

# DAY 3: DATA VISUALIZATION

BSTA 511/611, OHSU

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# GOALS FOR TODAY

- **Exploratory Data Analysis (EDA)**  
(Sections 1.4, 1.5, 1.6, 1.7.1)
  - Data visualization with ggplot
    - numerical & categorical variables, and relationships between variables
  - Summarizing numerical data
  - Frequency (two-way) tables
- Some **data wrangling** techniques along the way



Artwork by @allison\_horst

# INTERNATIONAL DAY OF WOMEN IN STATISTICS AND DATA SCIENCE

Tuesday, October 10, 2022

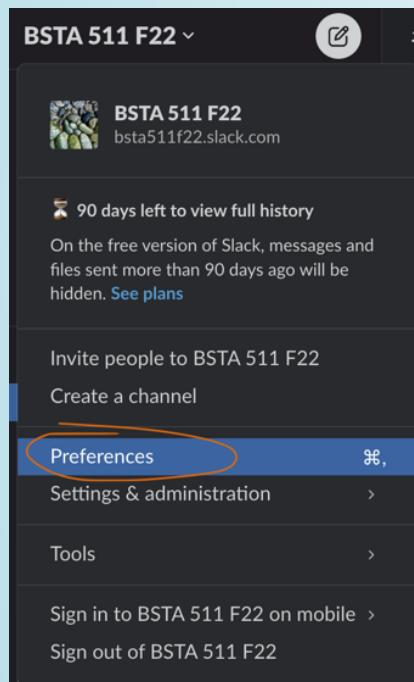
12 am - 11:59 pm UTC (5pm 10/9 to 4:59 pm 10/10 here)



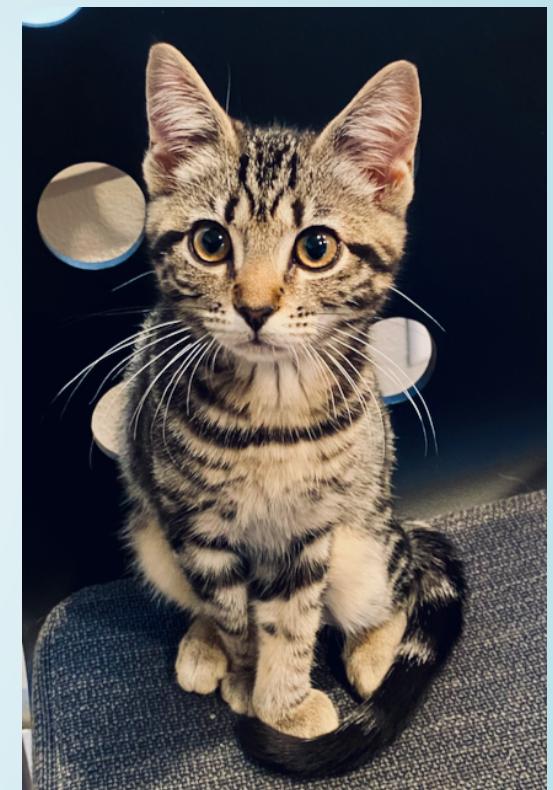
International Day of Women in Statistics and Data Science

# MIMI'S TIP OF THE DAY: SENDING MESSAGES IN SLACK

Are you frustrated that Slack sends a message when you press Enter? You can change that!



A screenshot of the Slack Preferences window under the 'Advanced' tab. The 'Input options' section contains two radio button options: 'Send the message' (unchecked) and 'Start a new line (use ⌘ Enter to send)' (checked). An orange oval highlights the checked option. A blue arrow points from the 'Preferences' button in the sidebar to this section in the preferences window.

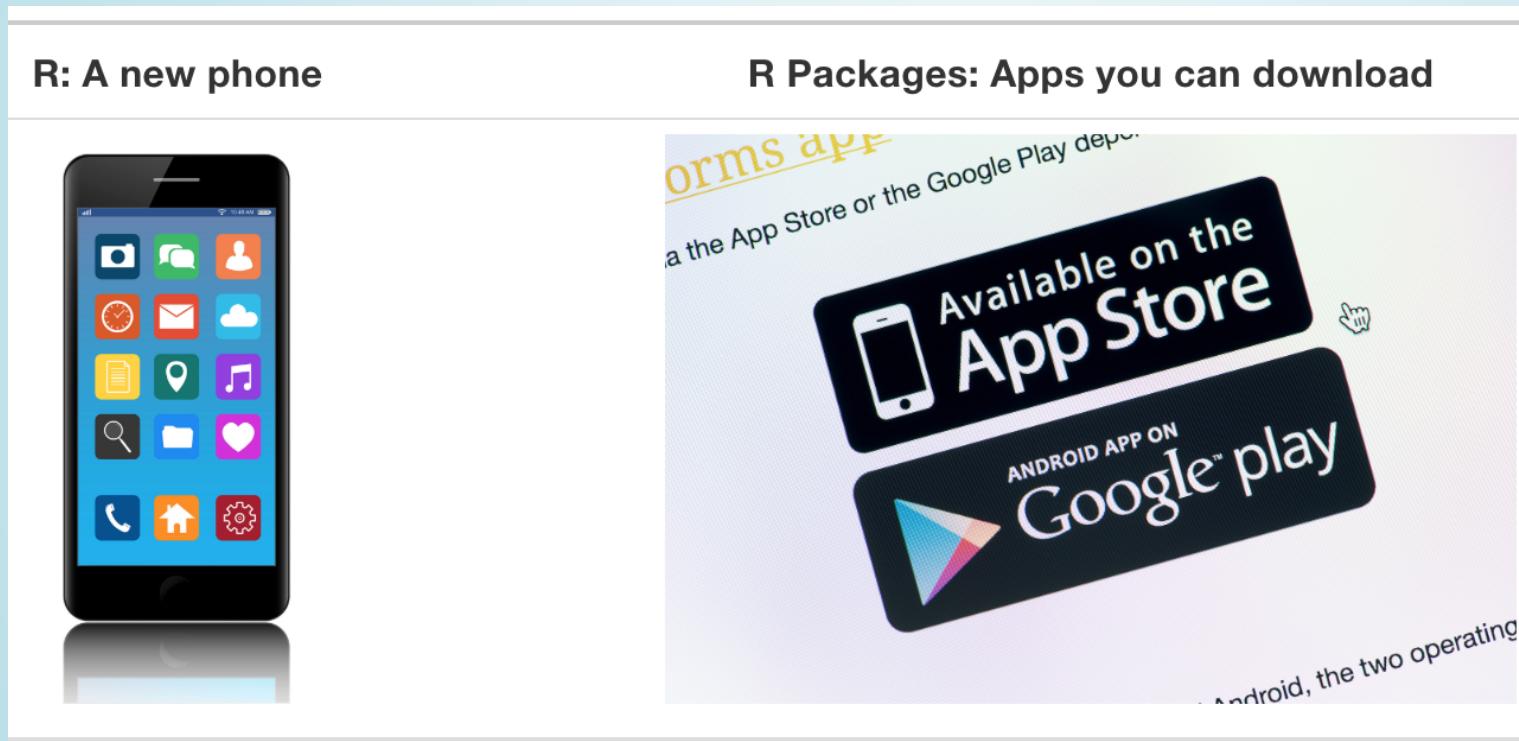


# RECAP OF LAST TIME

- (1.3) **Data collection principles**
  - Population vs. sample
  - Sampling methods
  - Experiments vs. Observational studies
- (1.2) **Intro to Data**
  - Data types
    - Numerical: discrete (integer in R), continuous (double or numeric in R)
    - Categorical: ordinal, nominal
      - character or factor in R
  - How are data stored in R? data frames, tibbles
  - Working with data in R: `dim()`, `nrow()`, `ncol()`, `names()`, `str()`, `summary()`, `head()`, `tail()`, `$`
- (1.4) **Summarizing numerical data**
  - `mean()`, `median()`, `sd()`, `quantile()`

# R PACKAGES

A good analogy for R packages is that they are like apps you can download onto a mobile phone:



ModernDive Figure 1.4

# FROM LAST TIME: INSTALL THE PACAKGES LISTED BELOW

- **knitr**
  - this might actually already be installed
  - check your packages list
- **tidyverse**
  - this is actually a bundle of packages
  - *Warning: it will take a while to install!!!*
  - see more info at <https://tidyverse.tidyverse.org/>
- **rstatix**
  - for summary statistics of a dataset
- **janitor**
  - for cleaning and exploring data
- **ggridges**
  - for creating ridgeline plots
- **devtools**
  - used to create R packages
  - for our purposes, needed to install some packages
- **oi\_biostat\_data**
  - this package is on github
  - **see the next slide for directions on how to install [oi\\_biostat\\_data](#)**

# DIRECTIONS FOR INSTALLING PACKAGE *oibio***stat**

- The textbook's datasets are in the R package **oibio***stat*
- Explanation of code below
  - Installation of **oibio***stat* package requires first installing **devtools** package
  - The code `devtools::install_github()` tells R to use the command `install_github()` from the **devtools** package without loading the entire package and all of its commands (which `library(devtools)` would do).

```
1 install.packages("devtools")
2 devtools::install_github("OI-Biostat/oi_biotest_data", force = TRUE)
```

- After running the code above, put `#` in front of the commands so that RStudio doesn't evaluate them when rendering.
- Now load the **oibio***stat* package
  - the code below needs to be run *every time* you restart R or render a Qmd file

```
1 library(oibio
```

# LOAD PACKAGES WITH `library()` COMMAND

- Tip: **at the top of your Qmd file**, create a chunk that loads all of the R packages you want to use in that file.
- Use the `library()` command to load each required package.
  - Packages need to be reloaded *every* time you open Rstudio.
  - `library()` commands to load needed packages *must* be in the Qmd file

```
1 # run these every time you open Rstudio
2 library(tidyverse)
3 library(oibiotstat)
4 library(ggridges)
5 library(janitor)
6 library(rstatix)
7 library(knitr)
8 library(gtsummary) # NEW!!
```

- You can check whether a package has been loaded or not
  - by looking at the Packages tab and
  - seeing whether it has been checked off or not

# CASE STUDY: DISCRIMINATION IN DEVELOPMENTAL DISABILITY SUPPORT (SECTION 1.7.1)

# CASE STUDY DESCRIPTION

- In the US, individuals with developmental disabilities typically receive services and support from state governments.
  - California allocates funds to developmentally disabled residents through the *Department of Developmental Services (DDS)*
  - Recipients of DDS funds are referred to as “consumers.”
- Dataset `dds.discr`
  - sample of 1,000 DDS consumers (out of a total of ~ 250,000)
  - data include **age, gender, race/ethnicity, and annual DDS financial support per consumer**
- **Previous research**
  - Researchers examined expenditures on consumers by ethnicity
  - Found that the mean annual expenditure on Hispanics was less than that on White non-Hispanics.
- Result: an allegation of ethnic discrimination was brought against the California DDS.
- **Question: Are the data sufficient evidence of ethnic discrimination?**
- See Section 1.7.1 in the textbook for more details

# LOAD `dds.discr` DATASET FROM `oibiostat` PACKAGE

- The textbook's datasets are in the R package `oibiostat`
- Make sure the `oibiostat` package is installed before running the code below.
- Load the `oibiostat` package and the dataset `dds.discr`

**the code below needs to be run *every time you restart R or render a Qmd file***

```
1 library(oibiostat)
2 data("dds.discr")
```

- After loading the dataset `dds.discr` using `data("dds.discr")`, you will see `dds.discr` in the Data list of the Environment window.

# GETTING TO KNOW THE DATASET

```
1 dim(dds.dscr)
[1] 1000      6

1 names(dds.dscr)
[1] "id"          "age.cohort"    "age"           "gender"        "expenditures"
[6] "ethnicity"

1 length(unique(dds.dscr$id)) # How many unique id's are there?
[1] 1000
```

# str() STRUCTURE

- We previously used the base R structure command `str()` to get information about variable types in a dataset.
- Note this dataset is a `tibble` instead of a `data.frame`

```
1 str(dds.dscr)      # base R

tibble [1,000 × 6] (S3: tbl_df/tbl/data.frame)
$ id              : int [1:1000] 10210 10409 10486 10538 10568 10690 10711 10778
10820 10823 ...
$ age.cohort     : Factor w/ 6 levels "0-5","6-12","13-17",...: 3 5 1 4 3 3 3 3 3 3
...
$ age            : int [1:1000] 17 37 3 19 13 15 13 17 14 13 ...
$ gender         : Factor w/ 2 levels "Female","Male": 1 2 2 1 2 1 1 2 1 2 ...
$ expenditures: int [1:1000] 2113 41924 1454 6400 4412 4566 3915 3873 5021 2887
...
$ ethnicity     : Factor w/ 8 levels "American Indian",...: 8 8 4 4 8 4 8 3 8 4 ...
- attr(*, "spec")=
.. cols(
..   ID = col_integer(),
..   `Age Cohort` = col_character(),
..   Age = col_integer(),
..   `Ethnicity` = col_factor(),
..   Sex = col_factor(),
..   Income = col_double()
```

# glimpse()

## New: glimpse()

- Use `glimpse()` from the `tidyverse` package (technically it's from the `dplyr` package) to get information about variable types.
- `glimpse()` tends to have nicer output for `tibbles` than `str()`

```
1 library(tidyverse)
2 glimpse(dds.dscr) # from tidyverse package (dplyr)
```

```
Rows: 1,000
Columns: 6
$ id          <int> 10210, 10409, 10486, 10538, 10568, 10690, 10711, 10778, 1...
$ age.cohort <fct> 13-17, 22-50, 0-5, 18-21, 13-17, 13-17, 13-17, 13-17, 13-...
$ age         <int> 17, 37, 3, 19, 13, 15, 13, 17, 14, 13, 13, 14, 15, 17, 20...
$ gender      <fct> Female, Male, Male, Female, Male, Female, Female, Male, F...
$ expenditures <int> 2113, 41924, 1454, 6400, 4412, 4566, 3915, 3873, 5021, 28...
$ ethnicity   <fct> White not Hispanic, White not Hispanic, Hispanic, Hispani...
```

# summary()

- We previously used the base R structure command `summary()` to get summary information about variables

```
1 summary(dds.dscr)      # base R
```

	id	age.cohort	age	gender	expenditures
Min.	:10210	0-5 : 82	Min.   : 0.0	Female:503	Min.   : 222
1st Qu.	:31809	6-12 :175	1st Qu.:12.0	Male   :497	1st Qu.: 2899
Median	:55384	13-17:212	Median  :18.0		Median : 7026
Mean	:54663	18-21:199	Mean    :22.8		Mean   :18066
3rd Qu.	:76135	22-50:226	3rd Qu.:26.0		3rd Qu.:37713
Max.	:99898	51+  :106	Max.   :95.0		Max.   :75098

	ethnicity
White not Hispanic	:401
Hispanic	:376
Asian	:129
Black	: 59
Multi Race	: 26
American Indian	:  4
Other	:  1

# tbl\_summary():SUMMARY TABLE

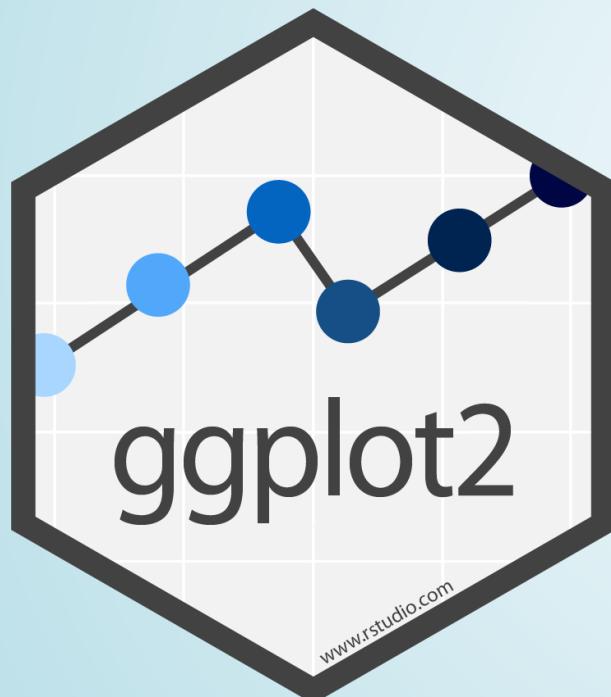
- **New:** Use `tbl_summary()` from the `gtsummary` package to get summary information

```
1 # library(gtsummary)
2 tbl_summary(dds.dscr)
```

Characteristic	N = 1,000 <sup>7</sup>
id	55,385 (31,809, 76,135)
age.cohort	
0-5	82 (8.2%)
6-12	175 (18%)
13-17	212 (21%)
18-21	199 (20%)
22-50	226 (23%)
51+	106 (11%)
age	18 (12, 26)
gender	
Female	503 (50%)
Male	497 (50%)
expenditures	7,026 (2,899, 37,713)
ethnicity	
American Indian	4 (0.4%)
Asian	129 (13%)
Black	59 (5.9%)
Hispanic	376 (38%)
Multi Race	26 (2.6%)
Native Hawaiian	3 (0.3%)
Other	2 (0.2%)
White not Hispanic	401 (40%)

<sup>7</sup> Median (IQR); n (%)

# VISUALIZE NUMERICAL VARIABLES WITH `ggplot`

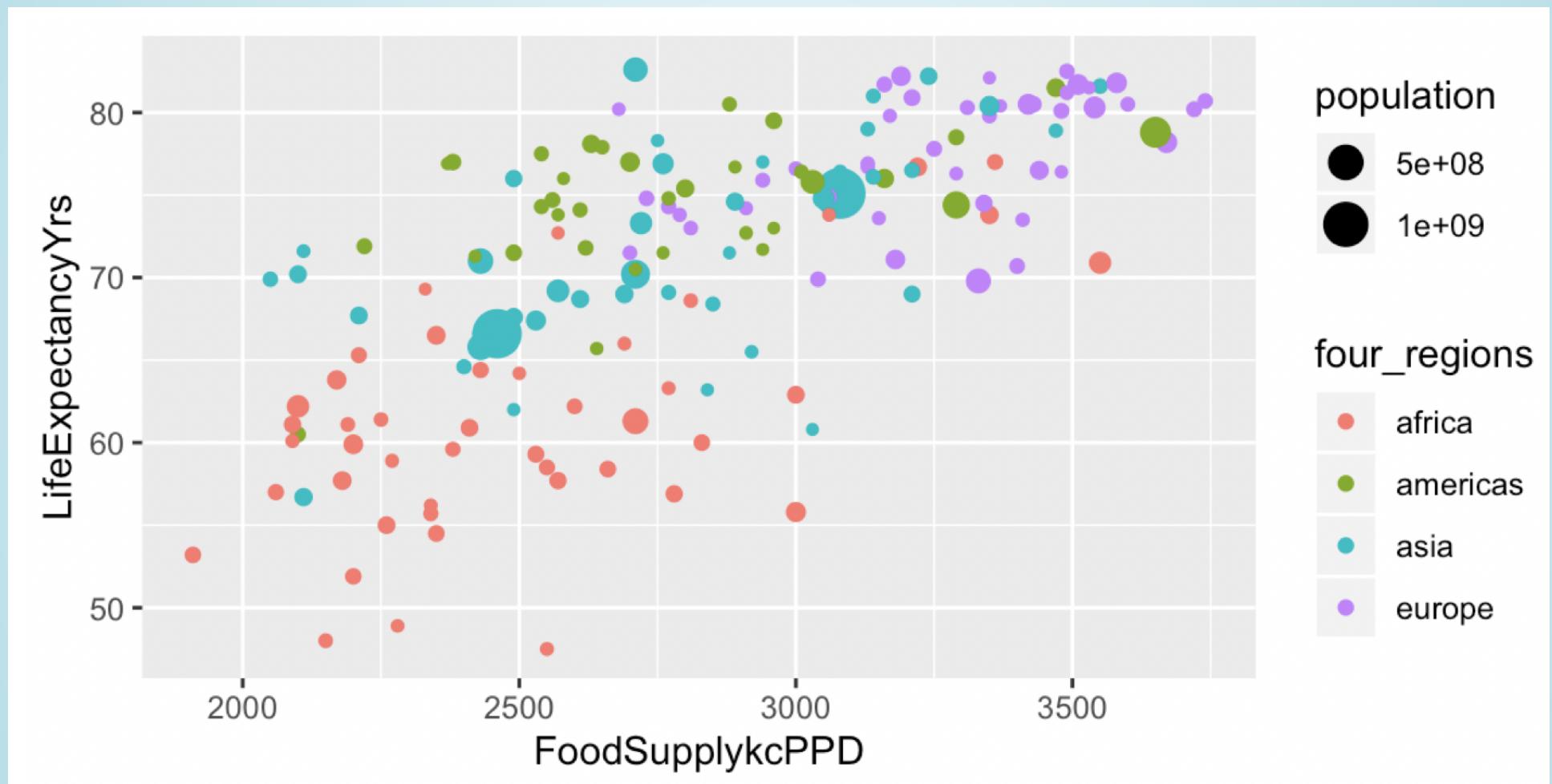


ggplot



Artwork by @allison\_horst

# WHAT DATA (VARIABLES) ARE INCLUDED IN THE PLOT BELOW?



# BASICS OF A GG PLOT

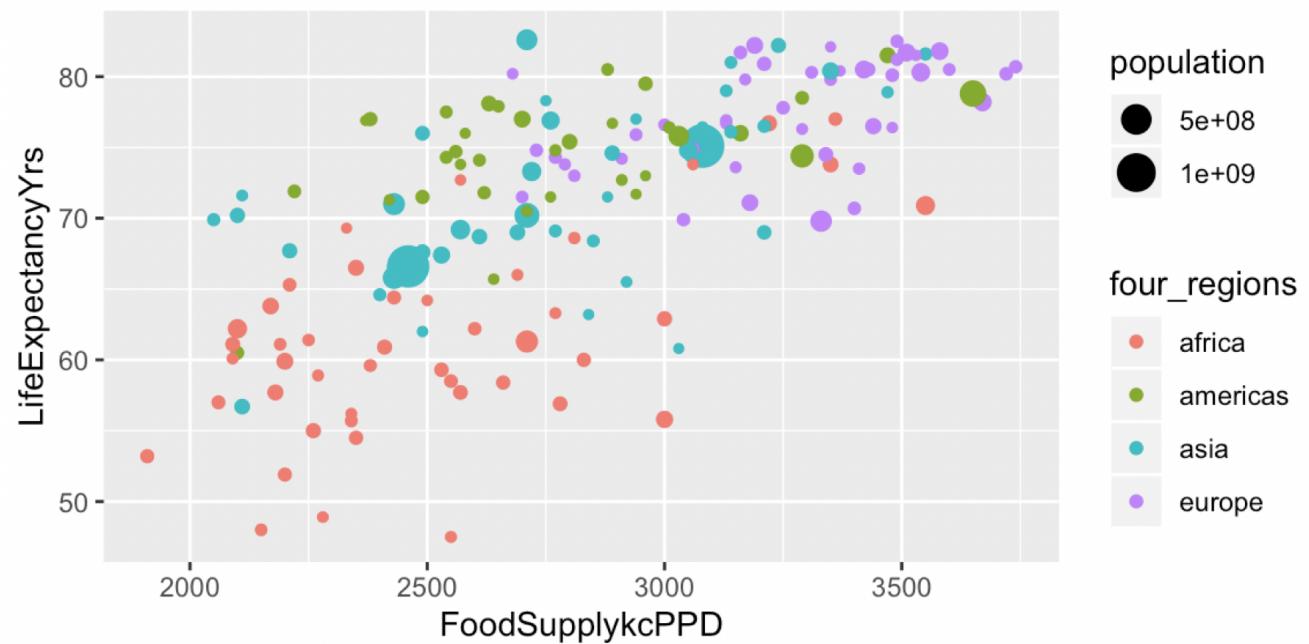
Function

```
ggplot(data = gapminder2011,  
       aes(x = FoodSupplykcPPD, y = LifeExpectancyYrs,  
            color = four_regions, size = population)) +  
       geom_point()
```

What kind of plot to make

Dataset

Which variables to plot



# GRAMMAR OF GG PLOT2

## 1. Tidy Data

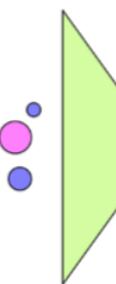
gdp	lifexp	pop	continent
340	65	31	Euro
227	51	200	Amer
909	81	80	Euro
126	40	20	Asia

```
ggplot(data = gapminder, mapping =  
       aes(x = gdp,  
            y = lifespan,  
            color = continent,  
            size = pop))
```

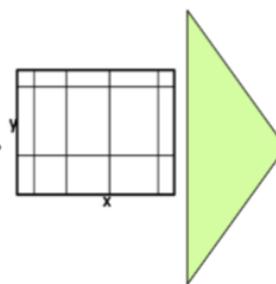
## 2. Mapping



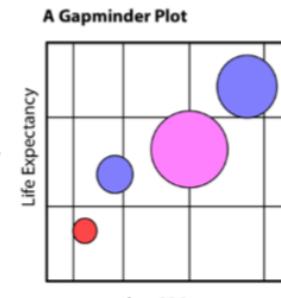
## 3. Geom



## 4. Co-ordinates, Scales

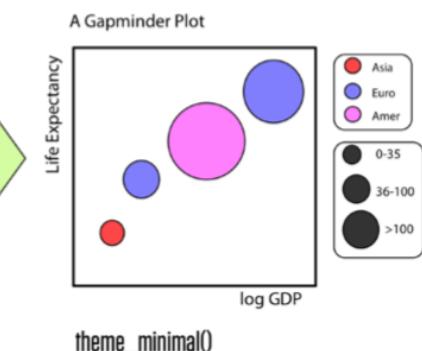


## 5. Labels & Guides



```
labs()  
guides()
```

## 6. Themes

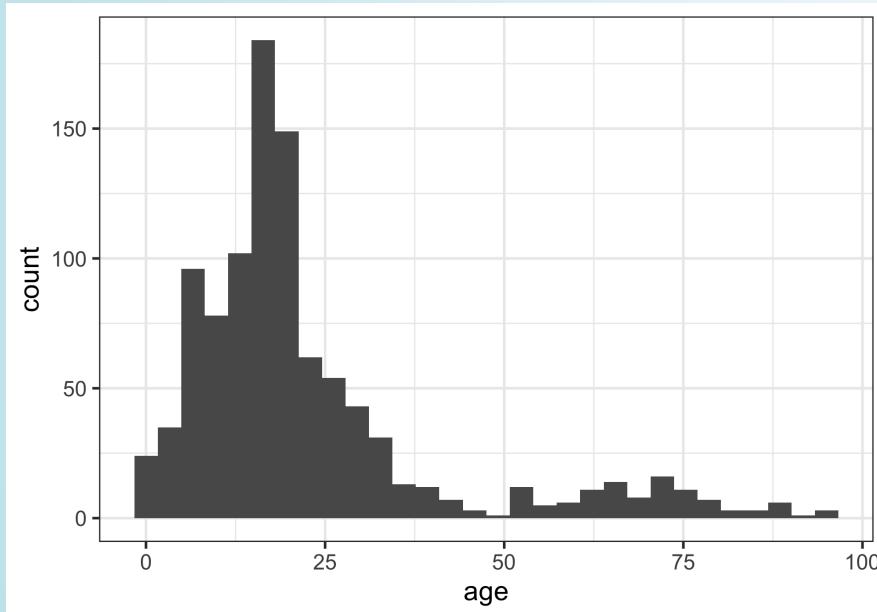


Kieran Healy

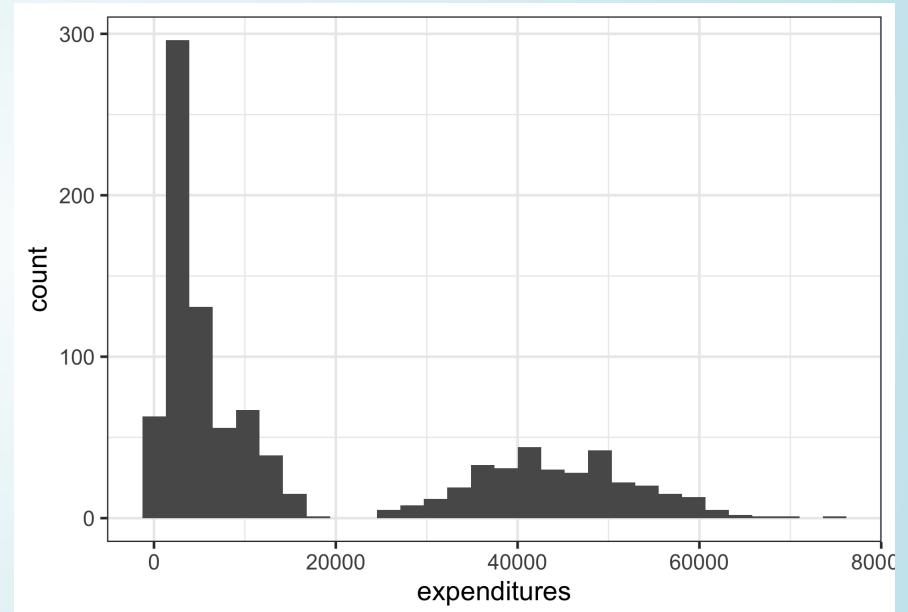
# HISTOGRAMS

What is being measured on the vertical axes?

```
1 ggplot(data = dds.discr,  
2         aes(x = age)) +  
3         geom_histogram()
```

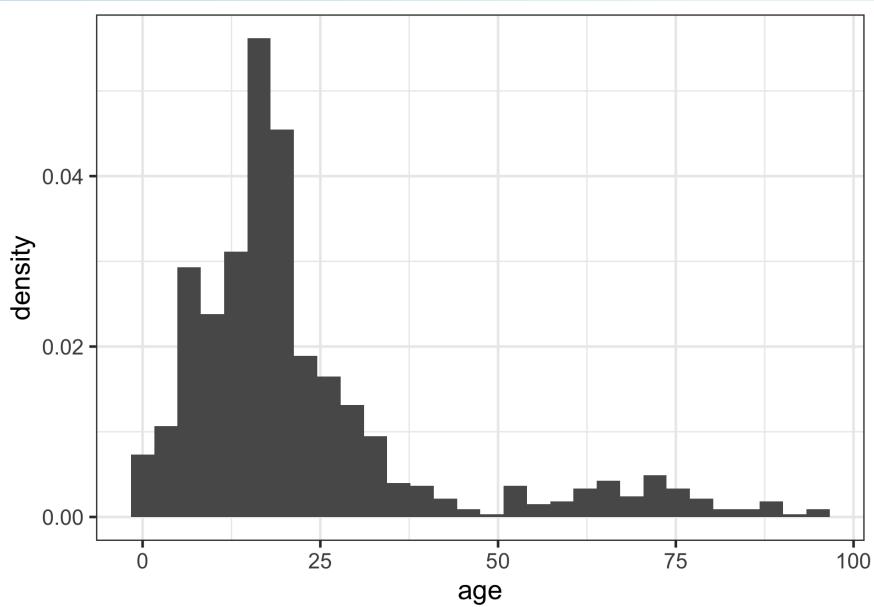


```
1 ggplot(data = dds.discr,  
2         aes(x = expenditures)) +  
3         geom_histogram()
```

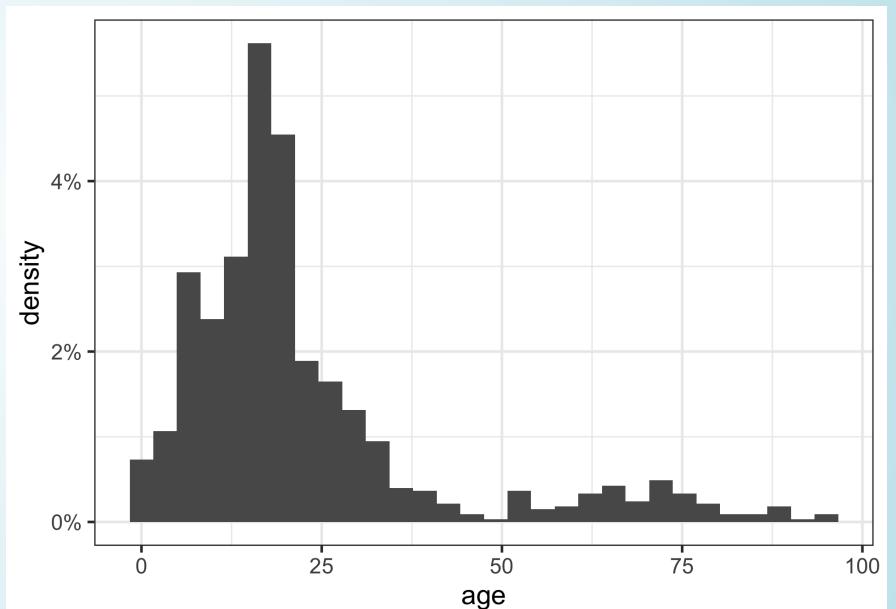


# HISTOGRAMS SHOWING PROPORTIONS

```
1 ggplot(data = dds.discr,
2         aes(x = age)) +
3   geom_histogram(
4     aes(y = stat(density)))
```



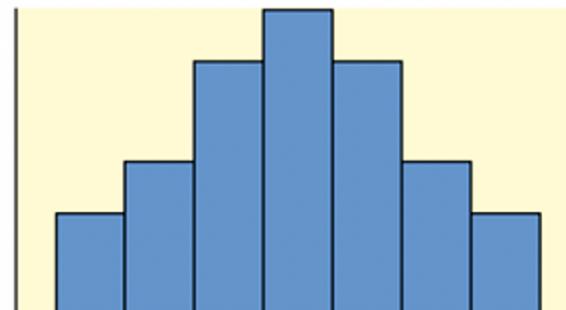
```
1 ggplot(data = dds.discr,
2         aes(x = age)) +
3   geom_histogram(
4     aes(y = stat(density))) +
5   scale_y_continuous(labels =
6     scales::percent_format())
```



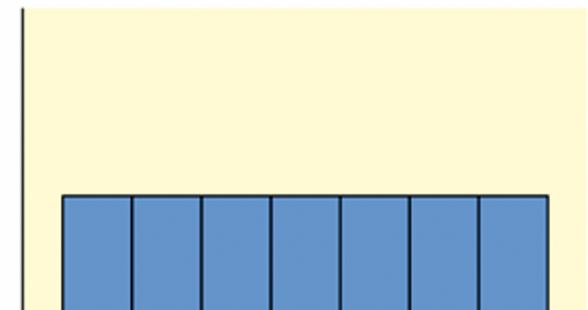
# DISTRIBUTION SHAPES

Common  
distribution  
shapes

symmetric

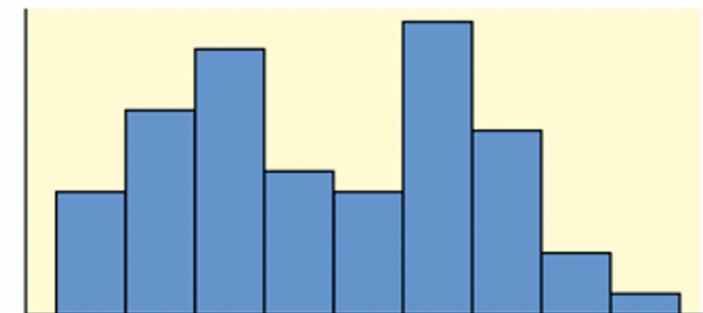
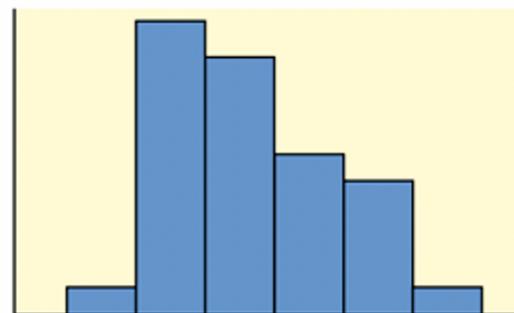
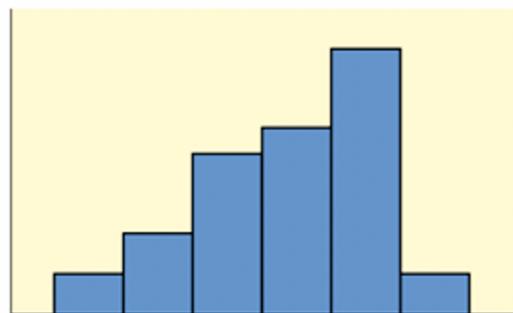


uniform



skewed left  
(negative)

skewed right  
(positive)

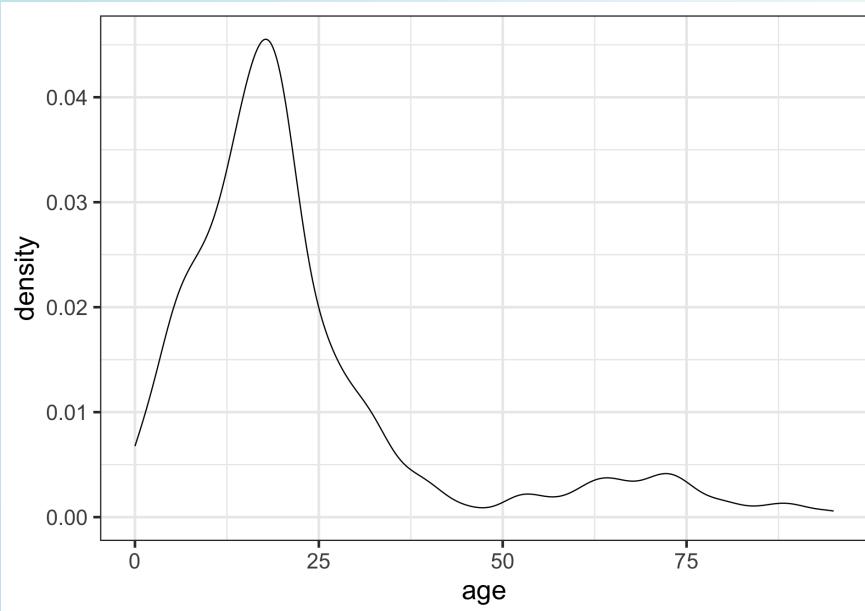


bimodal

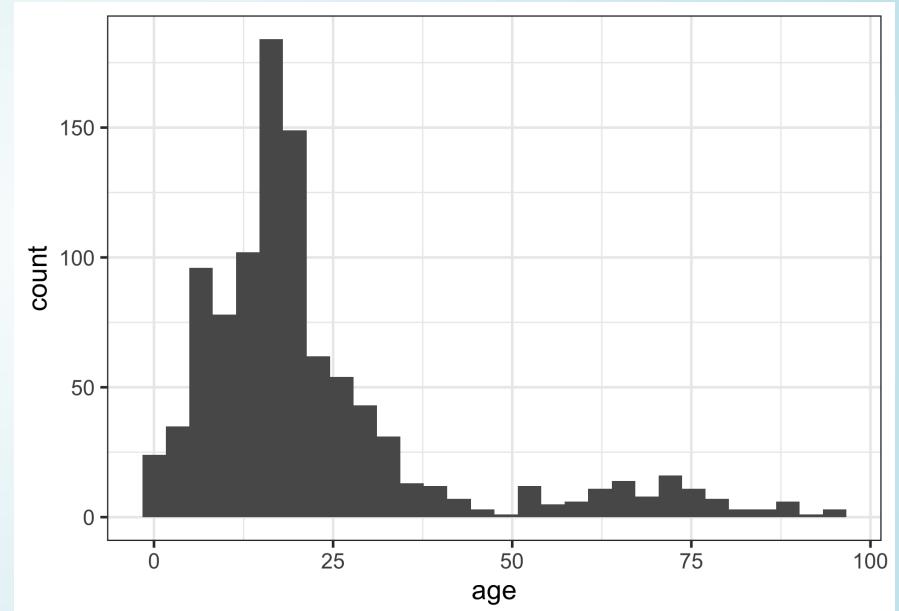
# DENSITY PLOTS

What is being measured on the vertical axes?

```
1 ggplot(data = dds.discr,  
2         aes(x = age)) +  
3         geom_density()
```



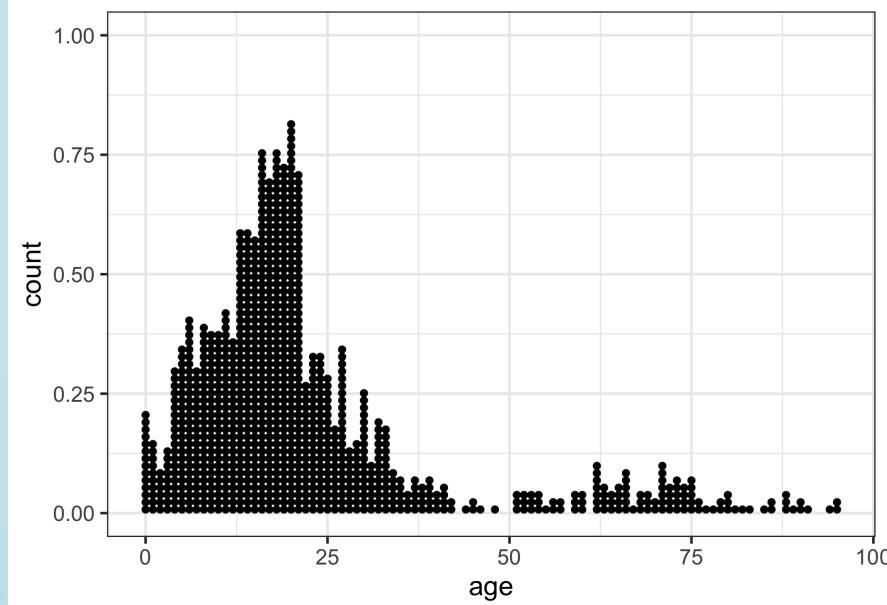
```
1 ggplot(data = dds.discr,  
2         aes(x = age)) +  
3         geom_histogram()
```



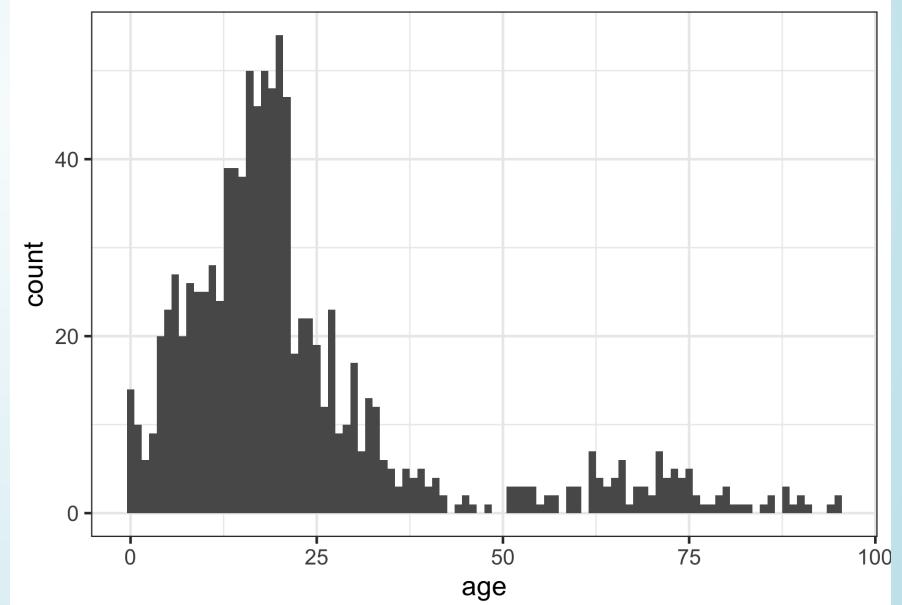
# DOT PLOTS

- Better for smaller samples
- What is being measured on the vertical axes?

```
1 ggplot(data = dds.dscr,  
2         aes(x = age)) +  
3         geom_dotplot(binwidth =1)
```

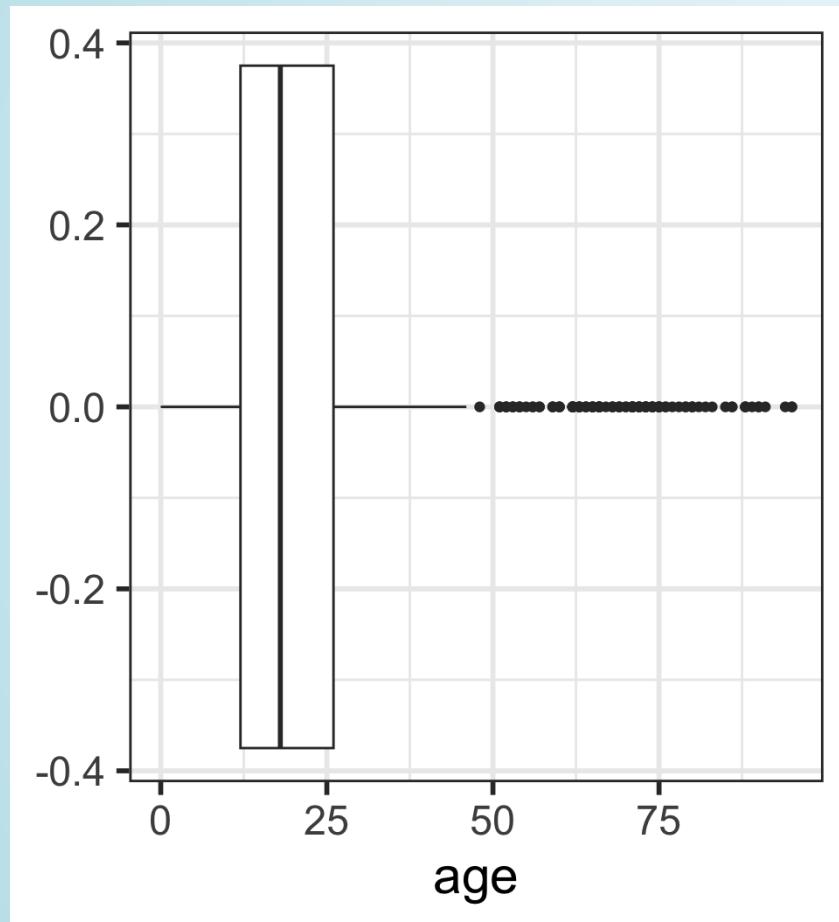


```
1 ggplot(data = dds.dscr,  
2         aes(x = age)) +  
3         geom_histogram(binwidth =1)
```

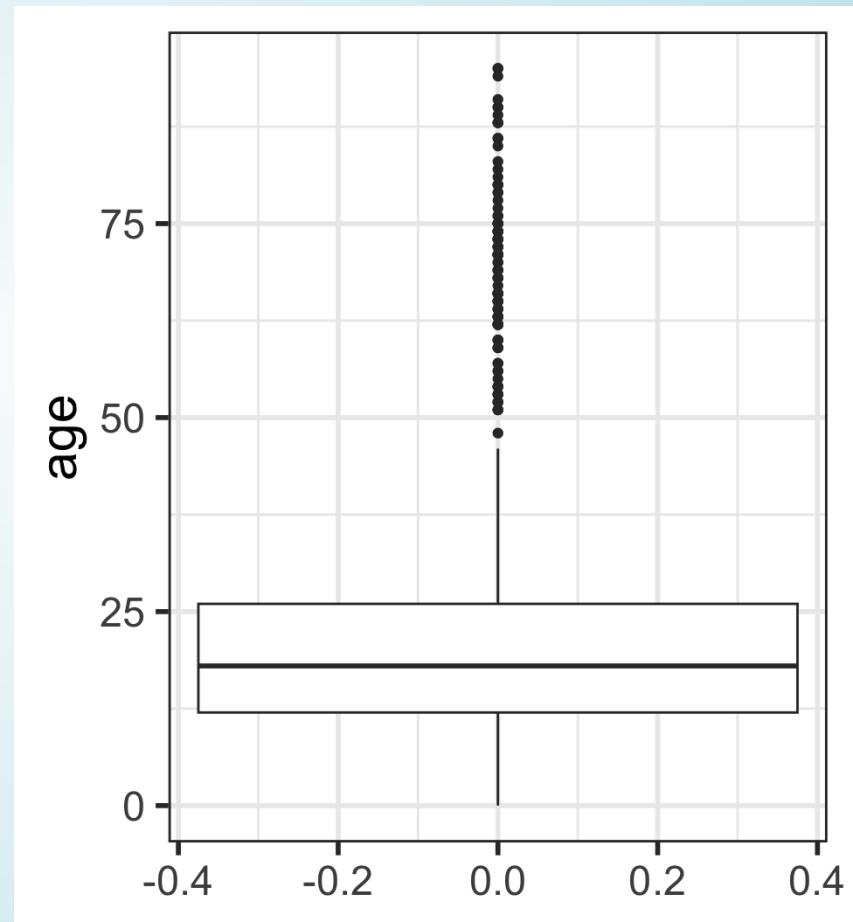


# BOXPLOTS

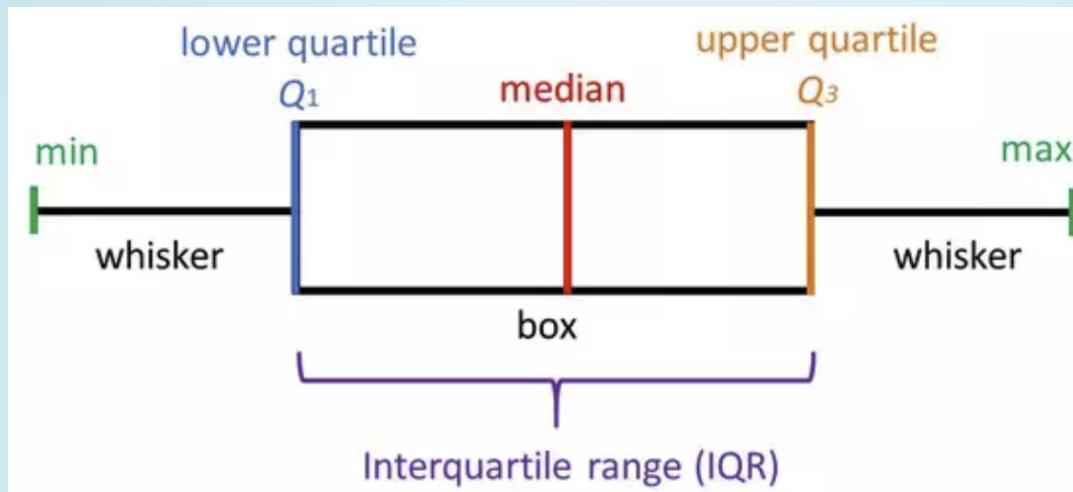
```
1 ggplot(data = dds.discr,  
2         aes(x = age)) +  
3     geom_boxplot()
```



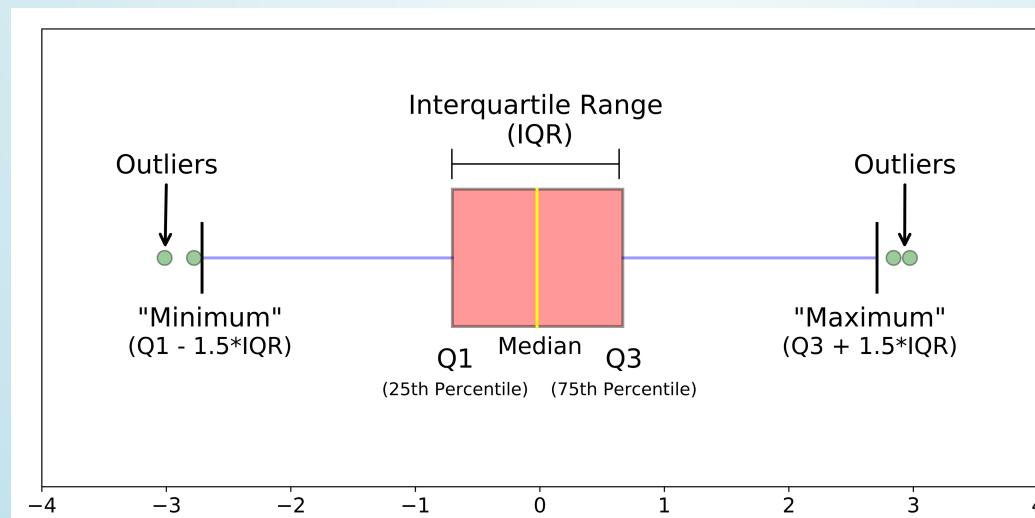
```
1 ggplot(data = dds.discr,  
2         aes(y = age)) +  
3     geom_boxplot()
```



# BOXPLOTS: 5 NUMBER SUMMARY VISUALIZATION



No outliers:



With outliers:

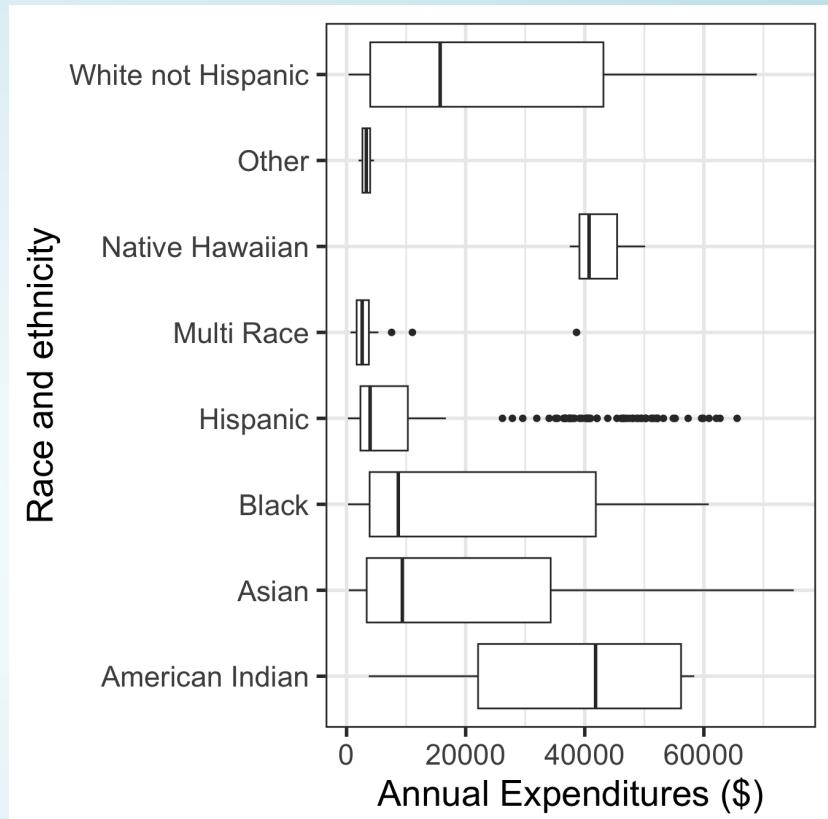
# VISUALIZING RELATIONSHIPS BETWEEN NUMERICAL AND CATEGORICAL VARIABLES (1.6.3)

# SIDE-BY-SIDE BOXPLOTS

```
1 ggplot(data = dds.discr,
2         aes(x = expenditures,
3               y = ethnicity)) +
4   geom_boxplot() +
5   labs(x = "Annual Expenditures ($)",
6        y = "Race and ethnicity")
```

Can you determine the following using boxplots?

- distribution shape
- sample size

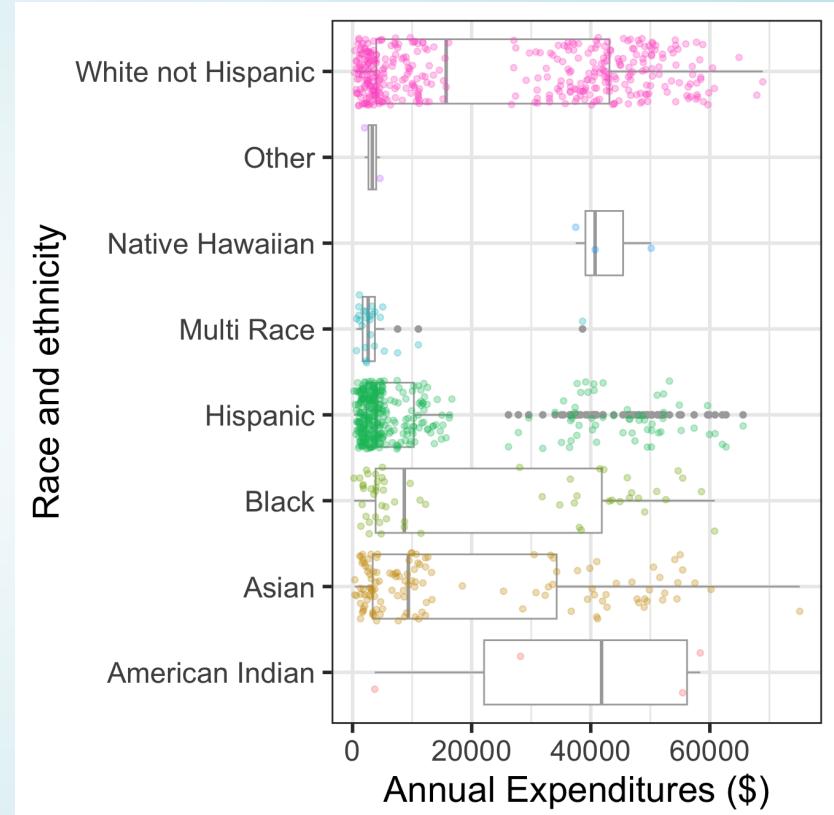


# SIDE-BY-SIDE BOXPLOTS WITH DATA POINTS

```
1 ggplot(data = dds.dscr,
2         aes(x = expenditures,
3               y = ethnicity)) +
4   geom_boxplot(color="darkgrey") +
5   labs(x = "Annual Expenditures ($)",
6        y = "Race and ethnicity") +
7   geom_jitter(
8     aes(color = ethnicity),
9     alpha = 0.3,
10    show.legend = FALSE,
11    position = position_jitter(
12      height = 0.4))
```

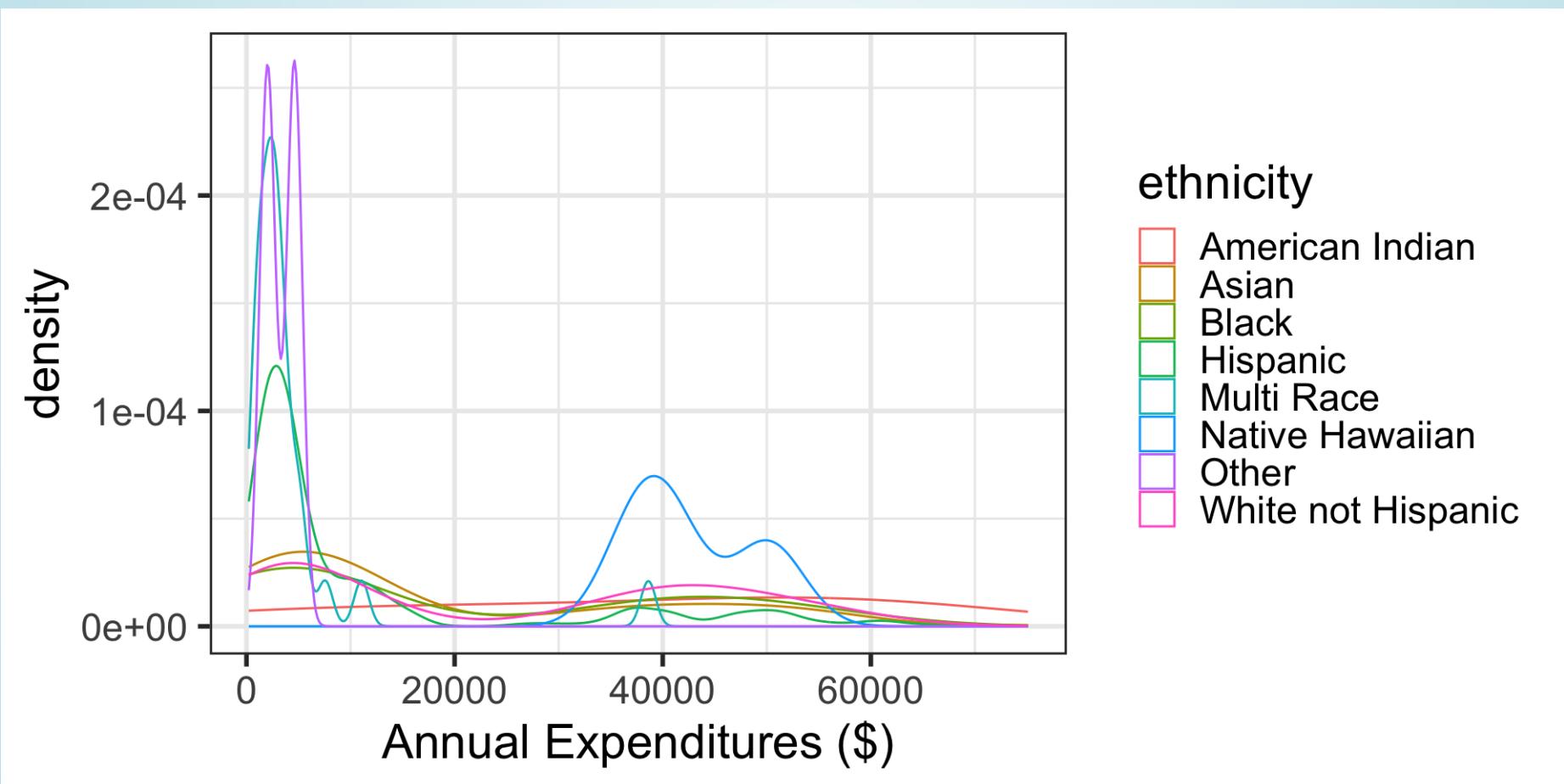
Can you determine the following using boxplots?

- distribution shape
- sample size



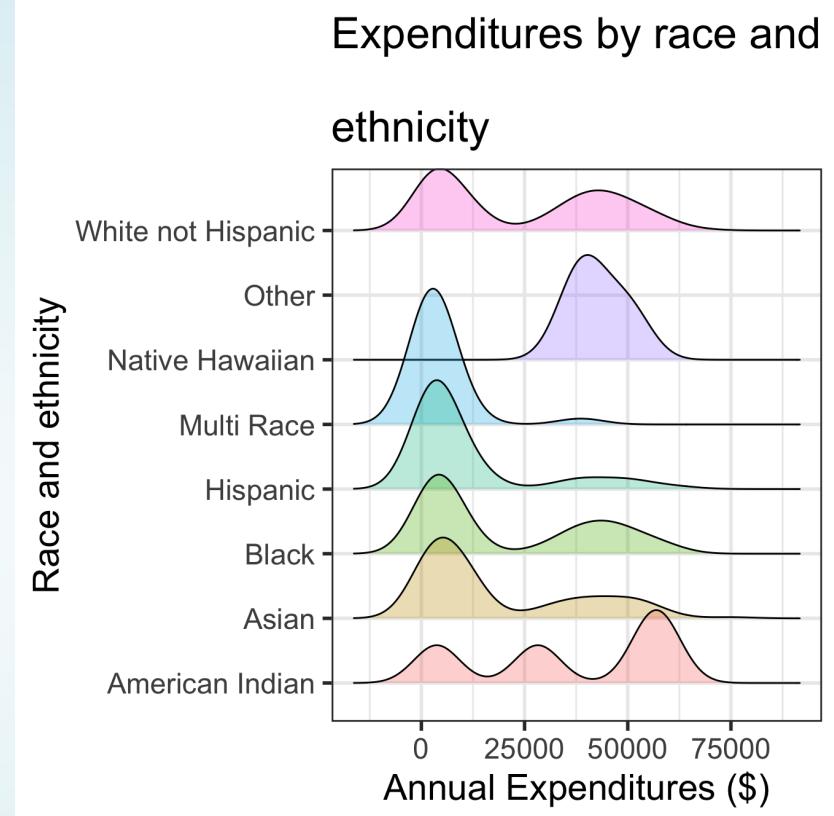
# DENSITY PLOTS BY GROUP

```
1 ggplot(data = dds.discr,
2         aes(x = expenditures,
3               color = ethnicity)) +
4   geom_density() +
5   labs(x = "Annual Expenditures ($)")
```



# RIDGE LINE PLOT

```
1 # library(ggridges)
2 ggplot(data = dds.dscr,
3         aes(x = expenditures,
4               y = ethnicity,
5               fill = ethnicity))
6 )
7 geom_density_ridges(
8     alpha = 0.3,
9     show.legend = FALSE) +
10 labs(x = "Annual Expenditures ($)",
11       y = "Race and ethnicity",
12       title =
13       "Expenditures by race and
14       \nethnicity")
```

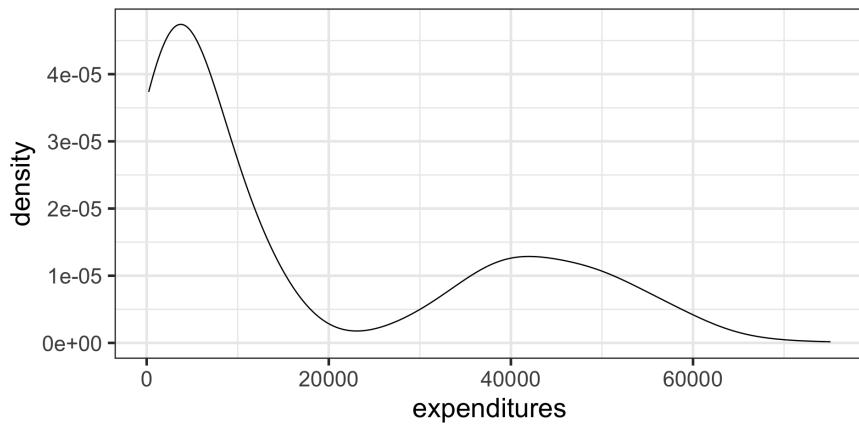


# TRANSFORMING DATA (1.4.5)

- We sometimes apply a transformation to highly skewed data to make it more symmetric
- Log transformations are often used for skewed right data

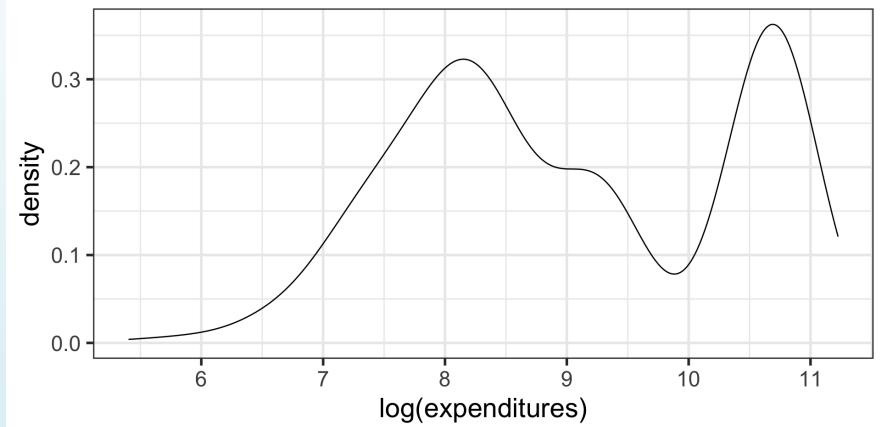
$x = \text{expenditures}$

```
1 ggplot(data = dds.discr,
2         aes(x = expenditures)) +
3     geom_density()
```



$x = \log(\text{expenditures})$

```
1 ggplot(data = dds.discr,
2         aes(x = log(expenditures))) +
3     geom_density()
```



# RELATIONSHIPS BETWEEN TWO NUMERICAL VARIABLES (1.6.1)

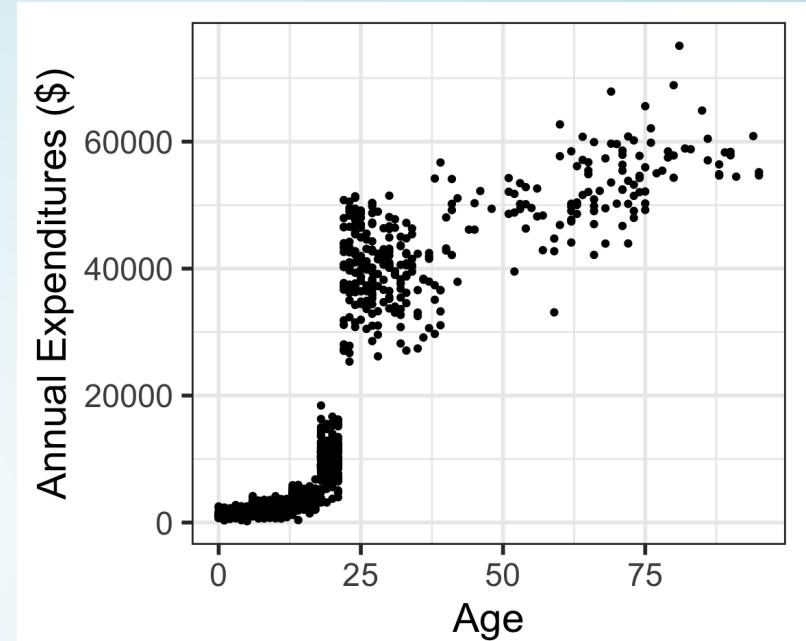
# SCATTERPLOTS

```
1 ggplot(data = dds.discr,
2         aes(x = age,
3               y = expenditures)) +
4   geom_point() +
5   labs(x = "Age",
6        y = "Annual Expenditures ($)")
```

Response vs. explanatory variables  
(Section 1.2.3)

- A **response variable** measures the outcome of interest in a study
- A study will typically examine whether the values of a response variable differ as values of an **explanatory variable** change

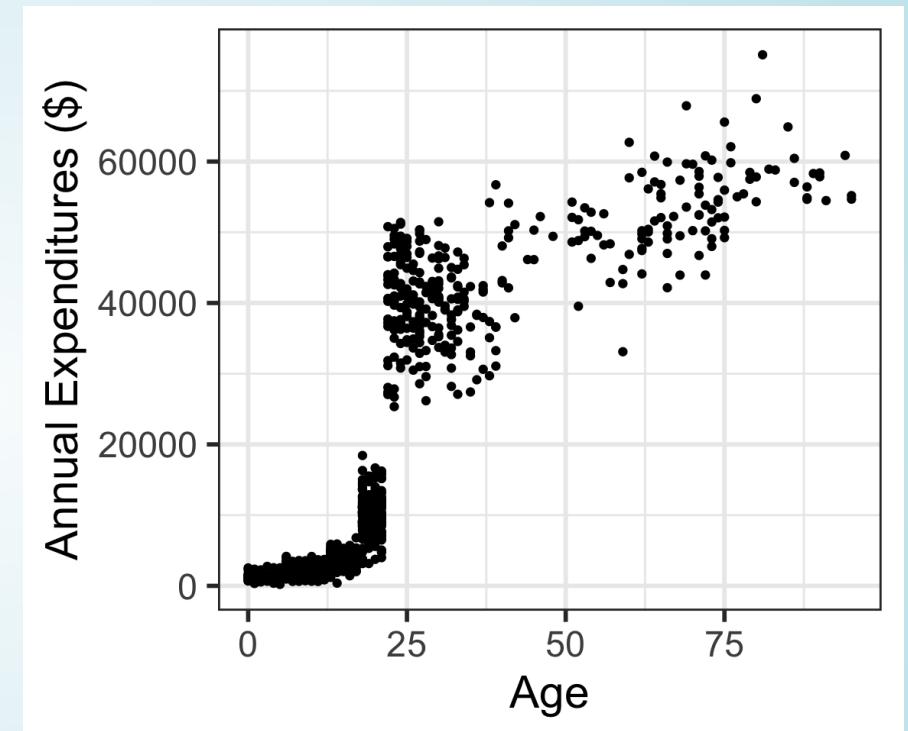
Describe the association between the variables



# DESCRIBING ASSOCIATIONS BETWEEN 2 NUMERICAL VARIABLES

Two variables  $x$  and  $y$  are

- **positively associated** if  $y$  increases as  $x$  increases.
- **negatively associated** if  $y$  decreases as  $x$  increases.
- If there is no association between the variables, then we say they are **uncorrelated** or **independent**.



- The term “association” is a very general term.
  - Can be used for numerical or categorical variables
  - Not specifically referring to linear associations

# (PEARSON) CORRELATION COEFFICIENT $R$

- $r = -1$  indicates a **perfect negative linear relationship**: As one variable increases, the value of the other variable tends to go down, following a *straight line*.
- $r = 0$  indicates **no linear relationship**: The values of both variables go up/down independently of each other.
- $r = 1$  indicates a **perfect positive linear relationship**: As the value of one variable goes up, the value of the other variable tends to go up as well in a linear fashion.
- The closer  $r$  is to  $\pm 1$ , the stronger the linear association.

# (PEARSON) CORRELATION COEFFICIENT (R): FORMULA

The (Pearson) correlation coefficient of variables  $x$  and  $y$  can be computed using the formula

$$r = \frac{1}{n - 1} \sum_{i=1}^n \left( \frac{x_i - \bar{x}}{s_x} \right) \left( \frac{y_i - \bar{y}}{s_y} \right)$$

where

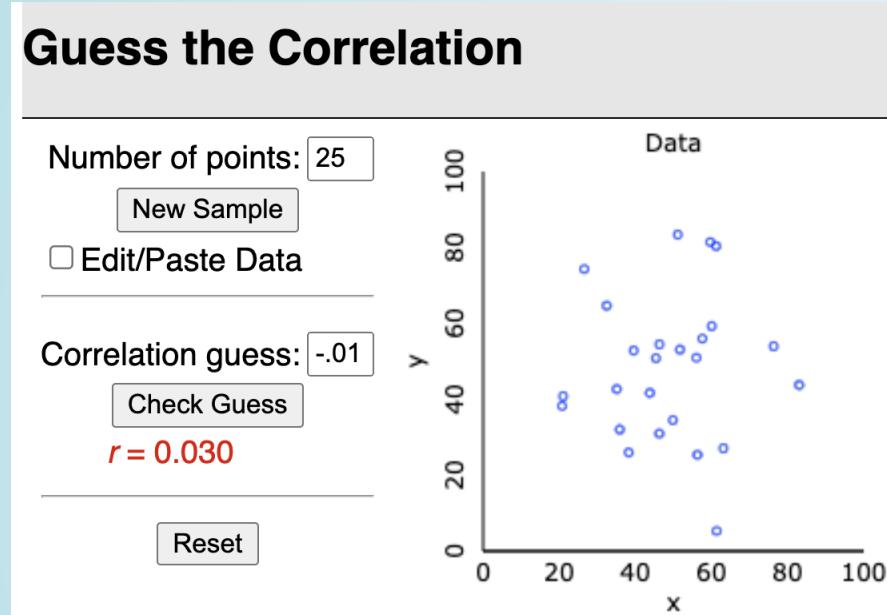
- $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$  are the  $n$  paired values of the variables  $x$  and  $y$
- $s_x$  and  $s_y$  are the sample standard deviations of the variables  $x$  and  $y$ , respectively

```
1 cor(dds.discr$age, dds.discr$expenditures)
```

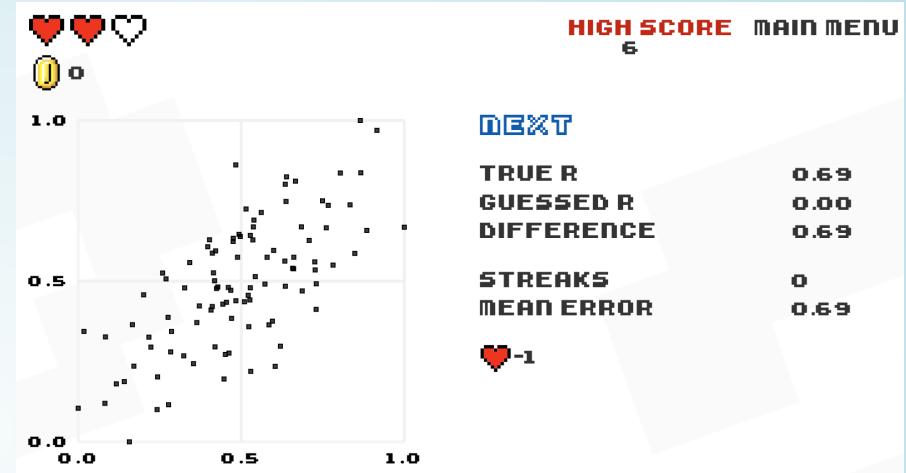
```
[1] 0.8432422
```

# GUESS THE CORRELATION GAME!

Rossmann & Chance's applet



Or, for the Atari-like experience



<http://guessthecorrelation.com/>

Tracks performance of guess vs. actual, error vs. actual, and error vs. trial

<http://www.rossmanchance.com/applets/GuessCorrelation.html>

# SCATTERPLOTS WITH COLOR-CODED DOTS

Describe the association between the variables

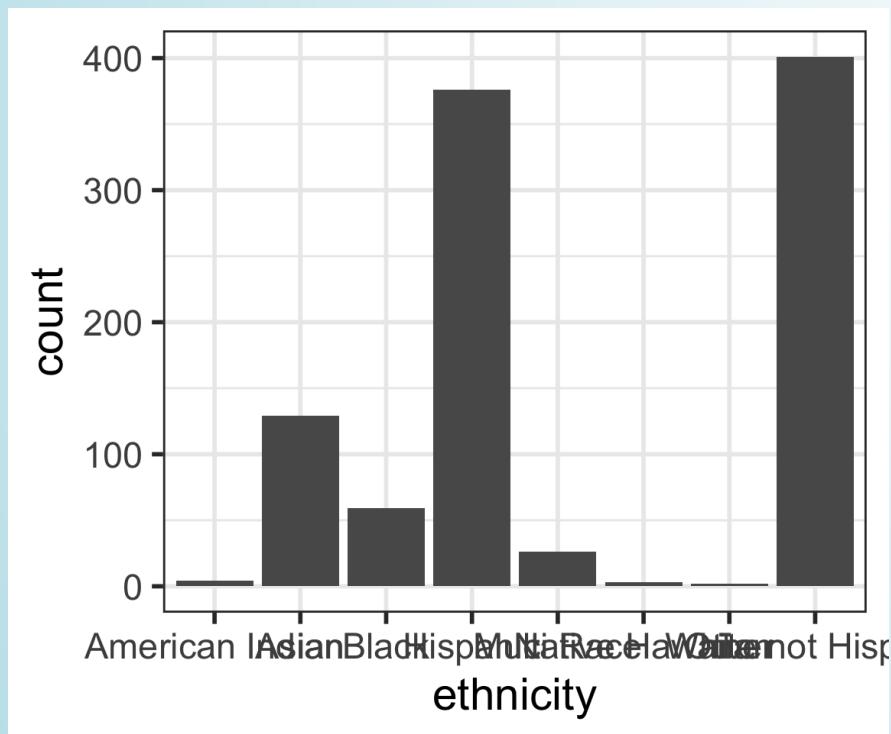
```
1 ggplot(data = dds.dscr,
2         aes(x = age, y = expenditures,
3               color = ethnicity)) +
4   geom_point(alpha = .5) +
5   labs(x = "Age", y = "Annual Expenditures ($)") +
6   theme(legend.position = "bottom")
```

CATEGORICAL DATA (1.5)  
AND RELATIONSHIPS  
BETWEEN TWO  
CATEGORICAL VARIABLES  
(1.6.2)

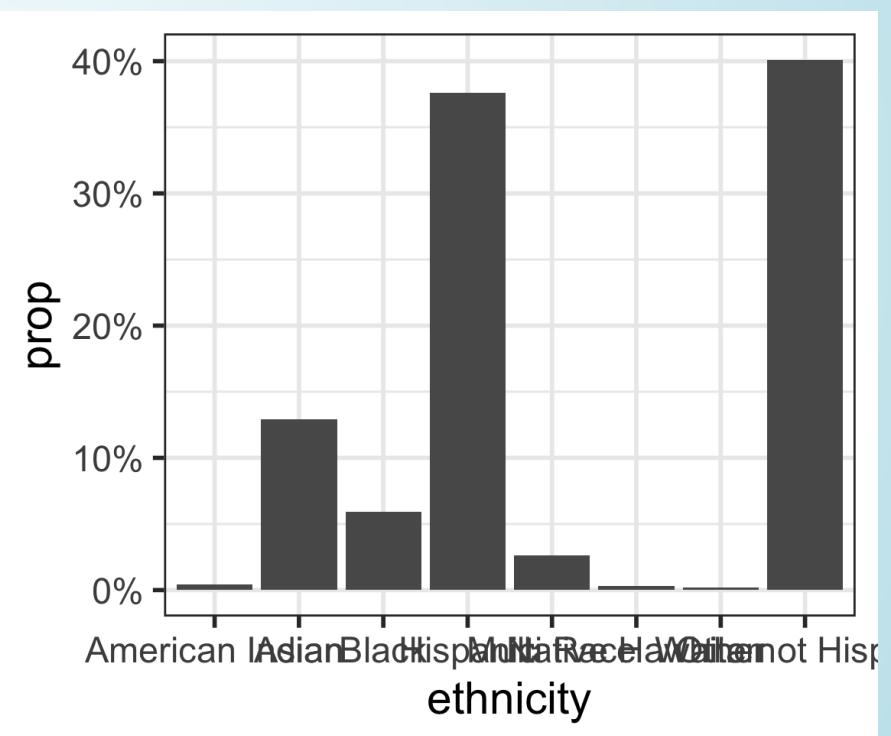
# BARPLOTS

Counts (below) vs.  
percentages (right)

```
1 ggplot(data = dds.dscr,  
2         aes(x = ethnicity)) +  
3     geom_bar()
```



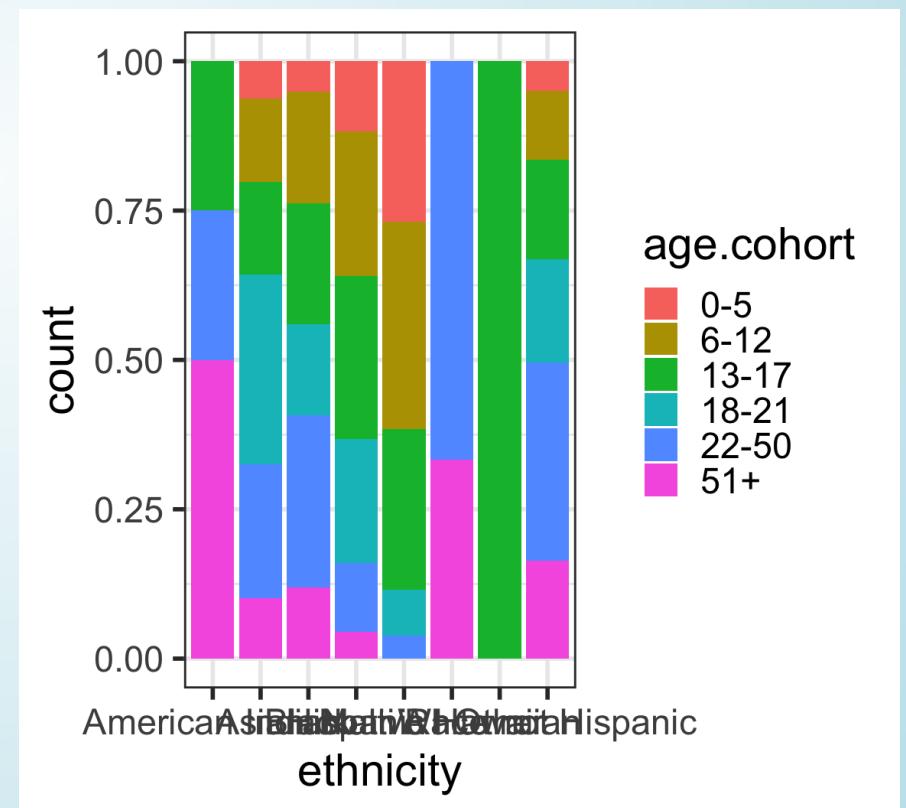
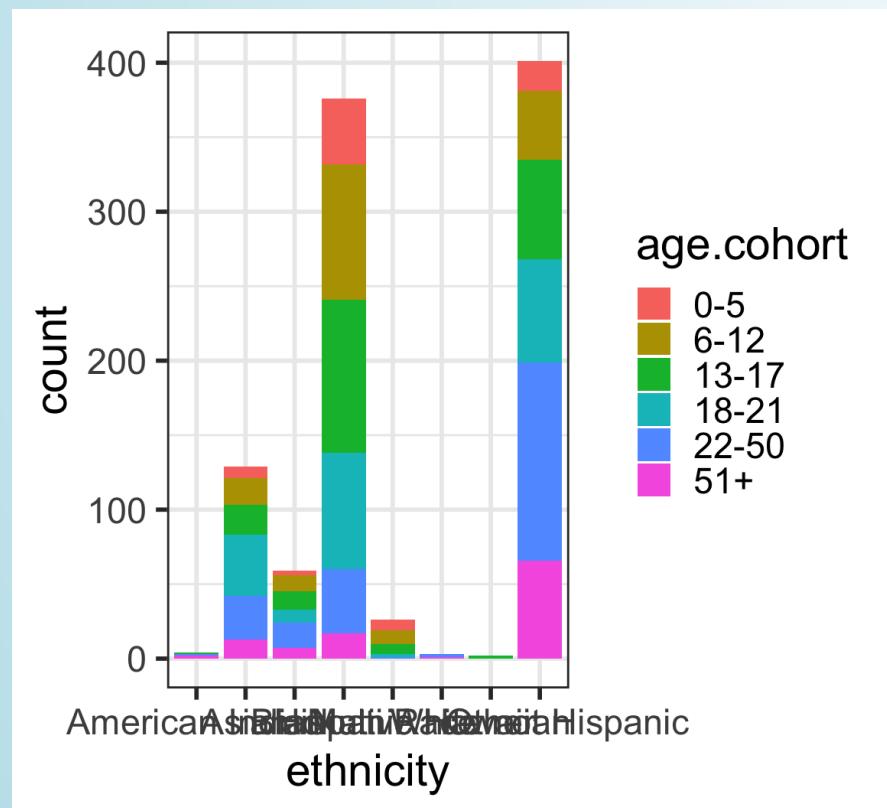
```
1 ggplot(data = dds.dscr,  
2         aes(x = ethnicity)) +  
3     geom_bar(aes(y = stat(prop)),  
4                group = 1)) +  
5     scale_y_continuous(labels =  
6                         scales::percent_format())
```



# BARPLOTS WITH 2 VARIABLES: SEGMENTED BAR PLOTS

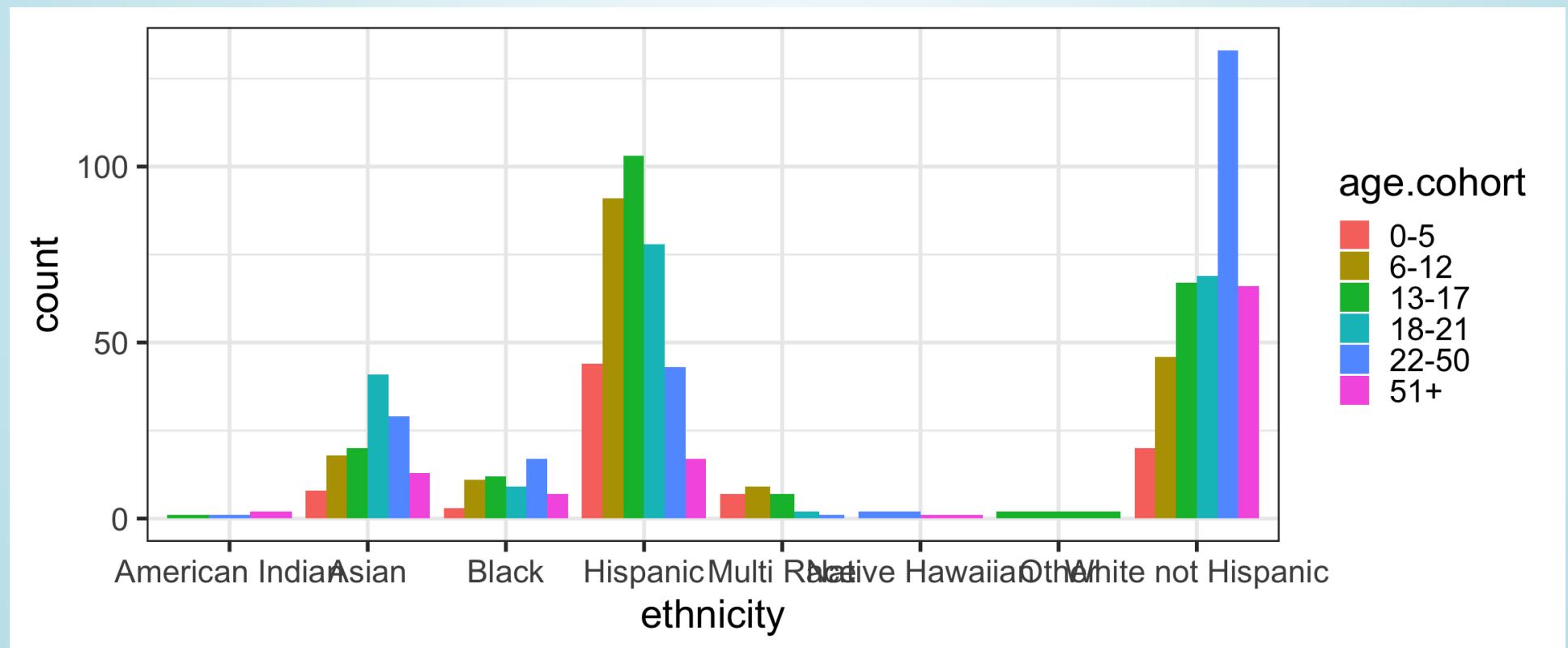
```
1 ggplot(data = dds.dscr,  
2         aes(x = ethnicity,  
3                  fill = age.cohort)) +  
4     geom_bar()
```

```
1 ggplot(data = dds.dscr,  
2         aes(x = ethnicity,  
3                  fill = age.cohort)) +  
4     geom_bar(position = "fill")
```

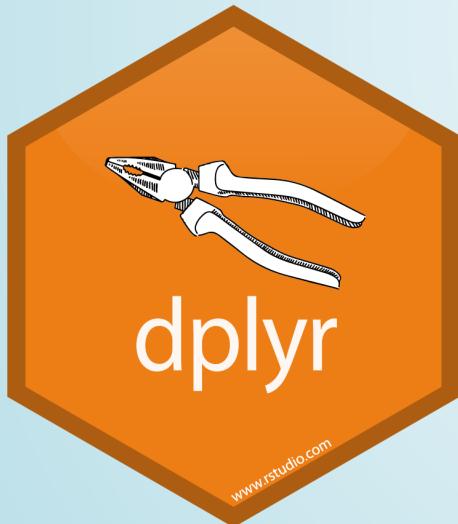


# BARPLOTS WITH 2 VARIABLES: SIDE-BY-SIDE BAR PLOTS

```
1 ggplot(data = dds.dscr,
2         aes(x = ethnicity,
3               fill = age.cohort)) +
4   geom_bar(position = "dodge")
```



# SUMMARIZING CATEGORICAL DATA AND SOME DATA WRANGLING



dplyr



magrittr



janitor

# FREQUENCY TABLES: `count()`

- `count` is from the `dplyr` package
- the output is a long tibble, and not a “nice” table

```
1 dds.dscr %>% count(ethnicity)

# A tibble: 8 × 2
  ethnicity      n
  <fct>     <int>
1 American     4
2 Asian        129
3 Black         59
4 Hispanic      376
5 Multi Race   26
6 Native Hawaiian 3
7 Other          2
8 White not Hispanic 401
```

```
1 dds.dscr %>%
2   count(ethnicity, age.cohort)

# A tibble: 35 × 3
  ethnicity      age.cohort     n
  <fct>          <fct>       <int>
1 American Indian 13–17           1
2 American Indian 22–50           1
3 American Indian 51+            2
4 Asian            0–5            8
5 Asian            6–12           18
6 Asian            13–17          20
7 Asian            18–21          41
8 Asian            22–50          29
9 Asian            51+            13
10 Black           0–5            3
# i 25 more rows
```

# HOW TO USE THE PIPE %>%

The pipe operator `%>%` strings together commands to be performed sequentially

```
1 dds.dscr %>% head(n=3)      # pronounce %>% as "then"

# A tibble: 3 × 6
  id age.cohort    age gender expenditures ethnicity
  <int> <fct>      <int> <fct>        <int> <fct>
1 10210 13-17       17 Female        2113 White not Hispanic
2 10409 22-50       37 Male          41924 White not Hispanic
3 10486 0-5         3 Male          1454 Hispanic
```

- Always *first list the tibble* that the commands are being applied to
- Can use **multiple pipes** to run multiple commands in sequence
  - What does the following code do?

```
1 dds.dscr %>% head(n=3) %>% summary()
```

# FREQUENCY TABLES: janitor PACKAGE'S tabyl FUNCTION

```
1 # default table  
2 dds.dscr %>%  
3 tabyl(ethnicity)
```

	ethnicity	n	percent
American Indian	Indian	4	0.004
	Asian	129	0.129
	Black	59	0.059
	Hispanic	376	0.376
	Multi Race	26	0.026
Native Hawaiian	Hawaiian	3	0.003
	Other	2	0.002
White not Hispanic	not Hispanic	401	0.401

## adorn\_ your table!

```
1 dds.dscr %>%  
2 tabyl(ethnicity) %>%  
3 adorn_totals("row") %>%  
4 adorn_pct_formatting(digits=2)
```

	ethnicity	n	percent
American Indian	Indian	4	0.40%
	Asian	129	12.90%
	Black	59	5.90%
	Hispanic	376	37.60%
	Multi Race	26	2.60%
Native Hawaiian	Hawaiian	3	0.30%
	Other	2	0.20%
White not Hispanic	not Hispanic	401	40.10%
	Total	1000	100.00%

# RELATIVE FREQUENCY TABLE

- A **relative frequency** table shows **proportions** (**or percentages**) instead of counts
- To the right I removed (deselected) the counts column (**n**) to create a relative frequency table

```
1 dds.discr %>%
2   tabyl(ethnicity) %>%
3   adorn_totals("row") %>%
4   adorn_pct_formatting(digits=2) %>%
5   select(-n)
```

	ethnicity	percent
American	Indian	0.40%
	Asian	12.90%
	Black	5.90%
	Hispanic	37.60%
	Multi Race	2.60%
Native	Hawaiian	0.30%
	Other	0.20%
White	not Hispanic	40.10%
	Total	100.00%

# CONTINGENCY TABLES (TWO-WAY TABLES)

- **Contingency tables** summarize data for two categorical variables
  - with each value in the table representing the number of times a particular combination of outcomes occurs
- **Row & column totals** are sometimes called **marginal totals**

```
1 dds.dscr %>%
2 tabyl(ethnicity, gender) %>%
3 adorn_totals(c("row", "col"))
```

	ethnicity	Female	Male	Total
American	Indian	3	1	4
	Asian	61	68	129
	Black	26	33	59
	Hispanic	192	184	376
	Multi Race	13	13	26
	Native Hawaiian	2	1	3
	Other	1	1	2
White not Hispanic		205	196	401
	Total	503	497	1000

# CONTINGENCY TABLES WITH PERCENTAGES

```
1 dds.dscr %>%
2   tabyl(ethnicity, age.cohort) %>%
3   adorn_totals(c("row")) %>%
4   adorn_percentages("row") %>%
5   adorn_pct_formatting(digits=0) %>%
6   adorn_ns()
```

ethnicity		0-5	6-12	13-17	18-21	22-50	51+
American Indian	0%	(0)	0%	(0)	25%	(1)	0%
Asian	6%	(8)	14%	(18)	16%	(20)	32%
Black	5%	(3)	19%	(11)	20%	(12)	15%
Hispanic	12%	(44)	24%	(91)	27%	(103)	21%
Multi Race	27%	(7)	35%	(9)	27%	(7)	8%
Native Hawaiian	0%	(0)	0%	(0)	0%	(0)	67%
Other	0%	(0)	0%	(0)	100%	(2)	0%
White not Hispanic	5%	(20)	11%	(46)	17%	(67)	17%
Total	8%	(82)	18%	(175)	21%	(212)	20%

# SUMMARIZING NUMERIC DATA

# MEAN ANNUAL DDS EXPENDITURES BY RACE/ETHNICITY

```
1 mean(dds.dscr$expenditures)
[1] 18065.79

1 dds.dscr %>%
2   summarize(
3     ave = mean(expenditures),
4     SD = sd(expenditures),
5     med = median(expenditures))

# A tibble: 1 × 3
  ave     SD    med
  <dbl>  <dbl> <dbl>
1 18066. 19543. 7026
```

```
1 dds.dscr %>%
2   group_by(ethnicity) %>%
3   summarize(
4     ave = mean(expenditures),
5     SD = sd(expenditures),
6     med = median(expenditures))

# A tibble: 8 × 4
  ethnicity          ave      SD    med
  <fct>            <dbl>    <dbl> <dbl>
1 American Indian  36438. 25694. 41818.
2 Asian             18392. 19209.  9369
3 Black              20885. 20549.  8687
4 Hispanic           11066. 15630.  3952
5 Multi Race        4457.   7332.  2622
6 Native Hawaiian  42782.  6576.  40727
7 Other              3316.   1836.  3316.
8 White not Hispanic 24698. 20604. 15718
```

# get\_summary\_stats() FROM rstatix PACKAGE

```
1 dds.dscr %>% get_summary_stats()  
  
# A tibble: 3 × 13  
  variable      n    min    max median     q1     q3    iqr    mad    mean     sd  
  <fct>      <dbl>  
1 id          1000 10210 99898 55384. 31809. 76135. 44326 3.27e4 5.47e4 2.56e4  
2 age         1000     0    95    18     12     26     14 1.04e1 2.28e1 1.85e1  
3 expenditures 1000    222 75098  7026   2899. 37713. 34814 7.76e3 1.81e4 1.95e4  
# i 2 more variables: se <dbl>, ci <dbl>  
  
1 dds.dscr %>%  
2   group_by(ethnicity) %>%  
3   get_summary_stats(expenditures, type = "common")  
  
# A tibble: 8 × 11  
  ethnicity variable      n    min    max median     iqr    mean     sd     se     ci  
  <fct>      <fct>      <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>  
1 American... expendi...     4    3726  58392  41818. 34085. 36438. 25694. 12847. 40885.  
2 Asian       expendi...   129    374   75098   9369   30892   18392. 19209. 1691.  3346.  
3 Black       expendi...    59    240   60808   8687   37987   20885. 20549. 2675.  5355.  
4 Hispanic    expendi...   376    222   65581   3952   7961.  11066. 15630.  806.  1585.  
5 Multi Ra... expendi...    26    669   38619   2622   2060.   4457.  7332.  1438.  2962.  
6 Native H... expendi...     3   37479  50141  40727   6331   42782.  6576.  3797. 16337.  
7 Other       expendi...     2    2018   4615   3316.   1298.   3316.  1836.  1298. 16499.  
8 White no... expendi...   401    340   68890  15718   39157   24698. 20604. 1029.  2023.
```

# HOW TO FORCE ALL OUTPUT TO BE SHOWN? (1/2)

Use `kable()` from the `knitr` package.

```
1 dds.discr %>% get_summary_stats() %>% kable()
```

variable	n	min	max	median	q1	q3	iqr
id	1000	10210	99898	55384.5	31808.75	76134.75	44326
age	1000	0	95	18.0	12.00	26.00	14
expenditures	1000	222	75098	7026.0	2898.75	37712.75	34814

# HOW TO FORCE ALL OUTPUT TO BE SHOWN? `knitr` (2/2)

Use `kable()` from the `knitr` package.

```
1 dds.dscr %>%
2   group_by(ethnicity) %>%
3   get_summary_stats(expenditures, type = "common") %>%
4   kable()
```

<b>ethnicity</b>	<b>variable</b>	<b>n</b>	<b>min</b>	<b>max</b>	<b>median</b>	<b>iqr</b>	<b>mean</b>
American Indian	expenditures	4	3726	58392	41817.5	34085.25	36438.250
Asian	expenditures	129	374	75098	9369.0	30892.00	18392.372
Black	expenditures	59	240	60808	8687.0	37987.00	20884.593
Hispanic	expenditures	376	222	65581	3952.0	7961.25	11065.569
Multi Race	expenditures	26	669	38619	2622.0	2059.75	4456.731
Native Hawaiian	expenditures	3	37479	50141	40727.0	6331.00	42782.333
Other	expenditures	2	2018	4615	3316.5	1298.50	3316.500

<b>ethnicity</b>	<b>variable</b>	<b>n</b>	<b>min</b>	<b>max</b>	<b>median</b>	<b>iqr</b>	<b>mean</b>
White not Hispanic	expenditures	401	340	68890	15718.0	39157.00	24697.549

BACK TO RESEARCH  
QUESTION

# CASE STUDY: DISCRIMINATION IN DEVELOPMENTAL DISABILITY SUPPORT (1.7.1)

- **Previous research**
  - Researchers examined DDS expenditures for developmentally disabled residents by ethnicity
  - Found that the mean annual expenditures on Hispanics was less than that on White non-Hispanics.
- **Result:** an allegation of ethnic discrimination was brought against the California DDS.
- **Question:** Are the data sufficient evidence of ethnic discrimination?

