



Tutorial: Geocomputation with R



Geographic vector data in R

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Find the slides and the code

https://github.com/jannes-m/erum18_geocompr



Simple features in R



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Simple feature access is a widely used **ISO standard**.



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```
library(sf)
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```

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data(random_points, package = "RQGIS")  
class(random_points)
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## [1] "sf"          "data.frame"
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```
## [1] "sf"          "data.frame"
```

This is a **data.frame**, i.e, an S3 object (as opposed to **SpatialObjects**).



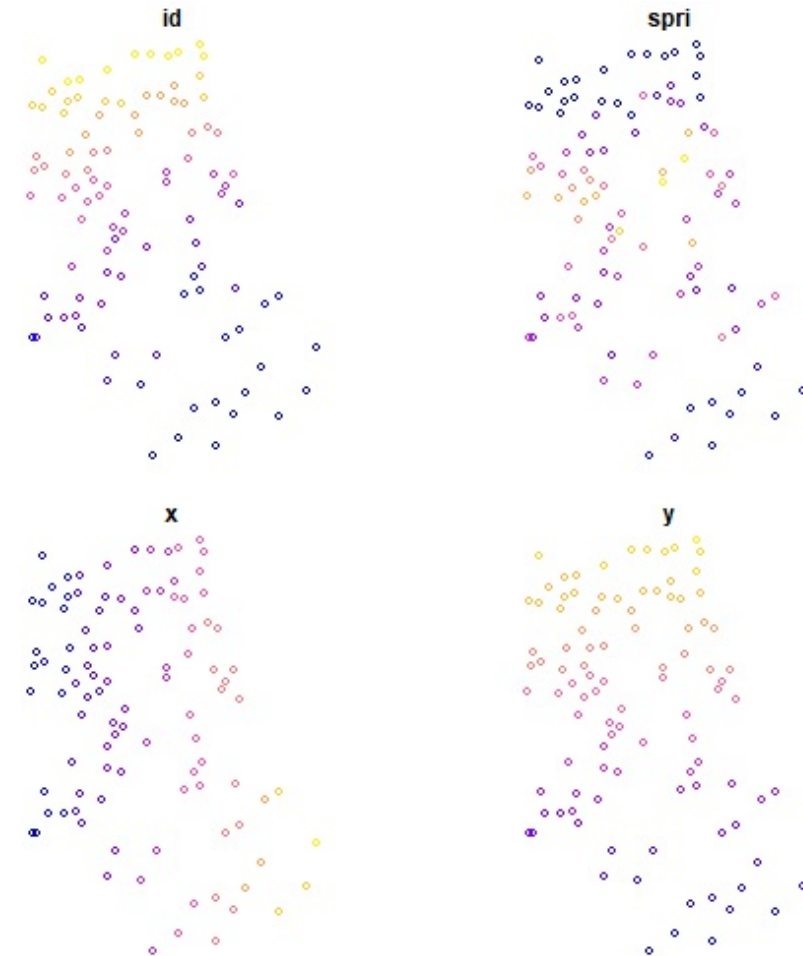
Simple features in R

```
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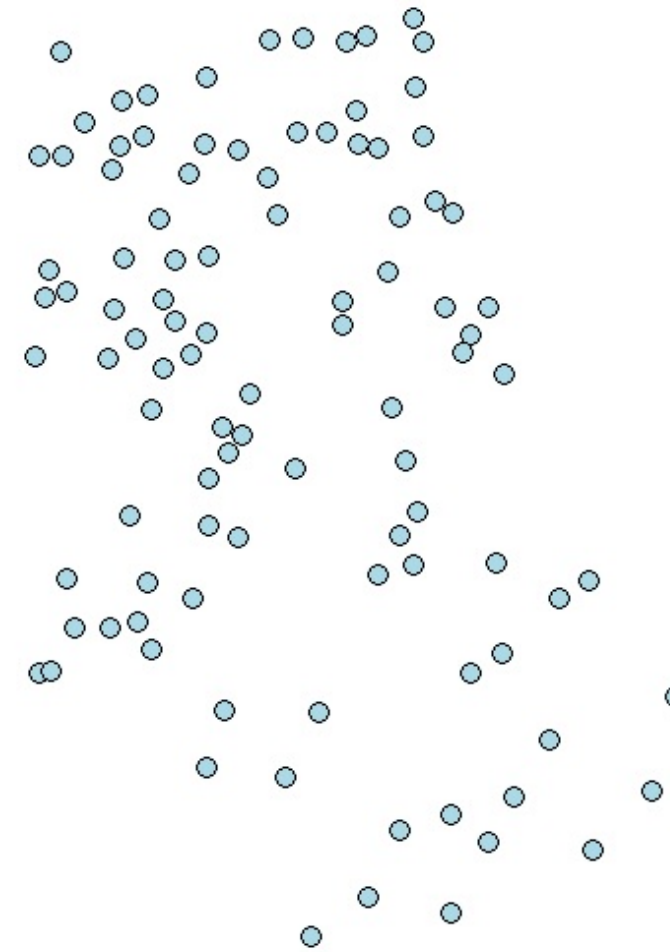
Simple features in R

```
plot(  
  st_geometry(random_points),  
  pch = 16, cex = 2,  
  col = "black"  
  bg = "lightblue"  
)
```



Simple features in R

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```





Simple features in R

```
library(dplyr)
select(random_points, 1:2) %>%
  head(2)
```

```
## Simple feature collection with 2 features and 2 fields
## geometry type: POINT
## dimension: XY
## bbox: xmin: 796749.3 ymin: 8932621 xmax: 797178.6 ymax: 8932755
## epsg (SRID): 32717
## proj4string: +proj=utm +zone=17 +south +datum=WGS84 +units=m +no_defs
##   id spri geometry
## 1  1     4 POINT (797178.6 8932755)
## 2  2     4 POINT (796749.3 8932621)
```

A few things to note:

- **sf** works with the **tidyverse**.



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- Geometry is **just** another column.



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A few things to note:

- **sf** works with the **tidyverse**.
- Geometry is **just** another column.
- The geometry column is **sticky**.



Things to note continued:

- Each observation (row) has a geometry (which can consist of multiple features, think of polygons with holes or multi-part polygons).



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- The geometry column is a so-called **list-column**.
- The geometry is build up of **simple** R structures.



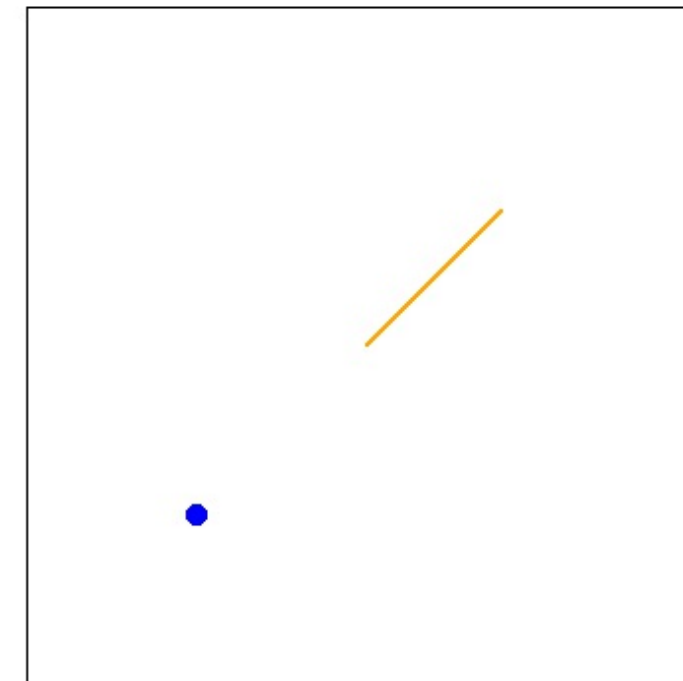
Geometries

```
# one point (a numeric vector)
p = st_point(c(1.25, 1.25))
# one line (a matrix consisting of at
# least two points)
mat = matrix(c(1.5, 1.5, 1.7, 1.7),
             ncol = 2, byrow = TRUE)
l = st_linestring(mat)
# one polygon
mat = matrix(c(1, 1, 1, 2, 2, 2,
              2, 1, 1, 1),
             ncol = 2, byrow = TRUE)
# a list of one or more matrices
# consisting of points
poly = st_polygon(list(mat))
# plot it
plot(poly)
plot(p, pch = 16, col = "blue",
      cex = 2, add = TRUE)
plot(l, cex = 2, col = "orange",
      lwd = 2, add = TRUE)
```



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Putting it all together

sf uses three classes to represent simple features in R:

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- **sf** is the **data.frame** with the attributes and the geometry list-column
- The geometry list column is of class **sfc**.

```
lc = random_points %>%  
  st_geometry  
class(lc)
```

```
## [1] "sfc_POINT" "sfc"
```



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- **sf** is the **data.frame** with the attributes and the geometry list-column
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```
lc = random_points %>%  
  st_geometry  
class(lc)
```

```
## [1] "sfc_POINT" "sfc"
```

- Each feature of the list column is of class **sfg**.

```
class(lc[[1]])
```

```
## [1] "XY"      "POINT" "sfg"
```



Putting it all together

sf uses three classes to represent simple features in R:

- **sf** is the **data.frame** with the attributes and the geometry list-column
- The geometry list column is of class **sfc**.

```
lc = random_points %>%  
  st_geometry  
class(lc)
```

```
## [1] "sfc_POINT" "sfc"
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- Each feature of the list column is of class **sfg**.

```
class(lc[[1]])
```

```
## [1] "XY"      "POINT" "sfg"
```

For more information, refer to `vignette("sf1", package = "sf")` and <https://geocompr.robinlovelace.net/spatial-class.html#vector-data>



Attribute operations



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- **sf** objects are basically dataframes and thus can be handled like any other R object.



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dim(random_points)
```

```
## [1] 100  5
```



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```
dim(random_points)
```

```
## [1] 100  5
```

```
str(random_points)
```

```
## Classes 'sf' and 'data.frame':  100 obs. of  5 variables:
## $ id      : int  1 2 3 4 5 6 7 8 9 10 ...
## $ spri     : int  4 4 3 2 4 5 6 2 3 3 ...
## $ x        : num  797179 796749 796816 797023 796647 ...
## $ y        : num  8932755 8932621 8932739 8932600 8932692 ...
## $ geometry:sfc_POINT of length 100; first list element: Classes 'XY', 'I
## - attr(*, "sf_column")= chr "geometry"
## - attr(*, "agr")= Factor w/ 3 levels "constant","aggregate",...: NA NA NA
## ..- attr(*, "names")= chr  "id" "spri" "x" "y"
```



Subsetting

```
# first 2 rows and first 2 columns  
random_points[1:2, 1:2]
```

```
## Simple feature collection with 2 features and 2 fields  
## geometry type: POINT  
## dimension: XY  
## bbox: xmin: 796749.3 ymin: 8932621 xmax: 797178.6 ymax: 8932755  
## epsg (SRID): 32717  
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##   id spri geometry  
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Tidyverse



- When **dplyr** is also attached to the global environment, a number of generic methods of the tidyverse become available for **sf**-objects, most notably the one-table verbs **select**, **slice**, **filter**, **arrange**, **mutate**, **summarize** (and **group_by**).



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- Piped operations are also supported (%>%).

```
select(random_points, 1:2) %>%  
  slice(1:2)
```

```
## Simple feature collection with 2 features and 2 fields  
## geometry type: POINT  
## dimension: XY  
## bbox: xmin: 795551.4 ymin: 8932370 xmax: 797242.3 ymax: 8934800  
## epsg (SRID): 32717  
## proj4string: +proj=utm +zone=17 +south +datum=WGS84 +units=m +no_defs  
## # A tibble: 2 x 3  
##       id spri geometry  
##   <int> <int> <POINT [m]>  
## 1     1     4 (797178.6 8932755)  
## 2     2     4 (796749.3 8932621)
```



Vector attribute operations

Further reading: <https://geocompr.robinlovelace.net/attr.html#vector-attribute-manipulation>



Your turn

- Select all observations of **random_points** (`data("random_points", package = "RQGIS")`) which have more than 10 species (column **spri**). Plot the geometry of all points and add your selection to the plot in another color.
- Based on **spri** add a categorical column to **random_points** with 0-5 corresponding to **low**, 5-10 to **medium** and >10 to **high**.
- Optional: create two points of class **sfg** and convert them into an object of class **sf** which has an **id** and a **geometry** column.



Spatial attribute operations



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- Spatial subsetting
- Topological or neighborhood operations
- Spatial joins (spatial overlay)



Spatial subsetting

```
# spData makes available  
# nz and nz_height  
library(spData)  
plot(st_geometry(nz))  
plot(st_geometry(nz_height),  
      pch = 16, col = "red2",  
      cex = 2, add = TRUE)
```



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```
# spData makes available  
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library(spData)  
plot(st_geometry(nz))  
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Spatial subsetting

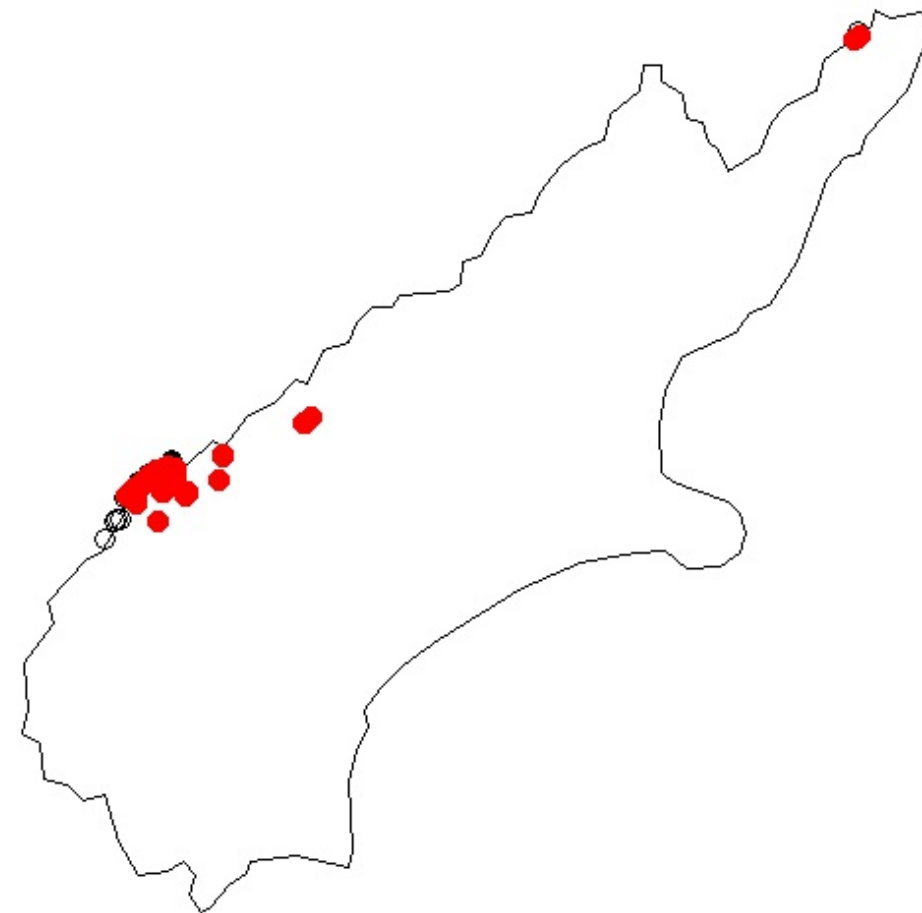
```
canterbury = nz %>%  
  filter(Name == "Canterbury")  
plot(st_geometry(canterbury))  
plot(st_geometry(nz_height),  
      cex = 2, add = TRUE)  
# spatial subsetting  
sel = nz_height[canterbury, ]  
plot(st_geometry(sel), cex = 2,  
      col = "red", pch = 16,  
      add = TRUE)
```





Spatial subsetting

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canterbury = nz %>%  
  filter(Name == "Canterbury")  
plot(st_geometry(canterbury))  
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      cex = 2, add = TRUE)  
# spatial subsetting  
sel = nz_height[canterbury, ]  
plot(st_geometry(sel), cex = 2,  
      col = "red", pch = 16,  
      add = TRUE)
```





Topological relations

Implicitly our subsetting used **st_intersects**, i.e. it returned all featured that touched or overlapped.

```
nz_height[canterbury, op = st_intersects]  
# see also  
?st_sf
```



Topological relations

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We can use **st_intersects** individually. This returns a boolean vector if there is an intersection.



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nz_height[canterbury, op = st_intersects]
# see also
?st_sf
```

We can use **st_intersects** individually. This returns a boolean vector if there is an intersection.

```
st_intersects(nz_height, canterbury, sparse = FALSE) %>% head
```

```
##      [,1]
## [1,] FALSE
## [2,] FALSE
## [3,] FALSE
## [4,] FALSE
## [5,]  TRUE
## [6,]  TRUE
```



aside from **st_intersects** there are further predicates:

- **st_disjoint**: the opposite of **st_intersects**
- **st_touches**: just touching
- ...
- have a look at **?st_intersects** for a complete list and description



Spatial join

Transfer the attribute of one spatial object to another spatial object based on intersecting geometries. For example, let us add the region name from **nz** to **nz_height** (so far consisting of columns **t50_fid**, **elevation** and **geometry**).



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```
join = st_join(nz_height, select(nz, Name))
```




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```
join = st_join(nz_height, select(nz, Name))
```

```
slice(join, 1:2)
```

```
## Simple feature collection with 2 features and 3 fields
## geometry type: POINT
## dimension: XY
## bbox: xmin: 1204143 ymin: 5048309 xmax: 1822492 ymax: 5650492
## epsg (SRID): 2193
## proj4string: +proj=tmerc +lat_0=0 +lon_0=173 +k=0.9996 +x_0=1600000 +y_0=5000000
## # A tibble: 2 x 4
##   t50_fid elevation Name          geometry
##   <int>    <int> <chr>          <POINT [m]>
## 1 2353944    2723 Southland (1204143 5049971)
## 2 2354404    2820 Otago      (1234725 5048309)
```



Spatial attribute operations on vector data

Further reading: <https://geocompr.robinlovelace.net/spatial-operations.html#spatial-vec>



Your turn

- Filter the Canterbury region from **nz**, and find all summits of **nz_height** that do not intersect with the Canterbury region (both datasets come with the **spData** package).
- What happens if we spatially join the elevation column of **nz_height** to **nz**?



Geometric operations



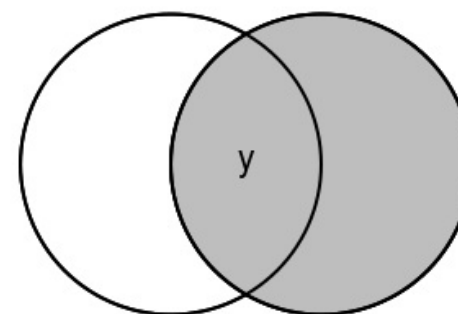
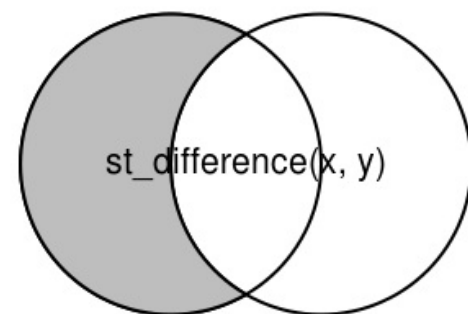
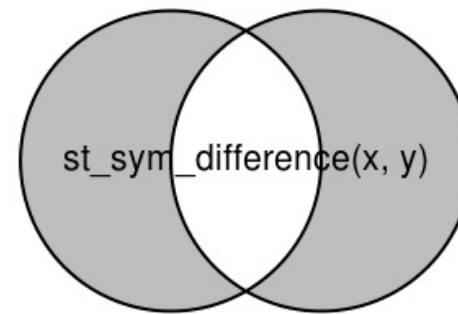
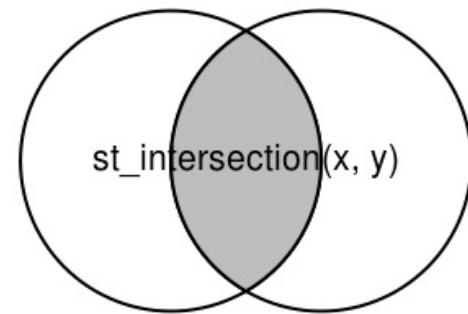
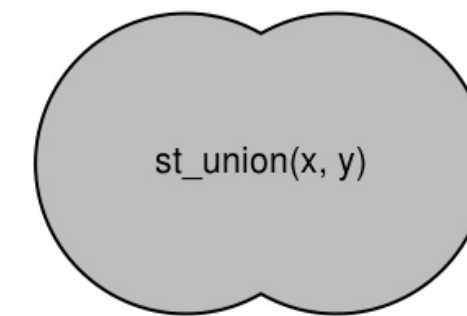
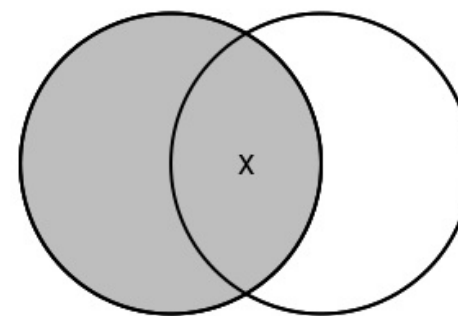
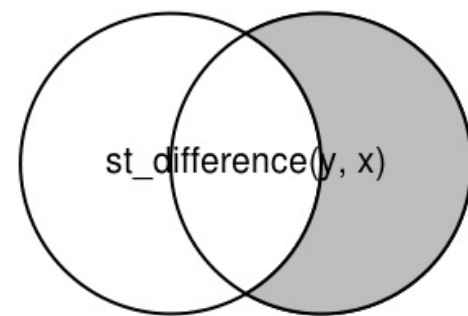
Geometric operations

What if we want the geometric intersection of two overlapping spatial objects instead of a boolean vector?



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Spatial aggregation (dissolving polygons)

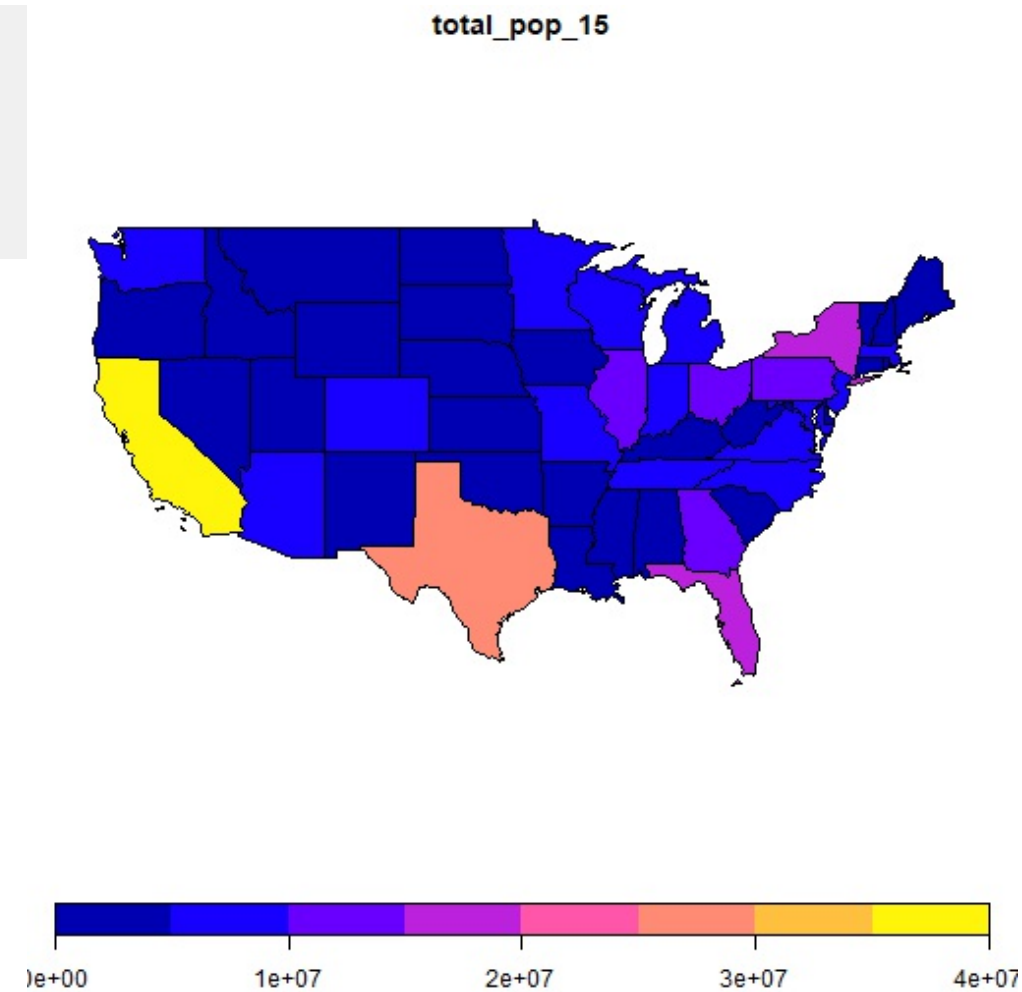


```
library(spData)
us_states %>%
  select(total_pop_15) %>%
  plot
```

Spatial aggregation (dissolving polygons)



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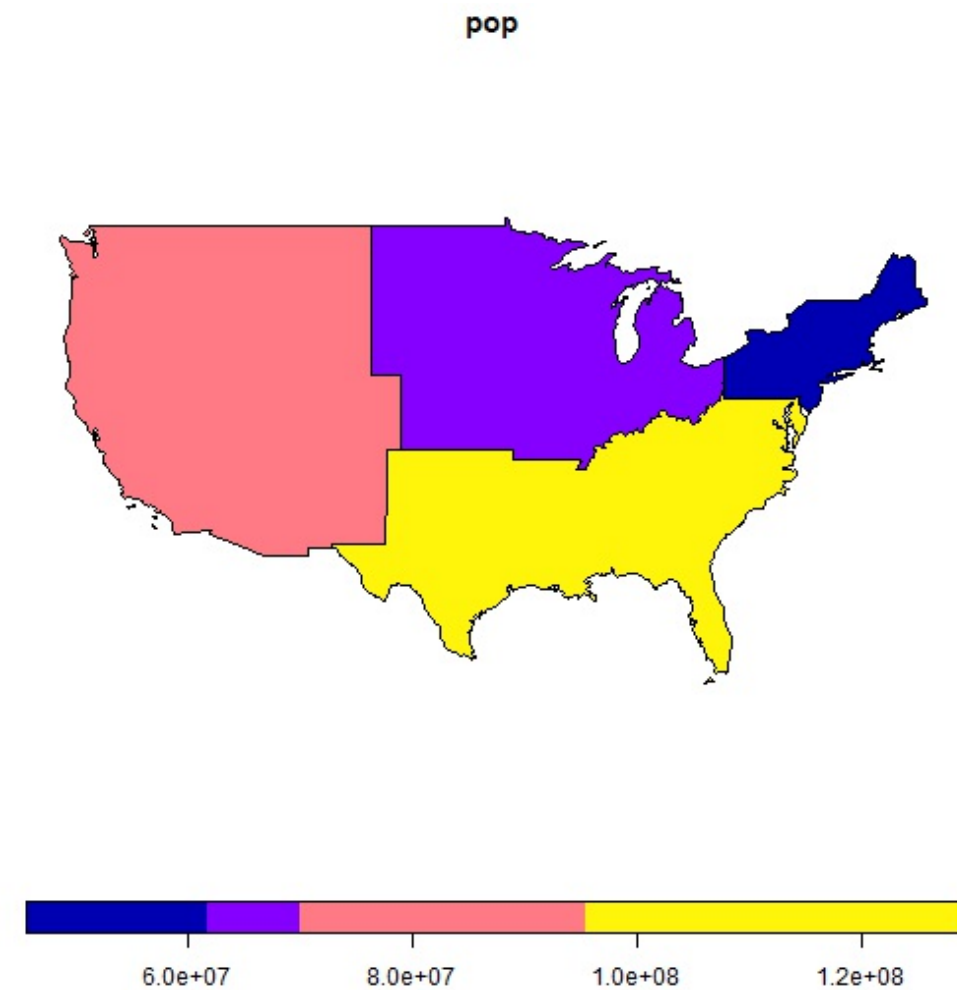


```
regions = us_states %>%  
  group_by(REGION) %>%  
  summarize(pop = sum(total_pop_15,  
                      na.rm = TRUE))  
regions %>%  
  select(pop) %>%  
  plot
```

Spatial aggregation (dissolving polygons)



```
regions = us_states %>%  
  group_by(REGION) %>%  
  summarize(pop = sum(total_pop_15,  
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regions %>%  
  select(pop) %>%  
  plot
```



CRS in sf



`sf` lets you use CRS and change CRS (reproject) through **Proj.4**.



CRS in sf

sf lets you use CRS and change CRS (reproject) through **Proj.4**.

```
st_crs(4326)
```

```
## Coordinate Reference System:  
##   EPSG: 4326  
##   proj4string: "+proj=longlat +datum=WGS84 +no_defs"
```



CRS in sf

Find out about a projection of a spatial object:

```
st_crs(us_states)
```

```
## Coordinate Reference System:  
##   EPSG: 4269  
##   proj4string: "+proj=longlat +datum=NAD83 +no_defs"
```



CRS in sf

Find out about a projection of a spatial object:

```
st_crs(us_states)
```

```
## Coordinate Reference System:  
##   EPSG: 4269  
##   proj4string: "+proj=longlat +datum=NAD83 +no_defs"
```

Change the CRS with the help of `st_transform()`:

```
st_transform(us_states, crs = 4326)
```

Further reading

Geometric operations on vector data





Your turn

- Create two overlapping circles (see below) and compute and plot their geometric intersection. Secondly union the circles.

```
pts = st_sfc(st_point(c(0, 1)), st_point(c(1, 1))) # create 2 points
# use the buffer function to create circles from points
circles = st_buffer(pts, dist = 1)
x = circles[1, ]
y = circles[2, ]
```



- Compute the average population (**total_pop_15**) for each **REGION** of **us_states**. Plot your result.
- Find out about the CRS of **nz**, reproject it into a geographic CRS (EPSG: 4326) and plot the original **nz** object next to your transformed **nz** object.



Recap

We have learned how to perform with **sf**-objects:

- Attribute operations
- Spatial attribute operations
- Geometric operations