# mapping\_R Markup

### Setup

The following are packages useful for working with GIS data in R. They only nee to be installed once!

install.packages(c("acs","choroplethr","choroplethrMaps","maptools","rgeos","mapproj","RColorBrewer","m

We start with a simple example plotting a cloropleth map using data from the American Community Survey (ACS), yearly census data collected by the U.S. Census Bureau. To access and plot this we need to load the following libraries:

library(acs)
library(choroplethr)

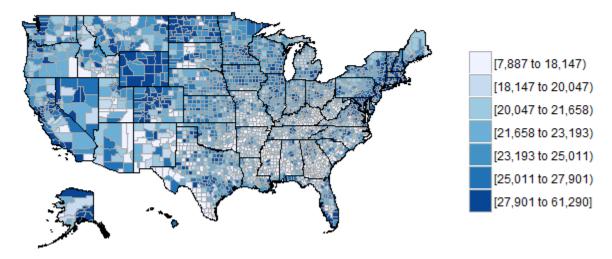
We need an api key to access the ACS data. Visit http://api.census.gov/data/key\_signup.html, request a key, and paste it into the line below:

api.key.install("e3dd607b83adce3268ef2bb723da22c68001e6f0")

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")

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



To see the description of a function and its arguments in R, place a "?" before its name: ?county\_choropleth\_acs

### Reading in Shapefiles

Load maptools, a library for reading and manipulating geographic data, including ESRI shapefiles.

library(maptools)

The following prompts you to select the provided county census shapefiles at the path . . . county\_census/Count\_2010Census\_DF Note: You will have to unzip the folder county\_census first!

```
"+proj=longlat +datum=WGS84" = unprojected prj4
```

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

Let's 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. So 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)</pre>



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 with library rgdal, a package for working with projections and transformations of geospatial data. We're going to read in a shape file of cultural points in Florida from the supplied data using function readShapeSpatial(). This function takes a prj4, a string containing the projection information of the shapefile, as an argument. We know already that our shapefile 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()</pre>
```

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
2236	# NAD83 / Florida East (ftUS)
2237	# NAD83 / Florida West (ftUS)
2238	# NAD83 / Florida North (ftUS)
2777	# NAD83(HARN) / Florida East
2778	# NAD83(HARN) / Florida West
2779	# NAD83(HARN) / Florida North
2881	# NAD83(HARN) / Florida East (ftUS)
2882	# NAD83(HARN) / Florida West (ftUS)
2883	# NAD83(HARN) / Florida North (ftUS)
3086	# NAD83 / Florida GDL Albers
3087	# NAD83(HARN) / Florida GDL Albers
3511	# NAD83(NSRS2007) / Florida East
3512	# NAD83(NSRS2007) / Florida East (ftUS)
3513	# NAD83(NSRS2007) / Florida GDL Albers
3514	# NAD83(NSRS2007) / Florida North

We see the code is 3087. Extract the prj4 string from this dataframe:

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

points(cultural\_proj)

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

### select cultural shapefile in cultural\_centers

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.

```
cultural <- readShapeSpatial(file.choose(),proj4string=CRS(prjstring))</pre>
```

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:

```
cultural_proj <- spTransform(cultural, CRS("+proj=longlat +datum=WGS84"))
plot(florida)</pre>
```



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

```
plot(florida)
points(cultural_proj, cex=.8, pch=24, col=554, bg="grey")
```



## join polygon data to points

county\_data <- over(cultural\_proj,florida)
cultural\_proj\$pop <- county\_data\$DP0010001</pre>

### set colors

```
library(RColorBrewer)

brks <- c(.5,1,1.5,2) * 10000000

cols <- brewer.pal(5,"Greens")

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

```
base R instructions for choropleth
brks <- c(25,30,35,40,45,50,55,60,65)
cols <- brewer.pal(8,"Purples")</pre>
mapcols <- cols[findInterval(florida$DP0020001, vec=brks)]</pre>
plot(florida,col=mapcols,border="white")
legend("bottomleft", legend = levels(cut(florida$DP0020001, brks)), fill = cols, title = "Median Age")
using ggplot2
library(ggplot2)
fl_shapes <- fortify(florida,region="GEOID10")</pre>
ggplot() + geom_map(data=as.data.frame(florida),aes(map_id = GEOID10,fill=DP0020001), map=fl_shapes) +
networky type example
library(maps)
library(geosphere)
library(reshape)
select - state shapes/tl 2014 us state.shp
state <- readShapeSpatial(file.choose())</pre>
select - /state_migrations_2014.csv
migration <- read.csv(file.choose())</pre>
centrs <- data.frame(as.character(state@data$NAME),coordinates(state))</pre>
colnames(centrs) <- c("name","long","lat")</pre>
migration <- migration[c(1,6:56)]
long_mig <- melt(migration,id.vars="from_state")</pre>
map("state")
define draw from state function
draw_from_state <- function(centrs, migrations, state_name, color=rgb(0,0,0,alpha=0.5)) {</pre>
    migrations$variable <- sub("."," ",migrations$variable,fixed=TRUE)</pre>
    migrations <- migrations[migrations$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),lw
}
}
draw_from_state(centrs, long_mig, "Florida", rgb(0,0,1,0.5))

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,.5))</pre>
```