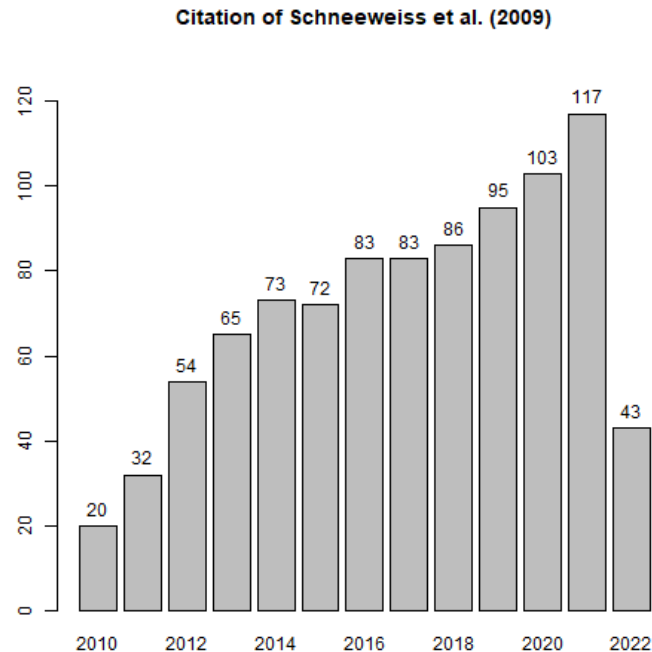
Regular epidemiological studies vs. **Proxies of underlying confounders**



| Unobserved confounder          | Observable proxy measurement   | Coding examples              |
|--------------------------------|--|------------------------------|
| Very frail health              | Use of oxygen canister   | CPT-4                        |
| Sick but not critical          | Code for hypertension during a hospital stay                             | ICD-9, ICD-10                |
| Health-seeking behavior        | Regular check-up visit; regular screening examinations                   | ICD-9, CPT-4, #PCP visits    |
| Fairly healthy senior          | Receiving the first lipid-lowering medication at age 70 years            | NDC, ATC, Read               |
| Chronically sick               | Regular visits with specialist, hospitalization; many prescription drugs | #specialist visits, NDC, ATC |
| Outcome surveillance intensity | General markers for health care utilization intensity                    | #visits, #different drugs    |

# High-dimensional proxy information

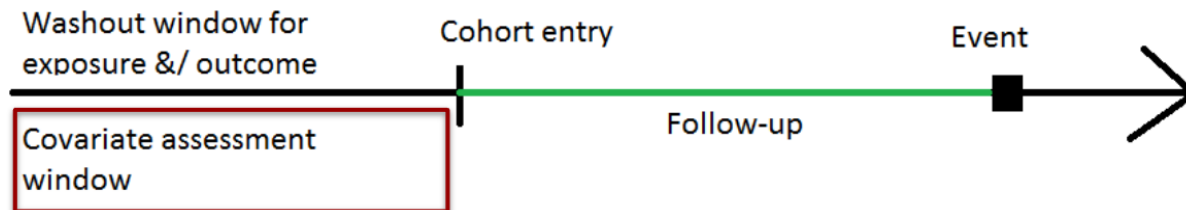
- Adjusting for something that **may not be interpretable** directly with the context of the research question.
- **Logic**: measures from same subject should be **correlated** = relevant proxy information





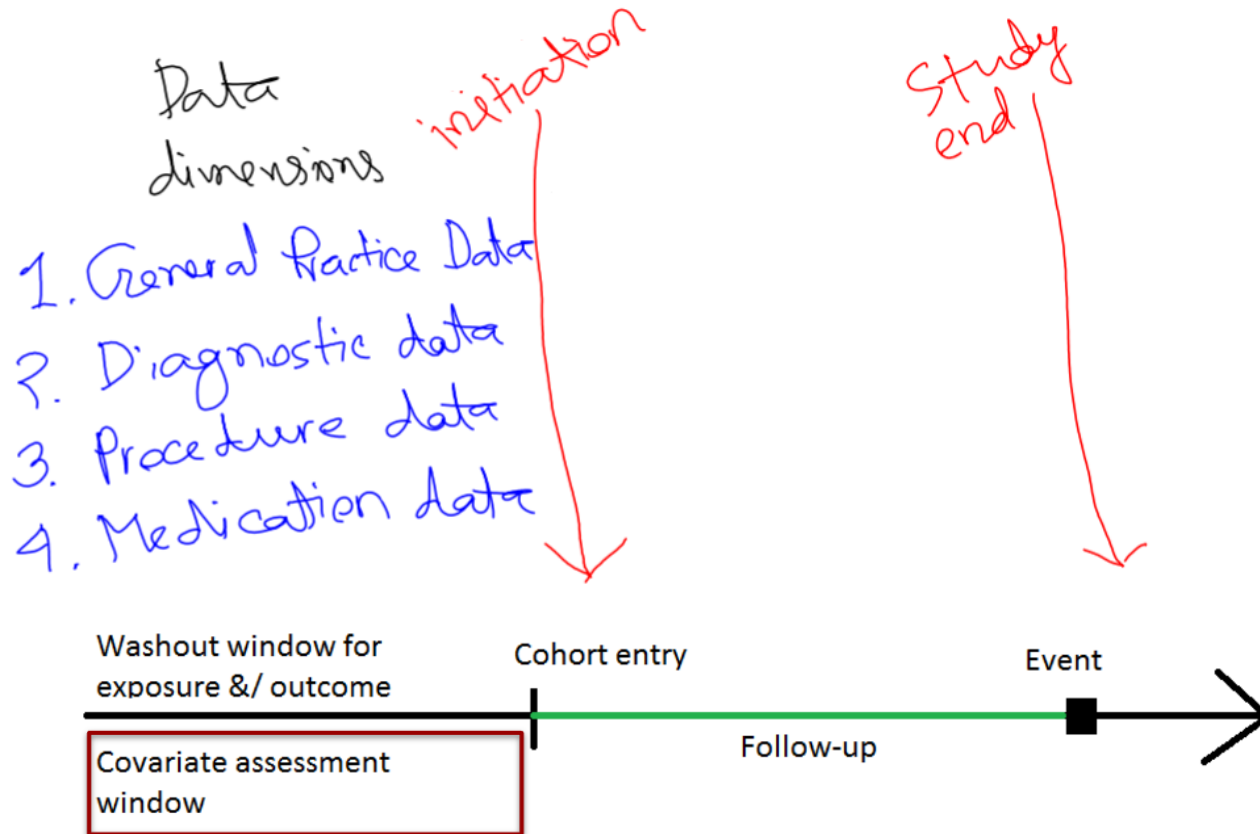
# hdPS: General Idea

[Karim et al. 2018](#) Epidemiology: Clinical Practice Research Datalink (1998–2012)



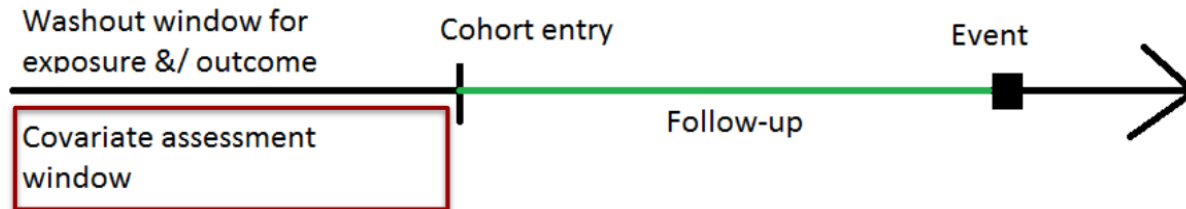
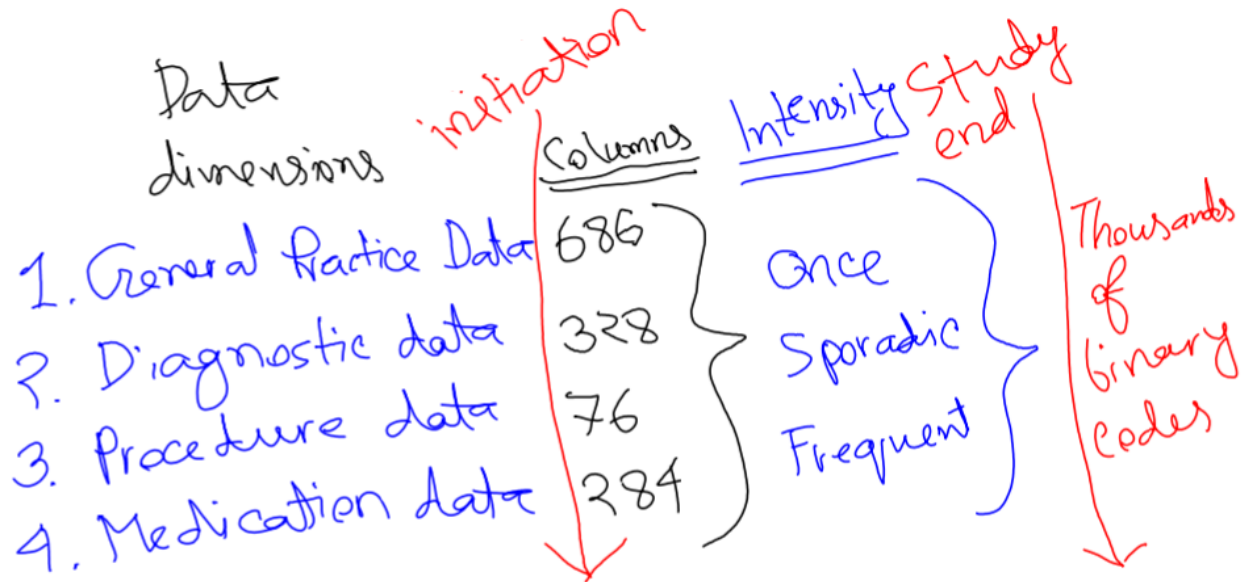
# hdPS: General Idea

[Karim et al. 2018](#) Epidemiology: Clinical Practice Research Datalink (1998–2012)



# hdPS: General Idea

[Karim et al. 2018](#) Epidemiology: Clinical Practice Research Datalink (1998–2012)



# hdPS: General Idea

List of additional proxy variables (**empirical covariates / EC**):

| Practice<br>(Dimension 1)     | Diagnostic<br>(Dimension 2)   | Procedure<br>(Dimension 3)   | Medication<br>(Dimension 4)   |
|-------------------------------|-------------------------------|------------------------------|-------------------------------|
| EC-dim1-1-once                | EC-dim2-1-once                | EC-dim3-1-once               | EC-dim4-1-once                |
| EC-dim1-1-sporadic            | EC-dim2-1-sporadic            | EC-dim3-1-sporadic           | EC-dim4-1-sporadic            |
| EC-dim1-1-frequent            | EC-dim2-1-frequent            | EC-dim3-1-frequent           | EC-dim4-1-frequent            |
| ...                           | ...                           | ...                          | ...                           |
| EC-dim1- <b>686</b> -frequent | EC-dim2- <b>328</b> -frequent | EC-dim3- <b>76</b> -frequent | EC-dim4- <b>284</b> -frequent |

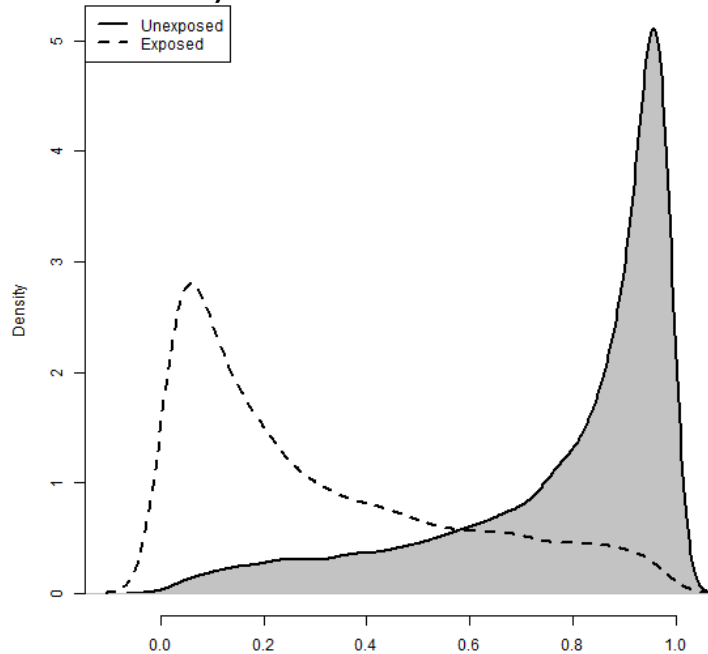
- Total  $(686+328+76+284)*3 = 4,122$  ECs
- 4 dimension  $\times$  3 intensity  $\times$  200 **most prevalent codes** [\*] = **2,400 ECs**
- [\*] [Schuster et al. \(2015\)](#), PDS recommended omitting prevalence-based selection

# hdPS: General Idea

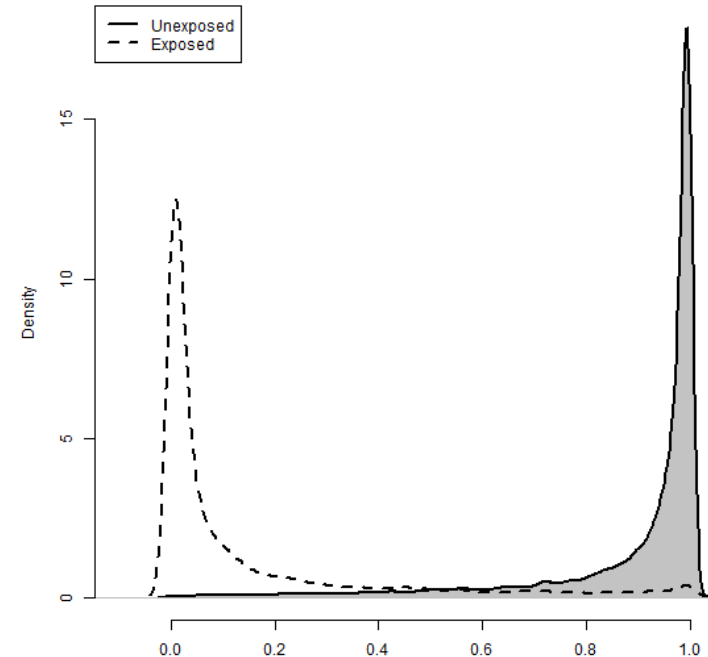
## Kitchen Sink Exposure Model ( $A \sim C + EC$ )

$$P(A = 1|C, EC) = \frac{1}{1 + \exp[\alpha_0 + \alpha_1 C_{\text{important}} + \alpha_2 C_{\text{potential confounder}} + \sum_{i=1}^{2,400} \alpha'_i EC_i]}$$

PS from only baseline confounders



PS from kitchen sink model!





# hdPS mechanism: find useful ECs

Assumption:

- $p_{u=1,a=1}$  = prevalence of unmeasured confounder among treated ( $A = 1$ )
- $p_{u=1,a=0}$  = prevalence of unmeasured confounder among untreated ( $A = 0$ )
- $p_{u=1,y=1}$  = prevalence of unmeasured confounder among dead ( $Y = 1$ )
- $p_{u=1,y=0}$  = prevalence of unmeasured confounder among alive ( $Y = 0$ )

Bross (1966) formula says, the amount of bias due to  $u$  is

$$\text{Bias}_M = \frac{p_{u=1,a=1} \times \left( \frac{p_{u=1,y=1}}{p_{u=1,y=0}} - 1 \right) + 1}{p_{u=1,a=0} \times \left( \frac{p_{u=1,y=1}}{p_{u=1,y=0}} - 1 \right) + 1}$$

In our example,

$U =$  smoking status

- [Bross \(1966\)](#) formula requires
  - binary  $U$
  - binary  $Y$
  - binary  $A$

# hdPS mechanism: find useful ECs

Assumption Calculate:

- $p_{EC=1,a=1}$  = prevalence of unmeasured confounder EC among treated ( $A = 1$ )
- $p_{EC=1,a=0}$  = prevalence of unmeasured confounder EC among untreated ( $A = 0$ )
- $p_{EC=1,y=1}$  = prevalence of unmeasured confounder EC among dead ( $Y = 1$ )
- $p_{EC=1,y=0}$  = prevalence of unmeasured confounder EC among alive ( $Y = 0$ )

Bross (1966) formula says, the amount of bias due to EC is

$$\text{Bias}_M = \frac{p_{EC=1,a=1} \times \left( \frac{p_{EC=1,y=1}}{p_{EC=1,y=0}} - 1 \right) + 1}{p_{EC=1,a=0} \times \left( \frac{p_{EC=1,y=1}}{p_{EC=1,y=0}} - 1 \right) + 1}$$

In our example,

$EC =$  EC-dim1-21-once  
= EC-dim2-95-once  
...  
= EC-dim4-64-once

- Bross (1966) formula requires
  - binary EC
  - binary Y
  - binary A

# hdPS mechanism: find useful ECs

Rank (descending) each EC by the magnitude of log-bias: Absolute  $\log Bias_M$

| Rank by bias | Absolute $\log Bias_M$ | EC                  |
|--------------|------------------------|---------------------|
| 1            | 0.42                   | EC-dim1-21-once     |
| 2            | 0.32                   | EC-dim2-95-once     |
| 3            | 0.25                   | EC-dim4-289-once    |
| ...          | ...                    | ...                 |
| 2,400        | 0.01                   | EC-dim4-64-frequent |

Take top **100** or **500** of these ECs. These are hdPS variables.

## hdPS Exposure Model ( $A \sim C + \text{top-ranked EC}$ )

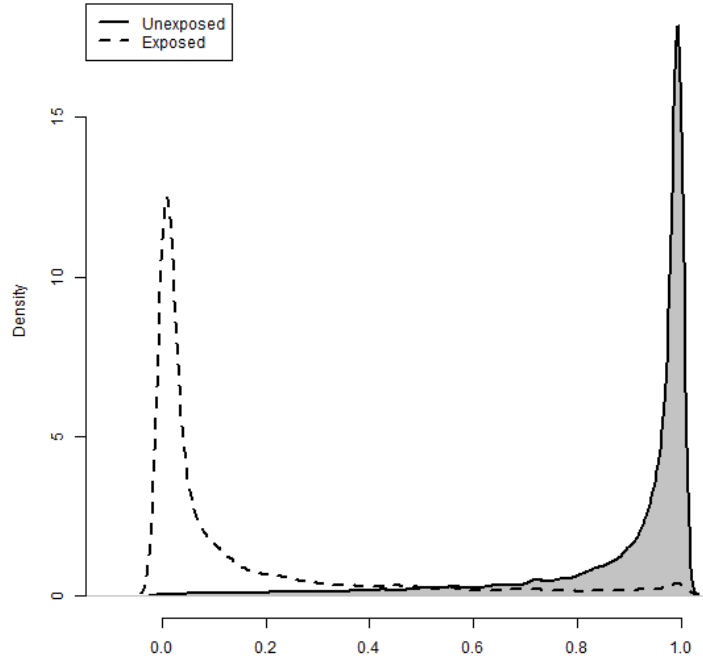
$$P(A = 1|C, EC) = \frac{1}{1 + \exp[\alpha_0 + \alpha_1 C_{\text{important}} + \alpha_2 C_{\text{potential confounder}} + \sum_{i=1}^{\text{top } 500} \alpha'_i EC_i]}$$

# hdPS: Assumption

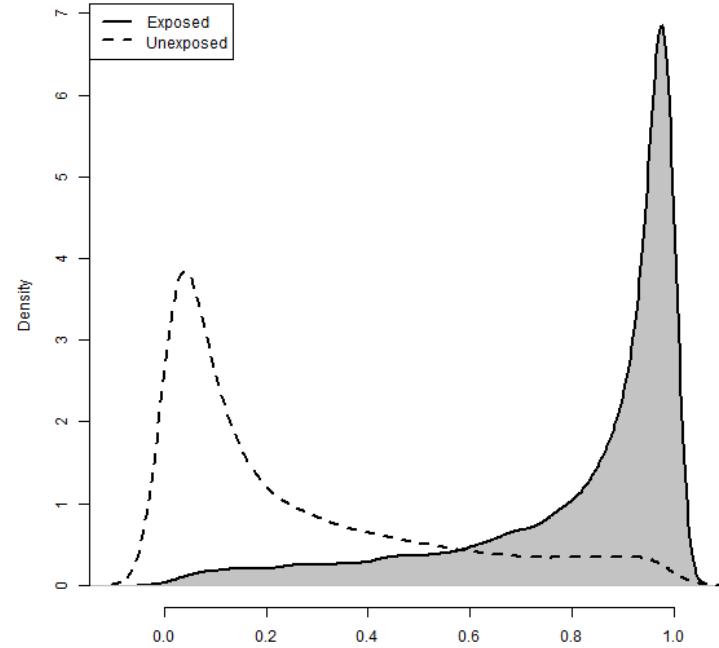
- The selected ECs collectively serve as **proxies of all unmeasured or residual confounding**
- **Implication:** an hdPS analysis may adjust for the unmeasured or residual confounding
- This assumption is strong and often not verifiable.
- Helpful in practice?

# hdPS: Balance

PS from kitchen sink model!



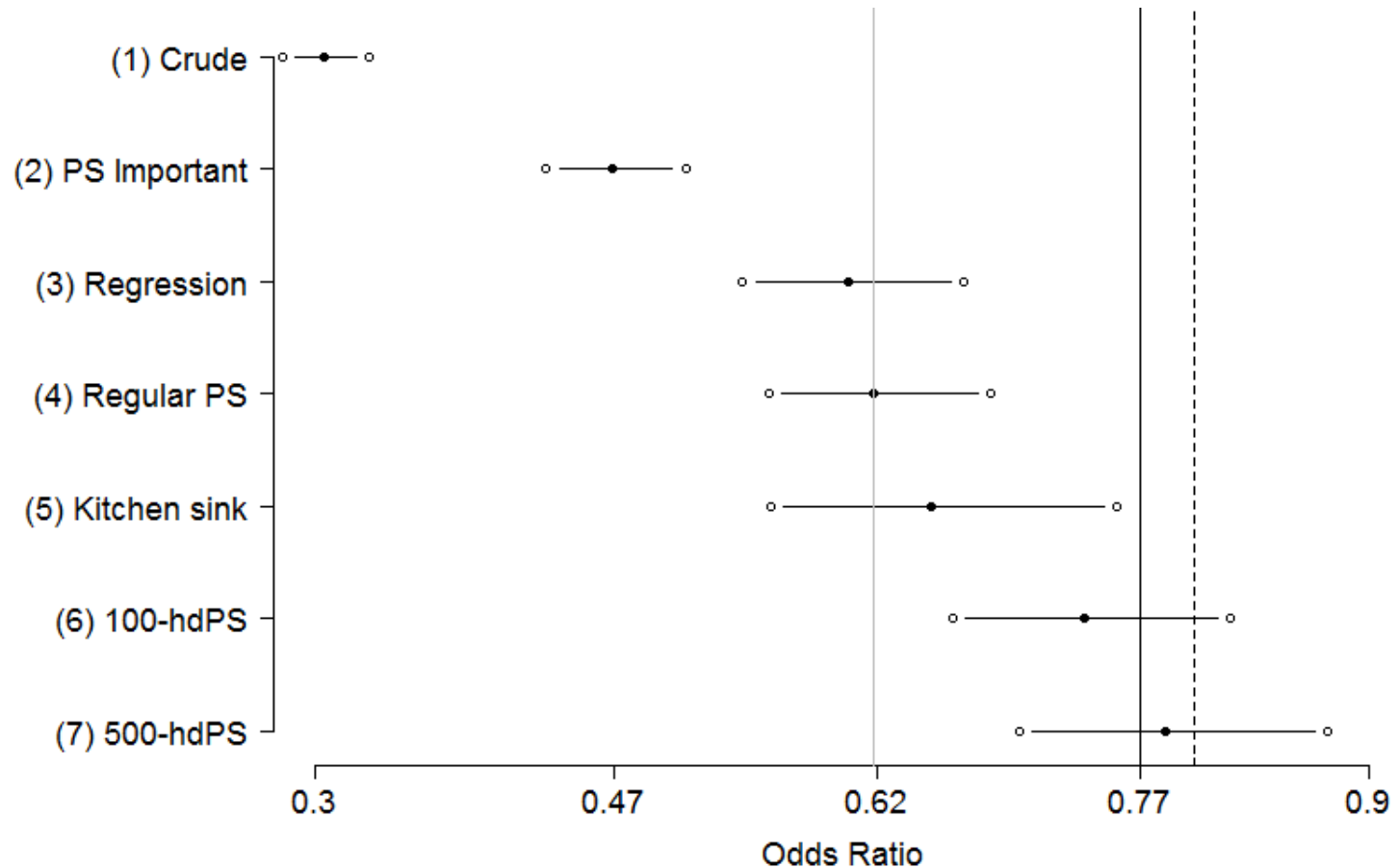
PS from 500-hdPS!





# hdPS: estimate treatment effect

- [Karim et al. 2018](#) Epidemiology
- Previous research: [Pang et al. \(2016\)](#): Epidemiology



# hdPS: Ways to improve

| Rank by bias | Absolute log $Bias_M$ | EC                  |
|--------------|-----------------------|---------------------|
| 1            | 0.42                  | EC-dim1-21-once     |
| 2            | 0.32                  | EC-dim2-95-once     |
| 3            | 0.25                  | EC-dim4-289-once    |
| ...          | ...                   | ...                 |
| 500          | 0.03                  | EC-dim4-63-frequent |

- ECs selected separately / **univariately** [VanderWeele et al. 2019](#) EJE
  - can be **correlated** (coming from same patient),
    - providing same information
    - **may not be useful anymore** in the presence of others
- **Multivariate** structure is good to consider
  - Model-specification

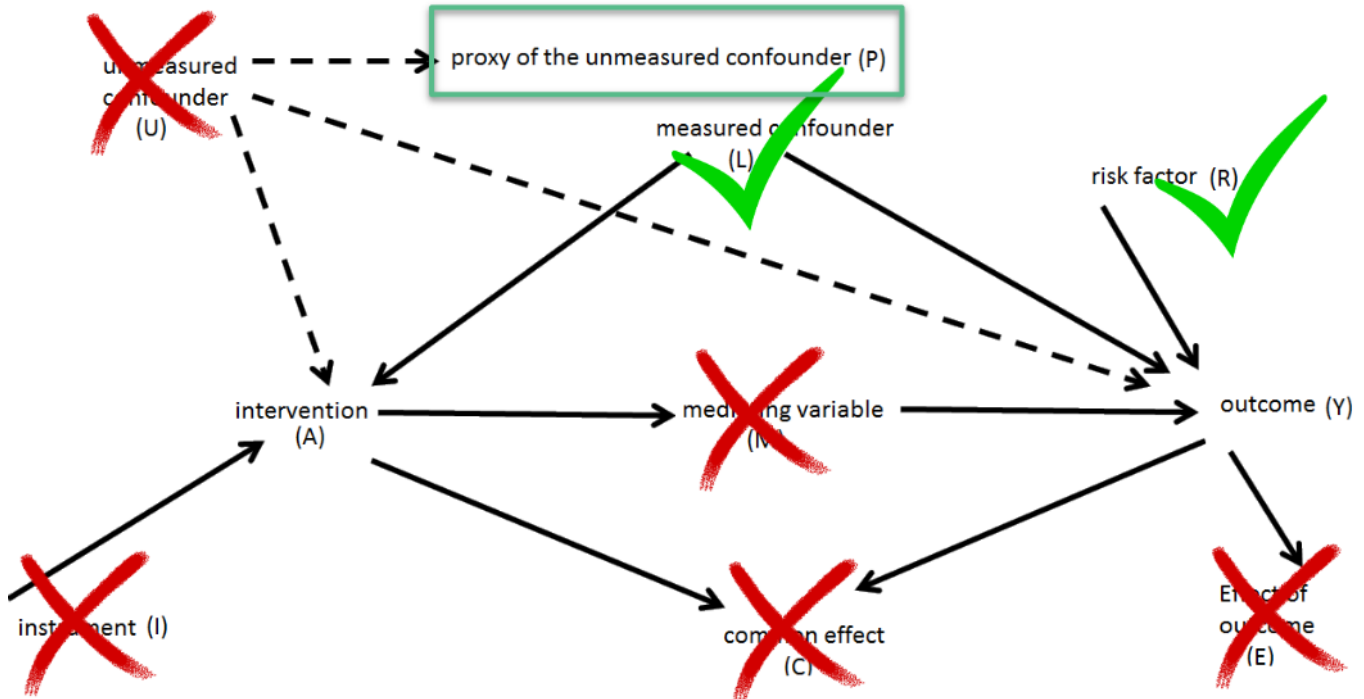
# Machine learning-based hdPS

# Variable selection in PS context

## Literature

- [Brookhart et al. \(2006\)](#), AJE
  - [Myers et al. \(2011\)](#), AJE
  - [Pearl \(2011\)](#), AJE
  - [Schuster et al \(2016\)](#), JCE
- bias amplification
  - inflated variance
  - overfitting

# Variable selection in PS context



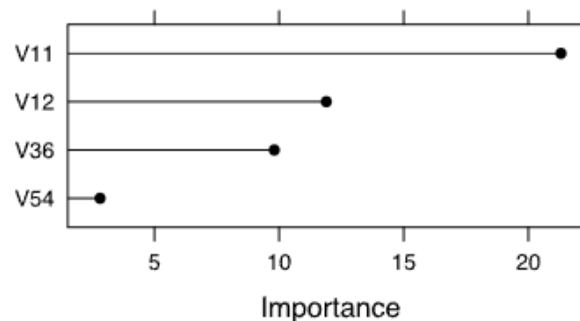
- How to select variables to adjust?
- Same idea for the proxies.
- **Pre-exposure** measurements (no mediator, collider, effect).
- **Associated with Y** (irrespective of association with A)



# Variable selection via ML

- **Jointly** consider in 1 model:
  - Perform variable selection based on **association with outcome**

| Approach   | Advantage  | Limitations  |
|--|--|--|
| LASSO <a href="#">Franklin et al. (2015)</a> , AJE                   | Variable selection by dropping <b>collinear</b> variables    | Tends to select one variable from a group, ignoring the rest |
| Elastic net  | More <b>stable</b> than LASSO                                | Non-linear and non-additive terms need to be specified       |
| Random forest <a href="#">Low et al. (2016)</a> , J. Comp. Eff. Res. | Automatically detect non-linearity and <b>non-additivity</b> | Only provides <b>variable importance</b> , but no cut-points |



# Machine learning-based hdPS

## Pure ML approach

Start with all ECs

### Outcome Model for EC selection ( $Y \sim C + ECs$ )

$$f(Y|C, EC) = \alpha_0 + \alpha_1 C_{\text{important}} + \alpha_2 C_{\text{potential confounder}} + \sum_{i=1}^{2,400} \alpha'_i EC_i$$

Say, 100 ECs (associated with Y) were selected by Elastic net approach

### Refined Exposure Model ( $A \sim C + \text{selected EC}$ )

$$P(A = 1|C, EC) = \frac{1}{1 + \exp[\alpha_0 + \alpha_1 C_{\text{important}} + \alpha_2 C_{\text{potential confounder}} + \sum_{i=1}^{\text{selected } 100} \alpha'_i EC_i]}$$

# Machine learning-based hdPS

## Hybrid approach (hdPS, then ML)

Start with top 500 ECs selected by Bross formula / prioritization

### Outcome Model for EC selection ( $Y \sim C + \text{top-500 ECs}$ )

$$f(Y|C, EC) = \alpha_0 + \alpha_1 C_{\text{important}} + \alpha_2 C_{\text{potential confounder}} + \sum_{i=1}^{500} \alpha'_i EC_i$$

Say, 100 ECs (associated with Y) were selected by Elastic net approach

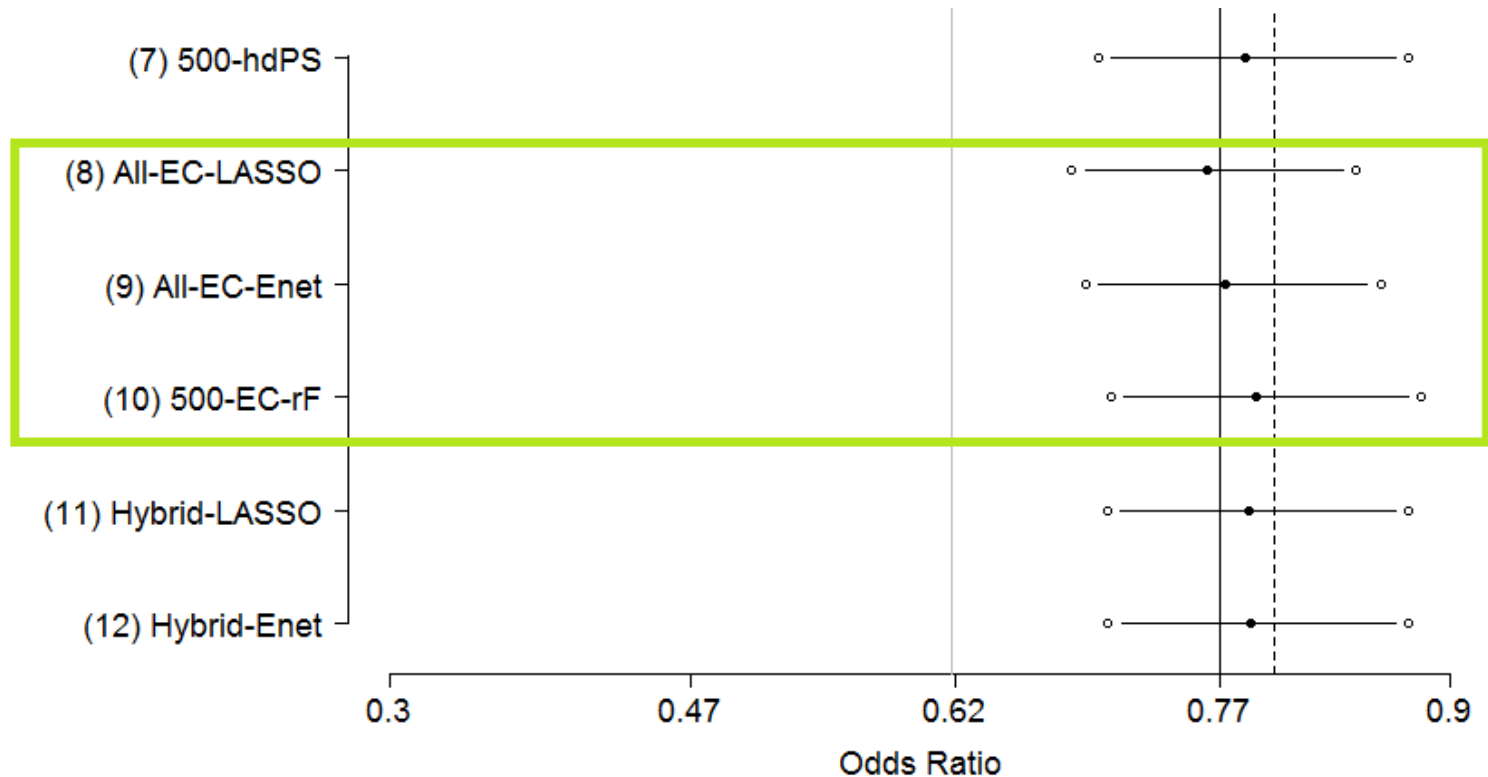
### Refined Exposure Model ( $A \sim C + \text{selected EC}$ )

$$P(A = 1|C, EC) = \frac{1}{1 + \exp[\alpha_0 + \alpha_1 C_{\text{important}} + \alpha_2 C_{\text{potential confounder}} + \sum_{i=1}^{\text{selected } 100} \alpha'_i EC_i]}$$

This approach is different than [Schneeweiss et al. \(2017\)](#), Epidemiology, where prioritization was used after applying LASSO.

# hdPS vs. ML: estimate treatment effect

[Karim et al. 2018](#) Epidemiology

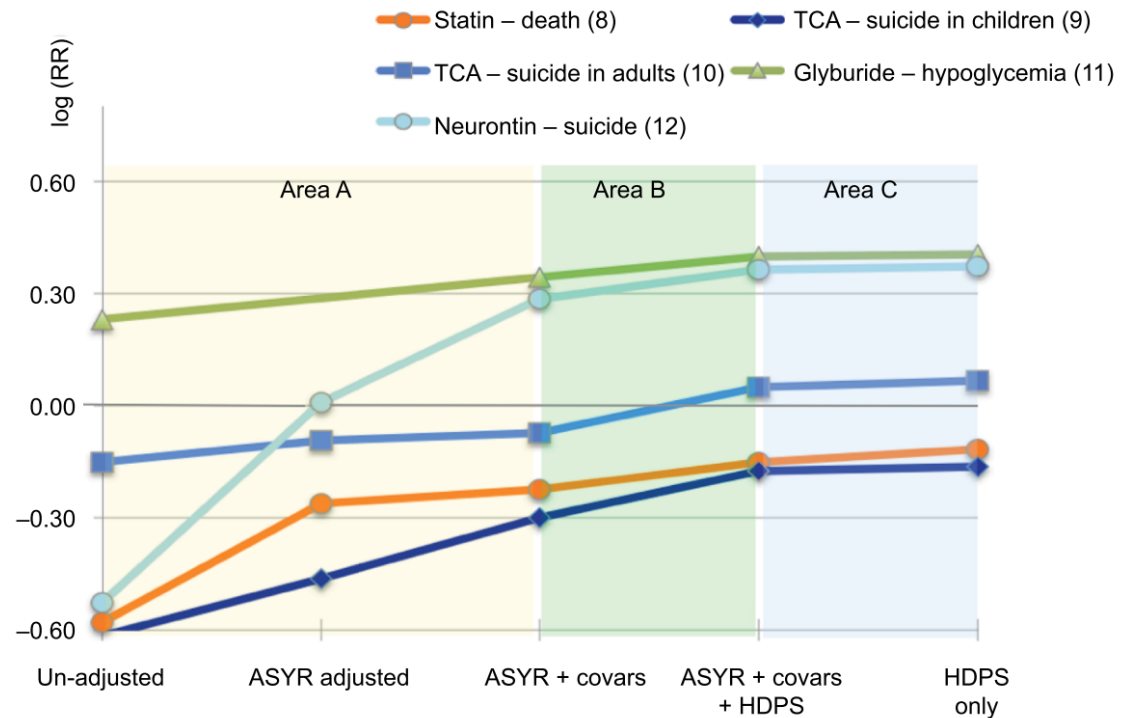


Only ~ 30% of the selected ECs were common.

# hdPS: estimate treatment effect

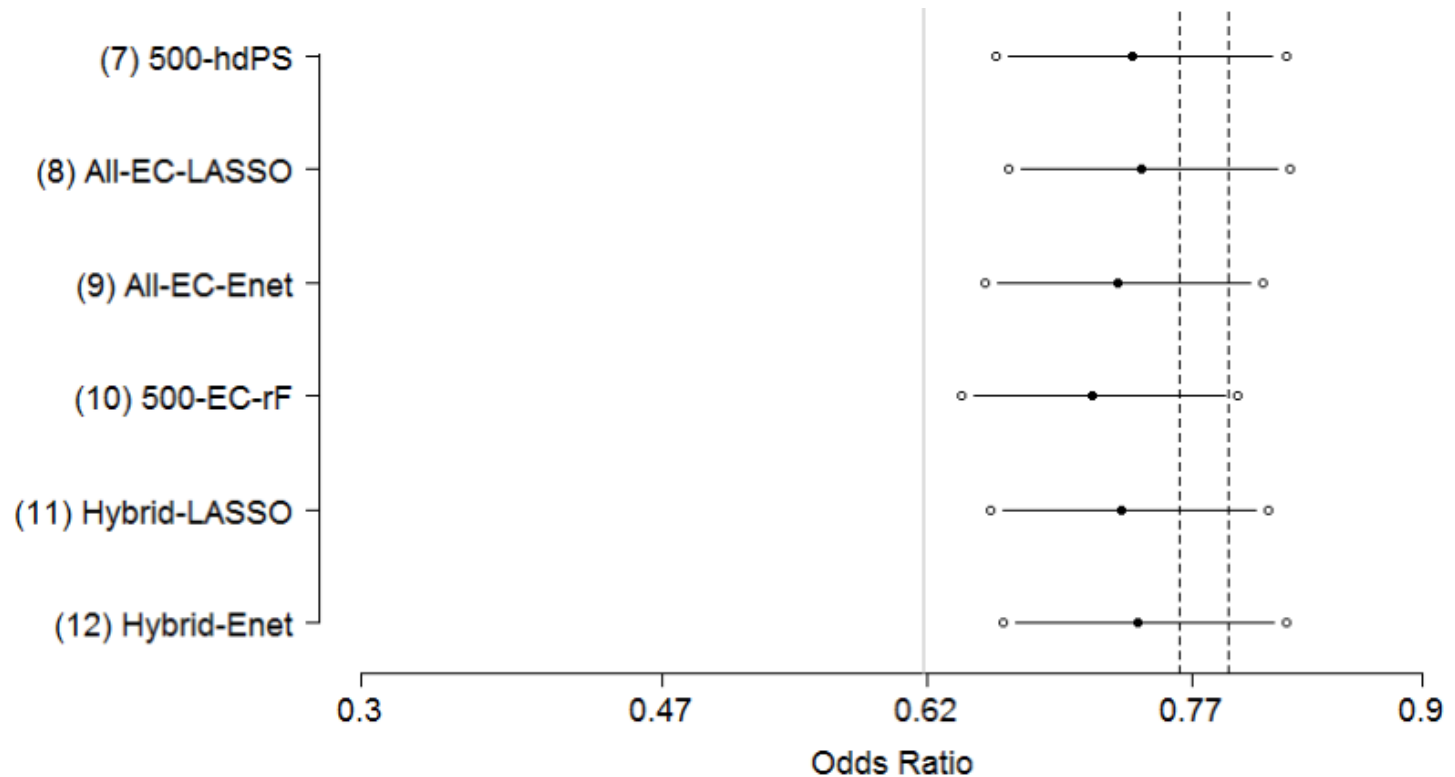
[Schneeweiss et al. 2018](#) Clinical Epidemiology: [CC BY NC license](#)

*"This strongly suggests that even **without the investigator-specifying covariates** for adjustment, the **algorithm alone** optimizes confounding adjustment."*



# hdPS vs. ML: estimate treatment effect

[Karim et al. 2018](#) Epidemiology



Quality of proxy information matters.

# Plasmode Simulation

Franklin et al. (2014) CSDA

| scenario | Multiplier of confounder effect | Exposure prevalence | Outcome prevalence | Unmeasured confounder |
|----------|---------------------------------|---------------------|--------------------|-----------------------|
| 1-U      | 1                               | 40                  | 5                  | Yes *                 |
| 2-U      | 3                               | 40                  | 5                  | Yes *                 |
| 3-U      | 5                               | 40                  | 5                  | Yes *                 |
| 4-U      | 1                               | 40                  | 10                 | Yes *                 |
| 5-U      | 3                               | 40                  | 10                 | Yes *                 |
| 6-U      | 5                               | 40                  | 10                 | Yes *                 |
| 7-U      | 1                               | 10                  | 5                  | Yes *                 |
| 8-U      | 3                               | 10                  | 5                  | Yes *                 |
| 9-U      | 5                               | 10                  | 5                  | Yes *                 |

Another baseline set with **no unmeasured confounding** (1-A to 9-A).

# Plasmode Simulation: Leaderboard

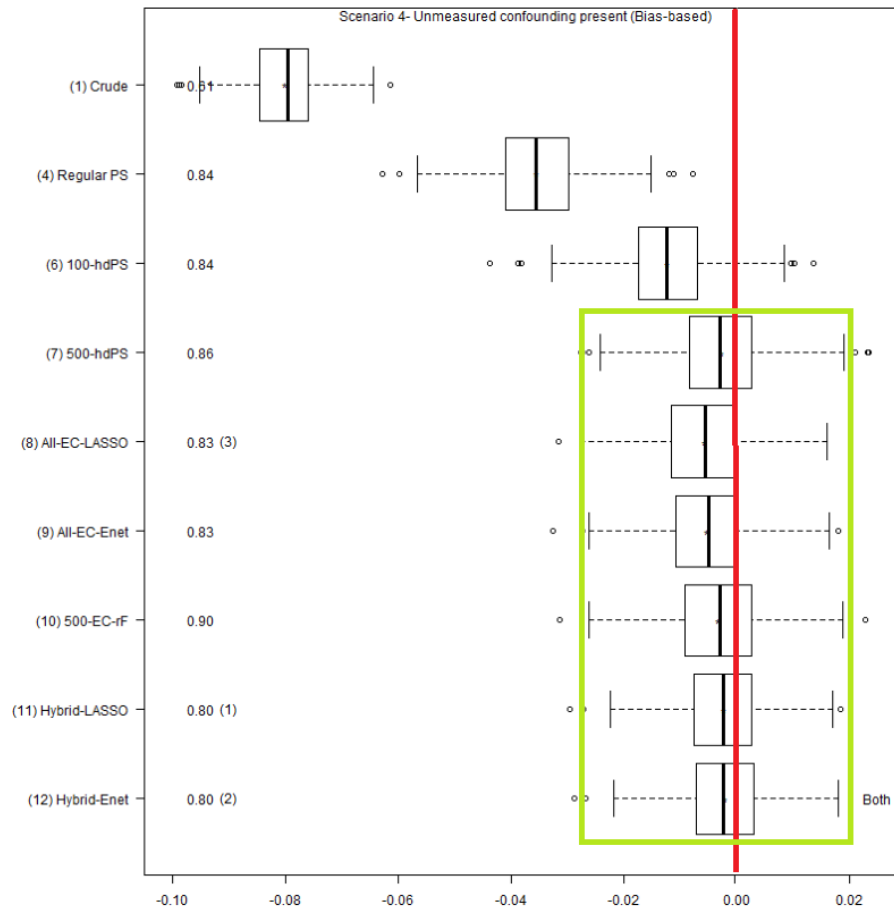
**Answer** to the question in the title of this talk (**bold** = pure ML):

| Scenario | Bias-Based       |                     | Exposure-Based      |                     |
|----------|------------------|---------------------|---------------------|---------------------|
|          | MSE              | Bias                | MSE                 | Bias                |
| 1-U      | Hybrid-Enet      | Hybrid-Enet         | <b>All-EC-Enet</b>  | <b>All-EC-Enet</b>  |
| 2-U      | Hybrid-LASSO     | <i>500-hdPS</i>     | <b>All-EC-Enet</b>  | <b>All-EC-Enet</b>  |
| 3-U      | Hybrid-LASSO     | <i>500-hdPS</i>     | <b>All-EC-Enet</b>  | <b>All-EC-Enet</b>  |
| 4-U      | Hybrid-Enet      | Hybrid-Enet         | <b>500-EC-rF</b>    | <b>500-EC-rF</b>    |
| 5-U      | <b>500-EC-rF</b> | <b>500-EC-rF</b>    | <b>500-EC-rF</b>    | <b>500-EC-rF</b>    |
| 6-U      | <b>500-EC-rF</b> | <b>500-EC-rF</b>    | <b>500-EC-rF</b>    | <b>500-EC-rF</b>    |
| 7-U      | Hybrid-Enet      | <i>500-hdPS</i>     | <b>All-EC-LASSO</b> | <b>All-EC-Enet</b>  |
| 8-U      | Hybrid-Enet      | <b>500-EC-rF</b>    | <b>All-EC-LASSO</b> | <b>All-EC-LASSO</b> |
| 9-U      | Hybrid-Enet      | <i>500-hdPS</i>     | <b>All-EC-LASSO</b> | <b>All-EC-Enet</b>  |
| 1-A      | Hybrid-LASSO     | <b>All-EC-LASSO</b> | <b>All-EC-Enet</b>  | <b>All-EC-LASSO</b> |
| 2-A      | Hybrid-LASSO     | Hybrid-LASSO        | <b>All-EC-Enet</b>  | <b>All-EC-Enet</b>  |
| 3-A      | Hybrid-Enet      | Hybrid-LASSO        | <b>All-EC-LASSO</b> | <b>All-EC-Enet</b>  |
| 4-A      | Hybrid-LASSO     | <b>All-EC-Enet</b>  | <b>All-EC-Enet</b>  | <b>All-EC-Enet</b>  |
| 5-A      | Hybrid-LASSO     | <b>500-EC-rF</b>    | <b>500-EC-rF</b>    | <b>500-EC-rF</b>    |
| 6-A      | Hybrid-Enet      | <b>500-EC-rF</b>    | <b>500-EC-rF</b>    | <b>500-EC-rF</b>    |
| 7-A      | Hybrid-Enet      | <i>500-hdPS</i>     | <b>All-EC-LASSO</b> | <b>All-EC-Enet</b>  |
| 8-A      | Hybrid-Enet      | <b>500-EC-rF</b>    | <b>All-EC-LASSO</b> | <b>All-EC-Enet</b>  |
| 9-A      | Hybrid-LASSO     | Hybrid-Enet         | <b>All-EC-LASSO</b> | <b>All-EC-Enet</b>  |



# Plasmode Simulation

Comparable if **adequate proxies** incorporated (RD estimates)



# Shared Limitations

- M-bias [Liu et al \(2012\)](#), AJE
- Z-bias [Myers et al. \(2011\)](#), AJE
- **EC interpretation** unclear vs. causal inference
  - not collected for research purposes
  - EC used in PS
- Primarily to deal with **residual confounding**
  - Not a straightforward extension to PS analysis
  - **Motivation of PS and hdPS are different to begin with**
- **No separation** of design and analysis stages in bias-based
  - exposure-based is OK; but has own issues
- post-selection bias [Taylor and Tibshirani \(2015\)](#)

# Advantage and Limitations

- Alternative ways to prioritize / rank
  - Automatic **cut-off** of how many variables
  - **Ranking**
- Pure ML methods can be used for **non-binary** outcomes and proxies
  - binary
  - categorized
  - continuous
  - survival
- **Coverage** not assessed [Morris et al. \(2019\)](#)
- Only a few ML methods assessed
- DR methods not covered

# Motivating Example

Basham et al. 2021 [EClinicalMedicine: CC BY license](#)

| Statistical Analysis <sup>a</sup>   | N         | Adjusted HR | 95% CI      |
|---|-----------|-------------|-------------|
| <i>Aim 1: analyzing post-TB airway disease risk</i>   |           |             |             |
| Covariate-adjusted (main analysis: respiratory TB vs controls)                                  | 1 005 328 | 2.08        | 1.91 – 2.28 |
| <i>Sensitivity analyses</i>   |           |             |             |
| Covariate-adjusted (removed ETOH, substance dependence, psychoses, and depression) <sup>b</sup> | 1 005 328 | 2.11        | 1.93 – 2.30 |
| Covariate-adjusted (van Walraven-weighted Elixhauser comorbidity score) <sup>c</sup>            | 1 005 328 | 2.06        | 1.89 – 2.26 |
| Covariate-adjusted (bronchiectasis and fibrosis added to the airway disease definition)         | 1 005 283 | 2.18        | 2.00 – 2.38 |
| Covariate-adjusted (removed respiratory TB patients with pleural samples; <i>n</i> = 55)        | 1 005 273 | 2.10        | 1.92 – 2.30 |
| <i>Different TB definitions</i>   |           |             |             |
| Covariate-adjusted (all forms of TB vs controls)  | 1 006 271 | 1.75        | 1.63 – 1.88 |
| Covariate-adjusted (non-respiratory TB vs controls) <sup>d</sup>                                | 1 004 733 | 1.36        | 1.20 – 1.53 |
| Age/sex-adjusted (pleural TB vs non-pleural TB)   | 1141      | 0.87        | 0.57 – 1.32 |
| <i>Aim 2: assessing potential unmeasured confounding</i>  |           |             |             |
| <i>PS methods</i>   |           |             |             |
| PS decile-adjusted (main covariates)  | 1 005 328 | 2.27        | 2.08 – 2.49 |
| hdPS decile-adjusted (main covariates + empirical covariates)                                   | 1 005 328 | 2.28        | 2.09 – 2.50 |
| LASSO-hdPS decile-adjusted (main covariates + LASSO-refined empirical covariates)               | 1 005 328 | 2.26        | 2.07 – 2.47 |
| <i>Adjustment for smoking behavior proxy variables</i>  |           |             |             |
| Covariate-adjusted subdata analysis (main covariates + tobacco use variable) <sup>e</sup>       | 31 063    | 1.53        | 1.37 – 1.71 |
| Covariate-adjusted (main covariates + personal health risk proxy variable)                      | 1 005 328 | 2.03        | 1.85 – 2.22 |

- Prefer to use hdPS / ML with ECs as a **secondary analysis**
- Proxy adjustment method (methods vs. subject area journals).

# JAMA Example

[Brown et al. \(2017\)](#)

| Method            | HR   | CI 95%     |
|-------------------|------|------------|
| Unadjusted        | 2.16 | 1.64-2.86  |
| Regression        | 1.59 | 1.17-2.17  |
| IPTW hdPS         | 1.61 | 0.997-2.59 |
| 1-1 hdPS matching | 1.64 | 1.07-2.53  |
| Pre-pregnancy     | 1.85 | 1.37-2.51  |

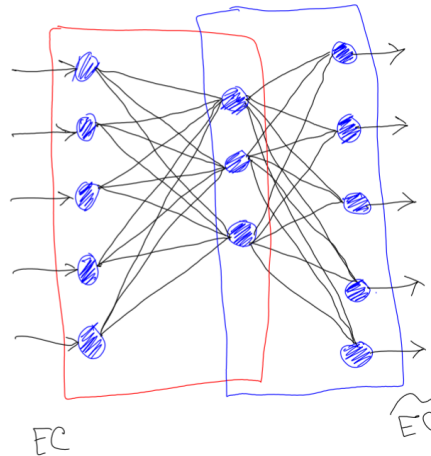
- Conclusion: **not associated**
- More discussion: [Amrhein, Trafimow, Greenland, 2019](#) The American Statistician

# Related research directions

# Related research directions

## AI : Autoencoders

[Weberpals et al. \(2021\)](#) used autoencoders (3, 5, 7 layers) to reduce EC dimensions.



- Autoencoder-based hdPS is useful.
- Shallow learning (less layers) had better MSE.
- **Did not perform better than LASSO.**

# Related research directions

## TMLE

Targetted learning approach [Pang et al. \(2016\)](#): Epidemiology

| Model                        | Max SW weight |
|------------------------------|---------------|
| Only important 5 confounders | 1.78          |
| 29 confounders               | 69.67         |
| 29 confounders + 400 ECs     | 390.77        |

- better covariate balance vs. overfitting
  - Varying number of covariates selected [Tazare et al. 2022](#)

[Haris and Platt \(2021\)](#), arxiv

- group importance score
- extension of the hdPS (**hdCS**) to non-binary outcome and confounders



# Related research directions

## Sample splitting

[Naimi et al. \(2021\)](#), AJE

SL, TMLE, AIPW and usefulness of sample splitting

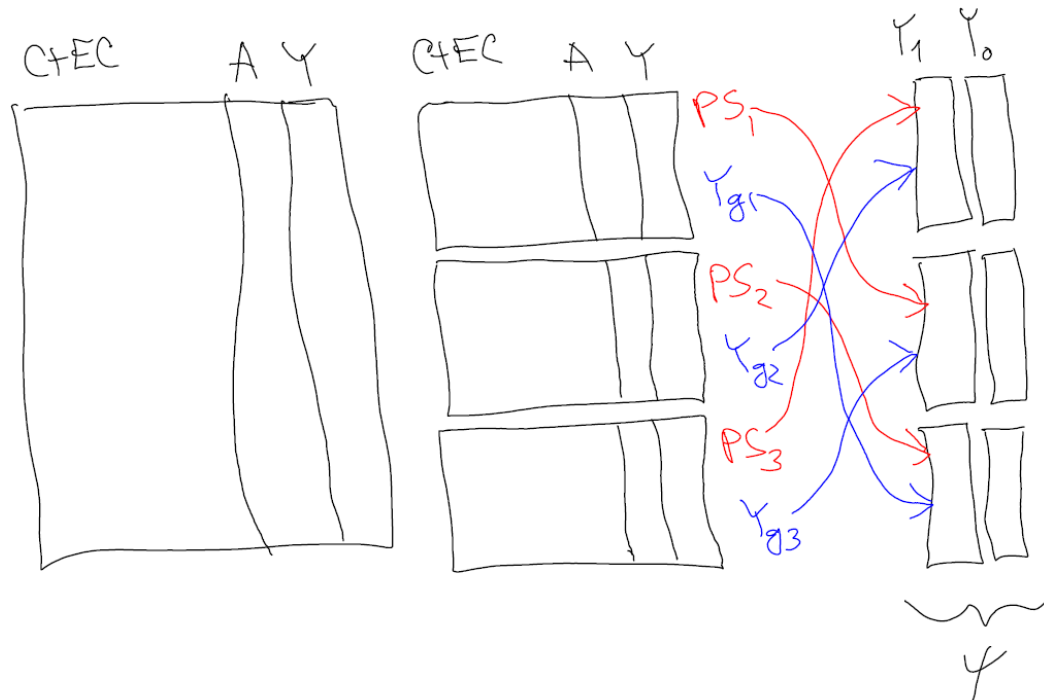
- ML based **singly robust methods** should be avoided
- Use **sample splitting**
- **rich SL library** of flexible regression as well as higher order interactions

# Related research directions

## Cross-fitting

[Zivich and Breskin \(2021\)](#), Epidemiology

- Cross-fitting + together with double-robust approaches



# Related research directions

## SL library

[Balzer and Westling \(2021\)](#), AJE

- TMLE without sample-splitting with a carefully chosen SL library

[Meng and Huang \(2021\)](#), arxiv

- SL with **smooth** (differentiable: LASSO, spline) learners outperform those that included non-smooth learners

# Take home message

- hdPS and ML alternatives generally reduces **residual confounding**
  - [\*] if **good proxies** available.
- hdPS: dependent on **Bross-formula** (all binary)
- Non-binary outcome: consider ML methods.
- **Hybrid-methods** performed better (MSE).
- Active area of research

# Thanks!

<http://ehsank.com/>