

Practical 324

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Regression analysis

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Simple regression

The simple regression analysis is a supervised machine learning approach to creating a model able to predict the value of one outcome variable Y based on one predictor variable X_1 , by estimating the intercept b_0 and coefficient (slope) b_1 , and accounting for a reasonable amount of error ϵ .

$$Y_i = (b_0 + b_1 * X_{i1}) + \epsilon_i$$

Least squares is the most commonly used approach to generate a regression model. This model fits a line to minimise the squared values of the **residuals** (errors), which are calculated as the squared difference between observed values the values predicted by the model.

$$redidual = \sum (observed - model)^2$$

A model is considered **robust** if the residuals do not show particular trends, which would indicate that “something” is interfering with the model. In particular, the assumption of the regression model are:

- **linearity**: the relationship is actually linear;
- **normality** of residuals: standard residuals are normally distributed with mean 0;
- **homoscedasticity** of residuals: at each level of the predictor variable(s) the variance of the standard residuals should be the same (*homo-scedasticity*) rather than different (*hetero-scedasticity*);
- **independence** of residuals: adjacent standard residuals are not correlated.

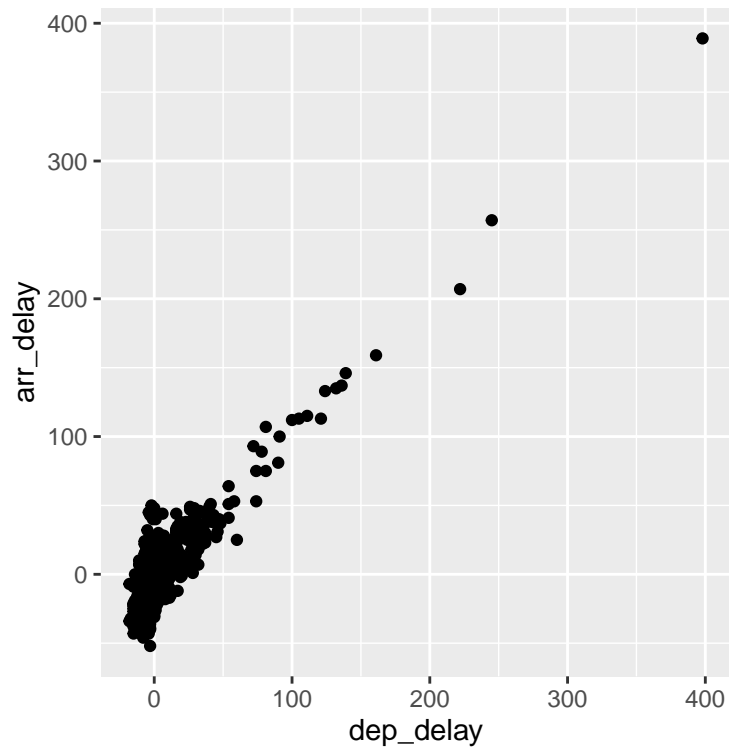
Example

The example that we have seen in the lecture illustrated how simple regression can be used to create a model to predict the arrival delay based on the departure delay of a flight, based on the data available in the `nycflights13` dataset for the flight on November 20th, 2013. The scatterplot below seems to indicate that the relationship is indeed linear.

$$arr_delay_i = (Intercept + Coefficient_{dep_delay} * dep_delay_{i1}) + \epsilon_i$$

```
# Load the library
library(nycflights13)
```

```
# November 20th, 2013
flights_nov_20 <- nycflights13::flights %>%
  dplyr::filter(!is.na(dep_delay), !is.na(arr_delay), month == 11, day == 20)
```



The code below generates the model using the function `lm`, and the function `summary` to obtain the summary of the results of the test. The model and summary are saved in the variables `delay_model` and `delay_model_summary`, respectively, for further use below. The variable `delay_model_summary` can then be called directly to visualise the result of the test.

```
# Classic R coding version
# delay_model <- lm(arr_delay ~ dep_delay, data = flights_nov_20)
# delay_model_summary <- summary(delay_model)
```

```
# Load magrittr library to use %$%
library(magrittr)
```

```
delay_model <- flights_nov_20 %$%
  lm(arr_delay ~ dep_delay)
```

```
delay_model_summary <- delay_model %>%
  summary()
```

```
delay_model_summary
```

```
##
## Call:
## lm(formula = arr_delay ~ dep_delay)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -43.906  -9.022  -1.758   8.678  57.052
```

```
##
## Coefficients:
##           Estimate Std. Error t value Pr(>|t|)
## (Intercept) -4.96717    0.43748  -11.35  <2e-16 ***
## dep_delay   1.04229    0.01788   58.28  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 13.62 on 972 degrees of freedom
## Multiple R-squared:  0.7775, Adjusted R-squared:  0.7773
## F-statistic: 3397 on 1 and 972 DF, p-value: < 2.2e-16
```

The image below highlights the important values in the output: the adjusted R^2 value; the model significance value **p-value** and the related F-statistic information **F-statistic**; the intercept and **dep_delay** coefficient estimates in the **Estimate** column and the related significance values of in the column **Pr(>|t|)**.

```
Call:
lm(formula = arr_delay ~ dep_delay)

Residuals:
    Min       1Q   Median       3Q      Max
-43.906  -9.022  -1.758   8.678  57.052

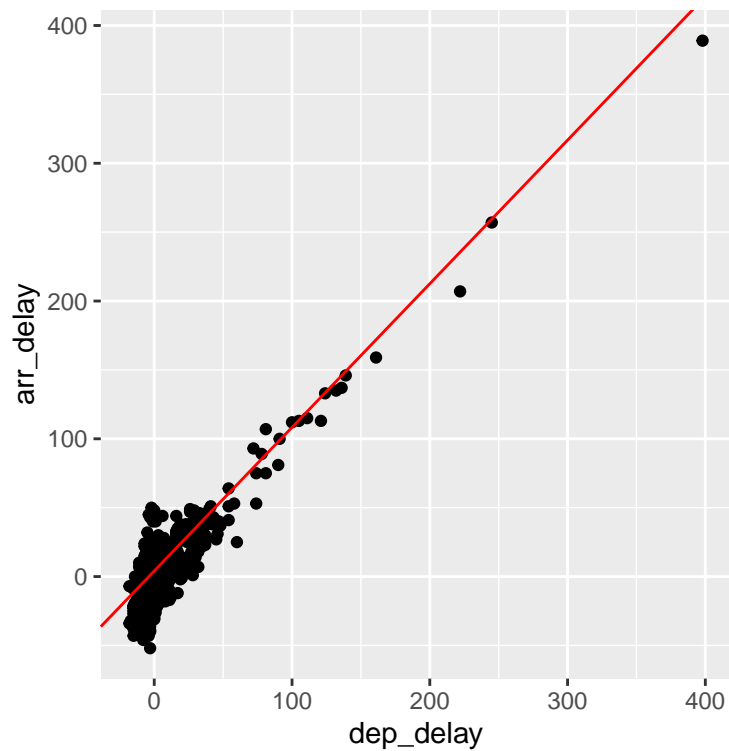
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) -4.96717    0.43748  -11.35  <2e-16 ***
dep_delay    1.04229    0.01788   58.28  <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 13.62 on 972 degrees of freedom
Multiple R-squared:  0.7775, Adjusted R-squared:  0.7773
F-statistic: 3397 on 1 and 972 DF, p-value: < 2.2e-16
```

The output indicates:

- **p-value: < 2.2e-16:** $p < .001$ the model is significant;
 - derived by comparing the calculated **F-statistic** value to F distribution 3396.74 having specified degrees of freedom (1, 972);
 - Report as: $F(1, 972) = 3396.74$
- **Adjusted R-squared: 0.7773:** the departure delay can account for 77.73% of the arrival delay;
- **Coefficients:**
 - Intercept estimate -4.9672 is significant;
 - **dep_delay** coefficient (slope) estimate 1.0423 is significant.

```
flights_nov_20 %>%
  ggplot2::ggplot(aes(x = dep_delay, y = arr_delay)) +
  ggplot2::geom_point() + ggplot2::coord_fixed(ratio = 1) +
  ggplot2::geom_abline(intercept = 4.0943, slope = 1.04229, color="red")
```

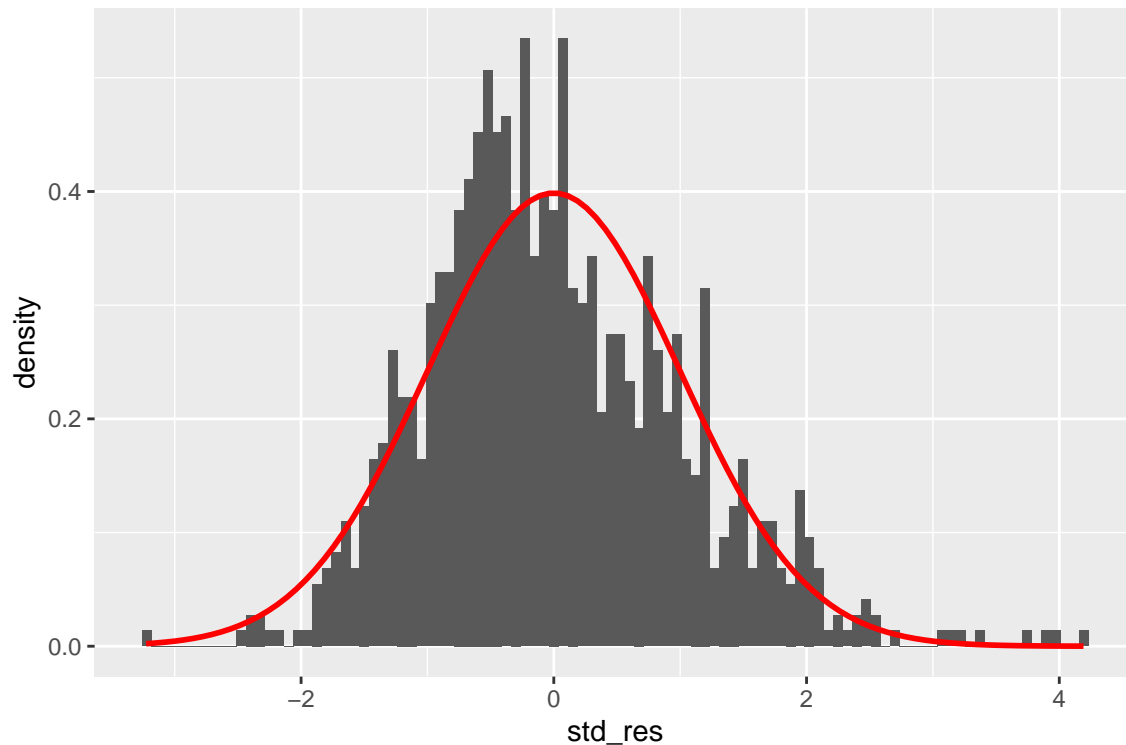


Checking assumptions

Normality The Shapiro-Wilk test can be used to check for the normality of standard residuals. The test should be not significant for robust models. In the example below, the standard residuals are *not* normally distributed. However, the plot further below does show that the distribution of the residuals is not far away from a normal distribution.

```
delay_model %>%
  stats::rstandard() %>%
  stats::shapiro.test()
```

```
##
##  Shapiro-Wilk normality test
##
## data:  .
## W = 0.98231, p-value = 1.73e-09
```



Homoscedasticity The Breusch-Pagan test can be used to check for the homoscedasticity of standard residuals. The test should be not significant for robust models. In the example below, the standard residuals are homoscedastic.

```
library(lmtest)

delay_model %>%
  lmtest::bptest()
```

```
##
## studentized Breusch-Pagan test
##
## data: .
## BP = 0.017316, df = 1, p-value = 0.8953
```

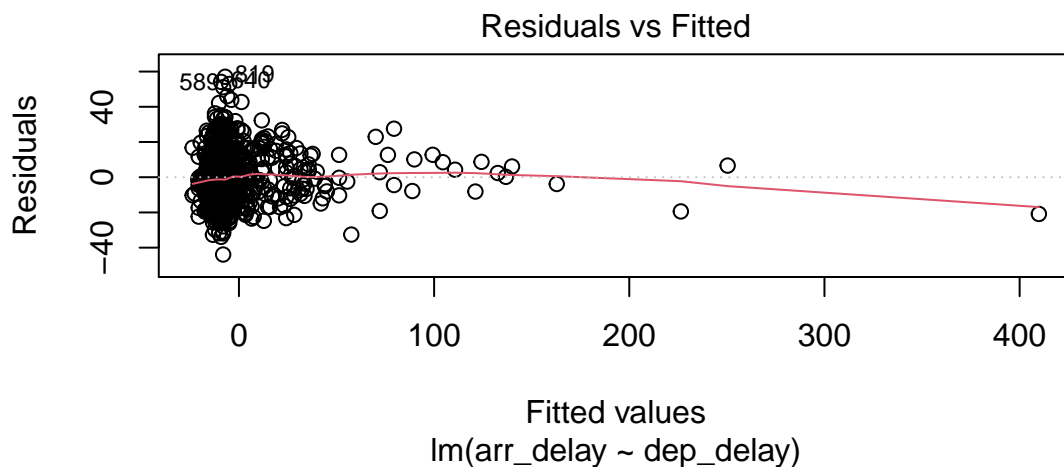
Independence The Durbin-Watson test can be used to check for the independence of residuals. The test should be statistic should be close to 2 (between 1 and 3) and not significant for robust models. In the example below, the standard residuals might not be completely independent. Note, however, that the result depends on the order of the data.

```
# Also part of the library lmtest
delay_model %>%
  lmtest::dwtest()
```

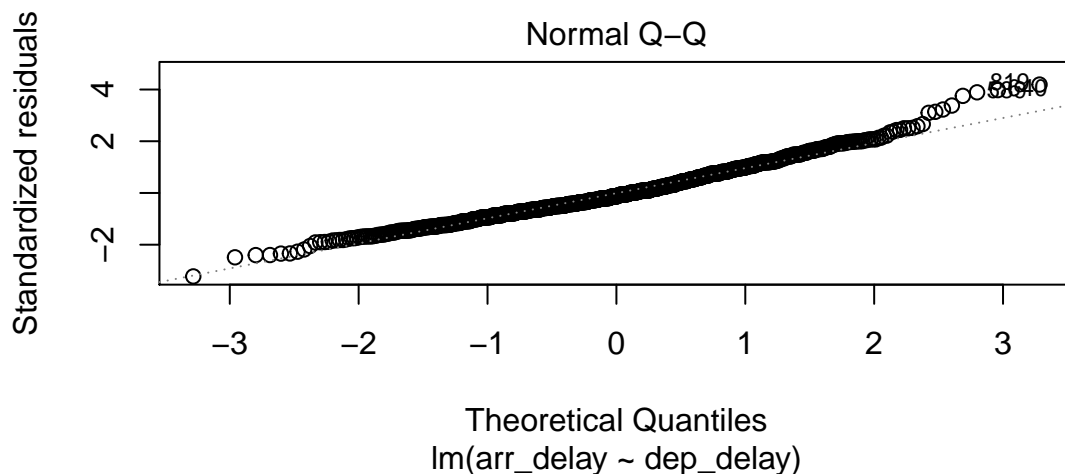
```
##
## Durbin-Watson test
##
## data: .
## DW = 1.8731, p-value = 0.02358
## alternative hypothesis: true autocorrelation is greater than 0
```

Plots The `plot.lm` function can be used to further explore the residuals visually. Usage is illustrated below. The *Residuals vs Fitted* and *Scale-Location* plot provide an insight into the homoscedasticity of the residuals, the *Normal Q-Q* plot provides an illustration of the normality of the residuals, and the *Residuals vs Leverage* can be useful to identify exceptional cases (e.g., Cook's distance greater than 1).

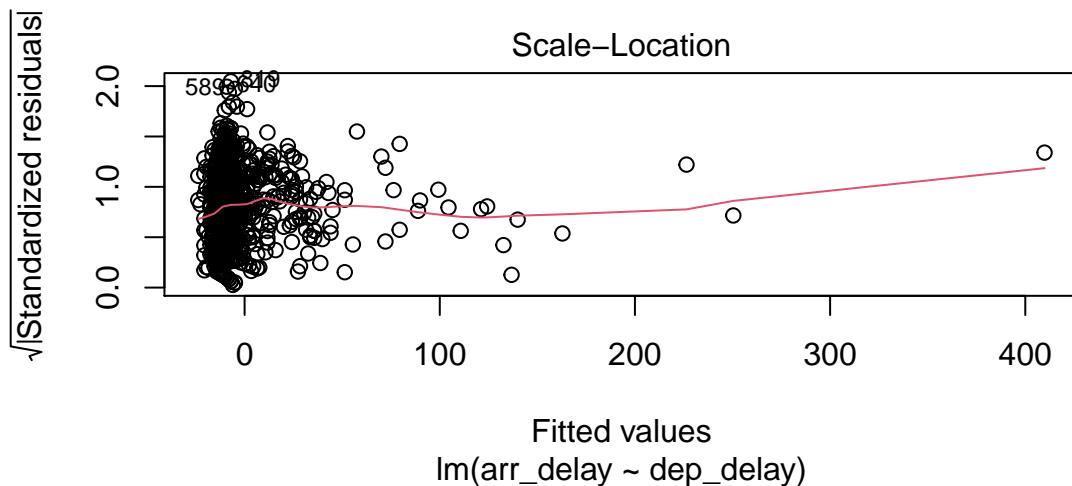
```
delay_model %>%
  plot(which = c(1))
```



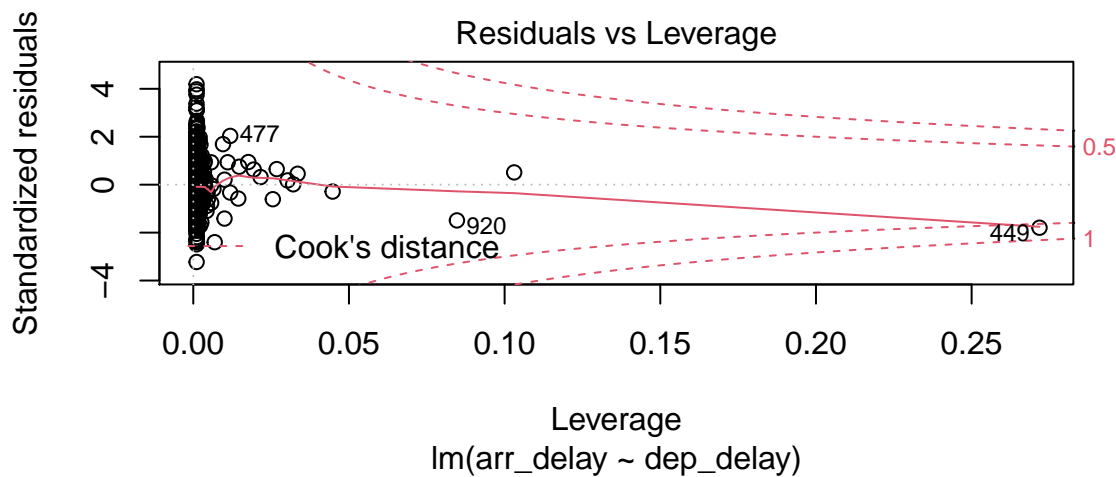
```
delay_model %>%
  plot(which = c(2))
```



```
delay_model %>%
  plot(which = c(3))
```



```
delay_model %>%
  plot(which = c(5))
```



How to report

Overall, we can say that the delay model computed above is fit ($F(1, 972) = 3396.74$, $p < .001$), indicating that the departure delay might account for 77.73% of the arrival delay. However the model is only partially robust. The residuals satisfy the homoscedasticity assumption (Breusch-Pagan test, $BP = 0.02$, $p = 0.9$), and the independence assumption (Durbin-Watson test, $DW = 1.87$, $p = 0.02$), but they are not normally distributed (Shapiro-Wilk test, $W = 0.98$, $p < .001$).

The `stargazer` function of the `stargazer` library can be applied to the model `delay_model` to generate a nicer output in RMarkdown PDF documents by including `results = "asis"` in the R snippet option.

```
# Install stargazer if not yet installed
# install.packages("stargazer")

library(stargazer)

# Not rendered in bookdown
stargazer(delay_model, header = FALSE)
```

Table 1:

<i>Dependent variable:</i>	
arr_delay	
dep_delay	1.042*** (0.018)
Constant	-4.967*** (0.437)
Observations	974
R ²	0.778
Adjusted R ²	0.777
Residual Std. Error	13.618 (df = 972)
F Statistic	3,396.742*** (df = 1; 972)
<i>Note:</i>	*p<0.1; **p<0.05; ***p<0.01

Multiple regression

The multiple regression analysis is a supervised machine learning approach to creating a model able to predict the value of one outcome variable Y based on two or more predictor variables $X_1 \dots X_M$, by estimating the intercept b_0 and the coefficients (slopes) $b_1 \dots b_M$, and accounting for a reasonable amount of error ϵ .

$$Y_i = (b_0 + b_1 * X_{i1} + b_2 * X_{i2} + \dots + b_M * X_{iM}) + \epsilon_i$$

The assumptions are the same as the simple regression, plus the assumption of **no multicollinearity**: if two or more predictor variables are used in the model, each pair of variables not correlated. This assumption can be tested by checking the variance inflation factor (VIF). If the largest VIF value is greater than 10 or the average VIF is substantially greater than 1, there might be an issue of multicollinearity.

Example

The example below explores whether a regression model can be created to estimate the number of people in Leicester commuting to work using private transport (**u121**) in Leicester, using the number of people in different industry sectors as predictors.

For instance, occupations such as electricity, gas, steam and air conditioning supply (**u144**) require to travel some distances with equipment, thus the related variable **u144** is included in the model, whereas people working in information and communication might be more likely to work from home or commute by public transport.

A multiple regression model can be specified in a similar way as a simple regression model, using the same `lm` function, but adding the additional predictor variables using a `+` operator.

```
leicester_2011OAC <- readr::read_csv("2011_OAC_Raw_uVariables_Leicester.csv")
```

```
# Select and
# normalise variables
leicester_2011OAC_transp <-
  leicester_2011OAC %>%
  dplyr::select(
    OA11CD,
    Total_Pop_No_NI_Students_16_to_74, Total_Employment_16_to_74,
```



```

    u121, u141:u158
  ) %>%
  # percentage method of travel
  dplyr::mutate(
    u121 = (u121 / Total_Pop_No_NI_Students_16_to_74) * 100
  ) %>%
  # percentage across industry sector columns
  dplyr::mutate(
    dplyr::across(
      u141:u158,
      function(x){ (x / Total_Employment_16_to_74) * 100 }
    )
  ) %>%
  # rename columns
  dplyr::rename_with(
    function(x){ paste0("perc_", x) },
    c(u121, u141:u158)
  )

# Selected variables

# perc_u120: Method of Travel to Work, Private Transport
# perc_u142: Industry Sector, Mining and quarrying
# perc_u144: Industry Sector, Electricity, gas, steam and air conditioning ...
# perc_u146: Industry Sector, Construction
# perc_u149: Industry Sector, Accommodation and food service activities

# Create model
commuting_model1 <-
  leicester_20110AC_transp %$%
  lm(
    perc_u121 ~
      perc_u142 + perc_u144 + perc_u146 + perc_u149
  )

# Print summary
commuting_model1 %>%
  summary()

```

```

##
## Call:
## lm(formula = perc_u121 ~ perc_u142 + perc_u144 + perc_u146 +
##     perc_u149)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -35.315  -6.598  -0.244   6.439  31.472
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  37.12690    0.94148   39.434 < 2e-16 ***
## perc_u142     3.74768    1.21255    3.091 0.00205 **
## perc_u144     1.16865    0.25328    4.614 4.48e-06 ***
## perc_u146     1.05408    0.09335   11.291 < 2e-16 ***

```

```
## perc_u149    -1.56948    0.08435 -18.606 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 9.481 on 964 degrees of freedom
## Multiple R-squared:  0.3846, Adjusted R-squared:  0.3821
## F-statistic: 150.6 on 4 and 964 DF,  p-value: < 2.2e-16
# Not rendered in bookdown
stargazer(commuting_model1, header=FALSE)
```

Table 2:

	<i>Dependent variable:</i>
	perc_u121
perc_u142	3.748*** (1.213)
perc_u144	1.169*** (0.253)
perc_u146	1.054*** (0.093)
perc_u149	-1.569*** (0.084)
Constant	37.127*** (0.941)
Observations	969
R ²	0.385
Adjusted R ²	0.382
Residual Std. Error	9.481 (df = 964)
F Statistic	150.622*** (df = 4; 964)
Note:	*p<0.1; **p<0.05; ***p<0.01

```
commuting_model1 %>%
  rstandard() %>%
  shapiro.test()
```

```
##
##  Shapiro-Wilk normality test
##
## data:  .
## W = 0.99889, p-value = 0.8307
```

```
commuting_model1 %>%
  bptest()
```

```
##
##  studentized Breusch-Pagan test
##
```

```
## data: .
## BP = 28.403, df = 4, p-value = 1.033e-05

commuting_model1 %>%
  dwtest()

##
## Durbin-Watson test
##
## data: .
## DW = 1.835, p-value = 0.004908
## alternative hypothesis: true autocorrelation is greater than 0

library(car)

commuting_model1 %>%
  vif()

## perc_u142 perc_u144 perc_u146 perc_u149
## 1.006906 1.016578 1.037422 1.035663
```

The output above suggests that the model is fit ($F(4, 964) = 150.62$, $p < .001$), indicating that a model based on the presence of people working in the four selected industry sectors can account for 38.21% of the number of people using private transportation to commute to work. However the model is only partially robust. The residuals are normally distributed (Shapiro-Wilk test, $W = 1$, $p = 0.83$) and there seems to be no multicollinearity with average VIF 1.02, but the residuals don't satisfy the homoscedasticity assumption (Breusch-Pagan test, $BP = 28.4$, $p < .001$), nor the independence assumption (Durbin-Watson test, $DW = 1.84$, $p < .01$).

The coefficient values calculated by the `lm` functions are important to create the model, and provide useful information. For instance, the coefficient for the variable `perc_u144` is 1.169, which indicates that if the presence of people working in electricity, gas, steam and air conditioning supply increases by one percentage point, the number of people using private transportation to commute to work increases by 1.169 percentage points, according to the model. The coefficients also indicate that the presence of people working in accommodation and food service activities actually has a negative impact (in the context of the variables selected for the model) on the number of people using private transportation to commute to work.

In this example, all variables use the same unit and are of a similar type, which makes interpreting the model relatively simple. When that is not the case, it can be useful to look at the standardized β , which provide the same information but measured in terms of standard deviation, which make comparisons between variables of different types easier to draw. For instance, the values calculated below using the function `lm.beta` of the library `lm.beta` indicate that if the presence of people working in construction has the highest impact on the outcome variable. If the presence of people working in construction increases by one standard deviation, the number of people using private transportation to commute to work increases by 0.29 standard deviations, according to the model.

```
# Install lm.beta library if necessary
# install.packages("lm.beta")
library(lm.beta)

lm.beta(commuting_model1)

##
## Call:
## lm(formula = perc_u121 ~ perc_u142 + perc_u144 + perc_u146 +
##     perc_u149)
##
## Standardized Coefficients::
```

```
## (Intercept)    perc_u142    perc_u144    perc_u146    perc_u149
##  0.00000000    0.07836017    0.11754058    0.29057993   -0.47841083
```

Exercise 324.1

Question 324.1.1: Create a model having as outcome variable the presence of people using private transport for commuting to work, and using a stepwise “*both*” approach, having all the variables created for the example above and related to the presence of people working in different industry sectors (`perc_u141` to `perc_u158`) as scope.

Question 324.1.2: Is the presence of people using public transportation to commute to work statistically, linearly related to mean age (`u020`)?

Question 324.1.3: Is the presence of people using public transportation to commute to work statistically, linearly related to (a subset of) the age structure categories (`u007` to `u019`)?