Lab 15

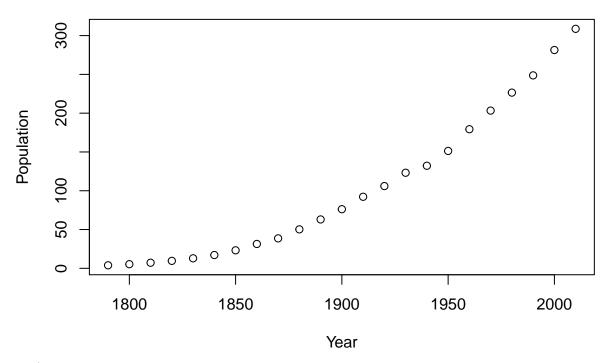
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Exercise 1 - Polynomial Regression. Predict US population

- a) Load the data USpop.csv and plot Population as a function of Year.
- We can see the curve line in the plot below.

```
USpop <- read_csv("./data/USpop.csv")</pre>
## Parsed with column specification:
## cols(
     Year = col_double(),
##
     Population = col_double()
## )
USpop
## # A tibble: 23 x 2
       Year Population
##
      <dbl>
                 <dbl>
##
    1 1790
                    3.9
##
   2 1800
                   5.3
   3 1810
                   7.2
##
   4 1820
                   9.6
##
##
   5 1830
                  12.9
##
    6 1840
                  17.1
      1850
                  23.2
##
    7
       1860
                  31.4
##
##
   9
       1870
                  38.6
## 10 1880
                  50.2
## # ... with 13 more rows
names(USpop)
## [1] "Year"
                    "Population"
attach(USpop)
plot(Year, Population)
```

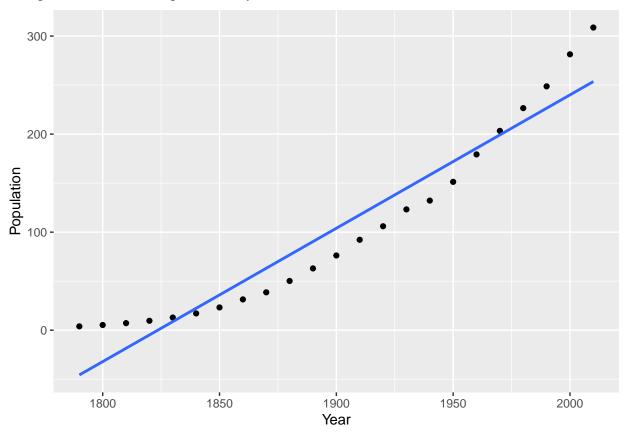


- b) Use a linear model to fit the data. Does a linear model provide a good fit?
- The p-value and R squared is good, maybe the liner model is a good model. However, according to the plot, the model does not provide a good fit. We considered using the polynomial regression model when comparing the regression line and observed data.
- If n is small, the F-stat should be big in order to reject the null (the F-statistic: 239.3 is large).

```
linearModel <- lm(Population~Year)
summary(linearModel)</pre>
```

```
##
## Call:
## lm(formula = Population ~ Year)
## Residuals:
##
       Min
                1Q
                   Median
                                3Q
                                       Max
  -27.774 -24.872
                    -6.295
                                    55.087
                            18.374
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
## (Intercept) -2.481e+03 1.672e+02
                                      -14.84 1.33e-12 ***
##
  Year
                1.360e+00 8.794e-02
                                       15.47 5.93e-13 ***
##
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 27.97 on 21 degrees of freedom
## Multiple R-squared: 0.9193, Adjusted R-squared: 0.9155
## F-statistic: 239.3 on 1 and 21 DF, p-value: 5.927e-13
augment(linearModel) %>%
  ggplot(aes(x = Year, y = Population)) +
  geom_point() +
  geom_smooth(method = lm, se = FALSE)
```

`geom_smooth()` using formula 'y ~ x'



- c) Calculate \mathbb{R}^2 . What do you observe? Does the value of R2 imply that a linear model is a good choice?
- Although R-squared 0.9193 is excellent, we must constantly compare the independent variable and dependent variable plots (overfitting). Then we'll understand why the linear model isn't the best option.
- d) Using the linear model, predict the US population for the year 2030. Is this a good prediction?
- Using linear model to predict Year 2030 is not make sense because the Population in Year 2010 is 308.7 million, but our prediction in Year 2030 is only 280.8202 million.

```
predict(linearModel, data.frame(Year = 2030))

##     1
## 280.8202

USpop %>%
    tail(1)

## # A tibble: 1 x 2
## Year Population
```

- e) Produce appropriate residual plots and decide whether or not an important predictor has been omitted. What do you observe?
 - Residuals vs Fitted plot: strong curve pattern , looks non-linearity

##

1

<dbl>

2010

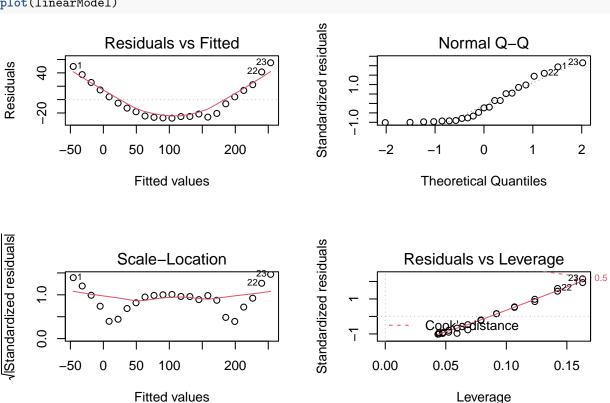
<dbl>

309.

 Normal Q-Q plot: The shape does not appear to follow a normal distribution, particularly the lower tail.

- Scale Location: We could say that the data in the middle is homoscedastic, while the data elsewhere is heteroscedastic.
- Residual vs Leverage: some potential outliers in observation 22 and 23 (high leverage).

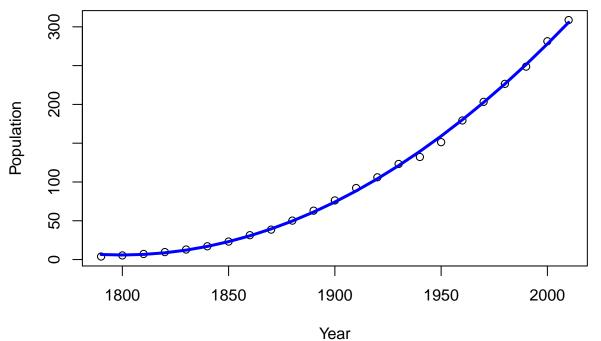
```
par(mfrow = c(2, 2))
plot(linearModel)
```



- f) Fit a quadratic model and plot the fitted curve. Is this a good fit?
- The plot shows that the curve is well fitted. The R-squared also excellent, and each independent variable is significant.

```
quadModel <- lm(Population ~ poly(Year, 2))</pre>
# or: quadModel2 <- lm(Population ~ Year + I(Year^2))
summary(quadModel)
##
## Call:
## lm(formula = Population ~ poly(Year, 2))
##
## Residuals:
##
       Min
                 1Q
                    Median
                                  3Q
                                         Max
   -7.8220 -0.7130
                     0.5961
                             1.8344
                                      3.7487
##
##
## Coefficients:
##
                   Estimate Std. Error t value Pr(>|t|)
                   103.9739
                                 0.6304
                                         164.94
## (Intercept)
                                                   <2e-16 ***
## poly(Year, 2)1 432.7557
                                 3.0231
                                         143.15
                                                   <2e-16 ***
## poly(Year, 2)2 127.4790
                                 3.0231
                                          42.17
                                                   <2e-16 ***
##
                     '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
```

```
##
## Residual standard error: 3.023 on 20 degrees of freedom
## Multiple R-squared: 0.9991, Adjusted R-squared: 0.999
## F-statistic: 1.113e+04 on 2 and 20 DF, p-value: < 2.2e-16
par(mfrow = c(1, 1))
plot(Year, Population)
Yhat = fitted.values(quadModel)
lines(Year, Yhat, col = "blue", lwd = 3)</pre>
```



- g) Predict the population for the year 2030. Is this a reasonable prediction?
- Compared to linear model: 280.8202 million people, the quadratic model is reasonable.

```
predict(quadModel, data.frame(Year = 2030))
##     1
## 365.4891
```

Exercise 2 - Regression Diagnostics

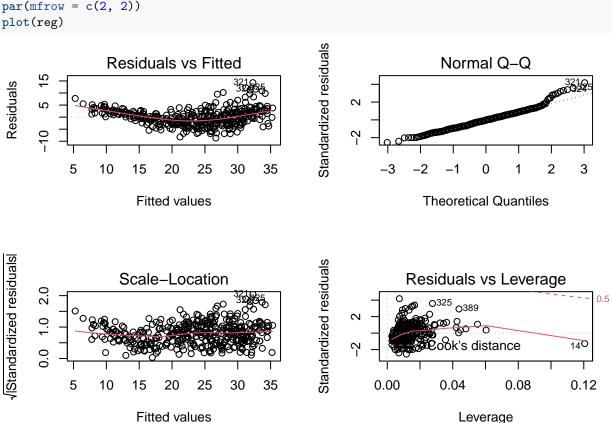
- a) Load the Auto.rda dataset. Predict mpg using the predictors year, accelerator, horsepower, weight. Generate different residual plots.
- Residuals vs Fitted plot: strong curve pattern , looks non-linearity
- Normal Q-Q plot: The shape does not follow a normal distribution, especially the higher tail.
- Scale Location: The data has a slight homoscedasticity to it.
- Residual vs Leverage: we focus on some high leverage observations, they may be potential outliers.

```
load("./data/Auto.rda")
attach(Auto)

## The following object is masked from package:ggplot2:
##
## mpg
```

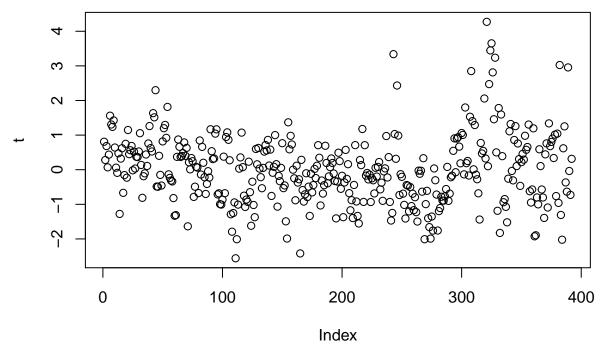
```
reg <- lm(mpg \sim year + acceleration + horsepower + weight)

par(mfrow = c(2, 2))
plot(reg)
```



b) Which of these residuals can be considered as outliers? Compare with the Bonferroni-adjusted quantile from t-distribution.

```
t <- rstudent(reg)
plot(t)
```



Look the summary first

```
summary(t)
               1st Qu.
                          Median
                                       Mean
                                              3rd Qu.
                                                            Max.
## -2.559162 -0.690663 -0.028693
                                  0.001774
                                             0.589435
t[abs(t) > 3]
        243
                 321
                           324
                                    325
                                             328
## 3.338459 4.272284 3.446234 3.651403 3.236226 3.024362
```

#

compute the upper quantile of the t-distribution.

```
qt(0.05/2/392, 387) # qt(alpha/ x/ , n-predictors-1)
## [1] -3.870293
```

compare the t-student residual vs critical value

• There is one outlier

```
• There is one outner

t[abs(t) > abs(qt(0.05/2/392, 387))]

## 321
## 4.272284
```

- c) Test Normality using Shapiro-Wilk normality test. Also look at the Normal Q-Q plot above. Shapiro-Wilk statistic W measures how close the graph is to a straight line.
- The small p-value indicates that there is non-normality (rejected the null).

```
shapiro.test(t)
```

```
##
## Shapiro-Wilk normality test
```

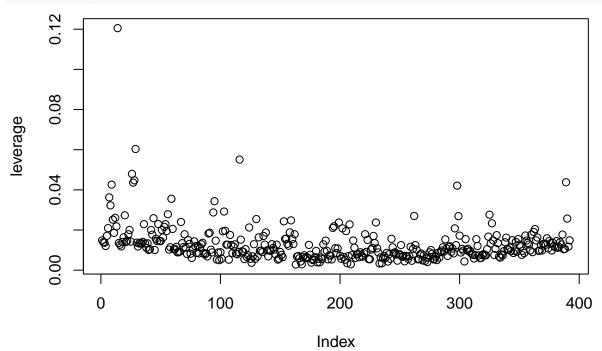
```
##
## data: t
## W = 0.97109, p-value = 5.101e-07
```

- d) Test HOMOSCEDASTICITY (constant variance) using the Breausch-Pagan test.
- A significantly different variance could overshadow the differences between means and lead to incorrect conclusions.
- HOMOSCEDASTICITY is rejected, meaning that there is evidence of equal variance (Heteroskedasticity).

```
library(car)
```

```
## Loading required package: carData
##
## Attaching package: 'car'
   The following object is masked from 'package:dplyr':
##
##
       recode
  The following object is masked from 'package:purrr':
##
##
       some
ncvTest(reg)
## Non-constant Variance Score Test
## Variance formula: ~ fitted.values
## Chisquare = 22.04621, Df = 1, p = 2.6616e-06
  e) Check for INFLUENTIAL DATA
```

infl <- influence(reg)
leverage <- infl\$hat
plot(leverage)</pre>



```
5/length(mpg) # average leverage: mean value
## [1] 0.0127551
summary(infl$hat)
       Min. 1st Qu. Median
                                  Mean 3rd Qu.
                                                     Max.
## 0.002781 0.007543 0.010638 0.012755 0.014735 0.120544
check_leverage <- leverage[leverage > 0.03] # 0.03 is between the third quantile and the max value
check_leverage
                                                                              28
##
                       8
                                  9
                                             14
                                                        26
                                                                   27
\#\#\ 0.03624109\ 0.03226743\ 0.04258253\ 0.12054403\ 0.04797419\ 0.04360256\ 0.04475796
                                                      298
           29
                      59
                                 95
                                           116
## 0.06035105 0.03555978 0.03434446 0.05510052 0.04212120 0.04379524
```