

# Assignment: Spatial Analysis in R

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## OVERVIEW

This exercise accompanies the lessons in Environmental Data Analytics (ENV872L) on spatial analysis.

## Directions

1. Change “Student Name” on line 3 (above) with your name.
2. Use the lesson as a guide. It contains code that can be modified to complete the assignment.
3. Work through the steps, **creating code and output** that fulfill each instruction.
4. Be sure to **answer the questions** in this assignment document. Space for your answers is provided in this document and is indicated by the “>” character. If you need a second paragraph be sure to start the first line with “>”. You should notice that the answer is highlighted in green by RStudio.
5. When you have completed the assignment, **Knit** the text and code into a single HTML file.
6. After Knitting, please submit the completed exercise (HTML file) to the dropbox in Sakai. Please add your last name into the file name (e.g., “Fay\_A09\_SpatialAnalysis.pdf”) prior to submission.

## DATA WRANGLING

### 1. Prepare the workspace

- Import: tidyverse, sf, and leaflet

```
library(tidyverse, quietly = TRUE)
library(sf, quietly = TRUE)
library(leaflet, quietly = TRUE)
library(mapview)
```

### 2. Read filtered county features into an sf dataframe and plot

In this exercise, we will be exploring stream gage height data in Nebraska, as there’s been recent floods there. First, we will import from the US Counties shapefile we’ve used in lab lessons, filtering it this time for just Nebraska counties. Nebraska’s state FIPS code is 31 (as North Carolina’s was 37).

- Read the cb\_2017\_us\_county\_20m.shp shapefile into an sf dataframe
- Filter for Nebraska counties (State FIPS = 31)
- Show the dataset’s coordinate reference system
- Plot the records as a map (in any format)

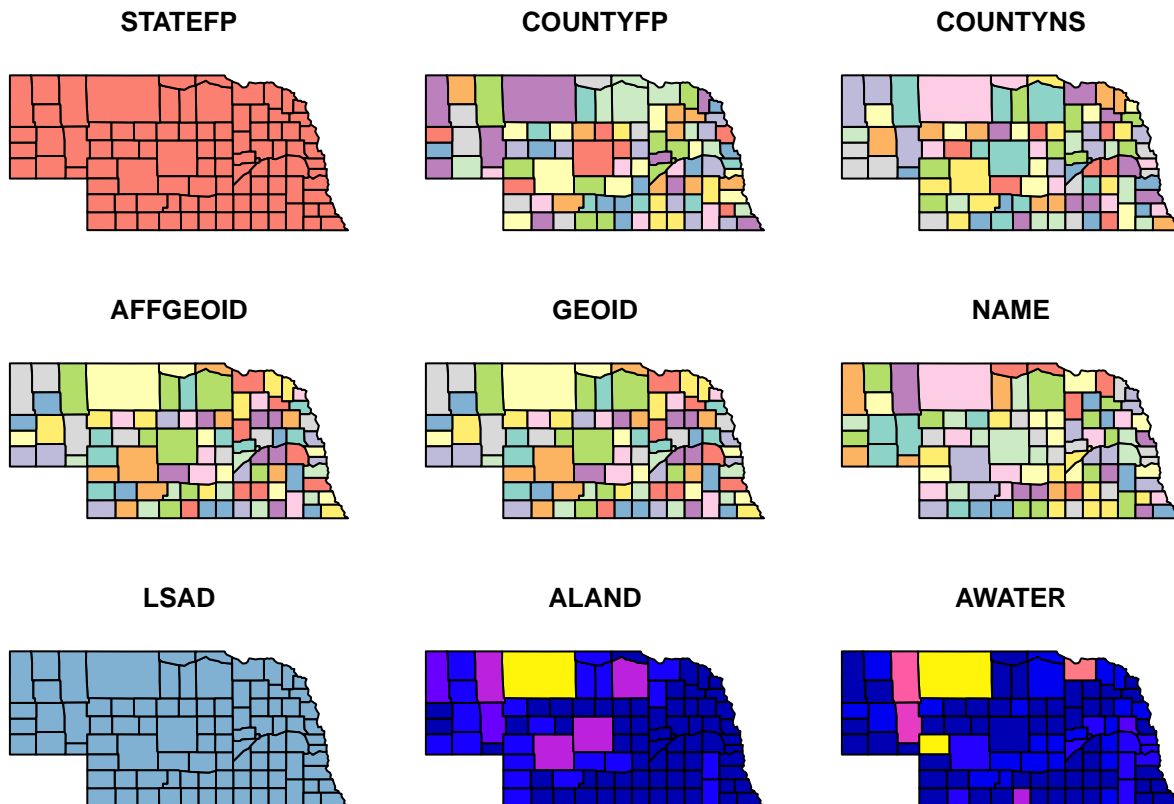
```
#Read in Counties shapefile into an sf dataframe, filtering for just NE counties
NE_counties <- st_read('../Data/Spatial/cb_2017_us_county_20m.shp') %>%
  filter(STATEFP == 31)
```

```
## Reading layer `cb_2017_us_county_20m' from data source `/Users/jakegreif/Environmental_Data_Analytic
## Simple feature collection with 3220 features and 9 fields
## geometry type:  MULTIPOLYGON
## dimension:      XY
## bbox:           xmin: -179.1743 ymin: 17.91377 xmax: 179.7739 ymax: 71.35256
## epsg (SRID):    4269
## proj4string:     +proj=longlat +datum=NAD83 +no_defs
```

```
#Reveal the CRS of the counties features
st_crs(NE_counties)
```

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

```
#Plot the data
plot(NE_counties)
```



> QUESTION: What is the EPSG code of the Counties dataset? Using <http://spatialreference.org>, is this a geographic or a projected coordinate system? (Or, does this CRS use angular or planar coordinate units?) To what datum is this CRS associated?

ANSWER: The EPSG code of the Counties dataset is 4269, which is a projected coordinate system. The CRS is associated with NAD 1983.

### 3. Read in gage locations csv as a dataframe, then display the column names it contains

Next we'll read in some USGS/NWIS gage location data I've added to the **Data/Raw** folder. These are in the `NWIS_SiteInfo_NE_RAW.csv` file. (See `NWIS_SiteInfo_NE_RAW.README.txt` for more info on this dataset.) \* Read the `NWIS_SiteInfo_NE_RAW.csv` file into a standard dataframe \* Display the column names of this dataset

```
#Read in gage locations csv as a dataframe
NE_gages <- read.csv("../Data/Raw/NWIS_SiteInfo_NE_RAW.csv")

#Reveal the names of the columns
names(NE_gages)

## [1] "site_no"          "station_nm"       "site_tp_cd"
## [4] "dec_lat_va"       "dec_long_va"      "coord_acy_cd"
## [7] "dec_coord_datum_cd"
```

QUESTION: What columns in the dataset contain the x and y coordinate values, respectively?

ANSWER: x = "dec\_long\_va", y = "dec\_lat\_va"

### 4. Convert the gage locations dataframe to an sf dataframe of points

- These data use the same coordinate reference system as the counties dataset
- Display the column names of the resulting sf dataframe

```
#Convert to an sf object
NE_gages_sf <- st_as_sf(NE_gages, coords = c("dec_long_va", "dec_lat_va"), crs=4269)

#Reveal the structure
names(NE_gages_sf)

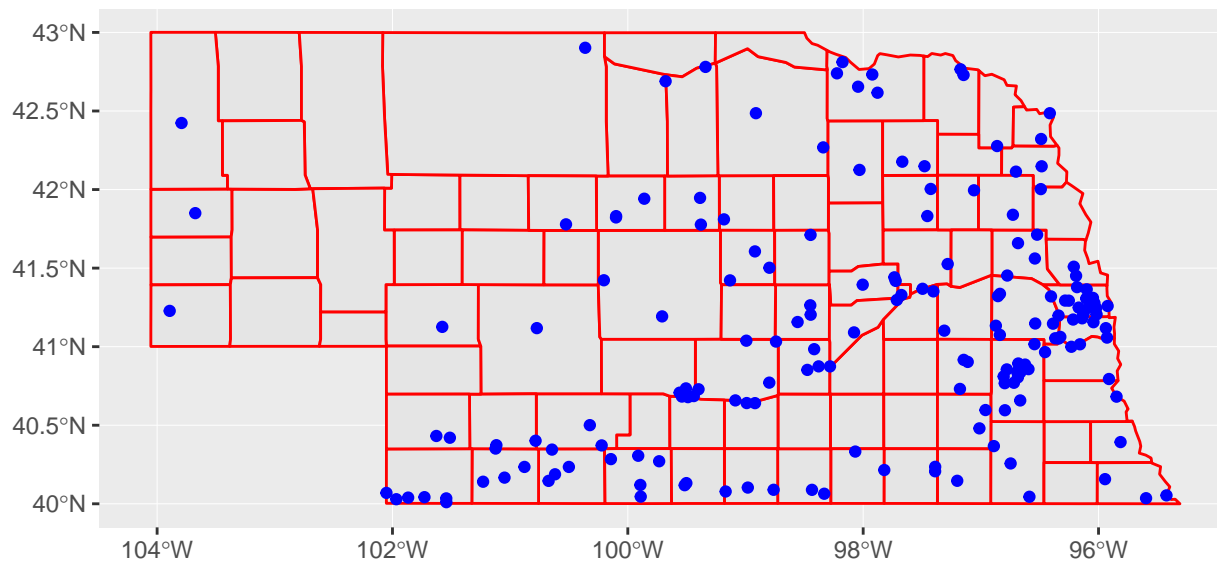
## [1] "site_no"          "station_nm"       "site_tp_cd"
## [4] "coord_acy_cd"     "dec_coord_datum_cd" "geometry"
```

QUESTION: What new field(s) appear in the sf dataframe created? What field(s), if any, disappeared? ANSWER: A new field called "geometry" has been created, and both "dec\_long\_va" and "dec\_lat\_va" disappeared.

### 5. Use ggplot to plot the gage locations on top of the counties

- Plot the different datasets in different colors

```
ggplot() +
  geom_sf(data = NE_counties, col='red') +
  geom_sf(data = NE_gages_sf, col='blue')
```



## 6. Read in the gage height data and join the site location data to it.

And finally, we want to attach some gage height data to our site locations. I've constructed a csv file listing many of the Nebraska gage sites, by station name and site number along with stream gage heights (in meters) recorded during the recent flood event. This file is titled `NWIS_SiteFlowData_NE_RAW.csv` and is found in the `Data/Raw` folder.

- Read this dataset in as a dataframe.
- Join our site information (already imported above) to these gage height data.
- The `site_no` and `station_nm` can both serve as joining attributes.
- Construct this join so that the result only includes records where both tables have data.
- Show the column names in this resulting dataframe
- Once joined, we will again have to convert this product (a dataframe) into a spatial dataframe. Do that.

```
#Read in the data
NE_gage_heights <- read.csv("../Data/Raw/NWIS_SiteFlowData_NE_RAW.csv")

#Show the column names
names(NE_gage_heights)

## [1] "site_no"      "station_nm"  "date"        "gage_ht"

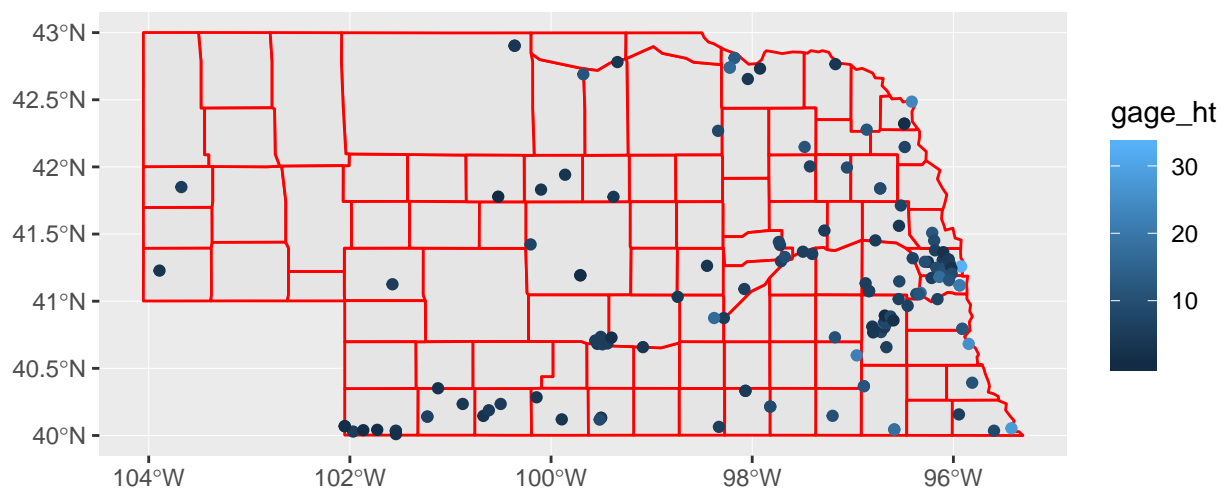
#Join location data to it
NE_gages_join <- NE_gages_sf %>%
  inner_join(y = NE_gage_heights, by = c("site_no" = "site_no"))
```

```
#Convert back to sf dataframe
NE_gages_join_sf <- st_as_sf(NE_gages_join,coords = c("dec_long_va","dec_lat_va"),crs=4269)
```

## 7. Map the pattern of gage height data

Now we can examine where the flooding appears most acute by visualizing gage heights spatially. \* Plot the gage sites on top of counties \* Show the magnitude of gage height by color, shape, other visualization technique.

```
#Plot the values
ggplot() +
  geom_sf(data = NE_counties,col='red') +
  geom_sf(data = NE_gages_join_sf, aes(color = gage_ht))
```



## SPATIAL ANALYSIS

Up next we will do some spatial analysis with our data. To prepare for this, we should transform our data into a projected coordinate system. We'll choose UTM Zone 14N (EPSG = 32614).

## 8. Transform the counties and gage site datasets to UTM Zone 14N

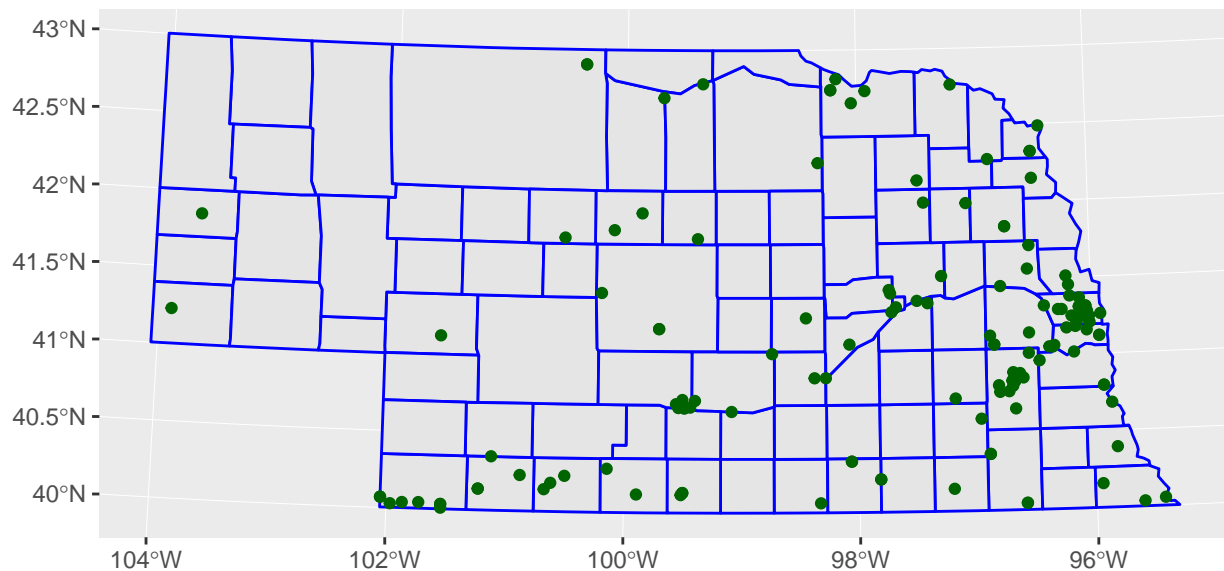
- Transform each dataset to crs 32614
- Using ggplot, plot the data so that each can be seen as different colors

```

#Transform the counties and gage location datasets to UTM Zone 14
NE_counties_utm <- st_transform(NE_counties, c=32614)
NE_gages_join_sf_utm <- st_transform(NE_gages_join_sf, c=32614)

#Plot the data
ggplot() +
  geom_sf(data = NE_counties_utm, col = 'blue') +
  geom_sf(data = NE_gages_join_sf_utm, col = 'darkgreen')

```



> QUESTION: The shape of Nebraska should look a bit different than the one created in Step 5? Why? >  
 ANSWER: The shape looks different because linear distances are different from angular distances. A flat map does not accurately represent linear distances, therefore the shape of the state needs to be distorted in order for linear distances to be preserved.

## 9. Select the gages falling within a given county

Now let's zoom into a particular county and examine the gages located there. \* Select Lancaster county from your county sf dataframe \* Select the gage sites falling **within** that county \* Remember you'll have to create a mask and then apply that mask \* Create a plot showing: \* all Nebraska counties, \* the selected county, \* and the gage sites in that county

```

#Select the county
Lancaster_county <- NE_counties_utm %>%
  filter(NAME == "Lancaster")

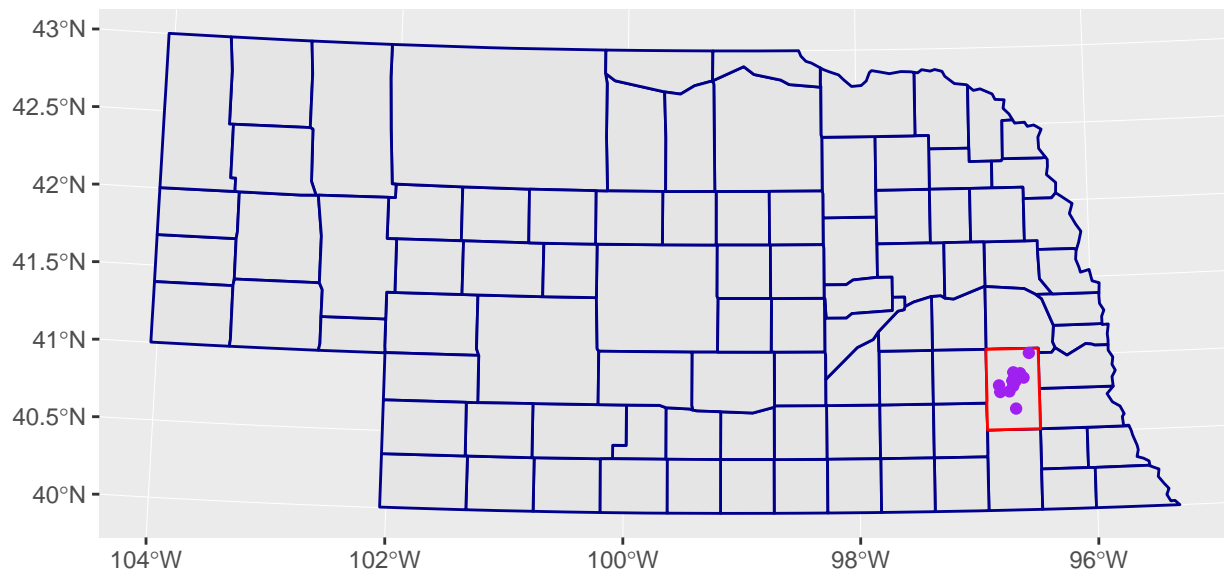
#Select gages within

```

```
Mask <- st_intersects(Lancaster_county,
                      NE_gages_join_sf_utm,
                      sparse = FALSE)

Lancaster_gages <- NE_gages_join_sf_utm[Mask,]

#Plot
ggplot() +
  geom_sf(data = NE_counties_utm, col = 'darkblue') +
  geom_sf(data = Lancaster_county, col = 'red') +
  geom_sf(data = Lancaster_gages, col = 'purple')
```



## 10. Tag each gage site with the name of the county in which it falls

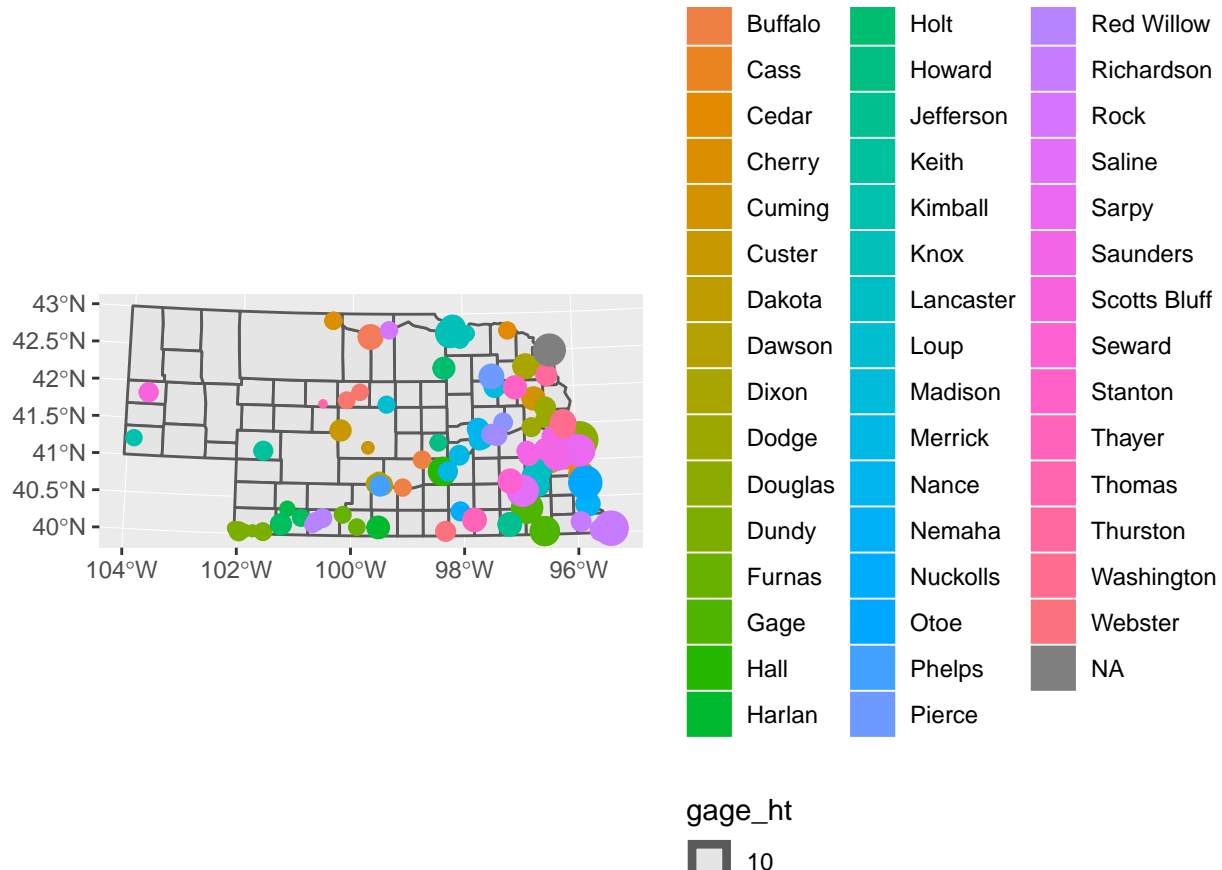
A spatial join (`st_join`) allows us to assign the attributes of an overlapping feature onto a another feature. We will use to assign each gage location the attributes of the county in which it is located. \* Spatially join the county features to the gage height features \* Display the list of fields in the resulting dataset \* Map the gage locations, \* Include county boundaries \* Displaying each gage locations county “NAME” as a different color. \* Display each gage size proportional to its “gage\_ht” value

```
#Join features
NE_gage_counties <- st_join(NE_gages_join_sf_utm, NE_counties_utm)

#Show column names
names(NE_gage_counties)
```

```
## [1] "site_no"          "station_nm.x"      "site_tp_cd"
## [4] "coord_acy_cd"     "dec_coord_datum_cd" "station_nm.y"
## [7] "date"             "gage_ht"           "STATEFP"
## [10] "COUNTYFP"        "COUNTYNS"         "AFFGEOID"
## [13] "GEOID"            "NAME"              "LSAD"
## [16] "ALAND"            "AWATER"            "geometry"
```

```
#Plot
ggplot() +
  geom_sf(data = NE_counties_utm) +
  geom_sf(data = NE_gage_counties, aes(size = gage_ht, fill = NAME, color = NAME))
```



## 11. Summarize data by county

Finally, we'll summarize our gage height data by county and then display each county by its mean gage height. \* Group the spatially joined gage location/county dataset on the county name \* Compute mean gage height \* Join (non-spatially) this result to our county sf dataframe \* Prior to joining, you'll need to drop the geometry column from the gage locations \* To do this, see the `st_drop_geometry` function \* Plot the counties showing mean gage heights for each county \* Not all counties will have data

```
#Group and summarize
NE_county_mean <- NE_gage_counties %>%
  group_by(NAME) %>%
  summarize(avg_gage_ht = mean(gage_ht))
```

```
## Warning: Factor `NAME` contains implicit NA, consider using
## `forcats::fct_explicit_na`
```



```
## Warning: Factor `NAME` contains implicit NA, consider using
## `forcats::fct_explicit_na`
```

```
#Convert result to a simple dataframe
```

```
NE_county_mean_sf <- st_as_sf(NE_county_mean)
```

```
#Join summary to County fc
```

```
NE_county_mean_drop <- st_drop_geometry(NE_county_mean_sf)
```

```
NE_county_mean_join <- NE_counties_utm %>%
```

```
  inner_join(y = NE_county_mean_drop, by = c("NAME" = "NAME"))
```

```
#Plot
```

```
ggplot() +
```

```
  geom_sf(data = NE_counties_utm) +
```

```
  geom_sf(data = NE_county_mean_join, aes(fill = avg_gage_ht, color = avg_gage_ht))
```

