

# Using the stepwise Neyman-orthogonal estimators in Stata

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# Overview

- This talk is about methods and software for estimating the causal impact of a few covariates on an outcome in a sparse high-dimensional model
- This talk
  - defines sparse high-dimensional models
  - discusses Neyman-orthogonal (NO) estimators for the parameters of interest
  - discusses why use BIC-based stepwise instead of the lasso for an NO estimator
  - discusses the the `swpo` command

# What's a high dimensional model?

- I have an extract of the data Sunyer et al. (2017) used to estimate the effect air pollution on the response time of primary school children

$$\mathbf{E}[\text{htime}_i | \text{no2\_class}_i, \mathbf{x}_i] = \exp(\text{no2\_class}_i \gamma + \mathbf{x}_i \beta)$$

htime            the response time on test of child  $i$  (hit time)

no2\_class       air-pollution level in the school of child  $i$

$\mathbf{x}_i$                 vector of control variables that might need to be included

- I want to estimate the effect no2\_class on htime and a confidence interval for the size of this effect

# High-dimensional models for inference

$$\mathbf{E}[\text{htime}_i | \text{no2\_class}, \mathbf{x}] = \exp(\text{no2\_class}_i \gamma + \mathbf{x}_i \beta)$$

- If the number of covariates in  $\mathbf{x}$  is small relative to the number of observations
  - I can simply include all the controls in  $\mathbf{x}$
- In **high-dimensional** models, there are too many potential control covariates in  $\mathbf{x}$  to reliably estimate  $\gamma$  when all the controls are included
- There are 252 controls in  $\mathbf{x}$ , but I only have 1,036 observations
- I cannot reliably estimate  $\gamma$  if I include all 252 controls

# Potential solutions

$$\mathbf{E}[\text{htime}_i | \text{no2\_class}, \mathbf{x}] = \exp(\text{no2\_class}_i \gamma + \mathbf{x}_i \boldsymbol{\beta})$$

- Suppose that  $\tilde{\mathbf{x}}$  contains the subset of  $\mathbf{x}$  that must be included to get a good estimate of  $\gamma$  for the sample size that I have
- If I knew  $\tilde{\mathbf{x}}$ , I could use the model

$$\mathbf{E}[\text{htime}_i | \text{no2\_class}, \tilde{\mathbf{x}}] = \exp(\text{no2\_class}_i \gamma + \tilde{\mathbf{x}}_i \boldsymbol{\beta})$$

- I am willing to assume the number of variables in  $\tilde{\mathbf{x}}_i$  is small relative to the sample size
  - This is a **sparsity** assumption

- A **high-dimensional** model is one in which there are too many potential covariates, given the sample size
- A **sparse** high-dimensional model is one in which, we only need to include a few of the many potential covariates
  - Few is defined relative to the sample size
- We must solve two problems to do estimation and inference in a sparse high-dimensional model
  - 1 How to select the few important covariates?
  - 2 How to get an estimator that is robust to the first stage covariate selection

# Theory-based model selection

- The traditional approach would be to use theory to determine which covariates should be included
  - Theory tells us to include controls  $\check{x}$
- Poisson quasi maximum likelihood (QML) of `htime` on `no2_class` and controls  $\check{x}$ 
  - Let  $\hat{\gamma}_{\check{x}}$  be estimator with theory-based controls
  - Let  $\hat{\gamma}_{\check{x}}$  be estimator with best-approximating-model controls
  - $\hat{\gamma}_{\check{x}}$  converges to  $\gamma$  but  $\hat{\gamma}_{\check{x}}$  does not converge to  $\gamma$ 
    - Live with large-sample bias from theory-based covariate selection

- Many researchers want to use the lasso and other data-based methods to perform the covariate selection
  - These methods should be able to remove the large-sample bias arising from theory-based covariate selection
- Some post-covariate-selection estimators provide reliable inference for the few parameters of interest

Some do not



# A naive approach

- Naive estimator:
  - ① Use covariate-selection to obtain estimate of which covariates in  $\mathbf{x}$  are in  $\tilde{\mathbf{x}}$   
Denote estimate by `xhat`
  - ② Use QML Poisson to estimate  $\gamma$  and  $\tilde{\beta}$   
`poisson htime no2_class xhat`

# Why naive approach fails

- Unfortunately, naive estimators that use the selected covariates as if they were  $\tilde{\mathbf{x}}$  provide unreliable inference in repeated samples
  - Covariate-selection methods make too many mistakes in estimating  $\mathbf{x}$  when some of the coefficients are small in magnitude
  - Here is an example of small coefficient
    - A nonzero coefficient with a magnitude between 1 and 3 times its standard error is small
  - If your model only approximates the process that generated the data, there are approximation terms
    - The coefficients on some of the approximating terms are probably small
- See Leeb and Pötscher (2005), Leeb and Pötscher (2006), Leeb and Pötscher (2008), and Pötscher and Leeb (2009)

# Missing small-coefficients covariates matters

- It might seem that not finding covariates with small coefficients does not matter
  - But it does
- When some of the covariates have small coefficients, the distribution of the covariate-selection method is not sufficiently concentrated on the set of covariates that best approximates the process that generated the data
  - Covariate-selection methods will frequently miss the covariates with small coefficients causing omitted variable bias
- The random inclusion or exclusion of these covariates causes
  - the distribution of the naive post-selection estimator to be not normal
  - it makes the usual large-sample theory approximation invalid in theory and unreliable in finite samples

# Let's get specific

- The regression function is

$$\mathbf{E}[y|\mathbf{d}, \mathbf{x}] = \exp(\mathbf{d}\boldsymbol{\alpha}' + \tilde{\mathbf{x}}\tilde{\boldsymbol{\beta}}') \quad (1)$$

where

- $\mathbf{d}$  includes the few covariates of interest
- $\tilde{\mathbf{x}}$  is the subset of  $\mathbf{x}$  that belong in the model
  - there are too many covariates in  $\mathbf{x}$  to use the quasi-maximum-likelihood (QML) Poisson estimator for the model

$$\mathbf{E}[y|\mathbf{d}, \mathbf{x}] = \exp(\mathbf{d}\boldsymbol{\alpha}' + \mathbf{x}\boldsymbol{\beta}')$$

- If you knew the subset  $\tilde{\mathbf{x}}$  you could estimate  $\boldsymbol{\alpha}$  and the  $\boldsymbol{\beta}$  the model in (1)

$$\mathbf{E}[y|\mathbf{d}, \mathbf{x}] = \exp(\mathbf{d}\boldsymbol{\alpha}' + \tilde{\mathbf{x}}\tilde{\boldsymbol{\beta}}')$$

A series of seminal papers

- Belloni, Chen, Chernozhukov, and Hansen (2012);
- Belloni, Chernozhukov, and Hansen (2014); and
- Belloni, Chernozhukov, and Wei (2016)  
derived a series of Neyman-orthogonal estimators that provide reliable inference about  $\boldsymbol{\alpha}$
- These estimators use a covariate-selection method to select  $\tilde{\mathbf{x}}$
- The cost of using a covariate-selection method is that these Neyman-orthogonal estimators do not produce estimates for  $\tilde{\boldsymbol{\beta}}$

- When you use two-step estimators, you usually have to adjust your standard errors to account for the parameters you estimated in the first step
  - When you estimate average partial effects, you have to adjust for estimating the coefficients in the first stage
  - Stack the moment conditions
- When you
  - 1 do model selection
  - 2 use the selected model

you have to use an estimator in the second stage that is robust to the model selection mistakes made in the first step

An NO estimator uses moment equations that have had the effect of the first stage model selection removed

- In a linear model NO estimators end up being an extension of the partialing out algorithm we all learned in first regression class
  - Stata calls NO estimators partialling-out estimators
- NO algorithm for

$$y_i = d_i\gamma + \mathbf{x}_i\boldsymbol{\beta} + \epsilon_i$$

- 1 Use selection method to find  $\mathbf{x}_y$  (subset of  $\mathbf{x}$ ) that should be included in model for  $y$
- 2 Let  $\tilde{y}$  be residuals from regressing  $y$  on  $\mathbf{x}_y$
- 3 Use selection method to find  $\mathbf{x}_d$  (subset of  $\mathbf{x}$ ) that should be included in model for  $d$
- 4 Let  $\tilde{d}$  be residuals from regressing  $d$  on  $\mathbf{x}_d$
- 5 Estimate  $\gamma$  from OLS of  $\tilde{y}_x$  on  $\tilde{y}_d$

# Covariate selection

- Methods for covariate selection
  - Best subset regression
    - Compute the BIC, or another IC, for all possible subsets of  $\mathbf{x}$
    - Select the model that minimizes the BIC
    - Infeasible at  $p$  gets large, cannot compute all  $2^p$  estimators
  - One can view the lasso as a feasible convex optimization problem that approximates the best-subset problem
    - The lasso has tuning parameters that must be selected
    - Each method of selecting the lasso tuning parameters is, in effect, a different version of the lasso
  - Stepwise algorithms are another way to approximate the best-subset problem



- Belloni, Chernozhukov, Hansen and coauthors use a particular version of the least absolute shrinkage and selection operator (lasso) to perform covariate selection
  - See Hastie et al. (2015) and Belloni et al. (2012) for introductions to the lasso and the form used by Belloni, Chernozhukov, Hansen and coauthors
- In our papers, we look at using different versions of the lasso and at using BIC-based stepwise

- BIC based stepwise algorithm

- 1 Let  $\mathbf{x}_f$  be the full set of potential covariates
- 2 Let  $\mathbf{x}_{in}$  be the covariates to include in the model
  - At the start let  $\mathbf{x}_{in}$  include the constant term
- 3 Let  $BIC_c$  be the BIC for the current model of QML of  $y$  on  $\mathbf{x}_{in}$
- 4 For each covariate  $j$  in  $\mathbf{x}_f$ , let  $BIC_j$  be the for the model of  $y$  on  $\mathbf{x}_{in}$  and  $x_j$
- 5 Let  $\tilde{j}$  the  $j$  that yields the smallest  $BIC_j$
- 6 If  $BIC_{\tilde{j}} < BIC_c$ , then
  - add  $x_{\tilde{j}}$  to  $\mathbf{x}_{in}$
  - remove  $x_{\tilde{j}}$  from  $\mathbf{x}_f$
  - let  $BIC_c = BIC_{\tilde{j}}$
  - go to step 4

else

exit

- See Drukker and Liu (2021) and citations therein for more details

# Why consider forward stepwise

- Drukker and Liu (2021)
  - discuss a family of data generating processes (DGPs) for which the lasso fails to select the covariates  $\tilde{\mathbf{x}}$  in finite samples
  - present simulation evidence that a BIC-based forward stepwise method **can** reliably select the  $\tilde{\mathbf{x}}$  from  $\mathbf{x}$  for DGPs in this family
  - present simulation evidence that a testing-based forward stepwise method **cannot** reliably select the  $\tilde{\mathbf{x}}$  from  $\mathbf{x}$  for DGPs in this family
- Using a BIC-based forward stepwise method takes longer than lasso-based methods
  - Can take **much** longer
  - You are trading time for selection accuracy for some DGPs

- Iterated sure independence screening (SIS) uses a first step that removes variables that have no marginal predictive power. The iterative process puts back the variables that have conditional predictive power and removes the ones that were false included in the first step.
- We are currently looking into using a version of iterated SIS to reduce the computation time of BIC-based forward-stepwise NO estimators
  - Fan and Lv (2008), Fan et al. (2009), and Fan and Song (2010) provide introductions to iterative SIS

## Use extract of data from Sunyer et al. (2017)

```
. use breathe7, clear  
. describe
```

Contains data from breathe7.dta

```
Observations:      1,089  
Variables:         20
```

22 Sep 2021 14:39

Variable name	Storage type	Display format	Value label	Variable label
htime	double	%10.0g		ANT: mean hit reaction time (ms)
no2_class	float	%9.0g		Classroom NO2 levels (g/m3)
sev_sch	float	%9.0g		School vulnerability index
noise_sch	float	%9.0g		Measured school noise (in dB)
age	float	%9.0g		Child's age (in years)
ppt	double	%10.0g		Daily total precipitation
grade	byte	%9.0g	grade	Grade in school
sex	byte	%9.0g	sex	Sex
age_start_sch	double	%4.1f		Age started school
oldsibl	byte	%1.0f		Older siblings living in house
youngsibl	byte	%1.0f		Younger siblings living in house
lbfeed	byte	%19.0f	bfeed	duration of breastfeeding
smokep	byte	%3.0f	noyes	1 if smoked during pregnancy
feduc4	byte	%17.0g	edu	Paternal education
meduc4	byte	%17.0g	edu	Maternal education
sev_home	float	%9.0g		Home vulnerability index
no2_home	float	%9.0g		Residential NO2 levels (g/m3)
overwt_who	byte	%32.0g	over_wt	WHO/CDC-overweight 0:no/1:yes
ndvi_mn	double	%10.0g		Home greenness (NDVI), 300m buffer
lbweight	float	%9.0g		1 if low birthweight

Sorted by:

# Potential Controls I

```
. local ccontrols "sev_home sev_sch age no2_home ppt ndvi_mn noise_sch"  
. local fcontrols "grade sex meduc4 "  
. local allcontrols "c.(`ccontrols`) i.(`fcontrols`) "  
. local allcontrols "`allcontrols` i.(`fcontrols`)#c.(`ccontrols`) "
```

## BIC-stepwise-based results

```
. posw htime no2_class, controls(`allcontrols`) model(poisson) method(bic)
select controls for htime using stepwise bic
select controls for no2_class using stepwise bic
Partialing-out stepwise bic
```

Number of obs	=	1,084
Number of controls	=	79
Number of selected controls	=	45
Wald chi2(1)	=	30.92
Prob > chi2	=	0.0000

Model: poisson

htime	Coefficient	Robust std. err.	z	P> z	[95% conf. interval]	
no2_class	.0034337	.0006175	5.56	0.000	.0022234	.0046439

Note: Chi-squared test is a Wald test of the coefficients of the variables of interest jointly equal to zero.

```
. nlcom exp(_b[no2_class])
      _nl_1: exp(_b[no2_class])
```

htime	Coefficient	Std. err.	z	P> z	[95% conf. interval]	
_nl_1	1.00344	.0006196	1619.47	0.000	1.002225	1.004654

Another microgram of NO<sub>2</sub> per cubic meter increases the mean reaction time by about 0.3%

## lasso-based results

```
. popoisson htime no2_class, controls(`allcontrols`) coef
```

```
Estimating lasso for htime using plugin
```

```
Estimating lasso for no2_class using plugin
```

```
Partialing-out Poisson model      Number of obs      =      1,084
                                   Number of controls     =          79
                                   Number of selected controls =          10
                                   Wald chi2(1)              =      29.40
                                   Prob > chi2               =      0.0000
```

	htime	Coefficient	Robust std. err.	z	P> z	[95% conf. interval]
no2_class		.0032534	.0006	5.42	0.000	.0020773 .0044294

Note: Chi-squared test is a Wald test of the coefficients of the variables of interest jointly equal to zero. Lasso selects controls for model estimation. Type `lassoinfo` to see number of selected variables in each lasso.

```
. nlcom exp(_b[no2_class])
```

```
  _nl_1: exp(_b[no2_class])
```

	htime	Coefficient	Std. err.	z	P> z	[95% conf. interval]
_nl_1		1.003259	.000602	1666.56	0.000	1.002079 1.004439

Another microgram of NO<sub>2</sub> per cubic meter increases the mean

23 / 24 ion time by about 0.20%



# Conclusions

- So far
  - Sparse high-dimensional models require covariate selection
  - You must use an NO estimator to account for covariate selection
  - There are DGPs for which an NO estimator that uses BIC-stepwise will perform well, but an NO estimator that uses lasso will not perform well
- Future
  - Use iterated SIS combined with BIC stepwise to get dramatically faster but just as accurate results

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