Heteroskedasticity

EC 339

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Motivation

The road so far

- Over the past three weeks, we have learned:
 - That omitting relevant variables from a model causes bias;
 - That deterministic/strong stochastic linear relationships between two independent variables harm regression standard errors, and, therefore, OLS inference;
 - That if the error term shows linear relationships across its own observations, OLS standard errors will be affected, also harming inference.

• This week, we will study the last violation of CLRM Assumptions: **Heteroskedasticity**.

Recall **CLRM Assumption V**:

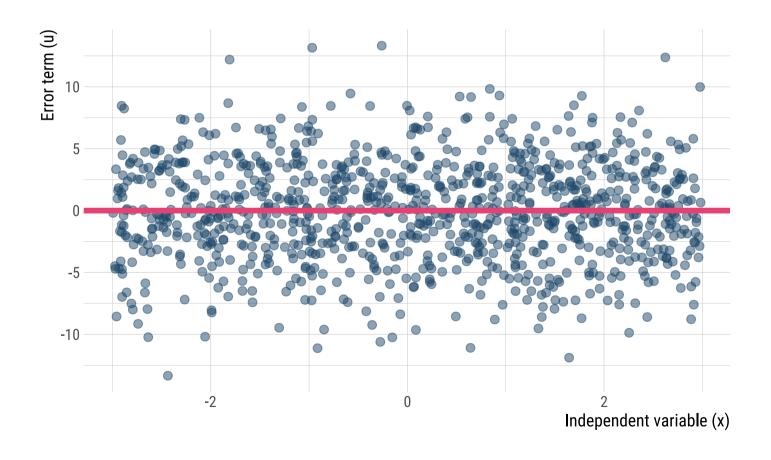
"The error term has a constant variance."

Mathematically...

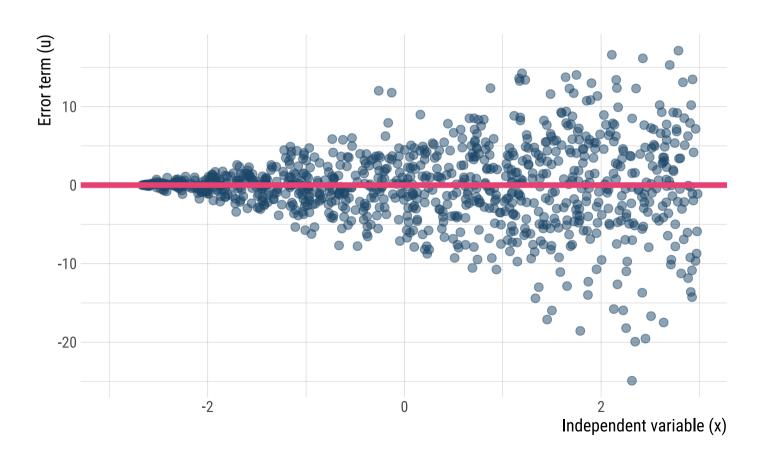
$$Var(u|x) = \sigma^2$$

In words, this assumption implies that the error term has the **same variance** for each value of the independent variable.

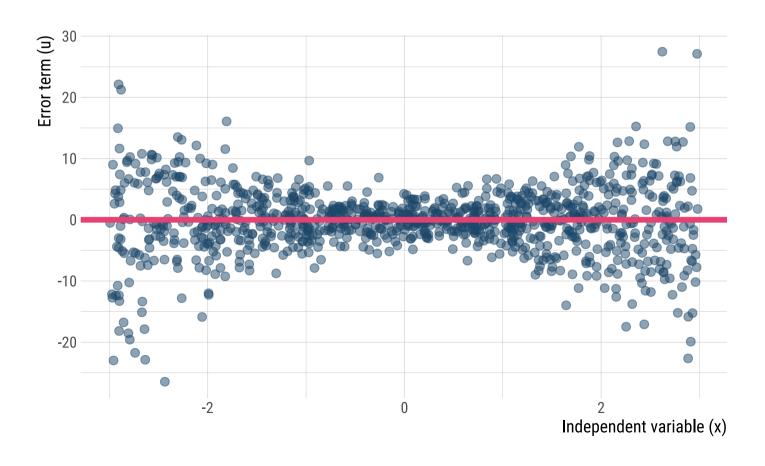
• Homoskedastic residuals:



• Heteroskedastic residuals (1):



• Heteroskedastic residuals (2):



Consequences of heteroskedasticity

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First of all, heteroskedasticity **does not** cause bias to OLS coefficients.

Similar to multicollinearity and serial correlation, heteroskedasticity affects OLS standard errors.

As a consequence, confidence intervals and hypothesis testing procedures become unreliable.

Therefore, how can we trust in our models' **inference**?

We can't!

Testing for heteroskedasticity

Testing for heteroskedasticity

Here, we will study **two** different statistical tests for heteroskedasticity.

- The Breusch-Pagan test;
- The **White** test.

We will study these procedures through an example.

As we have been studying for the past few weeks, all statistical tests involve **auxiliary regression models**.

For the **Breusch-Pagan** test, this is also the case. This time, it involves the regression's **squared residuals**.

The **recipe** 🐺 📡:

```
    Estimate the regression model via OLS, storing its residuals;
    Square the estimated residuals, obtaining û<sub>i</sub><sup>2</sup>;
    Estimate an auxiliary regression, with û<sub>i</sub><sup>2</sup> as the dependent variable, on all independent variables from the original model;
    Then, test the following null hypothesis:
    CLRM Assumption V is true
    H<sub>a</sub>: H<sub>0</sub> is not true
```

The Breusch-Pagan test's **test statistic** is given by

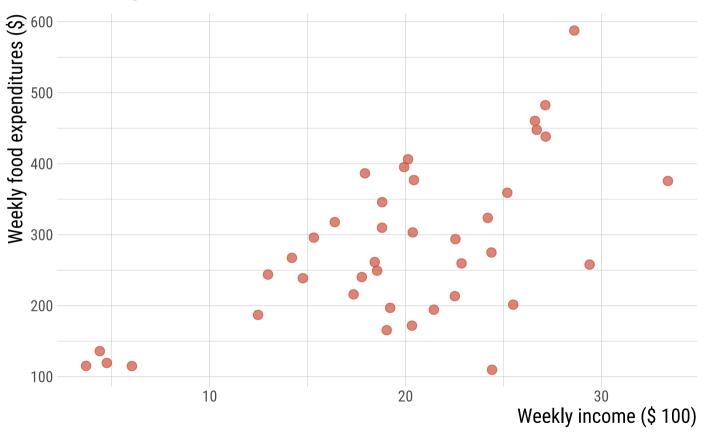
$$LM = n \cdot R_{\hat{u}^2}^2$$

Where n is the sample size, and $R^2_{\hat{u}^2}$ is the coefficient of determination from the auxiliary regression.

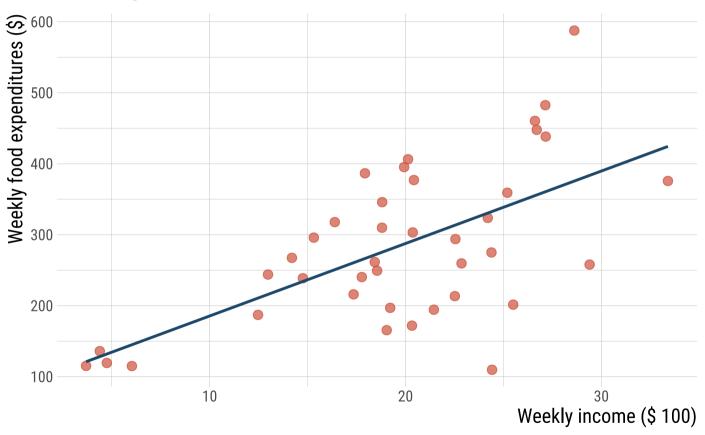
This LM test statistic is $\frac{\text{Chi-squared}}{\text{Chi-squared}}$ distributed, with k degrees-of-freedom.

In case we **reject** the null hypothesis, CLRM Assumption V is **violated** and we have **evidence** of heteroskedasticity in the model's residuals.









#> 1 7.38 0.00658 1 Koenker (studentised) greater

In R...

#>

```
food model ← lm(food exp ~ income, data = food data)
food_model %>% tidy()
#> # A tibble: 2 × 5
        estimate std.error statistic p.value
   term
  #> 1 (Intercept) 83.4 43.4 1.92 0.0622
#> 2 income
           10.2 2.09 4.88 0.0000195
food_model %>% breusch_pagan()
#> # A tibble: 1 × 5
   statistic p.value parameter method
                                          alternative
      <dbl> <dbl> <dbl> <chr>
                                          <chr>
```

What is our **inference**?

In Stata...

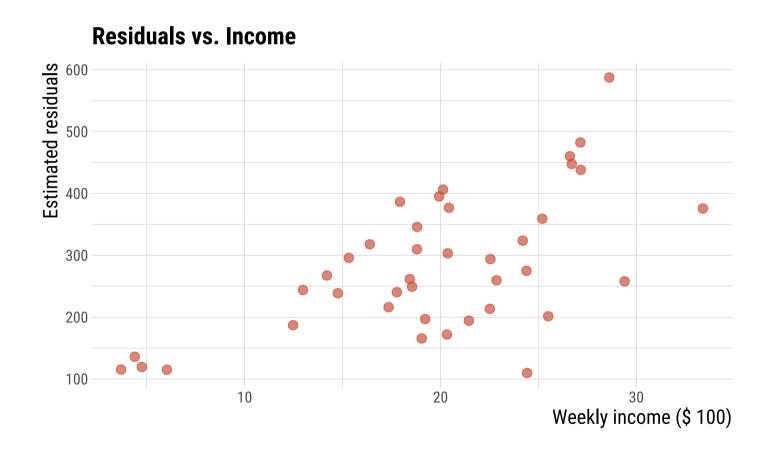
```
. reg food exp income
  Source | SS df MS Number of obs = 40
                           F(1, 38) = 23.79
  Residual | 304505.173 38 8013.29403
                           R-squared = 0.3850
                           Adj R-squared = 0.3688
  food_exp | Coefficient Std. err. t P>|t| [95% conf. interval]
  income | 10.20964 2.093263 4.88 0.000 5.972052 14.44723
   _cons | 83.41601 43.41016 1.92 0.062 -4.463272 171.2953
```

In Stata...

```
. estat hettest, iid
Breusch-Pagan/Cook-Weisberg test for heteroskedasticity
Assumption: i.i.d. error terms
Variable: Fitted values of food_exp
H0: Constant variance
        chi2(1) = 7.38
Prob > chi2 = 0.0066
```

What is our **inference**?

A quick look at this model's **residuals**:



Sometimes, a solution for heteroskedasticity is to log-transform the dependent variable.

- Why?
- It reduces the variable's variance.

Let's see.

```
food_model2 ← lm(log(food_exp) ~ income, data = food_data)
food_model2 %>% breusch_pagan()

#> # A tibble: 1 × 5
```

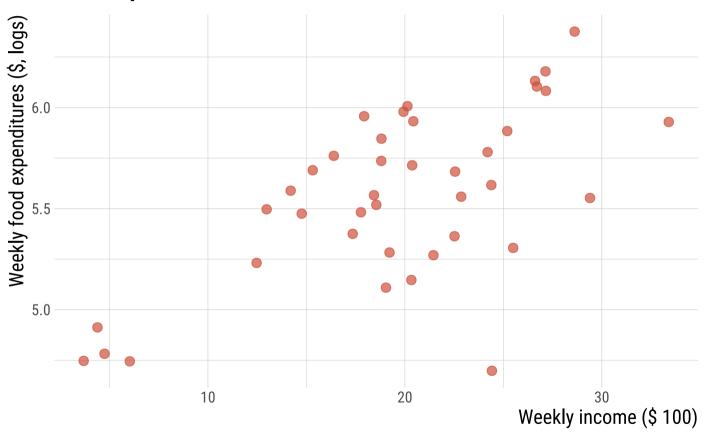
#> statistic p.value parameter method alternative
#> <dbl> <dbl> <dbl> <chr> #> 1 1.71 0.191 1 Koenker (studentised) greater

What happened?

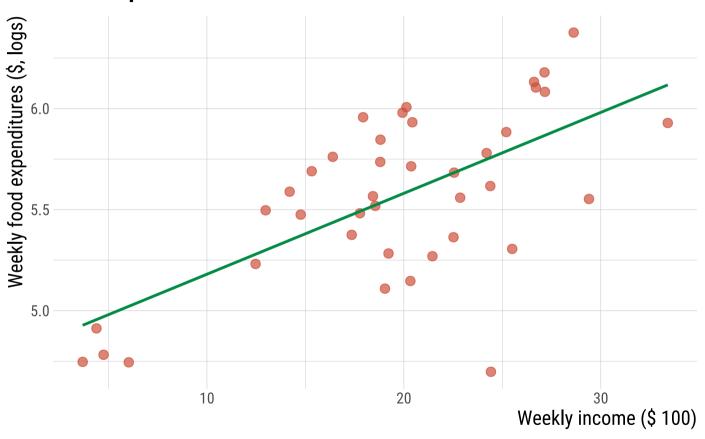
Sometimes, a solution for heteroskedasticity is to **log-transform** the dependent variable.

What happened?









The White test for heteroskedasticity is a more general form of the Breusch-Pagan test.

variables from the original model and desired functional forms;

Basically, it allows \hat{u}^2 to be correlated with further functional forms of the independent variables, such as squares, cubes, interactions, etc.

The **recipe** 🐺 📡:

- 3. Then, test the following null hypothesis:

 H_0 : CLRM Assumption V is true

 H_a : H_0 is not true

Now, let's **apply** this test to our food expenditure models:

• Original model (with food expenditures in levels):

```
food_model %>% white_lm(interactions = TRUE)

#> # A tibble: 1 × 5

#> statistic p.value parameter method alternative

#> <dbl> <dbl> <dbl> <chr> #> 1 7.56 0.0229 2 White's Test greater
```

What is our **inference**?

Now, let's **apply** this test to our food expenditure models:

• Original model (with food expenditures in levels):

```
. estat imtest, white

White's test
H0: Homoskedasticity
Ha: Unrestricted heteroskedasticity

    chi2(2) = 7.56

Prob > chi2 = 0.0229
```

What is our **inference**?

• Now, with food expenditures in logs:

```
food_model2 %>% white_lm(interactions = TRUE)

#> # A tibble: 1 × 5

#> statistic p.value parameter method alternative

#> <dbl> <dbl> <dbl> <chr> #> 1 1.76 0.416 2 White's Test greater
```

And **now**?

• Now, with food expenditures in logs:

```
. estat imtest, white

White's test
H0: Homoskedasticity
Ha: Unrestricted heteroskedasticity

    chi2(2) = 1.76
Prob > chi2 = 0.4156
```

And **now**?

Many times, however, log-transforming variables **does not** guarantee that heteroskedasticity will go away.

A nice solution is to use **heteroskedasticity-robust standard errors**.

By estimating these robust standard errors, we correct the **bias** in a model's standard errors, therefore improving **inference** from our models.

Consider the following model:

-0.954

#> 4 rooms 0.255

#> 5 stratio -0.0525

#> 3 log(dist) -0.134 0.0431 -3.12 1.93e- 3

0.00590

#> 2 lnox

0.117 -8.17 2.57e- 15

0.0185 13.7 1.15e- 36

-8.89 1.07e- 17

Consider the following model:

```
. reg lprice lnox log dist rooms stratio
   Source | SS df MS Number of obs = 506
                                  F(4, 501) = 175.86
   Residual | 35.1834663 501 .07022648
                                  R-squared = 0.5840
                                  Adj R-squared = 0.5807
   Total | 84.582225 505 .167489554 Root MSE = .265
   lprice | Coefficient Std. err. t P>|t| [95% conf. interval]
    lnox | -.9535388 .1167417 -8.17 0.000 -1.182902 -.7241751
  log_dist | -.1343395 .0431032 -3.12
                                0.002 -.2190247 -.0496542
    rooms .2545271 .0185303 13.74
                                0.000
                                     .2181203 .2909338
   stratio | -.0524511 .0058971 -8.89 0.000 -.0640372 -.040865
    cons | 11.08386 .3181113 34.84
                                0.000 10.45887 11.70886
```

Breusch-Pagan test:

```
price_model %>% breusch_pagan()

#> # A tibble: 1 * 5

#> statistic p.value parameter method alternative

#> <dbl> <dbl> <dbl> <chr> #> 1 69.9 2.42e-14 4 Koenker (studentised) greater
```

White test:

```
price_model %>% white_lm(interactions = TRUE)

#> # A tibble: 1 × 5

#> statistic p.value parameter method alternative
```

Breusch-Pagan test:

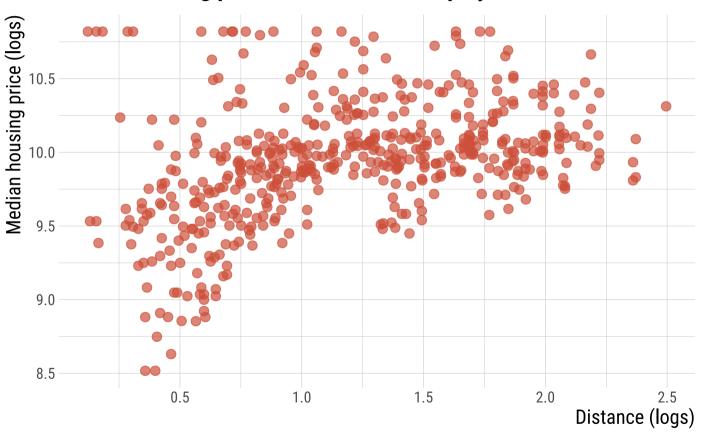
White test:

```
. estat imtest, white

White's test
H0: Homoskedasticity
Ha: Unrestricted heteroskedasticity

    chi2(14) = 143.98
Prob > chi2 = 0.0000
```

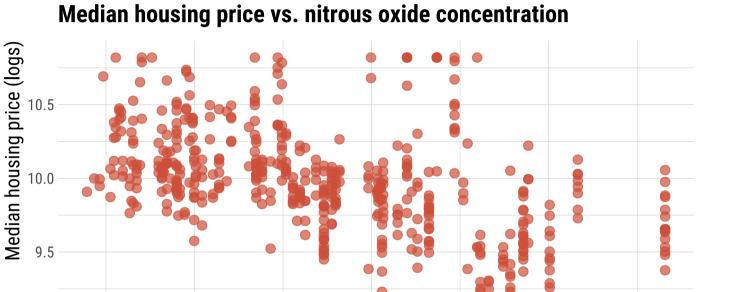




9.0

8.5

1.50

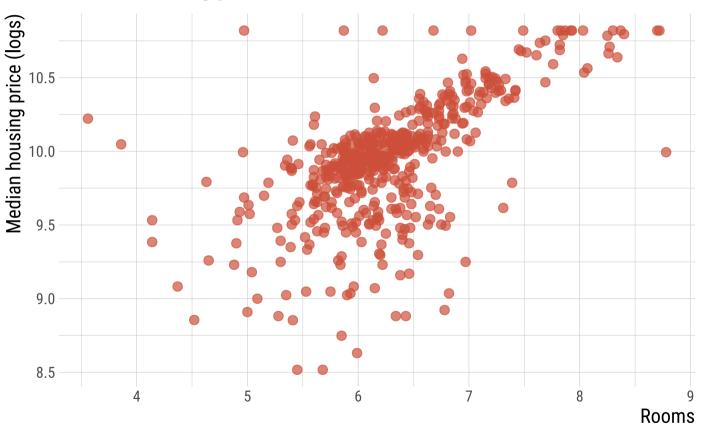


1.75

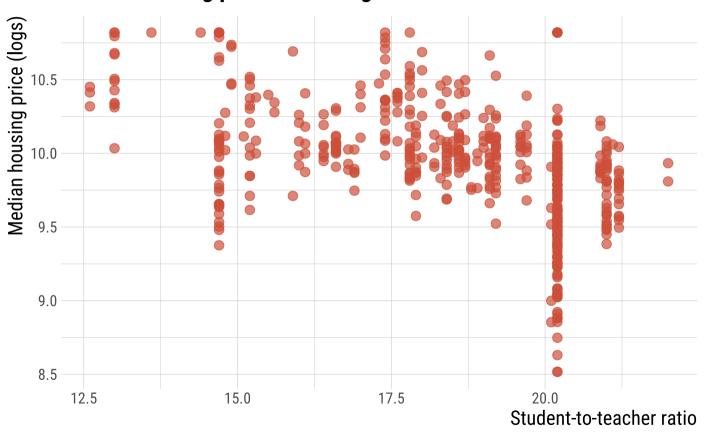
2.00

NO2 concentration (logs)









Robust (White) standard errors:

| Variable | Coefficient | Standard error | t-statistic | p-value |
|-------------|-------------|----------------|-------------|-----------|
| (Intercept) | 11.0838616 | 0.3772949 | 29.3771817 | 0.0000000 |
| lnox | -0.9535388 | 0.1268005 | -7.5199909 | 0.0000000 |
| log(dist) | -0.1343395 | 0.0535287 | -2.5096731 | 0.0123986 |
| rooms | 0.2545271 | 0.0247205 | 10.2962139 | 0.0000000 |
| stratio | -0.0524511 | 0.0046082 | -11.3821438 | 0.0000000 |

Robust (White) standard errors:

```
. reg lprice lnox log dist rooms stratio, robust
Linear regression
                                     Number of obs = 506
                                     F(4, 501) = 146.27
                                     Prob > F = 0.0000
                                     R-squared = 0.5840
                                     Root MSE = .265
                      Robust
    lprice | Coefficient std. err. t P>|t| [95% conf. interval]
     lnox | -.9535388 .1268005 -7.52 0.000 -1.202665 -.7044125
  log_dist | -.1343395 .0535287 -2.51
                                    0.012 -.2395078 -.0291711
     rooms .2545271 .0247205 10.30
                                    0.000
                                          .2059585 .3030956
   stratio | -.0524511 .0046082 -11.38 0.000 -.0615049 -.0433974
     cons | 11.08386
                    .3772949 29.38 0.000
                                          10.34259 11.82514
```

In **summary**, whenever interpreting a model with **heteroskedastic** residuals, use **robust standard errors** for inference purposes.

Otherwise, any inferential analysis from our models will not be valid, since violating **CLRM Assumption V** directly affects OLS standard errors.

Next time: Heteroskedasticity in practice