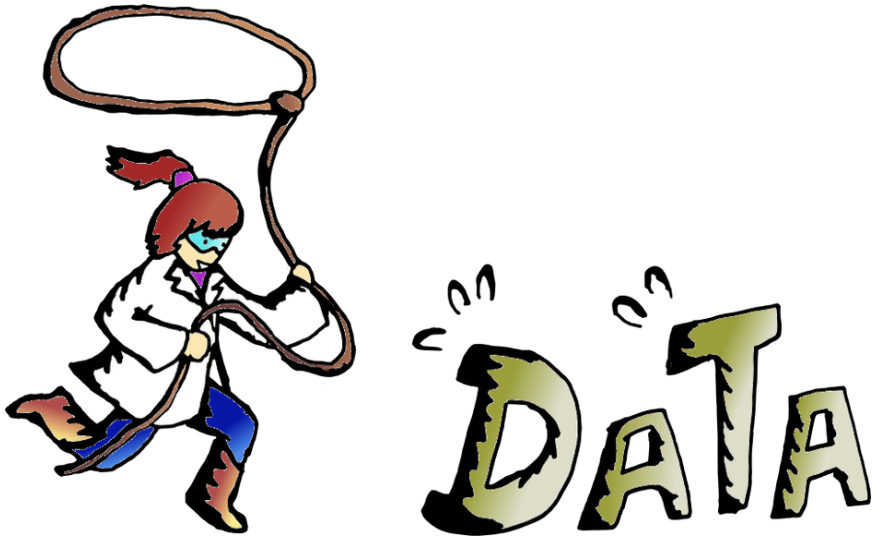
Joining, mapping, and reshaping data



Overview

Joining data frames continued

Creating maps

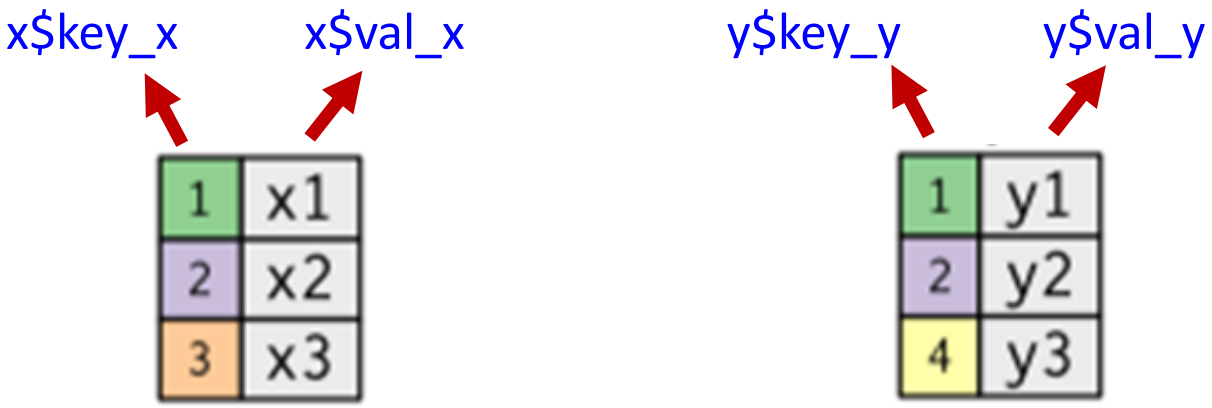
Reshaping data

Joining data frames

Left and right tables

Suppose we have two data frames called x and y

- x have two variables called key_x, and val_x
- y has two variables called key_y and val_y

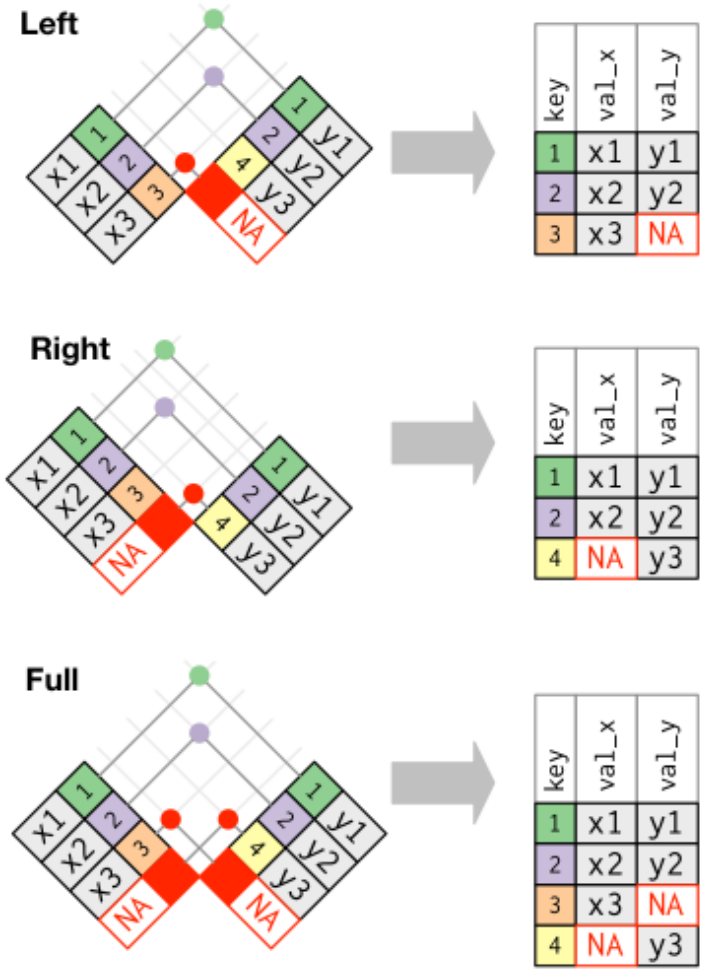


Data frame x **Data frame y**

Joins have the general form:

```
join(x, y, by = c("key_x" = "key_y"))
```

Joining data frames



Left joins keep all rows in the left table.

left_join(x, y, by = c("key_x" = "key_y"))

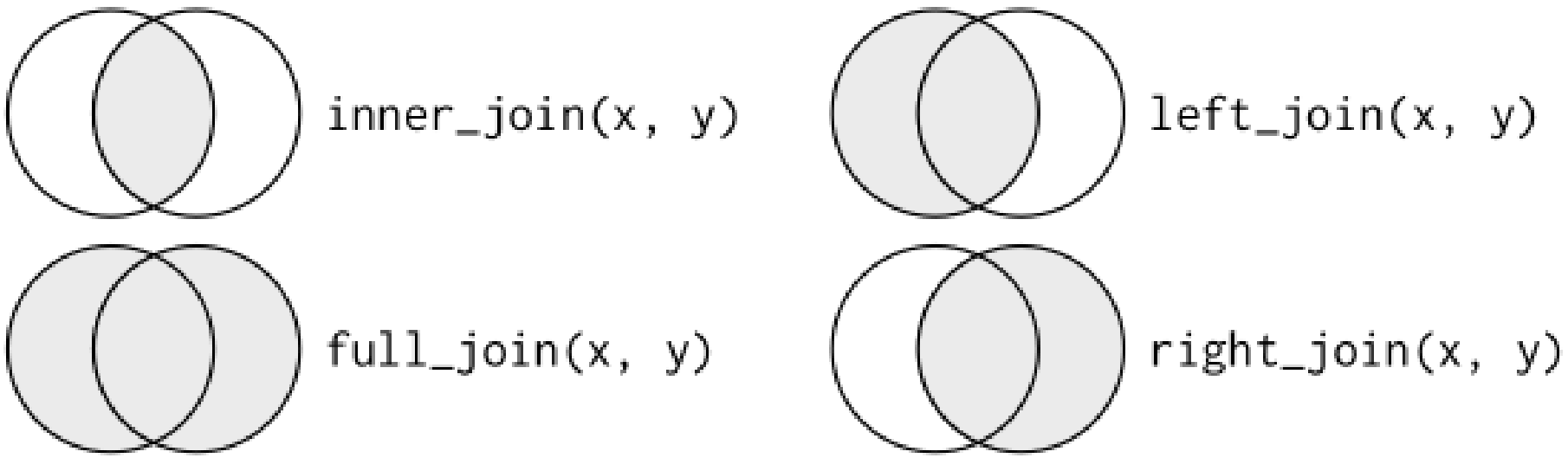
Right joins keep all rows in the right table.

right_join(x, y, by = c("key_x" = "key_y"))

Full joins keep all rows in both tables.

full_join(x, y, by = c("key_x" = "key_y"))

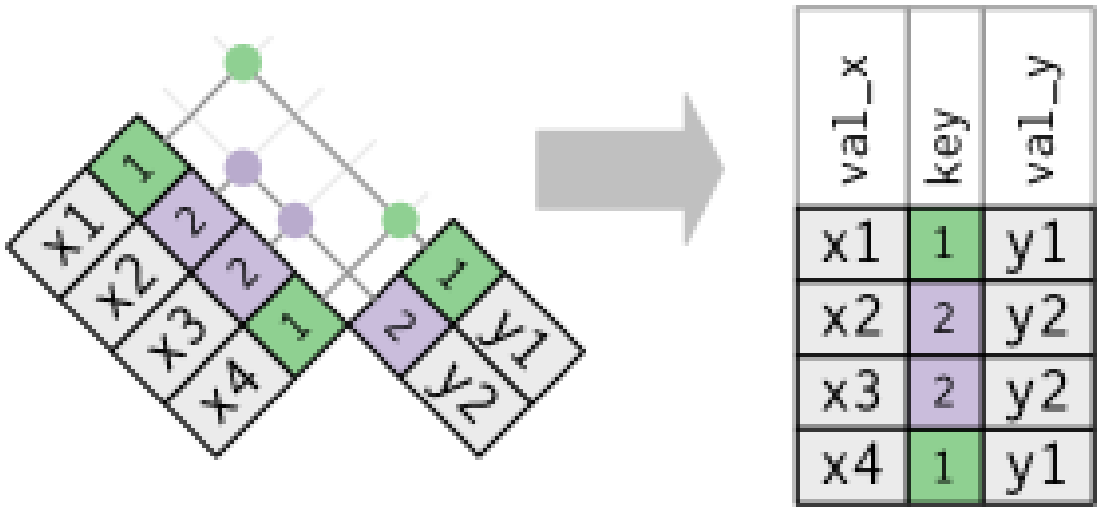
Summary



Duplicate keys

Duplicate keys are useful if there is a many-to-one relationship

- e.g., duplicates are useful in the left table when doing a left join

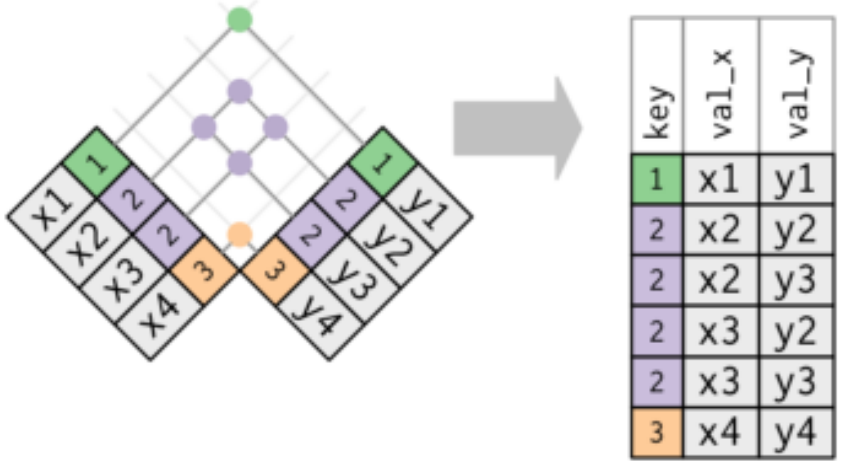


val_x	key	val_y
x1	1	y1
x2	2	y2
x3	2	y2
x4	1	y1

Duplicate keys

If both tables have duplicate keys you get all possible combinations of joined values (Cartesian product).

- **This is usually an error!**



key	val_x	val_y
1	x1	y1
2	x2	y2
2	x2	y3
2	x3	y2
2	x3	y3
3	x4	y4

Always check the output size after you join a table because even if there is not a syntax error you might not get the table you are expecting!

- You can check how many rows a data frame has using the nrow() function

Duplicate keys

To deal with duplicate keys in both tables, we can join the tables using multiple keys in order to make sure that each row is uniquely specified.

We can do this using the syntax:

```
join(x2, y2, by = c("key1_x" = "key1_y", "key2_x" = "key2_y"))
```

Duplicate keys

```
>  x2 <- data.frame(key1_x = c(1, 2, 2),
                    key2_x = c("a", "a", "b"),
                    val_x = c("y1", "y2", "y3"))

>  y2 <- y2 <- data.frame(key1_y = c(1, 2, 2, 3, 3),
                    key2_y = c("a", "a", "b", "a", "b"),
                    val_y = c("y1", "y2", "y3", "y4", "y5"))

> left_join(x2, y2, c("key1_x" = "key1_y"))

> left_join(x2, y2, c("key1_x" = "key1_y", "key2_x" = "key2_y"))
```

Structured Query Language

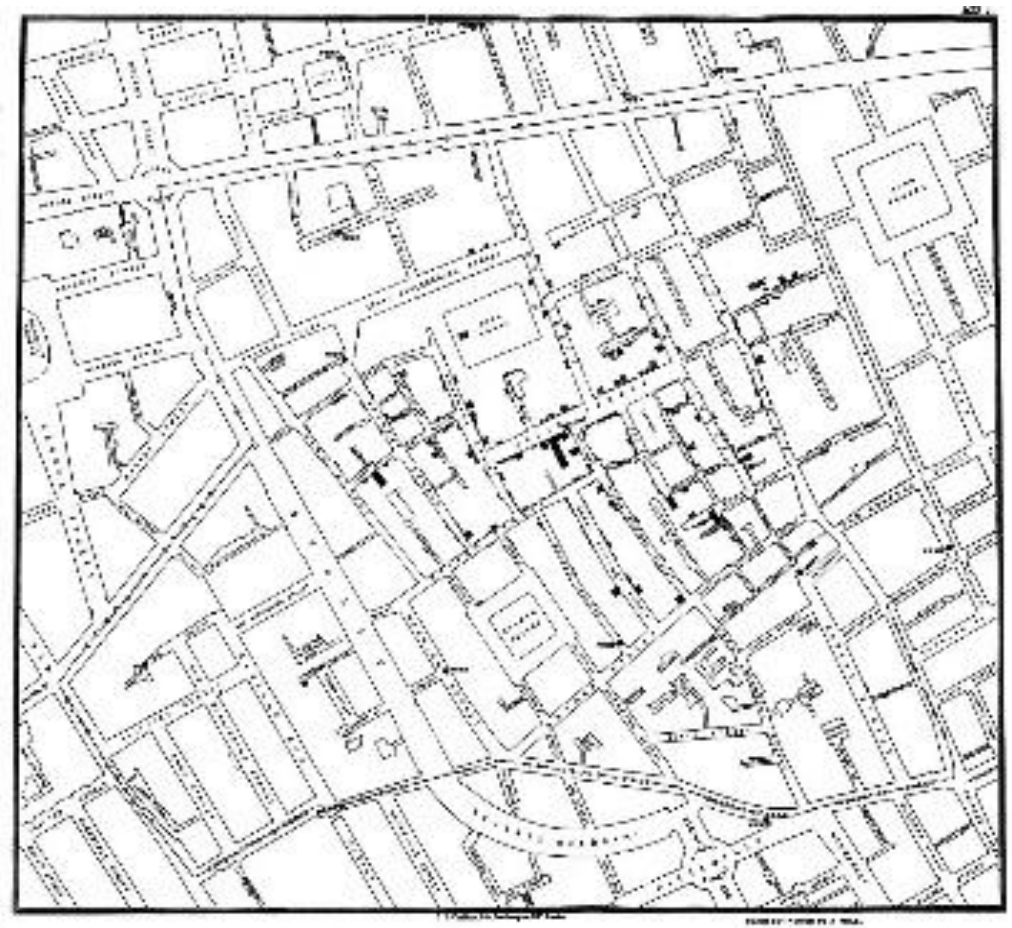
Having multiple tables that can be joined together is common in Relational Database Systems (RDBS).

- A common language used by RDBS is Structured Query Language (SQL)

dplyr	SQL
`inner_join(x, y, by = "z")`	`SELECT * FROM x INNER JOIN y USING (z)`
`left_join(x, y, by = "z")`	`SELECT * FROM x LEFT OUTER JOIN y USING (z)`
`right_join(x, y, by = "z")`	`SELECT * FROM x RIGHT OUTER JOIN y USING (z)`
`full_join(x, y, by = "z")`	`SELECT * FROM x FULL OUTER JOIN y USING (z)`

Let’s try it in R…

Spatial mapping

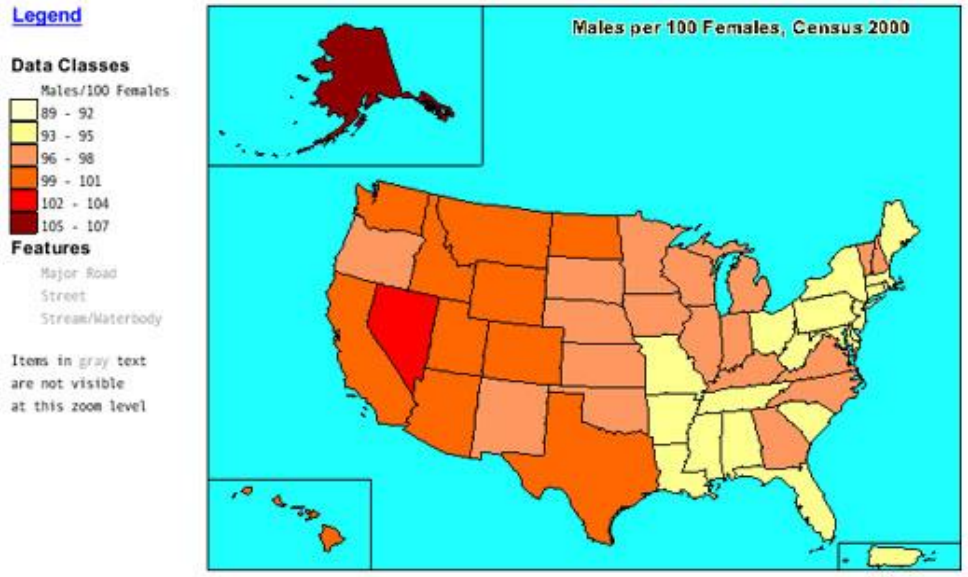


Maps

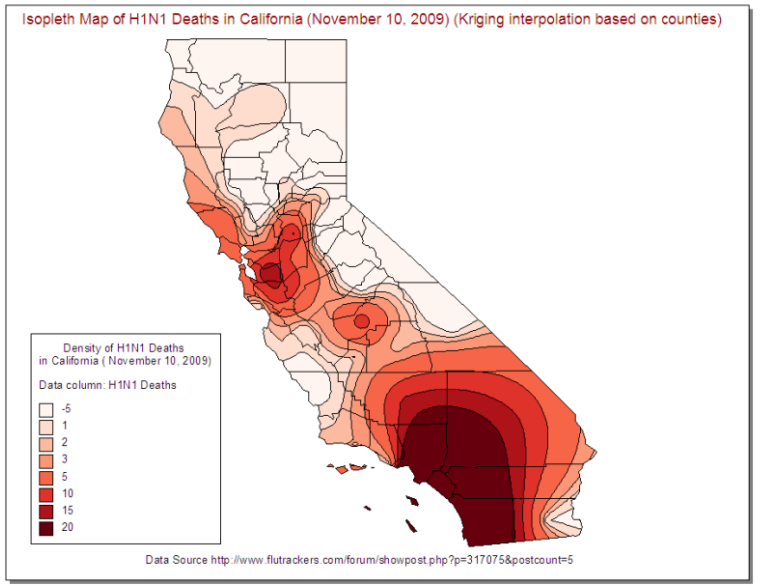
Choropleth maps: shades/colors in predefined areas based on properties of a variable

Isopleth maps: creates regions based on constant values

Choropleth map



Isopleth map



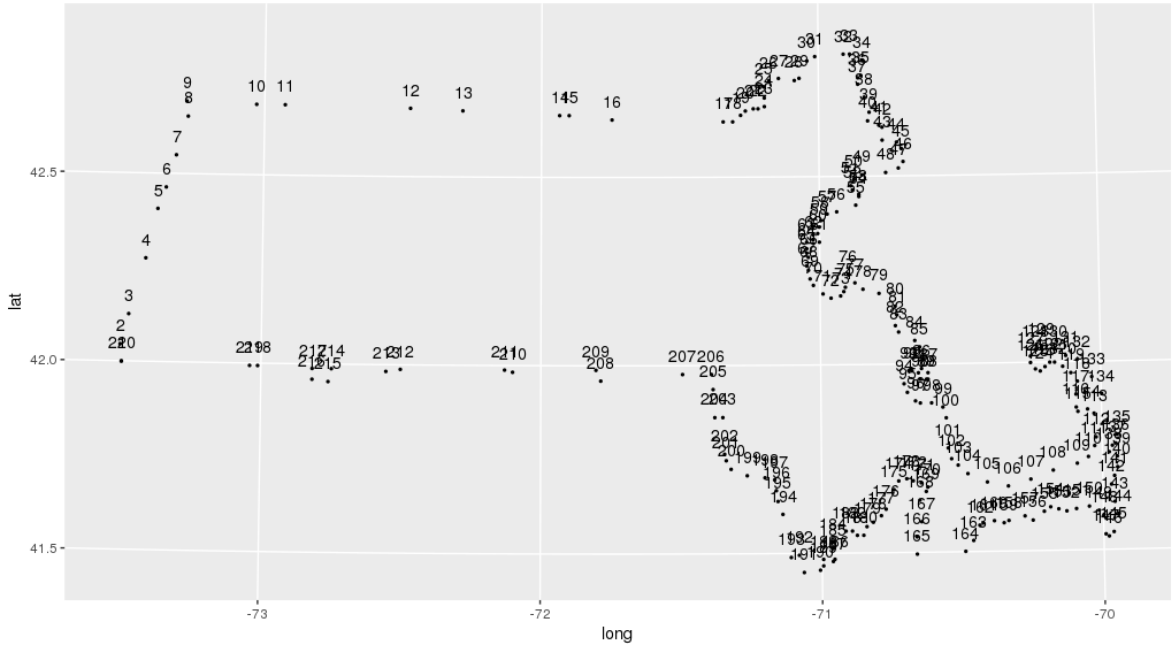
Choropleth maps

```
# has the coordinates for several maps
> library('maps')

# get a data frame with coordinates of states
> states_map <- map_data("state")
```

	long	lat	group	order	region	subregion
1	-87.46201	30.38968	1	1	alabama	*NA*
2	-87.48493	30.37249	1	2	alabama	*NA*
3	-87.52503	30.37249	1	3	alabama	*NA*
4	-87.53076	30.33239	1	4	alabama	*NA*
5	-87.57087	30.32665	1	5	alabama	*NA*

Choropleth maps

geom_polygon() works by connecting the dots:



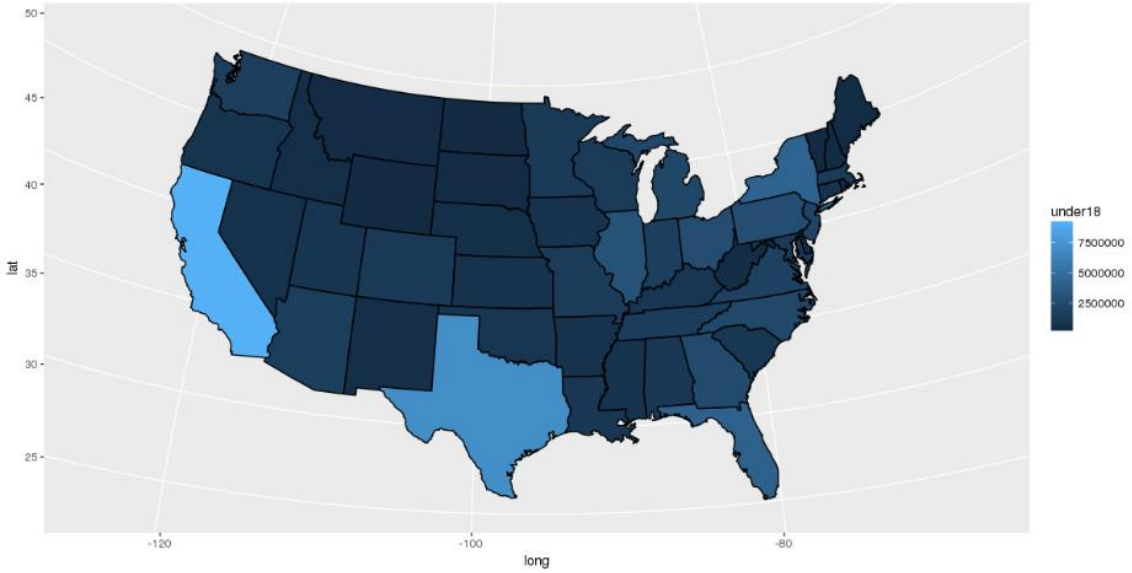
Often need to arrange points first: arrange(states_map, group, order)

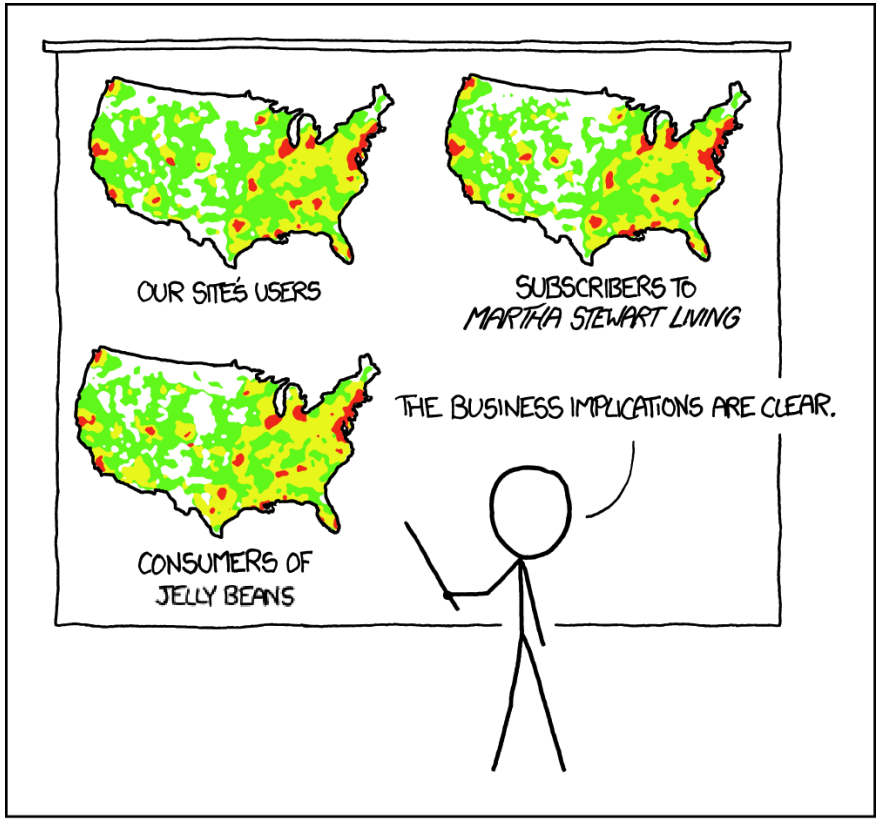
Choropleth maps

```
# has the coordinates for several maps
> library('maps')

# get a data frame with coordinates of states
> states_map <- map_data("state")

# filled white states with black borders
> ggplot(states_map,
              aes(x = long, y = lat, group = group)) +
              geom_polygon(fill = "white", color = "black")
```

Let’s try it in R!

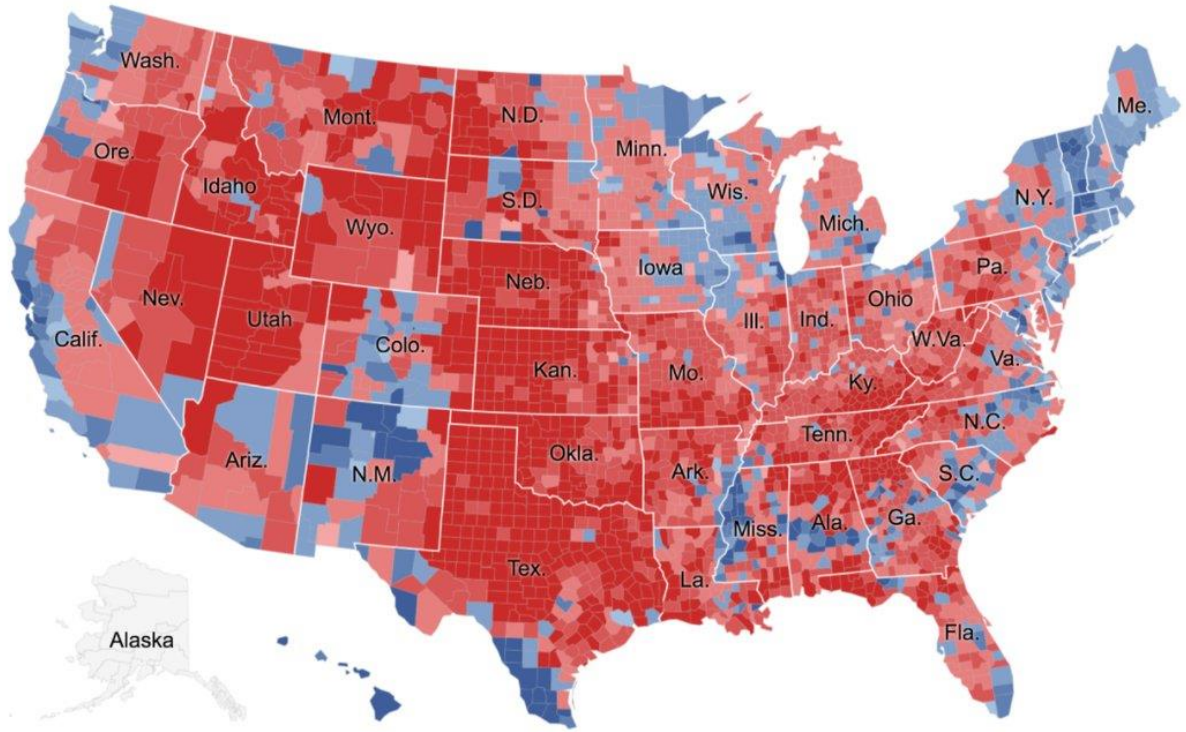
Pet Peeve #208





PET PEEVE #208:
GEOGRAPHIC PROFILE MAPS WHICH ARE BASICALLY JUST POPULATION MAPS

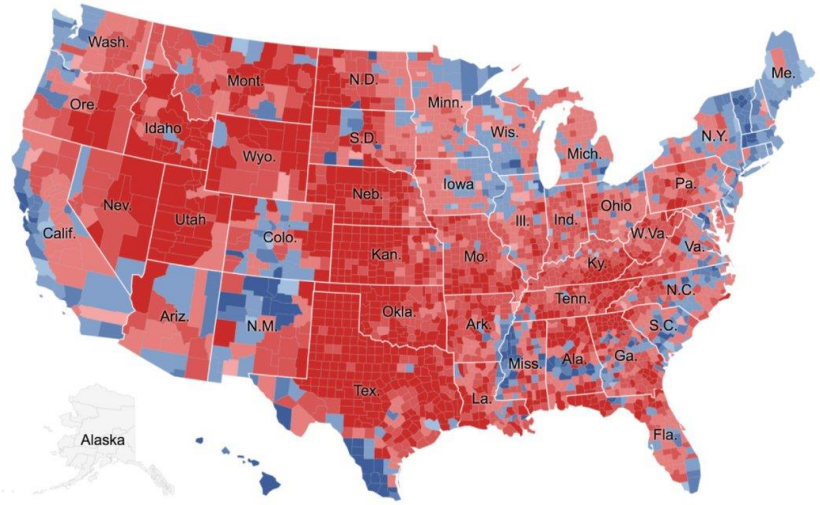
Question: in what way could this map be misleading?



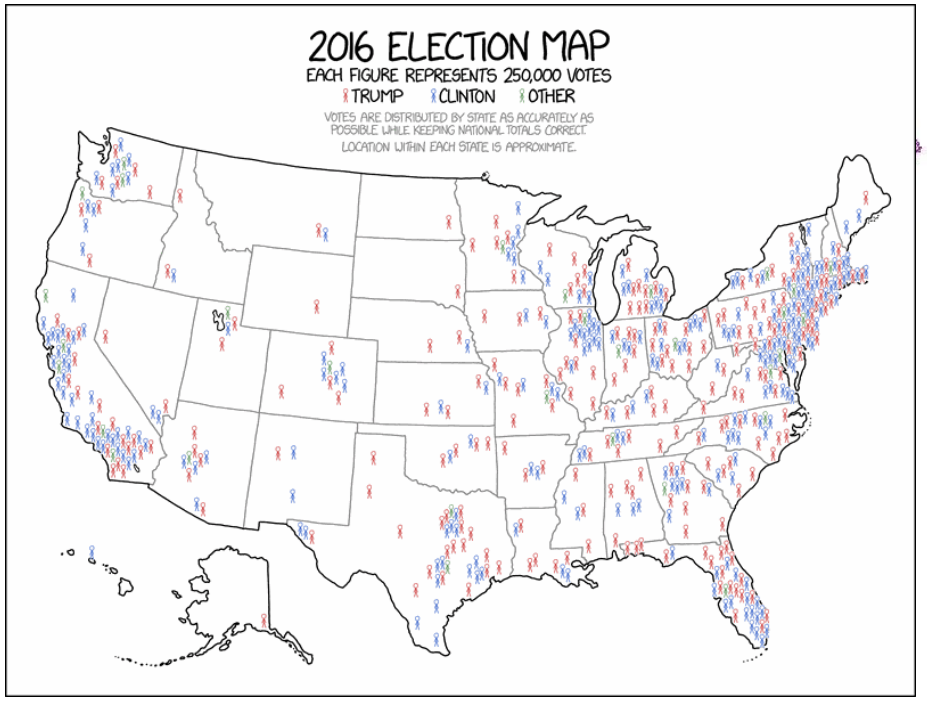
Darker red: county had higher % Trump vote

Darker blue: county had higher % Clinton vote

Cloropleth maps can be misleading



Looks like most of the country voted republican



Reshaping data

Wide vs. Long data

Plotting data using ggplot requires that data is in the right format

- i.e., requires data transformations.

Often this involves converting data from a **wide format** to **long format**

Wide data

Person	Age	Height
Bob	32	72
Alice	24	65
Steve	64	70

Long data

Person	name	value
Bob	Age	32
Bob	Height	72
Alice	Age	24
Alice	Height	65
Steve	Age	64
Steve	Height	70

```
library(tidyr)
```

tidyr::pivot_longer()

pivot_longer(df, cols) converts data from **wide** to **long**

- Takes multiple columns and converts them into two columns: name and value
  - Column names become categorical variable levels of a new variable called **name**
  - The data in rows become entries in a variable called **value**

Wide data

Person	Age	Height
Bob	32	72
Alice	24	65
Steve	64	70

Long data

Person	name	value
Bob	Age	32
Bob	Height	72
Alice	Age	24
Alice	Height	65
Steve	Age	64
Steve	Height	70

tidyr::pivot_wider()

pivot_wider(df, names_from, values_from) converts data from narrow to wide

- Turns the levels of categorical data into columns in a data frame

Narrow data

person	name	value
Bob	Age	32
Bob	Height	72
Alice	Age	24
Alice	Height	65
Steve	Age	64
Steve	Height	70

Wide data

Person	Age	Height
Bob	32	72
Alice	24	65
Steve	64	70

Let’s try it in R!