# 2023 554 R Notes on Spatial Data Analysis

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## 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 developments until that point.

To get R code alone then 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 is being increasingly used, details are here

These notes will focus on sf.

See Spatial Data Science with Applications in R by Pebesma and Biyand, for a more up to date version.

Another resource is the book Geocomputation with R

## The sf Representation

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

This 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 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/4.2-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</pre>
```

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:

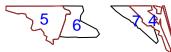
```
st_layers(file)
## Driver: GPKG
## Available layers:
## layer_name geometry_type features fields crs_name
## 1 nc.gpkg Multi Polygon 100 14 NAD27
```

It's 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.

```
# Subset the data
nc5 <- nc[1:5, ]
nc7 <- nc[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")</pre>
```



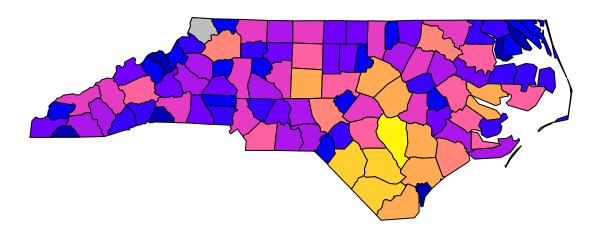


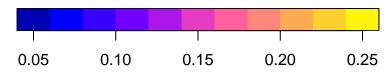
#### Visualizing Spatial Data

As you can see below, we can create a plot of the sizes in area of each county in North Carolina. 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 perimeters is 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
plot(nc[1,1], col = 'grey', add = TRUE)
```

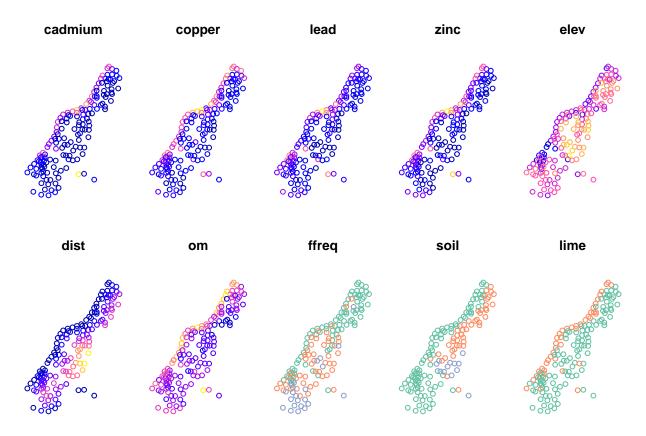
## **AREA**





We can also look at the meuse dataset from the sp package and convert the data into a sp object, namely a SpatialPointsDataFrame by using the coordinates() function. Then, using st\_as\_sf(), we can go from an sp object to an sf object.

```
library(sp)
data(meuse)
coordinates(meuse) = ~x+y
m.sf = st_as_sf(meuse)
opar = par(mar=rep(0,4))
plot(m.sf)
```



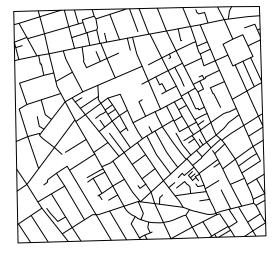
## John Snow Example

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

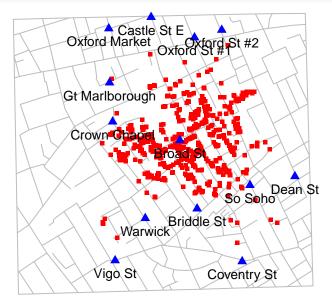
The Snow data consists of the relevant 1854 London streets, the location of 578 deaths from cholera, and the position of 13 water pumps (wells) that can be used to re-create John Snow's map showing deaths from cholera in the area surrounding Broad Street, London in the 1854 outbreak.

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

We first create a st\_multilinestring object containing the coordinates of the streets:



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



## 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 can be accessed here contains data that can be read into R.

As an example, consider ashington 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)
library(shapefiles)
```

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

The data can be downloaded from here: here

```
download.file("http://faculty.washington.edu/jonno/SISMIDmaterial/wacounty.shp",destfile =
download.file("http://faculty.washington.edu/jonno/SISMIDmaterial/wacounty.shx",destfile =
download.file("http://faculty.washington.edu/jonno/SISMIDmaterial/wacounty.dbf",destfile =
wacounty.sh
download.file("http://faculty.washington.edu/jonno/SISMIDmaterial/wacounty.dbf",destfile =
"wacounty.sh
wacounty = st_read(dsn=".",layer = "wacounty")

## Reading layer `wacounty' from data source `/Users/kpaulson/Repos/STAB' 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"
```

Let's see what these variables look like: we see county names and FIPS codes.

```
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)
```

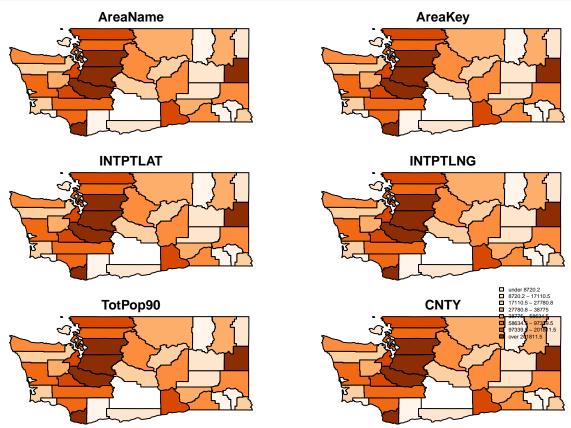
## Drawing a map

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
```

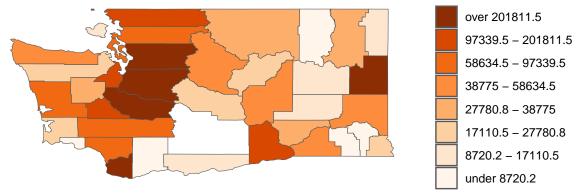
We look at some variables, and then set up the colors to map. We map 1990 Washington population counts by census tracts.

```
## Function to make the legend
## Adapted from maptools::leglabs()
make_legend <- function (vec, under = "under", over = "over", between = "-")
  x <- vec
  lx <- length(x)</pre>
  res <- character(lx - 1)
  res[1] \leftarrow paste(under, x[2])
  for (i in 2:(lx - 2)) res[i] <- paste(x[i], between, x[i +</pre>
  res[lx - 1] \leftarrow paste(over, x[lx - 1])
}
plotvar <- wacounty$TotPop90 # variable we want to map</pre>
summary(plotvar)
##
      Min. 1st Qu. Median
                                Mean 3rd Qu.
      2248 17110 38775 124787
                                      97340 1507319
##
nclr <- 8 # next few lines set up the color scheme for plotting
plotclr <- brewer.pal(nclr, "Oranges")</pre>
brks <- round(quantile(plotvar,probs=seq(0,1,1/(nclr))),digits=1)</pre>
colornum <- findInterval(plotvar,brks,all.inside=T)</pre>
colcode <- plotclr[colornum]</pre>
plot(wacounty,col=colcode)
legend("bottomright", legend=make_legend(round(brks, digits = 1)),
       fill=plotclr,cex=0.35,bty="n")
```

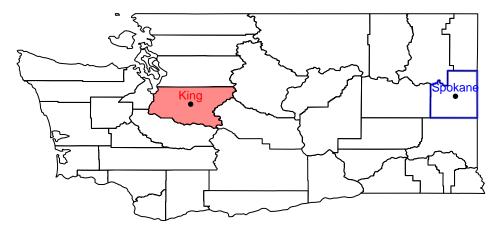


## In ggplot:

```
ggplot(data = wacounty) +
  geom_sf(aes(fill = colcode)) +
  scale_fill_manual(values = rev(plotclr), labels = rev(make_legend(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())
```



We now highlight a county



Now let's repeat 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/kpaulson/Repos/STAB' using driver `ESRI Shapefile'

## Simple feature collection with 1152 features and 7 fields (with 1 geometry empty)

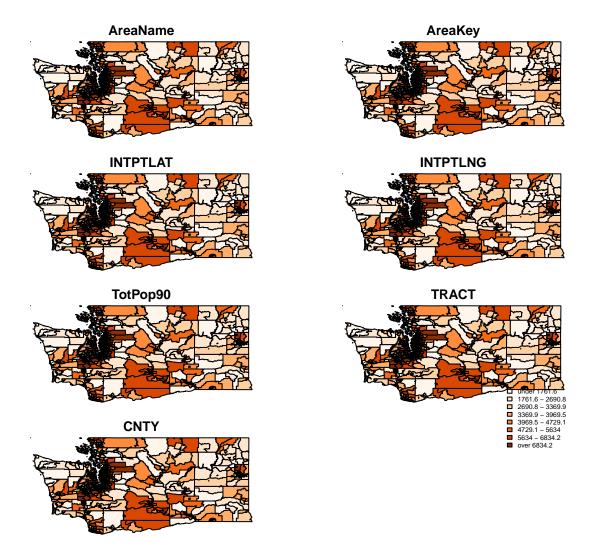
## Geometry type: MULTIPOLYGON

## Dimension: XY

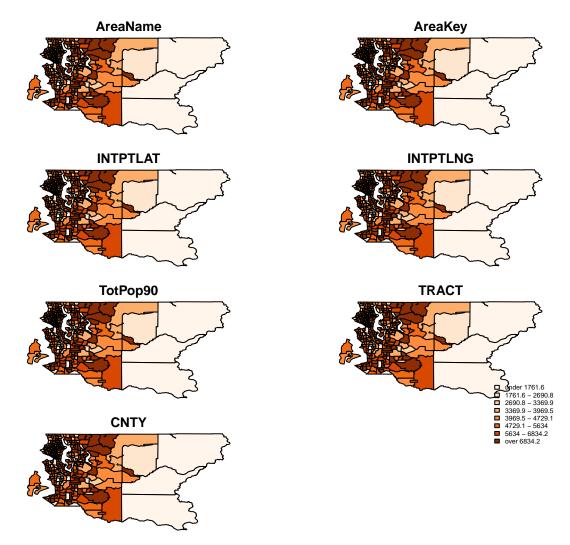
## Bounding box: xmin: -124.8485 ymin: 45.55 xmax: -116.9174 ymax: 49.0025

## CRS: NA
```

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

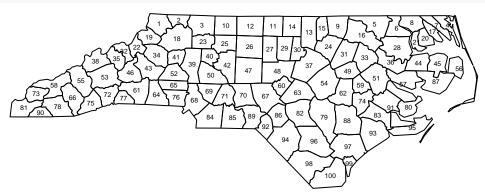


We zoom in on King County



A county map of North Carolina with text

```
nc <- read_sf(system.file("gpkg/nc.gpkg", package = "sf"))
plot(nc["BIR74"]$geom)
text(sf::st_coordinates(st_centroid(nc["BIR74"]$geom)), rownames(nc), cex=0.4, offset=0)</pre>
```



```
In ggplot:
```

```
centroids <- sf::st_coordinates(st_centroid(nc["BIR74"]$geom))
ggplot() +</pre>
```

