

SDS358: Applied Regression Analysis

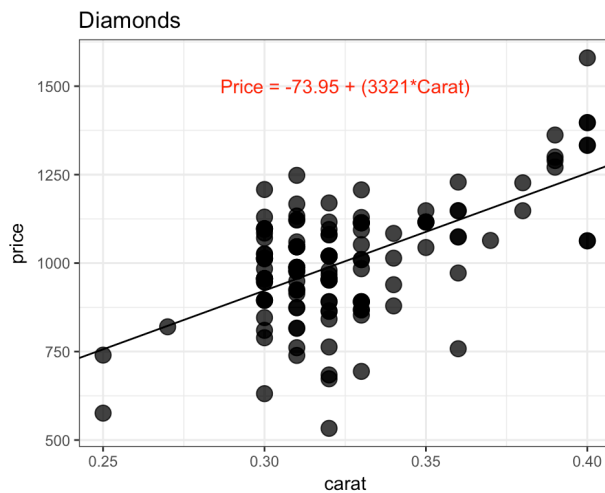
Day 6: SLR Continued

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Agenda for Today:

- Simple Linear Regression continued
 - Assumptions
 - Basis in residuals (e_i)
 - Outlier checks

Recall:



Interpretation:

- Slope
- Intercept

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Assumptions of Simple Linear Regression

- Quantitative Data for both variables.
- The relationship is linear in nature.
- Residuals
 - Independence: Error associated with each data point is independent of every other value.
 - For a given value of x , e has a normal distribution, meaning:
 - The population mean of e is 0.
 - For a given value of x , the population variance of e is σ_e^2
 - Homoscedasticity
 - No Outliers

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Residuals: Independence

- Every observation in the dataset must be unique.
- A subject (or record), cannot be recorded twice, and take two lines.
 - In other words, the same person cannot contribute twice (even with different data) to the prediction of y .

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Residuals: Population mean of zero

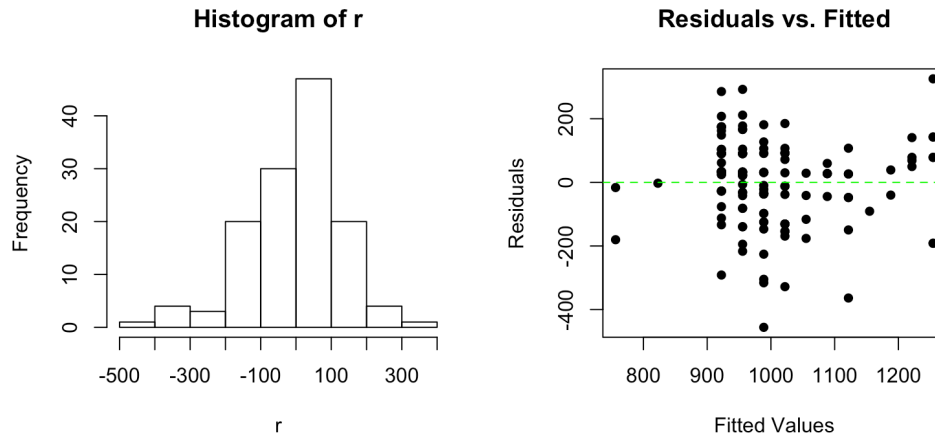
- This is easy: just record each observation's residual after fitting the model.
- Then, grab the mean...or better yet, a histogram

```
#Diamonds  
d_mod <- lm(price ~ carat, data=diamonds)  
r <- d_mod$residuals  
histogram(r)  
residFitted(d_mod)
```

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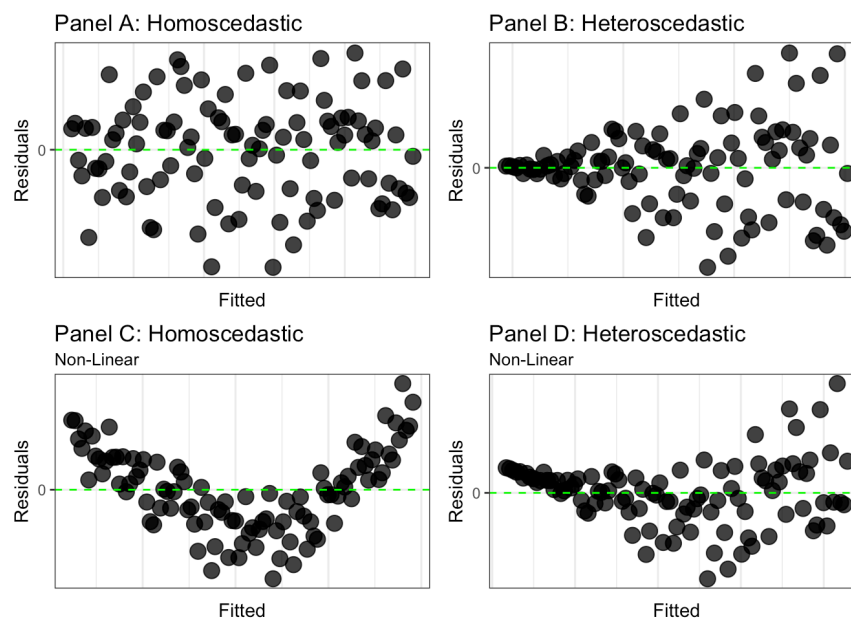
Residuals: Population mean of zero

- This is easy: just record each observation's residual after fitting the model.
- Then, grab the mean...or better yet, a histogram



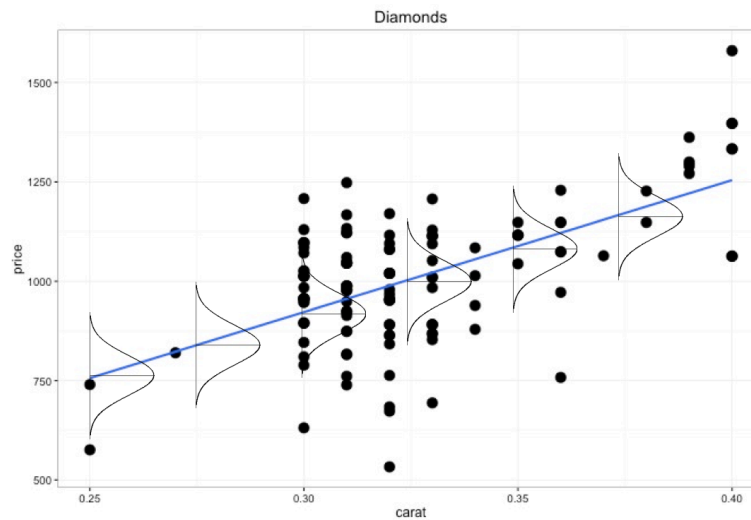
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Residuals: What to watch out for....



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Residuals: normal around each x



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Becuase Residuals are NORMAL:

- Measure of "overall error" in the regression model.

$$\text{Model Variance} = \sigma_e^2$$

- We can't actually measure this (it's a population value).
- But, we can estimate it with S_e^2 .

$$S_e^2 = \text{Estimate of } \sigma_e^2$$

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What is "sigma"?

- S_e^2 is the variance of the overall error
- S_e is the *average* of the overall error:

$$S_e = \sqrt{S_e^2}$$

- HINT: Think *standard deviation* but for regression model residuals.

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What is "sigma"?

- AGAIN: Think *standard deviation* but for regression model residuals.

$$S_e = \sqrt{S_e^2} = \sqrt{\frac{\sum(y - \hat{y})^2}{(n - p)}}$$

p = Number of "parameters"

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What is "sigma"?

- AGAIN: Think *standard deviation* but for regression model residuals.

$$S_e = \sqrt{S_e^2} = \sqrt{\frac{\sum (y - \hat{y})^2}{(n - k - 1)}}$$

k = Number of "predictors"

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What else are the residuals good for?

- What's the link between r , r^2 , and the linear model?
- Let's head out to RStudio and investigate....

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Proportion of Variance

- r^2 is defined as the "Proportion of variance accounted for."
- Literally:

"How much of the variance in y can be accounted for by x (or all x's), now that we know the relationship between y and x (or y and all x's)."

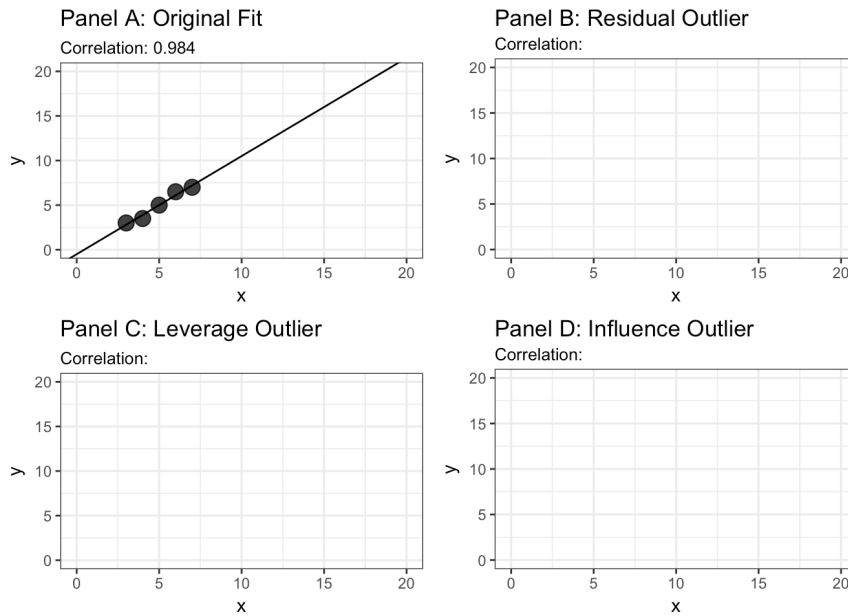
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SLR Diagnostics

- Residuals help show us if something is "off" in our model/data.
- *But* they, are just a part of a larger tool set to diagnose the validity of a model and/or look for problematic data
- We'll use *three* primary tools to look for bad data:
 - Studentized deleted residuals
 - Hat values (also known as leverage points)
 - Cook's Distance

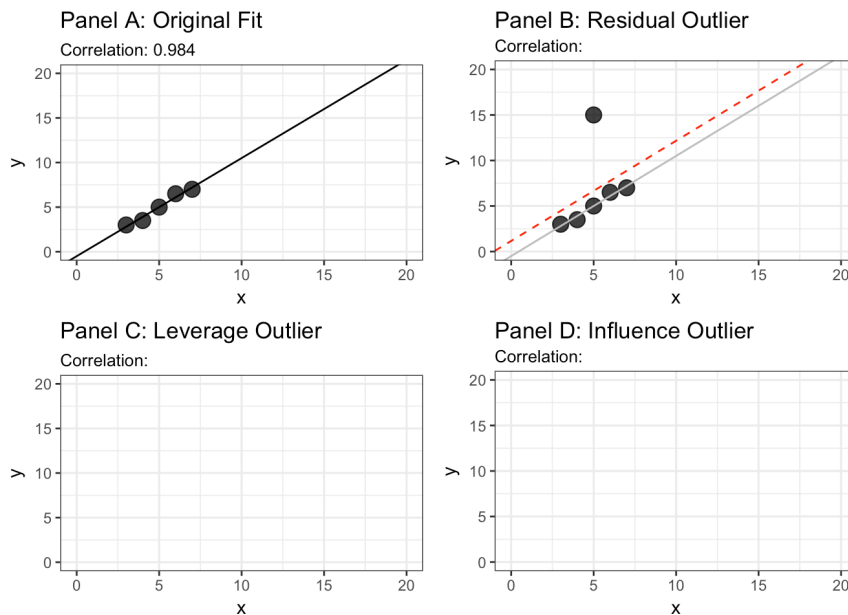
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Some invented data....



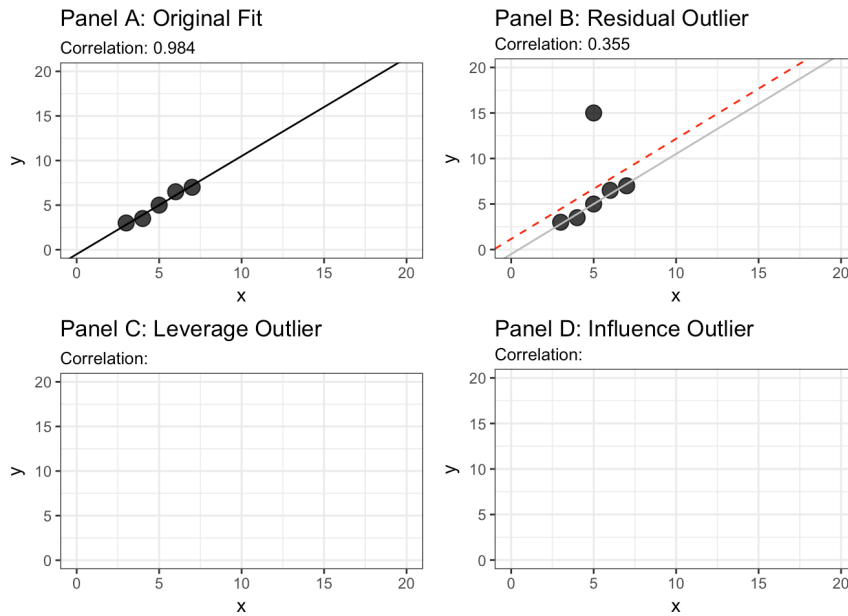
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Some invented data....You Guess



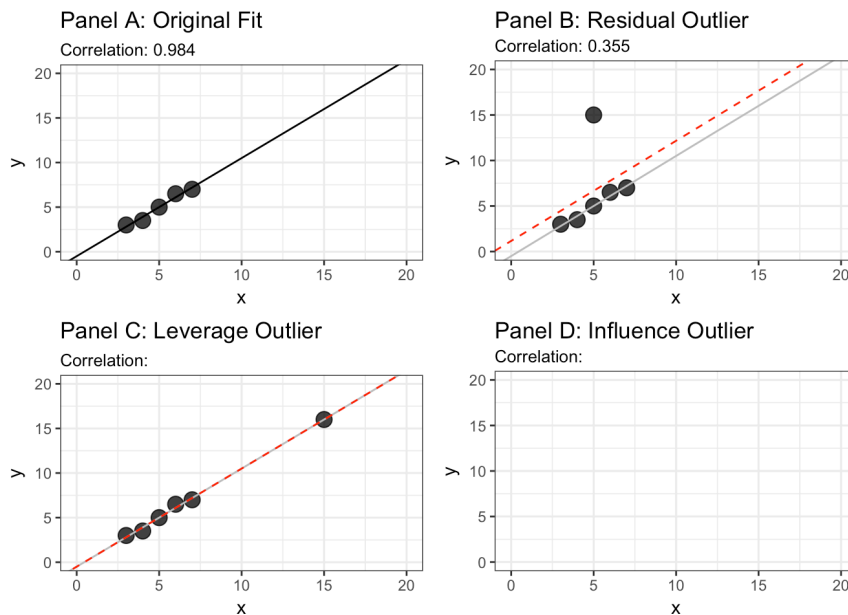
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Some invented data....



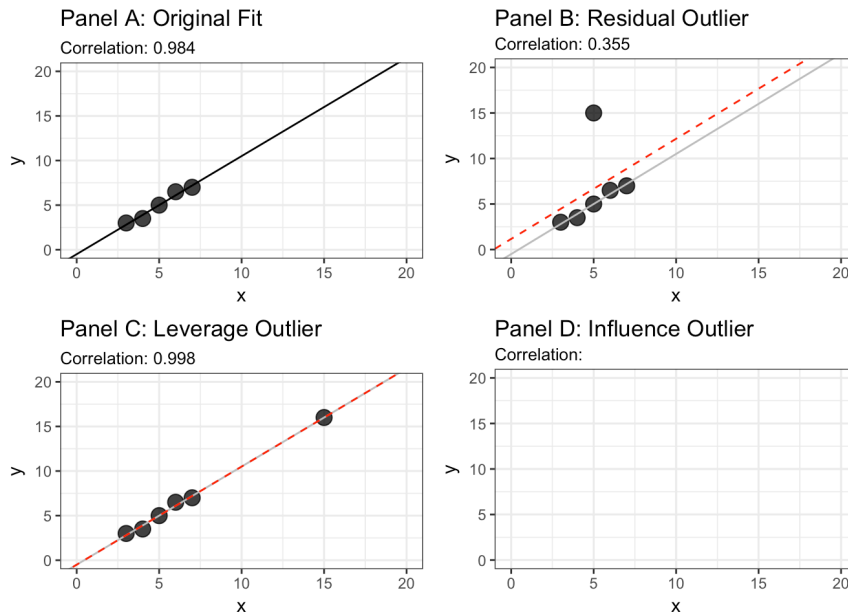
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Some invented data....You Guess



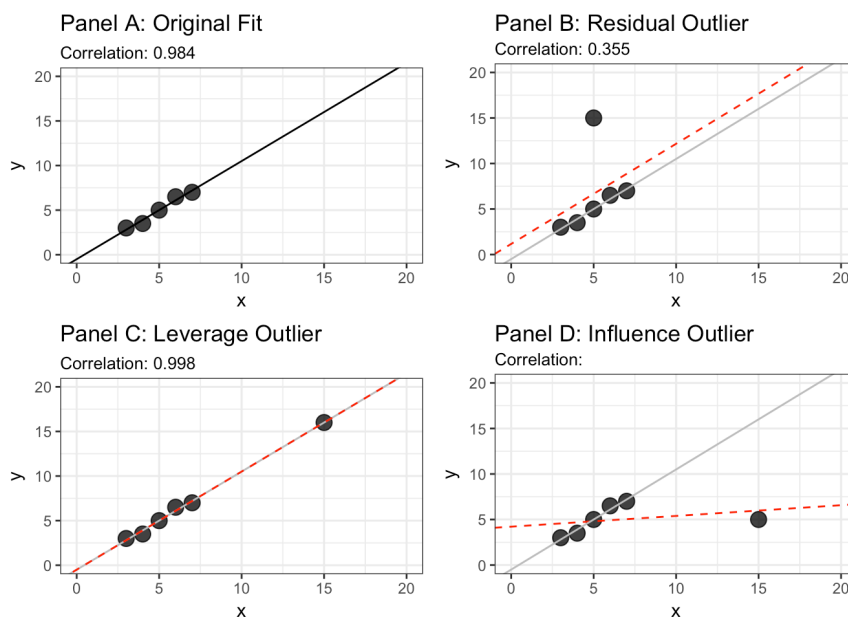
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Some invented data...



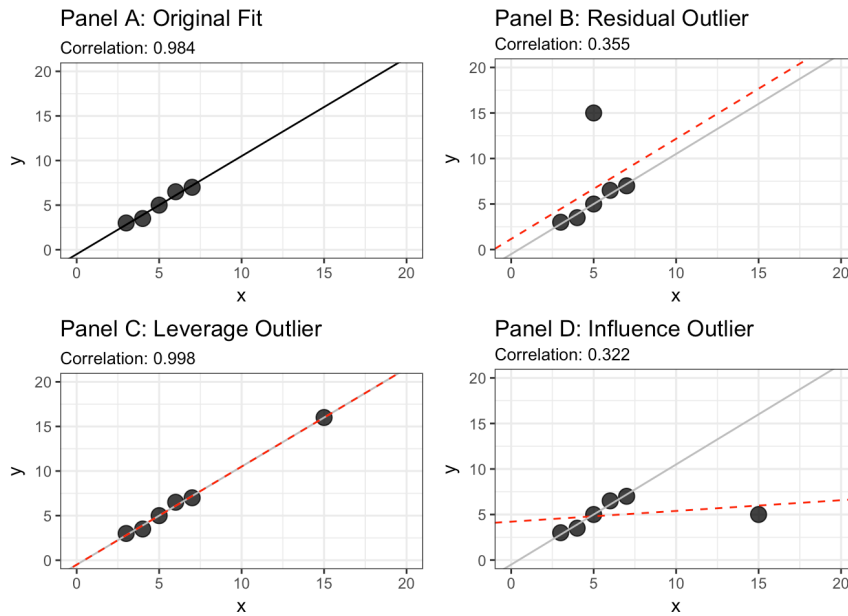
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Some invented data....You Guess



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Some invented data...



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SLR Diagnostics

- So which diagnostic do you choose?
 - Which is *most* important?

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SLR Diagnostics

- So which diagnostic do you choose?
 - Which is *most* important?
- Cook's distance
 - All three markers would take precedence
 - It's NOT black and white...

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SLR Diagnostics: Leverage

- The leverage of a single point is measured by the *hat* value.
- In simple regression:

$$\text{Leverage}_i = h_i = \frac{1}{n} + \frac{(x - \bar{x})^2}{\sum (x - \bar{x})^2}$$

- Think: "The farther away from the mean of x, the higher the hat value."
- Look for: 2 or 3 time the "average h_i "

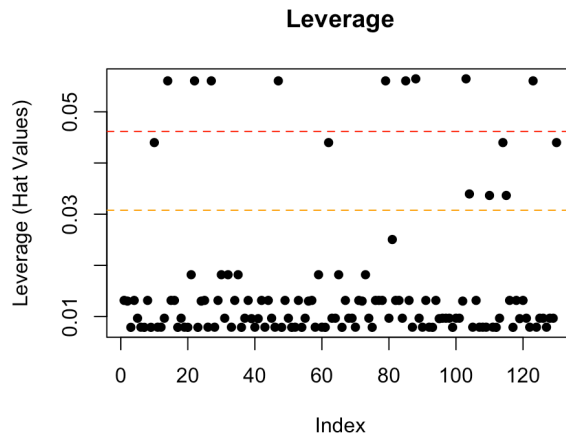
$$h_i > 2 * ((k + 1)/n)$$

$$h_i > 2 * \left(\frac{\text{Number of } \beta \text{ predictors in the model incl. } \beta_0}{n} \right)$$

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SLR Diagnostics: Leverage

```
levPlot(d_mod)
```



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SLR Diagnostics: Leverage

```
levPlot(d_mod, key.variable = "sampleDiamond",  
        print.obs=TRUE, sort.obs = TRUE)
```

##	sampleDiamond	price	carat	Predicted_Y	Hat_Values
## 1	88	740	0.25	756.2989	0.05644717
## 2	103	576	0.25	756.2989	0.05644717
## 3	14	1397	0.40	1254.4485	0.05604876
## 4	22	1333	0.40	1254.4485	0.05604876
## 5	27	1397	0.40	1254.4485	0.05604876
## 6	47	1580	0.40	1254.4485	0.05604876
## 7	79	1063	0.40	1254.4485	0.05604876
## 8	85	1333	0.40	1254.4485	0.05604876
## 9	123	1063	0.40	1254.4485	0.05604876

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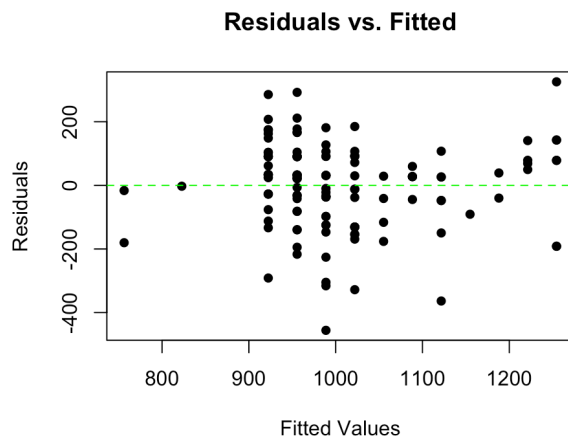
SLR Diagnostics: Residuals

- Looking at residuals is good too.
- Even looking at fitted values and examining values them in relation to some error of e (*Homoscedasticity*)

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SLR Diagnostics: Residuals

```
residFitted(d_mod)
```



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SLR Diagnostics: Standardized Residuals

- But, there's a **problem** with residuals: *how to evaluate them*.
- Comparing them to a value like a *z-score* might work.
- Then we can use the *sigma* of the model (the RMSE = 134.68):

$$s_e = \sqrt{\frac{(y - \hat{y})^2}{n - k - 1}}$$

- If we use this though, we can run into an issue: what happens to sigma with an observation with a large residual?

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SLR Diagnostics: Studentized Residuals

- On *TOP* of that, we know that Leverage plays a key role in influencing the SLR Model.
- So, we should recognize that the RMSE (s_e) is, *by itself* not great as a denominator.
- We can *incorporate* Leverage into the denominator and get:

$$\text{Studentized Residual}_i = E'_i = \frac{e_i}{s_e \sqrt{1 - h_i}}$$

with:

$$e_i = (y - \hat{y})^2$$

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SLR Diagnostics: Studentized DELETED Residuals

- Studentized Residuals are *better* than Standardized, but there's *still the issue* of a high residual outlier.
- To help "correct" for the effect of a single problematic residual, while still taking into account for Leverage, we can use a better calculation—through *DELETION*.

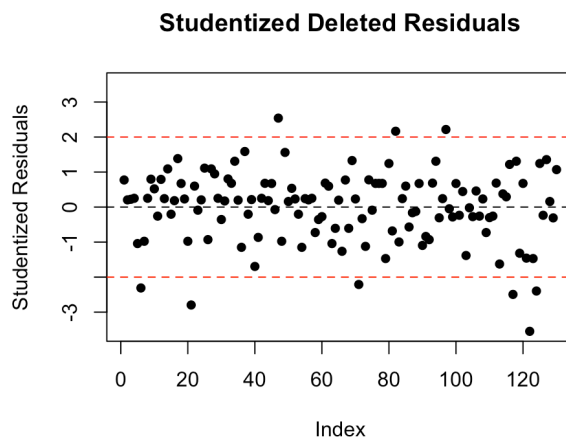
$$\text{Studentized Deleted Residual}_i = E_i^* = \frac{e_i}{S_{e(-i)} \sqrt{1 - h_i}}$$

- Look for: 2 times a "studentized" residual (absolute value).

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SLR Diagnostics: Studentized DELETED Residuals

```
studResidPlot(d_mod)
```



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SLR Diagnostics: Studentized DELETED Residuals

```
studResidPlot(d_mod, key.variable = "sampleDiamond",  
              print.obs = TRUE, sort.obs = TRUE)
```

##	sampleDiamond	price	carat	Predicted_Y	Student_Resid
## 1	47	1580	0.40	1254.4485	2.540426
## 2	97	1248	0.31	955.5587	2.215025
## 3	82	1208	0.30	922.3488	2.165628
## 4	71	631	0.30	922.3488	-2.210468
## 5	6	684	0.32	988.7687	-2.310115
## 6	124	673	0.32	988.7687	-2.397198
## 7	117	694	0.33	1021.9787	-2.494309
## 8	21	758	0.36	1121.6086	-2.796336
## 9	122	533	0.32	988.7687	-3.548077

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SLR Diagnostics: Cook' Distance

- Measure of "Influence": Leverage x "Outlyingness"

$$\text{Cook's Distance}_i = D_i = \frac{h_i}{1 - h_i} \times \frac{E_i'^2}{k + 1}$$

- Actually measures the change in estimates when i^{th} observation is removed.
- Rule of thumb:

$$D_i > 4/(n - k - 1)$$

$$D_i > 4/(n - p)$$

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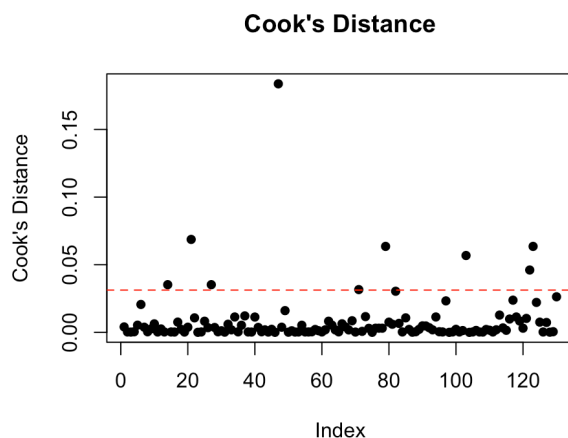
SLR Diagnostics: Cook' Distance

- Another good option for evaluating Cook's Distance: the F -distribution.
- Use (p) and $(n - p)$ degrees of freedom (that's $(k + 1)$ and $(n - k - 1)$).
- If D_i falls *NEAR* or *ABOVE* the 50th percentile, then it is *most likely* influential.
- If D_i falls *BELOW* the 50th percentile, but *ABOVE* the 20th percentile, then it *may be* influential.
- If D_i falls *BELOW* the 20th percentile, then it's *not* influential.
- But, really, a good visual examination takes care of it.

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SLR Diagnostics: Cook' Distance

```
cooksPlot(d_mod)
```



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SLR Diagnostics: Cook' Distance

```
cooksPlot(d_mod, key.variable = "sampleDiamond",  
          print.obs = TRUE, sort.obs = TRUE)
```

##	sampleDiamond	price	carat	Predicted_Y	Cooks_Distance	F_per
## 1	47	1580	0.40	1254.4485	0.18377167	0.16765506
## 2	21	758	0.36	1121.6086	0.06870973	0.06636795
## 3	79	1063	0.40	1254.4485	0.06355401	0.06154698
## 4	123	1063	0.40	1254.4485	0.06355401	0.06154698
## 5	103	576	0.25	756.2989	0.05681566	0.05520798
## 6	122	533	0.32	988.7687	0.04608736	0.04502562
## 7	14	1397	0.40	1254.4485	0.03523569	0.03461278
## 8	27	1397	0.40	1254.4485	0.03523569	0.03461278
## 9	71	631	0.30	922.3488	0.03160504	0.03110326

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SLR Diagnostics

- So which diagnostic do you choose?
- All three

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SLR Diagnostics

- So which diagnostic do you choose?
- All three

```
threeOuts(d_mod, key.variable = "sampleDiamond",)
```

##	sampleDiamond	Student_Resid	Hat_Values	Cooks_Distance	F_per
## 1	47	2.540426	0.05604876	0.18377167	0.16765506
## 2	21	-2.796336	NA	0.06870973	0.06636795
## 3	79	NA	0.05604876	0.06355401	0.06154698
## 4	123	NA	0.05604876	0.06355401	0.06154698
## 5	103	NA	0.05644717	0.05681566	0.05520798
## 6	122	-3.548077	NA	0.04608736	0.04502562
## 7	14	NA	0.05604876	0.03523569	0.03461278
## 8	27	NA	0.05604876	0.03523569	0.03461278
## 9	71	-2.210468	NA	0.03160504	0.03110326
## 10	6	-2.310115	NA	NA	NA

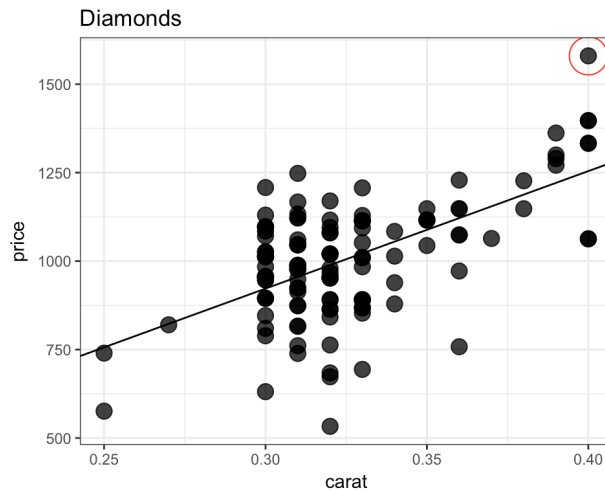
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"How much of an Influence?"

- When an observation has a high Cook's value (or is in general, an outlier), how much does it effect the model?
- Row 47 is our current "problem"
 - In all three measures of "outliers"
 - High Cook's value

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Observation 47



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Re-run without the observation:

- Remove the problem observation(s):

```
diamonds_noout <- diamonds %>%  
  filter(sampleDiamond != 47)
```

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Re-run without the observation:

- Re-run the model

```
d_mod_noout <- lm(price ~ carat, data=diamonds_noout)
d_mod_noout
```

```
##
## Call:
## lm(formula = price ~ carat, data = diamonds_noout)
##
## Coefficients:
## (Intercept)      carat
##      -4.152    3098.177
```

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Re-run without the observation:

- With the "outlier":

```
##
## Call:
## lm(formula = price ~ carat, data = diamonds)
##
## Coefficients:
## (Intercept)      carat
##      -73.95    3321.00
```

- Without the "outlier":

```
##
## Call:
## lm(formula = price ~ carat, data = diamonds_noout)
##
## Coefficients:
## (Intercept)      carat
##      -4.152    3098.177
```

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Re-run without the observation:

- What about σ_e ?

- With outlier:

```
summary(d_mod)$sigma
```

```
## [1] 134.6787
```

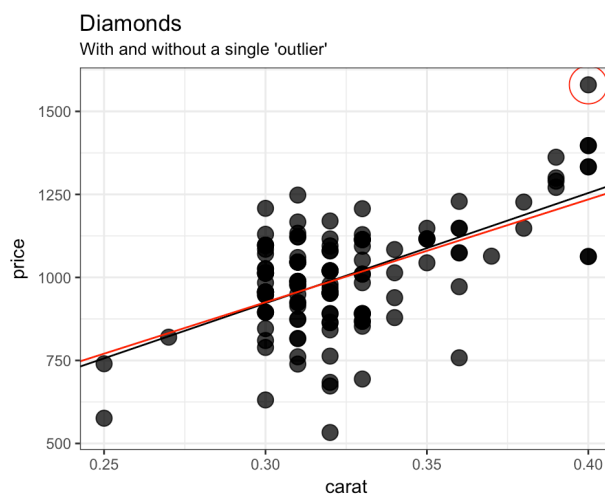
- Without outlier:

```
summary(d_mod_noout)$sigma
```

```
## [1] 131.8981
```

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Re-run without the observation:



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Summary

- Simple Linear Regression
- Interpretation and tie in to Pearson r
- Residuals and their usefulness
- Diagnostics Tools
- Types and Impact of Outliers
- Dealing Outliers with and Reporting