

CogLab: Making Inferences

WEEK 9

formative assignment #2

- descriptive statistics and plotting in R
 - due Nov 4

9	Tuesday, October 29, 2024	W9: Project Work
9	Thursday, October 31, 2024	W9 continued...
9	Monday, November 4, 2024	Formative Assignment (R Descriptive) Due
10	Tuesday, November 5, 2024	Weeks 10-12: Data Collection
10	Thursday, November 7, 2024	Weeks 10-12: Data Collection
10	Monday, November 11, 2024	Project Milestone #5 (Pre-Registration + Checklist) Due
11	Tuesday, November 12, 2024	Weeks 10-12: Data Collection
11	Thursday, November 14, 2024	Weeks 10-12: Data Collection
11	Sunday, November 17, 2024	Formative Assignment (R Descriptive) Resubmission Due
11	Monday, November 18, 2024	Formative Assignment (R Inferential) Due
12	Tuesday, November 19, 2024	Weeks 10-12: Data Collection
12	Thursday, November 21, 2024	Psychonomics Conference: NO CLASS
12	Friday, November 22, 2024	Project Milestone #6 (Analyses: Deadline 1) Due
13	Tuesday, November 26, 2024	THANKSGIVING BREAK!!! NO CLASS
13	Thursday, November 28, 2024	THANKSGIVING BREAK!!! NO CLASS
14	Monday, December 2, 2024	Project Milestone #6 (Analyses: Deadline 2) Due

project checklist

- project checklist

	Pilot 1	Pilot 2	Pilot 3
Which browser were you using?			
Which operating system (Mac / Windows / iPad, etc.)			
Date of piloting			
Were instructions clear? Please note down which instructions had typos / were unclear			
How long did the task take you?			
Was there a consent form?			
Was the demographic survey displayed correctly?			
Did you see the data being displayed at the end of the study?			
What do you think the experiment was about?			
Any other comments?			

CogLab Project Checklist

Task	Check if done
Sanity Check <ul style="list-style-type: none"> <input type="checkbox"/> Is the attention check response being recorded? <input type="checkbox"/> Is the free association response being recorded? <input type="checkbox"/> Can you differentiate between training / attention / association / prime / target? <input type="checkbox"/> Can you differentiate between prime and target trials? <input type="checkbox"/> Can you differentiate practice and test trials? <input type="checkbox"/> Is subject ID being recorded? <input type="checkbox"/> Is RT being recorded? 	<input type="checkbox"/>
For the demographic survey , how are you showing these questions? Are there multiple answers people can pick or is it a binary choice? Are people able to select multiple answers when they should not be?	<input type="checkbox"/>
For the demographic survey , what questions are being shown on the same screen? What questions should be on different screens?	<input type="checkbox"/>
For the demographic survey , how are the data being recorded and is it being recorded? Also, do you have all the questions you need?	<input type="checkbox"/>
Before Pre-Registration: <ul style="list-style-type: none"> <input type="checkbox"/> Are you providing accuracy feedback on priming practice trials? <input type="checkbox"/> Have you addressed ALL the feedback from Milestone 4? <ul style="list-style-type: none"> <input type="checkbox"/> Feedback 1 <input type="checkbox"/> Feedback 2 <input type="checkbox"/> Feedback 3 <input type="checkbox"/> Are you recording IP addresses? <input type="checkbox"/> Are you commenting the condition definition inside cognition.run <input type="checkbox"/> Have you piloted your experiment with Uma + other group + 5 friends) <input type="checkbox"/> Have they completed the pilot feedback sheet? <input type="checkbox"/> Have you sent the cognition.run link by Nov 10? <input type="checkbox"/> Have you finalized the analysis plan + sample size? <input type="checkbox"/> Have you created and submitted a pre-registration draft? 	<input type="checkbox"/>
Analysis <ul style="list-style-type: none"> <input type="checkbox"/> Did you confirm/correct all datatypes? <input type="checkbox"/> Did you figure out how to "filter" certain types of trials? <input type="checkbox"/> Did you fix all typos in attention responses? <input type="checkbox"/> Have you computed mean attention accuracy? <input type="checkbox"/> Have you applied exclusions based on accuracy AND RTs? <input type="checkbox"/> Have you created an RT bar graph? <input type="checkbox"/> Have you fit a statistical model? 	<input type="checkbox"/>

pre-registration + project checklist

- milestone #5:

pre-registration + project
checklist + piloting
(Nov 10)

1. **Data Collection:** Have any data been collected for this study already?
2. **Main Question:** What is the main question being asked or hypothesis being tested in this study?
3. **Dependent Variable(s):** Describe the key dependent variable(s) specifying how they will be measured.
4. **Condition(s):** How many and which conditions will participants be assigned to? Please include an example trial of each type of condition you have in your experiment. Please also specify which independent variable will be within-participants or between-participants.
5. **Analyses:** Specify exactly which analyses you will conduct to examine the main question/hypothesis.
6. **Outliers & Exclusions:** Describe exactly how outliers will be defined and handled, and your precise rule(s) for excluding observations.
7. **Predicted Plot:** Please submit a predicted plot for your study based on what you expect the pattern to look like for your main hypothesis.
8. **Sample Size:** How many observations will be collected or what will determine sample size? No need to justify the decision, but be precise about exactly how the number will be determined.
9. **Exploratory details:** Anything else you would like to pre-register? (e.g., secondary analyses, variables collected for exploratory purposes, unusual analyses planned?)

recap

- what we covered:
 - manipulating data using tidyverse verbs
 - project work
- your to-do's were:
 - *work on*: formative assignment #2 (R descriptive)
 - *work on*: project checklist + pre-registration

today's agenda

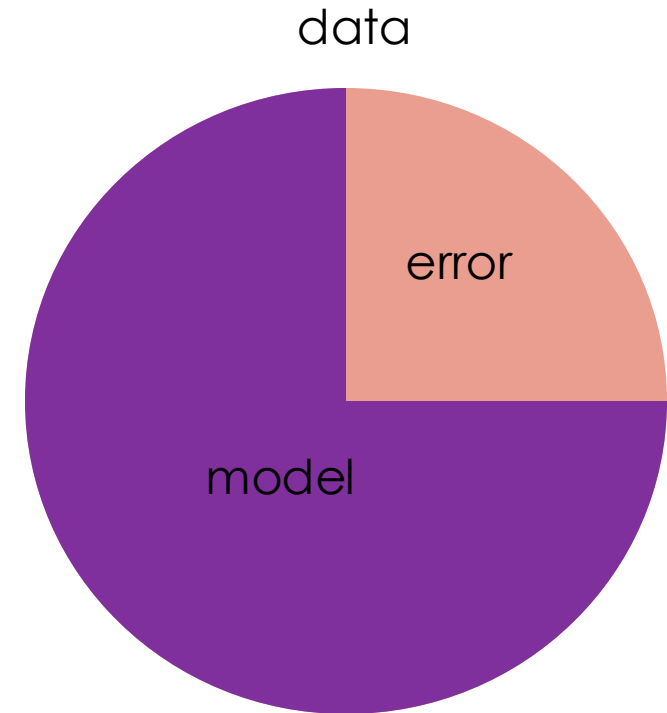
- making statistical inferences from data



what is the goal of statistics?

$$\text{data} = \text{model} + \text{error}$$

- the goal of statistics is to find a simple explanation to the observed data (Y, dependent variable), i.e., build a **model** of the data that approximates/explains it as well as possible –
- what is a **good** model? one that represents the data really well
- how do we start building models?



some simple models

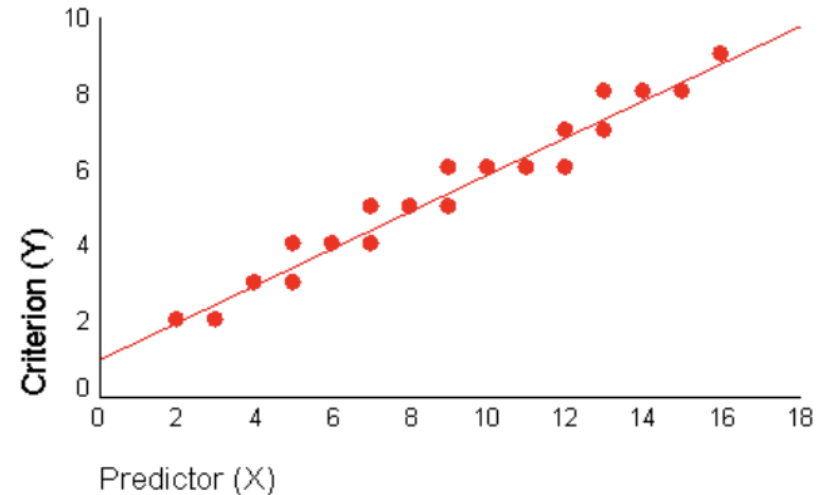
- **central tendencies** (mean/median/mode)
 - derived from the key dependent variable (data = Y) itself
 - no other variables needed for this
 - the mean is the best model if no other variables are available
- **measures of variation:** estimates of model fit
 - **sum of squared errors** (SSE or SS): $\sum_{i=1}^N (Y_i - \mu)^2$
 - **mean of squared errors** (MSE): $\frac{\sum_{i=1}^N (Y_i - \mu)^2}{N} = \frac{SS}{N}$
 - **root mean squared error** (RMSE): $\sqrt{\frac{\sum_{i=1}^N (Y_i - \mu)^2}{N}} = \sqrt{MSE}$

slightly more complex models...

- using one more variable (X) to “explain” the dependent variable/data (Y)
- how does knowing something about X impact what we know about Y ?
- what types of “models” are these?

linear regression

- a linear regression (or a linear model) is a model that fits a line to the data
- $Y = a + bX + \text{error}$
- slope: $b = r \frac{s_y}{s_x}$
- intercept: $a = M_y - bM_x$



exploring the data

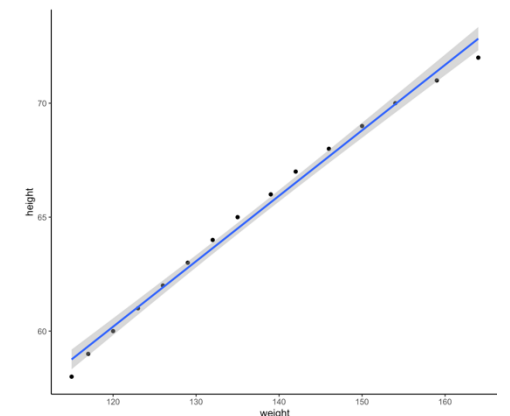
- open your project + Rmd
- new heading **# linear models**
- load the dataset **women**
- make a scatterplot of the data
 - x = weight
 - y = height
- fit a line to the data via `geom_smooth()`

```
# linear models
```

```
```${r}``  
data(women)
```

```
women %>%
 ggplot(aes(x= weight, y = height))+
 geom_point() +
 geom_smooth(method = "lm")+
 theme_classic()
```

	height	weight
1	58	115
2	59	117
3	60	120
4	61	123
5	62	126
6	63	129
7	64	132
8	65	135
9	66	139
10	67	142
11	68	146
12	69	150
13	70	154
14	71	159
15	72	164



# linear regression in R

- `predict` height by weight
- print the `summary` of the model
- what is the `equation` of the line?

```
women_model = lm(data = women, height ~ weight)
```

```
summary(women_model)
```

Call:

```
lm(formula = height ~ weight, data = women)
```

Residuals:

Min	1Q	Median	3Q	Max
-0.83233	-0.26249	0.08314	0.34353	0.49790

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	25.723456	1.043746	24.64	2.68e-12 ***
weight	0.287249	0.007588	37.85	1.09e-14 ***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

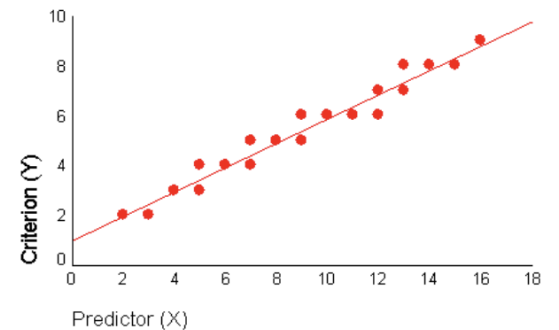
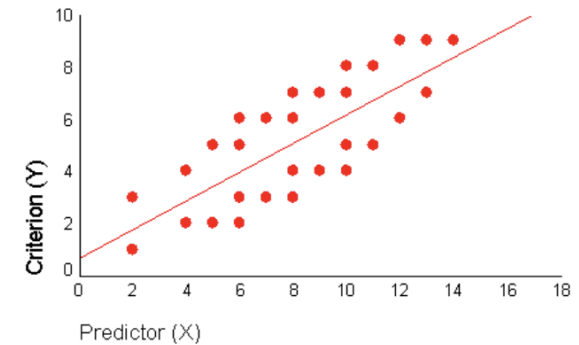
Residual standard error: 0.44 on 13 degrees of freedom

Multiple R-squared: 0.991, Adjusted R-squared: 0.9903

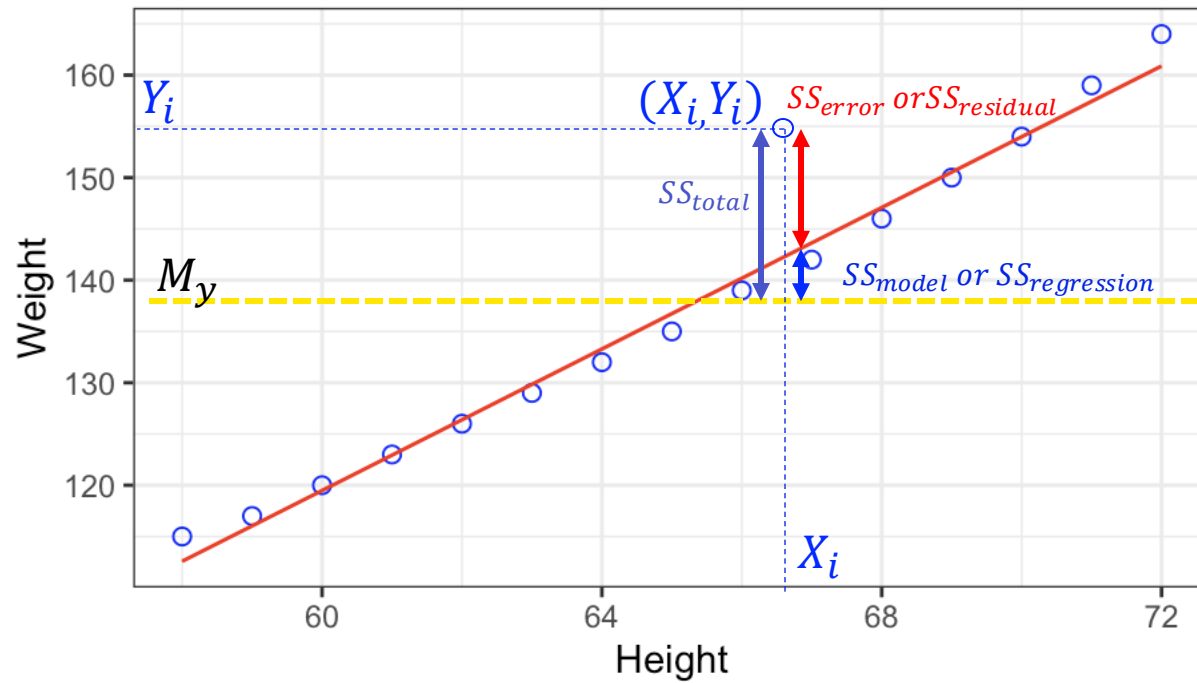
F-statistic: 1433 on 1 and 13 DF, p-value: 1.091e-14

# assessing model fit

- let's say we find a line of best fit
  - data = model + error
  - $Y = a + bX + \text{error}$
  - $\hat{Y} = a + bX = \text{predictions}$
  - $Y = \hat{Y} + \text{error}$
- how well does the line fit our data?
- $SS_{\text{error}} = \sum_{i=1}^n (y_i - a - bx_i)^2 = \sum (Y - \hat{Y})^2$



# understanding goodness/errors



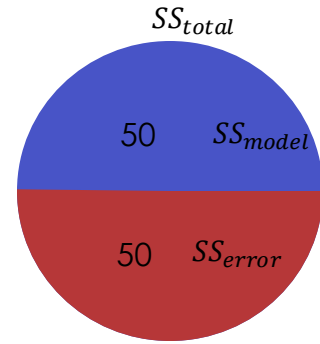
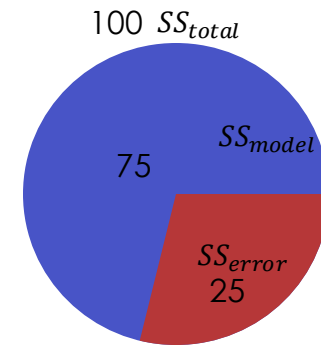
$$SS_{total} = SS_{model} + SS_{error}$$

$$SS_{total} = \sum (Y - M_y)^2$$

$$SS_{error} = \sum (Y - \hat{Y})^2$$

$$SS_{model} = \sum (\hat{Y} - M_y)^2$$

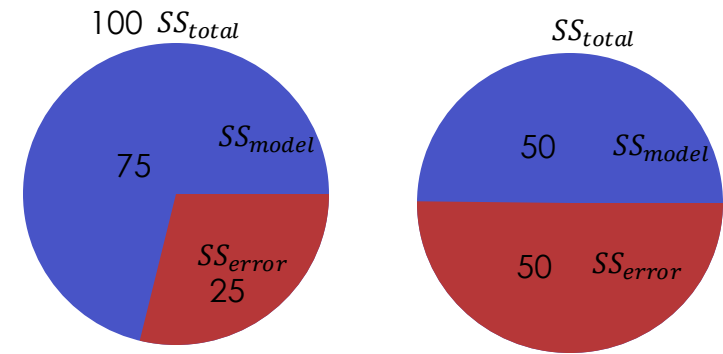
# overall test of model (ANOVA)



- **analysis of variance** assesses the overall fit of the model
- $SS_{total} = SS_{model} + SS_{error}$
- we calculate the ratio between the variance explained by the model and the natural variance expected/left over in the dependent variable
  - if  $\frac{SS_{model}}{SS_{error}}$  is high, the model explains **more** variance than expected
  - if  $\frac{SS_{model}}{SS_{error}}$  is low, the model explains **less** variance than expected
- typically, we want the “average” variance explained, so we also divide this by  $df$



# F ratio



- The F ratio compares the “average” squared error between **model (explained variance)** and the **natural (unexplained) variance (data = model + error)**

$$F = \frac{\text{explained variance}}{\text{unexplained variance}} = \frac{MS_{\text{model}}}{MS_{\text{error}}} = \frac{SS_{\text{model}}/df_{\text{model}}}{SS_{\text{error}}/df_{\text{error}}}$$

- obtaining  $SS_{\text{model}}$  and  $SS_{\text{error}}$ 
  - $SS_{\text{error}} = \sum(Y - \hat{Y})^2$  and  $SS_{\text{total}} = \sum(Y - M_y)^2$
  - $SS_{\text{model}} = SS_{\text{total}} - SS_{\text{error}}$
- obtaining  $df_{\text{model}}$  and  $df_{\text{error}}$ 
  - $k$  denotes the number of levels of the independent variable OR number of estimated parameters
  - $df_{\text{model}} = k - 1$
  - $df_{\text{error}} = n - k$

# ANOVA for women dataset

- install the `car` package
- use the `Anova()` function
- how do we report this F test?
- weight significantly predicted height,  $F(1,13) = 1433, p < .001$ .

```
car::Anova(women_model)
```

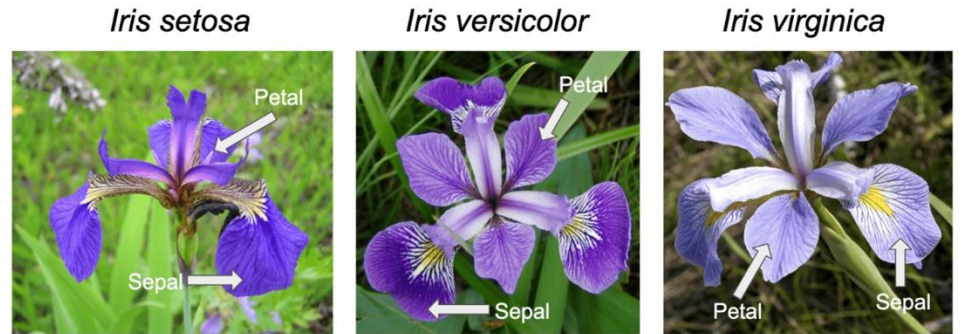
```
> car::Anova(women_model)
Anova Table (Type II tests)

Response: height
 Sum Sq Df F value Pr(>F)
weight 277.483 1 1433 1.091e-14 ***
Residuals 2.517 13

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

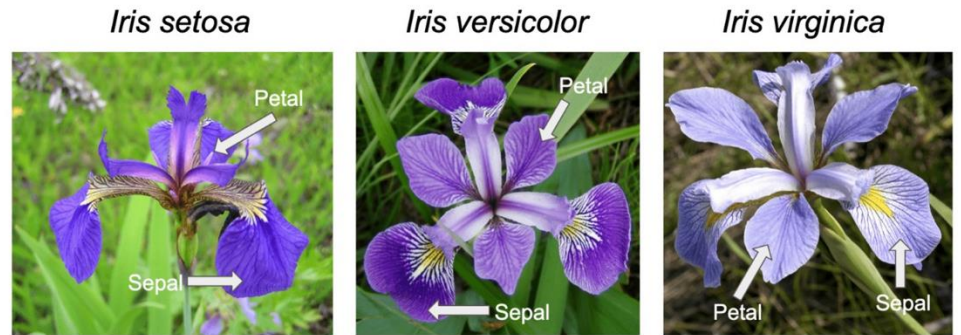
# ANOVAs for categorical IVs

- the same logic applies to problems where the independent variable is not continuous
- **research question**: what explains the variation in petal lengths (Y)?
  - data (Y) = model (X) + error
  - petal lengths (Y) = species (X) + error



# descriptive exercise

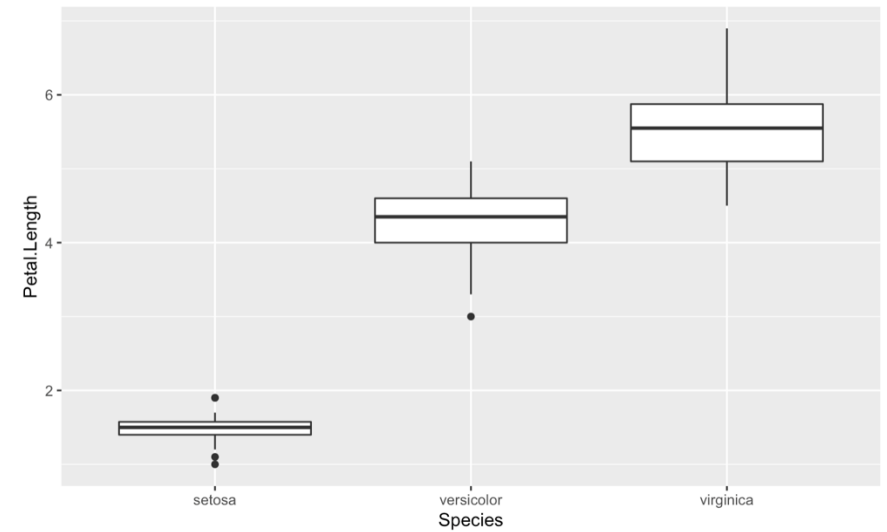
- obtain the mean petal length for each species in the `iris` dataset using `tidyverse` functions



```
A tibble: 3 × 2
 Species mean_length
 <fct> <dbl>
1 setosa 1.46
2 versicolor 4.26
3 virginica 5.55
```

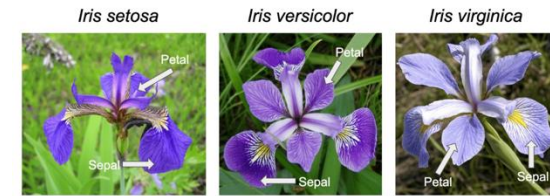
# plotting exercise

- make a **boxplot** of petal lengths by species



# ANOVA for iris

- load and view iris
- fit a model to petal lengths
- view `car::Anova()` results
- does species explain the variation in petal lengths?
- which species are different from each other?



```
anova
```

```
data(iris)
View(iris)
```

```
iris_model = lm(data=iris, Petal.Length ~ Species)
```

```
car::Anova(iris_model)
```

```
> car::Anova(iris_model)
```

```
Anova Table (Type II tests)
```

```
Response: Petal.Length
```

	Sum Sq	Df	F value	Pr(>F)
Species	437.10	2	1180.2	< 2.2e-16 ***
Residuals	27.22	147		

```

```

```
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

# follow-up tests

- when more than two groups are present, it can be useful to understand exactly **which groups differ** from each other
- install **emmeans** package
- load the package inline and compute pairwise differences

```
emmeans::emmeans(iris_model,
 pairwise ~ Species,
 adjust = "tukey")
```

```
$emmeans
```

Species	emmean	SE	df	lower.CL	upper.CL
setosa	1.46	0.0609	147	1.34	1.58
versicolor	4.26	0.0609	147	4.14	4.38
virginica	5.55	0.0609	147	5.43	5.67

Confidence level used: 0.95

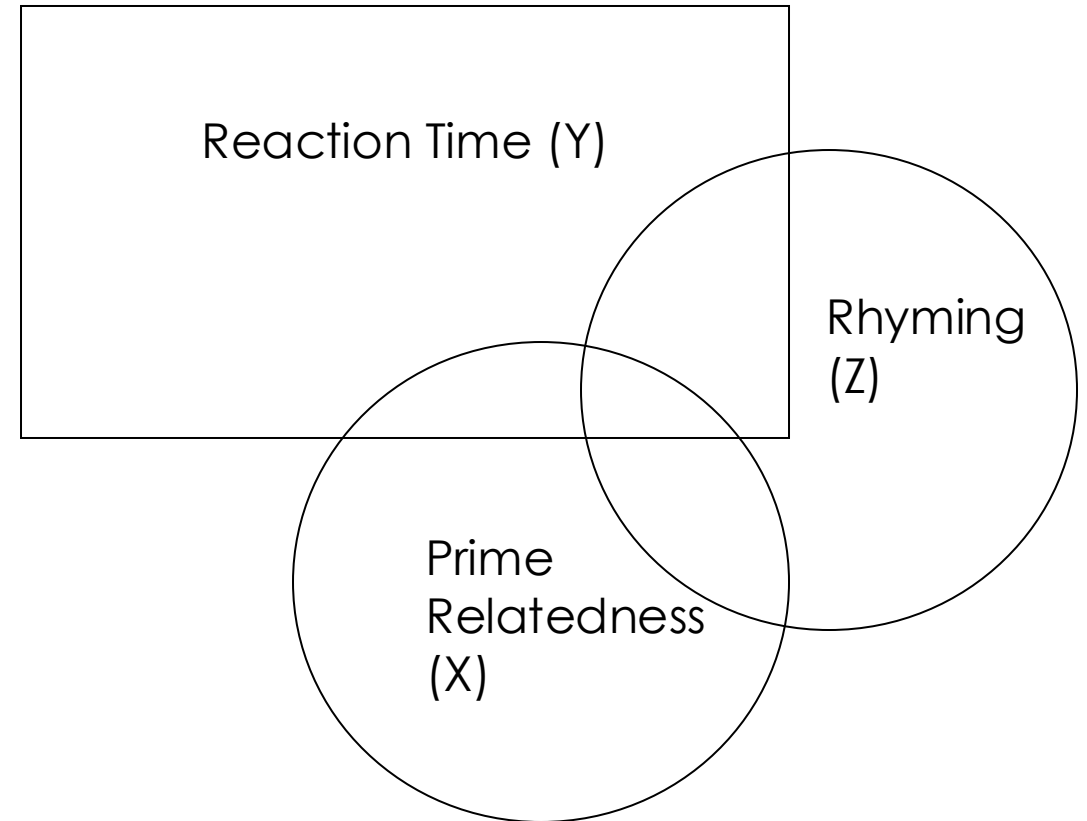
```
$contrasts
```

contrast	estimate	SE	df	t.ratio	p.value
setosa - versicolor	-2.80	0.0861	147	-32.510	<.0001
setosa - virginica	-4.09	0.0861	147	-47.521	<.0001
versicolor - virginica	-1.29	0.0861	147	-15.012	<.0001

P value adjustment: tukey method for comparing a family of 3 estimates

# even more complex models...

- what if the variation in our data (Y) could be explained further?
- **data = model + error**
  - *one IV*:  $Y = a + bX + \text{error}$
  - *multiple IVs*:  $Y = a + b_1X_1 + b_2X_2 + \dots + \text{error}$
- central idea remains the same, but more complex relationships are possible





# next class

- **before** class

- *apply*: formative assignment #2 (due Monday)
- *apply*: pre-registration + checklist (due Nov 10)

- **during** class

- complex models + project work