Influential Cases

• Could one or two data points throw off entire regression (leading to very different coefficients)?

- Would deleting a certain case result in . . .
 - Very different regression coefficients?
 - A different line of best fit?

Leverage vs. Influence

- **Leverage:** amount of *potential* each data point has to change the regression
- **Influence:** amount by which each data point *actually* changes the regression

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Leverage

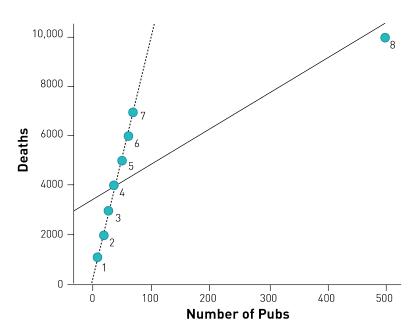
- A.k.a. "hat value"
- Written *h_{ii}*
- = a measure of how sensitive the regression is to a given data point i
 - Imagine moving a data point up and down on the Yaxis and observing how much the regression line follows it.
 - h_{ii} = ratio of how much the regression line moves compared to how much you move the data point
 - Range = 0–1; 0 = no influence, 1 = complete influence
- A data point with high leverage does not necessarily influence the regression; i.e., that point could be in line with the rest of the data.

Influence

- A data point needs a large residual to substantially change the regression coefficients.
- This means it lies far from the trend of the rest of the data.
- May indicate high influence
- Cook's distance measures the actual effect of deleting a given observation.
 - Each point with a large Cook's distance should be examined and potentially eliminated from the analysis.
 - General rule: Cook's distances >1 may be problematic (too much influence).

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Residuals vs. Influence



Example from Field et al. (2012). Discovering Statistics Using R

- Example: Eight different regions (boroughs) of London examined: number of deaths, number of pubs
 - The last case (Case 8) changes the line of best fit.
 - Interestingly, the residual for Case 8 is not large: the point is near the line.
 - Measures of influence (e.g., Cook's distance, hat value) indicate Case 8 has enormous influence.
 - Some cases can exert huge influence and still produce small residuals; thus it's important to look at both.

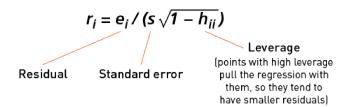
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Types of Residuals

- Unstandardized residuals
 - Measured in same units as outcome variable
 - Single model
 - Do not indicate which residual is "too large"
- Standardized residuals
 - Comparable across different regression models
 - Used to identify outliers

Standardized Residuals: Rough Method

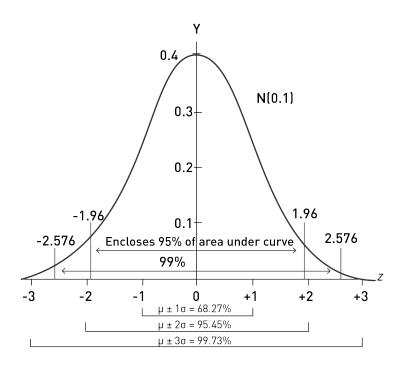
Standardized Residuals



- Take the normal residuals and divide by their standard error.
- Results look a lot like z scores (distribution close to a normal curve).
 - But not exactly a normal curve
- We standardize residuals in order to compare them across different models.
- Also enables us to use a consistent rule for large residuals
 - Analogy: z scores at tail end of a z distribution, with critical values >2
 - Outliers are significant and merit attention.

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Dealing With Standardized Residuals



- Standardized residuals give us a rough estimate of what scores are outliers.
- Values >3 are very large, not likely due to chance.
- If 1% (or more) of cases have residuals >2.5, model contains too much error.
- If 5% (or more) of cases have residuals >2, model has too much error *and* represents our data poorly.

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Studentized Residuals: Precise Method

Studentized Residuals

$$t_i = e_i / (s_{-i} \sqrt{1 - h_i})$$
New standard error

- Studentized residual looks similar to standardized residual.
- Difference: Before computing standard error, we remove one data point.
- Doing this makes numerator and denominator independent.
- Studentized residual now follows a Student's *t* distribution.
 - This lets us apply precise tests to identify significant outliers.

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