Seminar notebook

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Loading the required packages

```
library(tidyverse)
## -- Attaching packages -
                                                            --- tidyverse 1.3.1 --
## v ggplot2 3.3.5
                      v purrr
                                0.3.4
## v tibble 3.1.5
                      v dplyr
                                1.0.7
## v tidyr
            1.1.4
                      v stringr 1.4.0
## v readr
            2.0.2
                      v forcats 0.5.1
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                    masks stats::lag()
library(gapminder)
library(broom) #broom package takes the messy output of built-in functions in R, such as lm, #nls, or t.
theme_set(theme_minimal(base_size = 10)) #theme_set overides default qqplot theme
library(gapminder)
library(plotly)
##
## Attaching package: 'plotly'
## The following object is masked from 'package:ggplot2':
##
##
      last_plot
  The following object is masked from 'package:stats':
##
##
##
      filter
## The following object is masked from 'package:graphics':
##
##
```

Linear regression

layout

Simple linear regression

Estimation

Task 1. Filter year=2007 from the gapminder data and call the data frame object gm2007. 2. Create a regression object called model_lm that regresses lifeEXp on gdpPercap for 2007 data 3. Output the estimated coefficients, intercept and slope, only

```
gm2007 <- gapminder %>% filter(year == 2007)
model_lm <- lm(data = gm2007, lifeExp ~ gdpPercap)</pre>
coef(model_lm)
## (Intercept)
                    gdpPercap
## 5.956565e+01 6.371341e-04
```

Demo dplyr::summarise function

Summarise function displays the coefficient estimates in the regression model

```
• hat\_beta1 = cor(x,y) * sd(y) / sd(x)
• hat beta0 = mean(y) - hat beta1 * mean(x)
```

• Regression line passes through (mean(x), mean(y))

```
#output: correlation, sd, mean(x), mean(y), estimated intercept and slope coefficient using formula
gm2007 %>%
  summarize(cor_xy=cor(gdpPercap, lifeExp),
            sd_x = sd(gdpPercap),
           sd_y = sd(lifeExp),
           mean_x = mean(gdpPercap),
           mean_y = mean(lifeExp),
            hat_beta1 = cor_xy/sd_x*sd_y,
           hat_beta0 = mean_y - hat_beta1*mean_x)
## # A tibble: 1 x 7
```

```
cor_xy sd_x sd_y mean_x mean_y hat_beta1 hat_beta0
     <dbl> <dbl> <dbl> <dbl> <dbl> <
                                         <dbl>
                                                  <dbl>
                                                   59.6
## 1 0.679 12860. 12.1 11680. 67.0 0.000637
```

Inference

```
#Use summary() to obtain basic output of regression
summary(model lm)
##
## Call:
## lm(formula = lifeExp ~ gdpPercap, data = gm2007)
## Residuals:
      Min
               1Q Median
                               3Q
##
                                      Max
## -22.828 -6.316
                            6.898 13.128
                   1.922
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 5.957e+01 1.010e+00 58.95
                                            <2e-16 ***
## gdpPercap 6.371e-04 5.827e-05
                                   10.93
                                           <2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 8.899 on 140 degrees of freedom
## Multiple R-squared: 0.4606, Adjusted R-squared: 0.4567
## F-statistic: 119.5 on 1 and 140 DF, p-value: < 2.2e-16
(ISLR eq 3.8) p.66
```

```
se(beta hat) = sigma / sqrt(sum((x - mean(x))^2))
Estimated by:
se(beta hat) = sigma hat / sqrt(sum((x - mean(x))^2))
where (ISLR 3.15) p.69:
sigma hat = RSE = sqrt( RSS / (n-2) )
#Using augment() to extract residuals, compute the estimated se of the slope parameter by
#obtaining 1. RSS 2. MSE = (RSS/(n-p)) 3. estimated S.E of hat beta1
augment(model_lm) %>%
  summarize(RSS = sum(.resid^2),
            MSE = RSS /(n()-2),
            SE = sqrt(MSE)/sqrt(sum((gdpPercap - mean(gdpPercap))^2)))
## # A tibble: 1 x 3
##
        RSS
              MSE
                          SE
##
      <dbl> <dbl>
                       <dbl>
## 1 11086. 79.2 0.0000583
Model diagnostics
#qlance: Construct a single row summary "qlance" of a model, fit, or other object
#CODE
glance(model_lm)
## # A tibble: 1 x 12
     r.squared adj.r.squared sigma statistic p.value
                                                                               BIC
                                                            df logLik
                                                                         AIC
##
         <dbl>
                        <dbl> <dbl>
                                         <dbl>
                                                   <dbl> <dbl> <dbl> <dbl> <dbl> <
## 1
         0.461
                        0.457 8.90
                                          120. 1.69e-20
                                                             1 -511. 1028. 1037.
## # ... with 3 more variables: deviance <dbl>, df.residual <int>, nobs <int>
"Portion of variance in outcome explained by simple linear regression model"
                                         R^2 = \operatorname{cor}(x, y)^2
#compute correlation squared between qdpPercap and lifeExp
#CODE
cor(gm2007$gdpPercap, gm2007$lifeExp)^2
## [1] 0.4605827
                                         R^2 = 1 - \frac{\text{RSS}}{\text{TSS}}
#using residuals from augment(), obtain values of RSS, TSS, and R2
#CODE
augment(model_lm) %>%
  summarize(RSS = sum(.resid^2),
            TSS = sum((lifeExp - mean(lifeExp))^2),
            R2 = 1 - RSS/TSS)
```

A tibble: 1 x 3

TSS

<dbl> <dbl> <dbl> ## 1 11086. 20552. 0.461

R.2

RSS

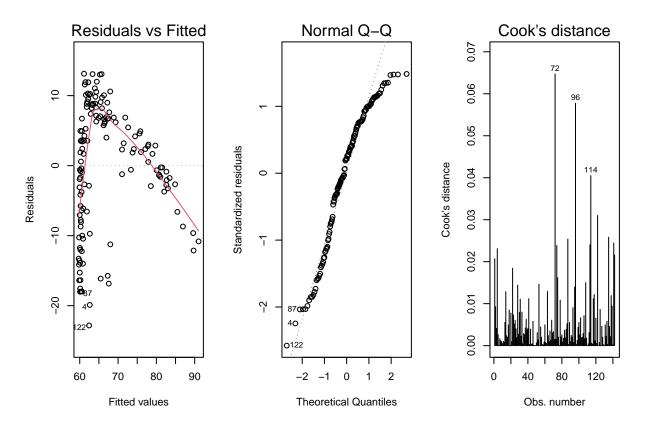
##

Diagnostic plots

Idea: look for patterns in residuals, which could indicate systematic errors (bias)

Outliers vs influential points: influential points affect the slope of the model line

```
#Plot residuals vs gdpPercap, using residuals from augment()
#CODE
par(mfrow=c(1,3))
plot(model_lm, which = c(1,2,4)) #plot.lm is plot on a regression generates 6 plots
```



There is a pattern in residuals. Suggets that the relationship is non-linear

Other diagnostics:

- Checking for (approximate) normality with quantile-quantile plot
- Checking for influential observations

Cook's distance, cooksd in the plots, measures how much the predictions for all other observations change if we leave out one observation

Point with high cooksd values