

Hierarchical Models for Spatial Data

Models for Socio-Environmental Data

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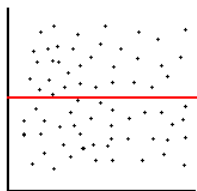


Of interest

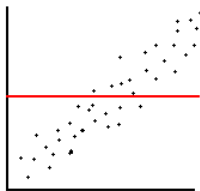
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Conn, P.B., D.S. Johnson, P.J. Williams , S.R. Melin, and M.B. Hooten. (In Press). A guide to Bayesian model checking for ecologists. Ecological Monographs.

$$\varepsilon_i = y_i - g(\boldsymbol{\theta}, \mathbf{x}_i)$$

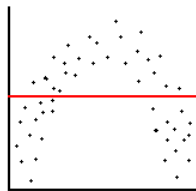
ε_i are iid



(a)



(b)



(c)

What goes wrong if we fail to account for autocorrelation?

The global issue is model checking. More specific issues include:

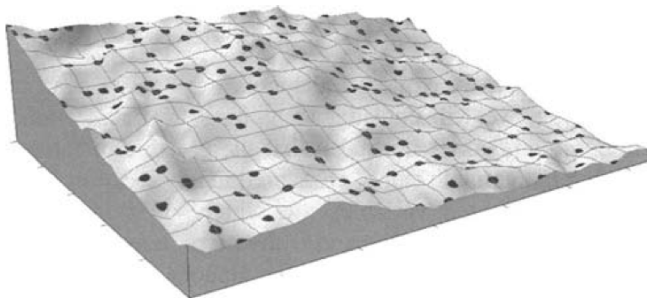
- ▶ Inference is excessively optimistic .
- ▶ Model selection favors over-parameterized models.
- ▶ Prediction errors increase.
- ▶ Your paper will not be published if it goes to a savvy reviewer.

Roadmap: Modeling spatial structure in data

- ▶ Continuous spatial processes
 - ▶ Detecting spatial dependence
 - ▶ Distance matrices
 - ▶ Semi-variograms
 - ▶ Modeling spatial dependence
- ▶ Areal spatial processes (briefly)
 - ▶ Detecting spatial dependence
 - ▶ Modeling spatial dependence

Most ecological data are spatial

Continuous spatial processes



Data for continuous spatial processes

All data points include a spatial reference.

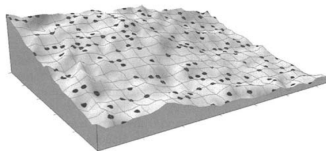
aspatial data point : y_i (1)

spatially referenced data point : $y(s_i)$ (2)

Where s_i is a vector of spatial coordinates of length 1, 2, or 3. The data are said to be continuous because they can occur at any point (s_i) in one, two, or three dimensional space. This does not mean that the value at that point ($y(s_i)$) can not be discrete.

Distance matrices

$n \times n$ matrix, i indexes rows, j indexes columns



$$\begin{pmatrix} 0 & d_{1,2} & d_{1,3} & . & . & d_{1,n} \\ d_{2,1} & 0 & d_{2,3} & . & . & d_{2,n} \\ d_{3,1} & d_{3,2} & 0 & . & . & d_{3,n} \\ . & . & . & 0 & . & . \\ . & . & . & . & 0 & . \\ d_{n,1} & d_{n,2} & . & . & d_{n,n-1} & 0 \end{pmatrix}$$

Assessing spatial correlation

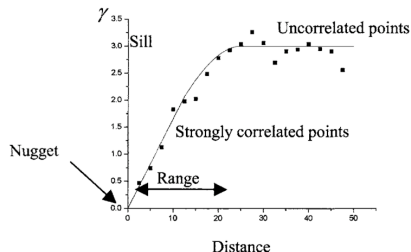
Let $\mu_i = g(\boldsymbol{\theta}, \mathbf{x}_i)$

1. Assume \mathbf{y} is measured at n spatial locations.
2. Compute the residuals: $\mathbf{e} = \mathbf{y} - \boldsymbol{\mu}$.
3. Examine the residuals \mathbf{e} for spatial correlation (i.e., autocorrelation).

Assessing spatial correlation

Empirical semi-variogram

$$\hat{\gamma}(d) = \frac{\sum_{i,j \in N(d)} (e_i - e_j)^2}{2N(d)}$$



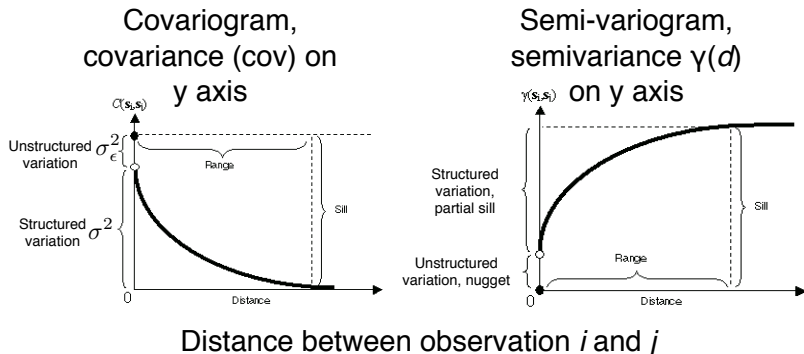
The y axis is the average squared difference between pairs of residuals at a given distance d divided by two times the number of pairs at distance d . The x axis is the distance between pairs. Distances are usually binned into categories. The notation $\sum_{i,j \in N(d)}$ means sum over the i, j pairs of data that are d distance apart.

In MCMC, compute residuals at each iteration, compute $\gamma(d)^{(k)}$ and plot variogram using posterior mean of $\gamma(d)$. Or, better, sample MCMC output for residuals in R, use R functions (geoR) to find variogram with credible intervals.

Modeling spatial structure with two sources of variation

1. **Correlated error:** The structured, process component. Varies with distance between points. Process variance here.
2. **Uncorrelated error:** The unstructured, site specific component. It includes effects of fine scale heterogeneity and measurement error.

Modeling spatial structure with two sources of variation



$$\text{cov}(d) = \text{cov}(0) - \gamma(d)$$

Figures modified from ESRI ArcGIS Desktop online manual

Remember the covariance matrix Σ

Imagine a vector of 3 random variables, $(z_1, z_2, z_3)'$. The covariance between any two of these random variables is simply an unstandardized version of the correlation between them— it is correlation measured in the units of the random variables. The covariance matrix (aka variance covariance matrix) of the random variable is:

$$\Sigma = \begin{pmatrix} \sigma_1^2 & \text{Cov}_{1,2} & \text{Cov}_{1,3} \\ \text{Cov}_{2,1} & \sigma_2^2 & \text{Cov}_{2,3} \\ \text{Cov}_{3,1} & \text{Cov}_{3,2} & \sigma_3^2 \end{pmatrix} \quad (3)$$

Generalizing, a $m \times m$ covariance matrix has the variances of the random variable on the diagonal and the covariance on the off diagonal. The covariance between random variable i and j is $\text{Cov}_{ij} = \rho \sigma_i \sigma_j$ where ρ is the correlation coefficient, which takes on values between -1 and 1 . Covariance can take on values between $-\infty$ and $+\infty$.

Remember the identity matrix \mathbf{I}

Using a 3×3 matrix to illustrate:

$$\mathbf{I} = \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix} \quad (4)$$

$$\sigma_{\varepsilon}^2 \mathbf{I} = \begin{pmatrix} \sigma_{\varepsilon}^2 & 0 & 0 \\ 0 & \sigma_{\varepsilon}^2 & 0 \\ 0 & 0 & \sigma_{\varepsilon}^2 \end{pmatrix} \quad (5)$$

Modeling spatial structure with two sources of error

$\mu_i = g(\boldsymbol{\theta}, \mathbf{x}_i)$, a model of an ecological process that can take on real values (for now).

$\boldsymbol{\mu} = g(\boldsymbol{\theta}, \mathbf{X})$, note that $\boldsymbol{\mu}$ is a vector with length = number of observations (n) and \mathbf{X} is a data matrix with number of rows = n and number of columns = number of predictor variables.

$$\mathbf{y} \sim \text{multivariate normal}(\boldsymbol{\mu}, \boldsymbol{\Sigma} + \sigma_{\varepsilon}^2 \mathbf{I}) \quad (6)$$

$\boldsymbol{\Sigma}$ is an $n \times n$ matrix with structured variance (σ^2) at distance 0 on the diagonal and the covariance between observation i and observation j on the off diagonals ($i \neq j$). \mathbf{I} an $n \times n$ matrix with ones on the diagonal and zeros elsewhere. σ_{ε}^2 is unstructured (uncorrelated) variance.

Alternative notation: random effects approach

$$\mathbf{y} = g(\boldsymbol{\theta}, X) + \boldsymbol{\eta} + \boldsymbol{\epsilon}$$

1. Correlated Error: $\boldsymbol{\eta} \sim \text{multivariate normal}(\mathbf{0}, \boldsymbol{\Sigma})$
2. Uncorrelated Error: $\boldsymbol{\epsilon} \sim \text{multivariate normal}(\mathbf{0}, \sigma_{\epsilon}^2 \mathbf{I})$

Alternative notation: hierarchical approach

$$\begin{aligned}\mathbf{y} &\sim \text{multivariate normal}(g(\boldsymbol{\theta}, \mathbf{X}) + \boldsymbol{\eta}, \sigma_{\varepsilon}^2 \mathbf{I}) \\ \boldsymbol{\eta} &\sim \text{normal}(0, \boldsymbol{\Sigma})\end{aligned}$$

1. Correlated Error: $\boldsymbol{\eta}$
2. Uncorrelated Error: σ_{ε}^2

Modeling spatial structure with two sources of error

These both imply:

$$\mathbf{y} \sim \text{multivariate normal}(g(\boldsymbol{\theta}, \mathbf{X}), \boldsymbol{\Sigma} + \sigma_{\varepsilon}^2 \mathbf{I})$$

Modeling spatial structure with two sources of error

Do we really need to predict $\frac{1}{2}(n^2 - n)$ covariances? No. Instead, we model¹ them as a function of distance using parametric covariance functions²:

► Exponential: $\Sigma_{i,j} = \sigma^2 \exp\left(-\frac{d_{i,j}}{\phi}\right)$

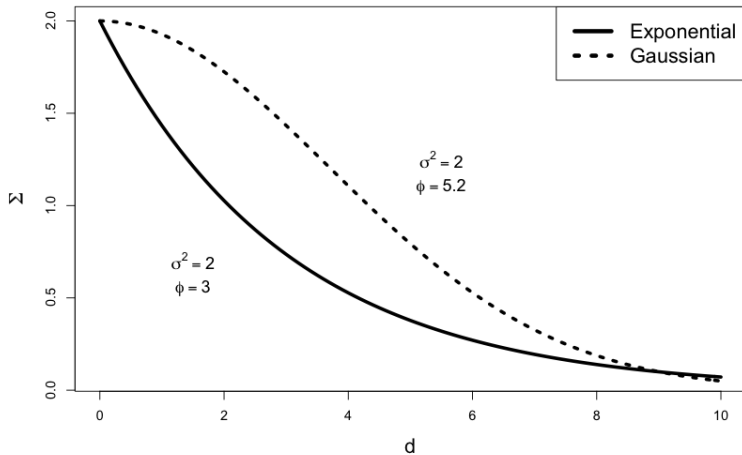
► Gaussian: $\Sigma_{i,j} = \sigma^2 \exp\left(-\frac{d_{i,j}^2}{\phi^2}\right)$

where $d_{i,j}$ = distance between locations i and j . Note that an aspatial model would require approximating the posterior distribution of a single variance parameter σ^2 . The spatial equivalent requires three: σ^2 , ϕ , and σ_ε^2 . Also note that when $i = j$ such that we are “at” a location, $d_{i,j} = 0$ and $\Sigma_{i,j} = \sigma^2$.

¹This is a great illustration of the main purpose of science: dimension reduction.

²There are many others, but these are used most frequently.

Modeling spatial structure with two sources of error



Important assumptions

- ▶ **Stationarity**: spatial structure does not vary with location, which means that the spatial correlation does not change within the area being analyzed.
- ▶ **Isotropy**: spatial structure does not vary with direction, which means the spatial correlation does not change with direction.

Toy illustration for 3 data points and simple linear regression

$$\begin{aligned}
 \mathbf{y} &= (y(\mathbf{s}_1), y(\mathbf{s}_2), y(\mathbf{s}_3)) \\
 \boldsymbol{\beta} &= (\beta_0, \beta_1) \\
 \mathbf{X} &= \begin{pmatrix} 1 & x_1 \\ 1 & x_2 \\ 1 & x_3 \end{pmatrix} \\
 \mathbf{D} &= \begin{pmatrix} d_{1,1} & d_{1,2} & d_{1,3} \\ d_{2,1} & d_{2,2} & d_{2,3} \\ d_{3,1} & d_{3,2} & d_{3,3} \end{pmatrix}
 \end{aligned}
 \quad
 \begin{aligned}
 \boldsymbol{\Sigma} &= \begin{pmatrix} \sigma^2 & \sigma^2 e^{-\frac{d_{1,2}}{\phi}} & \sigma^2 e^{-\frac{d_{1,3}}{\phi}} \\ \sigma^2 e^{-\frac{d_{2,1}}{\phi}} & \sigma^2 & \sigma^2 e^{-\frac{d_{2,3}}{\phi}} \\ \sigma^2 e^{-\frac{d_{3,1}}{\phi}} & \sigma^2 e^{-\frac{d_{3,2}}{\phi}} & \sigma^2 \end{pmatrix} \\
 \sigma_\varepsilon^2 \mathbf{I} &= \begin{pmatrix} \sigma_\varepsilon^2 & 0 & 0 \\ 0 & \sigma_\varepsilon^2 & 0 \\ 0 & 0 & \sigma_\varepsilon^2 \end{pmatrix} \\
 \boldsymbol{\Sigma} + \sigma_\varepsilon^2 \mathbf{I} &= \begin{pmatrix} \sigma^2 + \sigma_\varepsilon^2 & \sigma^2 e^{-\frac{d_{1,2}}{\phi}} & \sigma^2 e^{-\frac{d_{1,3}}{\phi}} \\ \sigma^2 e^{-\frac{d_{2,1}}{\phi}} & \sigma^2 + \sigma_\varepsilon^2 & \sigma^2 e^{-\frac{d_{2,3}}{\phi}} \\ \sigma^2 e^{-\frac{d_{3,1}}{\phi}} & \sigma^2 e^{-\frac{d_{3,2}}{\phi}} & \sigma^2 + \sigma_\varepsilon^2 \end{pmatrix}
 \end{aligned}$$

$$\begin{aligned}
 \mathbf{y} &\sim \text{multivariate normal}(\mathbf{X}\boldsymbol{\beta}, \boldsymbol{\Sigma} + \sigma_\varepsilon^2 \mathbf{I}) \\
 [\boldsymbol{\beta}, \sigma^2, \sigma_\varepsilon^2, \phi \mid \mathbf{y}] &\propto [\mathbf{y} \mid \mathbf{X}\boldsymbol{\beta}, \boldsymbol{\Sigma} + \sigma_\varepsilon^2 \mathbf{I}] \\
 &\times [\boldsymbol{\beta}][\sigma^2][\sigma_\varepsilon^2][\phi]
 \end{aligned}$$

Priors on ϕ

Choices for range parameter ϕ :

- ▶ $\phi \sim \text{gamma}(\gamma_1, \gamma_2)$
- ▶ $\log(\phi) \sim \text{normal}(\mu_\phi, \sigma_\phi^2)$
- ▶ $\phi \sim \text{Half-Cauchy}(\gamma)$

General spatial models

- ▶ Real valued, non-negative

$$g(\boldsymbol{\beta}, \mathbf{X}) = \exp(\mathbf{X}\boldsymbol{\beta})$$

$$\log(\mathbf{y}) \sim \text{multivariate normal}(\log(g(\boldsymbol{\beta}, \mathbf{X})), \boldsymbol{\Sigma} + \sigma_{\varepsilon}^2 \mathbf{I})$$

- ▶ Counts

$$g(\boldsymbol{\beta}, \mathbf{X}) = \exp(\mathbf{X}\boldsymbol{\beta})$$

$$\log(\boldsymbol{\lambda}) \sim \text{multivariate normal}(\log(g(\boldsymbol{\beta}, \mathbf{X})), \boldsymbol{\Sigma} + \sigma_{\varepsilon}^2 \mathbf{I})$$

$$y_i \sim \text{Poisson}(\lambda_i)$$

- ▶ Binary

$$g(\boldsymbol{\beta}, \mathbf{X}) = \text{logit}^{-1}(\mathbf{X}\boldsymbol{\beta})$$

$$\text{logit}(\mathbf{p}) \sim \text{multivariate normal}(\text{logit}(g(\boldsymbol{\beta}, \mathbf{X})), \boldsymbol{\Sigma} + \sigma_{\varepsilon}^2 \mathbf{I})$$

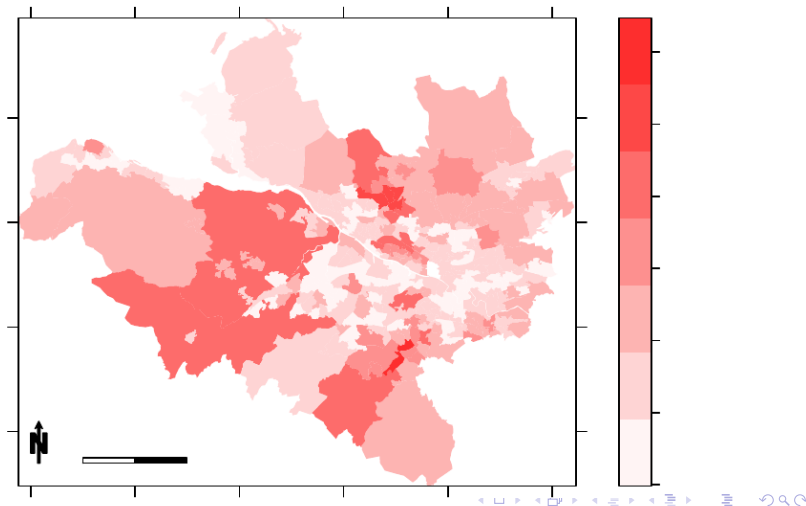
$$y_i \sim \text{Bernoulli}(p_i)$$

Simulating data for a continuous spatial process

1. Choose locations \mathbf{s}_i for $i = 1, \dots, n$.
2. Choose the mean $\boldsymbol{\mu}$. This could be a scalar or it could vary spatially. It could be the output of a model with parameter values that you choose and \mathbf{x} data.
3. Choose the unstructured variance σ_ε^2 .
4. Choose range parameter ϕ and variance component σ^2 .
5. Compute distance matrix \mathbf{D} between all n locations of interest.
6. Calculate covariance matrix $\boldsymbol{\Sigma} = \sigma^2 \exp\left(-\frac{\mathbf{D}}{\phi}\right)$.
7. Sample the n -dimensional vector $\mathbf{y} \sim \text{multivariate normal}(\boldsymbol{\mu}, \boldsymbol{\Sigma} + \sigma_\varepsilon^2 \mathbf{I})$.

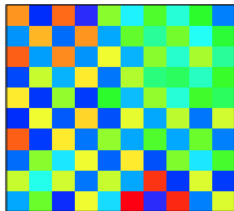
Most ecological data are spatial

Areal spatial processes

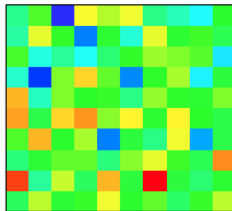


Regular, random, or clustered?

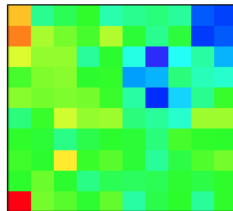
Regular



Random



Clustered



Measures of regularity, clustering

- ▶ Moran's I: similar to covariogram.

$$I = \frac{n}{(n-1)s^2w_{..}} \sum_{i=1}^n \sum_{j=1}^n w_{i,j} (y(a_i) - \bar{y})(y(a_j) - \bar{y})$$

- ▶ Geary's C: similar to variogram (or Durbin-Watson statistic in time series).

$$C = \frac{1}{2s^2w_{..}} \sum_{i=1}^n \sum_{j=1}^n w_{i,j} (y(a_i) - y(a_j))^2$$

s^2 = sample variance

Testing for regularity, clustering

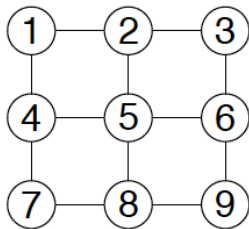
- ▶ Moran's I:

- ▶ $E(I) = -\frac{1}{n-1}$
- ▶ $I > E(I)$ implies clustering.
- ▶ $I < E(I)$ implies regularity.

- ▶ Geary's C:

- ▶ $E(C) = 1$
- ▶ $C > 1$ implies negative autocorrelation (regularity).
- ▶ $C < 1$ implies positive autocorrelation (clustering).
- ▶ $0 < C < 2$.

Areal data and proximity



$$\mathbf{W} = \begin{pmatrix} 0 & 1 & 0 & 1 & 0 & 0 & 0 & 0 & 0 \\ 1 & 0 & 1 & 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 0 & 1 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 & 1 & 0 & 1 & 0 & 0 \\ 0 & 1 & 0 & 1 & 0 & 1 & 0 & 1 & 0 \\ 0 & 0 & 1 & 0 & 1 & 0 & 0 & 0 & 1 \\ 0 & 0 & 0 & 1 & 0 & 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 & 1 & 0 & 1 \\ 0 & 0 & 0 & 0 & 0 & 1 & 0 & 1 & 0 \end{pmatrix}$$

Choices for elements of W , the “weights”

Possibilities include, but are not limited to:

- ▶ $w_{ij} = 1$ if i, j share a common boundary (possibly a common vertex)
- ▶ w_{ij} is an inverse distance between units
- ▶ $w_{ij} = 1$ if distance between units is $\leq K$
- ▶ $w_{ij} = 1$ for m nearest neighbors.

Modeling areal data

$$y(a_i) = \mathbf{x}(a_i)' \boldsymbol{\beta} + \varepsilon(a_i)$$

- ▶ Similar to that for continuous spatial modeling, except:
 - ▶ Covariance is not parameterized in terms of distance.
 - ▶ Usually not stationary (variance changes spatially).
- ▶ We need a modeling framework that accounts for these issues.

Modeling areal data

Two general types of *spatial autoregressive models*:

- ▶ Conditional autoregressive models (CAR): The probability of values estimated at any given location are conditional on neighboring values. Today.
- ▶ Simultaneous autoregressive models (SAR): See J. M. Ver Hoef, E. E. Peterson, M. B. Hooten, E. M. Hanks, and M. J. Fortin. Spatial autoregressive models for statistical inference from ecological data. *Ecological Monographs*, 88(1):36–59, 2018.

Conditional autoregressive model

$$\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}$$

- ▶ where, $\boldsymbol{\varepsilon} \sim \text{normal}(\mathbf{0}, \sigma^2(\mathbf{I} - \rho\mathbf{W})^{-1})$.
- ▶ ρ is an autocorrelation parameter.
- ▶ Note: Proximity matrix \mathbf{W} must be symmetric.
- ▶ Derivation requires Hamersley-Clifford theorem, Brooks lemma

Conditional autoregressive model with row standardization

$$\mathbf{y} \sim \text{normal}(\mathbf{X}\boldsymbol{\beta}, \sigma^2\mathbf{R})$$

- ▶ $\boldsymbol{\Sigma} = \sigma^2\mathbf{R}$
- ▶ $\mathbf{R} = (\text{diag}(\mathbf{W}\mathbf{1}) - \rho\mathbf{W})^{-1}$
 - ▶ $\mathbf{1}$ is a column vector of 1's.
 - ▶ $\mathbf{W}\mathbf{1}$ is the sums of the rows
 - ▶ $\text{diag}(\mathbf{W}\mathbf{1})$ is a matrix with the sums of the rows on the diagonal and zeros elsewhere.
 - ▶ Row standardization assures that $|\rho| < 1$
 - ▶ Equivalent to dividing each element of \mathbf{W} by the sum of the rows to obtain \mathbf{W}_+ and using $\boldsymbol{\Sigma} = \sigma^2(\mathbf{I} - \rho\mathbf{W}_+)^{-1}$
- ▶ $\rho \sim \text{Beta}(18, 2)$ to favor values close to 1
- ▶ $\sigma^2 \sim \text{IG}(r, q)$.

CAR for non-negative observations

Let $\mathbf{\Sigma} = \sigma^2 \text{diag}(\mathbf{W}\mathbf{1} - \rho\mathbf{W})^{-1}$

$$g(\boldsymbol{\beta}, \mathbf{X}) = \exp(\mathbf{X}\boldsymbol{\beta}) \quad (7)$$

$$\log(\boldsymbol{\mu}) \sim \text{multivariate normal}(\log(g(\boldsymbol{\beta}, \mathbf{X})), \mathbf{\Sigma}) \quad (8)$$

CAR for counts

Let $\mathbf{\Sigma} = \sigma^2 \text{diag}(\mathbf{W}\mathbf{1} - \rho\mathbf{W})^{-1}$

$$g(\boldsymbol{\beta}, \mathbf{X}) = \exp(\mathbf{X}\boldsymbol{\beta}) \quad (9)$$

$$y_i \sim \text{Poisson}(\lambda_i) \quad (10)$$

$$\log(\boldsymbol{\lambda}) \sim \text{multivariate normal}(\log(g(\boldsymbol{\beta}, \mathbf{X})), \mathbf{\Sigma}) \quad (11)$$

CAR for binary observations

Let $\Sigma = \sigma^2 \text{diag}(\mathbf{W}\mathbf{1} - \rho\mathbf{W})^{-1}$

$$g(\boldsymbol{\beta}, \mathbf{X}) = \text{logit}^{-1}(\mathbf{X}\boldsymbol{\beta}) \quad (12)$$

$$y_i \sim \text{Bernoulli}(p_i) \quad (13)$$

$$\text{logit}(\mathbf{p}) \sim \text{multivariate normal}(\text{logit}(g(\boldsymbol{\beta}, \mathbf{X})), \boldsymbol{\Sigma}) \quad (14)$$

Take home

- ▶ Data taken over space (or time) are likely to be structured by physical and biological processes.
- ▶ Our deterministic model may account for this structure. However, if the *residuals* show correlation over time and/ or space, then we are obliged to model their covariance to assure that iid assumptions are met.
- ▶ Doing so requires estimating only a few more parameters, in most cases two, relative to the aspatial model.
- ▶ Deciding whether to use a spatial or aspatial model could be sensibly treated as a problem in model selection.

Where next?

- ▶ M. B. Hooten, D. S. Johnson, B. T. McClintock, and J. Morales. Animal Movement: Statistical Models for Telemetry Data. Chapman and Hall/CRC., 2017.
- ▶ M. Blangiardo and M. Camelletti. Spatial and spatio-temporal Bayesian models with R - INLA. Wiley, West Sussex U.K., 2015.
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- ▶ C. K. Wikle. Hierarchical Bayesian models for predicting the spread of ecological processes. Ecology, 84(6):1282–1304,