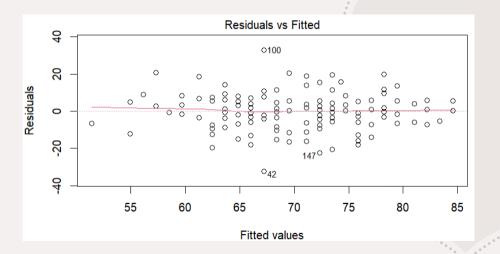


# Testing Model Assumptions

- Inspecting a residuals plots is a great way of seeing how the residuals of one's model are distributed and if there is a systematic pattern that appears to violate one of the model assumptions:
  - Correct form
  - Normally distributed residuals
  - Homogeneity of variances
  - Independence
- However, when examining these residuals plots for the first time, it can be difficult to tell what is a "big enough" diversion from one of these assumptions that it warrants being addressed
  - Especially with regards to the normality and homogeneity of variances assumptions as these are robust to some degree of violation.

# Testing Model Assumptions

- In addition to a visual examination of the residuals, it can be helpful to also run a test of each of these assumptions that provides a test statistic and corresponding p-value.
- For tests of model assumptions, it is desirable for the test to produce a **non-significant** *p*-value as a significant *p*-value typically means that the residuals significantly diverge from the model assumption.



## Testing the Normality Assumption using the Shapiro-Wilk Test

- Use the `shapiro.test()` function from the `stats` package
- Pass the function your model's standardized residuals
- Example:
  - ➤ shapiro.test(rstandard(model))
- A significant *p*-value suggests the model's residuals *significantly diverge* from being normally distributed (aka, the normality assumption is violated, the residuals are nonnormally distributed)

### Testing the Homogeneity of Variances Assumption using the Breusch-Pagan Test

- Use the `ncvTest()` function from the `car` package
- Pass the function your model
- Example:
  - > ncvTest(model)
- A significant *p*-value suggests the model's residuals *significantly diverge* from being homoscedastic (aka, the homogeneity of variances assumption is violated, heteroskedasticity is present)

## Testing the Independence Assumption using the Durbin-Watson Test

- Use the `durbinWatsonTest()` function from the `car` package
- Pass the function your model
- Example:
  - durbinWatsonTest(model)
- A significant *p*-value suggests the model's residuals *significantly diverge* from being independent (aka, the independence assumption is violated, the residuals are autocorrelated)