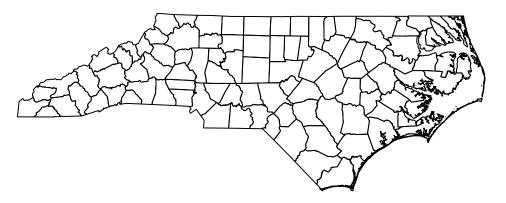
## Part 1 3 Variables

This is just for the sake of completeness.

```
library(sp)
library(spmodel)
## Warning: package 'spmodel' was built under R version 4.3.3
library(sf)
## Warning: package 'sf' was built under R version 4.3.3
## Linking to GEOS 3.11.0, GDAL 3.5.3, PROJ 9.1.0; sf_use_s2() is TRUE
library(spdep)
## Warning: package 'spdep' was built under R version 4.3.3
## Loading required package: spData
## Warning: package 'spData' was built under R version 4.3.3
## To access larger datasets in this package, install the spDataLarge
## package with: 'install.packages('spDataLarge',
## repos='https://nowosad.github.io/drat/', type='source')'
library(tmap)
## Warning: package 'tmap' was built under R version 4.3.3
library(corrplot)
## Warning: package 'corrplot' was built under R version 4.3.3
## corrplot 0.95 loaded
library(readr)
library(tidyverse)
```

```
## -- Attaching core tidyverse packages -----
                                                ----- tidyverse 2.0.0 --
               1.1.2
                        v purrr
## v dplyr
                                     1.0.2
## v forcats
              1.0.0
                         v stringr
                                     1.5.0
              3.5.1
                         v tibble
                                     3.2.1
## v ggplot2
## v lubridate 1.9.2
                         v tidyr
                                     1.3.0
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                    masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
library(tinytex)
library(ggcorrplot)
## Load in the demographic data and county data for North Carolina
ncdata=read.csv("NCcounty2024data.csv",header=T)
load("nc_counties.Rdata")
## Plot the counties
plot(nc_counties)
```



## ## Names of the counties nc\_counties[[7]]

## [1] "Alamance" "Alexander" "Alleghany" "Anson" "Ashe" ## [6] "Avery" "Beaufort" "Bertie" "Bladen" "Brunswick" ## [11] "Buncombe" "Burke" "Cabarrus" "Caldwell" "Camden" [16] "Carteret" "Caswell" "Catawba" "Chatham" "Cherokee" [21] "Chowan" "Clay" "Cleveland" "Columbus" "Craven" ## [26] "Cumberland" "Currituck" "Dare" "Davidson" "Davie" [31] "Duplin" "Durham" "Franklin" ## "Edgecombe" "Forsyth" [36] "Gaston" "Graham" ## "Gates" "Granville" "Greene" [41] "Guilford" "Henderson" ## "Halifax" "Harnett" "Haywood" ## [46] "Hertford" "Hoke" "Hvde" "Iredell" "Jackson" "Jones" "Lee" ## [51] "Johnston" "Lenoir" "Lincoln" ## [56] "Macon" "Madison" "Martin" "McDowell" "Mecklenburg" ## [61] "Mitchell" "Montgomery" "Moore" "Nash" "New Hanover" ## [66] "Northampton" "Onslow" "Orange" "Pamlico" "Pasquotank" [71] "Pender" "Pitt" "Polk" ## "Perquimans" "Person" "Rockingham" ## [76] "Randolph" "Richmond" "Robeson" "Rowan"

```
## [81] "Rutherford"
                                                        "Stanly"
                                                                        "Stokes"
                         "Sampson"
                                        "Scotland"
## [86] "Surry"
                         "Swain"
                                        "Transylvania" "Tyrrell"
                                                                        "Union"
## [91] "Vance"
                         "Wake"
                                                        "Washington"
                                        "Warren"
                                                                        "Watauga"
## [96] "Wayne"
                         "Wilkes"
                                        "Wilson"
                                                        "Yadkin"
                                                                        "Yancey"
## Obtain neighborhoods given the polygons
nc_neigh=poly2nb(nc_counties)
# Convert to sf object before merge
nc_counties_sf <- st_as_sf(nc_counties)</pre>
# Make sure the name matches
nc_counties_sf$County <- nc_counties_sf$NAME_2</pre>
nc_full <- left_join(nc_counties_sf, ncdata, by = "County")</pre>
#Neighborhoods for our analyses
# Define spatial neighbors using Queen's Contiquity
nc_nb <- poly2nb(nc_counties_sf, queen = TRUE)</pre>
# Convert to a spatial weights list object
nc_listw <- nb2listw(nc_nb, style = "W")</pre>
For Food Insecurity:
# Moran's I Test (Global Spatial Autocorrelation)
moran test Food <- moran.test(nc full$Food.Insecurity.2022, nc listw)
# Geary's C Test (Local Spatial Autocorrelation)
geary_test_Food <- geary.test(nc_full$Food.Insecurity.2022, nc_listw)</pre>
# Print results
print(moran_test_Food)
##
## Moran I test under randomisation
## data: nc_full$Food.Insecurity.2022
## weights: nc_listw
## Moran I statistic standard deviate = 5.9675, p-value = 1.205e-09
## alternative hypothesis: greater
## sample estimates:
## Moran I statistic
                            Expectation
                                                  Variance
         0.378017774
                           -0.010101010
##
                                              0.004230048
print(geary_test_Food)
```

```
## Geary C test under randomisation
##
## data: nc full$Food.Insecurity.2022
## weights: nc_listw
## Geary C statistic standard deviate = 5.0315, p-value = 2.433e-07
## alternative hypothesis: Expectation greater than statistic
## sample estimates:
## Geary C statistic
                           Expectation
                                                 Variance
         0.650030681
##
                           1.000000000
                                             0.004837982
# Moran's I Test (Global Spatial Autocorrelation)
moran_test_Overdose <- moran.test(nc_full$Overdose.Deaths.Per.100.000, nc_listw)
# Geary's C Test (Local Spatial Autocorrelation)
geary_test_Overdose <- geary.test(nc_full$Overdose.Deaths.Per.100.000, nc_listw)</pre>
# Print results
print(moran_test_Overdose)
##
## Moran I test under randomisation
## data: nc_full$Overdose.Deaths.Per.100.000
## weights: nc_listw
## Moran I statistic standard deviate = 3.2762, p-value = 0.000526
## alternative hypothesis: greater
## sample estimates:
## Moran I statistic
                           Expectation
                                                 Variance
        0.202493543
                          -0.010101010
##
                                             0.004210749
print(geary_test_Overdose)
##
## Geary C test under randomisation
## data: nc_full$Overdose.Deaths.Per.100.000
## weights: nc_listw
## Geary C statistic standard deviate = 3.4488, p-value = 0.0002816
## alternative hypothesis: Expectation greater than statistic
## sample estimates:
## Geary C statistic
                           Expectation
                                                 Variance
         0.757639361
                           1.000000000
                                             0.004938503
##
# Moran's I Test (Global Spatial Autocorrelation)
moran_test_Poverty <- moran.test(nc_full$Percent.Children.in.Poverty, nc_listw)</pre>
# Geary's C Test (Local Spatial Autocorrelation)
geary_test_Poverty <- geary.test(nc_full$Percent.Children.in.Poverty, nc_listw)</pre>
```

```
# Print results
print(moran_test_Poverty)
## Moran I test under randomisation
## data: nc_full$Percent.Children.in.Poverty
## weights: nc_listw
## Moran I statistic standard deviate = 6.5275, p-value = 3.343e-11
## alternative hypothesis: greater
## sample estimates:
## Moran I statistic
                           Expectation
                                                Variance
##
         0.417023892
                          -0.010101010
                                             0.004281657
print(geary_test_Poverty)
##
## Geary C test under randomisation
## data: nc_full$Percent.Children.in.Poverty
## weights: nc listw
##
## Geary C statistic standard deviate = 6.2864, p-value = 1.625e-10
## alternative hypothesis: Expectation greater than statistic
## sample estimates:
## Geary C statistic
                           Expectation
                                                Variance
         0.575068377
                           1.000000000
##
                                             0.004569172
moran_test_Uninsured <- moran.test(nc_full$Uninsured.Residents.2020, nc_listw)
# Geary's C Test (Local Spatial Autocorrelation)
geary_test_Uninsured <- geary.test(nc_full$Uninsured.Residents.2020, nc_listw)</pre>
# Print results
print(moran_test_Uninsured)
##
## Moran I test under randomisation
## data: nc_full$Uninsured.Residents.2020
## weights: nc_listw
## Moran I statistic standard deviate = 4.2552, p-value = 1.044e-05
## alternative hypothesis: greater
## sample estimates:
## Moran I statistic
                           Expectation
                                                Variance
         0.263314499
                          -0.010101010
                                             0.004128678
##
print(geary_test_Uninsured)
```

```
##
## Geary C test under randomisation
##
## data: nc_full$Uninsured.Residents.2020
## weights: nc_listw
##
## Geary C statistic standard deviate = 3.9943, p-value = 3.244e-05
## alternative hypothesis: Expectation greater than statistic
## sample estimates:
## Geary C statistic Expectation Variance
## 0.70740365 1.00000000 0.00536597
```