

# How to obtain valid tests and confidence intervals for treatment effects after confounder selection?

Stijn Vansteelandt

*Ghent University, Belgium*

*London School of Hygiene and Tropical Medicine, U.K.*

*joint work with Oliver Dukes, Vahe Avagyan*

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- 1 Introduction
- 2 A dissection of the problem
- 3 Proposal
- 4 Numerical results
- 5 Discussion

# Introduction (1)

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- Exposed and unexposed subjects in such studies usually differ in many observed (pre-exposure) characteristics  $L$ .
- This can make it difficult to make contrasts of the mean outcome between exposed and unexposed subjects with the same characteristics.

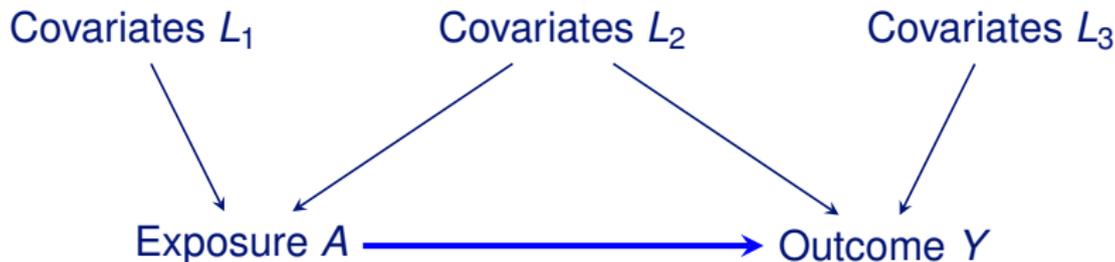
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- Exposed and unexposed subjects in such studies usually differ in many observed (pre-exposure) characteristics  $L$ .
- This can make it difficult to make contrasts of the mean outcome between exposed and unexposed subjects with the same characteristics.
- The **curse of dimensionality** thus forces us to adopt some form of **modelling**.
- E.g. a linear model

$$E(Y|A, L) = \psi A + \beta' L$$

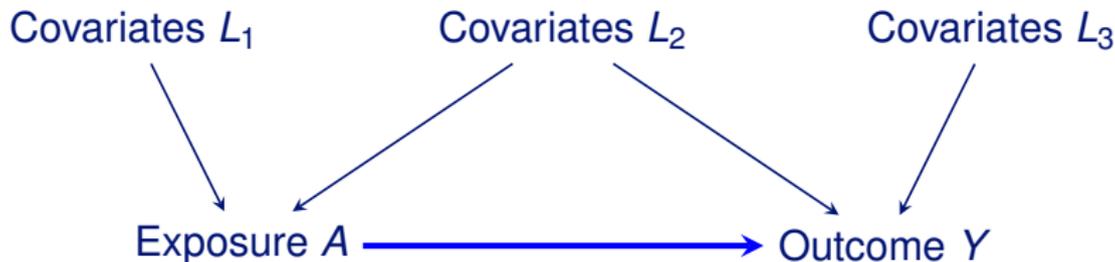
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- There may be more covariates than observations.
- This is not uncommon, considering the possible need for interactions or other higher-order terms...

## Introduction (3)

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- One common strategy is to adjust for  $L$  iff it is significantly associated with outcome, conditional on exposure, at e.g. the 5% level.
- A related common strategy is the lasso, without penalisation of the exposure effect.
- *How well does this work?*

# Outcome-based selection

- Suppose that the exposure has no effect.
- Suppose that  $L$  has a moderate effect on outcome, but a strong effect on exposure.
- Then when fitting model

$$E(Y|A, L) = \psi A + \beta L$$

one will typically have little power to detect that  $\beta \neq 0$ .

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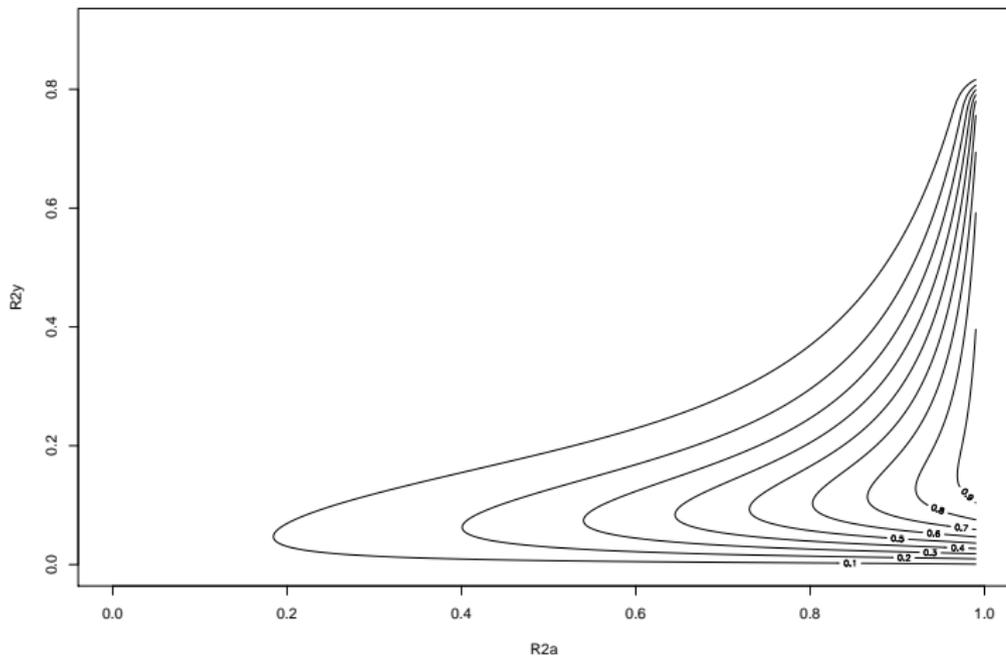
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- Upon removing  $L$  from the model, one is likely to find 'strong evidence' of an exposure effect.
- This can result in highly inflated Type I error rates

Type I error rate inflation ( $n = 100$ )

$R_{2y}$ :  $R^2$  of  $Y$ - $L$  association;  $R_{2a}$ :  $R^2$  of  $A$ - $L$  association



# Convergence with increasing sample size

- This problem **persists at all sample sizes**.
- No matter how large the sample size, one can always choose correlations between  $Y-L$  and  $A-L$ , at which outcome-based selection inflates Type I error rates.

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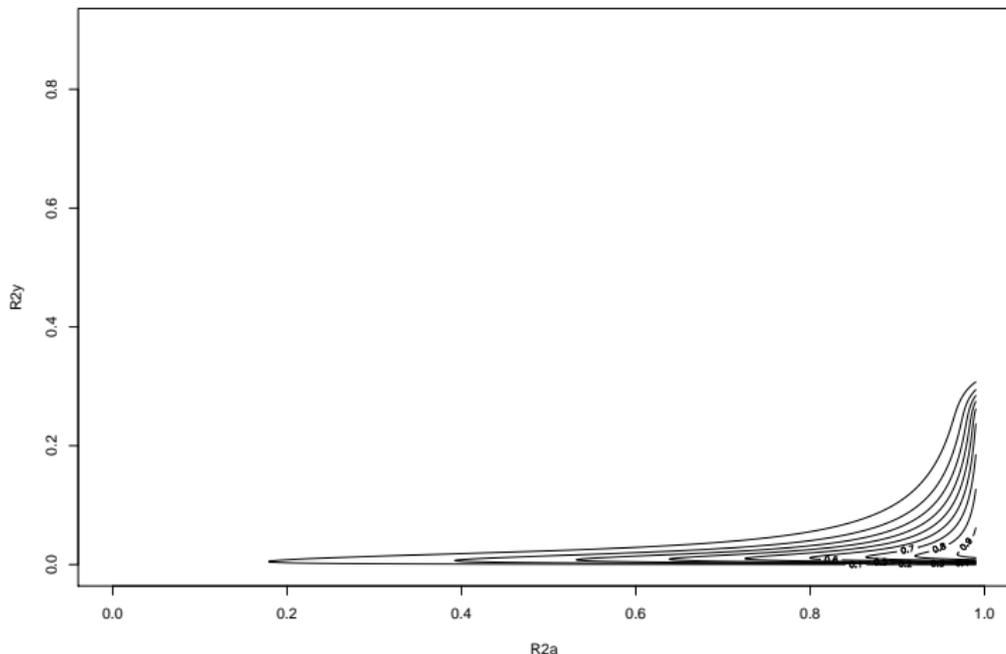
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- We therefore say that convergence of the test statistic to a normal limit (centered around the truth) is **non-uniform**.
- Lack of uniform convergence is a concern.
- It implies that we can never guarantee that the procedure will do well in finite samples.

Outcome-based selection ( $n = 1000$ )

$R_{2y}$ :  $R^2$  of  $Y$ - $L$  association;  $R_{2a}$ :  $R^2$  of  $A$ - $L$  association



# Propensity-score-based selection (1)

- One key reason why this procedure is problematic, is that it **prioritises the exposure**: it prioritises the elimination of covariates over the elimination of the exposure.

(Robins and Greenland, 1986)

- This problem can be overcome using **propensity scores**.

## Propensity-score-based selection (2)

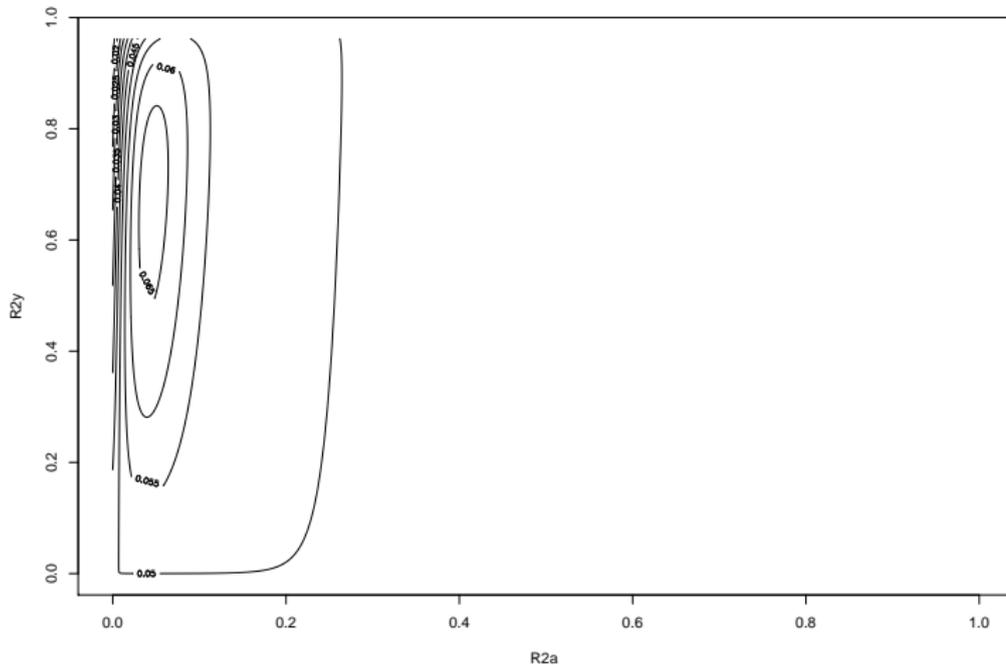
- Consider stepwise selection in a propensity score model, then regressing outcome on exposure and propensity score.
- By always adjusting for the propensity score, this strategy does not prioritise the exposure.

## Propensity-score-based selection (2)

- Consider stepwise selection in a propensity score model, then regressing outcome on exposure and propensity score.
- By always adjusting for the propensity score, this strategy does not prioritise the exposure.
- With linear models for  $Y$  and  $A$ , and a single covariate  $L$ , this strategy is tantamount to adjusting for  $L$  **iff it is significantly associated with exposure**, at e.g. the 5% level.

Type I error rate inflation ( $n = 100$ )

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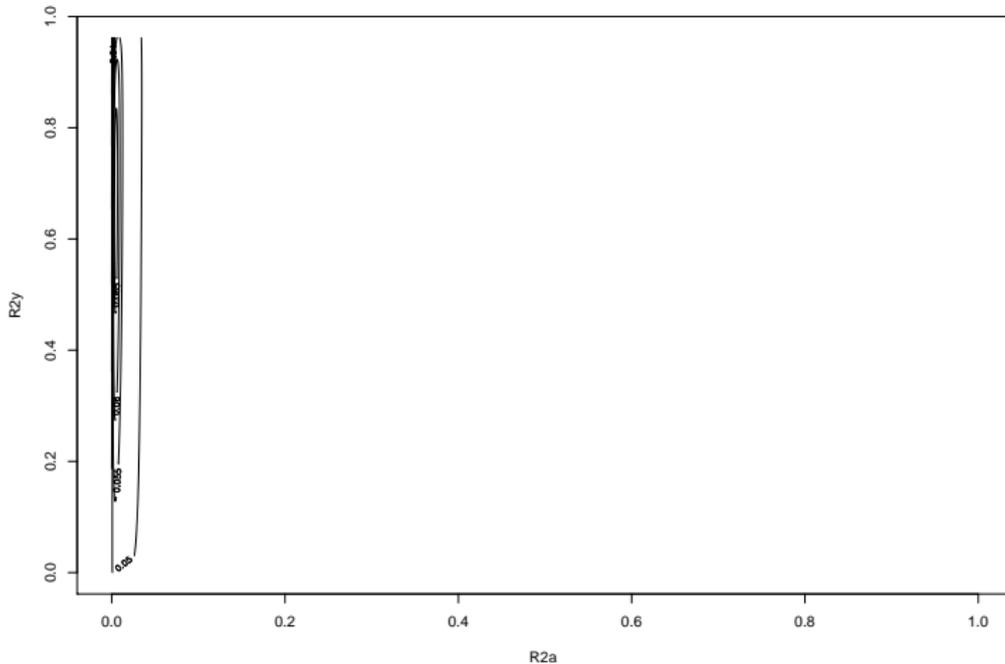
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## Propensity-score-based selection (2)

- By not prioritising the exposure, the problem of Type I error inflation is much less severe.
- In fact, ignoring the variable selection process often results in **conservative inferences**.
- This is line with the property that ignoring estimation of the propensity score typically results in conservative inferences.
- Also this persists at all sample sizes.

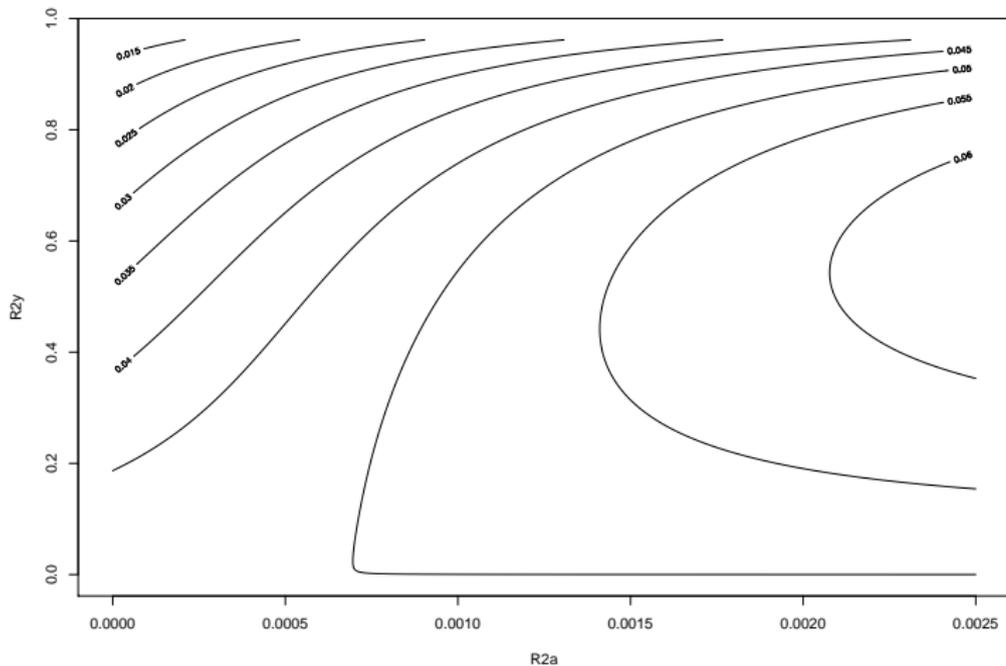
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- *What if the models are non-linear?*

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- *Problem solved?*
- Its typical conservatism implies a lack of power.
- *What if there are many covariates?*
- *What if the models are non-linear?*
- In view of this, the aim of this talk will be **to develop uniformly valid tests that incorporate selection.**
- The propensity score will continue to play a crucial role...

# Post-selection inference

- This problem of post-selection inference has been quite thoroughly studied for some selection strategies.  
(e.g. Leeb and Pötscher, 2005; Berk et al., 2013; Taylor et al., 2014; ...)
- Most proposed solutions infer the distribution of the estimator or test statistic after selection.  
(e.g. Claeskens and Hjört, 2006)
- This has the **disadvantage** that the results
  - are often complex,
  - not immediately accessible for routine data analysis,
  - and sometimes dependent on the choice of procedure.

- Inspired by others,

(Chernozhukov et al., 2017; Farrell, 2015)

I will instead propose **specific tests for treatment effect**  
in combination with **a specific selection strategy**.

- Their combination is such that the test statistic  
converges uniformly to a normal distribution centred at the truth.

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# Hypothesis-test-based selection

- Reconsider model  $E(Y|A, L) = \psi A + \beta L$   
(where  $A$  and  $L$  are mean centred).
- Perform a score test of  $\psi = 0$  based on the test statistic

$$\frac{1}{\sqrt{n}} \sum_{i=1}^n A_i (Y_i - \hat{\beta} L_i)$$

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- *What is the distribution of the test statistic?*
- Consider outcome-based selection...

# What is the distribution of the test statistic? (1)

By a Taylor expansion,

$$\begin{aligned} & \frac{1}{\sqrt{n}} \sum_{i=1}^n A_i \{Y_i - \hat{\beta}' L_i\} \\ &= \frac{1}{\sqrt{n}} \sum_{i=1}^n A_i \{Y_i - \beta' L_i\} + \sqrt{n}(\hat{\beta} - \beta) \left\{ \frac{1}{n} \sum_{i=1}^n A_i L_i \right\} \\ & \quad + \text{Remainder} \end{aligned}$$

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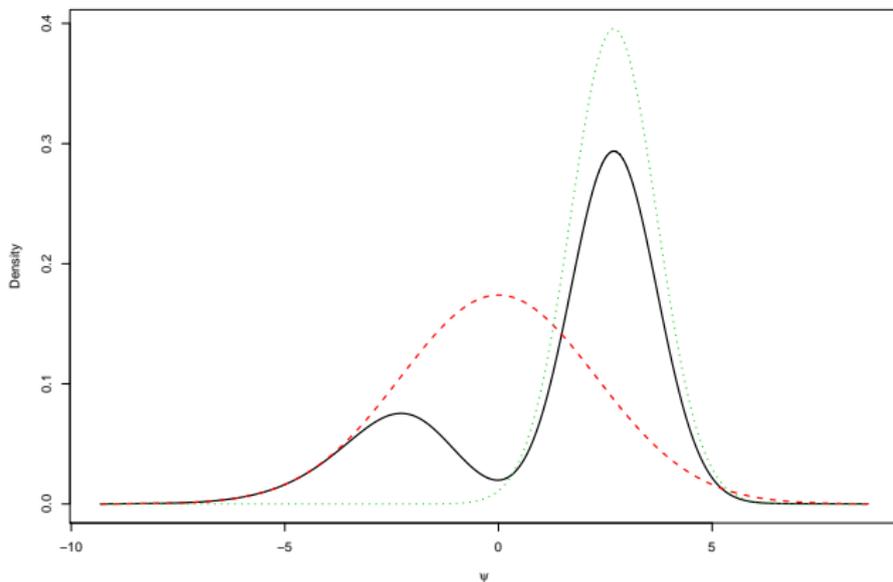
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- When  $\beta$  is of the order  $1/\sqrt{n}$ , we will often erroneously set  $\hat{\beta}$  to zero.
- This results in bias, which affects the score test.
- $\sqrt{n}(\hat{\beta} - \beta)$  then moreover has a complex distribution.

# Inference after variable selection (2)

This may cause bias, excess variability, and may invalidate inference.



## What is the distribution of the test statistic? (2)

- Convergence of

$$\frac{1}{\sqrt{n}} \sum_{i=1}^n A_i \{Y_i - \hat{\beta}' L_i\}$$

to a mean zero normal distribution is therefore non-uniform.

- We will remedy this  
using **bias-reduced double-robust estimators**.

(Vermeulen and Vansteelandt, 2015)

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# Double-robust estimation

- Consider the test statistic

$$\frac{1}{\sqrt{n}} \sum_{i=1}^n \{A_i - \pi(L_i; \gamma)\} \{Y_i - m(L_i; \beta)\}$$

where we use

- a parametric propensity score model  $\mathcal{A}$ :

$$E(A|L) = \pi(L; \gamma)$$

e.g.  $\text{expit}(\gamma' L)$  for binary  $A$ .

- a parametric outcome model  $\mathcal{B}$ :

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- This test statistic has mean zero under the null **when either model  $\mathcal{A}$  or model  $\mathcal{B}$  is correct.**
- We therefore call it **double-robust.**

(Robins and Rotnitzky, 2001; see Rotnitzky and Vansteelandt, 2014, for a review)

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- In practice, we need estimators of  $\gamma$  and  $\beta$ .
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 & \quad + \sqrt{n}(\hat{\beta} - \beta) \left\{ \frac{1}{n} \sum_{i=1}^n \frac{\partial}{\partial \beta} U_i(\gamma, \beta) \right\} + \text{Remainder}
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- **If we could set those gradients to zero,** then local changes in these estimators would not affect the double-robust test.

# Bias-reduced double-robust estimation

- **Bias-reduced double-robust estimators** achieve this by estimating  $\gamma$  by solving

$$\frac{1}{n} \sum_{i=1}^n \frac{\partial}{\partial \beta} U_i(\gamma, \beta) = 0$$

and  $\beta$  by solving

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- *Is this a valid proposal?*
- Suppose model  $\mathcal{A}$  is correct with true value  $\gamma^*$ .
- Then  $U_i(\gamma^*, \beta)$  has mean zero for all  $\beta$ , so that

$$E \left\{ \frac{\partial}{\partial \beta} U_i(\gamma^*, \beta) \right\} = 0$$

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- We will incorporate it by penalising the estimating equations with a bridge penalty:

$$0 = \frac{1}{n} \sum_{i=1}^n \frac{\partial}{\partial \beta} U_i(\gamma, \beta) + \lambda_{\beta} \delta |\beta|^{\delta-1} \circ \text{sign}(\beta)$$

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where  $\lambda_{\gamma} > 0$  and  $\lambda_{\beta} > 0$  are penalty parameters, and  $\delta \rightarrow 1+$ .

(Avagyan and Vansteelandt, 2017; Dukes, Avagyan and Vansteelandt, 2018)

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(Avagyan and Vansteelandt, 2017; Dukes, Avagyan and Vansteelandt, 2018)

- Standard choices of penalty (of the order  $\sqrt{\log(p)/n}$ ) make these gradients sufficiently close to zero.

# Example - $Y$ continuous, $A$ binary

- Consider models  $\pi(L; \gamma) = \text{expit}(\gamma' L)$  and  $m(L; \beta) = \beta' L$ .
- Then we estimate  $\gamma$  and  $\beta$  as the solutions to

$$0 = \frac{1}{n} \sum_{i=1}^n \{A_i - \text{expit}(\gamma' L_i)\} L_i + \lambda_\gamma \delta |\gamma|^{\delta-1} \circ \text{sign}(\gamma)$$

$$0 = \frac{1}{n} \sum_{i=1}^n w_i(\gamma) \{Y_i - \beta' L_i\} L_i + \lambda_\beta \delta |\beta|^{\delta-1} \circ \text{sign}(\beta)$$

where  $w_i(\gamma) = \text{expit}(\gamma' L_i) \{1 - \text{expit}(\gamma' L_i)\}$ .

# Example - $Y$ continuous, $A$ binary

- In practice, we let  $\delta \rightarrow 1+$  and solve the following problems:

$$\min_{\gamma} \mathcal{F}(\gamma) = \frac{1}{n} \sum_{i=1}^n \log\{1 + \exp(\gamma' L_i)\} - A_i(\gamma' L_i) + \lambda_{\gamma} \|\gamma\|_1$$

$$\min_{\beta} \mathcal{F}(\beta) = \frac{1}{2n} \sum_{i=1}^n [\hat{w}_i \{Y_i - \beta' L_i\}^2] + \lambda_{\beta} \|\beta\|_1$$

- Components of  $\hat{\eta}$  may be shrunk to zero, in view of which we recommend refitting the selected model.
- The test statistic is then

$$T_n = \frac{\frac{1}{n} \sum_{i=1}^n \{A_i - \text{expit}(\hat{\gamma}' L_i)\} \{Y_i - \hat{\beta}' L_i\}}{\sqrt{\frac{1}{n} \left\{ \frac{1}{n-1} \sum_{i=1}^n [\{A_i - \text{expit}(\hat{\gamma}' L_i)\} \{Y_i - \hat{\beta}' L_i\}]^2 \right\}}}$$

# Asymptotic properties - both models correct

- Let  $s_\gamma$  and  $s_\beta$  be the sparsity indices of models  $\mathcal{A}$  and  $\mathcal{B}$ .
- Suppose that (in addition to mild regularity conditions), the following sparsity assumptions hold:
  - (i)  $s_\gamma \log(p) = o(n)$
  - (ii)  $s_\beta \log(p) = o(n)$
  - (iii)  $s_\gamma s_\beta \log^2(p) = o(n)$ .

## Theorem

When model  $\mathcal{A}$  and  $\mathcal{B}$  are correct, the test statistic  $T_n$  converges **uniformly** to a standard normal distribution.

# A closer look at the conditions...

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  - (iii)  $s_\gamma s_\beta \log^2(p) = o(n)$determines the rate of convergence of the estimators.
- It suggests that if one model is sparse, the other can be more dense.
- When evaluating medical treatments, this is arguably satisfied as clinicians may use a limited number of variables to decide on treatment, whereas outcome may be affected by many more.

# Relationship to existing literature

Compared with other recent proposals from high-dimensional inference in GLMs:

(van de Geer et al., 2014; Belloni et al., 2016)

- We have weakened the assumptions on sparsity by making use of double robustness.

(see also Farrell, 2015, for the ATE)

- Other approaches usually require **ultra-sparsity**, e.g.  $s_\gamma \sqrt{\log(p)} = o(\sqrt{n})$  instead of  $s_\gamma \log(p) = o(n)$ .

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- Unlike others, we **do not require sample-splitting** to obtain weaker rates.

(Chernozhukov et al., 2017)

Suppose that (in addition to the previous conditions), the following sparsity assumptions hold:

- (iv) Either (a)  $s_\gamma \sqrt{\log(p)} = o(\sqrt{n})$  (if model  $\mathcal{A}$  is correct) or (b)  $s_\beta \sqrt{\log(p)} = o(\sqrt{n})$  (if model  $\mathcal{B}$  is correct).

### Theorem

When model  $\mathcal{A}$  or  $\mathcal{B}$  is correct, the test statistic  $T_n$  converges **uniformly** to a standard normal distribution.

Note the tradeoff between modelling and sparsity conditions.

# Relationship to existing literature

- Other proposals from high-dimensional inference in GLMs assume  $\mathcal{A}$  and  $\mathcal{B}$  to be linear, and  $\mathcal{B}$  to be correctly specified and ultra-sparse.

(van de Geer et al., 2014; Belloni et al., 2016; Shah and Bühlmann, 2017)

- By using specific bias-reduction strategies, our tests
  - allow arbitrary conditional mean models for  $\mathcal{A}$  and  $\mathcal{B}$ ,
  - remain valid when  $\mathcal{A}$  or  $\mathcal{B}$  is misspecified,
  - use weaker sparsity assumptions.

# Relationship to existing literature

- Other proposals from high-dimensional inference in GLMs assume  $\mathcal{A}$  and  $\mathcal{B}$  to be linear, and  $\mathcal{B}$  to be correctly specified and ultra-sparse.

(van de Geer et al., 2014; Belloni et al., 2016; Shah and Bühlmann, 2017)

- By using specific bias-reduction strategies, our tests
  - allow arbitrary conditional mean models for  $\mathcal{A}$  and  $\mathcal{B}$ ,
  - remain valid when  $\mathcal{A}$  or  $\mathcal{B}$  is misspecified,
  - use weaker sparsity assumptions.
- Weaker sparsity assumptions do not suffice for Wald tests.

- 1 Introduction
- 2 A dissection of the problem
- 3 Proposal
- 4 Numerical results**
- 5 Discussion

# Simulation study

- $n = 200$
- linear models with  $Z_1, \dots, Z_p$  for  $p = 140$  mutually independent, standard normal variates.
- 19 confounders, generally strongly associated with exposure, and more weakly with outcome.
- No pure exposure predictors.
- 40 pure outcome predictors.
- Covariates explain 80% of the variability in exposure and outcome.
- 1000 simulation experiments.
- Penalty parameters chosen via cross-validation (1 SE).

# Simulation results: $n = 200, p = 100$

## Correct outcome model

Method	Type I error
Standard naïve	0.212
hdm DS	0.470
hdm OI	0.451
Proposal	0.063
Proposal (Unweighted)	0.063

Simulation results:  $n = 200, p = 200$ 

## Correct outcome model

Method	Type I error
Standard naïve	0.399
hdm DS	0.454
hdm OI	0.435
Proposal	0.074
Proposal (Unweighted)	0.087

Simulation results:  $n = 200, p = 100$ 

## Misspecified outcome model

Method	Type I error
Standard naïve	0.156
hdm DS	0.194
hdm OI	0.191
Proposal	0.072
Proposal (Unweighted)	0.059

Simulation results:  $n = 200, p = 200$ 

## Misspecified outcome model

Method	Type I error
Standard naïve	0.266
hdm DS	0.233
hdm OI	0.233
Proposal	0.060
Proposal (Unweighted)	0.067

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- For testing the null, we have shown that **weaker conditions** are attainable without the need for sample-splitting.
- We have extended this to allow for model misspecification.
- This required the use of special ‘bias-reduced’ fitting strategies.

(Vermeulen and Vansteelandt, 2015)

# This talk was based on...

Avagyan, V. and Vansteelandt, S. (2017). Honest data-adaptive inference for the average treatment effect under model misspecification using penalised bias-reduced double-robust estimation. arXiv:1708.03787

Dukes, O., Avagyan, V. and Vansteelandt, S. (2018). High-Dimensional Doubly Robust Inference for Regression Parameters. Technical Report.

Vermeulen, K. and Vansteelandt, S. (2015). Bias-Reduced Doubly Robust Estimation. Journal of the American Statistical Association, 110(511):1024-1036.

# Bias in function of the nuisance parameters

