Week 14: Bootstrap

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Updated notes are here: https://clas.ucdenver.edu/marcelo-perraillon/ teaching/health-services-research-methods-i-hsmp-7607

Outline

- Review of standard errors
- The magic of bootstrapping
- Caveats

Standard error, reminder

- Back in the days (week 4 of this class) we covered **standard errors**
- We have an estimator, say, the mean of a sample $\bar{X} = \sum_{i=1}^{n} X_i / n$ or a proportion \hat{p}
- The parameter estimate has some error and a distribution (not the same as the standard deviation of the data)
- If we know the distribution of the parameter and its standard error, then we can build confidence intervals and do hypothesis testing
- In the context of linear regression, we know that $\hat{\beta}_j$ distributes normal and we have a formula for its standard error (the **variance-covariance** matrix, really). We know this because of **theory**

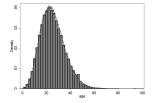
Derivation

- We use statistical theory to derive standard errors
- In the linear model, we use the central limit theorem, the law of large numbers, and the assumption of iid errors that are normally distributed
- We needed all that to come up with formulas for the standard error
- The logic of standard errors is a lot easier to understand using simulations

- Say that we have a population of 40,000 observations
- We will take a sample of 150 observations out of the 40,000 (recall, that in theory, we assume that the population is **infinitely** large)
- We will take the mean of the 150 observations
- If we could repeat this experiment many times, we could calculate the mean many times and see how it distributes (that's why this way of thinking about statistics is called frequentist)
- Keep in mind: we want to understand how the mean distributes, not the distribution of the 150 observations or the 40,000
- We will do this 1,000 times. In a real life example, we can't do this. We just get a sample of 150 observations. We can't repeat the experiments many times

- We will create a population of 40,000 people and simulate their "age" with (N(5,1))²
- I take the square to avoid negative ages; it will distribute Chi-square. Also, I'll remove the decimals

```
clear
    seed 1234567
set obs 40000
gen age = int((rnormal(5,1))^2)
sum age
    Variable |
                       Obs
                                          Std. Dev.
                                                           Min
                                  Mean
                                                                       Max
                                          10.12632
         age |
                   40,000
                              25,49595
                                                             1
                                                                        95
```



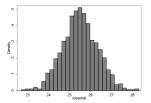
- Next, we take a sample of 150 and calculate the mean
- We repeat 1,000 times so we have a distribution for the mean and calculate the standard deviation of the means (i.e. the standard error)
- It will take a while...

```
postfile buffer meanhat using sampmean, replace
forvalues i=1/1000 {
    preserve
    sample 150, count
    qui sum age
    post buffer (r(mean))
    restore
  }
postclose buffer
```

I'm not bootstrapping here. This is about understanding the standard error

We can see how the means distribute and what is the standard deviation (standard error)

use sampmean hist meanhat		(med.gph	, replace)				
sum meanhat Variable	I -+	Obs	Mean	Std.	Dev.	Min	Max
meanhat graph export	•	1,000 , replac	25.49305 e	.842	0507	22.69333	28.30667



So what did we learn?

- Even though the data does not distribute normal (it had a Chi-square distribution) the means of the 150 do distribute normal
- The standard error (the standard deviation of the means) is 0.84
- With that information we could do hypothesis testing. For example, we know that 95% of the values are within 2 standard deviations
- If say, I use the first sample of 150 with mean 26.73, the 95 CI is [26.73 2 * 0.84, 26.73 + 2 * 0.84] = [25.05, 28.41]. We would reject the null that the mean is 29, for example
- (Recall, though, that we use the t-distribution because we have to estimate the standard error)

So what did we learn?

• Of course, we don't do simulations in practice since we can't and know that the **theoretical** SE of the mean is $\frac{\hat{\sigma}}{\sqrt{N}}$, where $\hat{\sigma}$ is the standard deviation of the sample

■ For example, we can use just 1 sample to get an approximation: Variable | Obs Mean Std. Dev. Min Max age | 150 26.73333 9.95673 4 54 di 9.957649 /sqrt(150) .81303864

Theory gives us a formula for the standard error and a distribution. With simulations, we found that it was 0.84. With theory, we got 0.81

What if we don't have theory?

- What happens when we don't have theory to tell us what is the standard error?
- We collect a sample and have an estimator but we don't know its standard error either because we don't know how to derive the theoretical SE or because there is no formula for it
- We can't use simulations because we do not know the true model; we just used simulations to understand the logic behind the theory
- This is when the **bootstrap** is truly like *magic*

Nonparametric bootstrap

- Suppose a new situation (that is slightly more realistic)
- We have a sample of 150 people and we calculate mean age but let's assume that we do not know the formula for the standard error of the mean
- How could we come up with an approximation for the standard error using the data?
- Enters the **bootstrap**
- I'll show you how the bootstrap works before we try to understand why it works

Nonparametric bootstrap

- We won't simulate from any distribution. We will resample with replacement. We will resample our sample of 150 observations
- We will use the 150 observations and obtain a sample with replacement so we have another set of 150 observations
- We will take the mean of the 150 observations and save it
- We will repeat this process 3000 times and use the 3000 means to calculate their standard deviation and distribution

Sampling with replacement

- Sampling with replacement can be confusing
- Suppose you have ten numbers: 2, 4, 6, 10, 3, 11, 20, 40, 13,1
- If we sample 10 numbers with replacement, we could get: 2, 4, 4, 4, 11, 1, 20, 6, 6, 2
- In other words, just a combination of the same numbers, some of them repeated but most likely not the same numbers
- Sampling 10 numbers out of those 10 numbers without replacement would imply getting the same exact 10 numbers

Stop here for a bit

- Make sure you understand what is different here from the simulations
- We are not drawing a random sample from a distribution
- We are using our **sample** to take other samples of the **same size**
- It can be hard to understand this distinction and even harder to understand why it works

Example bootstrap

- I saved one sample of 150 in a dataset called s150.dta
- We want to calculate the SE of the mean because we are pretending we don't know the formula for the standard error

use s150,clear sum age					
Variable	Obs	Mean	Std. Dev.	Min	Max
age	150	25.49595	10.70073	1	95
* Theoretical SE . di 10.70073/sqrt .87371095	(150)				

■ In this sample, the theoretical error is 0.87

Example bootstrap

- Resample from the 150 with replacement to get another sample of size 150
- Again, it's not going to be the same 150 observations, each will be different
- Take the mean and save it; repeat 3000 times

```
postfile buffer meanhat using sampmean_b, replace
forvalues i=1/3000 {
  preserve
  bsample 150
  qui sum age
  post buffer (r(mean))
restore
3
postclose buffer
use sampmean_b, clear
sum
   Variable |
                  Obs
                           Mean
                                  Std Dev
                                               Min
                                                        Max
meanhat
             3,000
                        25,4795
                                  .8323291
                                             22.52
                                                    28.56667
```

Our boostrapped SE is .8323291, which is close to theoretical SE.
 MAGIC

Another example

- You don't need to write your own program most of the time
- Stata has a bootstrap command
- We will use the auto dataset

Auto dataset

Auto dataset

```
sysuse auto, clear
(1978 Automobile Data)
```

reg price mpg turn

Source	SS SS	df	MS	Number of	obs =	74
	+			F(2, 71)	=	10.08
Model	140436412	2	70218206.1	Prob > F	=	0.0001
Residual	494628984	71	6966605.41	R-squared	=	0.2211
	+			Adj R-squa	red =	0.1992
Total	635065396	73	8699525.97	Root MSE	=	2639.4
price	Coef.	Std. Err.	t	P> t [95	% Conf.	Interval]
	+					
mpg	-259.6967	76.84886	-3.38	0.001 -41	2.929	-106.4645
turn	-38.03857	101.0624	-0.38	0.708 -239	.5513	163.4742
_cons	13204.27	5316.186	2.48	0.015 2	604.1	23804.45

■ We do have theory and we do have a formula for the SEs here...

Auto dataset

Let's bootstrap them anyway

bootstrap, reps(1000): regress price mpg turn
(running regress on estimation sample)

Bootstrap replications (1000) + 1+ 2+ 3+ 4+ 5 							
Linear regression				Wald chi Prob > c R-square	cions i2(2) chi2 ed quared	= = =	74 2,000 14.53 0.0007 0.2211 0.1992 2639.4328
 price mpg turn _cons	Coef. -259.6967 -38.03857		-2.48	0.013 0.769	[95% C -465.19 -291.63	Conf. 39 339	-

MAGIC!!!

Not impressed?

- Perhaps you are not too impressed because the SEs are not that close in this example
- But check the data more carefully; the auto dataset has only 74 observations
- What about if we try the same with more data?
- But first, why more data would be better? The way I make sense of it is that with more data we have more realizations or examples of values so it is like we were drawing random samples repeatedly, just like what we did getting at the beginning of the class
- Let's use the beauty dataset

Beauty dataset

Beauty dataset has more observations

reg lwage abvavg exper looks union

Source	SS	df	MS		r of obs	=	1,260
 Model Residual	59.4988269 385.481145	4 1,255	14.8747067 .307156291	Prob	> F	-	48.43 0.0000 0.1337
+- Total	444.979972	1,259	.353439215	Adj R	-squared	=	0.1310
lwage	Coef.	Std. Err.		P> t	[95% Co	onf.	Interval]
abvavg	1600523	.0618286		0.010	281351	2	0387534
exper	.0152176	.0013286	11.45	0.000	.01261	1	.0178241
looks	.1874543	.0413931	4.53	0.000	.10624	17	.2686616
union	.1986142	.0353455	5.62	0.000	.129271	15	.2679569
_cons	.7791512	.1212421	6.43	0.000	.541291	.7	1.017011

Beauty dataset

Bootstrap the SEs

bootstrap, reps(2000): reg lwage abvavg exper looks union (running regress on estimation sample)

Linear regression

Number of obs	=	1,260
Replications	=	2,000
Wald chi2(4)	=	216.74
Prob > chi2	=	0.0000
R-squared	=	0.1337
Adj R-squared	=	0.1310
Root MSE	=	0.5542

1	Observed	Bootstrap			Normal	-based
lwage	Coef.	Std. Err.	z	P> z	[95% Conf.	Interval]
+						
abvavg	1600523	.0632131	-2.53	0.011	2839476	036157
exper	.0152176	.0013484	11.29	0.000	.0125747	.0178604
looks	.1874543	.0429811	4.36	0.000	.1032128	.2716958
union	.1986142	.0322598	6.16	0.000	.1353861	.2618423
_cons	.7791512	.1255793	6.20	0.000	.5330203	1.025282

■ As I said, MAGIC!

When do we use bootstrapped SEs?

- We use them when we don't have theory to guide us
- The classic example: no theoretical SE for the median
- You will use them next semester when doing some versions of instrumental variables and propensity scores
- There are several variants of bootstrap (jackknife, parametric)
- Why does it work? Well, because, apparently, resampling from a sample with replacement is like sampling from a population; it works better when the sample itself is not small
- In other words, resampling with replacement from the 150 observations is like sampling from the 40,000 observations (the population)
- Active area of research

Median

- As I said, there is no good theoretical formula for the standard error of the median
- So bootstrapping is a good option. I'll do it the longer way, making my own program

```
seed seed 12354
sysuse auto, clear
* Write a command call mymedian
program mymedian, rclass
version 14.2
args x
qui sum 'x', det
return scalar med = r(p50)
end
bootstrap r(med), reps(1000): mymedian price
```

Median

Output

	ootstrap replications (1000) + 1+ 2+ 3+ 4+ 5						
				50			
••••				1000			
Bootstrap resul	ts			Number of	obs	=	74
				Replicati	ons	=	1,000
command:	mymedian pr	rice					
_bs_1:	r(med						
	01					Vormal·	d
-		Bootstrap					
I	Coei.	Std. Err.	z			Conf.	Intervalj
bs_1	5006.5	270.0553	18.54	0.000	4477	. 201	5535.799

Median - the shortest way

Using the summarize command directly

bootstrap r(p50), reps(1000): summarize price, detail (running summarize on estimation sample) Bootstrap replications (1000) + 1 + 3 50							
Bootstrap resul	ts			Number of Replicati			74 1,000
	summarize p r(p50)	orice, detail					
	Observed	Bootstrap				lormal	 -based
	Coef.	Std. Err.			[95%	Conf.	Interval]
		257.0563					

Pesky details

. estat bootstrap. all

But what about the distribution of the estimate? We know the SE but we need to know how it distributes as well

Bootstrap results Number of obs 74 Replications = 1000 command: summarize price, detail bs 1: r(p50) Observed Bootstrap Coef Bias Std. Err. [95% Conf. Interval] _bs_1 | 5006.5 7.681 257.05629 4502.679 5510.321 (N) 4603.5 5708.5 (P) 4647 5719 (BC) (N) normal confidence interval (P) percentile confidence interval bias-corrected confidence interval (BC)

Summary

- Active area of research
- Extremely useful in some situations, although for most applied research we don't need it because there is theory
- The idea is so simple yet so powerful