Inference and Simulation

EC 607, Set 04

Edward Rubin

Prologue

Schedule

Last time

The CEF and least-squares regression

Today

Inference

Read MHE 3.1

Upcoming

Lab: TBD

Problem set 002 coming soon.

Class project, step 1 due on May 1.

Why?

Q What's the big deal with inference?

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A We rarely know the CEF or the population (and its regression vector).

We can draw statistical inferences about the population using samples.

Important The issue/topic of statistical inference is separate from causality.

Separate questions

- 1. How do we interpret the estimated coefficient $\hat{\beta}$?
- 2. What is the sampling distribution of $\hat{\beta}$?

Moving from population to sample

Recall The population-regression function gives us the best linear approximation to the CEF.

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Moving from population to sample

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$$eta = E\left[\mathbf{X}_i\mathbf{X}_i'\right]^{-1}E[\mathbf{X}_i\mathbf{Y}_i]$$

which we estimate via the ordinary least squares (OLS) estimator[†]

$$\hat{eta} = \left(\sum_i \mathrm{X}_i \mathrm{X}_i'
ight)^{-1} \left(\sum_i \mathrm{X}_i \mathrm{Y}_i
ight)^{-1}$$

† MHE presents a method-of-moments motivation for this derivation, where $\frac{1}{n}\sum_i \mathbf{X}_i \mathbf{X}_i'$ is our sample-based estimated for $E[\mathbf{X}_i \mathbf{X}_i']$. You've also seen others, e.g., minimizing MSE of \mathbf{Y}_i given \mathbf{X}_i .

A classic

However you write it, this OLS estimator

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Note I'm following MHE in defining $e_i = \mathrm{Y}_i - \mathrm{X}_i' \beta$.

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has asymptotic covariance

$$E\left[\mathrm{X}_{i}\mathrm{X}_{i}^{\prime}
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which we estimate by (1) replacing e_i with $\hat{e}_i = Y_i - X_i'\hat{\beta}$ and (2) replacing expectations with sample means, e.g., $E\left[X_iX_i'e_i^2\right]$ becomes $\frac{1}{n}\sum\left[X_iX_i'\hat{e}_i^2\right]$.

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Standard errors of this flavor are known as heteroskedasticity-consistent (or -robust) standard errors (or Eicker-Huber-White).

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Now, returning to to the asym. covariance matrix of $\hat{\beta}$,

$$egin{aligned} E\left[\mathrm{X}_{i}\mathrm{X}_{i}^{\prime}
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If the CEF is nonlinear, then our linear approximation (linear regression) generates heteroskedasticity.

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Thus, even if $Y_i \mid X_i$ has contant variance, $e_i \mid X_i$ is heteroskedastic.

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Thus, even if $\mathbf{Y}_i \mid \mathbf{X}_i$ has contant variance, $e_i \mid \mathbf{X}_i$ is heteroskedastic. Unless you want to assume the CEF is *linear*.

Two notes

1. Heteroskedasticity is **not our biggest concern** in inference.

...as an empirical matter, heteroskedasticity may matter very little... If heteroskedasticity matters a lot, say, more than a 30 percent increase or any marked decrease in standard errors, you should worry about possible programming errors or other problems. (MHE, p.47)

2. Notice that we've **avoided "standard" stronger assumptions**, *e.g.*, normality, fixed regressors, linear CEF, homoskedasticity.

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2. Notice that we've **avoided "standard" stronger assumptions**, e.g., normality, fixed regressors, linear CEF, homoskedasticity.

Following (2): We only have large-sample, asymptotic results (consistency) rather than finite-sample results (unbiasedness).

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One practical way we can study the behavior of an estimator: **simulation**.

Note You need to make sure your simulation can actually test/respond to the question you are asking (e.g., bias vs. consistency).

Simulation

Let's compare false- and true-positive rates[†] for

- 1. Homoskedasticity-assuming standard errors $\left(\operatorname{Var}[e_i | \mathrm{X}_i] = \sigma^2 \right)$
- 2. Heteroskedasticity-robust standard errors

[†] The false-positive rate goes by many names; another common name: type-I error rate.

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Simulation outline

- 1. Define data-generating process (DGP).
- 2. Choose sample size n.
- 3. Set seed.
- 4. Run 10,000 iterations of
 - a. Draw sample of size n from DGP.
 - b. Conduct inference.
 - c. Record inferences' outcomes.

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$$\mathrm{Y}_i = 1 + e^{0.5 \mathrm{X}_i} + arepsilon_i$$

where $\mathrm{X}_i \sim \mathrm{Uniform}(0,10)$ and $arepsilon_i \sim N(0,1)$.

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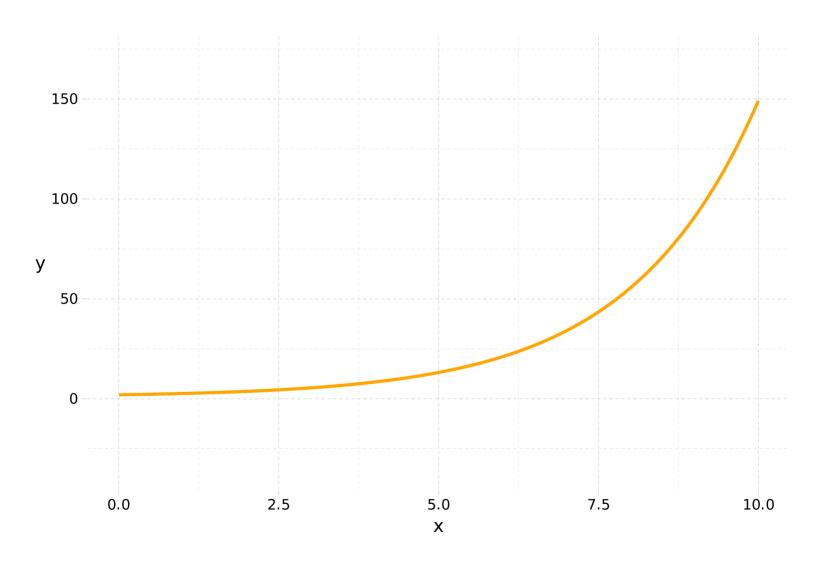
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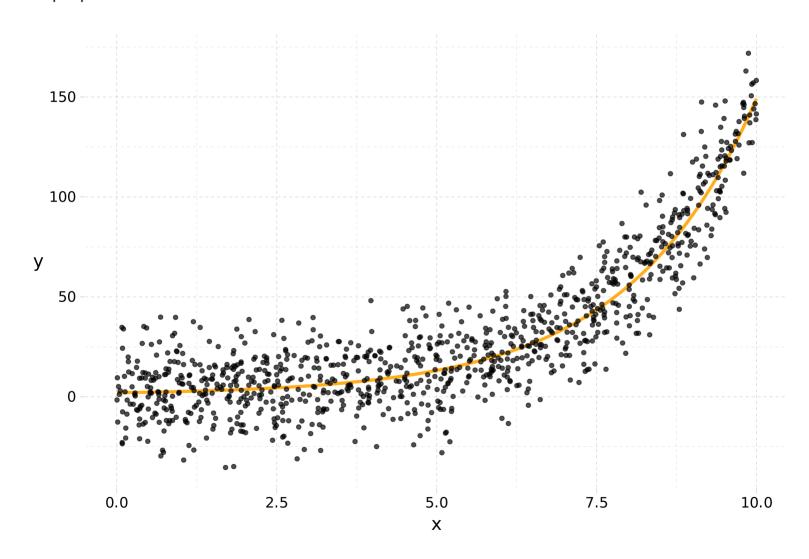
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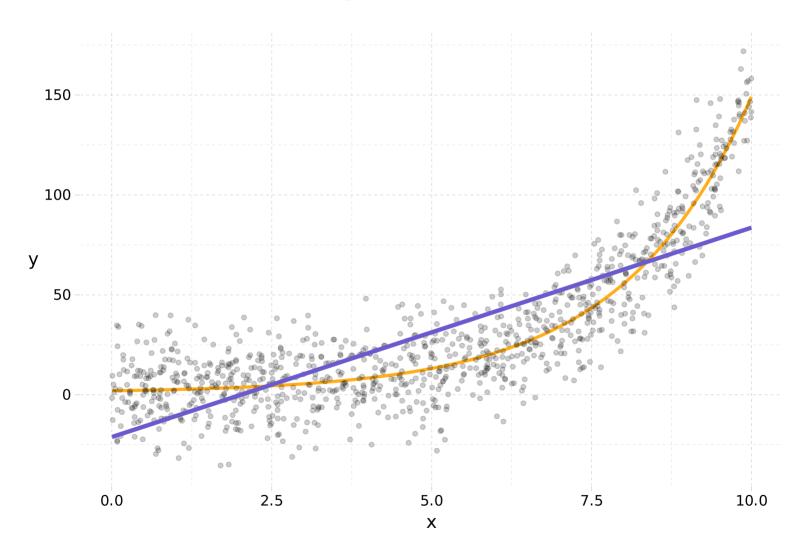
Our CEF



Our population



The population least-squares regression line



Iterating

To make iterating easier, let's wrap our DGP in a function.

We still need to run a regression and draw some inferences.

Note We're defaulting to size-30 samples.

We will use Im_robust() from the estimatr package for OLS and inference.

- se_type = "classical" provides homoskedasticity-assuming SEs
- se_type = "HC2" provides heteroskedasticity-robust SEs

† lm() works for "spherical" standard errors but cannot calculate het.-robust standard errors.

Inference

Now add these estimators to our iteration function...

```
fun iter = function(iter, n = 30) {
  # Generate data
  iter df = tibble(
    \epsilon = rnorm(n, sd = 15),
    x = runif(n, min = 0, max = 10),
    v = 1 + \exp(0.5 * x) + \epsilon
  # Estimate models
  lm1 = lm robust(y ~ x, data = iter df, se type = "classical")
  lm2 = lm_robust(y ~ x, data = iter df, se type = "HC2")
  # Stack and return results
  bind rows(tidy(lm1), tidy(lm2)) %>%
    select(1:5) \%>\% filter(term = "x") \%>\%
    mutate(se_type = c("classical", "HC2"), i = iter)
```

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There are a lot of ways to run a single function over a list/vector of values.

- lapply(), e.g., lapply(X = 1:3, FUN = sqrt)
- for(), *e.g.*, for (x in 1:3) sqrt(x)
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We're going to go with map() from the purrr package because it easily parallelizes across platforms using the furrr package.

Run it!

Run our function 10,000 times

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# Packages
p_load(purrr)
# Set seed
set.seed(12345)
# Run 10,000 iterations
sim_list = map(1:1e4, fun_iter)
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# Packages
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# Set options
set.seed(123)
# Tell R to parallelize
plan(multisession)
# Run 10,000 iterations
sim_list = future_map(
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    .options = furrr_options(seed = T)
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The furrr package (future + purrr) makes parallelization easy and fun!

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The furr package (future + purrr) makes parallelization easy and fun!

Note Use multisession or multicore instead of multiprocess.

Run it!!

Our fun_iter() function returns a data.frame, and future_map() returns a list (of the returned objects).

So sim_list is going to be a list of data.frame objects. We can bind them into one data.frame with bind_rows().

```
# Bind list together
sim_df = bind_rows(sim_list)
```

Run it!!

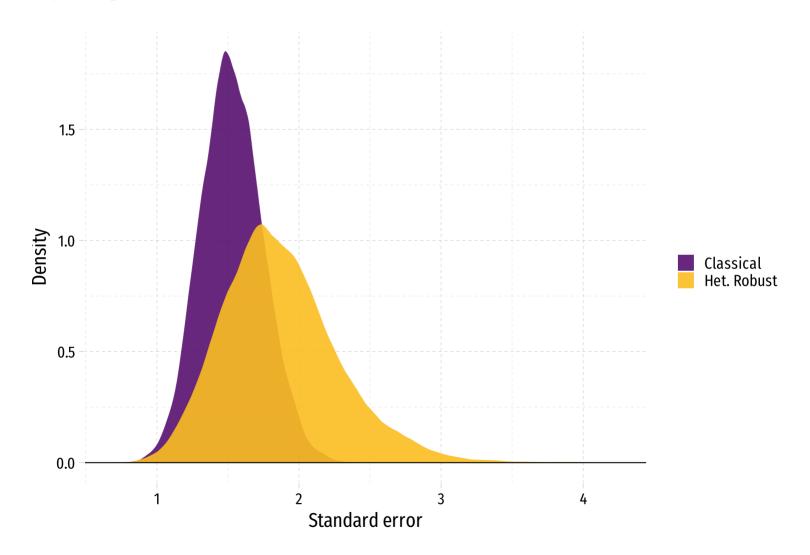
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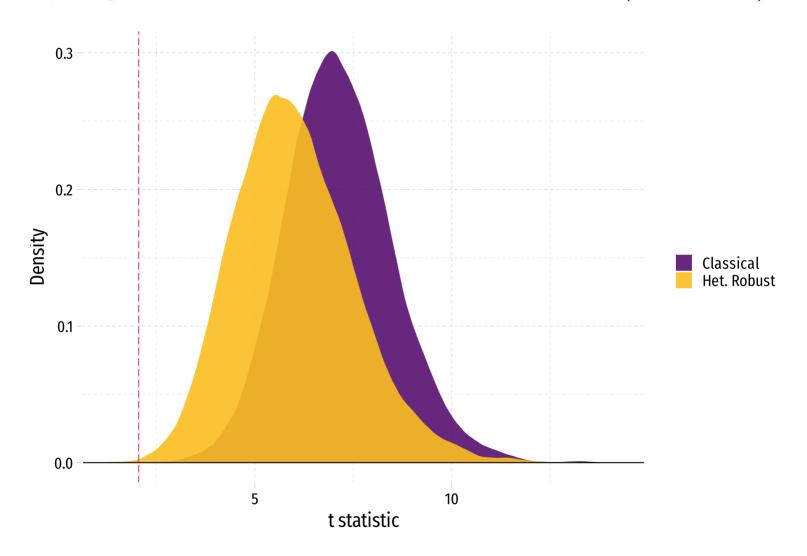
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So what are the results?

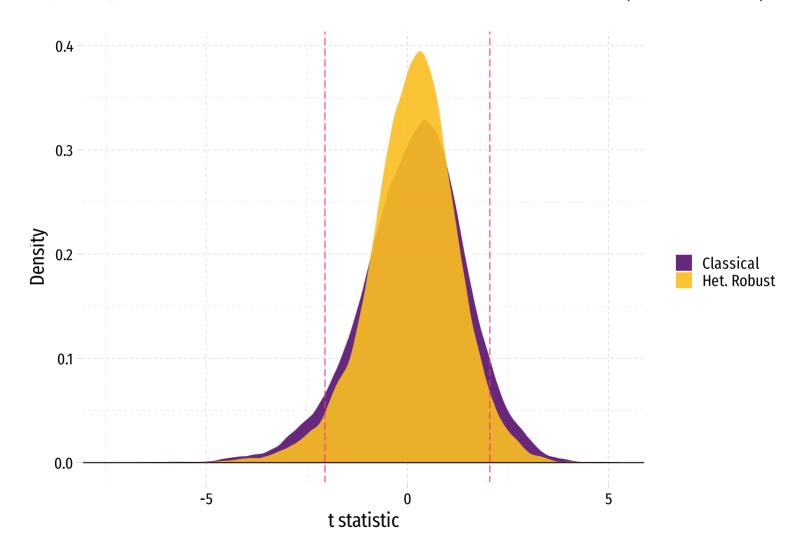
Comparing the distributions of standard errors for the coefficient on \boldsymbol{x}



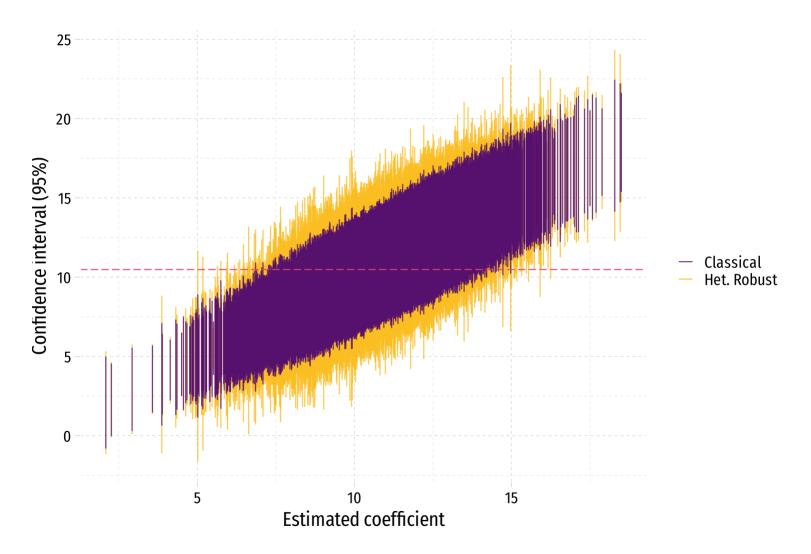
Comparing distributions of t stats for the coefficient on x $(H_o: \beta_1 = 0)$



Comparing distributions of t stats for the coefficient on x $(H_o: \beta_1 = \beta)$



Comparing the confidence intervals for the coefficient on \boldsymbol{x}



How did it go?

For a 5% test the classical SEs

- reject the **true value** in 11.38% of samples
- reject zero in 99.98% of samples

For a 5% test the **het.-robust** SEs

- reject the **true value** in 6.97% of samples
- reject **zero** in 99.93% of samples

All of these test are for a false H_0 .

Q How would the simulation change to enforce a *true* null hypothesis?

Updating to enforce the null

Let's update our simulation function to take arguments γ and δ such that

$$\mathrm{Y}_i = 1 + e^{\gamma \mathrm{X}_i} + arepsilon_i$$

where $arepsilon_i \sim \mathrm{N}(0, \sigma^2 \mathrm{X}_i^\delta)$.

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$$\mathbf{Y}_i = 1 + e^{\gamma \mathbf{X}_i} + arepsilon_i$$

where $arepsilon_i \sim \mathrm{N}(0, \sigma^2 \mathrm{X}_i^\delta)$.

In other words,

- $\gamma = 0$ implies no relationship between Y_i and X_i .
- $\delta = 0$ implies homoskedasticity.

Updating to enforce the null

Updating the function...

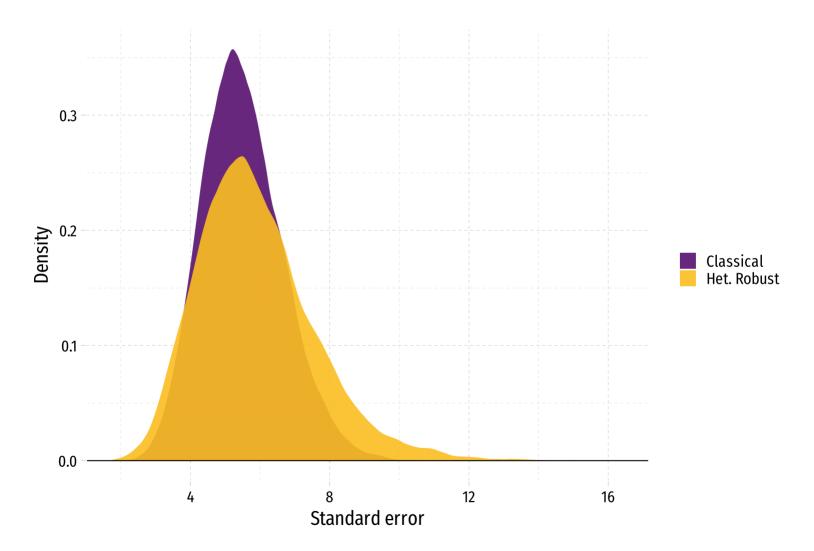
```
flex iter = function(iter, y = 0, \delta = 1, n = 30) {
  # Generate data
  iter df = tibble(
    x = runif(n, min = 0, max = 10),
    \varepsilon = \text{rnorm}(n, \text{sd} = 15 * x^{\delta}),
    v = 1 + exp(v * x) + \varepsilon
  # Estimate models
  lm1 = lm robust(y ~ x, data = iter df, se type = "classical")
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  bind rows(tidy(lm1), tidy(lm2)) %>%
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    mutate(se_type = c("classical", "HC2"), i = iter)
```

Run again!

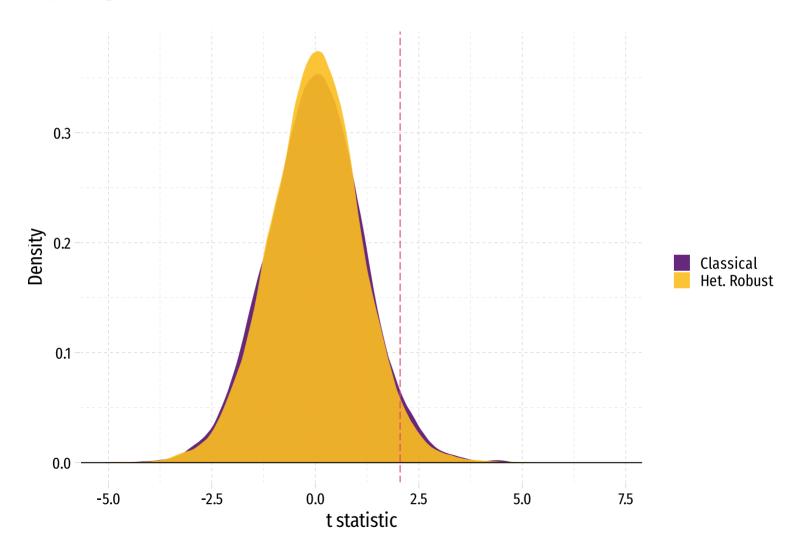
Now we run our new function flex_iter() 10,000 times

```
# Packages
p_load(purrr, furrr)
# Set options
set.seed(123)
# Tell R to parallelize
plan(multisession)
# Run 10,000 iterations
null_df = future_map(
  1:1e4, flex iter,
  # Enforce the null hypothesis
  y = 0,
  # Specify heteroskedasticity
  \delta = 1.
  .options = furrr_options(seed = T)
) %>% bind_rows()
```

Comparing the distributions of standard errors for the coefficient on \boldsymbol{x}



Comparing the distributions of t statistics for the coefficient on x



Distributions of p-values: both methods slightly over-reject the (true) null

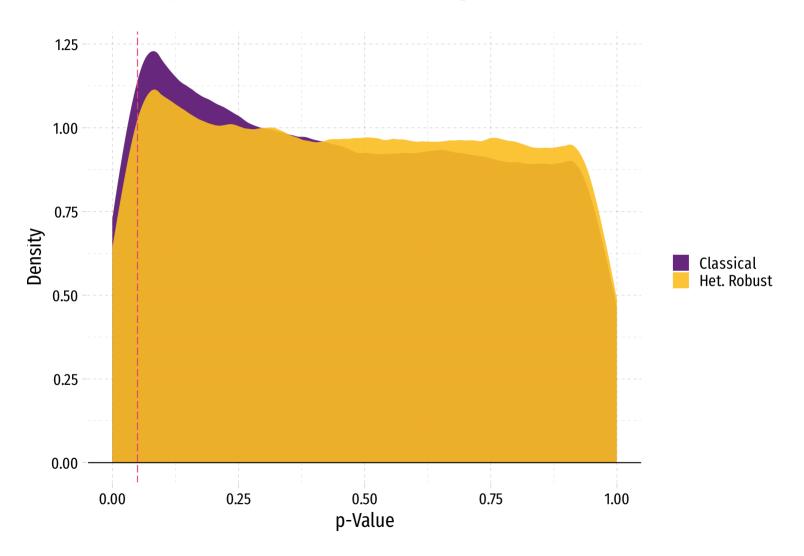


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