## Increase Your Impact: Visualizations in R

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### Visualizing Survey Data

Social scientists often work with data collected using a complex survey design. Survey instruments may be stratified by region or some other characteristic, contain replicate weights to make them comparable to a reference population, have a clustered structure, and so on. In Chapter 4 of the book of Healy, you can check how to calculate and then plot frequency tables of categorical variables, using some data from the General Social Survey (GSS). However, if we want accurate estimates of US households from the GSS, we will need to take the survey's design into account, and use the survey weights provided in the dataset. Thomas Lumley's survey library provides a comprehensive set of tools for addressing these issues. The tools and the theory behind them are discussed in detail in Lumley (2010), and an overview of the package is provided in Lumley (2004). While the functions in the survey package are straightforward to use and return results in a generally tidy form, the package predates the tidyverse and its conventions by several years. This means we cannot use survey functions directly with dplyr. However, Greg Freedman Ellis has written a helper package, srvyr, that solves this problem for us, and lets us use the survey library's functions within a data analysis pipeline in a familiar way.

For example, the 'gss\_lon data contains a small subset of measures from every wave of the GSS since its inception in 1972. It also contains several variables that describe the design of the survey and provide replicate weights for observations in various years. These technical details are described in the GSS documentation. Similar information is typically provided by other complex surveys. Here we will use this design information to calculate weighted estimates of the distribution of educational attainment by race, for selected survey years from 1976 to 2016.

To begin, we load the survey and srvyr libraries.

```
# remember, if you have not installed the packages yet, use install.packages()
library(survey)
library(tidyverse)
library(socviz)
```

Next, we take our gss\_lon dataset and use the survey tools to create a new object that contains the data, as before, but with some additional information about the survey's design:

The two options set at the beginning provide some information to the survey library about how to behave. You should consult Lumley (2010) and the survey package documentation for details. The subsequent operations create gss\_wt, an object with one additional column (stratvar), describing the yearly sampling strata. We use the interaction() function to do this. It multiplies the vstrat variable by the year variable to get a vector of stratum information for each year. We have to do this because of the way the GSS codes

its stratum information. In the next step, we use the as\_survey\_design() function to add the key pieces of information about the survey design. It adds information about the sampling identifiers (ids), the strata (strata), and the replicate weights (weights). With those in place we can take advantage of a large number of specialized functions in the survey library that allow us to calculate properly weighted survey means or estimate models with the correct sampling specification. For example, we can easily calculate the distribution of education by race for a series of years from 1976 to 2016. We use survey\_mean() to do this:

```
out_grp <- gss_wt %>%
  filter(year %in% seq(1976, 2016, by = 4)) %>%
  group_by(year, race, degree) %>%
  summarize(prop = survey_mean(na.rm = TRUE))
```

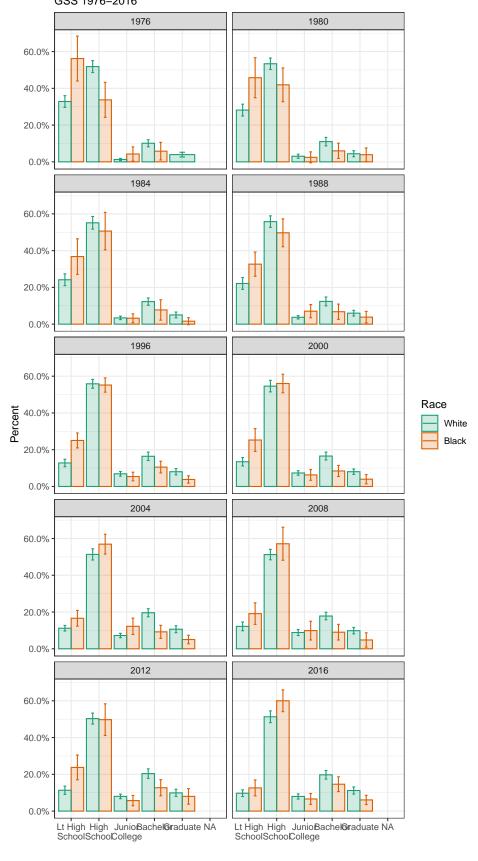
This gives us the numbers that we want and returns them in a tidy data frame. The interaction() function produces variable labels that are a compound of the two variables we interacted, with each combination of categories separated by a period, (such as White.Graduate). However, perhaps we would like to see these categories as two separate columns, one for race and one for education, as before. Because the variable labels are organized in a predictable way, we can use one of the convenient functions in the tidyverse's tidyr library to separate the single variable into two columns while correctly preserving the row values. Appropriately, this function is called separate().

```
out_mrg <- gss_wt %>%
  filter(year %in% seq(1976, 2016, by = 4)) %>%
  mutate(racedeg = interaction(race, degree)) %>%
  group_by(year, racedeg) %>%
  summarize(prop = survey_mean(na.rm = TRUE)) %>%
  separate(racedeg, sep = "\\.", into = c("race", "degree"))
```

The call to separate() says to take the racedeg column, split each value when it sees a period, and reorganize the results into two columns, race and degree. This gives us a tidy table much like 'out\_grp, but for the marginal frequencies.

Reasonable people can disagree over how best to plot a small multiple of a frequency table while faceting by year, especially when there is some measure of uncertainty attached. A barplot is the obvious approach for a single case, but when there are many years it can become difficult to compare bars across panels. This is especially the case when standard errors or confidence intervals are used in conjunction with bars. This is sometimes called a *dynamite plot*, not because it looks amazing but because the t-shaped error bars on the tops of the columns make them look like cartoon dynamite plungers. An alternative is to use a line graph to join up the time observations, faceting on educational categories instead of year.

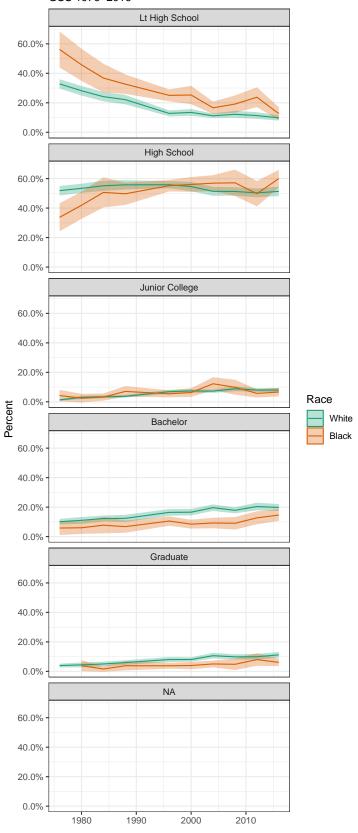
# Educational Attainment by Race GSS 1976–2016



This plot has a few cosmetic details and adjustments that you can look up in Healy's book (Chapter 8). Try to replicate the plot by peeling back the plot from the bottom, one instruction at a time, to see what changes. One useful adjustment to notice is the new call to the scales library to adjust the labels on the x-axis. The adjustment on the y-axis is scales::percent to convert the proportion to a percentage. On the x-axis, the issue is that several of the labels are rather long. If we do not adjust them they will print over one another. The scales::wrap\_format() function will break long labels into lines. It takes a single numerical argument (here 10) that is the maxmimum length a string can be before it is wrapped onto a new line.

A graph like this is true to the categorical nature of the data, while showing the breakdown of groups within each year. But you should experiment with some alternatives. For example, we might decide that it is better to facet by degree category instead, and put the year on the x-axis within each panel. If we do that, then we can use geom\_line() to show a time trend, which is more natural, and geom\_ribbon() to show the error range. This is perhaps a better way to show the data, especially as it brings out the time trends within each degree category, and allows us to see the similarities and differences by racial classification at the same time. Try to replicate the plot below too.

Educational Attainment by Race GSS 1976–2016



#### Additional Assignment to Practice with Making Graphs out of Survey Data

Lineplots are used to display time series. Use the economics dataset available in ggplot2 for this exercise. Show how the unemployment rate has changed since the 1960s in the US, and especially since the beginning of the financial crisis.

- 1. Based on the economics dataset, generate a new dataset (econ) that includes a variable for unemployment rate and a dummy variable that separates the time before the beginning of the financial crisis from the period after the crisis (as.Date(date) < "2008-10-01").
- 2. Draw a line plot that shows the unemployment rate over time.
- 3. Display years in 5- year steps on the x-axis.
- 4. Add a colored vertical line at October 2008 (more precisely: intercept = as.numeric(as.Date("2008-10-01"))).
- 5. Add the text "financial crisis" to the horizontal line.
- 6. Remove the title on x-axis

#### Visualizing Maps

R has a huge and growing number of spatial data packages. We recommend taking a quick browse on R's main website to see the spatial packages available: http://cran.r-project.org/web/views/Spatial.html.

In this tutorial we will use the following packages:

- ggmap: extends the plotting package ggplot2 for maps
- rgdal: R's interface to the popular C/C++ spatial data processing library gdal
- rgeos: R's interface to the powerful vector processing library geos
- maptools: provides various mapping functions

## Integer64 fields read as strings: Pop\_2001

- dplyr and tidyr: fast and concise data manipulation packages
- tmap: a new packages for rapidly creating beautiful maps

```
x <- c("ggmap", "rgdal", "rgeos", "maptools", "tidyverse", "tmap")
#install.packages(x) # warning: uncommenting this may take a number of minutes
lapply(x, library, character.only = TRUE) # load the required packages</pre>
```

The first file we are going to load into R Studio is the london\_sport shapefile. The data can be downloaded from: https://github.com/Robinlovelace/Creating-maps-in-R. It is worth looking at this input dataset in your file browser before opening it in R. You will notice that there are several files named "london\_sport", all with different file extensions. This is because a shapefile is actually made up of a number of different files, such as .prj, .dbf and .shp. You could also try opening the file"london\_sport.shp" file in a conventional GIS such as QGIS to see what a shapefile contains. Once you think you understand the input data, it's time to open it in R. There are a number of ways to do this, the most commonly used and versatile of which is readOGR. This function, from the rgdal package, automatically extracts the information regarding the data. rgdal is R's interface to the "Geospatial Abstraction Library (GDAL)" which is used by other open source GIS packages such as QGIS and enables R to handle a broader range of spatial data formats.

```
lnd <- readOGR(dsn = "/Users/velden/Desktop/GSSS/data", layer = "london_sport")

## OGR data source with driver: ESRI Shapefile

## Source: "/Users/velden/Desktop/GSSS/data", layer: "london_sport"

## with 33 features

## It has 4 fields</pre>
```

readOGR is a function which accepts two arguments: dsn which stands for "data source name" and specifies the directory in which the file is stored, and layer which specifies the file name (note that there is no need to include the file extention .shp). The arguments are separated by a comma and the order in which they are specified is important. You do not have to explicitly type dsn = or layer = as R knows which order they appear, so readOGR("data", "london\_sport") would work just as well. For clarity, it is good practice to include argument names when learning new functions so we will continue to do so. The file we assigned to the lnd object contains the population of London Boroughs in 2001 and the percentage of the population participating in sporting activities. This data originates from the Active People Survey. The boundary data is from the Ordnance Survey. For information about how to load different types of spatial data, see the help documentation for readOGR. This can be accessed by typing ?readOGR. For another worked example, in which a GPS trace is loaded, please see Cheshire and Lovelace (2014).

Spatial objects like the lnd object are made up of a number of different slots, the key slots being <code>@data</code> (non geographic attribute data) and <code>@polygons</code> (or <code>@lines</code> for line data). The data slot can be thought of as an attribute table and the geometry slot is the polygons that make up the physical boundaries. Specific slots are accessed using the <code>@ symbol</code>. Let's now analyse the sport object with some basic commands:

```
head(lnd@data, n = 2)
mean(lnd$Partic_Per)
```

Now we have seen something of the structure of spatial objects in R, let us look at plotting them. Note, that plots use the geometry data, contained primarily in the **@polygons** slot.

#### plot(lnd)



plot is one of the most useful functions in R, as it changes its behaviour depending on the input data (this is called *polymorphism* by computer scientists). Inputting another object such as plot(lnd@data) will generate an entirely different type of plot. Thus R is intelligent at guessing what you want to do with the data you provide it with.

R has powerful subsetting capabilities that can be accessed very concisely using square brackets, as shown in the following example:

```
# select rows of lnd@data where sports participation is less than 15
lnd@data[lnd$Partic_Per < 15, ] # we don't use the tidyverse verb select,</pre>
```

##		ons_label		name	Partic_Per	Pop_2001
##	17	OOAQ		Harrow	14.8	206822
##	21	OOBB		Newham	13.1	243884
##	32	AAOO	City of	London.	9.1	7181

#### # because it doesn't work well with polygons

The above line of code asked R to select only the rows from the 1nd object, where sports participation is lower than 15, in this case rows 17, 21 and 32, which are Harrow, Newham and the city centre respectively. The square brackets work as follows: anything before the comma refers to the rows that will be selected, anything after the comma refers to the number of columns that should be returned. For example if the data frame had 1000 columns and you were only interested in the first two columns you could specify 1:2 after the comma. The: symbol simply means "to", i.e. columns 1 to 2. Try experimenting with the square brackets notation (e.g. guess the result of lnd@data[1:2, 1:3] and test it). So far we have been interrogating only the attribute data slot (@data) of the lnd object, but the square brackets can also be used to subset spatial objects, i.e. the geometry slot. Using the same logic as before try to plot a subset of zones with high sports participation. Try to replicate the following graph:

```
plot(lnd, col = "lightgrey") # plot the london_sport object
sel <- lnd$Partic_Per > 25
plot(lnd[ sel, ], col = "turquoise", add = TRUE)
```



You have just interrogated and visualised a spatial object: where are areas with high levels of sports participation in London? The map tells us. Do not worry for now about the intricacies of how this was achieved.

As a bonus stage, select and plot only zones that are close to the centre of London. Programming encourages rigorous thinking and it helps to define the problem more specifically:

**Challenge**: Select all zones whose geographic centroid lies within 10 km of the geographic centroid of inner London.<sup>1</sup>

<sup>&</sup>lt;sup>1</sup>To see how this map was created, see the code in README.Rmd. This may be loaded by typing file.edit("README.Rmd") or online at github.com/Robinlovelace/Creating-maps-in-R/blob/master/README.Rmd.



Figure 1: Zones in London whose centroid lie within 10 km of the geographic centroid of the City of London. Note the distinction between zones which only touch or 'intersect' with the buffer (light blue) and zones whose centroid is within the buffer (darker blue).

The code below should help understand the way spatial data work in R.

```
# Find the centre of the london area
easting_lnd <- coordinates(gCentroid(lnd))[[1]]
northing_lnd <- coordinates(gCentroid(lnd))[[2]]
# arguments to test whether or not a coordinate is east or north of the centre
east <- sapply(coordinates(lnd)[,1], function(x) x > easting_lnd)
north <- sapply(coordinates(lnd)[,2], function(x) x > northing_lnd)
# test if the coordinate is east and north of the centre
lnd$quadrant <- "unknown" # prevent NAs in result
lnd$quadrant[east & north] <- "northeast"</pre>
```

**Challenge**: Based on the the above code as reference try and find the remaining 3 quadrants and colour them. Hint - you can use the **llgridlines** function in order to overlay the long-lat lines. Try to desolve the quadrants so the map is left with only 4 polygons.

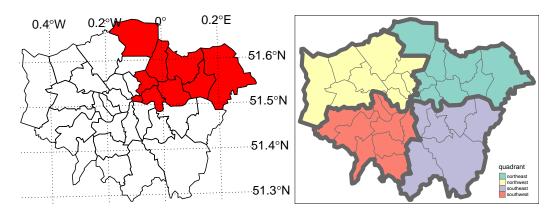


Figure 2: The 4 quadrants of London and dissolved borders. Challenge: recreate a plot that looks like this.

Alongside visualisation and interrogation, a GIS must also be able to create and modify spatial data. R's spatial packages provide a very wide and powerful suite of functionality for processing and creating spatial data.

Reprojecting and joining/clipping are fundamental GIS operations, so in this section we will explore how these operations can be undertaken in R. Firstly, we will join non-spatial data to spatial data so it can be mapped. Finally we will cover spatial joins, whereby information from two spatial objects is combined based on spatial location.

R objects can be created by entering the name of the class we want to make. vector and data.frame objects for example, can be created as follows:

```
vec <- vector(mode = "numeric", length = 3)
df <- data.frame(x = 1:3, y = c(1/2, 2/3, 3/4))</pre>
```

We can check the class of these new objects using class():

class(vec)

```
## [1] "numeric"
```

class(df)

## [1] "data.frame"

The same logic applies to spatial data. The input must be a numeric matrix or data.frame:

```
sp1 <- SpatialPoints(coords = df)</pre>
```

We have just created a spatial points object, one of the fundamental data types for spatial data. (The others are lines, polygons and pixels, which can be created by SpatialLines, SpatialPolygons and SpatialPixels, respectively.) Each type of spatial data has a corollary that can accepts non-spatial data, created by adding DataFrame. SpatialPointsDataFrame(), for example, creates points with an associated data.frame. The number of rows in this dataset must equal the number of features in the spatial object, which in the case of sp1 is 3.

```
class(sp1)
```

```
## [1] "SpatialPoints"
## attr(,"package")
## [1] "sp"
spdf <- SpatialPointsDataFrame(sp1, data = df)
class(spdf)</pre>
```

```
## [1] "SpatialPointsDataFrame"
## attr(,"package")
## [1] "sp"
```

The above code extends the pre-existing object sp1 by adding data from df. To see how strict spatial classes are, try replacing df with mat in the above code: it causes an error. All spatial data classes can be created in a similar way, although SpatialLines and SpatialPolygons are much more complicated (Bivand et al. 2013). More frequently your spatial data will be read-in from an externally-created file, e.g. using readOGR(). Unlike the spatial objects we created above, most spatial data comes with an associate 'CRS'.

The Coordinate Reference System (CRS) of spatial objects defines where they are placed on the Earth's surface. You may have noticed 'proj4string 'in the summary of 1nd above: the information that follows represents its CRS. Spatial data should always have a CRS. If no CRS information is provided, and the correct CRS is known, it can be set as follow:

```
proj4string(lnd) <- NA_character_ # remove CRS information from lnd
proj4string(lnd) <- CRS("+init=epsg:27700") # assign a new CRS</pre>
```

R issues a warning when the CRS is changed. This is so the user knows that they are simply changing the CRS, not *reprojecting* the data. An easy way to refer to different projections is via EPSG codes.

Under this system 27700 represents the British National Grid. 'WGS84' (epsg:4326) is a very commonly used CRS worldwide. The following code shows how to search the list of available EPSG codes and create a new version of 1nd in WGS84:<sup>2</sup>

Above, spTransform converts the coordinates of lnd into the widely used WGS84 CRS. Now we've transformed lnd into a more widely used CRS, it is worth saving it. R stores data efficiently in .RData or .Rds formats. The former is more restrictive and maintains the object's name, so we use the latter.

#### References

Healy, K. (2018). Data visualization: a practical introduction. Princeton University Press. (https://socviz.co/)
Robin Love Lace's Github https://github.com/Robinlovelace/Creating-maps-in-R

<sup>&</sup>lt;sup>2</sup>Note: entering projInfo() provides additional CRS options. spatialreference.org provides more information about EPSG codes.