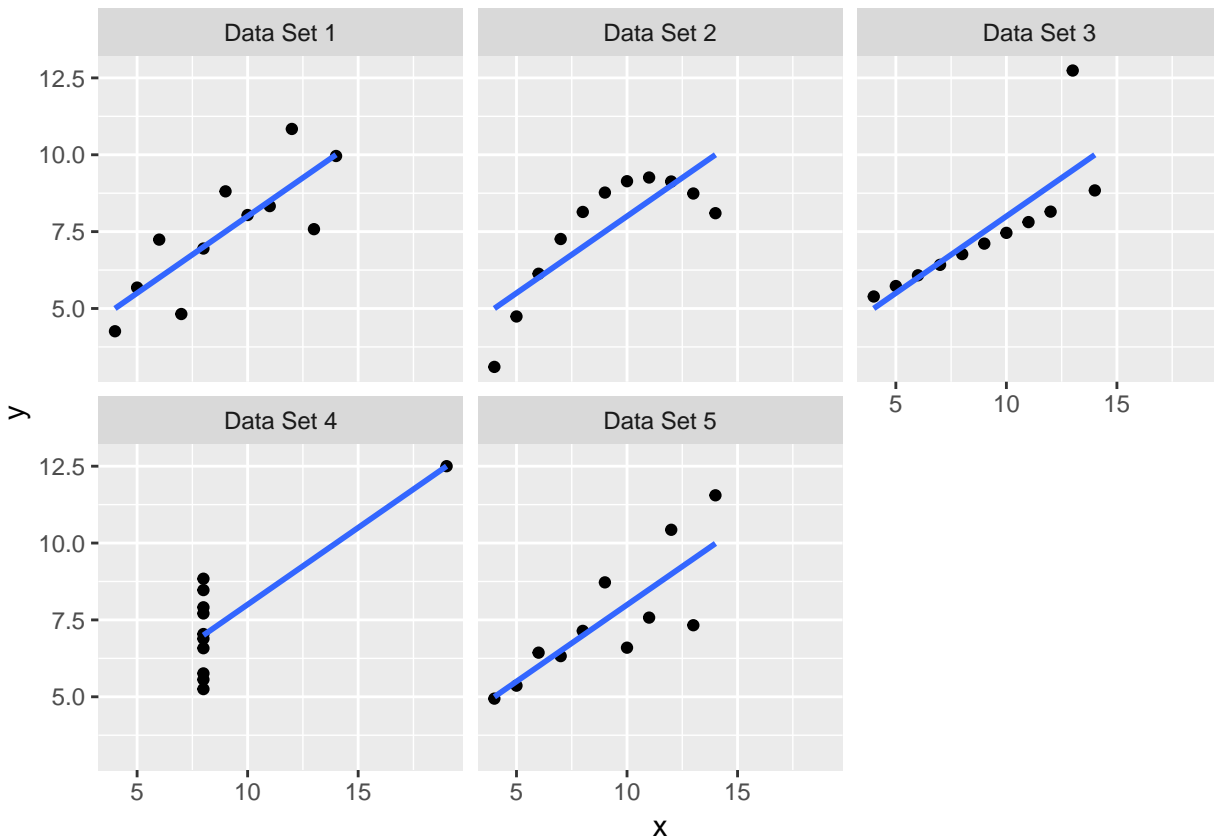


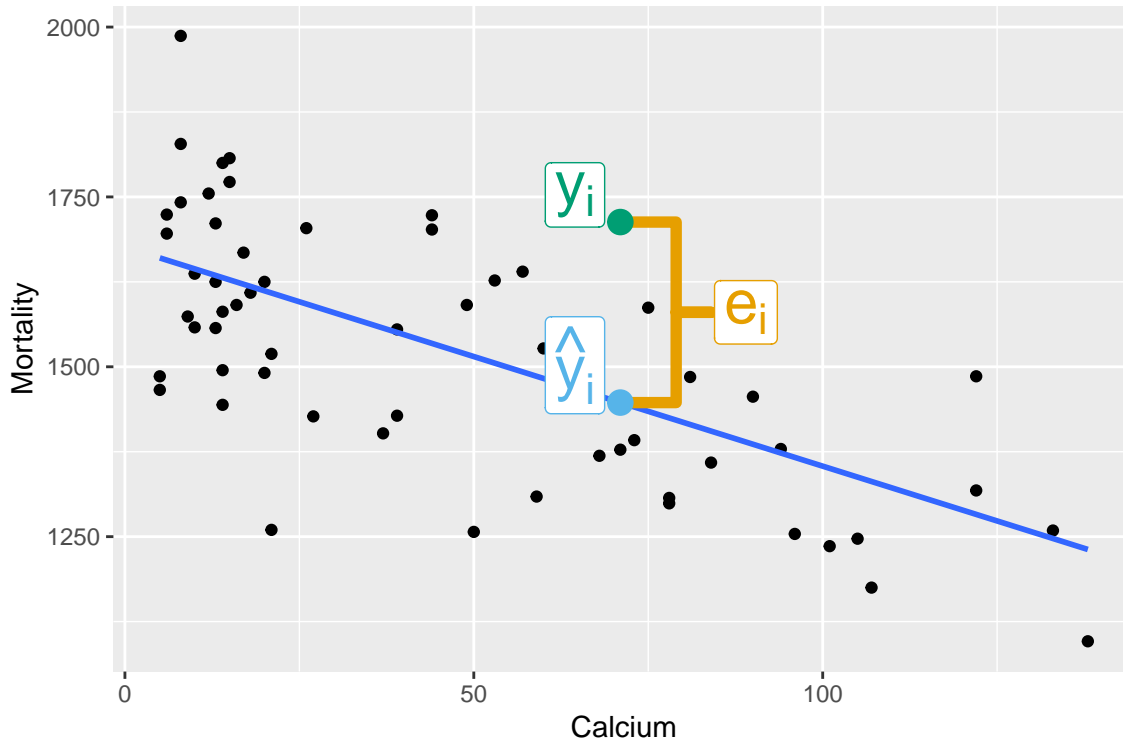
Linear Regression: Conditions for Inference, Residual Diagnostics

All 5 Have Essentially the Same Estimated Intercept, Slope, R^2 , and Residual Standard Deviation!



- Briefly, **conditions for linear regression** (see last page for more detail):
 - Sample **representative** of population
 - No **outliers** (points that don't fit the trend)
 - **Linear** relationship
 - **Independent** observations
 - **Normally** distributed residuals (or large enough sample size)
 - **Equal variability** of residuals
- Use **plots** to help diagnose the appropriateness of a linear model:
 - Scatter plot of explanatory (x axis) vs. response (y axis)
 - Scatter plot of predicted (x axis) vs. residual (y axis)
 - Histogram or density plot of residuals (x axis)
- Checks of whether the sample is representative and whether the observations are independent come from thinking about data collection process, not plots.

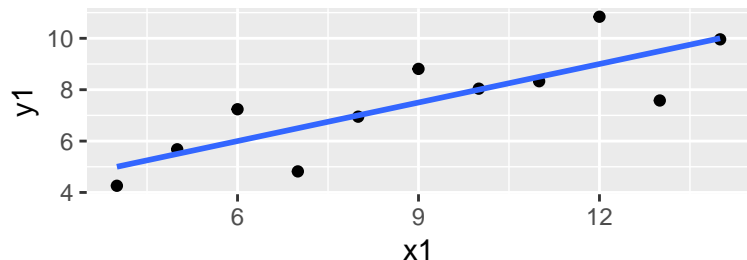
A Reminder about Residuals



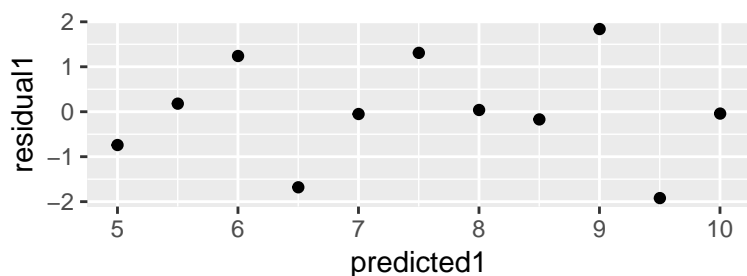
- Residuals give the vertical distance between a data point and the line of best fit
- Positive if point above line, negative otherwise
- $\text{Residual} = \text{Observed} - \text{Predicted}$
- $e_i = y_i - \hat{y}_i$ (e stands for error)

Anscombe Quintet: Data Set 1 (All Is Well)

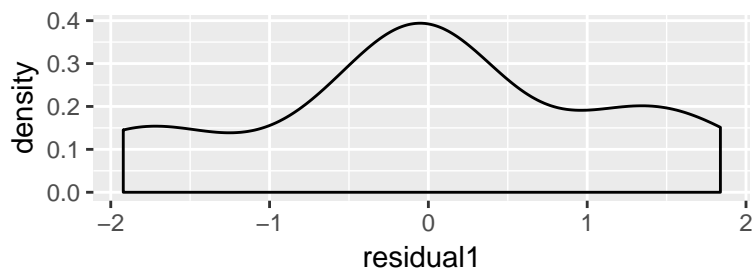
```
ggplot(data = anscombe, mapping = aes(x = x1, y = y1)) +  
  geom_point() +  
  geom_smooth(method = "lm", se = FALSE)
```



```
linear_fit1 <- lm(y1 ~ x1, data = anscombe)  
anscombe <- anscombe %>% mutate(  
  predicted1 = predict(linear_fit1),  
  residual1 = residuals(linear_fit1)  
)  
ggplot(data = anscombe, mapping = aes(x = predicted1, y = residual1)) +  
  geom_point()
```



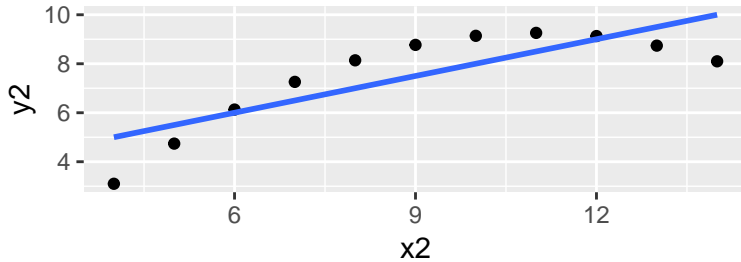
```
ggplot(data = anscombe, mapping = aes(x = residual1)) +  
  geom_density()
```



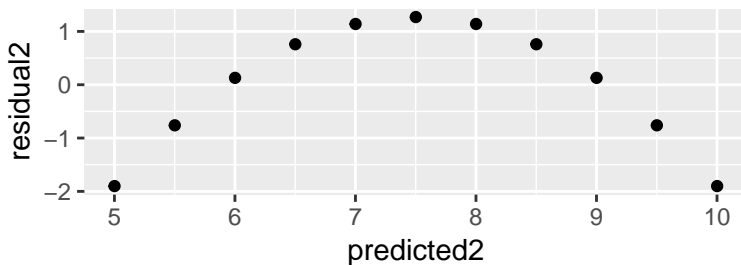
- **Outliers?** No
- **Linear** relationship? Yes
- **Normally** distributed residuals? Good enough
- **Equal variability** of residuals? Yes

Anscombe Quintet: Data Set 2 (Nonlinear)

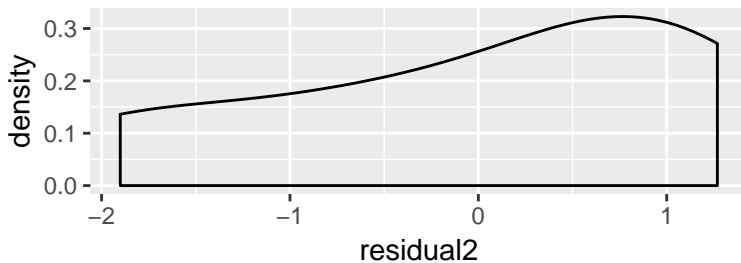
```
ggplot(data = anscombe, mapping = aes(x = x2, y = y2)) +  
  geom_point() +  
  geom_smooth(method = "lm", se = FALSE)
```



```
linear_fit2 <- lm(y2 ~ x2, data = anscombe)  
anscombe <- anscombe %>% mutate(  
  predicted2 = predict(linear_fit2),  
  residual2 = residuals(linear_fit2)  
)  
ggplot(data = anscombe, mapping = aes(x = predicted2, y = residual2)) +  
  geom_point()
```



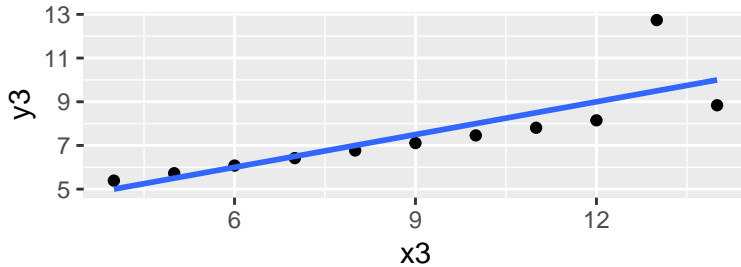
```
ggplot(data = anscombe, mapping = aes(x = residual2)) +  
  geom_density()
```



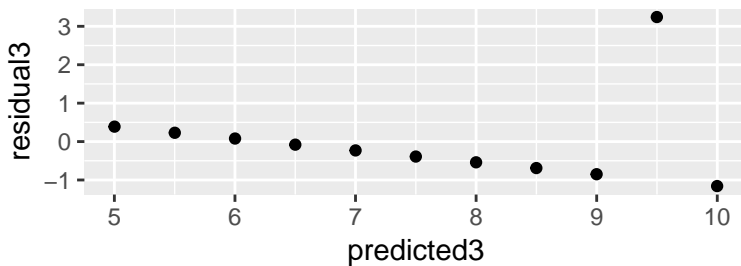
- **Outliers?** No
- **Linear relationship?** No (**this is a problem!!**)
- **Normally distributed residuals?** No perfect, probably good enough
- **Equal variability of residuals?** Yes

Anscombe Quintet: Data Set 3 (Outlier)

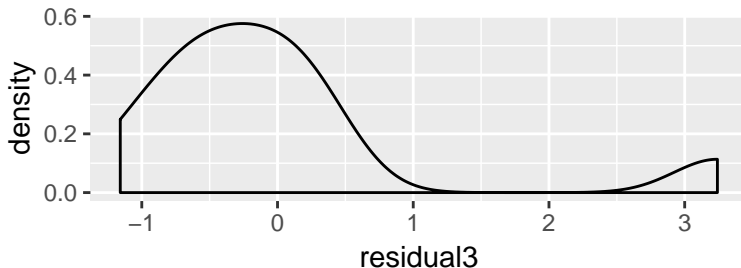
```
ggplot(data = anscombe, mapping = aes(x = x3, y = y3)) +  
  geom_point() +  
  geom_smooth(method = "lm", se = FALSE)
```



```
linear_fit3 <- lm(y3 ~ x3, data = anscombe)  
anscombe <- anscombe %>% mutate(  
  predicted3 = predict(linear_fit3),  
  residual3 = residuals(linear_fit3)  
)  
ggplot(data = anscombe, mapping = aes(x = predicted3, y = residual3)) +  
  geom_point()
```



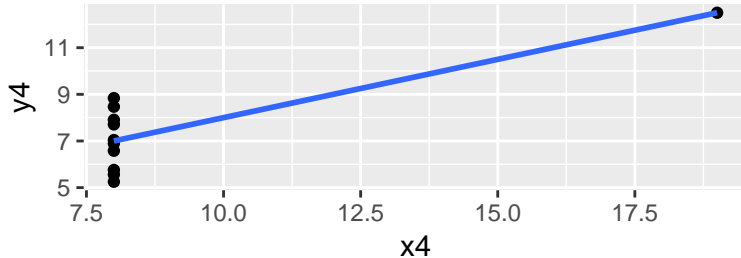
```
ggplot(data = anscombe, mapping = aes(x = residual3)) +  
  geom_density()
```



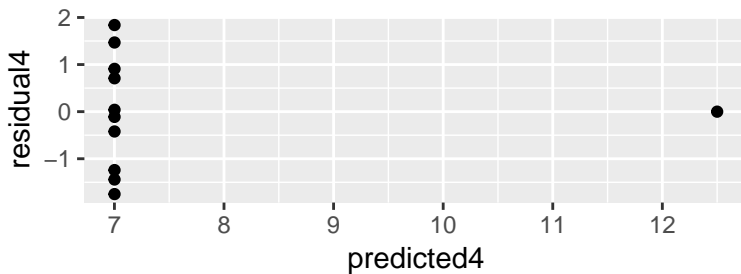
- **Outliers?** Yes (this is a problem!!)
- **Linear relationship?** Yes (other than the outlier)
- **Normally distributed residuals?** No, there is an outlier
- **Equal variability of residuals?** Yes (other than the outlier)

Anscombe Quintet: Data Set 4 (Outlier)

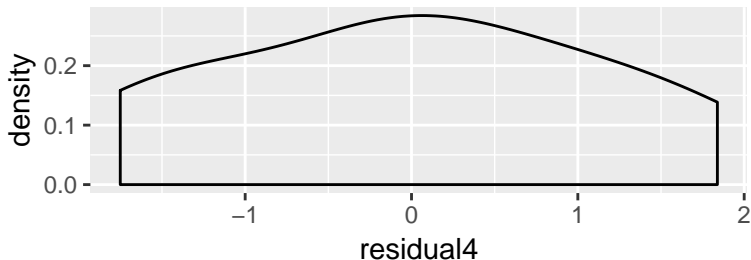
```
ggplot(data = anscombe, mapping = aes(x = x4, y = y4)) +  
  geom_point() +  
  geom_smooth(method = "lm", se = FALSE)
```



```
linear_fit4 <- lm(y4 ~ x4, data = anscombe)  
anscombe <- anscombe %>% mutate(  
  predicted4 = predict(linear_fit4),  
  residual4 = residuals(linear_fit4)  
)  
ggplot(data = anscombe, mapping = aes(x = predicted4, y = residual4)) +  
  geom_point()
```



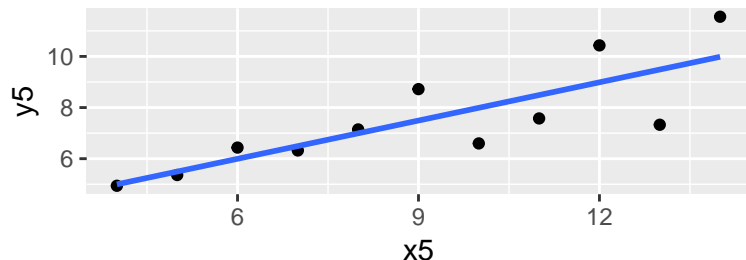
```
ggplot(data = anscombe, mapping = aes(x = residual4)) +  
  geom_density()
```



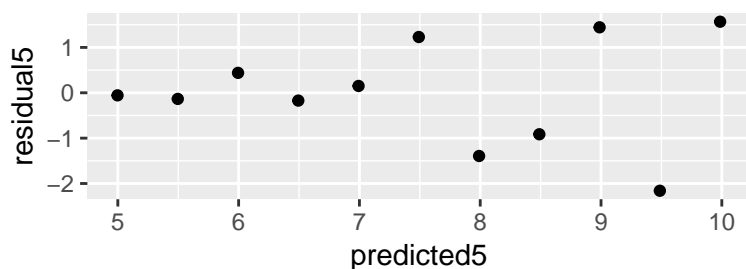
- **Outliers?** Yes (**this is a problem!!**)
- **Linear** relationship? Difficult to assess
- **Normally** distributed residuals? Yes
- **Equal variability** of residuals? Difficult to assess

Anscombe Quintet: Data Set 5 (Lack of Equal Variability of Residuals)

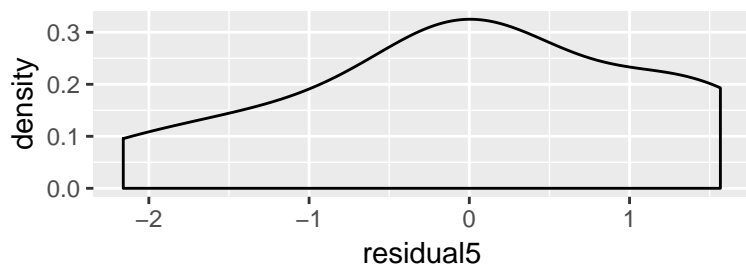
```
ggplot(data = anscombe, mapping = aes(x = x5, y = y5)) +  
  geom_point() +  
  geom_smooth(method = "lm", se = FALSE)
```



```
linear_fit5 <- lm(y5 ~ x5, data = anscombe)  
anscombe <- anscombe %>% mutate(  
  predicted5 = predict(linear_fit5),  
  residual5 = residuals(linear_fit5)  
)  
ggplot(data = anscombe, mapping = aes(x = predicted5, y = residual5)) +  
  geom_point()
```



```
ggplot(data = anscombe, mapping = aes(x = residual5)) + geom_density()
```



- **Outliers?** No
- **Linear relationship?** Yes
- **Normally distributed residuals?** Yes
- **Equal variability of residuals?** No (**this is a problem!!**)

Regression Conditions

Think of a helpful leprechaun named **Robert O'Line**:



- Sample **representative** of population
- No **outliers** (points that don't fit the trend)
- **Linear** relationship
- **Independent** observations
- **Normally** distributed residuals (or large enough sample size)
- **Equal variability** of residuals

Condition	How Important?	How to Check?
R epresentative	Critical	Think about data collection (randomization?)
No O utliers	Very Important	<ul style="list-style-type: none"> • Scatter Plot of explanatory variable vs response variable • Scatter plot of predicted value vs residuals • histogram or density plot of residuals
L inear relation-ship	Very Important	<ul style="list-style-type: none"> • Scatter Plot of explanatory variable vs response variable (pattern is linear) • Scatter plot of predicted value vs residuals (no curved patterns)
I ndependent observations	Very Important	<p>Think about data collection (randomization?) Situations where observations are not independent:</p> <ul style="list-style-type: none"> • Observations collected over time (e.g., monthly unemployment measurements over time) • Multiple observations on the same person (e.g., baseline and follow-up measurements of health in a clinical trial)
N ormally distributed residuals	Somewhat Important	<ul style="list-style-type: none"> • histogram or density plot of residuals (unimodal, approximately symmetric, no outliers) • ...or large enough sample size
E qual variability of residuals	Somewhat Important	<ul style="list-style-type: none"> • Scatter Plot of explanatory variable vs response variable (same amount of vertical spread around line for all values of x) • Scatter plot of predicted value vs residuals (same amount of vertical spread for all values of x)