### MODULE 9: Spatial Statistics in Epidemiology and Public Health Lecture 4: Spatial regression

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- Waller and Gotway (2004, Chapter 9) Applied Spatial Statistics for Public Health Data. New York: Wiley.
- Haining, R. (2003). Spatial Data Analysis: Theory and Practice. Cambridge: Cambridge University Press.
- Banerjee, S., Carlin, B.P., and Gelfand, A.E. (2014) Hierarchical Modeling and Analysis for Spatial Data, 2nd Ed. Boca Raton, FL: CRC/Chapman & Hall.
- Blangiardo, M. and Cameletti, M. (2015) Spatial and Spatio-temporal Bayesian Models with R-INLA. Chichester: Wiley.

#### What do we have so far?

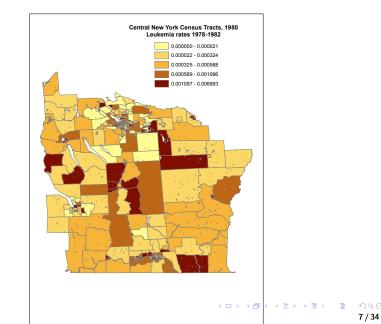
- ► Tension between statistical precision (want large local sample sizes → big regions), and geographic precision (want small regions for more detail in map).
- Disease mapping approaches use small area estimation techniques to borrow information from all areas and from neighboring areas to improve local estimation in each area.
- But what about local covariates?
- Can we adjust for those (say, using regression models)?
- And still borrow information?
- With *independent* observations we know how to use *linear* and generalized linear models such as linear, Poisson, logistic regression.
- What happens with dependent observations?

"...all models are wrong. The practical question is how wrong do they have to be to not be useful." Box and Draper (1987, p. 74)

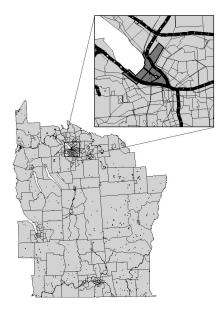
- In statistical modeling, we are often trying to describe the mean of the outcome as a function of covariates, assuming error terms are mutually independent.
- Where do correlated errors come from?
- Perhaps outcomes truly correlated (infectious disease).
- Perhaps we omitted an important variable that has spatial structure itself.

- NY leukemia data and covariates (Waller and Gotway, 2004).
- 281 census tracts (1980 Census).
- 8 counties in central New York.
- ▶ 592 cases for 1978-1982.
- 1,057,673 people at risk.

## Crude Rates (per 100,000)



#### Outliers, where are the top 3 rates?



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- Let  $Y_i$  = count for region *i*.
- Let  $E_i = expected$  count for region *i*.
- Let (x<sub>i,TCE</sub>, x<sub>i,65</sub>, x<sub>i,home</sub>) be the associated covariate values.
   Poisson regression:

$$Y_i \sim Poisson(E_i\zeta_i)$$

where

$$\log(\zeta_i) = \beta_0 + x_{i,TCE}\beta_{TCE} + x_{i,65}\beta_{65} + x_{i,home}\beta_{home}.$$

#### Poisson distribution for counts.

- Link function: Natural log of mean of Y<sub>i</sub> is a linear function of covariates.
- βs represent multiplicative increases in expected counts, e<sup>β</sup> a measure of relative risk associated with one unit increase in covariate.
- *E<sub>i</sub>* an *offset*, what we expect if the covariates have no impact.
- Age, race, sex adjustments in either *E<sub>i</sub>* (standardization) or covariates.

#### Adding spatial correlation: New York data

- Assume E<sub>i</sub> known, perhaps age-standardized, or based on global (external or internal) rates.
- Our model is

$$Y_i|\boldsymbol{\beta},\psi_i \stackrel{ind}{\sim} \mathsf{Poisson}(E_i \exp(\boldsymbol{x}_i'\boldsymbol{\beta}+\psi_i)),$$

 $\log(\zeta_i) = \beta_0 + x_{i,TCE}\beta_{TCE} + x_{i,65}\beta_{65} + x_{i,home}\beta_{home} + \psi_i.$ 

- The  $\psi_i$  represent the random intercepts.
- Add overdispersion via  $\psi_i \overset{ind}{\sim} N(0, v_{\psi})$ .
- Add spatial correlation via

$$\boldsymbol{\psi} \sim MVN(\mathbf{0}, \boldsymbol{\Sigma}).$$

- Overdispersion model (i.i.d. ψ<sub>i</sub>) results in each estimate being a compromise between the *local* SMR and the *global average* SMR.
- "Borrows information (strength)" from other observations to improve precision of local estimate.
- "Shrinks" estimate toward global mean. (Note: "shrink" does not mean "reduce", rather means "moves toward").

- Spatial model (correlated ψ<sub>i</sub>) results in each estimate begin a compromise between the *local* SMR and the *local average* SMR.
- Shrinks each  $\psi_i$  toward the average of its *neighbors*.
- Can also include *both* global and local shrinkage (Besag, York, and Mollié 1991).
- How do we fit these models?

# Bayesian inference regarding model parameters based on *posterior distribution*

 $\Pr[m{eta}, m{\psi} | m{Y}]$ 

proportional to the product of the likelihood times the prior

 $\Pr[\mathbf{Y}|\boldsymbol{\beta}, \boldsymbol{\psi}]\Pr[\boldsymbol{\psi}]\Pr[\boldsymbol{\beta}].$ 

Defers spatial correlation to the prior rather than the likelihood.

Could model *joint* distribution

$$\psi \sim MVN(\mathbf{0}, \Sigma).$$

Could also model conditional distribution

$$\psi_i | \psi_{j \neq i} \sim N\left(\frac{\sum_{j \neq i} c_{ij} \psi_j}{\sum_{j \neq i} c_{ij}}, \frac{1}{v_{CAR} \sum_{j \neq i} c_{ij}}\right), i = 1, \dots, N.$$

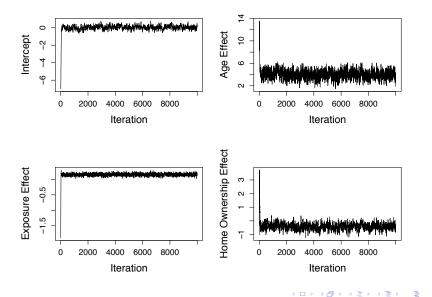
where c<sub>ij</sub> are weights defining the neighbors of region *i*.
Adjacency weights: c<sub>ij</sub> = 1 if *j* is a neighbor of *i*.

- The conditional specification defines the conditional autoregressive (CAR) prior (Besag 1974, Besag et al. 1991).
- Under certain conditions on the c<sub>ij</sub>, the CAR prior defines a valid multivariate joint Gaussian distribution.
- Variance covariance matrix a function of the *inverse* of the matrix of neighbor weights.

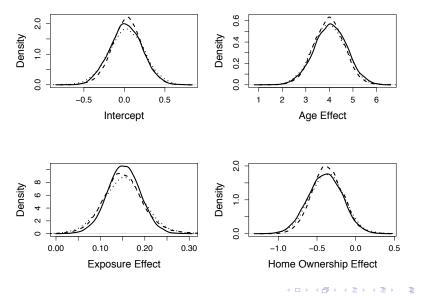
- Posterior often difficult to calculate mathematically.
- Markov chain Monte Carlo: Iterative simulation approach to model fitting.
- Given *full conditional* distributions, simulate a new value for each parameter, holding the other parameter values fixed.
- The set of simulated values converges to a sample from the posterior distribution.
- Alternative: integrated nested Laplace analysis using the inla package (example code).

$$Y_{i}|\beta,\psi_{i} \stackrel{ind}{\sim} \text{Poisson}(E_{i} \exp(\mathbf{x}_{i}'\beta + \psi_{i})),$$
  
$$\log(\zeta_{i}) = \beta_{0} + x_{i,TCE}\beta_{TCE} + x_{i,65}\beta_{65} + x_{i,home}\beta_{home} + \psi_{i}.$$
  
$$\beta_{k} \sim \text{Uniform.}$$
  
$$\psi_{i}|\psi_{j\neq i} \sim N\left(\frac{\sum_{j\neq i}c_{ij}\psi_{j}}{\sum_{j\neq i}c_{ij}}, \frac{1}{v_{CAR}\sum_{j\neq i}c_{ij}}\right), i = 1, \dots, N.$$
  
$$\frac{1}{v_{CAR}} \sim \text{Gamma}(0.5, 0.0005).$$

#### MCMC trace plots



#### Posterior densities



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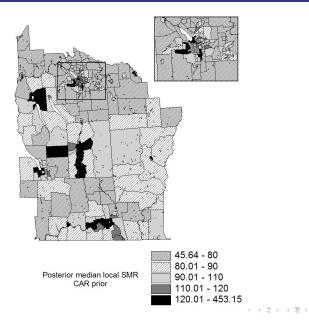
Covariate	Posterior	95% Credible
	Median	Set
$\beta_0$	0.048	(-0.355, 0.408)
$\beta_{65}$	3.984	(2.736, 5.330)
$\beta$ TCE	0.152	(0.066, 0.226)
$\beta_{home}$	-0.367	(-0.758, 0.049)

A nifty thing about MCMC estimates:

We get posterior samples from any function of model parameters by taking that function of the sampled posterior parameter values.

- Gives us posterior inference for  $SMR_i = Y_{i,fit}/E_i$ .
- Also can get Pr[SMR<sub>i</sub> > 200|Y] and map these exceedence probabilities.

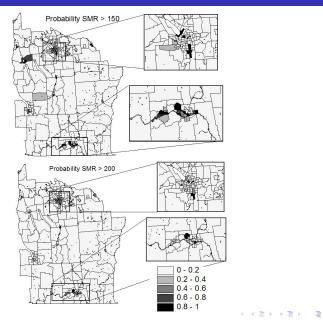
#### Posterior median SMRs



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#### Posterior exceedence probabilities



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- Associations between local covariates and local outcomes (counts and rates).
- Spatial correlation between random intercepts (inside the link function).
- (Aside: This is a clever idea since we can use a multivariate Gaussian distribution for correlation...).
- Result: Local rates adjusted for covariates and smoothed by borrowing information.
- Many examples in the literature, and many extensions, we'll start with one tomorrow!

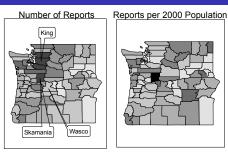
#### Bonus Example

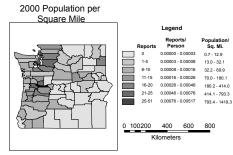
 Cryptozoology Example: Waller and Carlin (2010) Disease Mapping. In *Handbook of Spatial Statistics*, Gelfand et al. (eds.). Boca Raton: CRC/Chapman and Hall.



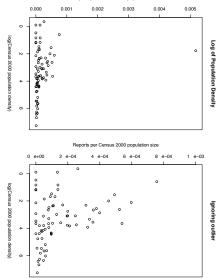
- County-specific reports of encounters with Sasquatch (Bigfoot).
- Data downloaded from www.bfro.net
- Sightings from counties in Oregon and Washington (Pacific Northwest).
- Probability of report related to population density?
- (Hopefully) rare events in small areas.
- Perhaps spatial smoothing will stabilize local rate estimates.

#### Sasquatch Data



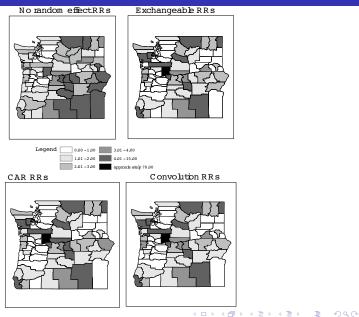


#### Reports vs. Population Density



Reports per Census 2000 population size

#### Mapped relative risks

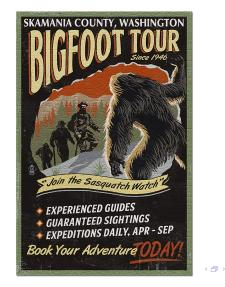


#### Skamania Sasquatch Ordinances

- http://www.skamaniacounty.org/commissioners/ homepage/ordinances-2/
- Big Foot Ordinance 69-1: "THEREFORE BE IT RESOLVED that any premeditated, willful and wonton slaying of any such creature shall be deemed a felony punishable by a fine not to exceed Ten Thousand Dollars (\$10,000.00) and/or imprisonment in the county jail for a period not to exceed Five (5) years. ADOPTED this 1st day of April, 1969."
- ▶ Big Foot Ordinance 1984-2:
  - Repealed felony and jail sentence.
  - Established a Sasquatch Refuge (Skamania County).
  - Clarified penalty (gross misdemeanor vs. misdemeanor) and penalty (fine and jail time), disallowed insanity defense, and clarified distinction between coroner designation of victim as humanoid (murder) or anthropoid (this ordinance).



# www.amazon.com/Skamania-County-Washington-Bigfoot-Vintage/dp/B076PWN7ZM



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- What method to use depends on what you want data you have and what question you want to answer.
- All methods try to balance trend (fixed effects) with correlation (here, with random effects).
- All models wrong, some models useful.
- Trying more than one approach often sensible.

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