HW3_boyuj

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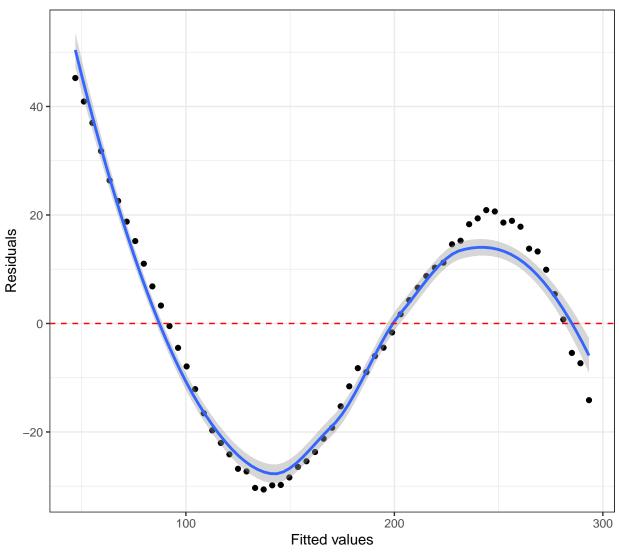
Problem 3 and 4

```
library(data.table)
covid_raw <- fread("https://opendata.ecdc.europa.eu/covid19/casedistribution/csv")</pre>
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])</pre>
fit < -lm(\Cumulative_number_for_14_days_of_COVID-19_cases_per_100000\cdot\cdot\cdot\cdot\cdot
        data=us_filtered)
## augment the data as previous
library(broom)
fit.diags <- broom::augment(fit)</pre>
library(ggplot2)
library(gridExtra)
plot1 <- ggplot(fit, aes(.fitted, .resid))+</pre>
  geom_point()+
  stat_smooth(method="loess")+
  geom_hline(yintercept=0, col="red", linetype="dashed")+
  xlab("Fitted values")+ylab("Residuals")+
  ggtitle("Residuals vs Fitted")+theme bw()
plot2 <- ggplot(fit, aes(qqnorm(.stdresid, plot.it=FALSE)[[1]], .stdresid))+</pre>
  geom_point(na.rm = TRUE)+
  xlab("Theoretical Quantiles")+
  ylab("Standardized Residuals")+
  ggtitle("Normal Q-Q")+theme_bw()
plot3 <- ggplot(fit, aes(.fitted, sqrt(abs(.stdresid))))+</pre>
  geom_point(na.rm=TRUE)+
  stat_smooth(method="loess", na.rm = TRUE)+
  xlab("Fitted Value")+
  ylab(expression(sqrt("|Standardized residuals|")))+
  ggtitle("Scale-Location")+theme_bw()
plot4 <- ggplot(fit, aes(.hat, .stdresid))+</pre>
  geom_point(aes(size=.cooksd), na.rm=TRUE)+
  stat_smooth(method="loess", na.rm=TRUE)+
  xlab("Leverage")+
```

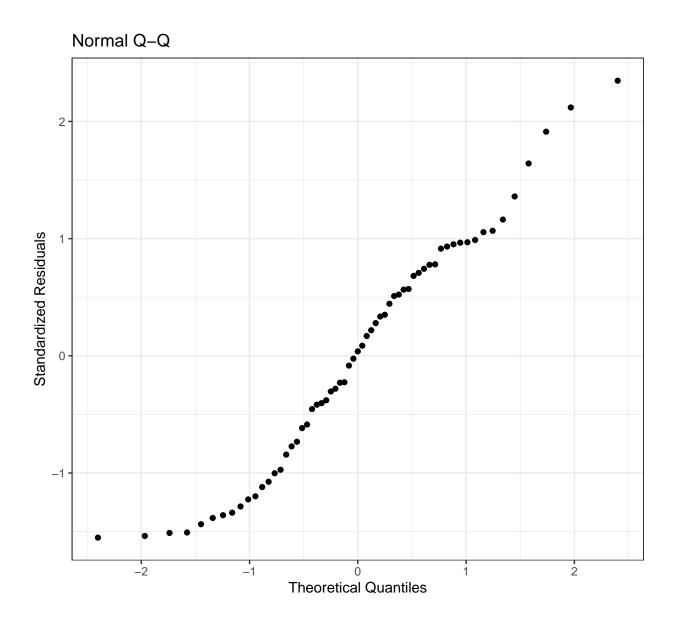
```
ylab("Standardized Residuals")+
ggtitle("Residuals vs Leverage")+
scale_size_continuous("Cook's Distance", range=c(1,5))+
theme_bw()+theme(legend.position="bottom")
plot1
```

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

Residuals vs Fitted

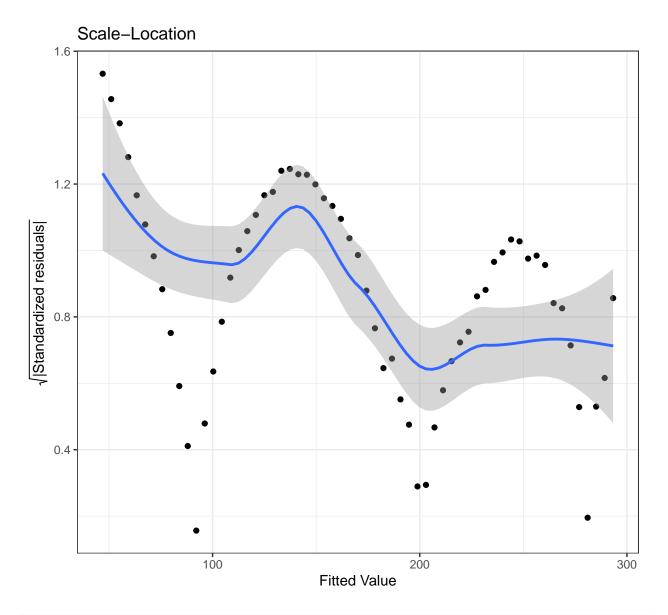


plot2



plot3

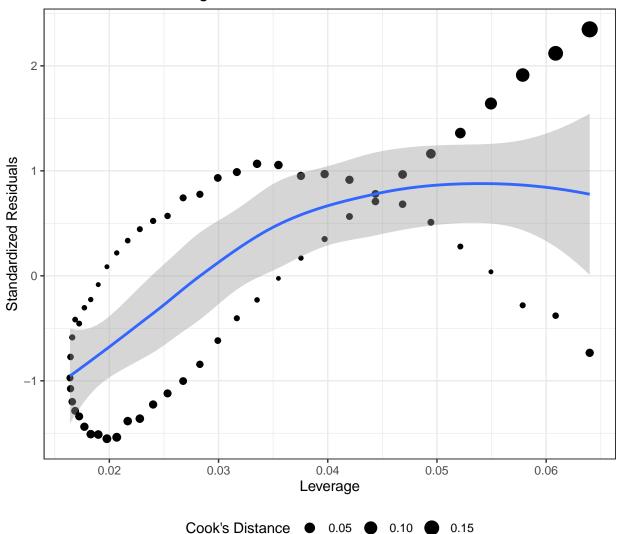
'geom_smooth()' using formula 'y ~ x'



plot4

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

Residuals vs Leverage



grid.arrange(plot1, plot2, plot3, plot4, ncol = 2, nrow = 2)

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

