#### Adventures in Simulation in R

EC 425/525, Lab 6

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# Prologue

# Schedule

#### Last time

Plotting

### Today

Simulation

#### Motivation

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You just need to be careful to **ask a clear, answerable question** and then **run a simulation** that corresponds/answers this question.

In addition, simulations can be computationally intense—they are often the first time you have to really think about efficiency in coding.

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  - **Sample** from your population.
  - Construct estimates/inferences that relate to your original question.

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- 2. Iterate. In each iteration:
  - **Sample** from your population.
  - Construct estimates/inferences that relate to your original question.
- 3. **Summarize** results.

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- 3. Writing a function for a single iteration can be helpful (see above).
- 4. There is a (big) difference between unbiasedness and consistency.
- 5. You build simulations/DGPs with assumptions.
- 6. Analytical results can inform and/or replace simulations.

# Example simulation

#### The question

Q We've shown that instrumental variables (IV) is consistent, how does it perform (*i.e.*, is it unbiased) in finite (small) samples?

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Q We've shown that instrumental variables (IV) is consistent, how does it perform (*i.e.*, is it unbiased) in finite (small) samples?

Note This question is definitely answerable analytically.

Nevertheless, let's see how IV performs at several small-ish sample sizes.

While we're at it, let's confirm OLS is indeed biased in this setting.

#### **DGP**

We want a valid instrument for a setting in which treatment is endogenous.

$$Y_i = \alpha + \tau D_i + \varepsilon_i$$

So we want

- 1. Endogenous treatment:  $Cov(D_i, \, \varepsilon_i) \neq 0$
- 2. Predictive:  $Cov(Z_i, D_i) \neq 0$
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  eq 0$
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- 3. Excludability:  $Cov(\mathbf{Z}_i, \, \varepsilon_i) = 0$

where (2) and (3) imply  $\mathbf{Z}_i$  is a valid instrument.

#### **DGP**

In other words, the variance-covariance matrix of  $D_i$ ,  $\varepsilon_i$ , and  $Z_i$  is

$$\Sigma = egin{bmatrix} \sigma_{ ext{D}}^2 & \sigma_{ ext{D},arepsilon} & \sigma_{ ext{D}, ext{Z}} \ \sigma_{ ext{D},arepsilon} & \sigma_{arepsilon}^2 & 0 \ \sigma_{ ext{D}, ext{Z}} & 0 & \sigma_{ ext{Z}}^2 \end{bmatrix}$$

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If we assume unit variances and covariances are 0.6, then

$$\Sigma = egin{bmatrix} 1 & 0.6 & 0.6 \ 0.6 & 1 & 0 \ 0.6 & 0 & 1 \end{bmatrix}$$

#### **DGP**

To simplify our lives, let's assume that  $D_i$ ,  $\varepsilon_i$ , and  $Z_i$  come from a multivariate normal distribution.

We defined their covariance matrix. We need to define their means.

$$\mu_{
m D}=10$$
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m D}=10$$
,  $\mu_{arepsilon}=0$ , and  $\mu_{
m Z}=3$ .

Finally, we need to define the way in which  $D_i$  and  $\varepsilon_i$  affect  $Y_i$ .

$$Y_i = 7 + 1 \times D_i + \varepsilon_i$$

i.e., 
$$au=1$$
.

#### **DGP**

Lucky for us, R's MASS package has a function mvrnorm() that draws n random observations from a multivariate normal distribution with means mu and variance-covariance matrix Sigma.

### Sampling from our DPG

We're ready to write a function that performs one iteration.

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```
sim_iter \leftarrow function(n) {
  # Define our variance-covariance matrix (D, \varepsilon, Z)
  \Sigma \leftarrow \text{matrix}(\text{data} = c(1, 0.6, 0.6, 0.6, 1, 0, 0.6, 0, 1), ncol = 3)
  # Our vector of means (D, \varepsilon, Z)
  \mu = c(10, 0, 3)
  # Draw n observations; convert to tibble
  sample_df \leftarrow MASS::mvrnorm(n = n, mu = \mu, Sigma = \Sigma) %>% tibble()
  # Name variables
  names(sample_df) \leftarrow c("D", "\epsilon", "Z")
  # Calculate Y
  sample_df %\diamond% mutate(Y = 7 + 1 * D + \epsilon)
```

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Now we just need to estimate  $\beta_{\rm IV}$  and  $\beta_{\rm OLS}$ . We'll use estimatr.

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Previous OLS estimates of the effect of x on y

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lm_robust(y \sim x)
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Now we just need to estimate  $\beta_{IV}$  and  $\beta_{OLS}$ . We'll use estimatr.

Previous OLS estimates of the effect of x on y

```
lm_robust(y ~ x)
```

New IV estimates of the effect of x on y with instrument z

```
iv_robust(y ~ x | z)
```

```
sim iter \leftarrow function(n) {
  # Define our variance-covariance matrix (D, \varepsilon, Z)
  \Sigma \leftarrow \text{matrix}(\text{data} = c(1, 0.6, 0.6, 0.6, 1, 0, 0.6, 0, 1), \text{ncol} = 3)
  # Our vector of means (D, \varepsilon, Z)
  \mu = c(10, 0, 3)
  # Draw n observations; convert to tibble
  smpl df \leftarrow MASS::mvrnorm(n = n, mu = \mu, Sigma = \Sigma) %>% data.frame()
  # Name variables
  names(smpl df) \leftarrow c("D", "\epsilon", "Z")
  # Calculate Y
  smpl_df \%\% mutate(Y = 7 + 1 * D + \epsilon)
  # Fstimates
  est df \leftarrow bind rows(
    # The OLS estimates
    lm_robust(Y ~ D, data = smpl_df) %>% tidy() %>% mutate(est = "OLS"),
    # The TV estimates
    iv robust(Y ~ D | Z, data = smpl df) %>% tidy() %>% mutate(est = "IV")
  return(est_df)
```

#### Repeat

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The output of sim\_iter() is a data frame, so we can actually use a function from furrr that expects outputted data frames, namely, future\_map\_dfr.

The suffix \_dfr means the function will row-bind the data frames returned by individual iterations.

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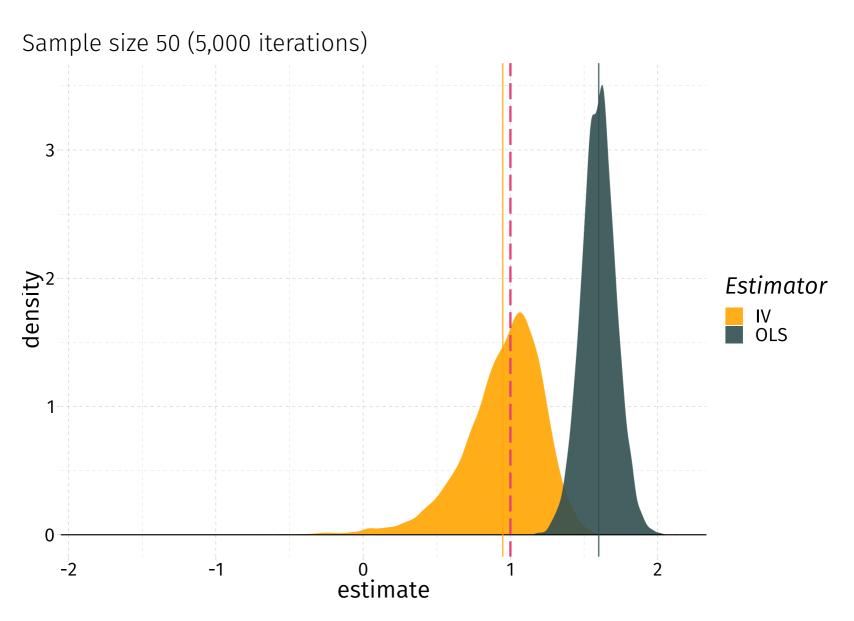
The output of sim\_iter() is a data frame, so we can actually use a function from furrr that expects outputted data frames, namely, future\_map\_dfr.

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We'll also use the rep() function which repeats things, e.g., rep("a", 3) repeats "a" three times.

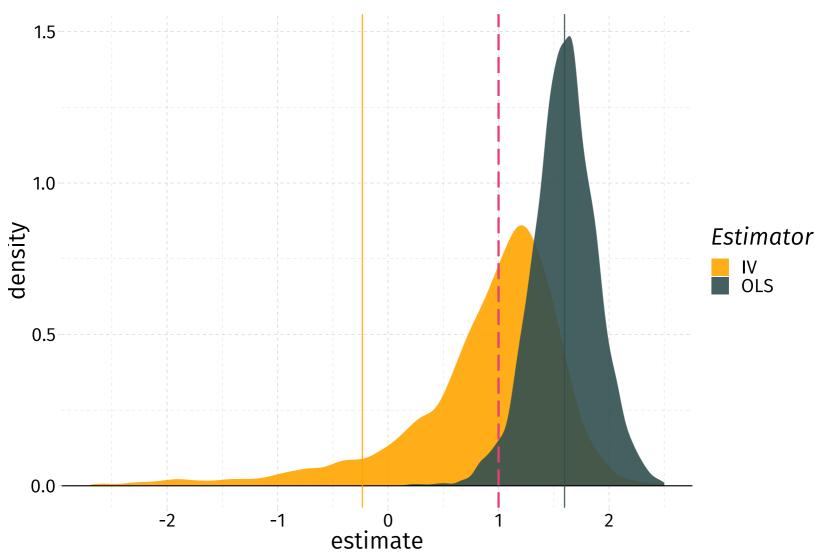
Assuming we've already entered sim\_iter() into memory, we can run our simulation 5,000 times, each with sample size 50—in parallel!

```
# Load furrr
p load(furrr)
# Tell R to parallelize with 4 cores
plan(multiprocess, workers = 4)
# Set a seed
set.seed(12345)
# Run simulation with sample size 50
sim50 ← future map dfr(
  # Repeat sample size 50 for 5000 times
  rep(50, 5000),
  # Our function
  sim iter,
  # Let furrr know we want to set a seed
  .options = future options(seed = T)
```

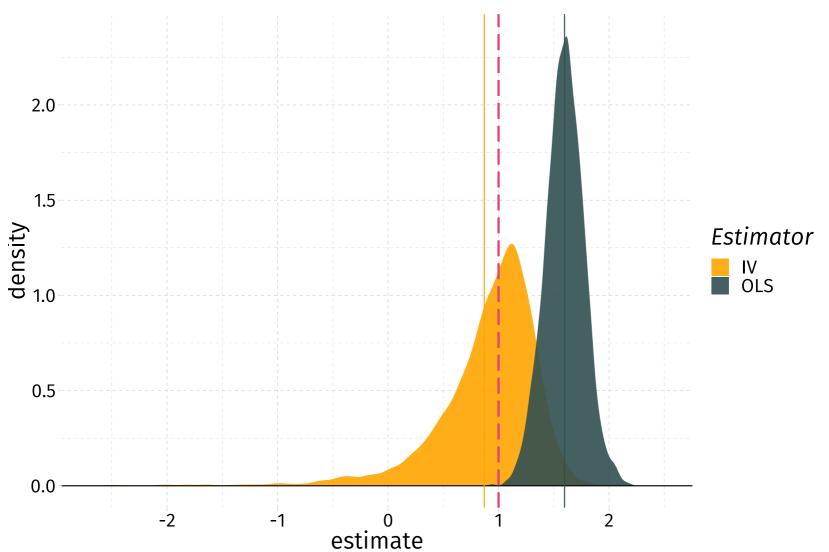


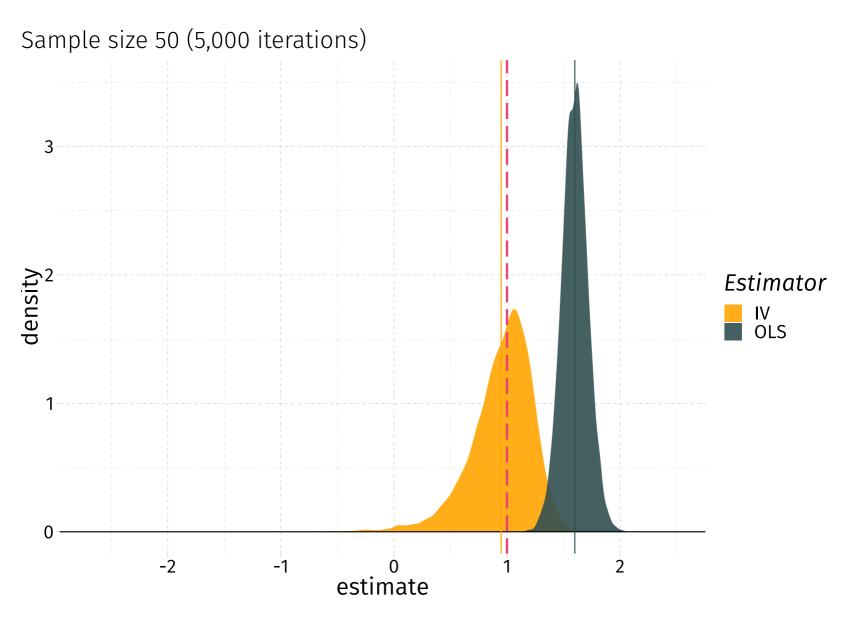
Let's vary the sample size and see what happens.

#### Sample size 10 (5,000 iterations)

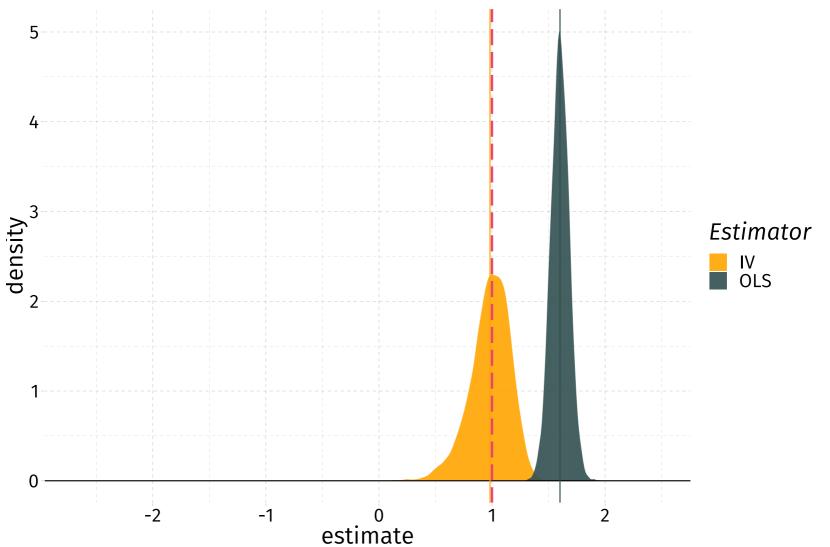


#### Sample size 25 (5,000 iterations)





#### Sample size 100 (5,000 iterations)



### **Assumptions**

Keep in mind that we made several assumptions about

- the distribution (joint normality is very restrictive)
- variance (all equal, independent, and homoskedastic)
- covariances (again, all equal)
- strong instrument

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