

# Homework 3

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## Setup

```
knitr::opts_chunk$set(echo = TRUE)
library(data.table)

## Warning: package 'data.table' was built under R version 4.0.5
library(broom)

## Warning: package 'broom' was built under R version 4.0.5
library(gridExtra)

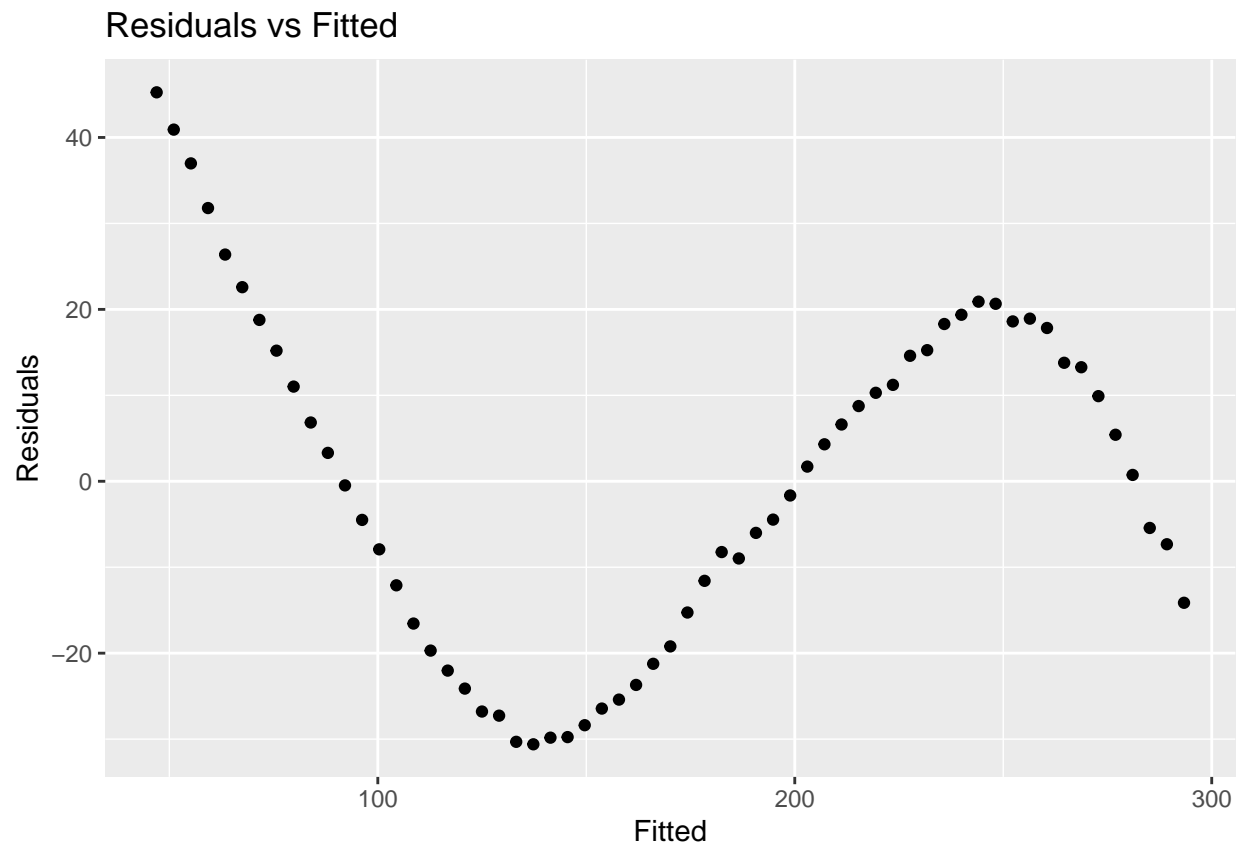
## Warning: package 'gridExtra' was built under R version 4.0.5
library(ggplot2)

## Warning: package 'ggplot2' was built under R version 4.0.5
covid_raw <- fread("https://opendata.ecdc.europa.eu/covid19/casedistribution/csv")
us <- covid_raw[covid_raw$countriesAndTerritories == 'United_States_of_America',]
us_filtered <- us[us$month %in% c(6:7),]
us_filtered$index <- rev(1:dim(us_filtered)[1])
fit<-lm(~Cumulative_number_for_14_days_of_COVID-19_cases_per_100000~index, data=us_filtered)

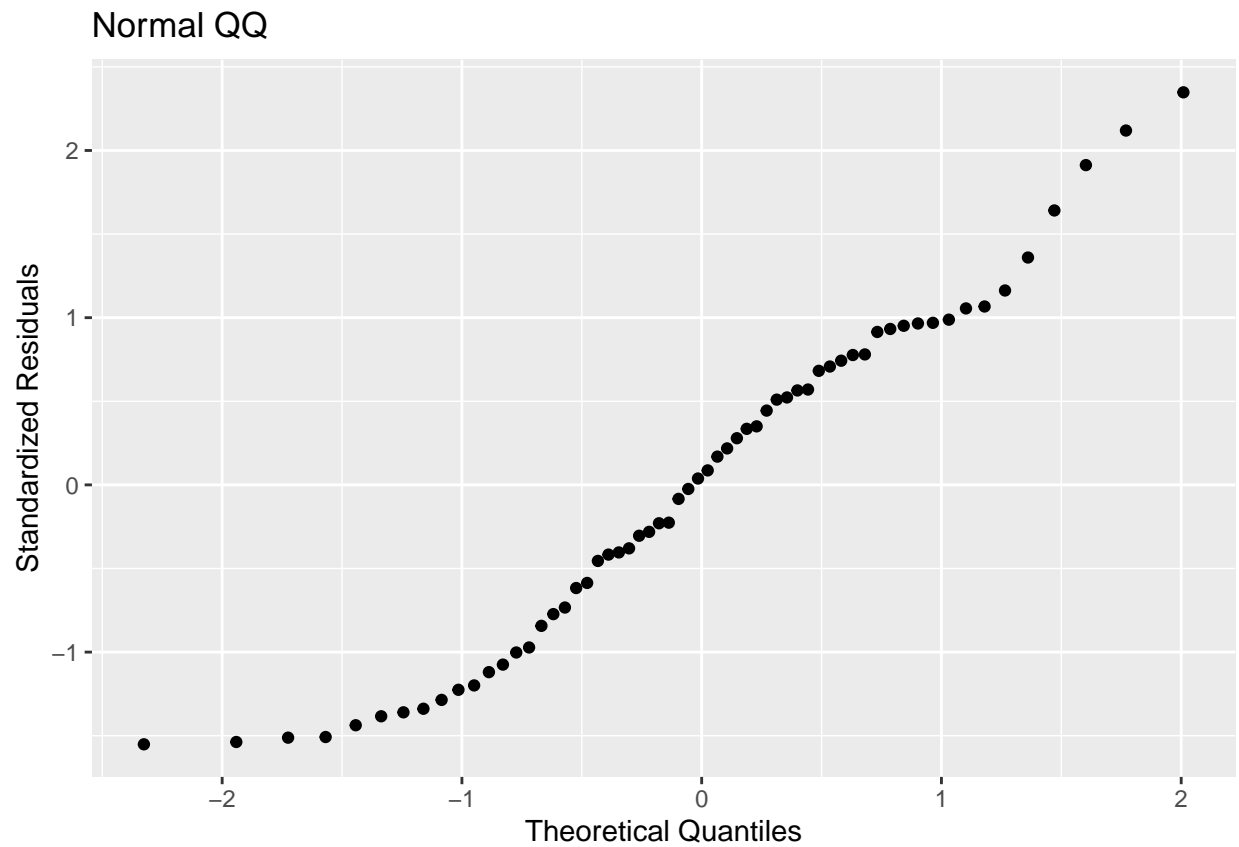
fit.diags <- broom::augment(fit)
```

## Problem 3

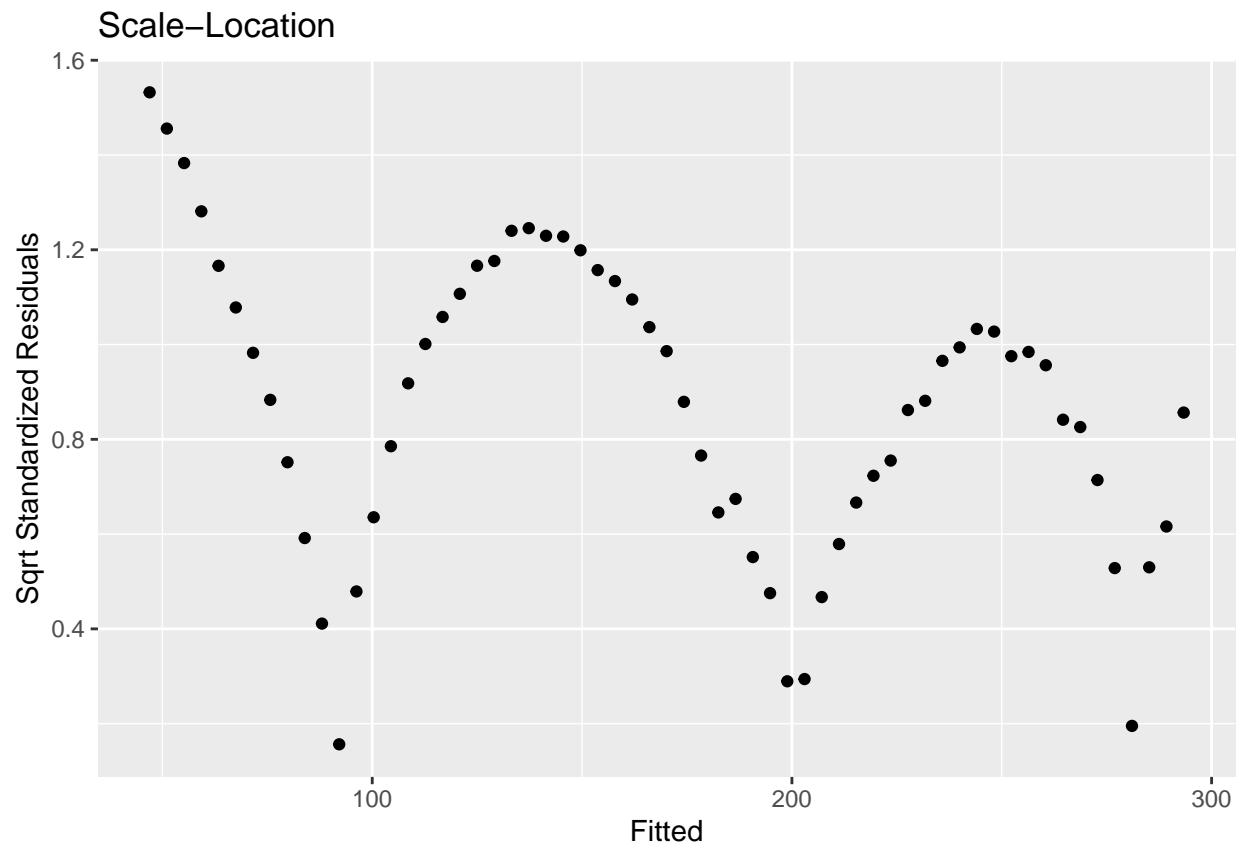
```
rvf <- ggplot(fit.diags, aes(x=.fitted, y=.resid))+
  geom_point() +
  labs(x="Fitted", y="Residuals", title="Residuals vs Fitted")
rvf
```



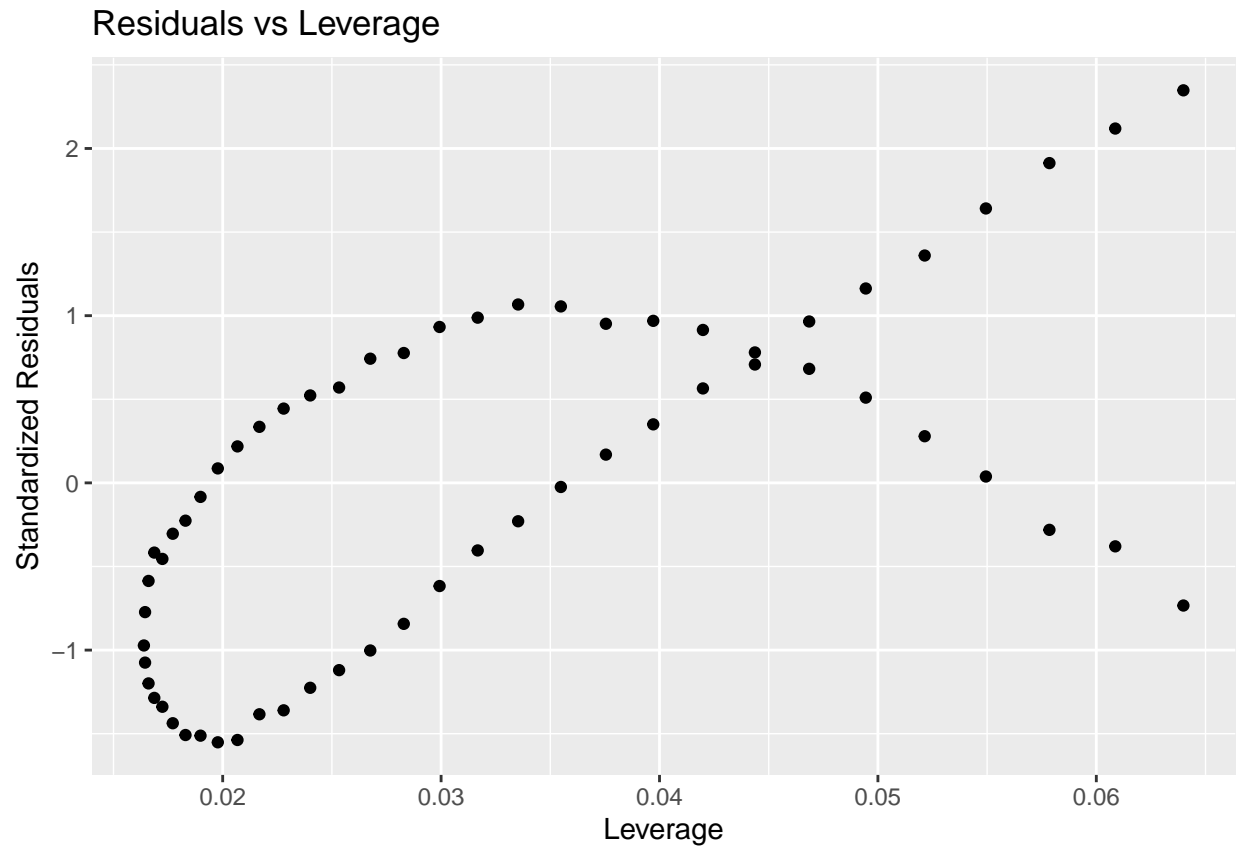
```
qq <- ggplot(fit.diagnostics, aes(x=qnorm(seq(0.01,0.99,1/(nrow(fit.diagnostics) + 1))), y=.std.resid[order(.std.resid)])) +  
  geom_point() +  
  labs(x="Theoretical Quantiles", y="Standardized Residuals", title="Normal QQ")  
qq
```



```
s1 <- ggplot(fit.diagnostics, aes(x=.fitted, y=sqrt(abs(.std.resid))))+  
  geom_point() +  
  labs(x="Fitted", y="Sqrt Standardized Residuals", title="Scale-Location")  
  
s1
```



```
rvl <- ggplot(fit.diags, aes(x=.hat, y=.std.resid))+  
  geom_point() +  
  labs(x="Leverage", y="Standardized Residuals", title="Residuals vs Leverage")  
rvl
```



# Problem 4

```
grid.arrange(rvf, qq, sl, rvl, ncol = 2, nrow = 2)
```

