

2.1 — Data 101 & Descriptive Statistics

ECON 480 • Econometrics • Fall 2020

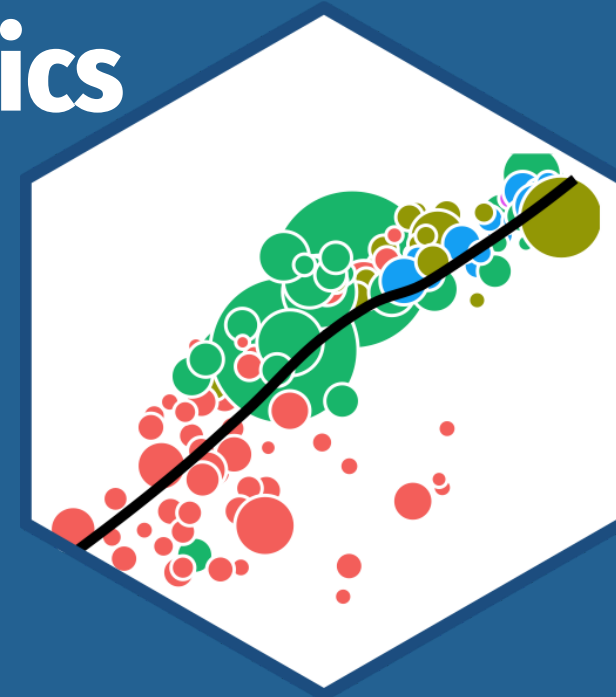
Ryan Safner

Assistant Professor of Economics

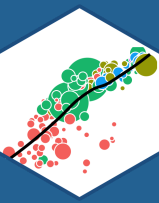
✉ safner@hood.edu

🔗 ryansafner/metricsF20

🌐 metricsF20.classes.ryansafner.com



Outline



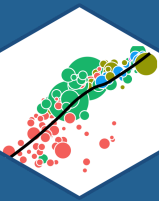
The Two Big Problems with Data

Data 101

Descriptive Statistics

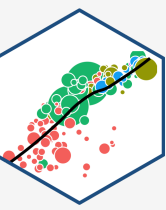
Measures of Center

Measures of Dispersion



The Two Big Problems with Data

Two Big Problems with Data



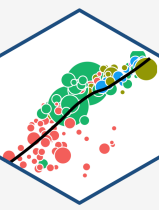
- We want to use econometrics to **identify** causal relationships and make **inferences** about them

1. Problem for **identification**: **endogeneity**

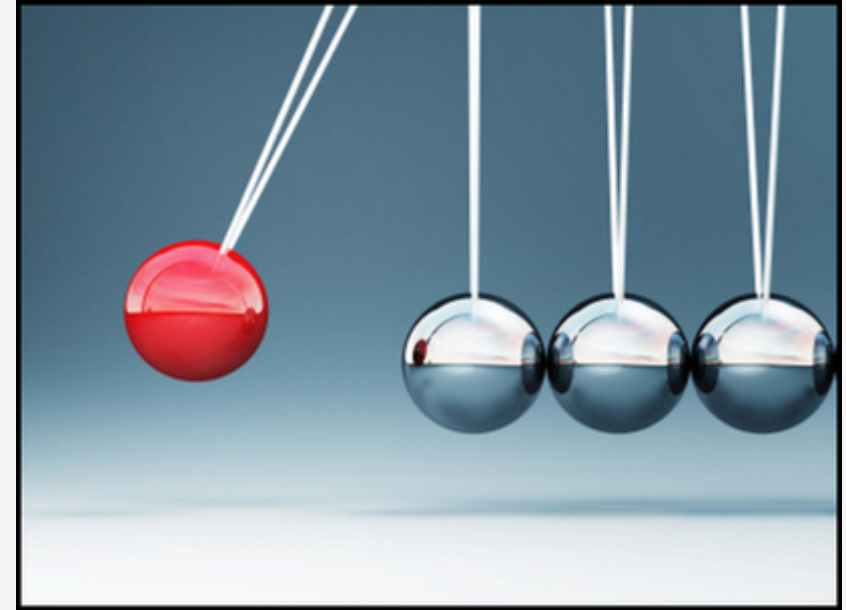
2. Problem for **inference**: **randomness**



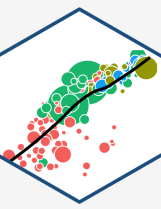
Identification Problem: Endogeneity



- An independent variable (X) is **exogenous** if its variation is *unrelated* to other factors that affect the dependent variable (Y)
- An independent variable (X) is **endogenous** if its variation is *related* to other factors that affect the dependent variable (Y)

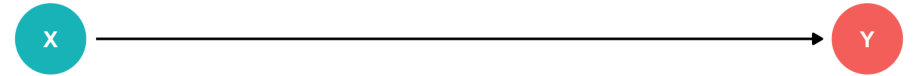


Identification Problem: Endogeneity

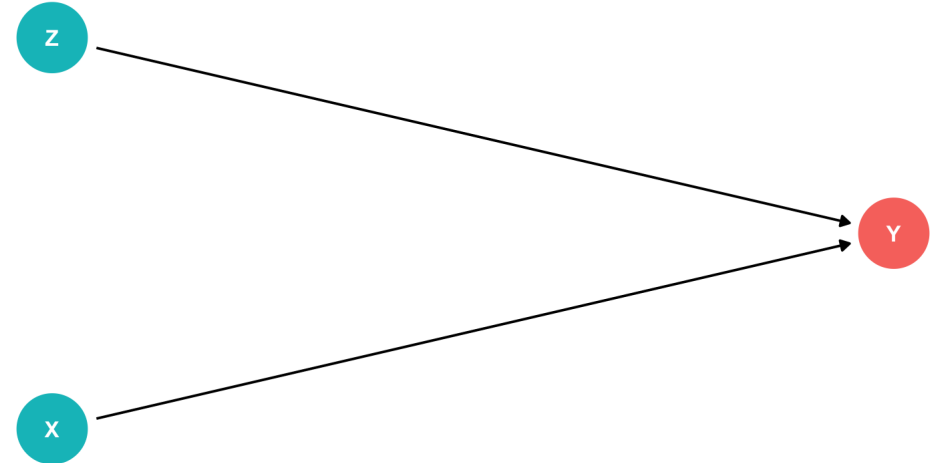


- An independent variable (X) is **exogenous** if its variation is *unrelated* to other factors that affect the dependent variable (Y)

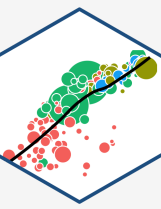
X causes Y



X and Z (independently) cause Y

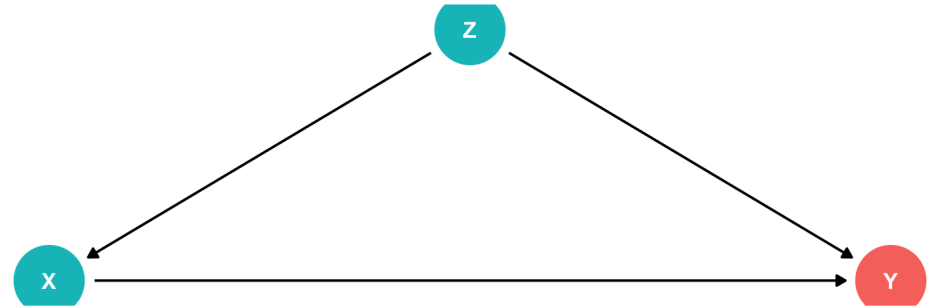


Identification Problem: Endogeneity



- An independent variable (X) is **endogenous** if its variation is *related* to other factors that affect the dependent variable (Y)

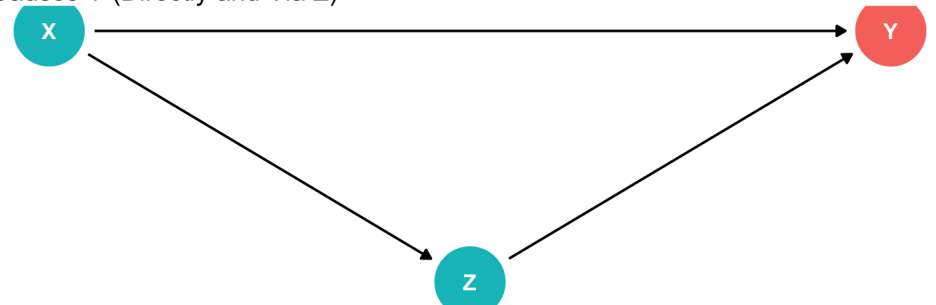
Z causes X and Y



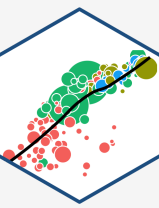
X Causes Y Indirectly Through Z



X Causes Y (Directly and Via Z)



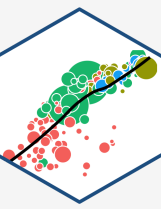
Inference Problem: Randomness



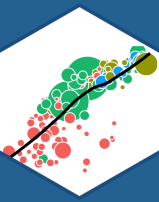
- Data is **random** due to **natural sampling variation**
 - Taking one sample of a population will yield slightly different information than another sample of the same population
- Common in statistics, *easy to fix*
- **Inferential Statistics**: making claims about a wider population using sample data



The Two Problems: Where We're Heading...Ultimately

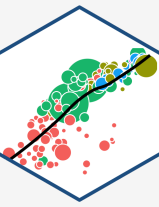


- We want to **identify** causal relationships between **population** variables
 - Logically first thing to consider
 - .hi-purple[Endogeneity problem]
- We'll use **sample** *statistics* to **infer** something about population *parameters*
 - In practice, we'll only ever have a finite *sample distribution* of data
 - We *don't* know the *population distribution* of data
 - .hi-purple[Randomness problem]



Data 101

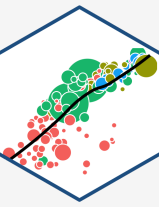
Data 101



- **Data** are information with context
- **Individuals** are the entities described by a set of data
 - e.g. persons, households, firms, countries



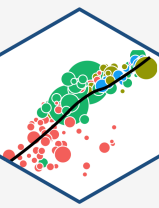
Data 101



- **Variables** are particular characteristics about an individual
 - e.g. age, income, profits, population, GDP, marital status, type of legal institutions
- **Observations** or **cases** are the separate individuals described by a collection of variables
 - e.g. for one individual, we have their age, sex, income, education, etc.
- individuals and observations are *not necessarily* the same:
 - e.g. we can have multiple observations on the same individual over time



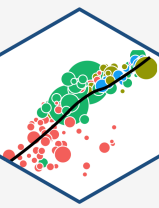
Categorical Data



- **Categorical data** place an individual into one of several possible *categories*
 - e.g. sex, season, political party
 - may be responses to survey questions
 - can be quantitative (e.g. age, zip code)
- In R: `character` or `factor` type data
 - `factor` \implies specific possible categories

Question	Categories or Responses
Do you invest in the stock market?	__ Yes __ No
What kind of advertising do you use?	__ Newspapers __ Internet __ Direct mailings
What is your class at school?	__ Freshman __ Sophomore __ Junior __ Senior
I would recommend this course to another student.	__ Strongly Disagree __ Slightly Disagree __ Slightly Agree __ Strongly Agree
How satisfied are you with this product?	__ Very Unsatisfied __ Unsatisfied __ Satisfied __ Very Satisfied

Categorical Data: Visualizing I



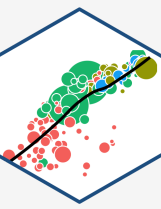
```
diamonds %>%  
  count(cut) %>%  
  mutate(frequency = n / sum(n),  
         percent = round(frequency * 100, 2))
```

Summary of diamonds by cut

cut	n	frequency	percent
Fair	1610	0.0298480	2.98
Good	4906	0.0909529	9.10
Very Good	12082	0.2239896	22.40
Premium	13791	0.2556730	25.57
Ideal	21551	0.3995365	39.95

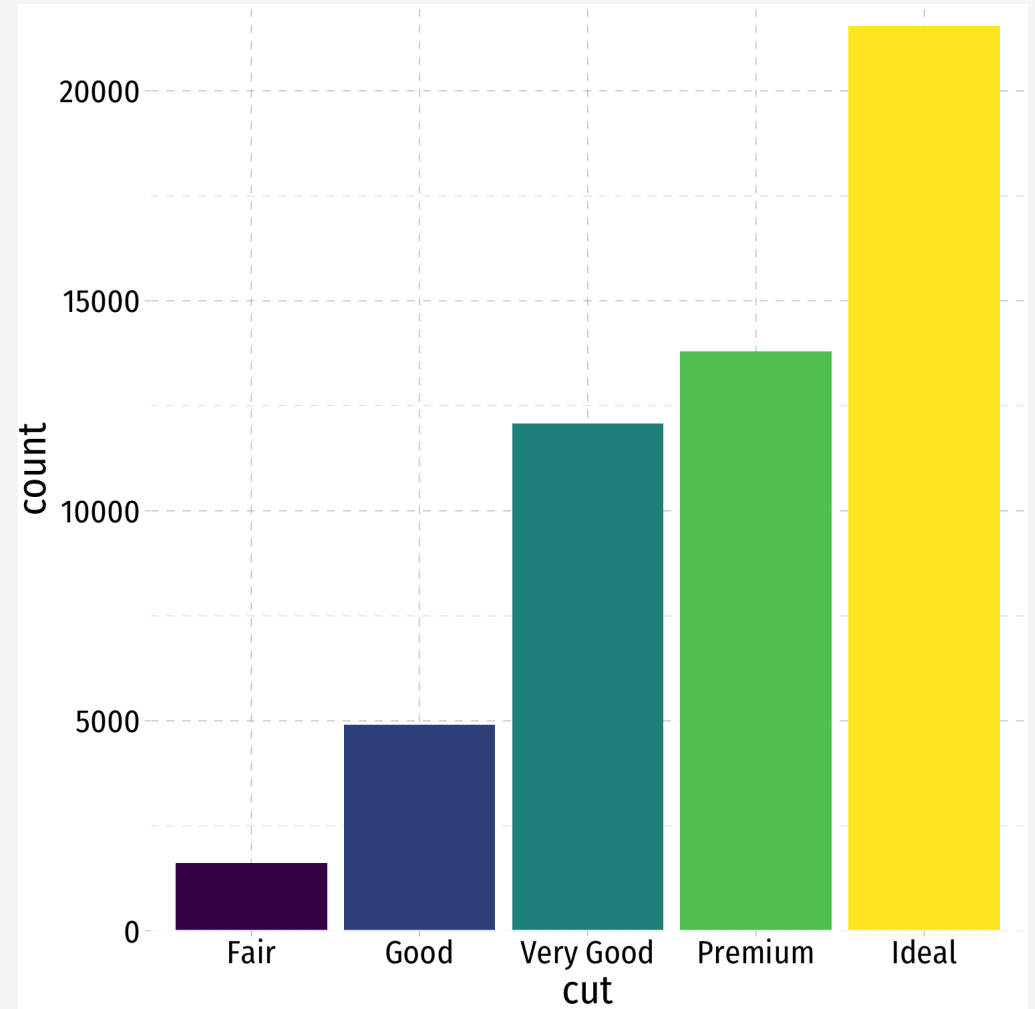
- Good way to represent categorical data is with a **frequency table**
- **Count (n)**: total number of individuals in a category
- **Frequency: proportion** of a category's occurrence relative to all data
 - Multiply proportions by 100% to get **percentages**

Categorical Data: Visualizing II

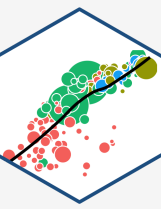


- Charts and graphs are *always* better ways to visualize data
- A **bar graph** represents categories as bars, with lengths proportional to the count or relative frequency of each category

```
ggplot(diamonds, aes(x=cut,  
                    fill=cut))+  
  geom_bar()+  
  guides(fill=F)+  
  theme_pander(base_family = "Fira Sans Condens",  
              base_size=20)
```



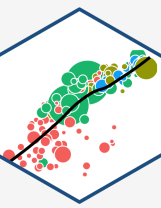
Categorical Data: Visualizing III



- Avoid pie charts!
- People are *not* good at judging 2-d differences (angles, area)
- People *are* good at judging 1-d differences (length)

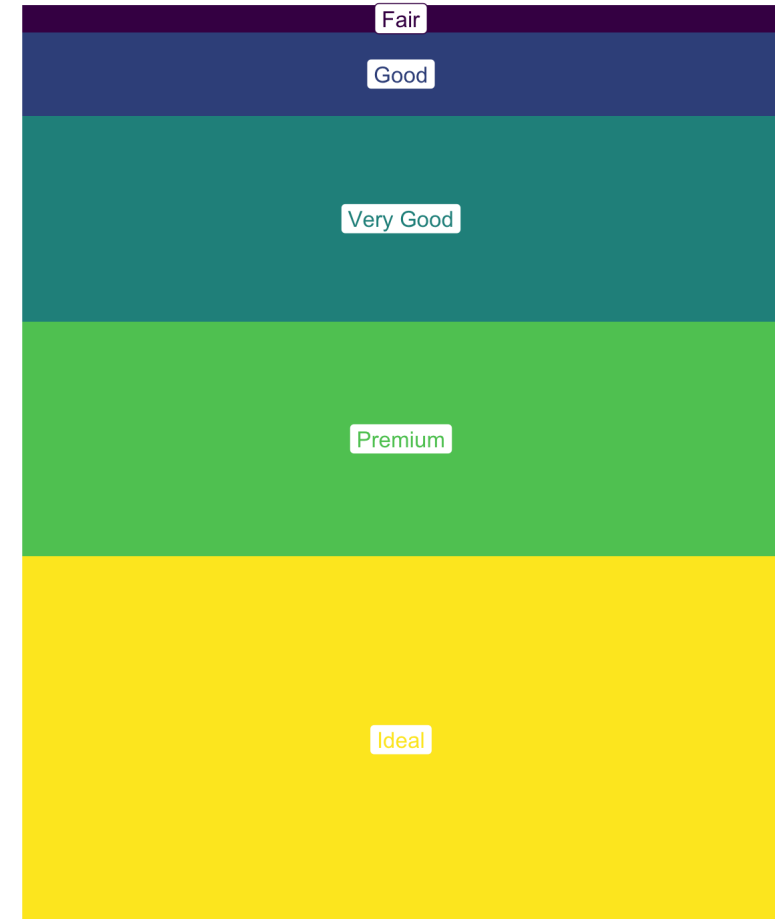


Categorical Data: Visualizing IV

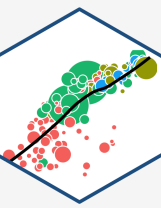


- Maybe a *stacked bar chart*

```
diamonds %>%  
  count(cut) %>%  
  ggplot(data = .)+  
  aes(x = "",  
       y = n)+  
  geom_col(aes(fill = cut))+  
  geom_label(aes(label = cut,  
                 color = cut),  
            position = position_stack(vjust =  
            )+  
  guides(color = F,  
         fill = F)+  
  theme_void()
```

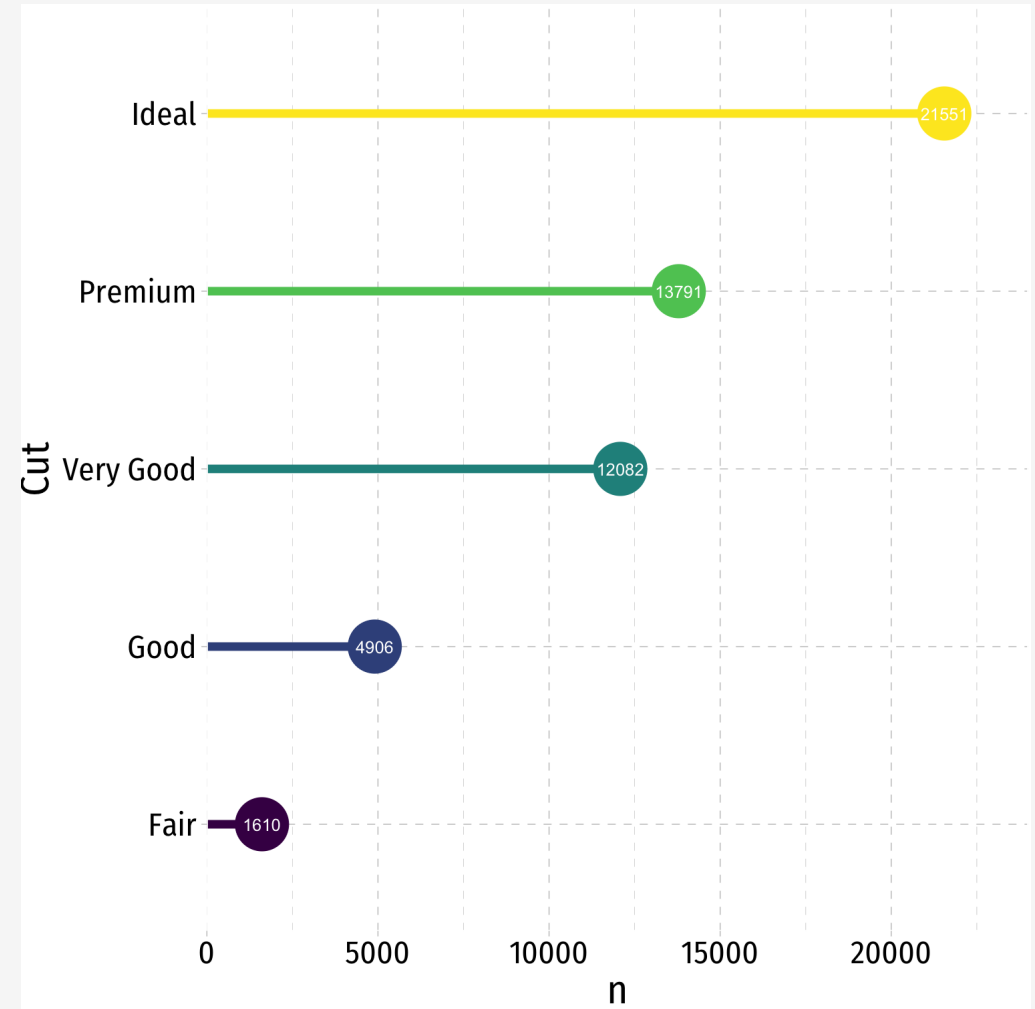


Categorical Data: Visualizing IV

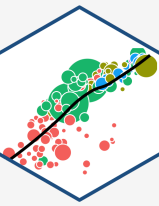


- Maybe *lollipop chart*

```
diamonds %>%
  count(cut) %>%
  mutate(cut_name = as.factor(cut)) %>%
ggplot(., aes(x = cut_name, y = n, color = cut))
  geom_point(stat="identity",
            fill="black",
            size=12) +
  geom_segment(aes(x = cut_name, y = 0,
                  xend = cut_name,
                  yend = n), size = 2)+
  geom_text(aes(label = n),color="white", size=
coord_flip()+
labs(x = "Cut")+
theme_pander(base_family = "Fira Sans Condens
              base_size=20)+
guides(color = F)
```



Categorical Data: Visualizing IV

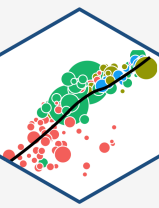


- Maybe a *treemap*

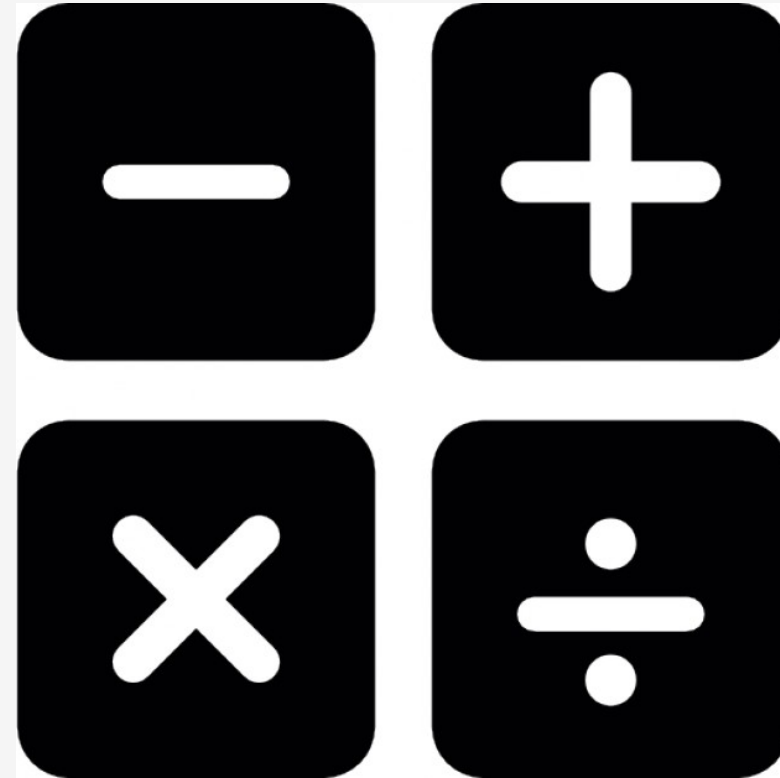
```
library(treemapify)
diamonds %>%
  count(cut) %>%
  ggplot(., aes(area = n, fill = cut)) +
  geom_treemap() +
  guides(fill = FALSE) +
  geom_treemap_text(aes(label = cut),
                    colour = "white",
                    place = "topleft",
                    grow = TRUE)
```



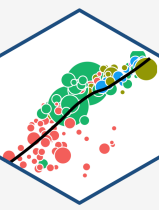
Quantitative Data I



- **Quantitative variables** take on numerical values of equal units that describe an individual
 - Units: points, dollars, inches
 - Context: GPA, prices, height
- We can mathematically manipulate *only* quantitative data
 - e.g. sum, average, standard deviation
- In R: `numeric` type data



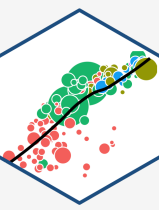
Discrete Data



- **Discrete data** are finite, with a countable number of alternatives
- **Categorical**: place data into categories
 - e.g. letter grades: A, B, C, D, F
 - e.g. class level: freshman, sophomore, junior, senior
- **Quantitative**: integers
 - e.g. SAT Score, number of children, age (years)



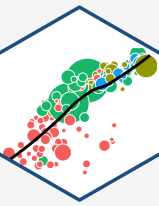
Continuous Data



- **Continuous data** are infinitely divisible, with an uncountable number of alternatives
 - e.g. weight, length, temperature, GPA
- Many discrete variables may be treated as if they are continuous
 - e.g. SAT scores (whole points), wages (dollars and cents)



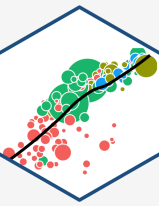
Spreadsheets



ID	Name	Age	Sex	Income
1	John	23	Male	41000
2	Emile	18	Male	52600
3	Natalya	28	Female	48000
4	Lakisha	31	Female	60200
5	Cheng	36	Male	81900

- The most common data structure we use is a **spreadsheet**
 - In *R*: a `data.frame` or `tibble`
- A **row** contains data about all variables for a single **individual**
- A **column** contains data about a single **variable** across all individuals

Spreadsheets



ID	Name	Age	Sex	Income
1	John	23	Male	41000
2	Emile	18	Male	52600
3	Natalya	28	Female	48000
4	Lakisha	31	Female	60200
5	Cheng	36	Male	81900

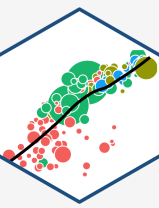
- Each **cell** can be referenced by its row and column (in that order!),
`df[row, column]`

```
example[3,2] # value in row 3, column 2
```

```
## # A tibble: 1 x 1  
##   Name  
##   <chr>  
## 1 Natalya
```

- Recall [how to “subset” data frames](#) from 1.2; though it's now much easier with `filter()` and `select()`!

Spreadsheets II



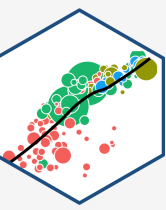
- It is common to use some notation like the following:
- Let $\{x_1, x_2, \dots, x_n\}$ be a simple data series on variable X
 - n individual observations
 - x_i is the value of the i^{th} observation for $i = 1, 2, \dots, n$

Quick Check: Let x represent the score on a homework assignment:

75, 100, 92, 87, 79, 0, 95

1. What is n ?
2. What is x_1 ?
3. What is x_6 ?

Datasets: Cross-Sectional



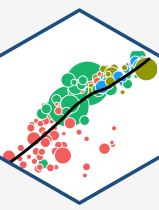
ID	Name	Age	Sex	Income
1	John	23	Male	41000
2	Emile	18	Male	52600
3	Natalya	28	Female	48000
4	Lakisha	31	Female	60200
5	Cheng	36	Male	81900

- **Cross-sectional data:** observations of individuals at a given point in time
- Each observation is a unique individual

x_i

- Simplest and most common data
- A "**snapshot**" to compare differences across individuals

Datasets: Time-Series



Year	GDP	Unemployment	CPI
1950	8.2	0.06	100
1960	9.9	0.04	118
1970	10.2	0.08	130
1980	12.4	0.08	190
1985	13.6	0.06	196

- **Time-series data**: observations of the *same* individual(s) over time

- Each observation is a time period

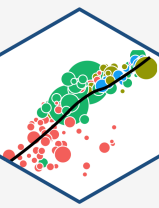
x_t

- Often used for macroeconomics, finance, and forecasting

- Unique challenges for time series

- A **"moving picture"** to see how individuals change over time

Datasets: Panel

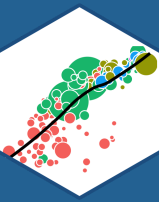


City	Year	Murders	Population	UR
Philadelphia	1986	5	3.700	8.7
Philadelphia	1990	8	4.200	7.2
D.C.	1986	2	0.250	5.4
D.C.	1990	10	0.275	5.5
New York	1986	3	6.400	9.6

- **Panel**, or **longitudinal** dataset: a time-series for *each* cross-sectional entity
 - Must be *same* individuals over time
- Each obs. is an individual in a time period

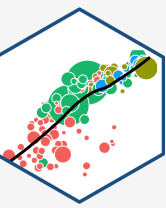
$$x_{it}$$

- More common today for serious researchers; unique challenges and benefits
- A **combination** of "snapshot" comparisons over time



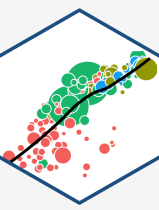
Descriptive Statistics

Variables and Distributions



- Variables take on different values, we can describe a variable's **distribution** (of these values)
- We want to *visualize* and *analyze* distributions to search for meaningful patterns using **statistics**

Two Branches of Statistics

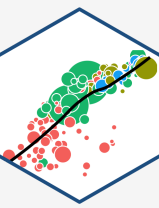


- Two main branches of statistics:
 1. **Descriptive Statistics:** describes or summarizes the properties of a sample
 2. **Inferential Statistics:** infers properties about a larger population from the properties of a sample[†]

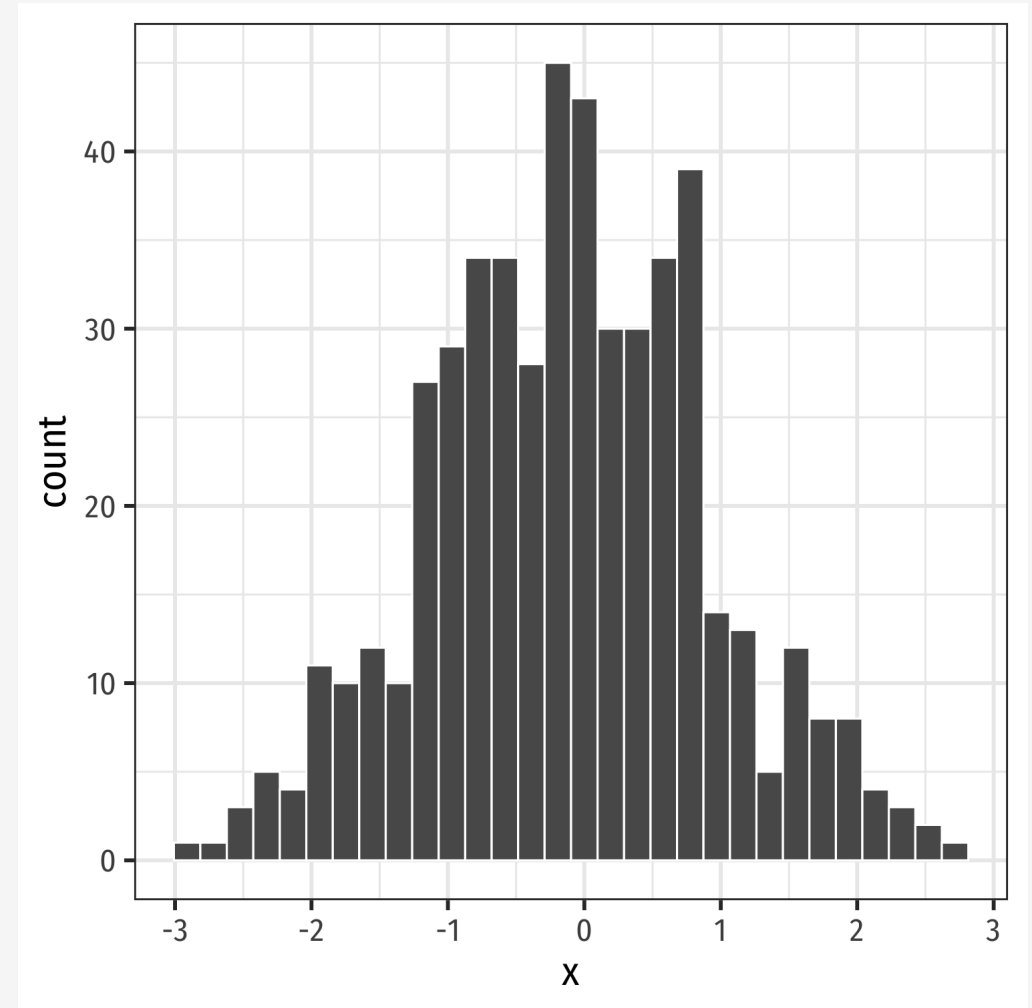


[†] We'll encounter inferential statistics mainly in the context of regression later.

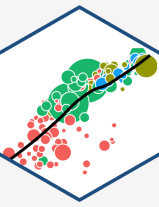
Histograms



- A common way to present a *quantitative* variable's distribution is a **histogram**
 - The quantitative analog to the bar graph for a categorical variable
- Divide up values into **bins** of a certain size, and count the number of values falling within each bin, representing them visually as bars



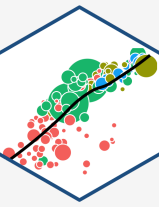
Histogram: Example



Example: a class of 13 students takes a quiz (out of 100 points) with the following results:

$\{0, 62, 66, 71, 71, 74, 76, 79, 83, 86, 88, 93, 95\}$

Histogram: Example

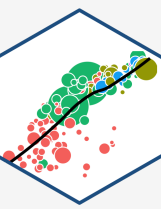


Example: a class of 13 students takes a quiz (out of 100 points) with the following results:

{0, 62, 66, 71, 71, 74, 76, 79, 83, 86, 88, 93, 95}

```
quizzes<-tibble(scores = c(0,62,66,71,71,74,76,79,83,86,88,93
```

Histogram: Example

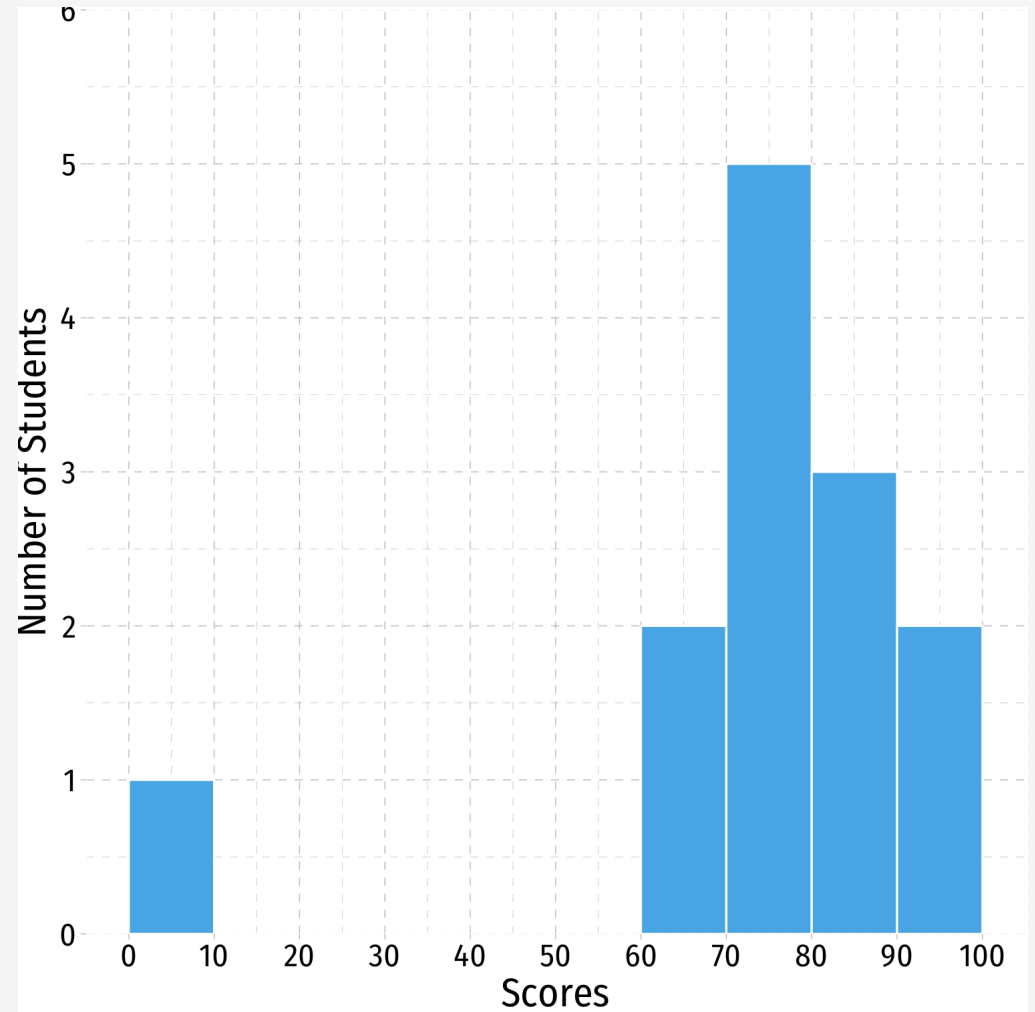


Example: a class of 13 students takes a quiz (out of 100 points) with the following results:

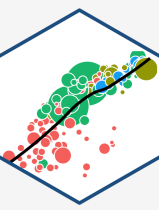
{0, 62, 66, 71, 71, 74, 76, 79, 83, 86, 88, 93, 95}

```
h<-ggplot(quizzes,aes(x=scores))+  
  geom_histogram(breaks = seq(0,100,10),  
                color = "white",  
                fill = "#56B4E9")+  
  scale_x_continuous(breaks = seq(0,100,10))+  
  scale_y_continuous(limits = c(0,6), expand = c(0,0))+  
  labs(x = "Scores",  
       y = "Number of Students")+  
  ggthemes::theme_pander(base_family = "Fira Sans Condensed",  
                          base_size=20)
```

h

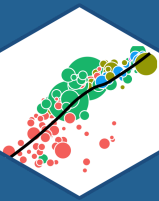


Descriptive Statistics



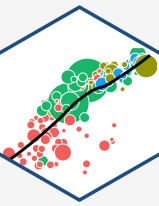
- We are often interested in the *shape* or *pattern* of a distribution, particularly:
 - Measures of **center**
 - Measures of **dispersion**
 - **Shape** of distribution





Measures of Center

Mode

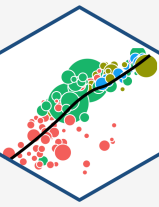


- The **mode** of a variable is simply its most frequent value
- A variable can have multiple modes

Example: a class of 13 students takes a quiz (out of 100 points) with the following results:

$\{0, 62, 66, 71, 71, 74, 76, 79, 83, 86, 88, 93, 95\}$

Mode

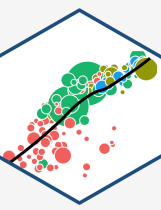


- There is no dedicated `mode()` function in R, surprisingly
- A workaround in `dplyr`:

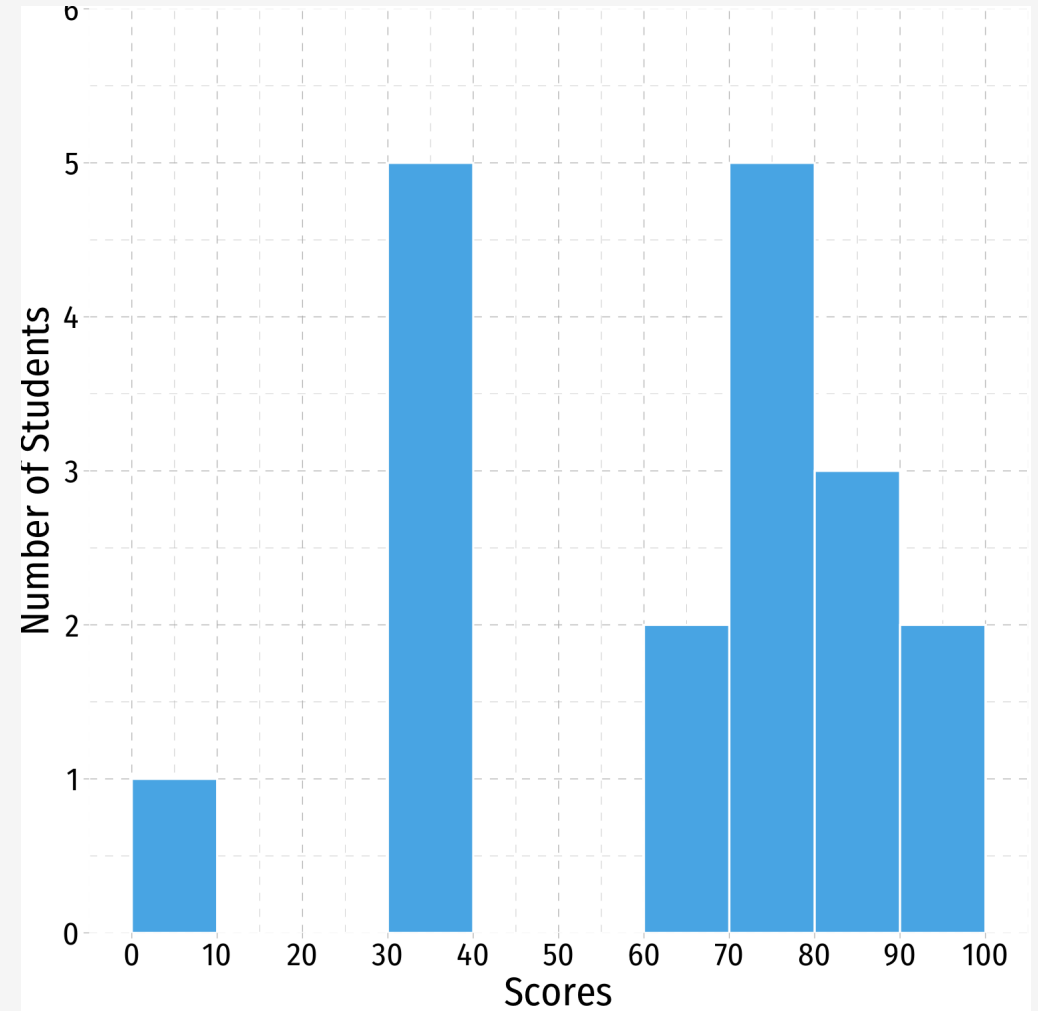
```
quizzes %>%  
  count(scores) %>%  
  arrange(desc(n))
```

```
## # A tibble: 12 x 2  
##   scores     n  
##   <dbl> <int>  
## 1     71     2  
## 2      0     1  
## 3     62     1  
## 4     66     1  
## 5     74     1  
## 6     76     1  
## 7     79     1  
## 8     83     1  
## 9     86     1  
## 10    88     1  
## 11    93     1  
## 12    95     1
```

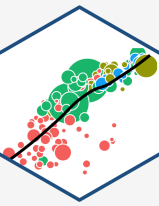
Multi-Modal Distributions



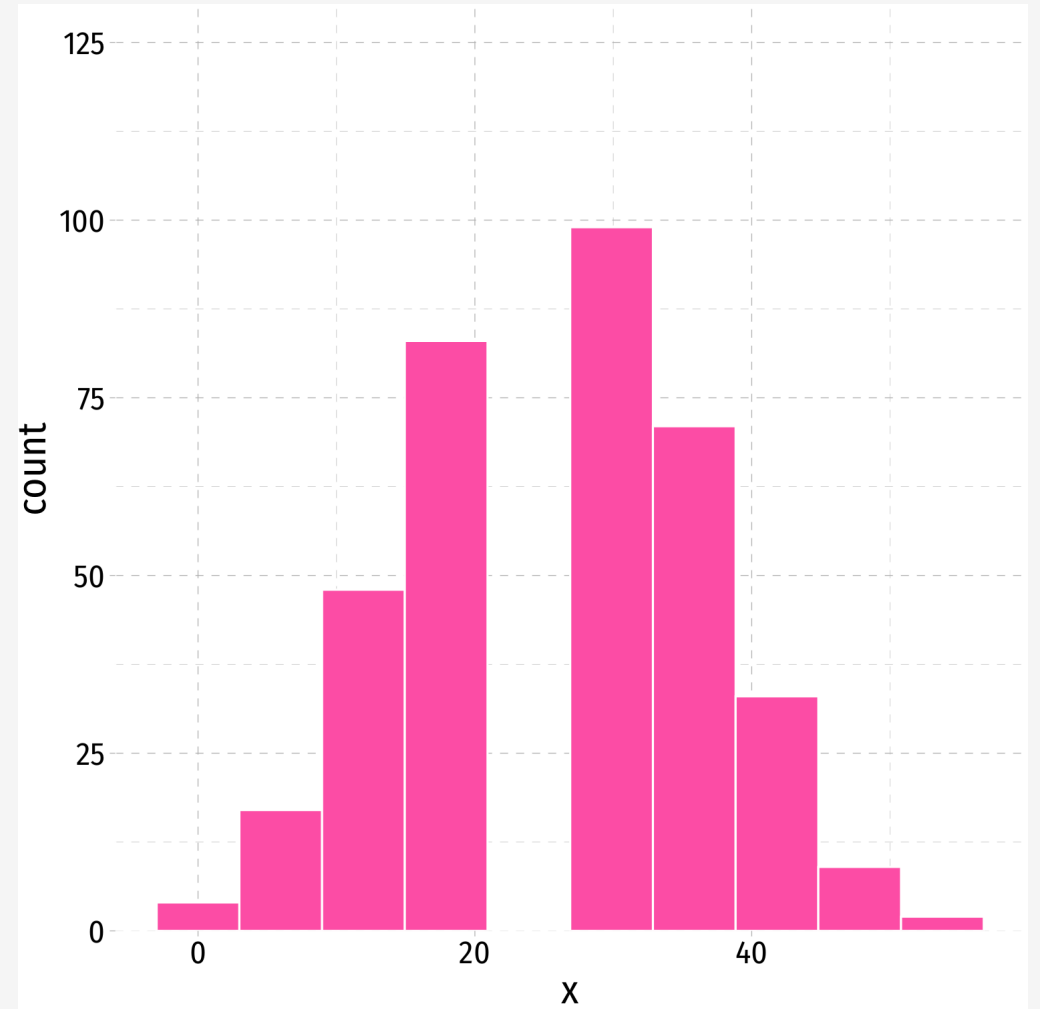
- Looking at a histogram, the modes are the "peaks" of the distribution
 - Note: depends on how wide you make the bins!
- May be unimodal, bimodal, trimodal, etc



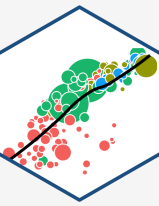
Symmetry and Skew I



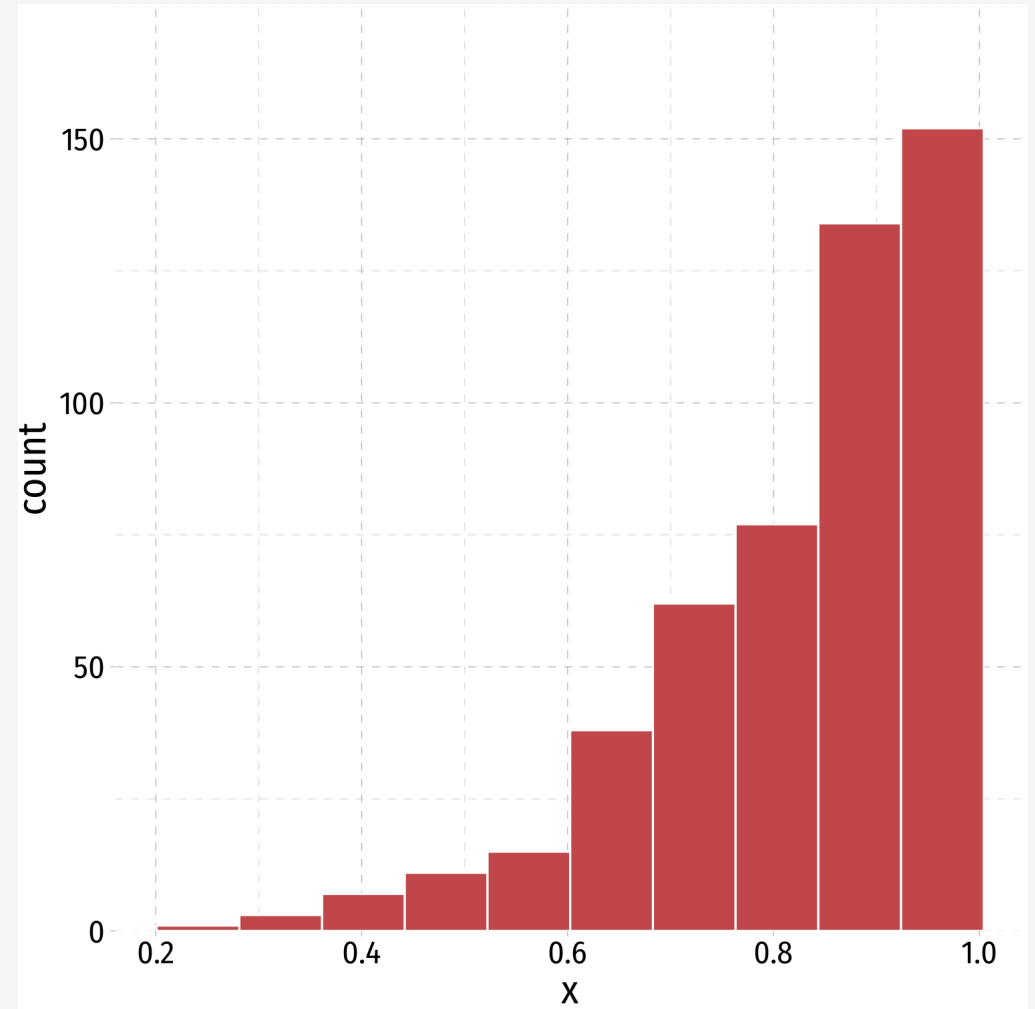
- A distribution is **symmetric** if it looks roughly the same on either side of the "center"
- The thinner ends (far left and far right) are called the **tails** of a distribution



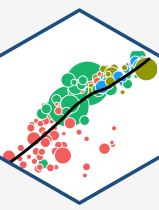
Symmetry and Skew I



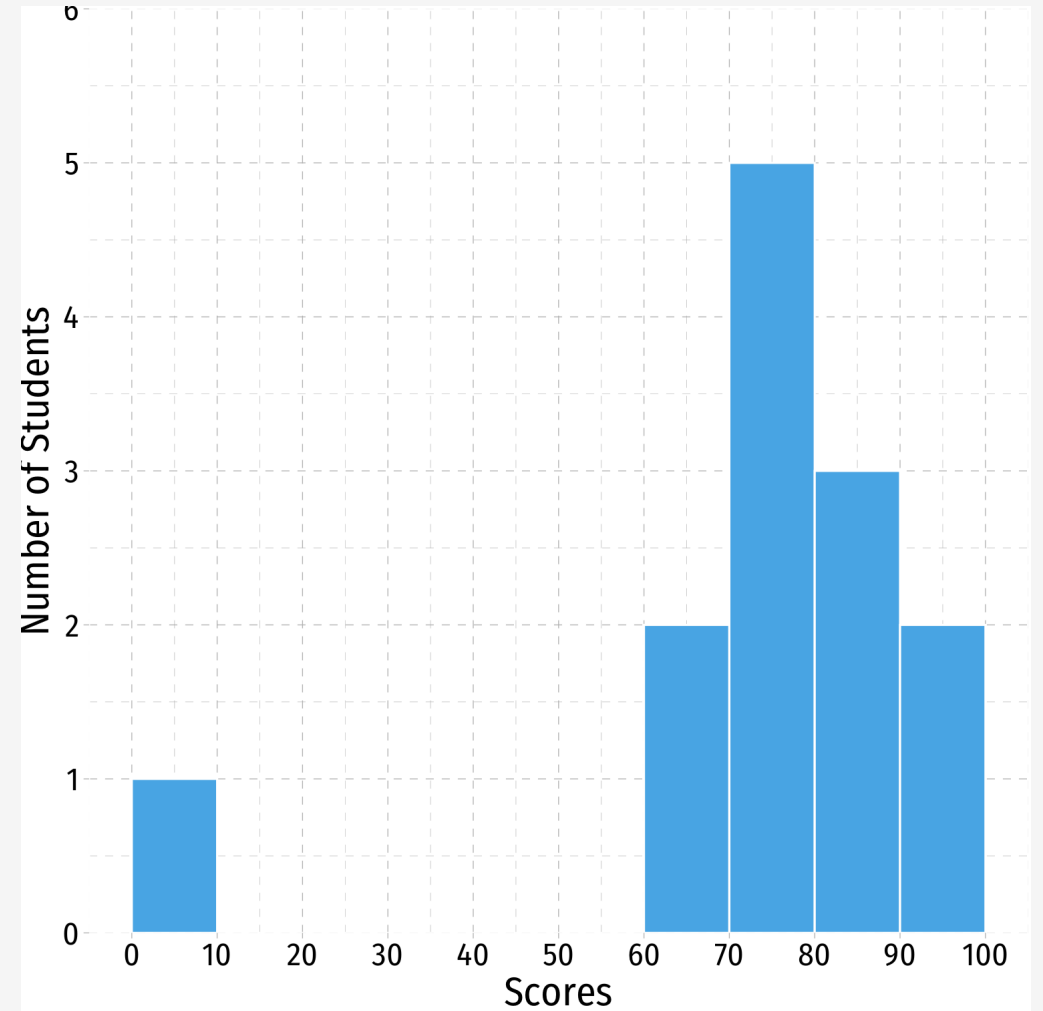
- If one tail stretches farther than the other, distribution is **skewed** in the direction of the longer tail



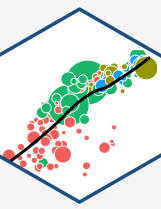
Outliers



- **Outlier**: extreme value that does not appear part of the general pattern of a distribution
- Can strongly affect descriptive statistics
- Might be the most informative part of the data
- Could be the result of errors
- Should always be explored and discussed!



Arithmetic Mean (Population)



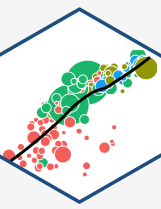
- The natural measure of the center of a *population's* distribution is its **"average"** or **arithmetic mean (μ)**

$$\mu = \frac{x_1 + x_2 + \dots + x_N}{N} = \frac{1}{N} \sum_{i=1}^N x_i$$

- For N values of variable x , "mu" is the sum of all individual x values (x_i) from 1 to N , divided by the N number of values[†]
- See [today's class notes](#) for more about the **summation operator, Σ** , it'll come up again!

[†] Note the mean need not be an actual value of the data!

Arithmetic Mean (Sample)



- When we have a *sample*, we compute the **sample mean** (\bar{x})

$$\bar{x} = \frac{x_1 + x_2 + \dots + x_n}{n} = \frac{1}{n} \sum_{i=1}^n x_i$$

- For n values of variable x , "x-bar" is the sum of all individual x values (x_i) divided by the n number of values

Example:

{0, 62, 66, 71, 71, 74, 76, 79, 83, 86, 88, 93, 95}

$$\bar{x} = \frac{1}{13}(0 + 62 + 66 + 71 + 71 + 74 + 76 + 79 + 83 + 86 + 88 + 93 + 95)$$

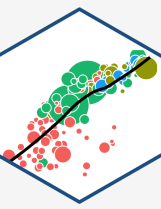
$$\bar{x} = \frac{944}{13}$$

$$\bar{x} = 72.62$$

```
quizzes %>%  
  summarize(mean=mean(scores))
```

```
## # A tibble: 1 x 1  
##   mean  
##   <dbl>  
## 1 72.6
```

Arithmetic Mean: Affected by Outliers



- If we drop the outlier (0)

Example:

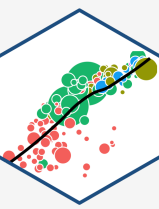
{62, 66, 71, 71, 74, 76, 79, 83, 86, 88, 93, 95}

$$\begin{aligned}\bar{x} &= \frac{1}{12}(62 + 66 + 71 + 71 + 74 + 76 + 79 + 83 + 86 + 88 + 93 + 95) \\ &= 78.67\end{aligned}$$

```
quizzes %>%  
  filter(scores>0) %>%  
  summarize(mean=mean(scores))
```

```
## # A tibble: 1 x 1  
##   mean  
##   <dbl>  
## 1  78.7
```

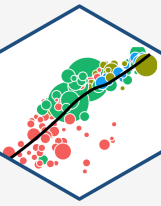
Median



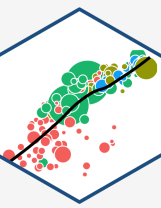
{0, 62, 66, 71, 71, 74, **76**, 79, 83, 86, 88, 93, 95}

- The **median** is the midpoint of the distribution
 - 50% to the left of the median, 50% to the right of the median
- Arrange values in numerical order
 - For odd n : median is middle observation
 - For even n : median is average of two middle observations

Mean, Median, and Outliers



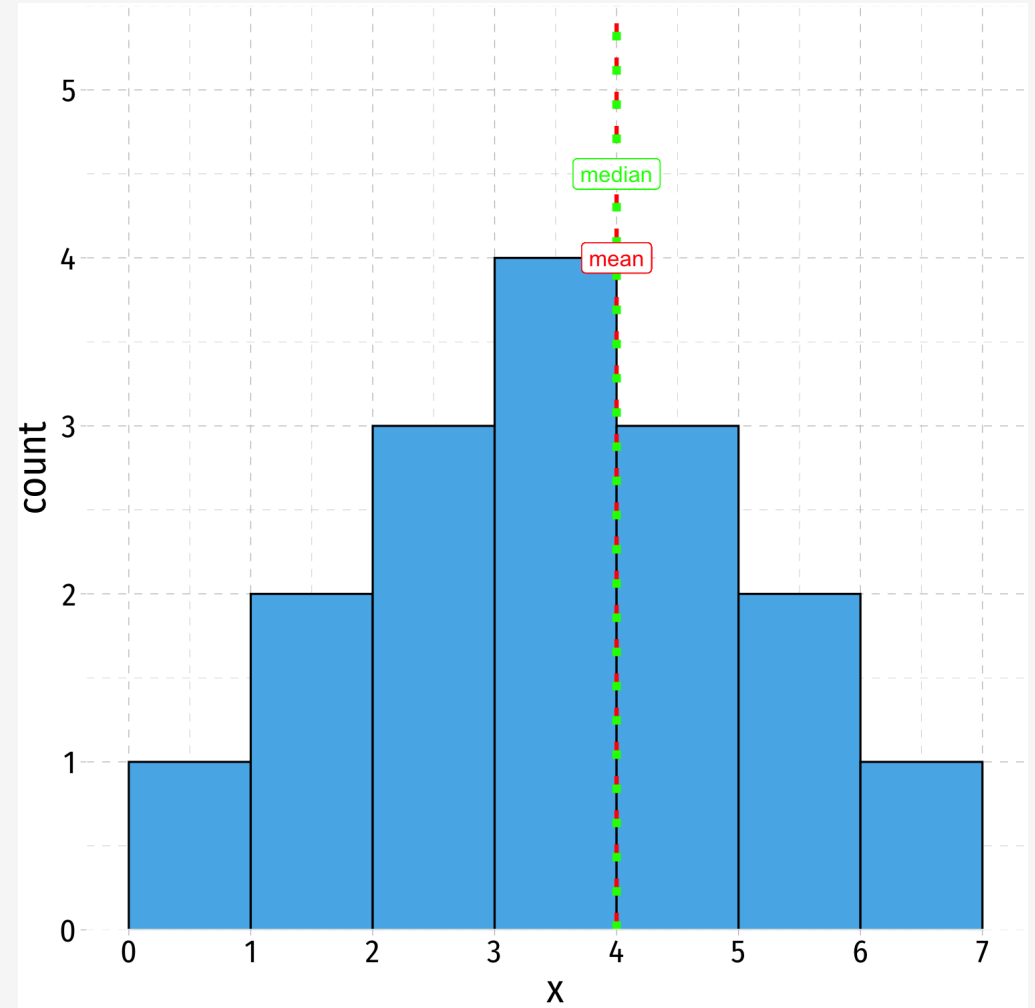
Mean, Median, Symmetry, Skew I



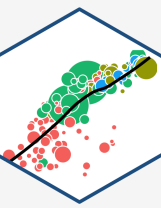
- Symmetric distribution: mean \approx median

```
symmetric %>%  
  summarize(mean = mean(x),  
            median = median(x))
```

```
## # A tibble: 1 x 2  
##   mean median  
##   <dbl> <dbl>  
## 1     4     4
```



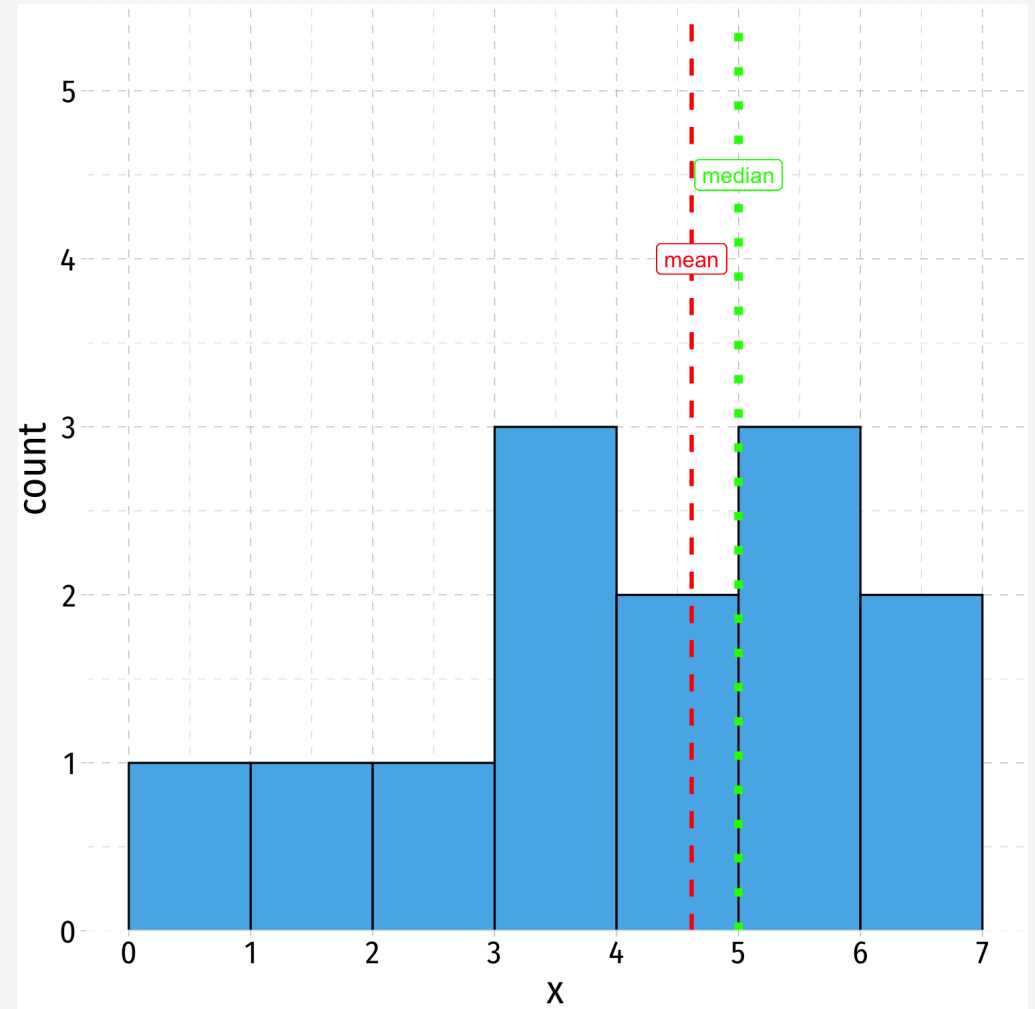
Mean, Median, Symmetry, Skew II



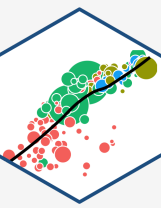
- Left-skewed: mean < median

```
leftskew %>%  
  summarize(mean = mean(x),  
            median = median(x))
```

```
##      mean median  
## 1 4.615385      5
```



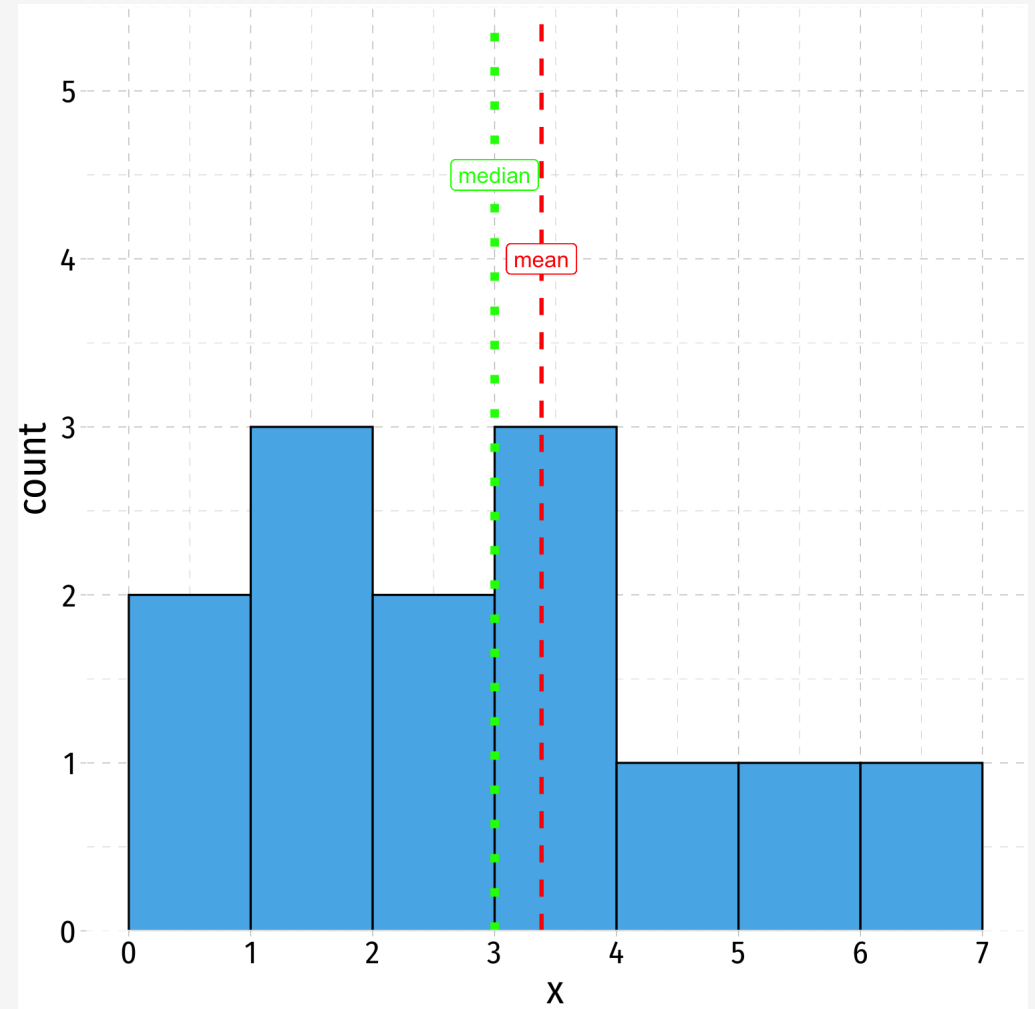
Mean, Median, Symmetry, Skew III

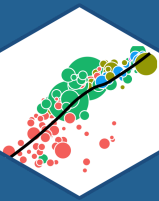


- Right-skewed: mean $>$ median

```
rightskew %>%  
  summarize(mean = mean(x),  
            median = median(x))
```

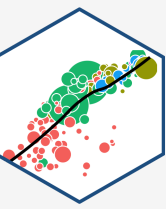
```
## # A tibble: 1 x 2  
##   mean median  
##   <dbl> <dbl>  
## 1  3.38     3
```





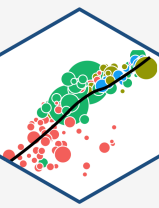
Measures of Dispersion

Measures of Dispersion: Range



- The more *variation* in the data, the less helpful a measure of central tendency will tell us
- Beyond just the center, we also want to measure the spread
- Simplest metric is **range** = $max - min$

Measures of Dispersion: 5 Number Summary I



- Common set of summary statistics of a distribution: **"five number summary"**:

1. Minimum value
2. 25th percentile (Q_1 , median of first 50% of data)
3. 50th percentile (median, Q_2)
4. 75th percentile (Q_3 , median of last 50% of data)
5. Maximum value

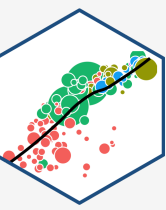
```
# Base R summary command (includes Mean)
summary(quizzes$scores)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##      0.00   71.00   76.00   72.62   86.00   95.00
```

```
quizzes %>% # dplyr
  summarize(Min = min(scores),
            Q1 = quantile(scores, 0.25),
            Median = median(scores),
            Q3 = quantile(scores, 0.75),
            Max = max(scores))
```

```
## # A tibble: 1 x 5
##   Min     Q1 Median     Q3    Max
##   <dbl> <dbl> <dbl> <dbl> <dbl>
## 1     0    71    76    86    95
```

Measures of Dispersion: 5 Number Summary II

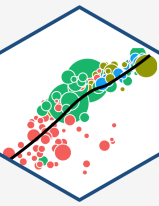


- The n^{th} **percentile** of a distribution is the value that places n percent of values beneath it

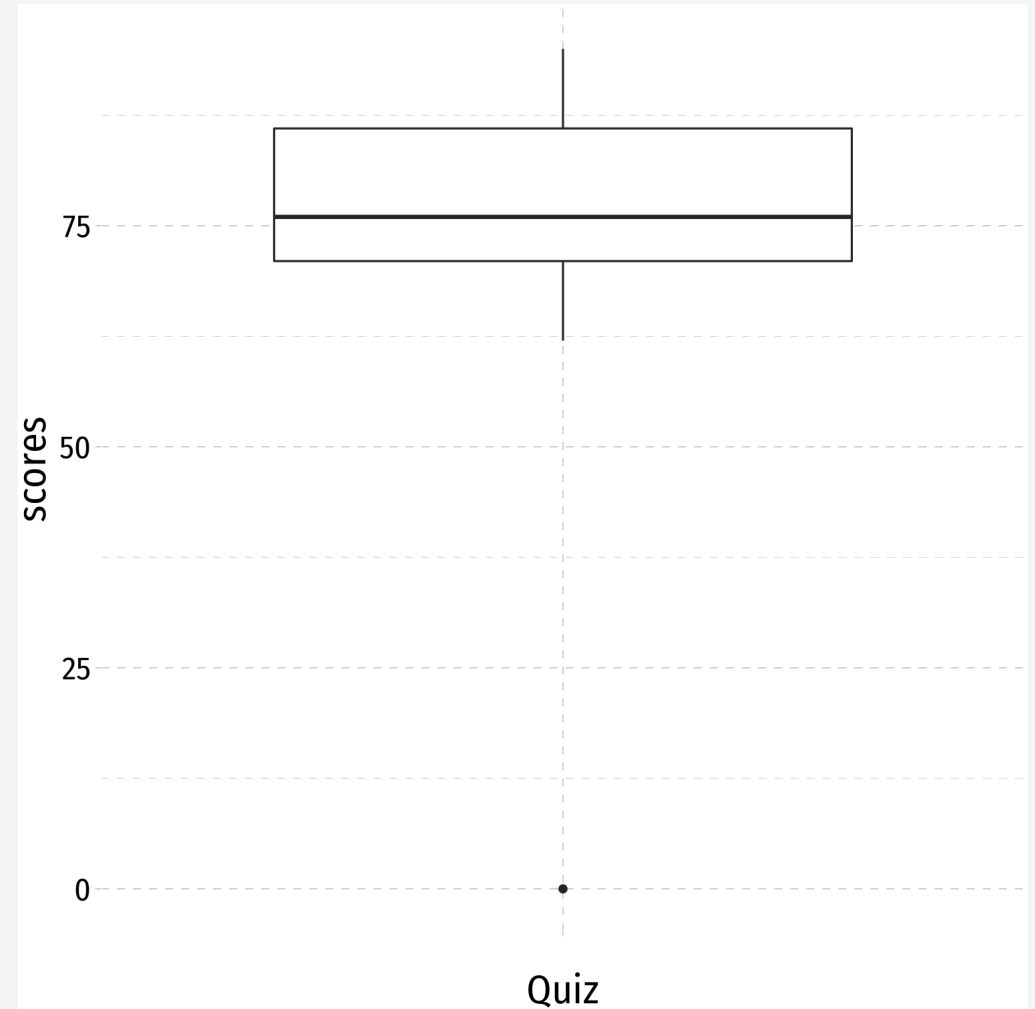
```
quizzes %>%  
  summarize("37th percentile" = quantile(scores,0.37))
```

```
## # A tibble: 1 x 1  
##   `37th percentile`  
##           <dbl>  
## 1           72.3
```

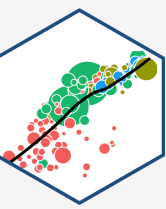
Boxplots I



- **Boxplots** are a great way to visualize the 5 number summary
- **Height of box:** Q_1 to Q_3 (known as **interquartile range (IQR)**, middle 50% of data)
- **Line inside box:** median (50th percentile)
- **"Whiskers"** identify data within $1.5 \times IQR$
- Points *beyond* whiskers are **outliers**
 - common definition:
 $Outlier > 1.5 \times IQR$



Comparisons I



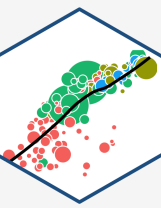
- Boxplots (and five number summaries) are great for comparing two distributions

Example:

Quiz 1 : {0, 62, 66, 71, 71, 74, 76, 79, 83, 86, 88, 93, 95}

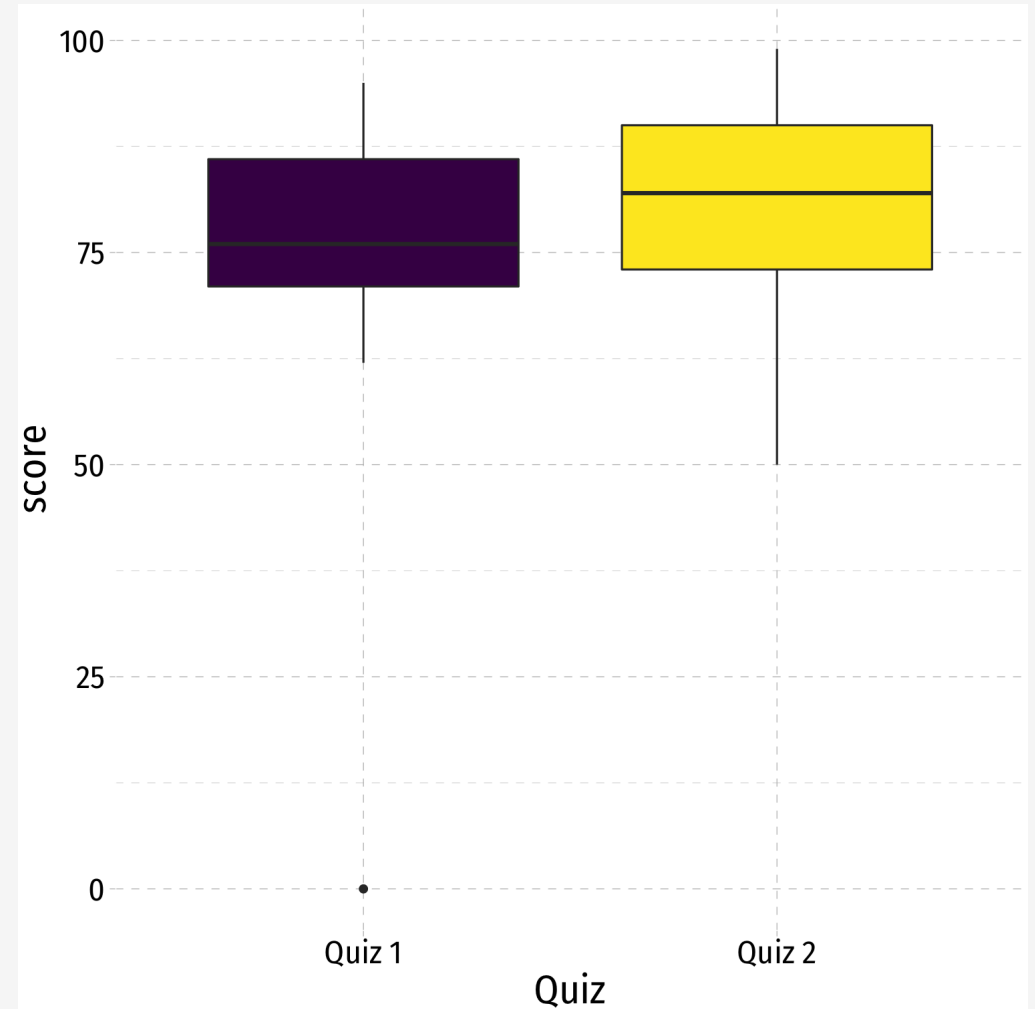
Quiz 2 : {50, 62, 72, 73, 79, 81, 82, 82, 86, 90, 94, 98, 99}

Comparisons II

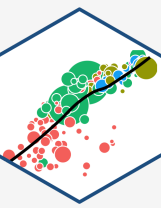


```
quizzes_new %>% summary()
```

##	student	quiz_1	quiz_2
##	Min. : 1	Min. : 0.00	Min. : 50.00
##	1st Qu.: 4	1st Qu.: 71.00	1st Qu.: 73.00
##	Median : 7	Median : 76.00	Median : 82.00
##	Mean : 7	Mean : 72.62	Mean : 80.62
##	3rd Qu.: 10	3rd Qu.: 86.00	3rd Qu.: 90.00
##	Max. : 13	Max. : 95.00	Max. : 99.00



Aside: Making Nice Summary Tables I



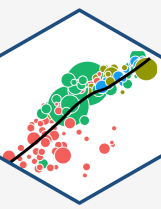
- I don't like the options available for printing out summary statistics
- So I wrote my own R function called `summary_table()` that makes nice summary tables (it uses `dplyr` and `tidyr`!). To use:
 1. Download the `summaries.R` [file](#) from the website[†] and move it to your working directory/project folder
 2. Load the function with the `source()` command:[‡]

```
source("summaries.R")
```

[†] One day I'll make this part of a package I'll write.

[‡] If it *was* a package, then you'd load with `library()`. But you can run a single `.R` script with `source()`.

Aside: Making Nice Summary Tables II



3) The function has at least 2 arguments: the `data.frame` (automatically piped in if you use the pipe!) and then all variables you want to summarize, separated by commas[†]

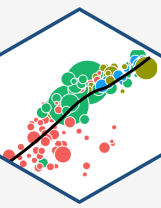
```
mpg %>%
  summary_table(hwy, cty, cyl)
```



```
## # A tibble: 3 x 9
##   Variable  Obs   Min    Q1 Median    Q3   Max  Mean `Std. Dev.`
##   <chr>    <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>      <dbl>
## 1 cty      234     9    14    17    19    35  16.9      4.26
## 2 cyl      234     4     4     6     8     8   5.89     1.61
## 3 hwy      234    12    18    24    27    44  23.4     5.95
```

[†] There is one restriction: No variable name can have an underscore (`_`) in it. You will have to rename them or else you will break the function!

Aside: Making Nice Summary Tables II



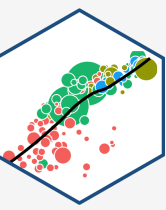
4) When `knit`ted in R `markdown`, it looks nicer:

```
mpg %>%  
  summary_table(hwy, cty, cyl) %>%  
  knitr::kable(., format="html")
```

Variable	Obs	Min	Q1	Median	Q3	Max	Mean	Std. Dev.
cty	234	9	14	17	19	35	16.86	4.26
cyl	234	4	4	6	8	8	5.89	1.61
hwy	234	12	18	24	27	44	23.44	5.95

- We'll talk more about using `markdown` and making final products nicer when we discuss your paper project (have you forgotten?)

Measures of Dispersion: Deviations

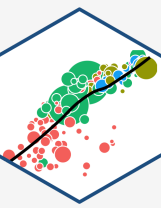


- Every observation i **deviates** from the mean of the data:

$$\text{deviation}_i = x_i - \mu$$

- There are as many deviations as there are data points (n)
- We can measure the *average* or **standard deviation** of a variable from its mean
- Before we get there...

Variance (Population)

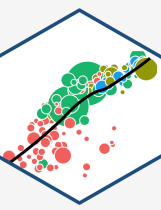


- The **population variance** (σ^2) of a *population* distribution measures the average of the *squared* deviations from the *population* mean (μ)

$$\sigma^2 = \frac{1}{N} \sum_{i=1}^N (x_i - \mu)^2$$

- Why do we square deviations?
- What are these units?

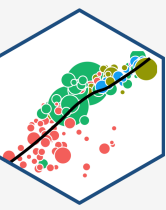
Standard Deviation (Population)



- Square root the variance to get the **population standard deviation** (σ), the average deviation from the population mean (in same units as x)

$$\sigma = \sqrt{\sigma^2} = \sqrt{\frac{1}{N} \sum_{i=1}^N (x_i - \mu)^2}$$

Variance (Sample)

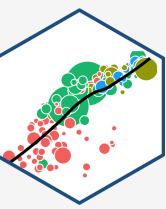


- The **sample variance** (s^2) of a *sample* distribution measures the average of the *squared* deviations from the *sample* mean (\bar{x})

$$\sigma^2 = \frac{1}{n - 1} \sum_{i=1}^n (x_i - \bar{x})^2$$

- Why do we divide by $n - 1$?

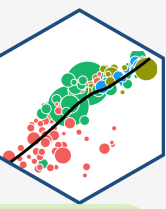
Standard Deviation (Sample)



- Square root the sample variance to get the **sample standard deviation** (s), the average deviation from the *sample* mean (in same units as x)

$$s = \sqrt{s^2} = \sqrt{\frac{1}{n-1} \sum_{i=1}^n (x_i - \bar{x})^2}$$

Sample Standard Deviation: Example



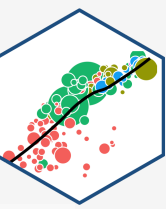
Example: Calculate the sample standard deviation for the following series:

$\{2, 4, 6, 8, 10\}$

```
sd(c(2,4,6,8,10))
```

```
## [1] 3.162278
```

The Steps to Calculate `sd()`, Coded I



```
# first let's save our data in a tibble  
sd_example<-tibble(x=c(2,4,6,8,10))
```

```
# first find the mean (just so we know)
```

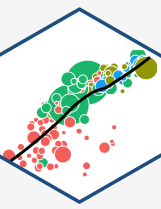
```
sd_example %>%  
  summarize(mean(x))
```

```
## # A tibble: 1 x 1  
##   `mean(x)`  
##     <dbl>  
## 1         6
```

```
# now let's make some more columns:
```

```
sd_example <- sd_example %>%  
  mutate(deviations = x-mean(x), # take deviations from mean  
         deviations_sq = deviations^2) # square them
```

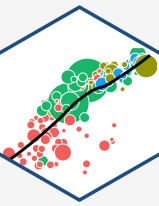
The Steps to Calculate `sd()`, Coded II



```
sd_example # see what we made
```

```
## # A tibble: 5 x 3
##       x deviations deviations_sq
##   <dbl>     <dbl>         <dbl>
## 1     2        -4             16
## 2     4        -2              4
## 3     6         0              0
## 4     8         2              4
## 5    10         4             16
```

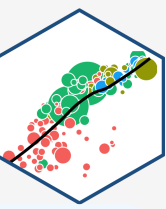
The Steps to Calculate `sd()`, Coded III



```
sd_example %>%  
  # sum the squared deviations  
  summarize(sum_sq_devs = sum(deviations_sq),  
            # divide by n-1 to get variance  
            variance = sum_sq_devs/(n()-1),  
            # square root to get sd  
            std_dev = sqrt(variance))
```

```
## # A tibble: 1 x 3  
##   sum_sq_devs variance std_dev  
##   <dbl>      <dbl> <dbl>  
## 1         40         10   3.16
```

Sample Standard Deviation: You Try



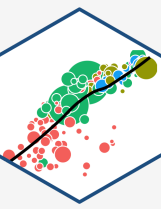
You Try: Calculate the sample standard deviation for the following series:

$\{1, 3, 5, 7\}$

```
sd(c(1,3,5,7))
```

```
## [1] 2.581989
```

Descriptive Statistics: Populations vs. Samples



Population parameters

- **Population size:** N
- **Mean:** μ

- **Variance:** $\sigma^2 = \frac{1}{N} \sum_{i=1}^N (x_i - \mu)^2$

- **Standard deviation:** $\sigma = \sqrt{\sigma^2}$

Sample statistics

- **Population size:** n
- **Mean:** \bar{x}

- **Variance:** $s^2 = \frac{1}{n-1} \sum_{i=1}^n (x_i - \bar{x})^2$

- **Standard deviation:** $s = \sqrt{s^2}$