# Spatial modeling of gun-related homicide rates

Phillip Yates 10/29/2020

Storing this here as a reference because it's a good example of Modeling for Spatial Correlation and it hasn't been covered in any class yet so I will want to look at this later.

First we will load the tidyverse library for data reshaping and visualization.

```
library(tidyverse)
```

The data from this analysis come from "Firearm legislation and firearm morality in the USA: a cross-sectional, state-level study" by Kalesan et al. (2016). The response variable,  $Y_i$ , is the log firearm-related death rate (i.e., the log of the number of deaths divided by the population) in 2010 in state i. This is regressed onto five potential cofounders.

- 1. log 2009 firearm death rate per 10,000 people
- 2. Firearm ownership rate quartile
- 3. Unemployment rate quartile
- 4. Non-firearm homicide rate quartile
- 5. Firearm export rate quartile

The covariate of interest is the number of gun control laws in effect in the state. This gives p=6 covariates.

We fit the linear model

$$Y_i = \beta_0 + \sum_{j=1}^p X_i \beta_j + \epsilon_i.$$

We compute the usual non-spatial model with  $\epsilon_i \sim \text{Normal}(0, \sigma^2)$  with the spatial model  $\text{Cov}(\epsilon_1, \dots, \epsilon_n) \sim \text{Normal}(\mathbf{0}, \mathbf{\Sigma})$ . The covariance

$$\mathbf{\Sigma} = \tau^2 \mathbf{S} + \sigma^2 \mathbf{I}_n$$

is decomposed into a spatial covariance  $\tau^2 \mathbf{S}$  and a non-spatial covariance  $\sigma^2 \mathbf{I}_n$  where  $\mathbf{I}_n$  is the  $n \times n$  identity matrix. The spatial covariance follows the conditionally-autoregressive model  $\mathbf{S} = (\mathbf{M} - \rho \mathbf{A})^{-1}$ , where  $\mathbf{A}$  is the adjacency matrix with (i, j) element equal 1 if states i and j are neighbors and zero otherwise, and M is the diagonal matrix with  $i^{th}$  diagonal element equal to the number of states that neighbor state i.

#### Load the data

```
load("~/StatisticsMasters/bayesian/guns.rdata")
Y <- log(10000*Y/N)
Z[,1] <- log(Z[,1]) # Z is also the 2nd column in X
X <- cbind(1,Z,rowSums(X))

# Remove Alaska and Hawaii
Y <- Y[-c(2,11)]
X <- X[-c(2,11),]
n <- length(Y)
p <- ncol(X)</pre>
```

#### Fit the non-spatial model

```
ns_model <- "model{</pre>
  # Likelihood
  for(i in 1:n){
    Y[i] ~ dnorm(mu[i],taue)
    mu[i] <- inprod(X[i,],beta[])</pre>
 # Priors
  for(j in 1:p){
    beta[j] ~ dnorm(0,0.01)
 taue ~ dgamma(0.1,0.1)
  sig <- 1/sqrt(taue)</pre>
library(rjags)
dat <- list(Y=Y,n=n,X=X,p=p)</pre>
init <- list(beta=rep(0,p))</pre>
model1 <- jags.model(textConnection(ns_model),inits=init,data=dat,quiet=TRUE)</pre>
update(model1,10000,progress.bar="none")
samp1 <- coda.samples(model1, variable.name=c("beta", "sig"), n.iter=20000,</pre>
                      progress.bar="none")
summary(samp1)
##
## Iterations = 10001:30000
## Thinning interval = 1
## Number of chains = 1
## Sample size per chain = 20000
## 1. Empirical mean and standard deviation for each variable,
##
      plus standard error of the mean:
##
                Mean
                            SD Naive SE Time-series SE
## beta[1] 0.018309 0.082986 5.868e-04
                                              0.0042871
## beta[2] 0.749915 0.082229 5.814e-04
                                              0.0028864
## beta[3] -0.001870 0.019956 1.411e-04
                                              0.0007665
## beta[4] -0.013559 0.015900 1.124e-04
                                              0.0004122
## beta[5] 0.019993 0.017777 1.257e-04
                                              0.0005595
## beta[6] 0.018221 0.018292 1.293e-04
                                              0.0006310
## beta[7] -0.007745 0.004451 3.147e-05
                                              0.0001011
## sig
            0.101305 0.011458 8.102e-05
                                              0.0001262
##
## 2. Quantiles for each variable:
##
##
               2.5%
                           25%
                                     50%
                                               75%
                                                        97.5%
## beta[1] -0.13831 -0.038264 0.015396 0.072071 0.1884339
## beta[2] 0.59194 0.694109 0.748453 0.804361 0.9164515
## beta[3] -0.04113 -0.014979 -0.002198 0.011079 0.0384936
## beta[4] -0.04513 -0.024034 -0.013632 -0.002925 0.0176630
## beta[5] -0.01591 0.008396 0.020181 0.031929 0.0542824
```

```
## beta[6] -0.01760 0.006136 0.018304 0.030585 0.0538376
## beta[7] -0.01641 -0.010733 -0.007719 -0.004753 0.0008937
## sig 0.08177 0.093231 0.100326 0.108193 0.1268669
```

#### Create an adjacency matrix for the states in the US

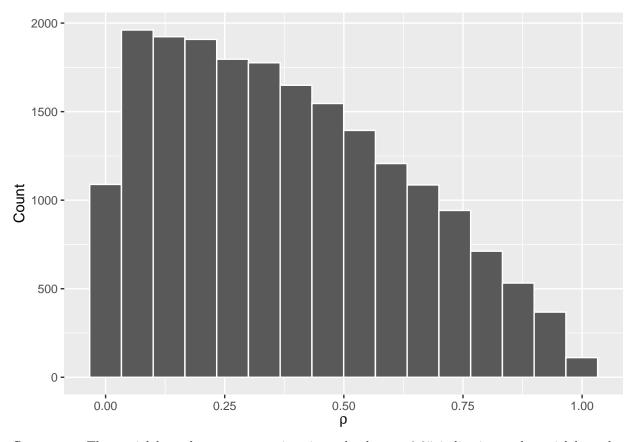
```
library(maps)
library(spdep)
library(maptools)
usa.state <- map(database="state",fill=TRUE,plot=FALSE)
# The strsplit function splits the elements of a character vector into substrings
state.ID <- sapply(strsplit(usa.state$names,":"),function(x) x[1])
usa.poly <- map2SpatialPolygons(usa.state,IDs=state.ID)
# The poly2nb function builds a neighbors list from a "polygon" list
usa.nb <- poly2nb(usa.poly)
# The nb2mat function builds a spatial weights matrix for neighbors list
A <- nb2mat(usa.nb,style="B")
A <- A[-8,] # Take out DC
A <- A[-8]
M <- diag(rowSums(A))</pre>
```

### Fit the spatial model

```
sp_model <- "model{</pre>
  # Likelihood
    for(i in 1:n){
      Y[i] ~ dnorm(mu[i]+S[i],taue)
    S[1:n] ~ dmnorm(zero[1:n],taus*0mega[1:n,1:n])
    for(i in 1:n){
      mu[i] <- inprod(X[i,],beta[])</pre>
      zero[i] <- 0
    Omega[1:n,1:n] \leftarrow M[1:n,1:n]-rho*A[1:n,1:n]
  # Priors
    for(j in 1:p){
      beta[j] ~ dnorm(0,0.01)
    taue \sim dgamma(0.1,0.1)
    taus ~ dgamma(0.1,0.1)
    rho \sim dunif(0,1)
    sig[1] <- 1/sqrt(taue)</pre>
    sig[2] <- 1/sqrt(taus)</pre>
}"
dat <- list(Y=Y,n=n,X=X,A=A,M=M,p=p)</pre>
init <- list(rho=0.99,beta=lm(Y~X-1)$coef)</pre>
model2 <- jags.model(textConnection(sp_model),inits=init,data=dat,quiet=TRUE)</pre>
```

```
update(model2,10000,progress.bar="none")
samp2 <- coda.samples(model2,variable.names=c("beta","rho","sig"),n.iter=20000,</pre>
                     progress.bar="none")
summary(samp2)
##
## Iterations = 11001:31000
## Thinning interval = 1
## Number of chains = 1
## Sample size per chain = 20000
##
## 1. Empirical mean and standard deviation for each variable,
##
     plus standard error of the mean:
##
##
                          SD Naive SE Time-series SE
               Mean
## beta[1]
          0.045755 0.112939 7.986e-04
                                            0.0076482
## beta[2] 0.760519 0.106056 7.499e-04
                                            0.0044500
## beta[3] -0.007284 0.026648 1.884e-04
                                            0.0012876
## beta[4] -0.010997 0.021054 1.489e-04
                                            0.0007331
## beta[5]
          0.015524 0.023492 1.661e-04
                                            0.0009524
## beta[6]
           0.015186 0.025590 1.810e-04
                                            0.0011993
## beta[7] -0.008098 0.005755 4.069e-05
                                            0.0001607
## rho
           0.379281 0.250670 1.773e-03
                                            0.0035509
## sig[1]
           0.104950 0.014132 9.993e-05
                                            0.0001960
           0.149645 0.026389 1.866e-04
## sig[2]
                                            0.0005220
##
## 2. Quantiles for each variable:
##
##
              2.5%
                          25%
                                    50%
                                              75%
                                                     97.5%
## beta[1] -0.16014 -0.0304465 0.039668 0.116492 0.280788
## beta[2] 0.55968 0.6896451 0.756840 0.828257 0.979974
## beta[3] -0.06210 -0.0242202 -0.006783
                                        0.010293 0.044540
## beta[4] -0.05394 -0.0245636 -0.010777
                                        0.003069 0.029439
## beta[5] -0.03155  0.0004606  0.015772  0.030816  0.062237
## beta[6] -0.03703 -0.0011419 0.015435 0.032379 0.063756
## beta[7] -0.01920 -0.0119231 -0.008219 -0.004373 0.003521
## rho
           0.01558 0.1675387
                               0.349020 0.563487 0.895657
           0.08132 0.0950696 0.103605 0.113300 0.136616
## sig[1]
           ## sig[2]
rho <- samp2[[1]][,8]
ggplot(NULL,aes(x=rho))+geom_histogram(bins=16,color="white")+
 labs(x=expression(rho),y="Count")
```

## Don't know how to automatically pick scale for object of type mcmc. Defaulting to continuous.

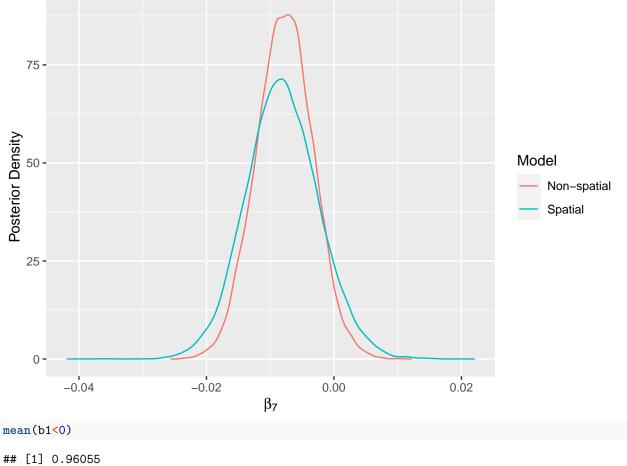


Summary: The spatial dependence parameter is estimated to be near 0.35, indicating weak spatial dependence

## Compare the results across models

The objective is to determine if the coefficient corresponding to the number of gun laws,  $\beta_7$ , is non-zero. Below we compare its posterior distribution for the spatial and non-spatial models.

```
b1 <- samp1[[1]][,7]
b2 <- samp2[[1]][,7]
d1 <- density(b1)
d2 <- density(b2)
Model <- c(rep("Non-spatial",length(d1$x)),rep("Spatial",length(d2$x)))
x <- c(d1$x,d2$x)
y <- c(d1$y,d2$y)
density.data <- as.data.frame(x=x,y=y,Model=Model)
ggplot(density.data,aes(x=x,y=y,color=Model))+geom_line()+
  labs(x=expression(beta[7]),y="Posterior Density")</pre>
```



mean(b2<0)

## [1] 0.9194

Summary: Both models provide evidence of a negative relationship between the number of gun laws and the firearm-related death rate. However, there is more uncertainty in the spatial model, which is likely more realistic.