

Right Map, Right Model

Two examples using the BYM2 model

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Talk Overview

- The Besag York Mollié model: workhorse of spatial epidemiology.
- Computational Concerns:
 - Coding the BYM2 model.
 - Getting the map you want from the geo-coordinates you have.
- Example: accounting for time plus roads, rails, and airplanes.
- Example: accounting for disconnected regions.

Spatial Smoothing For Areal Data

- Counts of (rare) events in small-population regions are noisy
 - Borrow information from *neighboring* areas
- Besag, 1973, 1974: Conditional Auto-Regressive Model (CAR), and **Intrinsic Conditional Auto-Regressive (ICAR)** model
 - Gaussian Markov Random Field (GMRF)
- CAR model has parameter α for the amount of spatial dependence;
 - CAR model requires computing matrix determinants
 - Cubic operation (ON^3) (calculated at every *step* of the sampler)
- ICAR simplification: let $\alpha == 1$ (complete spatial dependence)
 - ICAR model computes pairwise distance between neighbors
 - Linear operation (ON)

Neighbor Relationship

- The binary neighbor relationship (written $i \sim j$ where $i \neq j$) is encoded as
 - 1 if regions n_i and n_j are neighbors
 - 0 otherwise
- For ICAR models, neighbor relationship is
 - symmetric - if $i \sim j$ then $j \sim i$
 - not reflexive - a region is not its own neighbor ($i \not\sim i$)
- Many possible definitions of “neighbor”
 - ‘rook’ - areas share a bounding line
 - ‘queen’ - areas share a boundary point
 - *mobility networks* - amount of travel between areas

Intrinsic Conditional Auto-Regressive (ICAR) Model

- Conditional specification: multivariate normal random vector ϕ , each ϕ_i is conditional on the values of its neighbors
- Joint specification rewrites to *Pairwise Difference*:

$$p(\phi) \propto \exp \left\{ -\frac{1}{2} \sum_{i \sim j} (\phi_i - \phi_j)^2 \right\}$$

- Centered at 0, assuming common variance for all elements of ϕ .
- Each $(\phi_i - \phi_j)^2$ penalizes the distance between the *values* of neighboring regions.
- ϕ is non-identifiable - adding a constant to ϕ washes out of $\phi_i - \phi_j$
- Sum-to-zero constraint centers ϕ

Besag York Mollié (1991) BYM Model

- Basic regression Lognormal Poisson regression plus 2 components for spatial smoothing
 - spatial effects: ICAR component
 - random effect: per-region standard Normal
- Formula: $\mu + x\beta + \phi + \theta$
 - μ is the fixed intercept.
 - x is the design matrix, β is vector of regression coefficients.
 - ϕ is an ICAR spatial component
 - θ is an vector of ordinary random-effects components.

BYM2 model: Riebler et al, 2016

- *Penalized Complexity* (Simpson et al, 2014)
 - **spatial and random effects must have** $\text{Var}(\phi_i) \approx 1$
 - combined spatial and random effects have mixing parameter, scale parameter, (following Leroux, 2000)

- BYM2 model:

$$\mu + x\beta + \left((\sqrt{\rho/s}) \phi^* + (\sqrt{1-\rho}) \theta^* \right) \sigma$$

where:

- $\sigma \geq 0$ is the overall standard deviation.
- $\rho \in [0, 1]$ - proportion of spatial variance.
- ϕ^* is the ICAR component.
- $\theta^* \sim N(0, 1)$ is the vector of ordinary random effects
- s is a scaling factor s.t. $\text{Var}(\phi_i) \approx 1$, computed from neighbor graph, i.e. s is *data*, not a parameter.

Coding Challenge: from Math to Model

$$\mu + x\beta + \left((\sqrt{\rho/s}) \phi^* + (\sqrt{1-\rho}) \theta^* \right) \sigma$$

```
parameters {  
  real beta0; // intercept  
  vector[K] betas; // covariates  
  real<lower=0, upper=1> rho; // proportion of spatial variance  
  sum_to_zero_vector[N] phi; // spatial effects  
  vector[N] theta; // heterogeneous random effects  
  real<lower = 0> sigma; // scale of combined effects  
}  
model {  
  vector[N] gamma = sqrt(rho / tau) * phi + sqrt(1 - rho) * theta;  
  y ~ poisson_log(log_E + beta0 + xs_centered * betas + gamma * sigma);  
  target += -0.5 * dot_self(phi[neighbors[1]] - phi[neighbors[2]]); // ICAR prior  
  beta0 ~ std_normal();  betas ~ std_normal();  
  rho ~ beta(0.5, 0.5);  theta ~ std_normal();  sigma ~ std_normal();  
}
```

Stan ICAR Model

Encode neighbor information as graph edgeset, i.e. pairs of indices for neighbors i, j :

```
data {  
  int<lower = 0> N;    // number of areal regions  
  int<lower = 0> N_edges; // number of neighbor pairs  
  array[2, N_edges] int<lower = 1, upper = N> neighbors; // columnwise adjacent  
  ...  
}
```

Use Stan's `sum_to_zero_vector` to identify ϕ

```
parameters {  
  sum_to_zero_vector[N] phi; // spatial effects  
  ...  
}
```

Use Stan's vectorized operations and multi-indexing to compute ICAR prior

```
model {  
  target += -0.5 * dot_self(phi[neighbors[1]] - phi[neighbors[2]]);  
  ...  
}
```

Stan Constrained Parameter `sum_to_zero_vector`

```
sum_to_zero_vector[K] beta;
```

- On the unconstrained scale, beta is length $K - 1$ (because the last is determined by the first $K - 1$).
 - Stan computes on the *unconstrained* scale.
 - Constraining transform keeps variance of all elements equal.
- Outperforms “hard” and “soft” sum-to-zero constraints
 - “hard” sum-to-zero: N^{th} element == - sum(elements 1 : N-1)
 - “soft” sum-to-zero: `constrain sum(beta) ~ Normal(0, epsilon)` `epsilon = 0.001`
- The larger, more complex the model, the bigger the difference.
- See Stan case study [The Sum-to-Zero Constraint in Stan](#)

From Geospatial Maps to Neighbor Graphs

- GIS data: points, lines, and bounding polygons
 - Common format: [shapefile](#)
 - Python: [GeoPandas](#) - add support for geographic data to pandas objects.
 - R: [sf](#) Simple Features for R - “spatial analysis simplified”
- Neighbor graph
 - Graph nodes denote regions, edges denote neighbors
 - [GDAL](#)-based utilities compute neighbor graph from geo-dataframe geometry column.
- Shapefiles may not line up with your data.
 - Shapefile region IDs may differ from dataset IDs.
 - Map boundaries may differ from idealized boundaries.
- Neighbor graphs are difficult to edit.
 - Easy to get node indices wrong.
 - Difficult to maintain symmetry

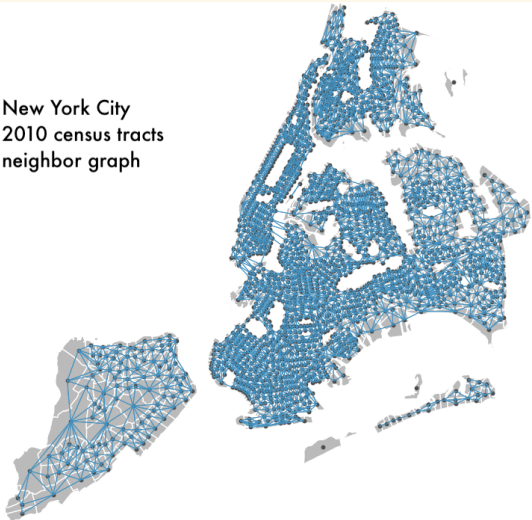
Example Dataset: NYC traffic accidents

Dataset taken from [Bayesian Hierarchical Spatial Models: Implementing the Besag York Mollié Model in Stan](#).

- Aggregated counts of car vs. child pedestrian traffic accidents, localized to US Census tract. Per-tract data includes:
 - raw counts of accidents, population
 - measures of foot traffic, car traffic
 - socio-economic indicators: median income, neighborhood transiency
- Starting point: [NYC Planning Census Blocks](#)
 - " These boundary files are derived from the US Census Bureau's TIGER data products and have been geographically modified to fit the New York City base map."

NYC Census Blocks

New York City
2010 census tracts
neighbor graph



- Study focus is on *pedestrians*
- Map doesn't respect geography
 - East River separates Manhattan from Brooklyn, Queens
- Geography doesn't respect ICAR model
 - Graph must be fully connected - else elements of ϕ are undefined.

Ward et al Mobility Network Graph

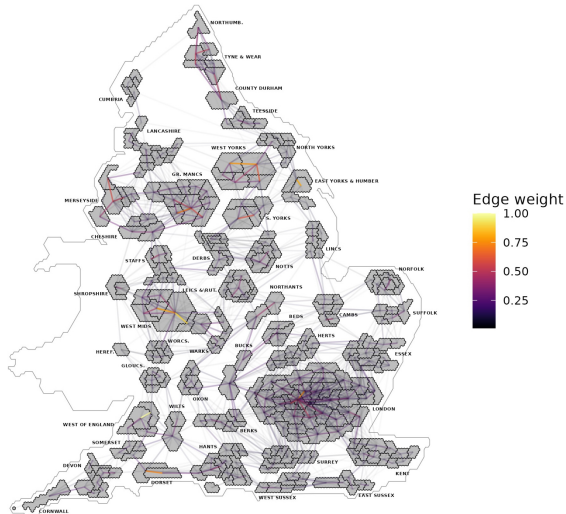


Fig 1. The mobility network overlaid on a hex map of England, where the edge weight describes the rate of journeys between different LTLA's, normalised to take values in $[0,1]$. Created from geographical files using the House of Commons Library, which is under the Open Parliament License v3.0.

- Online: [Fig 1](#)
- Northern england shows divide between east and west coasts

BYM2 Model for Spatio-Temporal Smoothing

PLOS COMPUTATIONAL BIOLOGY

 OPEN ACCESS  PEER-REVIEWED

RESEARCH ARTICLE

Bayesian spatial modelling of localised SARS-CoV-2 transmission through mobility networks across England

Thomas Ward , Mitzi Morris, Andrew Gelman, Bob Carpenter, William Ferguson, Christopher Overton, Martyn Fyles

Version 2

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- Neighbor graph reflects *mobility networks*
- Extend the BYM2 model to include a daily-level random effect, i.e. three components:
 - Spatial effect θ_i
 - Ordinary per-region random effect ϕ_i
 - Ordinary per-region, per_day random effect $\phi_{i,t}$
 - Proportion of variance ρ is a simplex

Ward et al Simulation Study

Bayesian spatial modelling of localised SARS-CoV-2 transmission through mobility networks across England

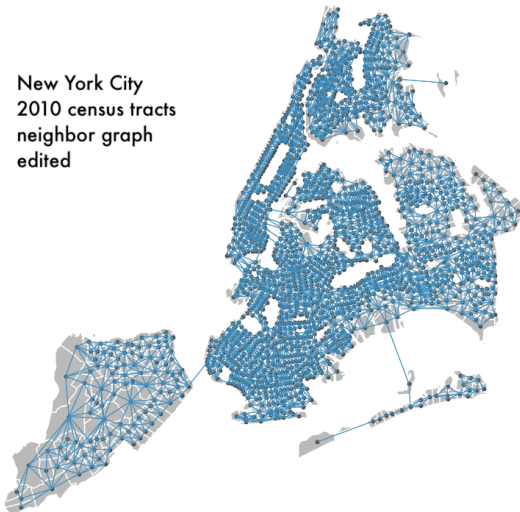
Table 3

The average error for the naïve estimate (%), BYM2 (%), and total reduction in error (%) for the simulation scenarios.

Scenario	Average Error for Naïve Estimate (%)	Average Error for BYM2 Estimate (%)	Reduction in Error (%)
1	7.41	2.42	67.3
1.1	15.11	3.48	77.0
1.2	10.04	5.81	42.1
1.3	16.76	6.30	62.4
2	5.79	3.63	37.3
2.1	11.03	5.20	52.9
2.2	8.03	5.53	31.1
2.3	12.73	6.46	49.2
3	2.66	2.00	24.6
3.1	5.50	3.21	41.6
3.2	3.82	2.90	24.2
3.3	5.45	3.41	37.4

2017 BYM2 Case Study - Fully Connected Map

New York City
2010 census tracts
neighbor graph
edited



- Map editing process
 - Identify pairs of tracts to connect
 - Write utility functions to munge data structures
- Extremely tedious and error prone
- Must be scripted for reproducibility

BYM2 Multicomp Model

- Make the model respect the geography
- Extends the BYM2 model to account for disconnected graphs and islands, following the recommendations from [A note on intrinsic Conditional Autoregressive models for disconnected graphs](#), Freni-Sterrantino et.al. 2018.
 - Component nodes are given the BYM2 prior
 - Singleton nodes (islands) are given a standard Normal prior
 - Compute per-connected component scaling factor
 - Impose a sum-to-zero constraint on each connected component
- See R and Python notebooks from GeoMED 2024 Workshop
 - GitHub Repository: [mitzimorris/GeoMED_2024](#)
 - Notebook “h6_bym2_multicomp”

Scaling factor, following Riebler

- Geometric mean of the variances of the spatial covariance matrix
- Neighborhood structure matrix is the *precision* matrix
- Expensive to compute, but as data, is only done once

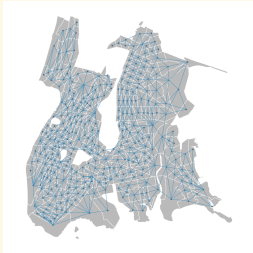
```
get_scaling_factor = function(nbs) {  
  # Create ICAR precision matrix  
  N = length(nbs)  
  adj_matrix = nb2mat(nbs, style="B")  
  Q = Diagonal(N, rowSums(adj_matrix)) - adj_matrix  
  
  # Get covariance matrix, compute the geometric mean of diagonal  
  Q_pert = Q + Diagonal(N) * max(diag(Q)) * sqrt(.Machine$double.eps)  
  Q_inv = q_inv_dense(Q_pert, adj_matrix)  
  return(exp(mean(log(diag(Q_inv)))))  
}
```

Different Maps Have Difference Scaling Factors

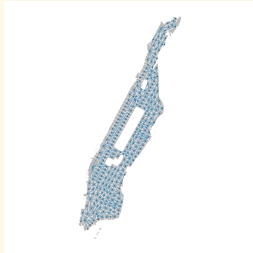
Brklyn-Queens: 0.71



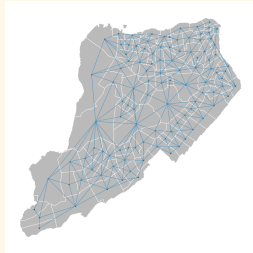
Bronx: 0.57



Manhattan: 0.8



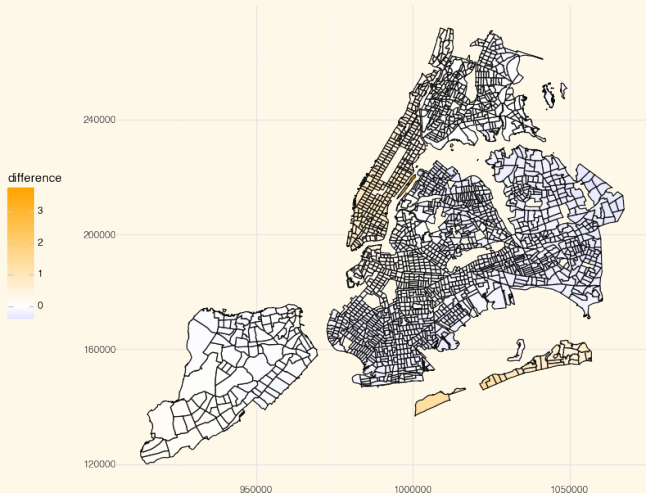
Staten Island: 0.36



Fully connected graph has scaling factor of 1.3

Comparison of ICAR component estimates

Changes in spatial component ϕ
BYM2_multicomp_fit - BYM2_1comp_fit



- Systematic differences due to per-component scaling factors
- Minor differences from removing spurious connections

References, Thanks!, and Questions?

- Simpson et al., 2014: Penalising model component complexity: A principled, practical approach to constructing priors
- Riebler et al., 2016: An intuitive Bayesian spatial model for disease mapping that accounts for scaling
- Freni-Sterrantino et al., 2018: A note on intrinsic conditional autoregressive models for disconnected graphs
- Morris et al., 2019: Bayesian Hierarchical Spatial Models: Implementing the Besag York Mollié Model in Stan
- Ward et. al., 2023 Bayesian spatial modelling of localised SARS-CoV-2 transmission through mobility networks across England