

Psychological Statistics Lab

3

PSYC 2020-A01 / PSYC 6022-A01 | 2025-09-04 | Descriptive Statistics II

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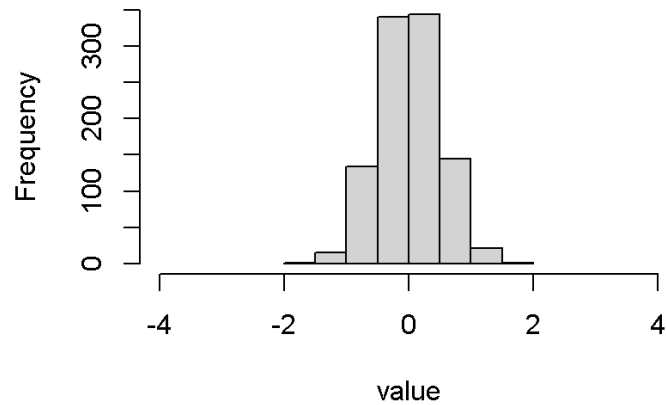
Outline

- Assignment 2 Review
- More on types of data
- Labeling data
- Aggregating data
- Descriptives by group
- Weighted mean

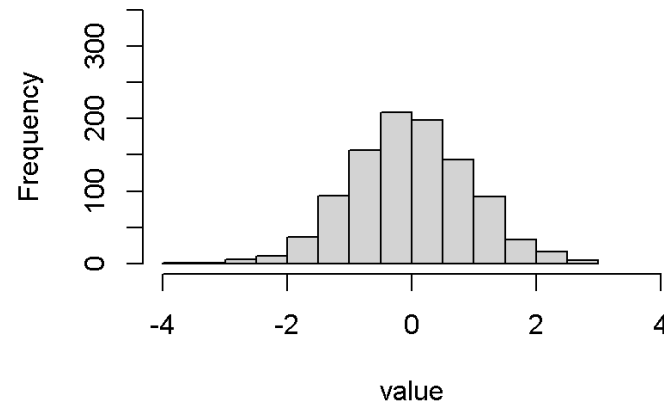
Measures of Variability Review

Describe the “spread” or “dispersion” of the data

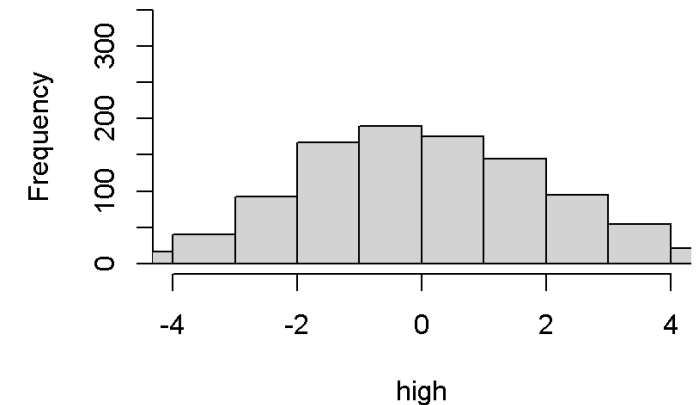
low variance



standard



high variance



Measures of Variability Functions

```
1 testscores <- c(88, 93, 92, 99, 96)
```

Variance: `var()` function

Standard Deviation: `sd()` function

Interquartile Range: `IQR()` function

- Difference between the 3rd and 1st quantile (so 50% of the data within this range)
- 25% of the data lower than the 1st quantile
- 75% of the data lower than the 3rd quantile
- $IQR = 3rd - 1st$

R Code [↺ Start Over](#)

[Run Code](#)

```
1 # let's look at some variances!
```

Quantiles: `quantile()` function

Object Types: Vector

We've already seen and used vectors

```
1 c(2, 3, 4, 2)
```

```
[1] 2 3 4 2
```

```
1 c("Cat", "Dog", "Bird")
```

```
[1] "Cat" "Dog" "Bird"
```

We know they can only hold one type of data

```
1 c("Cat", "Dog", 4)
```

```
[1] "Cat" "Dog" "4"
```

Shorthand for sequenced numbers:

```
1 1:10
```

```
[1] 1 2 3 4 5 6 7 8 9 10
```

Object Types: Matrix

Like a vector, but multidimensional

Can only hold one type of data

- Worse than a dataframe?
- Okay! But pop up in function output often
- So we need to know how to work with them

```
1 matrix(c(1, 2, 3, 4), nrow = 2)
```

```
      [,1] [,2]  
[1,]    1    3  
[2,]    2    4
```

Default is to fill in columns first

```
1 matrix(c(1, 2, 3, 4), nrow = 2, byrow = T)
```

```
      [,1] [,2]  
[1,]    1    2  
[2,]    3    4
```

Object Types: Matrix

Can give names

colnames() Function

```
1 mymatrix <- matrix(c(0, 1, 2, 1, 2, 3, 2, 3,
2                       nrow = 3, byrow = T)
3 colnames(mymatrix) <- c("Var1", "Var2", "Var3")
4 mymatrix
```

	Var1	Var2	Var3
[1,]	0	1	2
[2,]	1	2	3
[3,]	2	3	4

Retrieves object column names,
then assigns them something else

dimnames Argument

```
1 mymatrix <- matrix(c(0, 1, 2, 1, 2, 3, 2, 3,
2                       nrow = 3, byrow = T,
3                       dimnames = list(NULL, c("Var1", "Var2", "Var3")))
4 mymatrix
```

	Var1	Var2	Var3
[1,]	0	1	2
[2,]	1	2	3
[3,]	2	3	4

Sets names while creating the
matrix Takes a `list(row_names,
column_names)`

Object Types: List

Limitations of vectors:

- Can only hold one type of data

Enter the `list`

Can create by calling the `list()` function and giving it some objects

```
1 list(1, 2, "cat")
```

```
[[1]]  
[1] 1  
  
[[2]]  
[1] 2  
  
[[3]]  
[1] "cat"
```

```
1 list(1:4, c("a", "b"))
```

```
[[1]]  
[1] 1 2 3 4  
  
[[2]]  
[1] "a" "b"
```


Object Types: Dataframes

More review

Like matrices, but can hold different types of data

```
1 matrix(c("cat", 2, "dog", 2),  
2       nrow = 2, byrow = T,  
3       dimnames = list(NULL, c("Pet", "Count"))
```

	Pet	Count
[1,]	"cat"	"2"
[2,]	"dog"	"2"

```
1 mydataframe <- data.frame(  
2   pet = c("cat", "dog"),  
3   count = c(2, 2))  
4 mydataframe
```

	pet	count
1	cat	2
2	dog	2

Indexing

To access some subset of an object (vector, list, dataframe, etc.)

We already know one! The `$` operator indexes by selecting a column

```
1 mydataframe$pet
```

```
[1] "cat" "dog"
```

Indexing

To access some subset of an object (vector, list, dataframe, etc.)

Can also use brackets `[]`

```
1 myvector <- c(2, 4, 6, 8)
2
3 myvector[1]
```

```
[1] 2
```

Can select more than one position at a time

```
1 myvector[c(1, 2, 3)]
```

```
[1] 2 4 6
```

```
1 myvector[1:3]
```

```
[1] 2 4 6
```

Indexing

Need to account for multiple dimensions

Matrix: `[element position]` or `[row, column]`

```
1 mymatrix
```

	Var1	Var2	Var3
[1,]	0	1	2
[2,]	1	2	3
[3,]	2	3	4

```
1 mymatrix[1]
```

```
[1] 0
```

```
1 mymatrix[4]
```

```
[1] 1
```

```
1 mymatrix[2, ]
```

Var1	Var2	Var3
1	2	3

```
1 mymatrix[3, 2]
```

```
Var2
```

Indexing

Dataframe: `[column]` or `[row, column]`

```
1 mydataframe[1]
```

```
pet  
1 cat  
2 dog
```

```
1 mydataframe["pet"]
```

```
pet  
1 cat  
2 dog
```

```
1 mydataframe$pet
```

```
[1] "cat" "dog"
```

```
1 mydataframe[2, "pet"]
```

```
[1] "dog"
```

Can also do more than one element at a time:

```
1 mydataframe[1:2, "count"]
```

```
[1] 2 2
```

Labeling Data

Adding meaningful labels or categories to your dataset.

Useful for organizing and interpreting your data.

- E.g., Adding labels like “Pass” or “Fail” based on test scores.

	id	order	rating
1	1	1	3
2	2	2	3
3	3	3	4
4	4	1	3
5	5	1	2

	id	order	rating
1	1	Pizza	Fair
2	2	Pasta	Fair
3	3	Salad	Good
4	4	Pizza	Fair
5	5	Pizza	Poor

Key Functions for Labeling Data

```
scores <- c(85, 60, 45, 70, 50)
```

```
gender <- c(1, 2, 1, 2, 2)
```

`ifelse(test, yes, no)`

For simple conditional labeling

```
# Labeling as Pass/Fail
labels <- ifelse(scores >= 60,
                 "Pass", "Fail")

# Combining into a data frame
data <- data.frame(Scores = scores,
                  Label = labels)

data
```

	Scores	Label
1	85	Pass
2	60	Pass
3	45	Fail
4	70	Pass
5	50	Fail

`cut()`

For creating bins or ranges

```
# Creating bins
categories <- cut(scores,
                  breaks = c(0, 50, 70, 100),
                  labels = c("Low", "Medium",
                             "High"))

# Combining into a data frame
data <- data.frame(Scores = scores,
                  Category = categories)

data
```

	Scores	Category
1	85	High
2	60	Medium
3	45	Low
4	70	Medium
5	50	Low

`factor()`

Convert numeric or character data into labeled categories

```
# Converting to factor with labels
gender_labeled <- factor(gender,
                        levels = c(1, 2),
                        labels = c("Male",
                                   "Female"))

# Combine into a data frame
data <- data.frame(gender,
                  gender_labeled)

data
```

	gender	gender_labeled
1	1	Male
2	2	Female
3	1	Male
4	2	Female
5	2	Female

Descriptives by Group

Aggregation involves summarizing or grouping data based on certain variables.

Commonly used to calculate summary statistics like mean, variance, or sum for each group.


Benefits:

- Data summary
- Data simplification
- Statistical analysis
- Trend identification
- Data privacy
- Data-driven decisions

```
aggregate(x ~ group, FUN)
```


Descriptives by Group Example

R Code

 Start Over

Run Code

```
1 # average Sepal Length by iris species?
```

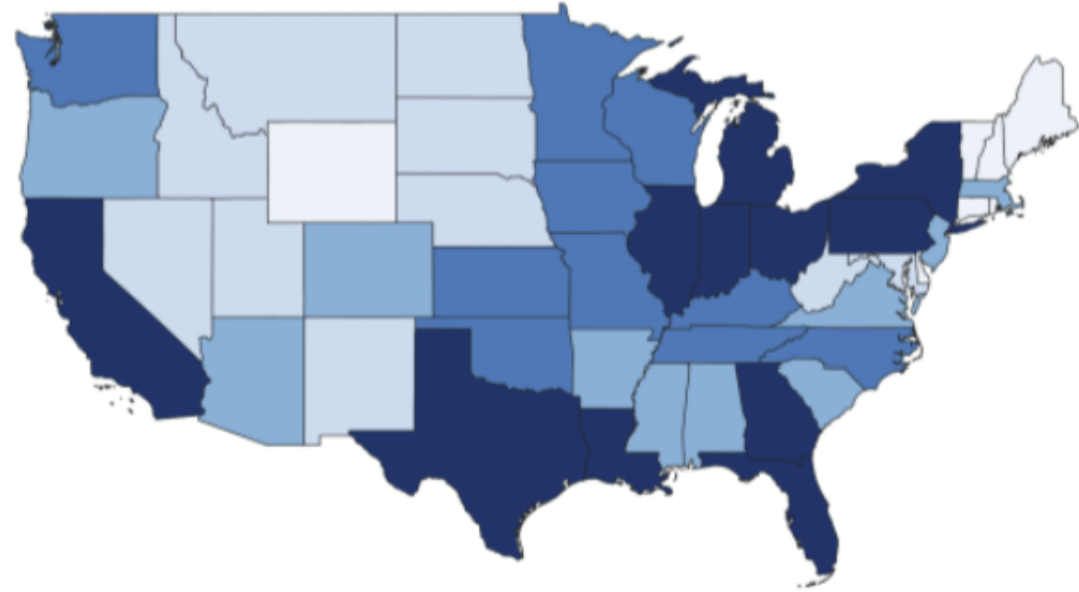
Descriptives by Group Visualization

Why visualize aggregated data?

- Makes group-level insights easier to interpret.
- Highlights patterns, trends, and outliers in grouped data
- Helps communicate findings effectively



(a) Point map of US hospitals



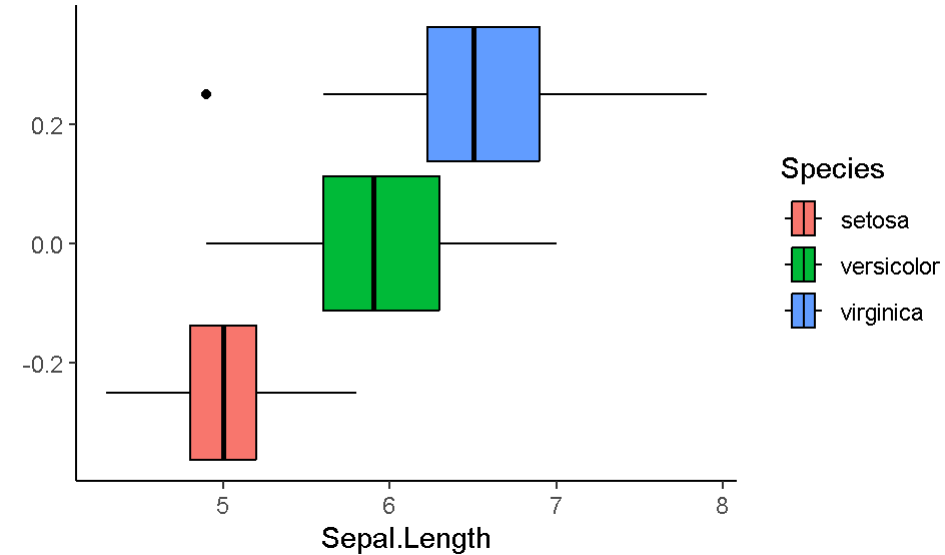
(b) Same data, but aggregated by state into a polygon (choropleth) map

Visualizations: Boxplots

Back to the `iris` dataset,
distribution of Sepal Length by
species

Anatomy of a boxplot:

- “Minimum”
- 25th Quantile (Q1)
- Median
- 75th Quantile (Q3)
- “Maximum”
- Points representing outliers



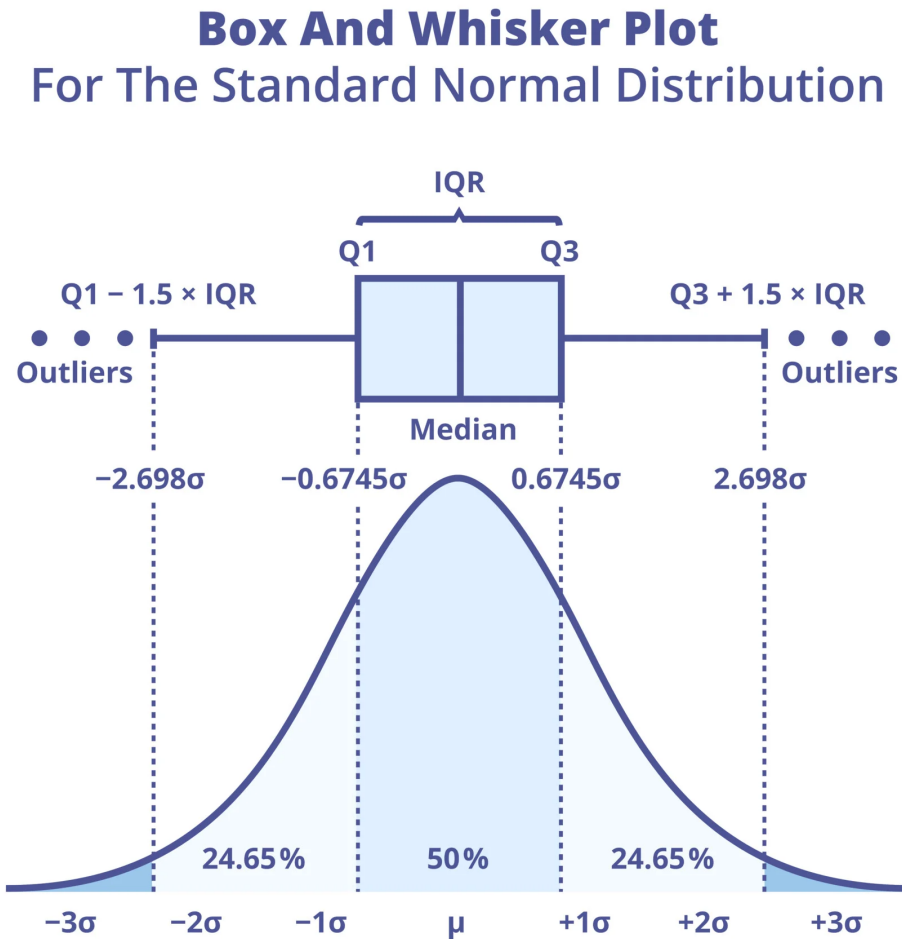
“Minimum” and “maximum” are not the *true* min and max

- Minimum: $Q1 - 1.5 * IQR$
- Maximum: $Q3 + 1.5 * IQR$

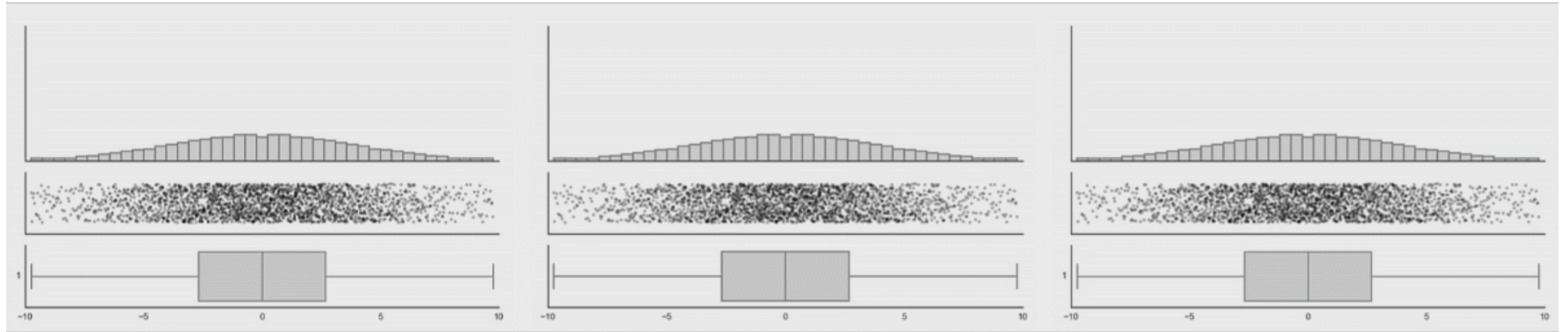
Means that the whiskers contain ~99% of the data,
rest are outliers

Visualizations: Boxplots

One more resource for boxplot anatomy



Visualizations: Boxplots...



Datasaurus Dozen Boxplots

Let's Do Some Visualization

Base R Graphics: Boxplot

`boxplot()` function

Its arguments are:

Required arguments:

- `x` = vector (variable) you want to plot

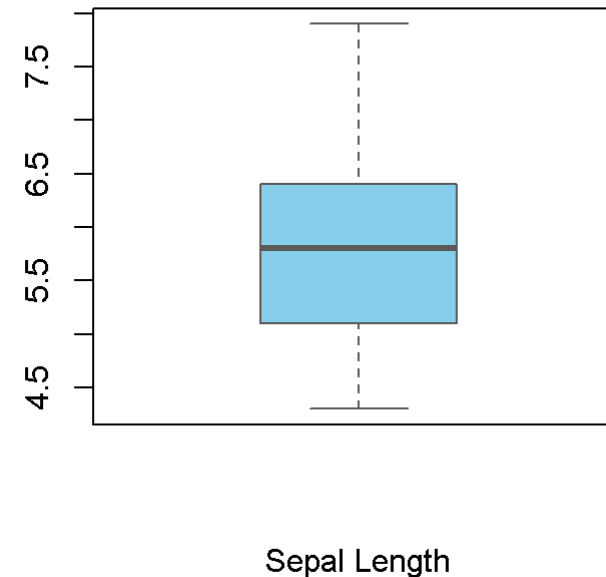
Optional arguments:

- `main`: title
- `xlab`: label for x-axis
- `ylab`: label for y-axis
- `border`: color for bar borders
- `col`: color for bars
- `horizontal`: T/F to switch

Plot

Code

Boxplot of Sepal Length



Let's Do Some Visualization

Base R Graphics: Boxplot

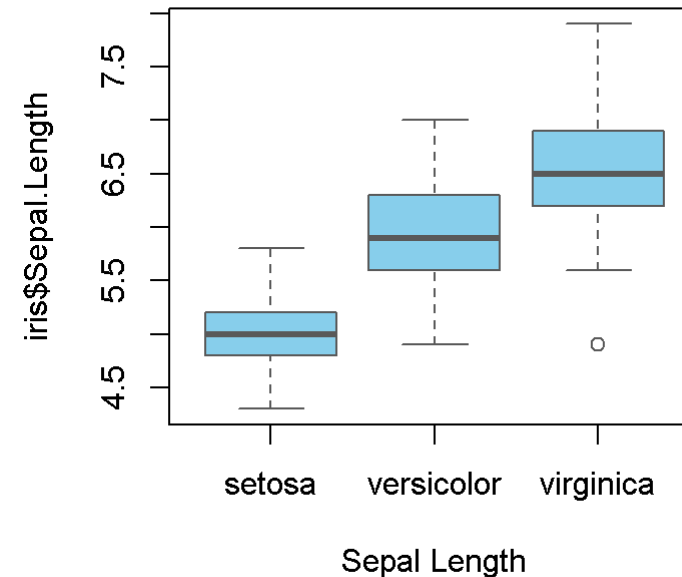
To group them, you can change the `x` to a “formula”

`outcome_var ~ group_var`

Plot

Code

Boxplot of Sepal Length by Species



Visualizations: Boxplot

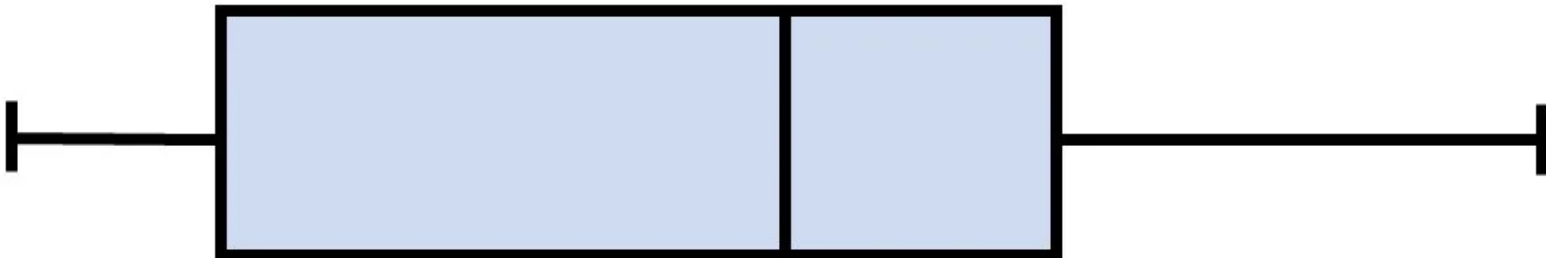
Normal Distribution



Positive Skew



Negative Skew



No Assignment!