

## Overview

Review of ggplot

Review of material covered in the class so far

If there is time/interest

• Bonus features of ggplot: special geoms, animation, interactive graphics

## Announcements

#### Midterm exam is on Thursday

- Bring a pen and a pencil
- One page (2 sides) with code and equations only!
  - You will turn in this page of notes with your exam (put your name on it)
  - You can write down conditions for hypothesis tests. Have SE formulas, etc.

#### Office hours this week

- Stephan and Nathan are doing another review session tonight
- No TA office hours since no homework

# Review of the grammar of graphics and ggplot



# The grammar of graphics

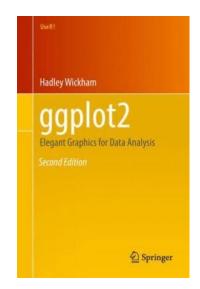
Leland Wilkinson noticed similarities between many graphs and tried to generate a 'grammar' that could be used to express a graph

• i.e., a list elements that can be combined together to create a graph

Leland Wilkinson

The Grammar of Graphics
Second Edition

Hadley Wickham implemented these ideas in R in the ggplot2 package



## Graphs are composed of...

A Frame: Coordinate system on which data is placed

• ggplot() +

**Glyphs**: basic graphic unit representing cases or statistics

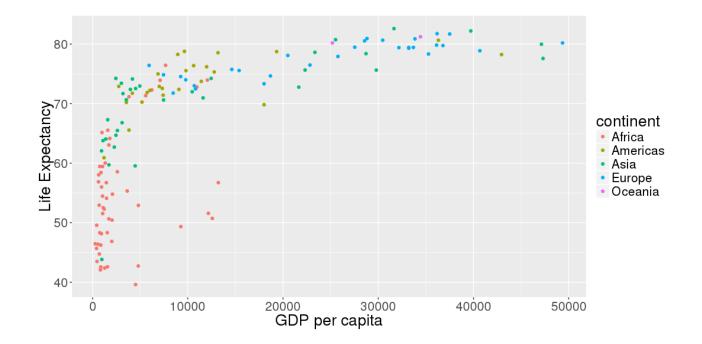
- Data is mapped onto these aesthetics such as: shape, color, size, etc. and/or aesthetics can be set to a fixed value
  - geom\_point(aes(x = gdpPercap, y = lifeExp, color = continent))

geom point(aes(x = gdpPercap, y = lifeExp), color = "red")

**Scales and guides**: shows how to interpret axes and other properties of the glyphs

scale x continuous(trans = "log10")

scale color brewer(type = "qua", palette = 2)



## Plots can also contain...

Facets: allows for multiple side-by-side graphs based on a categorical variable

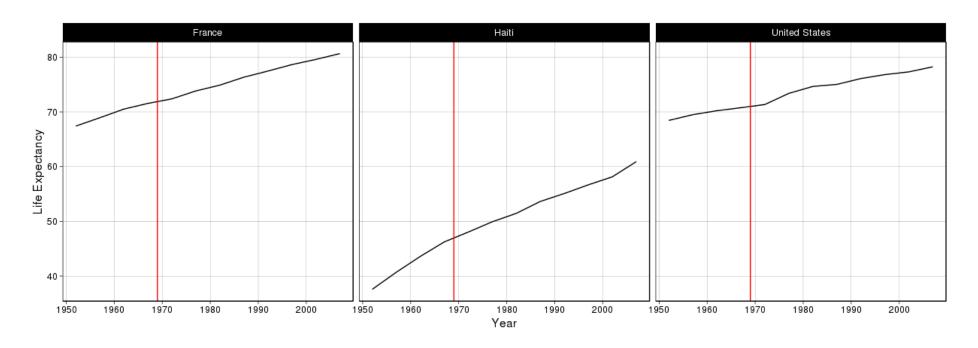
facet\_wrap(~country)

Layers: allows for more than one types of data to be mapped onto the same figure

geom\_vline(xintercept = 1969, col = "red")

**Theme**: contains finer points of display (e.g., font size, background color, etc.)

theme\_wsj()



## Questions?

# ggplot2 cheat sheet

## Data visualization with ggplot2:: CHEAT SHEET

**Basics** 

data

and y locations.

size = A

ggplot (data = <DATA>) +

<FACET\_FUNCTION> +

<SCALE\_FUNCTION> +

<THEME\_FUNCTION>

function per layer.

6 = "twodash")

<COORDINATE FUNCTION> +

last\_plot() Returns the last plot.

Matches file type to file extension.

Aes Common aesthetic values.

color and fill - string ("red", "#RRGGBB")

ggplot2 is based on the grammar of graphics, the idea

and geoms-visual marks that represent data points.

To display values, map variables in the data to visual

properties of the geom (aesthetics) like size, color, and x

coordinate

<GEOM\_FUNCTION>(mapping = aes(<MAPPINGS>

ggplot(data = mpg, aes(x = cty, y = hwy)) Begins a plot

ggsave("plot.png", width = 5, height = 5) Saves last plot

as 5' x 5' file named "plot.png" in working directory.

linetype - integer or string (0 = "blank", 1 = "solid",

lineend - string ("round", "butt", or "square")

linejoin - string ("round", "mitre", or "bevel")

Studio

2 = "dashed", 3 = "dotted", 4 = "dotdash", 5 = "longdash",

size - integer (line width in mm) 0 1 2 3 4 5 6 7 8 9 10 11 12

shape - integer/shape name or 13 14 15 16 17 18 19 20 21 22 23 24 25

a single character ("a") ⊠⊠□○△♦○●□♦△♥

that you finish by adding layers to. Add one geom

Complete the template below to build a graph.

stat = <STAT>, position = <POSITION>) +

that you can build every graph from the same components: a data set, a coordinate system, Geoms Use a geom function to represent data points, use the geom's aesthetic properties to represent variables. Each function returns a layer.

#### GRAPHICAL PRIMITIVES

a <- ggplot(economics, aes(date, unemploy)) b <- ggplot(seals, aes(x = long, y = lat))

> a + geom\_blank() and a + expand\_limits() Ensure limits include values across all plots.

b + geom\_curve(aes(yend = lat + 1, xend = long + 1), curvature = 1) - x, xend, y, yend, alpha, angle, color, curvature, linetype, size

a + geom\_path(lineend = "butt", linejoin = "round", linemitre = 1) x, y, alpha, color, group, linetype, size

> a + geom\_polygon(aes(alpha = 50)) - x, y, alpha, color, fill, group, subgroup, linetype, size b + geom\_rect(aes(xmin = long, ymin = lat,

xmax = long + 1, ymax = lat + 1)) - xmax, xmin, ymax, ymin, alpha, color, fill, linetype, size

a + geom\_ribbon(aes(ymin = unemploy - 900. ymax = unemploy + 900)) - x, ymax, ymin, alpha, color, fill, group, linetype, size

#### LINE SEGMENTS

common aesthetics: x, y, alpha, color, linetype, size

b + geom\_abline(aes(intercept = 0, slope = 1)) b + geom\_hline(aes(yintercept = lat)) b + geom\_vline(aes(xintercept = long))

b + geom\_segment(aes(vend = lat + 1, xend = long + 1)) b + geom\_spoke(aes(angle = 1:1155, radius = 1))

#### ONE VARIABLE continuous

c <- ggplot(mpg, aes(hwy)); c2 <- ggplot(mpg)

c + geom\_area(stat = "bin")

x, y, alpha, color, fill, linetype, size

c + geom\_density(kernel = "gaussian") x, y, alpha, color, fill, group, linetype, size, weight

c + geom\_dotplot() x, y, alpha, color, fill

c + geom\_freqpoly()

x, y, alpha, color, group, linetype, size

c + geom\_histogram(binwidth = 5) x, y, alpha, color, fill, linetype, size, weight

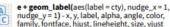
c2 + geom\_qq(aes(sample = hwy)) x, y, alpha, color, fill, linetype, size, weight

d <- ggplot(mpg, aes(fl))

d + geom\_bar() x, alpha, color, fill, linetype, size, weight

#### TWO VARIABLES

both continuous e <- ggplot(mpg, aes(cty, hwy))



e + geom\_point() x, y, alpha, color, fill, shape, size, stroke

e + geom\_quantile() x, y, alpha, color, group, linetype, size, weight

e + geom\_rug(sides = "bl") x, y, alpha, color, linetype, size

e + geom\_smooth(method = lm) x, y, alpha, color, fill, group, linetype, size, weight

e + geom\_text(aes(label = cty), nudge\_x = 1, nudge\_y = 1) - x, y, label, alpha, angle, color, family, fontface, hjust, lineheight, size, vjust

#### one discrete, one continuous f <- ggplot(mpg, aes(class, hwy))

f + geom col() x, y, alpha, color, fill, group, linetype, size

x, y, lower, middle, upper, ymax, ymin, alpha, color, fill, group, linetype, shape, size, weight

f + geom\_dotplot(binaxis = "y", stackdir = "center") x, y, alpha, color, fill, group

f + geom\_violin(scale = "area") x, y, alpha, color, fill, group, linetype, size, weight

#### both discrete

g <- ggplot(diamonds, aes(cut, color))

g + geom\_count()

x, y, alpha, color, fill, shape, size, stroke

e + geom\_jitter(height = 2, width = 2) x, y, alpha, color, fill, shape, size

#### continuous bivariate distribution h <- ggplot(diamonds, aes(carat, price))

h + geom bin2d(binwidth = c(0.25, 500)) x, y, alpha, color, fill, linetype, size, weight

ggplot.

h + geom\_density\_2d()

x, y, alpha, color, group, linetype, size

h + geom\_hex() x, y, alpha, color, fill, size

continuous function

i <- ggplot(economics, aes(date, unemploy))

i + geom area()

x, y, alpha, color, fill, linetype, size

/ i + geom line() x, y, alpha, color, group, linetype, size

i + geom\_step(direction = "hv") x, y, alpha, color, group, linetype, size

visualizing error

df <- data.frame(grp = c("A", "B"), fit = 4:5, se = 1:2) j <- ggplot(df, aes(grp, fit, ymin = fit - se, ymax = fit + se))

j + geom\_crossbar(fatten = 2) - x, y, ymax,

ymin, alpha, color, fill, group, linetype, size

j + geom\_errorbar() - x, ymax, ymin, alpha, color, group, linetype, size, width Also geom\_errorbarh().

j + geom\_linerange() x, ymin, ymax, alpha, color, group, linetype, size

j + geom\_pointrange() - x, y, ymin, ymax, alpha, color, fill, group, linetype, shape, size

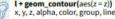
data <- data.frame(murder = USArrests\$Murder, state = tolower(rownames(USArrests))) map <- map data("state")

k <- ggplot(data, aes(fill = murder))

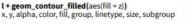
k + geom\_map(aes(map\_id = state), map = map) + expand\_limits(x = map\$long, y = map\$lat) map\_id, alpha, color, fill, linetype, size

#### THREE VARIABLES

seals\$z <- with(seals, sqrt(delta\_long^2 + delta\_lat^2)); l <- ggplot(seals, aes(long, lat))



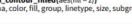
x, y, z, alpha, color, group, linetype, size, weight



I + geom\_raster(aes(fill = z), hjust = 0.5, vjust = 0.5, interpolate = FALSE) x, y, alpha, fill

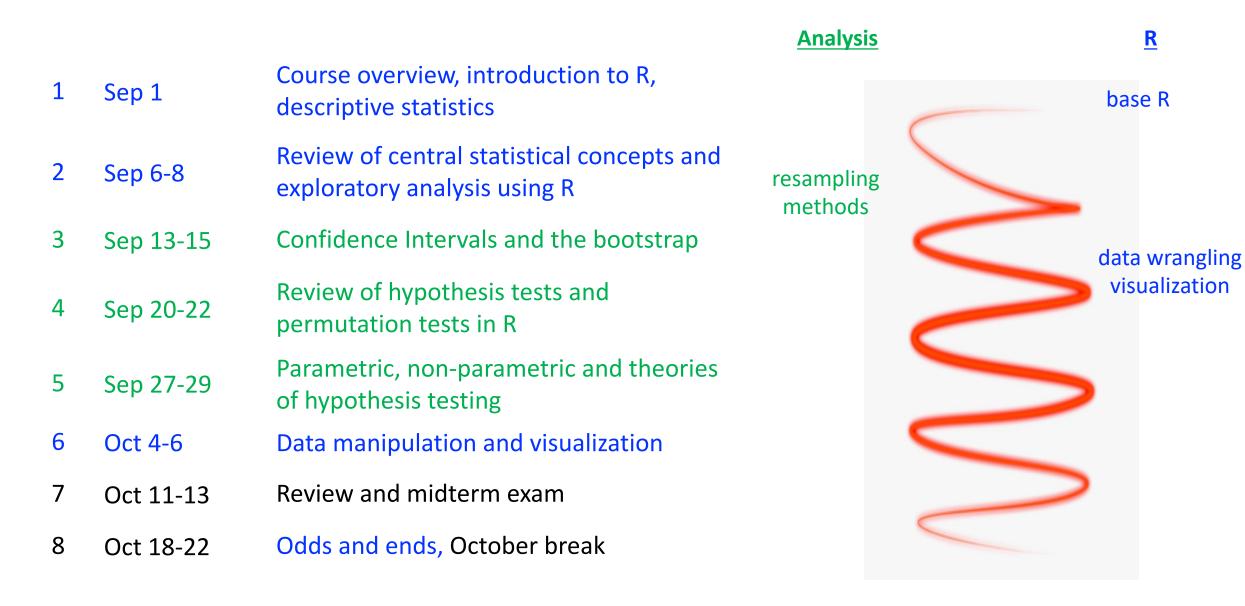


l + geom\_tile(aes(fill = z)) x, y, alpha, color, fill, linetype, size, width

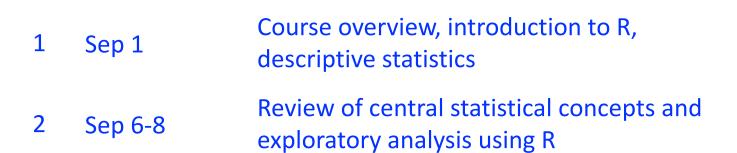


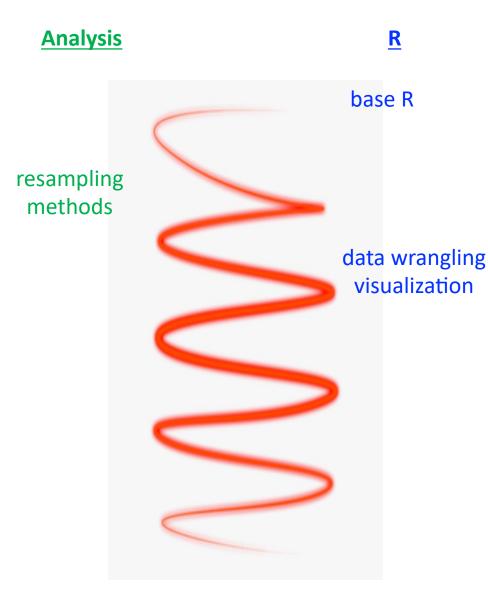


## What we have covered so far...

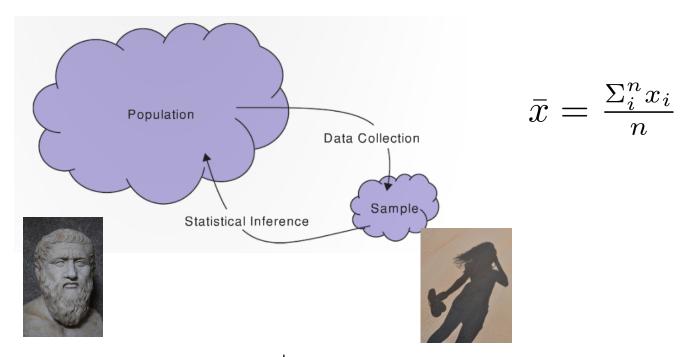


## What we have covered so far...





# Parameters and statistics commonly used symbols



	Population parameter (Plato)	Sample statistic (shadow)
Mean	μ	χ̄
Standard deviation	σ	S
Proportion	π	ĝ
Correlation	ρ	r
Regression slope	β	b

## Base R

#### Basics of R

```
> my_vec <- c(5, 28, 19)
```

> inds\_less\_than\_10 <- 10

#### How to plot data in base R

- > drinks\_table <- table(profiles\$drinks)
- > barplot(drinks\_table)
- > pie(drinks\_table)
- > hist(profiles\$height)

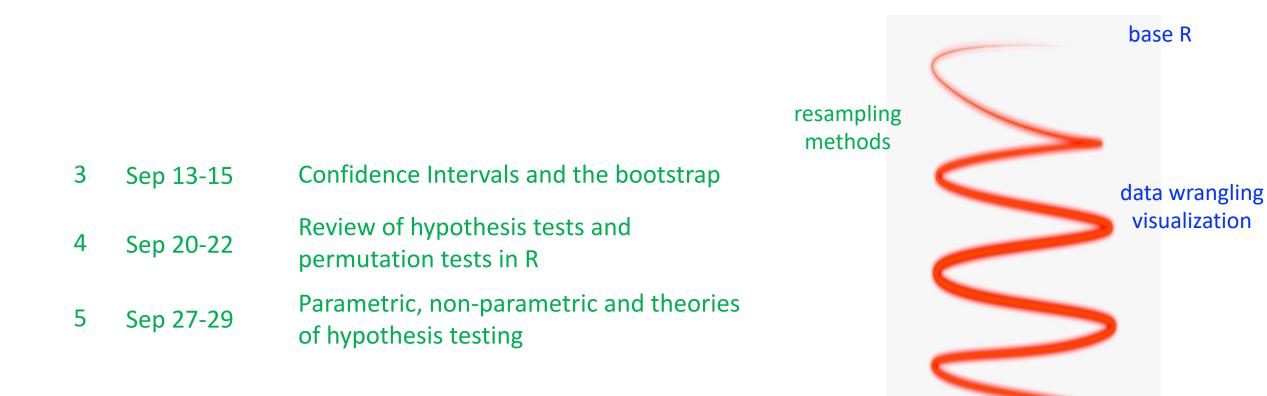
```
For loops

my_results <- NULL

for (i in 1:100) {

    my_results[i] <- i^2
}
```

## What we have covered so far...



**Analysis** 

R

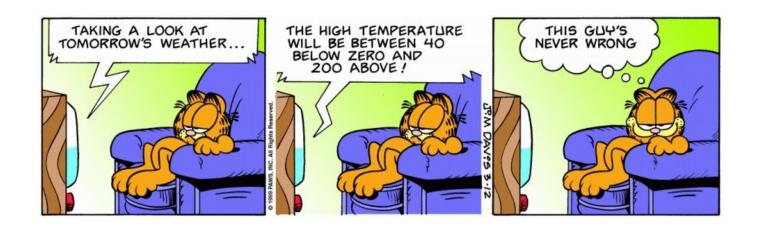
## Probability and confidence intervals

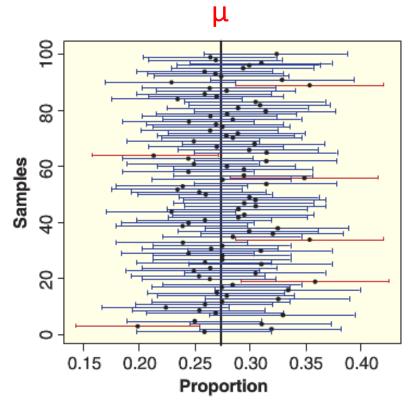
Probability functions; e.g., rnorm, pnorm, dnorm, qnorm

#### Confidence intervals:

$$Cl_{95} = stat \pm 2 \cdot SE$$

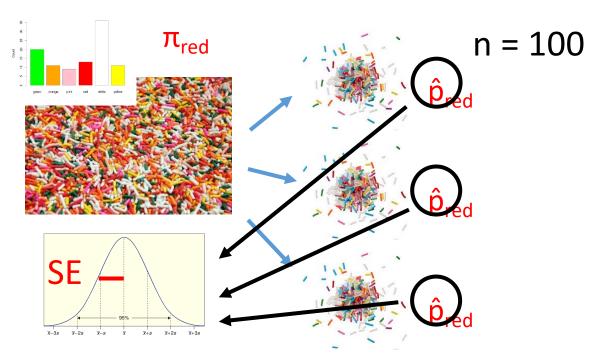






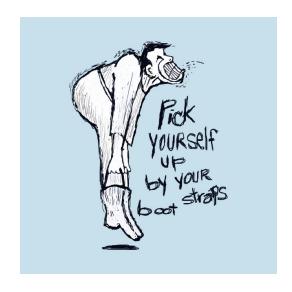
# Sampling and bootstrap distributions

### Sampling distribution



Sampling distribution!

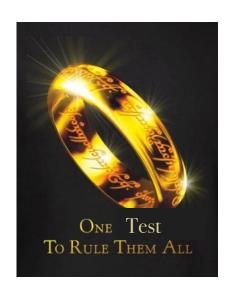
#### Bootstrap distribution

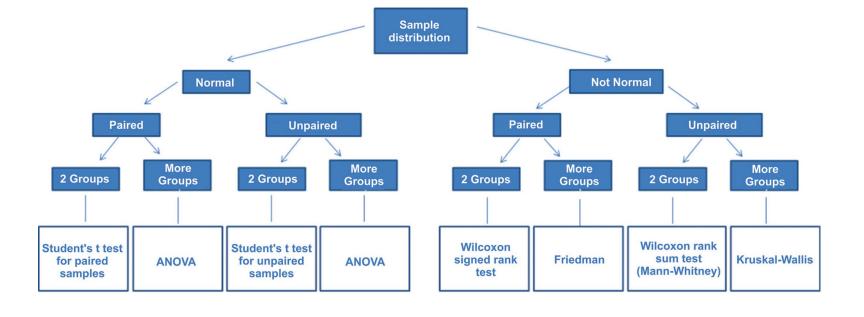


Sample with replacement from our original sample to mimic a sampling distribution

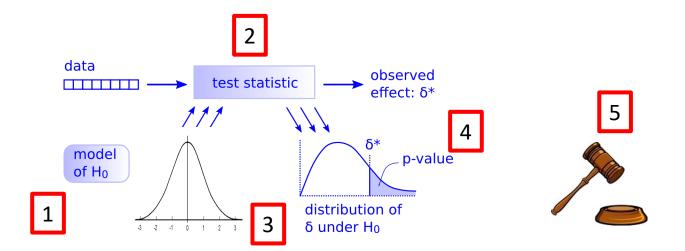
$$Cl_{95} = stat \pm 2 \cdot SE^*$$

## Hypothesis tests

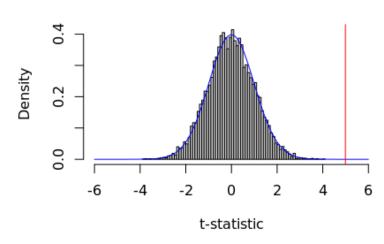




#### Just need to follow 5 steps!



#### Null distribution



# Randomization/permutation tests

#### Create a null distribution through computational simulations/shuffling

rbinom(), sample(), etc.

 $H_0$ :  $\pi = 0.5$ 

 $H_A$ :  $\pi > 0.5$ 

 $H_0$ :  $\mu_T - \mu_C = 0$ 

 $H_A: \mu_T - \mu_C > 0$ 

$$H_0: \mu_i = \mu_j \dots = \dots \mu_k$$

 $H_A$ :  $\mu_i \neq \mu_j$  for some i, j







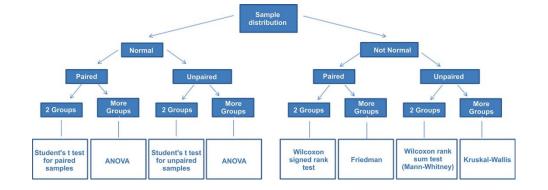
Data	1 Sample	2 Samples	> 2 Samples
Categorical data	H <sub>0</sub> : $\pi = p_0$ H <sub>A</sub> : $\pi \neq p_0$ Flip "coins" rbinom()	$H_0$ : $\pi_1 = \pi_2$ $H_A$ : $\pi_1 \neq \pi_2$ Flip "coins" rbinom()	$H_0$ : $\pi_1 = p_1$ , $\pi_2 = p_2$ ,, $\pi_k = p_k$ $H_A$ : At least one $p_i$ is different than specified  Flip coins  rmultinom()
Quantitative data	H <sub>0</sub> : $\mu = v_0$ H <sub>A</sub> : $\mu \neq v_0$ resample	H <sub>0</sub> : $\mu_1 = \mu_2$ H <sub>A</sub> : $\mu_1 \neq \mu_2$ Shuffle data	H <sub>0</sub> : $\mu_1 = \mu_2 = = \mu_k$ H <sub>A</sub> : At least one $\mu_i$ is different Shuffle data
	sample(, replace = TRUE)	sample()	sample()

## Parametric tests

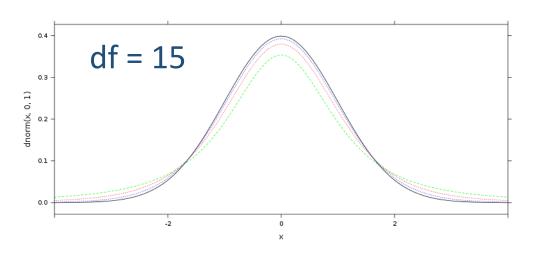
#### Use mathematical density functions for the null distribution

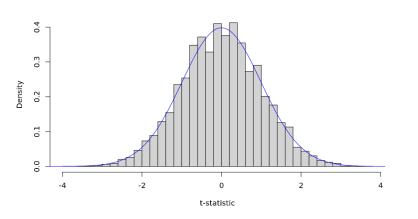
$$H_0: \mu_T - \mu_C = 0$$
  
 $H_A: \mu_T - \mu_C > 0$ 

$$t = \frac{\bar{x}_t - \bar{x}_c}{\sqrt{\frac{s_t^2}{n_t} + \frac{s_c^2}{n_c}}}$$









Data	1 Sample	2 Samples	> 2 Samples
	$H_0$ : π = $p_0$ $H_A$ : π ≠ $p_0$	$H_0: \pi_1 = \pi_2$ $H_A: \pi_1 \neq \pi_2$	$H_0$ : $\pi_1 = p_1$ , $\pi_2 = p_2$ ,, $\pi_k = p_k$ $H_A$ : At least one $p_i$ is different than specified
Categorical data	<u>z-test</u>	z-test or a chi-square	<u>chi-square test</u>
	$z = \frac{\hat{p} - p_0}{\sqrt{\frac{p_0(1 - p_0)}{n}}}$	$z = \frac{\hat{p_1} - \hat{p_2}}{\sqrt{\frac{\hat{p_1}(1-\hat{p_1})}{n_1} + \frac{\hat{p_2}(1-\hat{p_2})}{n_2}}}$	$\chi^2 = \sum_{i=1}^k \frac{(Observed_i - Expected_i)^2}{Expected_i}$
	$H_0$ : $\mu = v_0$ $H_A$ : $\mu \neq v_0$	$H_0: \mu_1 = \mu_2$ $H_A: \mu_1 \neq \mu_2$	$H_0$ : $\mu_1 = \mu_2 = = \mu_k$ $H_A$ : At least one $\mu_i$ is different
Quantitative data	One sample t-test	Two sample t-test	Analysis of Variance
	$t = \frac{\bar{x} - v_0}{s / \sqrt{n}}$	$t = \frac{\bar{x}_1 - \bar{x}_2}{\sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}}$	$F = \frac{\frac{1}{K-1} \sum_{i=1}^{K} n_i (\bar{x}_i - \bar{x}_{tot})^2}{\frac{1}{N-K} \sum_{i=1}^{K} \sum_{j=1}^{n_i} (x_{ij} - \bar{x}_i)^2}$
			$df_1 = k,  df_2 = n - k$

Data	1 Sample	2 Samples
Categorical Data	$SE = \sqrt{\frac{\pi(1-\pi)}{n}}$	$SE = \sqrt{\frac{\pi_1(1-\pi_1)}{n_1} + \frac{\pi_2(1-\pi_2)}{n_2}}$
	$\hat{p} \pm z^* \cdot \sqrt{\frac{\hat{p}(1-\hat{p})}{n}}$	$\hat{p}_1 - \hat{p}_2 \pm z^* \sqrt{\frac{\hat{p}_1(1-\hat{p}_1)}{n_1} + \frac{\hat{p}_2(1-\hat{p}_2)}{n_2}}$
Quantitative Data	$SE = \frac{s}{\sqrt{n}}$ $x \pm t^* \frac{s}{\sqrt{n}}$	$SE = \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}$ $(\overline{x_1} - \overline{x_2}) \pm t^* \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}$

## Theories of hypothesis testing



Fisher (1890-1962)

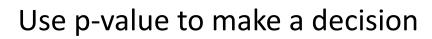


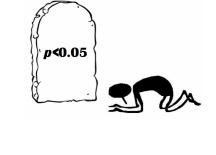
Neyman (1894-1981)

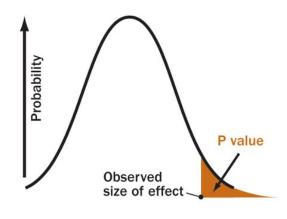


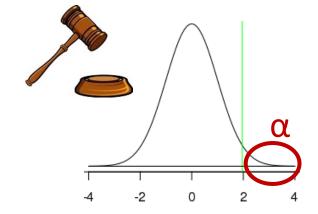
Pearson (1895-1980)

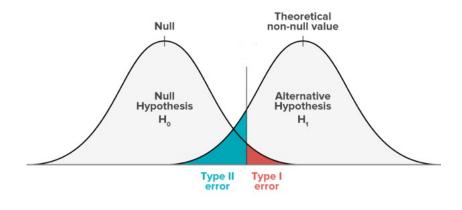
#### p-value a strength of evidence



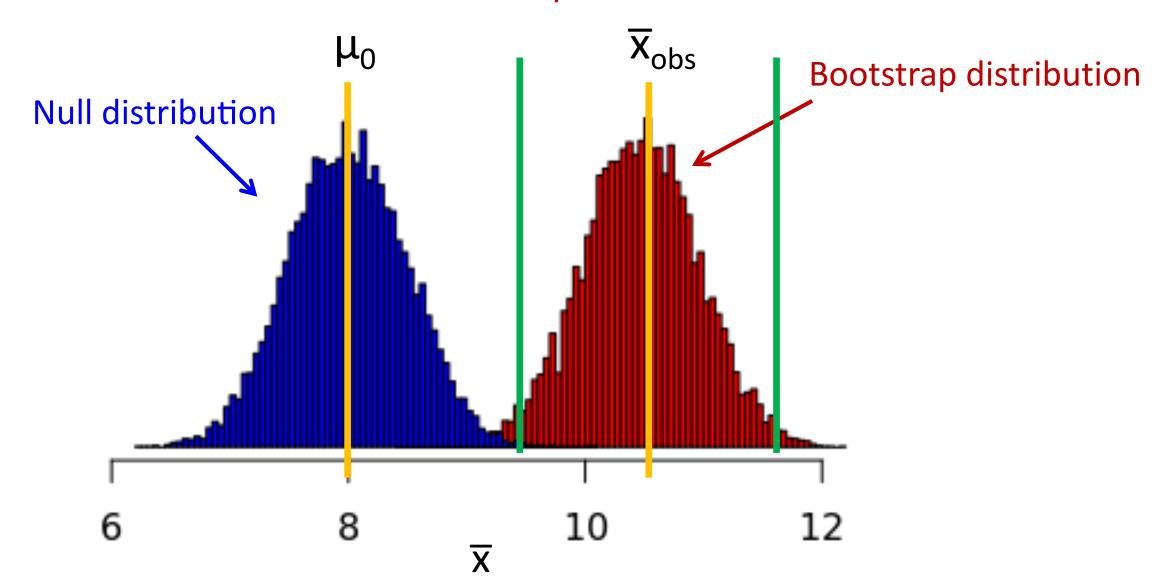








# Relationship between null and bootstrap distributions



# Data manipulation with dplyr

# **dplyr** is a package that has a set of verbs for transformations data

- All these function take a data frame and other arguments and return a data frame
- 1. filter()
- 2. select()
- 3. mutate()
- 4. arrange()
- 5. summarize()
- 6. group\_by()

age <sup>‡</sup>	body_type	diet <sup>‡</sup>
22	a little extra	strictly anything
35	average	mostly other
38	thin	anything
23	thin	vegetarian
29	athletic	NA
29	average	mostly anything

```
film results <- movies |>
   filter(title type == "Feature Film") |>
   select(critics_score, audience_score, genre) |>
   mutate(audience prefers =
         audience score - critics score) |>
   group_by(genre) |>
    summarize(mean audience prefers =
          mean(audience prefers)) |>
     arrange(desc(mean_audience_prefers))
head(film results)
```

# Grammar of graphics with ggplot

A Frame: Coordinate system on which data is placed

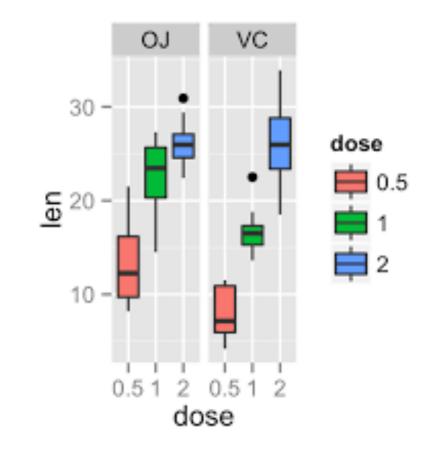
**Glyphs**: basic graphic unit representing cases or statistics

**Scales and guides**: shows how to interpret axes and other properties of the glyphs

**Facets**: allows for multiple side-by-side graphs based on a categorical variable

**Layers:** allows for more than one types of data to be mapped onto the same figure

**Theme**: contains finer points of display (e.g., font size, background color, etc.)



## Questions

