mapping_R Markup

Introduction

This tutorial walks through three examples demonstrating how geospatial data can be manipulated and visualized in R. We assume you have downloaded R and know how to run R commands. For those unfamiliar with R or GIS mapping, below are a few introductory resources for exploration:

- R
    - *Tutorials for learning R* on r-bloggers.com
    - *Quick R* by Robert I. Kabacoff, Ph.D.
    - *R for Data Science* by Garrett Grolemund and Hadly Wickham
- GIS mapping
    - *A Gentle Introduction to GIS* by QGIS; map projections in particular
    - *Introduction to GIS* by the International Centre for Integrated Mountain Development; information about map projections on p.72

As you go through the tutorial, any time you'd like to see more information about a function, place a "?" before its name:

```
`?`(county_choropleth_acs)
```

R packages for geospatial data

Each example within this tutorial introduces a variety of R packages to visualize maps in R. Below we provide short descriptions of the R packages used in each example throughout this tutorial, which can to be installed with the following code:

```
install.packages(c("acs", "choroplethr", "choroplethrMaps", "maptools", "rgeos",
    "mapproj", "RColorBrewer", "maps", "geosphere", "rgdal", "reshape", "ggplot2"))
```

Packages used in:

- EXAMPLE 1: Mapping Census Data
    - **ACS** & **choroplethr** - can be used together to easily make choroplethGCT-PEPANNRES maps using data from the American Community Survey (ACS), yearly census data collected by the U.S. Census Bureau. To access ACS data you need an API key. Visit http://api.census.gov/data/key_signup.html, request a key, and paste it into the line below:
    - **choroplethrMaps** - contains a global map and maps of the USA used by the choroplethr package.
- EXAMPLE 2: Working with Shapefiles, Projections, and Visualization
    - **maptools** - contains functions for reading and manipulating geographic data, including ESRI shapefiles.
    - **rgdal** - geospatial data abstraction and projection / transformation.
    - **RColorBrewer** - provides color schemes that are especially useful for creating thematic maps.
    - **ggplot2** - package for creating and customizing graphics is R.
    - **rgeos** - contains functions for performing geometric analysis. For example `gLength()` calculates the length of input geometry, while `gBuffer()` adds a buffer to an input feature.

- **mapproj** - simple package for converting from latitude and logitude into projected coordinates.

- EXAMPLE 3: Migration Distances Map

    - **maps** another simple set of tools for creating maps, with links to several databases of spatial data.
    - **geosphere** - supports trigonometric calculations for geographic applications. For example, computing distance to the horizon from a given location and altitude.
    - **reshape** - reshapes data from 'wide' format (where repeated measurements are located across multiple columns) to 'long' format (where repeated measurements are spread across unique rows)
    - **mapproj** - simple package for converting from latitude and logitude into projected coordinates.

EXAMPLE 1: Mapping Census Data

This example introduces how to access and plot American Community Survey (ACS) yearly census data by the U.S. Census Bureau through choropleth maps.

Packages used: **ACS**, **choroplethr**, **choroplethrMaps**
After installing packages (see introduction), load needed packages:

```
library(acs)
library(choroplethr)
library(choroplethrMaps)
```

We need an api key to access the ACS data. Visit http://api.census.gov/data/key_signup.html, request a key, and paste it into the line below:

```
api.key.install("<ACS API key>")
```

Great, now we have access to the census data. Table B19301 contains per capita income for the year of 2011. Lets plot it!

```
county_choropleth_acs(tableId = "B19301")
```

To see the description of a function and its arguments in R, remember you can place a "?" before its name:

```
`?`(county_choropleth_acs)
```

You can explore the ACS data on the Census Bureau's website: http://factfinder.census.gov/faces/nav/jsf/pages/searchresults.xhtml Be sure to set the dataset to one of the ACS sources under Topics -> dataset. Use the ID column to find your tableID variable. Here's another example to get you started playing around with these functions:

```
state_choropleth_acs("B01002", num_colors = 1, zoom = c("texas", "louisiana",
    "mississippi", "alabama", "georgia", "florida"))
```

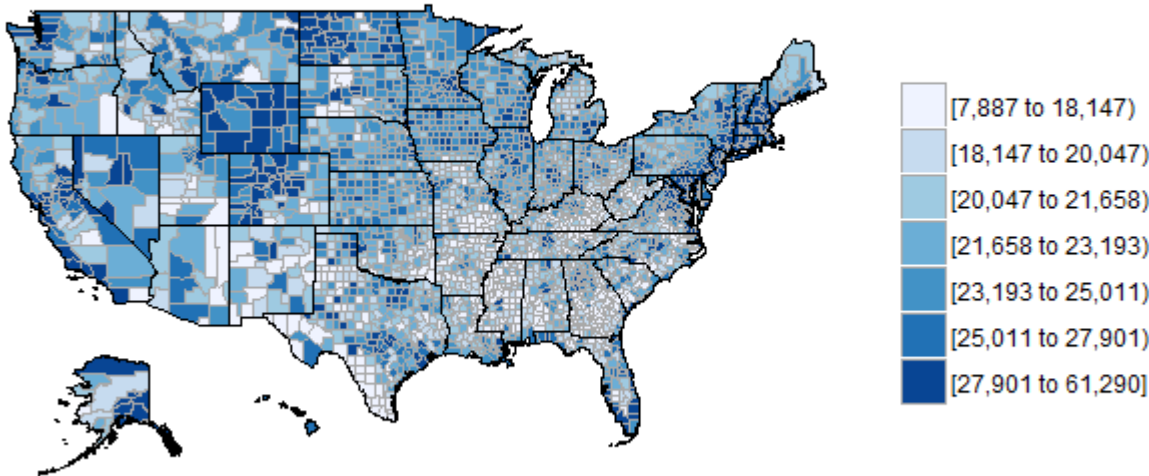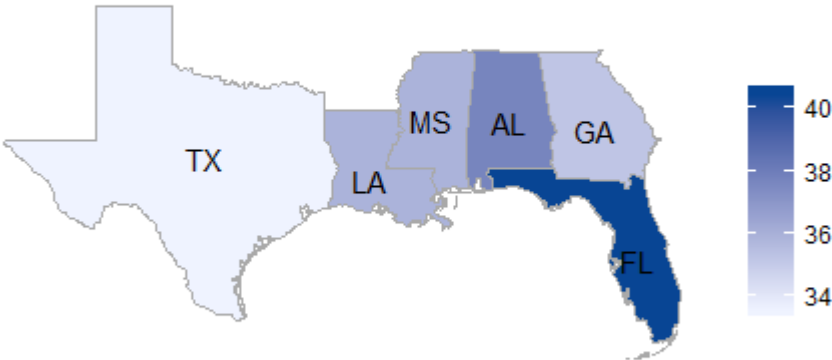**Per capita income in the past 12 months (in 2011 inflation-adjusted dollars)**

	[7,887 to 18,147)
	[18,147 to 20,047)
	[20,047 to 21,658)
	[21,658 to 23,193)
	[23,193 to 25,011)
	[25,011 to 27,901)
	[27,901 to 61,290]

Figure 1: US county chloropleth using ACS per capita income data

Median Age by Sex: Median age -- Total:

EXAMPLE 2: Working with Shapefiles, Projections, and Visualization

This example loads and plots both polygon and point shapefiles, uses the `rgdal` library to change projections, uses the `over` function to join datasets, and provides a few visualization examples.

Packages used: **maptools, rgdal, RColorBrewer, ggplot2**
After installing packages (see introduction), load needed packages:

```
library(maptools)
library(rgdal)
library(RColorBrewer)
library(ggplot2)
```

In this example we will work with the data provided along with this tutorial. **Make sure you have unzipped the folder county_census before proceeding!** The following prompts you to select the provided county census shapefiles at the path . . . county_census/County_2010Census_DP1.shp.

```
counties <- readShapeSpatial(file.choose(), proj4string = CRS("+proj=longlat +datum=WGS84"))
```

Note that the second parameter of the maptools function `readShapeSpatial` was a string representing the projection of the data, called a `prj4` string. So far we've used "+proj=longlat +datum=WGS84", which is simply unprojected longitude and latitude coordinates.

Inspect the first few rows of the counties data to get a feel for its structure:

```
head(counties@data)
```

Census data assigns codes to counties using the Federal Information Processing Standard (FIPS). A FIPS code starts with two digits representing the state, and is followed by three digits representing the county. For example, Florida is 12 and Clay County Florida is 12019. To select all the counties in Florida we can use a regular expression matching all codes that start with "12":

```
florida <- counties[substring(counties$GEOID10, 1, 2) == "12", ]
plot(florida)
```


You can look up other state and county codes using the U.S. Census Bureau site: https://www.census.gov/geo/reference/codes/cou.html

Projection and Layering with RGDAL

Next we'll work more with projections using library `rgdal`. We're going to read in a shape file of cultural points in Florida from the supplied data, again using function `readShapeSpatial()`. We know already that our cultural centers layer uses NAD83(HARN) / Florida GDL Albers. We can make the EPSG data frame of projections to find the `prj4` string for this projection (use `?make_EPSG()` to find out more about this table):

```
library(rgdal)
EPSG <- make_EPSG()
```

We can use regular expressions to search the note field of `EPSG` for any that refer to Florida:

```
EPSG[grep("florida", EPSG$note, ignore.case = TRUE), 1:2]
```

```
##        code                                  note
## 705    2236            # NAD83 / Florida East (ftUS)
## 706    2237            # NAD83 / Florida West (ftUS)
## 707    2238            # NAD83 / Florida North (ftUS)
## 1245   2777              # NAD83(HARN) / Florida East
## 1246   2778              # NAD83(HARN) / Florida West
## 1247   2779             # NAD83(HARN) / Florida North
## 1349   2881        # NAD83(HARN) / Florida East (ftUS)
## 1350   2882        # NAD83(HARN) / Florida West (ftUS)
## 1351   2883       # NAD83(HARN) / Florida North (ftUS)
## 1553   3086               # NAD83 / Florida GDL Albers
## 1554   3087         # NAD83(HARN) / Florida GDL Albers
## 1978   3511          # NAD83(NSRS2007) / Florida East
## 1979   3512  # NAD83(NSRS2007) / Florida East (ftUS)
## 1980   3513   # NAD83(NSRS2007) / Florida GDL Albers
## 1981   3514         # NAD83(NSRS2007) / Florida North
## 1982   3515 # NAD83(NSRS2007) / Florida North (ftUS)
## 1983   3516          # NAD83(NSRS2007) / Florida West
## 1984   3517  # NAD83(NSRS2007) / Florida West (ftUS)
## 3074   6437             # NAD83(2011) / Florida East
## 3075   6438       # NAD83(2011) / Florida East (ftUS)
## 3076   6439        # NAD83(2011) / Florida GDL Albers
## 3077   6440            # NAD83(2011) / Florida North
## 3078   6441      # NAD83(2011) / Florida North (ftUS)
## 3079   6442             # NAD83(2011) / Florida West
## 3080   6443       # NAD83(2011) / Florida West (ftUS)
## 3747  26758                  # NAD27 / Florida East
## 3748  26759                  # NAD27 / Florida West
## 3749  26760                 # NAD27 / Florida North
## 3895  26958                  # NAD83 / Florida East
## 3896  26959                  # NAD83 / Florida West
## 3897  26960                 # NAD83 / Florida North
```

We see the code we're looking for is 3087. Extract the `prj4` string from this dataframe:

```
subset(EPSG, code == 3087)
prjstring <- subset(EPSG, code == 3087)$prj4
```

Inspect our `prjstring` variable if you want to see the format of the `prj4` variable.

Now that we have the appropriate `prj4` we can read in the cultural centers data. The following prompts you to select the shape file. Select the actual `.shp` file in the provided data from ...cultural_centers/gc_culturecenter_oct15.shp.

```r
cultural <- readShapeSpatial(file.choose(), proj4string = CRS(prjstring))
```

Before we overlay the cultural points, we need to transform this layer to match that of the Florida counties layer - simple longitude and latitude in WGS84:

```r
cultural_proj <- spTransform(cultural, CRS("+proj=longlat +datum=WGS84"))

plot(florida)
points(cultural_proj)
```


You can play around with the symbology for your map with some additional arguments in the `points` function. For example:

```r
plot(florida)
points(cultural_proj, cex = 0.8, pch = 24, col = 554, bg = "grey")
```
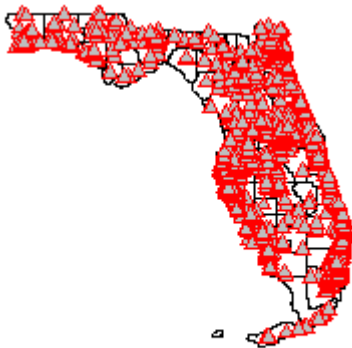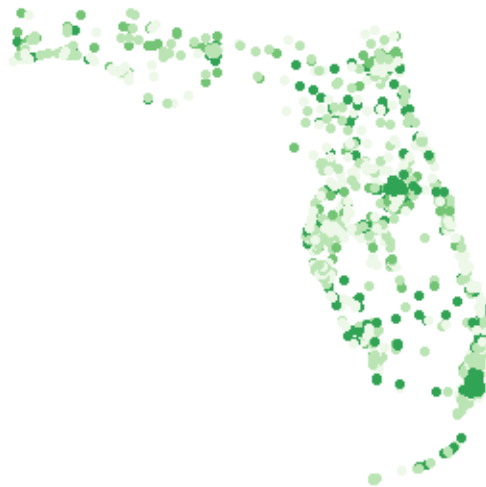
join polygon data to points

```
county_data <- over(cultural_proj, florida)
cultural_proj$pop <- county_data$DP0010001
```

set colors

```
library(RColorBrewer)
```

```
brks <- c(0.5, 1, 1.5, 2) * 1e+06
cols <- brewer.pal(5, "Greens")

mapcols <- cols[findInterval(cultural_proj$pop, vec = brks)]
plot(cultural_proj, col = mapcols, pch = 20)
```
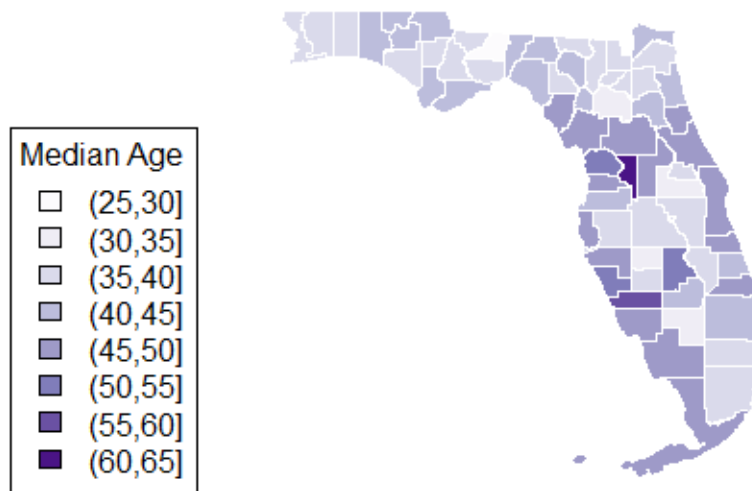

base R instructions for choropleth

```
brks <- c(25, 30, 35, 40, 45, 50, 55, 60, 65)
cols <- brewer.pal(8, "Purples")

mapcols <- cols[findInterval(florida$DP0020001, vec = brks)]
plot(florida, col = mapcols, border = "white")

legend("bottomleft", legend = levels(cut(florida$DP0020001, brks)), fill = cols,
    title = "Median Age")
```
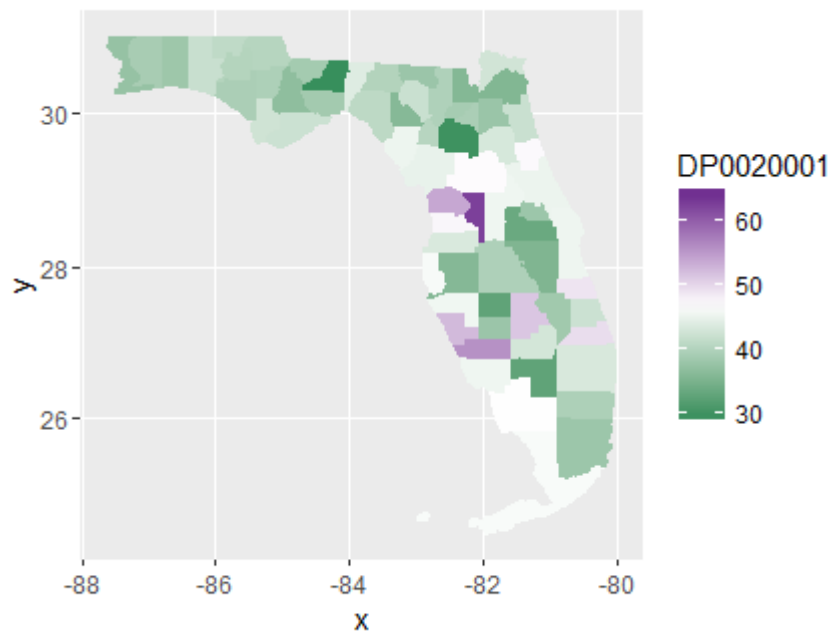
using ggplot2

```
library(ggplot2)
```

```
fl_shapes <- fortify(florida, region = "GEOID10")

ggplot() + geom_map(data = as.data.frame(florida), aes(map_id = GEOID10, fill = DP0020001),
    map = fl_shapes) + expand_limits(x = fl_shapes$long, y = fl_shapes$lat) +
    scale_fill_gradient2(low = "seagreen", mid = "white", high = "darkorchid4",
        midpoint = 47, limits = c(29, 65)) + coord_map(projection = "mercator")
```

EXAMPLE 3: Migration Distances Map

This example loads both a shapefile and .csv file, transforms the data set with a melt function, and writes a function that maps the distance data contained in the .csv file.

Packages used: **maps, geosphere, reshape, maptools**
After installing packages (see introduction), load needed packages:

```
library(maps)
library(geosphere)
library(reshape)
library(maptools)
```

Import shapefile and .csv file

Import shapefile of map of continental United States. Choose `state_shapes/tl_2014_us_state.shp` file when prompted.

```
state <- readShapeSpatial(file.choose())
```

Import data file of migration distances between U.S.A. states. Choose `state_migrations_2014.csv` file when prompted.

```
migration <- read.csv(file.choose())
```

Extract state names and geographic coordinate (latitude and longitude) information from `state` shapefile; save it into a data frame called `centrs`.

```
centrs <- data.frame(as.character(state@data$NAME), coordinates(state))
colnames(centrs) <- c("name", "long", "lat")
```

Reshape data with melt function

Redefine the `migration` data set to only include columns 1 & 6-through-56 of data. Then use `melt` function from `reshape` package to transform data set into rows representing unique instances of data, based on a selected variable id (in our case, the `from_state` variable). *For more on the `melt` function, see https://www.r-bloggers.com/melt/.*

```
migration <- migration[c(1, 6:56)]
long_mig <- melt(migration, id.vars = "from_state")
```

Create a mapping function

Create a function named `data_from_state` that maps migration distances from any state selected. NOTE: To use this function, a map must be drawn (i.e. a new plot must be called) first.

```
draw_from_state <- function(centrs, migrations, state_name, color = rgb(0, 0,
    0, alpha = 0.5)) {
    migrations$variable <- sub(".", " ", migrations$variable, fixed = TRUE)
    migrations <- migrations[migrations$variable == state_name & migrations$from_state !=
        state_name, ]
    for (i in 1:nrow(migrations)) {
        if (nrow(centrs[centrs$name == as.character(migrations[i, ]$from_state),
            ]) > 0) {
            from_long <- centrs[centrs$name == as.character(migrations[i, ]$from_state),
                ]$long
            from_lat <- centrs[centrs$name == as.character(migrations[i, ]$from_state),
                ]$lat
            to_long <- centrs[centrs$name == as.character(migrations[i, ]$variable),
                ]$long
            to_lat <- centrs[centrs$name == as.character(migrations[i, ]$variable),
                ]$lat
            number <- migrations[i, ]$value
            lines(gcIntermediate(c(from_long, from_lat), c(to_long, to_lat),
                n = 50, addStartEnd = TRUE), lwd = sqrt(number)/50, col = color)
        }
    }
}
```
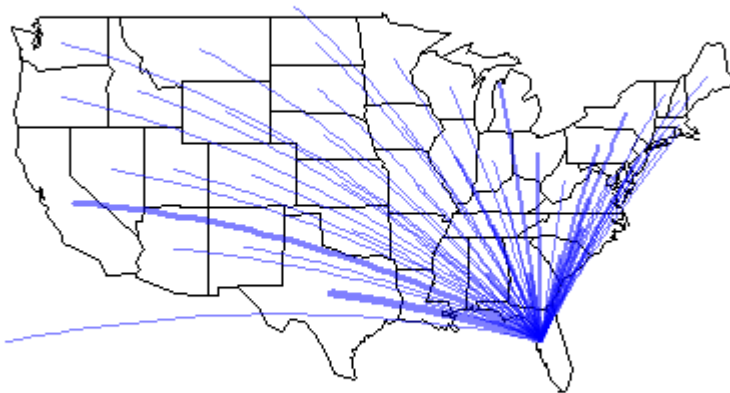
Using our mapping function: example 1

First draw imported `state` shapefile.

```
map("state")
```

Now use written function to map migration distances from Florida.

```
draw_from_state(centrs, long_mig, "Florida", rgb(0, 0, 1, 0.5))
```

Using our mapping function: example 2

First draw a world map (limited to North and Central America by creating x- and y-coordinate limits), and subsquently use written function to map migration distances from Wyoming onto map.

```r
xlim <- c(-171.738281, -56.601563)
ylim <- c(12.039321, 71.856229)
map("world", col = "#f2f2f2", fill = TRUE, bg = "white", lwd = 0.05, xlim = xlim,
    ylim = ylim)
draw_from_state(centrs, long_mig, "Wyoming", rgb(1, 0, 0, 0.5))
```