

week10-pdfgen

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Data Import and Cleaning

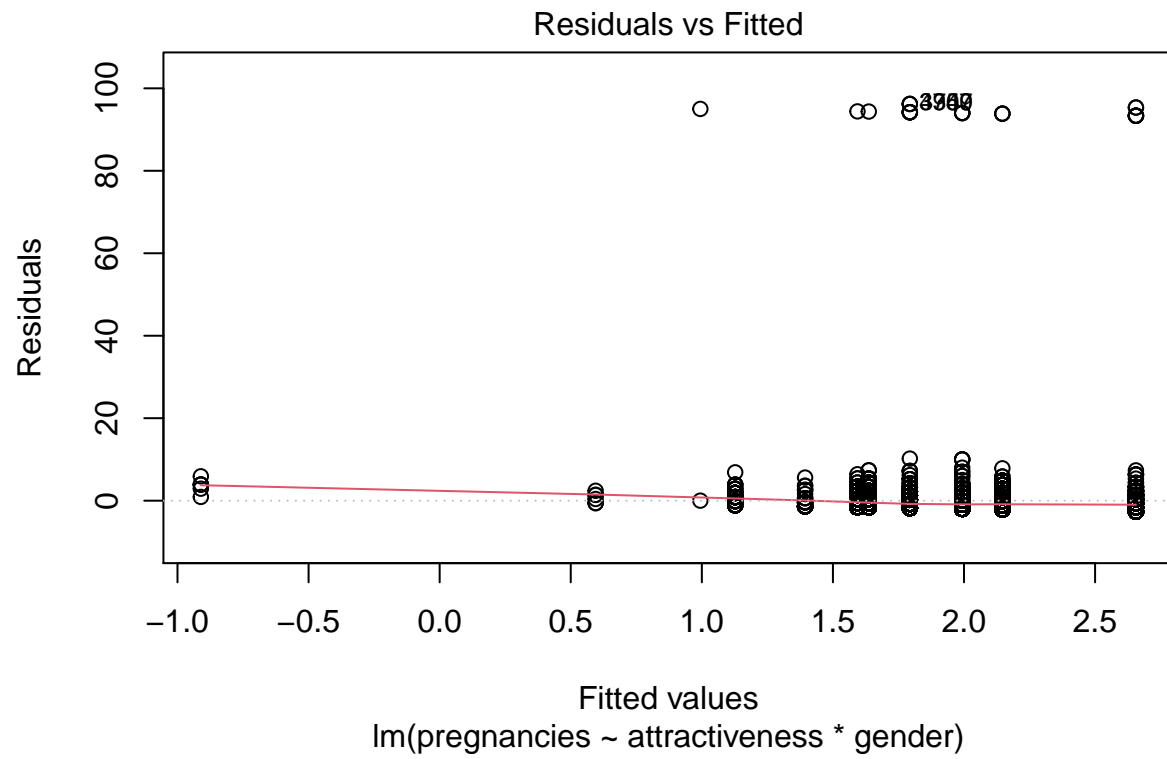
In this block I imported the National Longitudinal Study of Adolescent to Adult Health Wave IV, 2008 dataset from the UNC Dataverse website. I selected three variables and converted gender to a factor variable.

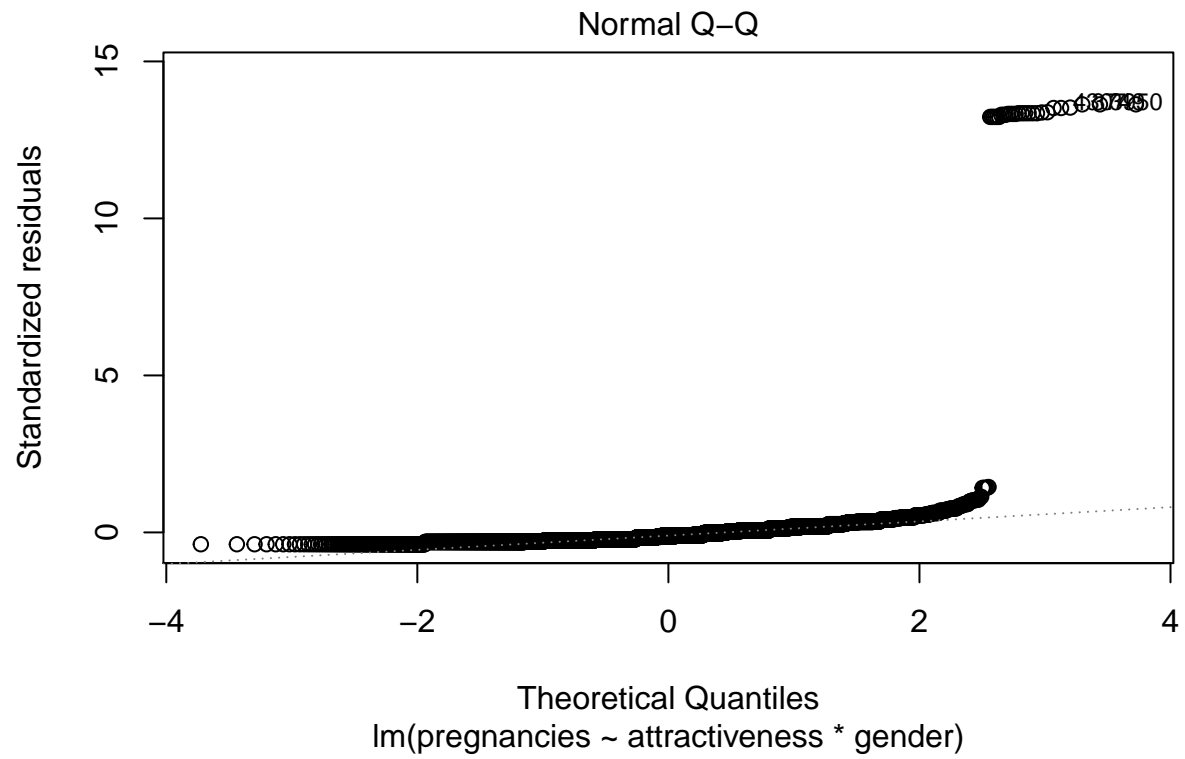
```
health_tbl <- read_tsv("../data/w4inhome_dvn.tab") %>%  
  select(attractiveness = H4MH8,  
         pregnancies = H4TR9,  
         gender = BIO_SEX4) %>%  
  mutate(gender = factor(gender, levels = c(1,2), labels = c("Male", "Female")))
```

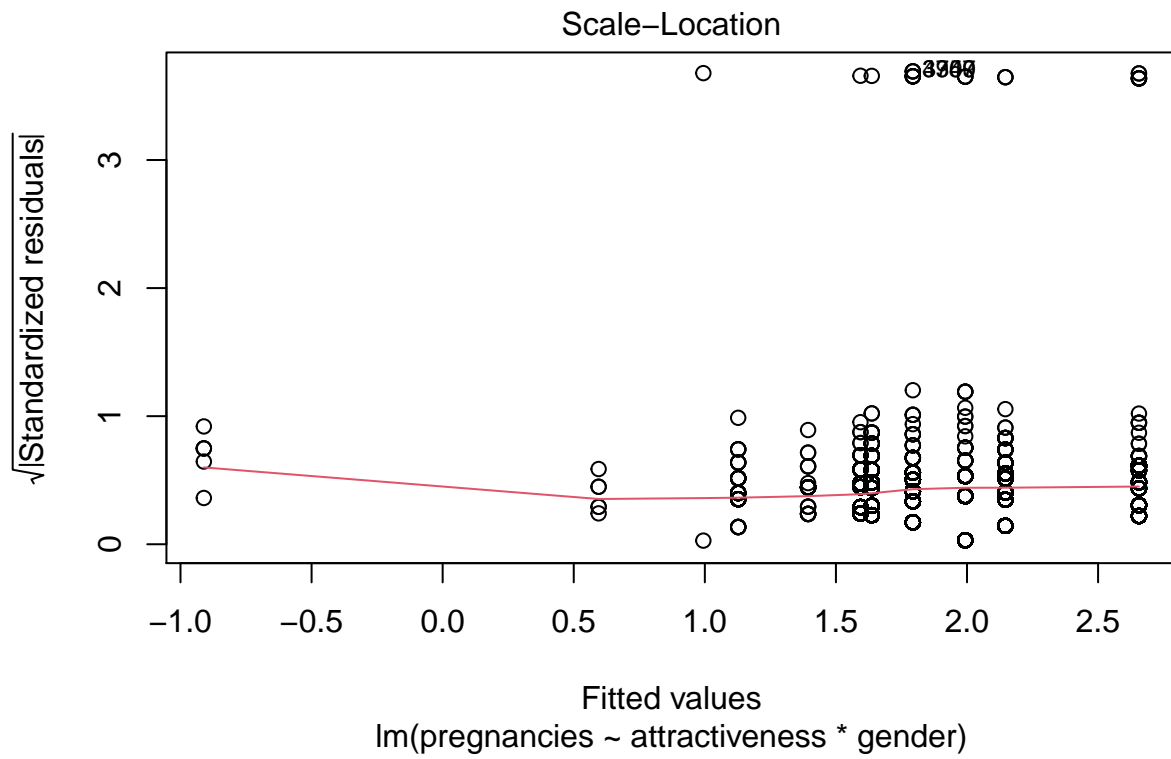
Analysis

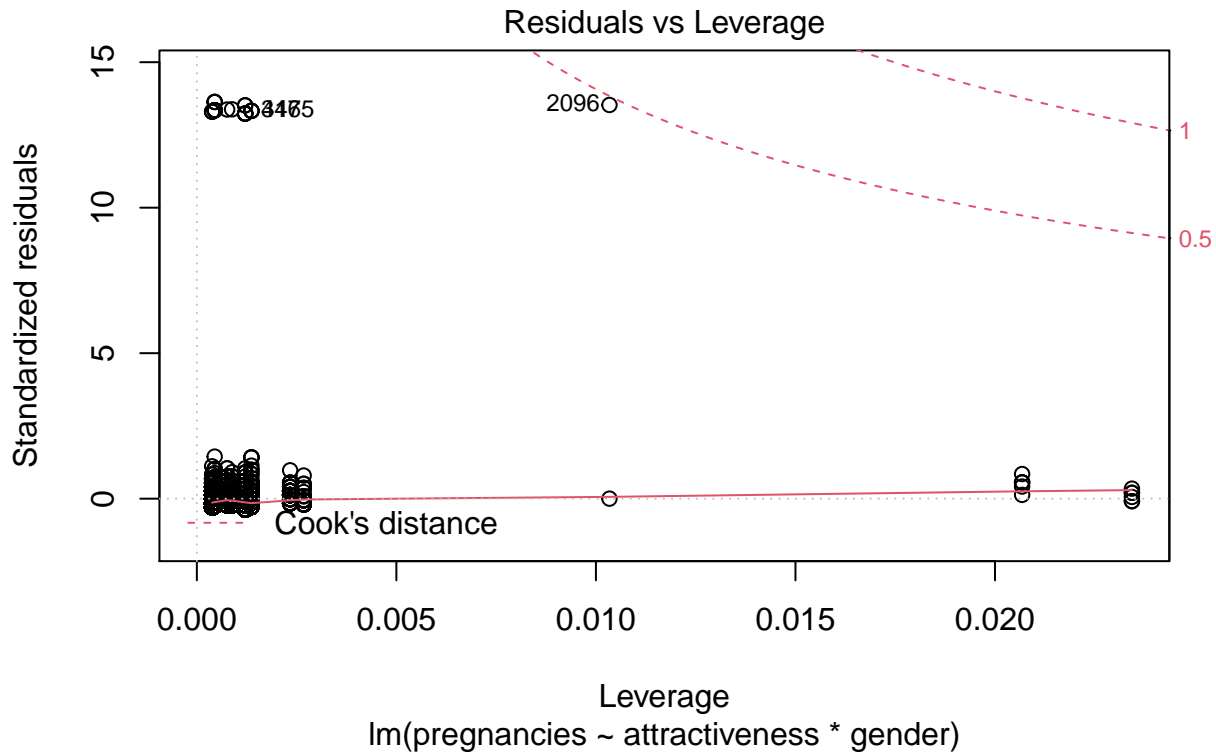
In this block I conducted a multiple linear regression of pregnancies on the two independent variables attractiveness and gender, checked assumptions and examined a coefficients table.

```
model <- lm(pregnancies ~ attractiveness * gender, data = health_tbl) #The specification first*second i  
plot(model)
```









```
summary(model)
```

```
##
## Call:
## lm(formula = pregnancies ~ attractiveness * gender, data = health_tbl)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -2.656 -1.793 -0.793  0.363  96.207
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      2.1929    0.4263   5.145 2.78e-07 ***
## attractiveness  -0.1998    0.1839  -1.086   0.277
## genderFemale      0.9724    0.5858   1.660   0.097 .
## attractiveness:genderFemale -0.3098    0.2526  -1.227   0.220
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 7.058 on 5110 degrees of freedom
## Multiple R-squared:  0.002349, Adjusted R-squared:  0.001764
## F-statistic: 4.011 on 3 and 5110 DF, p-value: 0.007314
```

Attractiveness was statistically significant ($t = 0.2521$, $p = 0.012$) but gender and interaction were not.

#Visualization I visualize the interaction using the fitted variables.

```
ggplot(health_tbl, aes(x = attractiveness, y = pregnancies, color = gender)) +
  geom_smooth(method = "lm") +
  geom_point()
```

