LECTURE 20: AREAL DATA MODEL FITTING

CLASS INTRO

INTRO QUESTIONS

Last Time:

• Discuss the mechanism for include spatial random effects in for areal data. How is this simulated / how can models account for this?

Today:

 Model Fitting for Areal Data

SPDEP

ADJACENCY MATRIX

Using the code below, create an adjacency matrix for Montana. Then identify the neighbors for Gallatin county.

These functions also work with general shape files.

MORAN'S I / GEARY'S C

Recall:

• Moran's I

$$I = \frac{n \sum_{i} \sum_{j} w_{ij} (Y_i - \bar{Y}) (Y_j - \bar{Y})}{(\sum_{i \neq j} w_{ij}) \sum_{i} (Y_i - \bar{Y})^2}.$$

This is a spatial analogue measuring the lagged autocorrelation.

• Geary's C

$$C = \frac{(n-1)\sum_{i}\sum_{j}w_{ij}(Y_{i} - Y_{j})^{2}}{2(\sum_{i \neq j}w_{ii})\sum_{i}(Y_{i} - \bar{Y})^{2}}$$

MORAN'S I / GEARY'S C

- The R package spdep contains built in functions for Moran's I and Geary's C.
- moran.test() and geary.test() both take a numeric vector (response) and a listw object created by nb2listw as arguments.

MORAN'S I / GEARY'S C

Using the Tester - Rosendale election results, compute and interpret Moran's I and Geary's C with the proportion voting for Tester.

```
Tester <- read_csv('Tester_Results.csv')
Tester <- Tester %>%
   mutate(Tester_Prop = TESTER / (TESTER + ROSENDALE + BRECKENRIDGE))

#drop Yellowstone National Park
mt.poly.noYNP <- mt.poly[1:56,]
mt.nb.noYNP <- poly2nb(mt.poly.noYNP)
mt.listw <- nb2listw(mt.nb.noYNP, style = 'W')</pre>
```

SAR / CAR

The spdep package also contains the functionality to fit SAR / CAR models.

Follow the include tutorial code and answer these four questions.

- 1. Summarize the data set, note Z is a standardized (standard normal) response for PROPCAS
- 2. What is nyadjmat
- 3. Summarize the results from SAR
- 4. Choose a model between lmo, SAR, and CAR

BAYESIAN MODELS FOR AREAL DATA

CARBAYES

Similar to earlier functionality, there are R packages for analyzing areal data using Bayesian methods. We will look at CARBayes <u>Tutorial</u>

PROPERTY VALUES IN GLASGOW TUTORIAL

- Using the CARBayes package, answer the following questions.
- 1. Describe the data set
- 2. What are the results of moran.mc? What is the purpose of using resid.model as the response?
- 3. Interpret and describe the results of the S.CARleroux() model call.

JAGS

- Again JAGS is a possibility for any situation, it just requires sampling model and prior along with explicit documentation.
- Recall

$$Y_i | \psi_i \sim Poisson(E_i \exp(\psi_i))$$

 $\psi_i = x_i^T \beta + \theta_i + \phi_i$

where x_i are spatial covariates, θ_i corresponds to region wide heterogeneity, and ψ_i captures local clustering.

JAGS MODEL CODE

```
car model <- "model {</pre>
 for (i in 1 : regions) {
    O[i] ~ dpois(mu[i])
    log(mu[i]) \leftarrow log(E[i]) + beta0 + beta1*x1 + phi[i] + theta[i]
    theta[i] \sim dnorm(0.0, tau.h)
    xi[i] <- theta[i] + phi[i]</pre>
    SMRhat[i] <- 100 * mu[i] / E[i]</pre>
    SMRraw[i] <- 100 * O[i] / E[i]
  phi[1:regions] ~ car.normal(adj[], weights[], num[], tau.c)
  beta0 \sim dnorm(0.0, 1.0E-5)
  beta1 ~ dnorm(0.0, 1.0E-5)
  tau.h ~ dgamma(1.0E-3, 1.0E-3)
  tau.c ~ dgamma(1.0E-3, 1.0E-3)
  sd.h <- sd(theta[]) #marginal SD of heterogeneity effects</pre>
  sd.c <- sd(phi[]) # marginal SD of clustering effects</pre>
  alpha <- sd.c / (sd.h + sd.c)
```