Probabilistic Modeling of Spreading / Contracting Point Pattern using Hierarchical Bayesian Diffusion Equation

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Dec 2, 2022

Introduction and Project Goal

Goal

- Spatio-temporal modeling of the spread (= diffusion) of disease
- Practically, extrapolate the spatial point pattern of the next time period.
- Main interest : diffusion pattern
 - How fast the disease diffuses to the population
 - Whether the number of infected patients tend to increase or decrease

Possible Approaches

- Direct modeling by spatio-temporal point process : contagion ≈ parent points generating offspring points
 - Need a separate point process from the point processes learnt in class mainly focused on attraction / repulsion
 - Main goal of point process model is Exploratory Data Analysis (EDA).
- Binary regression model: discretize the domain and find if the virus has spreaded in that region using spatial information.
 - Apply Hefley et al. (2017): hierarchical Bayesian regression model containing ecological diffusion model
 - Original application: Chronic Wasting Disease (CWD) in cervids in Wisconsin.



Model Illustration

- $\bullet \ Y_i, i=1,...,n \ : \ \text{Bernoulli response of the} \ i^{th} \ \text{observation at location} \ \boldsymbol{s_i} = (s_{1i},s_{2i}) \in \Omega = [a,b] \times [c,d] \ \text{and time} \ t_i.$
- ullet $p_i:=Pr(Y_i=1)$, relies on spatio-temporal effect $u(s_i,t_i)$ and the individual covariate information x_i .

Data Model :
$$Y_i|p_i \sim Ber(p_i)$$
 ; $p_i = g^{-1}(u(\boldsymbol{s_i}, t_i)e^{\boldsymbol{x_i'}\beta})$ (1)

$$\text{Diffusion Model}: \ \frac{\partial}{\partial t}u(\boldsymbol{s},t) = \Big(\frac{\partial^2}{\partial s_1^2} + \frac{\partial^2}{\partial s_2^2}\Big)[\mu(\boldsymbol{s})u(\boldsymbol{s},t)] + \lambda(\boldsymbol{s})u(\boldsymbol{s},t) \ \ \textbf{(2)}$$

$$log(\mu(s)) = z(s)^T \alpha ; \lambda(s) = w(s)^T \gamma$$
 (3)

Prior Model :
$$\beta \sim MVN(\mathbf{0}, 10^6 \mathbf{I})$$
 ; $\alpha \sim MVN(\mathbf{0}, 10^6 \mathbf{I})$; $\gamma \sim MVN(\mathbf{0}, 10^6 \mathbf{I})$; $\phi \sim TN(0, 10^6)$; $\theta \sim TN(0, 10^6)$ (4)

Boundary Condition :
$$u(s,t) = 0, \forall t > 0$$
 (5)

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Initial Condition :
$$u(s,0) = \frac{\theta e^{\frac{-|s-a|}{\phi^2}}}{\int_{s} e^{\frac{-|s-d|}{\phi^2}}}$$
 (6)

- Link function $g:[0,\infty)\to [0,1]$. Use $g^{-1}(x)=\sqrt{\frac{2}{\pi}}\int_0^x e^{x^2/2}dx$
- spatio-temporal effect $u(s_i, t_i)$ is modeled by the diffusion equation in (2).
 - ullet $\mu(m{s})>0$ is the diffusion rate of location $m{s}$ and $\lambda(m{s})$ is the growth rate of location $m{s}$
 - $oldsymbol{v}(s)$ and w(s) are vectors containing spatial covariates (e.g, forests and population).



- Ecological diffusion widely used to model the temporal movements of transient spatial processes
- Interpretable model : diffusion rate, growth rate, ...
- Bayesian hierarchical regression model : easy uncertainty quantification

Blocked Gibbs Sampler with Metropolis-Hastings within Gibbs

- $oldsymbol{\Theta} := (oldsymbol{lpha}, oldsymbol{\gamma}, oldsymbol{eta}, \phi, heta)$: the parameter of interest.
- Sample from **joint posterior** $p(\Theta|y)$ from the following blocked Gibbs sampler.
- ullet Since Y_i is Bernoulli and priors for each of Θ : MVN or TN, no semi-conjugacy o Metropolis-Hastings within Gibbs

Step 1 : Set initial values for Θ

Step 2 : For $i < n_{mcmc}$, iterate the following:

Step 3 : Update u(s,t) and sample (θ,ϕ) jointly from $p(\theta,\phi|\alpha,\beta,\gamma)$ using random walk Metropolis-Hastings

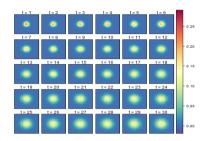
Step 4 : Update u(s,t) and sample α from $p(\alpha|\theta,\phi,\beta,\gamma)$ using random walk Metropolis-Hastings

Step 5 : Update u(s,t) and sample $oldsymbol{eta}$ from $p(oldsymbol{eta}| heta,\phi,oldsymbol{lpha},oldsymbol{\gamma})$ using random walk Metropolis-Hastings

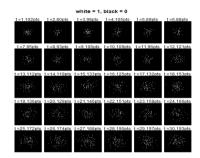
Step 6 : Update u(s,t) and sample γ from $p(\gamma|\theta,\phi,\alpha,\beta)$ using random walk Metropolis-Hastings

Solving the PDE

Simulation Study



(a) True probability of binary event



(b) True binary event