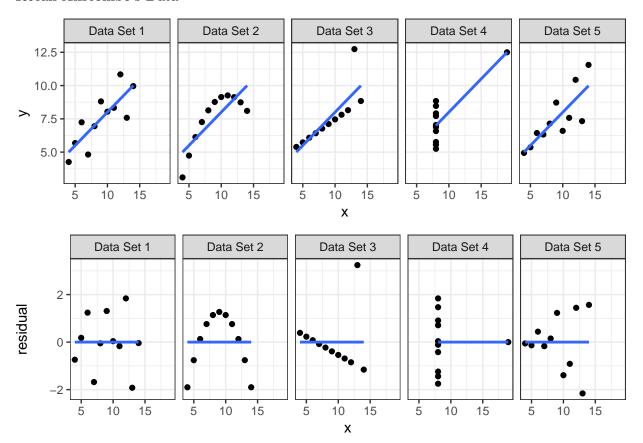
Chapter 11: Outliers and Influential Observations

Recall Anscombe's Data



- For today, let's focus on Data Sets 3 and 4
- Definitions (note: there are not universally agreed on definitions for these terms):
 - An **outlier** is an observation that "doesn't fit" with the patterns in the rest of the data
 - * Both data set 3 and data set 4 have outliers
 - An **influential observation** is an observation whose removal from the data set would substantially change the model fit (coefficient estimates)
 - * Both data set 3 and data set 4 have influential observations
 - * The point in data set 4 is more influential
 - A high leverage observation is one whose explanatory variable values are far from the explanatory variable values of other observations
 - * Data set 4 has a high leverage observation
 - * Data set 3 does not
- Note that residual plots exactly fail to identify very influential/high leverage observations!!!

Leverage

• If the model has 1 X variable, the leverage of observation i is defined to be

$$h_i = \frac{(X_i - \bar{X})^2}{\sum_{j=1}^n (X_j - \bar{X})^2} + \frac{1}{n}$$

- Basically, how far is X_i from \bar{X} , standardized in an obsure way.

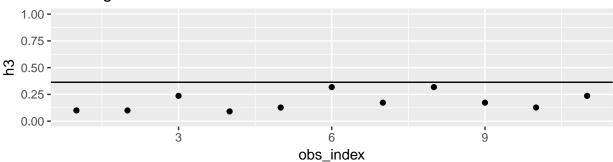
 - $-\frac{1}{n} \le h_i \le 1$ Average of leverages is p/n (where p is the number of parameters for the mean in the model).
- As a very rough guide, $h_i > 2p/n$ indicates an observation is worth looking into more

Plots of leverage vs. observation index (code will be shown later)

2p/n; p = 2 since we have beta_0 and beta_1 in our simple linear regression model 2 * 2 / nrow(anscombe)

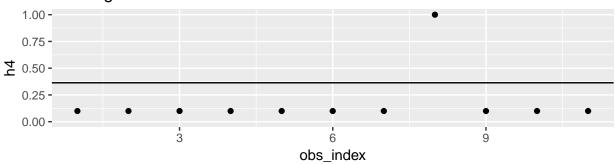
[1] 0.3636364

Leverage - Data Set 3



Looks OK

Leverage - Data Set 4



confirm observation 8 is the one with a big X! anscombe\$x4[8]

[1] 19

Studentized Residuals

• Observations with high leverage tend to have small residuals!

$$-SD(res_i) = \sigma\sqrt{1 - h_i}$$

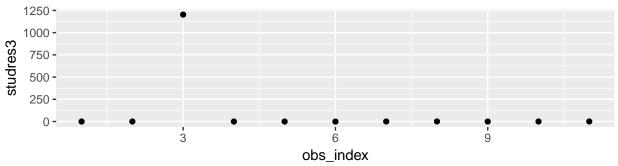
- Looking at just the residuals can be misleading
- The studentized residuals adjust by dividing residual by its estimated standard deviation

$$studres_i = \frac{res_i}{\hat{\sigma}\sqrt{1-h_i}}$$

- A studentized residual less than -2 or greater than 2 could indicate problems if other diagnostics also indicate issues
- We expect about 5% of studentized residuals to be less than -2 or greater than 2.

Plots of studentized residuals vs. observation index (code will be shown later)

Studentized Residuals – Data Set 3

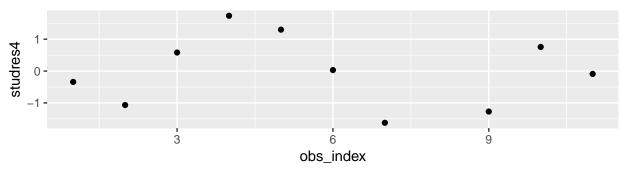


confirm observation 3 is the one with a big Y!
anscombe\$y3[3]

[1] 12.74

Warning: Removed 1 rows containing missing values (geom_point).

Studentized Residuals – Data Set 4



Looks OK... what's up with that warning?

anscombe\$studres4

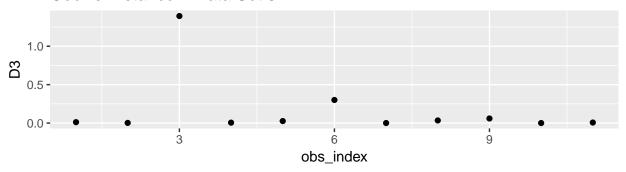
anscombe\$h4

Cook's Distance

- Measures how different predicted values for all observations are when observation i is or is not used for model estimation
- Fit model using all observations; get predicted values \hat{Y}_j for each $j=1,\ldots,n$
- Fit model using all observations other than i; get predicted values $\hat{Y}_{j(i)}$ for each $j=1,\ldots,n$
- $D_i = \frac{\sum_{j=1}^n (\hat{Y}_{j(i)} \hat{Y}_j)^2}{p\hat{\sigma}^2}$ As a very rough guide, $D_i > 1$ indicates an observation is worth looking into more

Plots of Cook's distance vs. observation index (code will be shown later)

Cook's Distance – Data Set 3

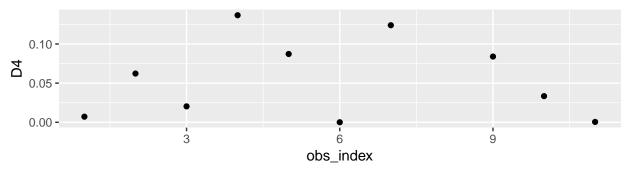


confirm observation 3 is the one with a big Y! anscombe\$y3[3]

[1] 12.74

Warning: Removed 1 rows containing missing values (geom_point).

Cook's Distance - Data Set 4



Looks OK... what's up with that warning?

anscombe \$D4

anscombe\$h4

1 2 3 4 5 6 7 8 9 10 11

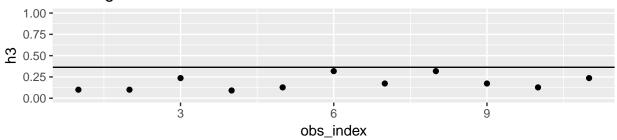
R Code: Manual Plots

- Every statistical software package will give you different plots by default
- Our book suggests the plots we've looked at so far, which are not the defaults for R/require more code to create:

```
anscombe <- anscombe %>%
  mutate(
   obs_index = row_number(),
   h3 = hatvalues(fit3),
   studres3 = rstudent(fit3),
   D3 = cooks.distance(fit3)
)

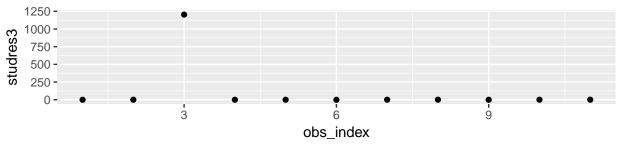
ggplot(data = anscombe, mapping = aes(x = obs_index, y = h3)) +
  geom_point() +
  geom_hline(yintercept = 2 * 2 / nrow(anscombe)) +
  ylim(0, 1) +
  ggtitle("Leverage - Data Set 3")
```

Leverage – Data Set 3



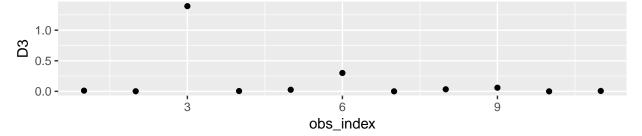
```
ggplot(data = anscombe, mapping = aes(x = obs_index, y = studres3)) +
geom_point() +
ggtitle("Studentized Residuals - Data Set 3")
```

Studentized Residuals – Data Set 3



```
ggplot(data = anscombe, mapping = aes(x = obs_index, y = D3)) +
geom_point() +
ggtitle("Cook's Distance - Data Set 3")
```

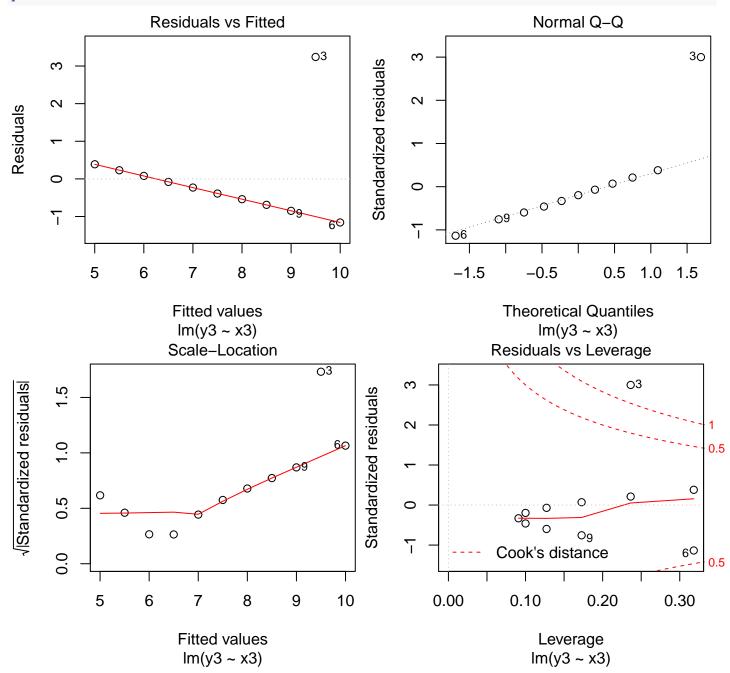
Cook's Distance - Data Set 3



R Code: Default Plots

You can get a set of different diagnostic plots more easily, but I find the plot involving Cook's distance and Leverage less intuitive:





Note: to get the plots to all show up in the knitted pdf, I had to set figure height and width in the code chunk declaration: ```{r, fig.height = 4, fig.width = 4}