

2025 554 R Notes on Spatial Data Analysis

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2024-12-19

R for Spatial Analysis

R has extensive spatial capabilities, the Spatial task view is [here](#)

Some of the notes that follow are build on Roger Bivand's notes taken from the latter site, and these are based on Bivand et al. (2013), which is a good reference book of the spatial capabilities of R at that point.

See [Spatial Data Science: With Applications in R](#) by Pebesma and Bivand, for a more up to date version.

Another resource is the book [Geocomputation with R](#)

To get R code alone from this .Rmd file, load the `knitr` library and then type

```
purl("2023-554-Spatial-Classes.Rmd")
```

from the directory with this file in.

Overview of Spatial Classes

Class definitions are objects that contain the formal definition of a class of R objects, and are usually referred to as an S4 class.

Spatial classes were defined to represent and handle spatial data, so that data can be exchanged between different classes - they are different from regular classes since they need to contain information about spatial locations and their coordinate reference systems

- The `sp` library was traditionally the workhorse for representing spatial data.
- Now `sf` has been replaced, details are [here](#)

These notes will focus on `sf`.

The `sf` Representation

The `sf` is the successor package to R packages formerly used in spatial data including `sp`, `rgeos` and the vector parts of `rgdal`, providing an interface to certain tidyverse packages.

The `sf` package reads and writes data through GDAL, and uses GEOS, s2geometry, and PROJ.

The `sf` package has `sf` objects, a sub-class of a `data.frame` or a `tibble`. These objects contain at least one geometry list-column of class `sfc`, where each list element contains the geometry as an R object of class `sfg`.

All functions are prefixed with `st_` to indicate “spatial type” to make them more easily searchable on the command line.

You can load the package by running `library(sf)`, but note that if you've never used this package before, you should run `install.packages('sf')` in your R console (and then load the package with the `library(sf)` command).

Creating a Spatial Object

We can use the `st_read` function (part of the `sf` package) to read in a spatial file.

```
file <- system.file("gpkg/nc.gpkg", package = "sf")
nc <- st_read(file)
## Reading layer `nc.gpkg' from data source
##   `/Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/sf/gpkg/nc.gpkg'
##   using driver `GPKG'
## Simple feature collection with 100 features and 14 fields
## Geometry type: MULTIPOLYGON
## Dimension:      XY
## Bounding box:   xmin: -84.32385 ymin: 33.88199 xmax: -75.45698 ymax: 36.58965
## Geodetic CRS:   NAD27
```

Note that the `st_read()` function takes in two arguments, the `dsn`, or data source name, and the layer. Our provided data has only one layer, so that argument is omitted in our example. You can check the available layers by querying:

```
st_layers(file)
## Driver: GPKG
## Available layers:
##   layer_name geometry_type features fields crs_name
## 1   nc.gpkg Multi Polygon      100      14   NAD27
```

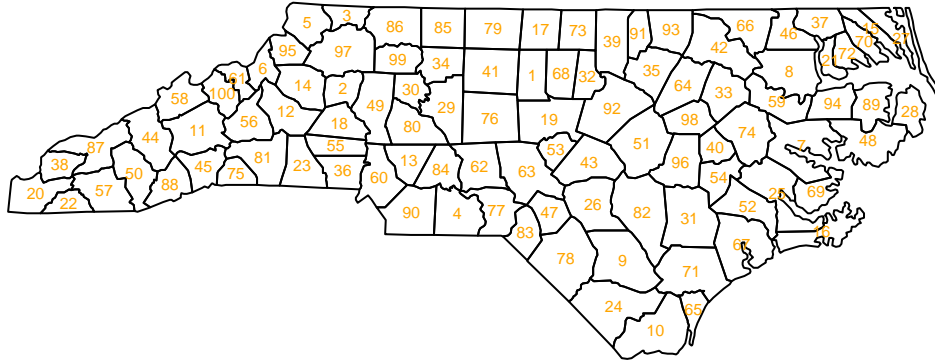
Let's look at the first few lines, define the centroids, and convert `nc` into an `sf` object. We need the centroids so that we can extract coordinates.

```
library(maps)
map("county", "North Carolina", fill = TRUE, plot = FALSE) |>
  st_as_sf() |>
  st_make_valid() -> nc1

head(nc1,n=3)
## Simple feature collection with 3 features and 1 field
## Geometry type: MULTIPOLYGON
## Dimension:      XY
## Bounding box:   xmin: -81.37147 ymin: 35.78121 xmax: -79.23434 ymax: 36.57763
## Geodetic CRS:   +proj=longlat +ellps=clrk66 +no_defs +type=crs
##                                     ID
## north carolina,alamance   north carolina,alamance
## north carolina,alexander  north carolina,alexander
## north carolina,alleghany  north carolina,alleghany
##                                     geom
## north carolina,alamance MULTIPOLYGON (((-79.23434 3...
## north carolina,alexander MULTIPOLYGON (((-81.15374 3...
## north carolina,alleghany MULTIPOLYGON (((-81.37147 3...
centroids = st_centroid(st_geometry(nc1)) |> st_coordinates()
```

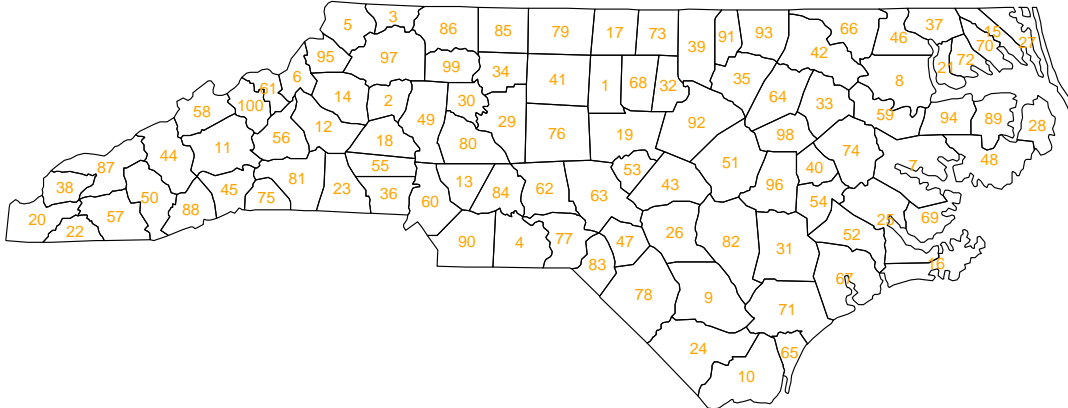
Let's plot the county boundaries and add a county label. Notice that the labeling here is according to alphabetical order, as seen in the `head(nc1)` call in the previous chunk.

```
plot(st_geometry(nc1))
text(centroids, labels = 1:nrow(nc1), col = "orange", cex=.5)
```



Another way of labeling, but in ggplot:

```
ggplot() +
  geom_sf(data = nc1, col = "black", fill = NA) +
  geom_text(aes(x = centroids[, 1], y = centroids[, 2],
               label = 1:100), size = 2, color = "orange") +
  theme_bw() +
  coord_sf() +
  xlab("") +
  ylab("") +
  theme(legend.title = element_blank(),
        panel.grid = element_blank(),
        panel.border = element_blank(),
        axis.ticks = element_blank(),
        axis.text = element_blank())
```



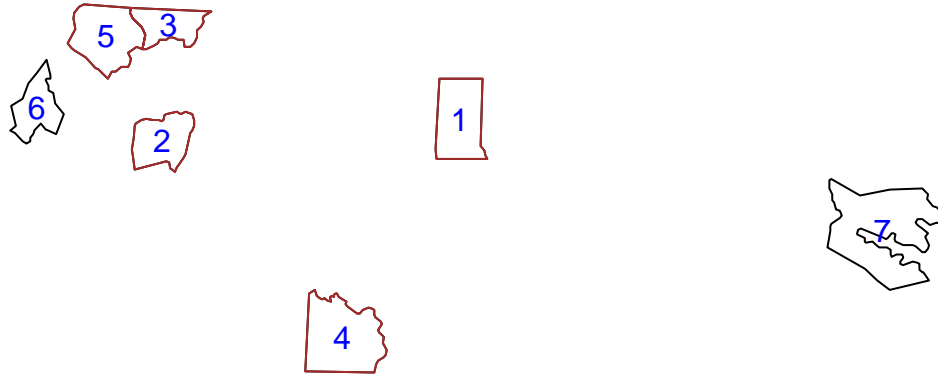
It is also possible to subset this data and plot it. You can run each line sequentially in R to see what each line of code adds to the plot. Note that this is written in base R, not with `ggplot()` where each line of code builds upon the plot.

We plot 7 areas only, and then highlight the first 5.

```
# Subset the data
nc5 <- nc1[1:5, ]
nc7 <- nc1[1:7, ]

# Plot the data
plot(st_geometry(nc7))
plot(st_geometry(nc5), add = TRUE, border = "brown")
cc <- st_coordinates(st_centroid(st_geometry(nc7)))
```

```
text(cc, labels = 1:nrow(nc7), col = "blue")
```

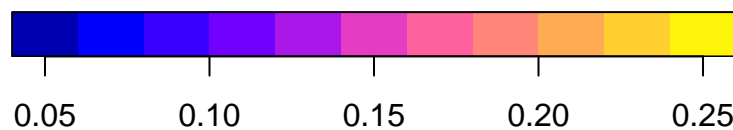
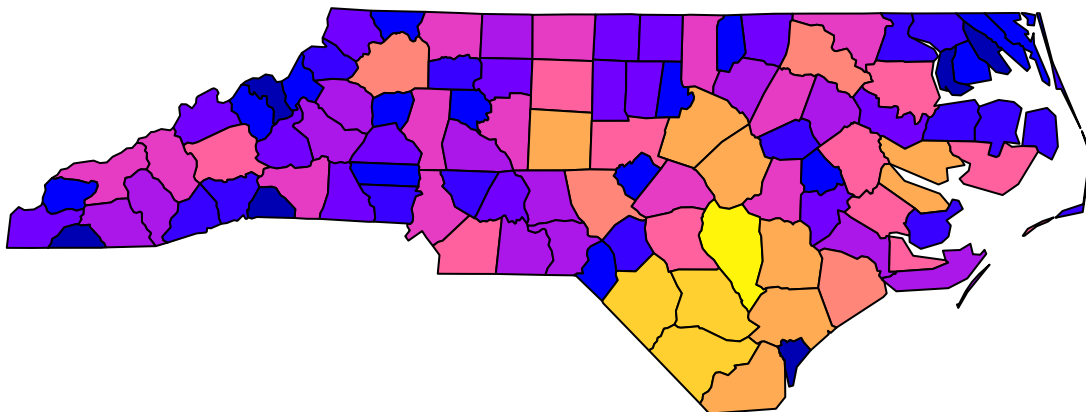


Visualizing Spatial Data

As you can see below, we can create a plot of the area sizes in each county in North Carolina (this is the first feature in the data frame). The area part comes into play when we index the `nc` object with a 1. If you change this to a 2, you'll see that this now plots the perimeter of each county instead, as the perimeters are the second feature.

```
par(mar = c(0,0,1,0))
plot(nc[1], reset = FALSE) # reset = FALSE: we want to add to a plot with a legend
```

AREA

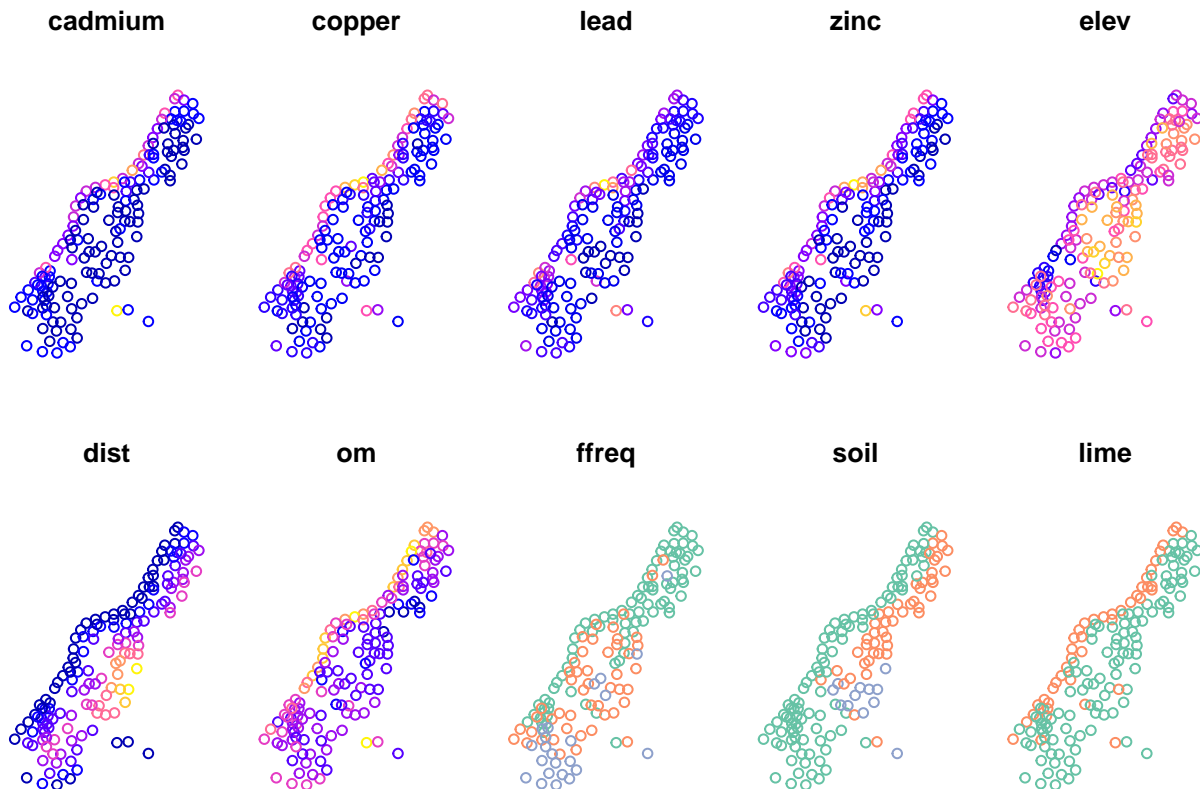


```
# plot(nc[1,1], col = 'grey', add = TRUE) This would grey out the first county
```

We can also look at the `meuse` dataset from the `sp` package. When converting this `sp` object into an `sf` object

using `st_as_sf`, notice that we must specify the coordinate columns.

```
data(meuse, package = "sp")
m.sf <- st_as_sf(meuse, coords = c("x","y"))
head(m.sf,n=3)
## Simple feature collection with 3 features and 12 fields
## Geometry type: POINT
## Dimension: XY
## Bounding box: xmin: 181025 ymin: 333537 xmax: 181165 ymax: 333611
## CRS: NA
##   cadmium copper lead zinc elev      dist   om ffreq soil lime landuse dist.m
## 1    11.7     85  299 1022 7.909 0.00135803 13.6     1    1    1    Ah     50
## 2     8.6     81  277 1141 6.983 0.01222430 14.0     1    1    1    Ah     30
## 3     6.5     68  199  640 7.800 0.10302900 13.0     1    1    1    Ah    150
##
##          geometry
## 1 POINT (181072 333611)
## 2 POINT (181025 333558)
## 3 POINT (181165 333537)
opar = par(mar=rep(0,4))
plot(m.sf)
```



Reading Shapefiles

ESRI (a company one of whose products is ArcGIS) shapefiles consist of three files, and this is a common form.

- The first file (`*.shp`) contains the geography of each shape.
- The second file (`*.shx`) is an index file which contains record offsets.
- The third file (`*.dbf`) contains feature attributes with one record per feature.

The Washington state Geospatial Data Archive (wagda) can be accessed [here](#) and contains data that we can read in.

As an example, consider Washington county data that was downloaded from wagda.

The data consists of the three files: `wacounty.shp`, `wacounty.shx`, `wacounty.dbf`.

The following code reads in these data and then draws a county level map of 1990 populations, and a map with centroids.

First load the libraries.

```
library(maps) # for background map outlines
```

Note that there are problems with the files, which are sorted by using the `repair=T` argument.

The data can be downloaded from [here](#), but we read in directly:

```
download.file("http://faculty.washington.edu/jonno/SISMIDmaterial/wacounty.shp",destfile = "wacounty.shp")
download.file("http://faculty.washington.edu/jonno/SISMIDmaterial/wacounty.shx",destfile = "wacounty.shx")
download.file("http://faculty.washington.edu/jonno/SISMIDmaterial/wacounty.dbf",destfile = "wacounty.dbf")
wacounty = st_read(dsn=".",layer = "wacounty")
## Reading layer `wacounty' from data source
##   `/Users/jonno/Dropbox/554+555/554-25-R-Lectures' using driver `ESRI Shapefile'
## Simple feature collection with 39 features and 6 fields (with 1 geometry empty)
## Geometry type: MULTIPOLYGON
## Dimension:      XY
## Bounding box:   xmin: -124.7312 ymin: 45.5434 xmax: -116.915 ymax: 49.0026
## CRS:            NA
class(wacounty)
## [1] "sf"           "data.frame"
head(wacounty,n=3)
## Simple feature collection with 3 features and 6 fields
## Geometry type: MULTIPOLYGON
## Dimension:      XY
## Bounding box:   xmin: -119.8756 ymin: 45.8358 xmax: -116.915 ymax: 47.2616
## CRS:            NA
##           AreaName AreaKey INTPTLAT INTPTLNG TotPop90 CNTY
## 1 WA, Adams County   53001 46.98899 -118.5569   13603    1
## 2 WA, Asotin County  53003 46.18248 -117.1850   17605    3
## 3 WA, Benton County  53005 46.24764 -119.5015  112560    5
##           geometry
## 1 MULTIPOLYGON (((-118.9784 4...
## 2 MULTIPOLYGON (((-117.2276 4...
## 3 MULTIPOLYGON (((-119.8728 4...
```

Let's see what these variables look like: we see county names and FIPS codes. When linking datasets, note that it's a good idea to use one of these variables, as they are generally reliable.

```
names(wacounty)
## [1] "AreaName" "AreaKey"  "INTPTLAT" "INTPTLNG" "TotPop90" "CNTY"      "geometry"
wacounty$AreaName[1:3]
## [1] "WA, Adams County" "WA, Asotin County" "WA, Benton County"
wacounty$AreaKey[1:3]
## [1] "53001" "53003" "53005"
# head(wacounty)
```

Drawing a map

We look at some variables.

```
wacounty$INTPTLAT[1:3] # latitude
## [1] 46.98899 46.18248 46.24764
wacounty$INTPTLNG[1:3] # longitude
## [1] -118.5569 -117.1850 -119.5015
wacounty$CNTY[1:3]
## [1] "1" "3" "5"
wacounty$TotPop90[1:3]
## [1] 13603 17605 112560
```

We look at some variables, and then set up the colors to map. We map 1990 Washington population counts by census tracts.

First, define a utility function `leglabs` which will help us create labels for our legends.

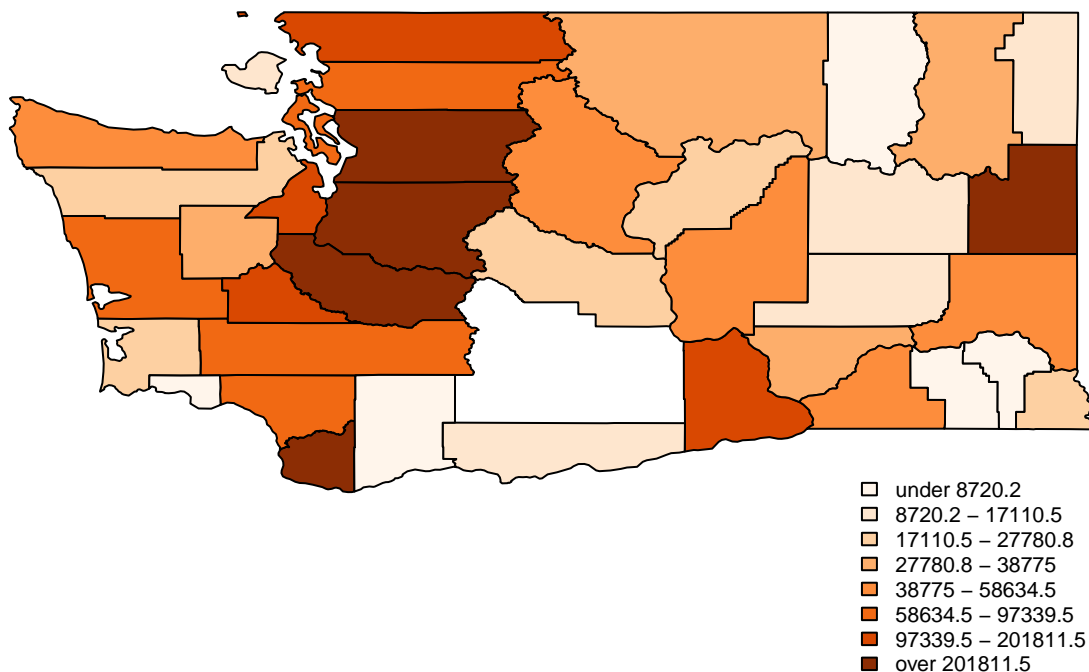
```
# utility function to make legend; copied from maptools package
leglabs <- function(vec, under = "under", over = "over", between = "-") {
  x <- vec
  lx <- length(x)
  if (lx < 3)
    stop("vector too short")
  res <- character(lx - 1)
  res[1] <- paste(under, x[2])
  for (i in 2:(lx - 2)) res[i] <- paste(x[i], between, x[i + 1])
  res[lx - 1] <- paste(over, x[lx - 1])
  res
}
```

Method 1

The first option for drawing maps is to use base R as follows:

```
# setup
plotvar <- wacounty$TotPop90 # variable we want to map
summary(plotvar)
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##      2248  17110   38775  124787   97340 1507319
nclr <- 8 # next few lines set up the color scheme for plotting
plotclr <- brewer.pal(nclr, "Oranges")
brks <- round(quantile(plotvar, probs=seq(0,1,1/(nclr))), digits=1)
colnum <- findInterval(plotvar, brks, all.inside=T)
colcode <- plotclr[colnum]

# plot
plot(wacounty["TotPop90"], col=colcode, main="")
legend("bottomright", legend=leglabs(round(brks, digits=1)), fill=plotclr, cex=0.65, bty="n", inset = c(0, -0
```

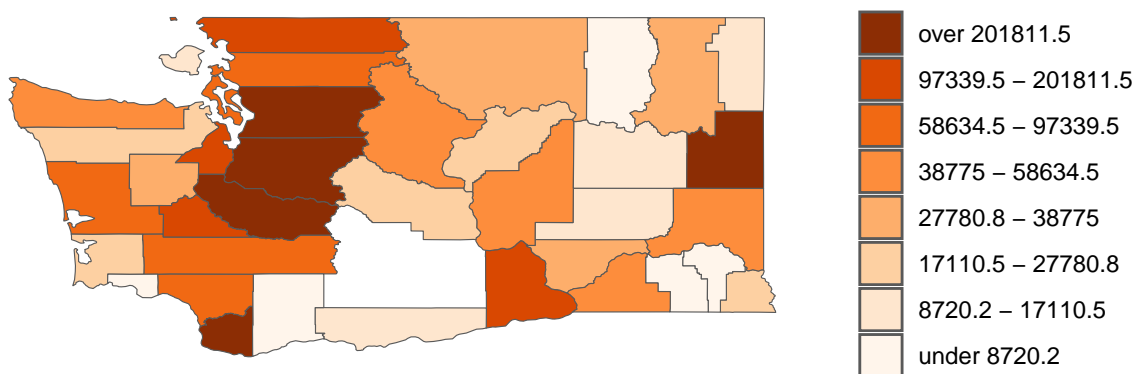


Method 2

This is effectively the same as method 1, but instead of using the base R `plot()` function, we use `ggplot()` here.

```

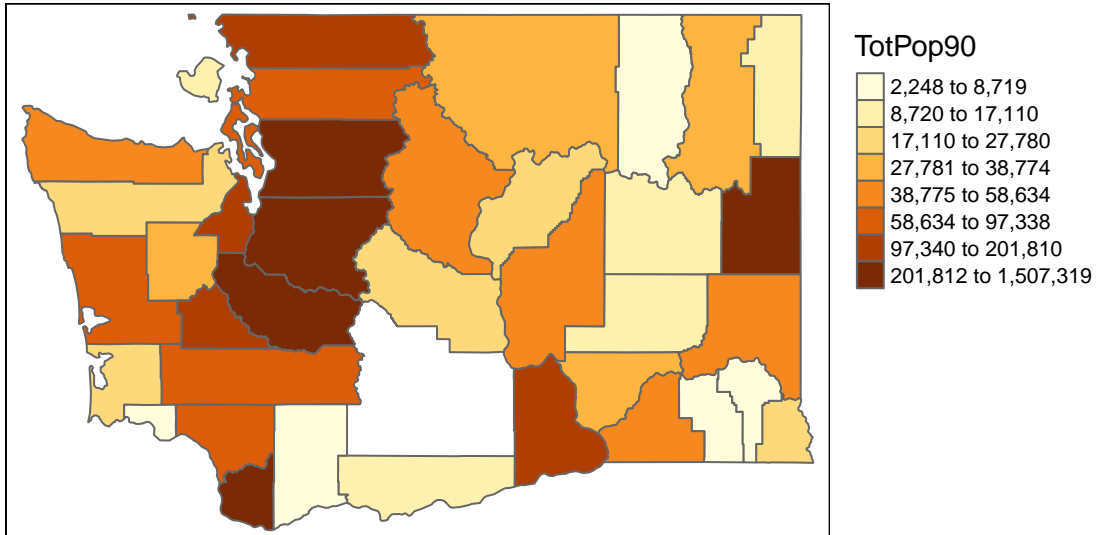
ggplot(data = wacounty) +
  geom_sf(aes(fill = colcode)) +
  scale_fill_manual(values = rev(plotclr), labels = rev(leglabs(round(brks,digits=1)))) +
  theme_bw() +
  coord_sf() +
  xlab("") +
  ylab("") +
  theme(legend.title = element_blank(),
        panel.grid = element_blank(),
        panel.border = element_blank(),
        axis.ticks = element_blank(),
        axis.text = element_blank())
  
```



Method 3

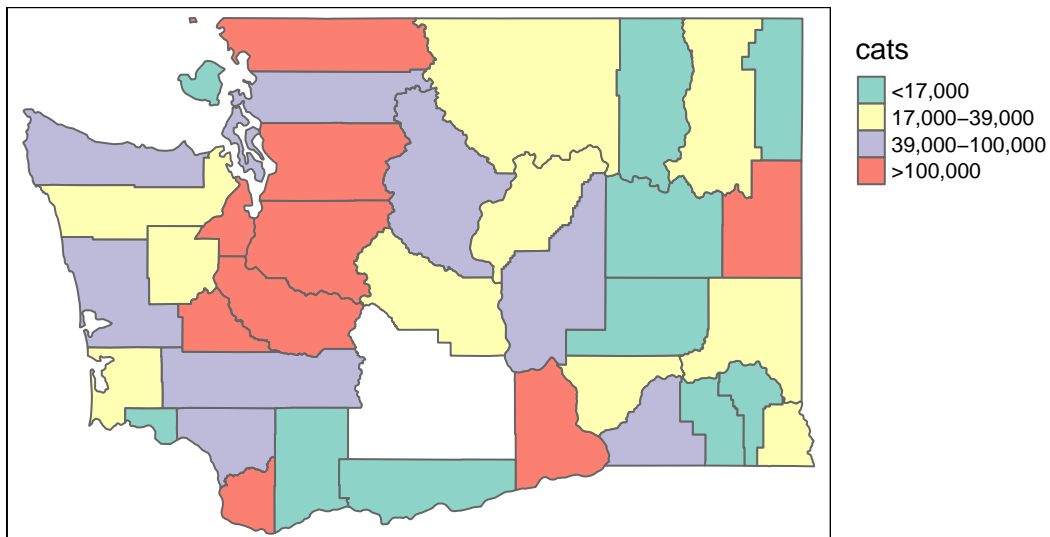
We can also use `tmap`:


```
library(tmap)
sel = !st_is_empty(wacounty)
tm_shape(st_set_crs(wacounty, 4326)[sel,]) +
  tm_polygons("TotPop90", breaks = brks) +
  tm_layout(legend.outside = TRUE)
```



For illustration, we define our own cutpoints by hand.

```
summary(wacounty$TotPop90)
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##      2248  17110   38775 124787  97340 1507319
wacounty$cats <- cut(wacounty$TotPop90,breaks=c(0,17000,39000,100000,1520000),
  labels=c("<17,000","17,000-39,000","39,000-100,000",>100,000"))
tm_shape(st_set_crs(wacounty, 4326)[sel,]) +
  tm_polygons("cats") +
  tm_layout(legend.outside = TRUE)
```



We now highlight a couple of counties: King County and Spokane County.

```

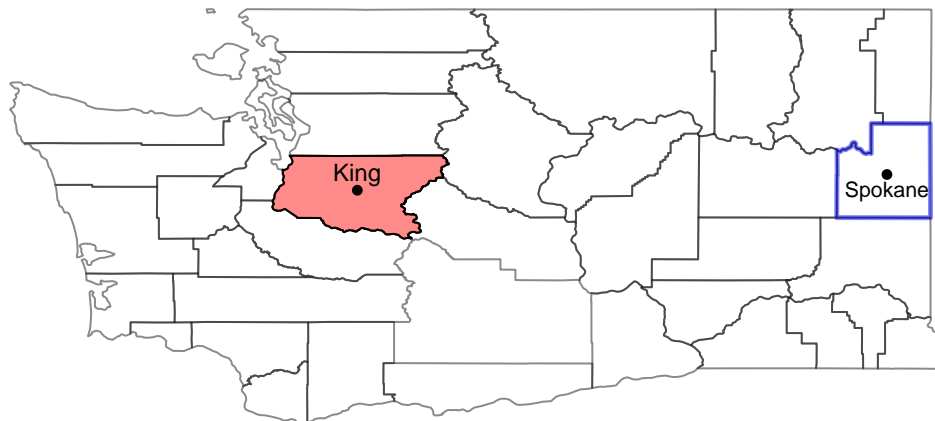
# identify IDs for counties of interest
xx <- which(wacounty$AreaName == "WA, King County")
xx2 <- which(wacounty$AreaName == "WA, Spokane County")

# plot the whole state
plot(st_geometry(wacounty), border="#00000075")

# highlight counties of interest
plot(st_geometry(wacounty[xx, ]), col="#ff000075", add=T)
plot(st_geometry(wacounty[xx2, ]), col=NA, border="#0000ff75", add=T, lwd=2.5)

# Add some labels
text(st_coordinates(st_centroid(st_geometry(wacounty[xx, ]))),
     "King", cex=0.75, pos=3, offset=0.25)
text(st_coordinates(st_centroid(st_geometry(wacounty[xx2, ]))),
     "Spokane", cex=0.7, pos=1, offset=0.25)
points(st_coordinates(st_centroid(st_geometry(wacounty[c(xx, xx2), ]))),
       pch=16, cex=0.75)

```



Now let's examine for census tracts.

```

## Reading Shapefiles
download.file("http://faculty.washington.edu/jonno/SISMIDmaterial/watract.shp", destfile = "watract.shp")
download.file("http://faculty.washington.edu/jonno/SISMIDmaterial/watract.shx", destfile = "watract.shx")
download.file("http://faculty.washington.edu/jonno/SISMIDmaterial/watract.dbf", destfile = "watract.dbf")
watract=st_read(dsn=".", layer = "watract")
## Reading layer `watract' from data source
##   `/Users/jonno/Dropbox/554+555/554-25-R-Lectures' using driver `ESRI Shapefile'
## Simple feature collection with 1152 features and 7 fields (with 1 geometry empty)
## Geometry type: MULTIPOLYGON
## Dimension:      XY
## Bounding box:   xmin: -124.8485 ymin: 45.55 xmax: -116.9174 ymax: 49.0025
## CRS:            NA

```

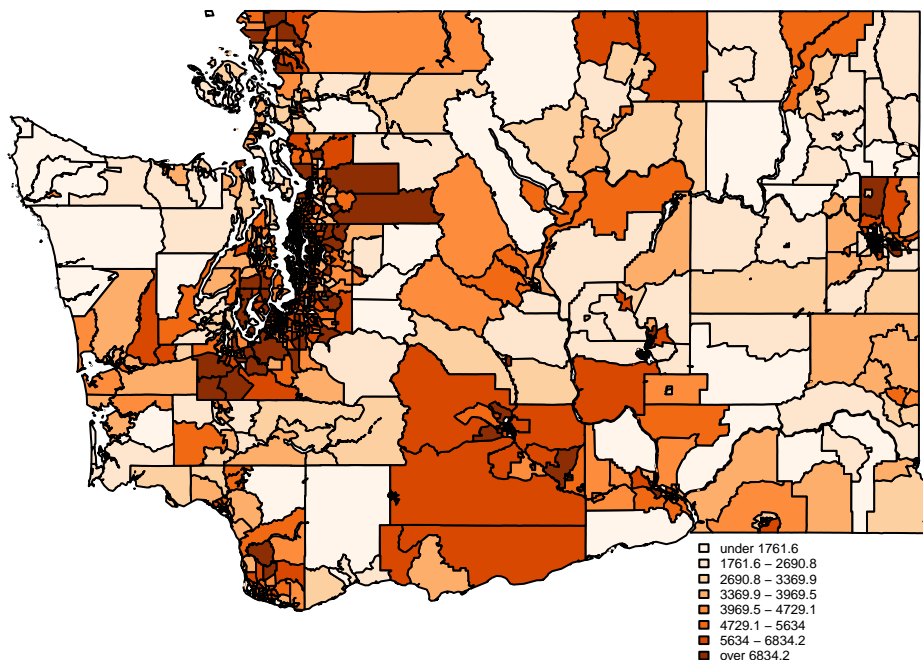
Drawing a census tract map: we repeat but now map populations at the census tract level.

```

watract <- st_read("watract.shp") |> st_make_valid() |> st_set_crs(4326)
## Reading layer `watract' from data source
##   `/Users/jonno/Dropbox/554+555/554-25-R-Lectures/watract.shp'
##   using driver `ESRI Shapefile'
## Simple feature collection with 1152 features and 7 fields (with 1 geometry empty)

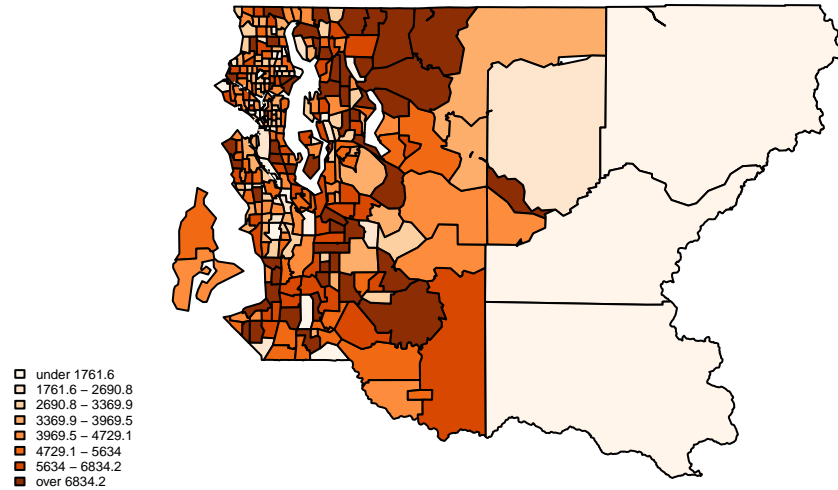
```

```
## Geometry type: MULTIPOLYGON
## Dimension:      XY
## Bounding box:   xmin: -124.8485 ymin: 45.55 xmax: -116.9174 ymax: 49.0025
## CRS:            NA
names(watract)
## [1] "AreaName" "AreaKey"  "INTPTLAT" "INTPTLNG" "TotPop90" "TRACT"    "CNTY"
## [8] "geometry"
plotvar <- watract$TotPop90 # variable we want to map
brks <- round(quantile(plotvar, probs=seq(0,1,1/(nclr))), digits=1)
colnum <- findInterval(plotvar, brks, all.inside=T)
colcode <- plotclr[colnum]
plot(st_geometry(watract), col=colcode)
legend(-119, 46, legend=leglabs(round(brks, digits=1)),
      fill=plotclr, cex=0.4, bty="n")
```



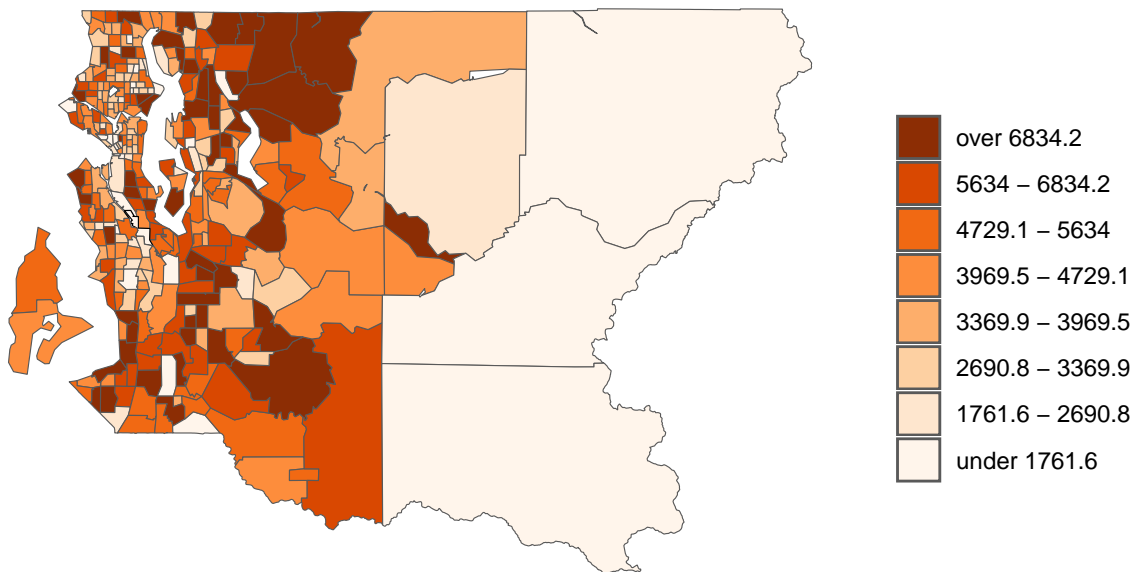
Notice that we can also choose to zoom in on King county. Here is the Base R version:

```
xx <- which(watract$CNTY == 33)
plot(st_geometry(watract[xx, ]), col = colcode[xx])
legend("bottomleft",
      legend = leglabs(round(brks, digits = 1)),
      fill = plotclr, cex = 0.4, bty = "n"
)
```



Here is the `ggplot()` version:

```
ggplot(data = wattract[xx, ]) +
  geom_sf(aes(fill = colcode[xx])) +
  scale_fill_manual(values = rev(plotclr), labels = rev(leglabs(round(brks, digits = 1)))) +
  theme_bw() +
  coord_sf() +
  xlab("") +
  ylab("") +
  theme(
    legend.title = element_blank(),
    panel.grid = element_blank(),
    panel.border = element_blank(),
    axis.ticks = element_blank(),
    axis.text = element_blank()
  )
```



John Snow Example

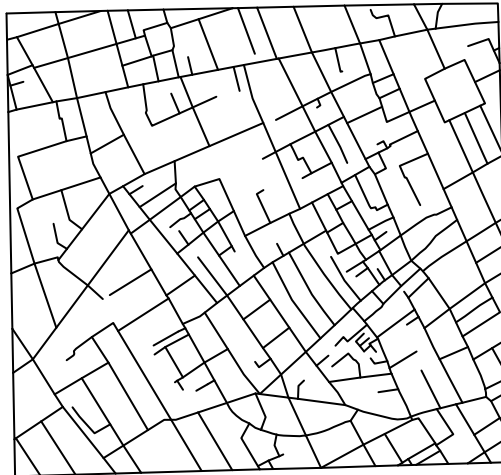
For fun, let's look at the poster child of health mapping.

The Snow data consists of the relevant 1854 London streets, the location of 578 deaths from cholera, and the position of 13 water pumps (wells) that can be used to re-create John Snow's map showing deaths from cholera in the area surrounding Broad Street, London in the 1854 outbreak.

```
library(HistData)
data(Snow.deaths)
data(Snow.pumps)
data(Snow.streets)
# data(Snow.polygons)
```

We first create an `sfc` object containing the coordinates of the streets using the `st_linestring()` function; `st_sfc()` combines all the individual `LINESTRING` pieces:

```
# Streets
slist <- split(Snow.streets[,c("x", "y")],
              as.factor(Snow.streets[, "street"]))
m <- lapply(slist, as.matrix) |> lapply(st_linestring) |> st_sfc()
plot(m)
```



Display the streets and then add the deaths and pumps (with labels). The red squares are deaths, blue triangles are pumps

```
plot(m, col="gray")
# deaths:
p <- st_as_sf(Snow.deaths, coords = c("x", "y"))
plot(p, add=TRUE, col='red', pch=15, cex=0.6)
# pumps:
spp <- st_as_sf(Snow.pumps, coords = c("x", "y"))
plot(spp, add=TRUE, col='blue', pch=17, cex=1.0)
text(Snow.pumps[,c("x", "y")], labels=Snow.pumps$label,
     pos=1, cex=0.8)
```

