

# . Introduction

Fire plays an important role in Canada's forested ecosystems. It helps to maintain forest health and diversity; however, it can have undesirable negative effects on public safety, health and property. Forest fire has numerous causes such as dry weather and human behavior. Moreover, large areas of dead forest due to mountain pine beetle outbreak in British Columbia may lead to more severe wildfires. Therefore, it is important to study the distribution of forest fire and its relationship to these factors.

# 2. Data Structure

Study Area: British Columbia (divided into I = 1712 homogeneous grid cells)

Response:  $N_i$ , total fire counts in each region over 44 years Covariates (regional specific): area affected by mountain pine beetle (MPB) outbreak, area of forest covering, area of pine leading stands, number of roadways and drought climate Spatial Information: an adjacency matrix W, coding adjacencies of partitioning grid cells

# 3. Exploratory Analysis



#### Figure 4: Deviance of Res





• Figure 1-3: spatial distribution of response and covariates

(larger circle indicates a higher value)

- Figure 4: deviance of residuals under the standard log-linear Poisson regression model  $\rightarrow$  not randomly distributed  $\rightarrow$  data are spatially correlated
- Overdispersed count data (Alexander *et al.*, 2000)

# **Modeling Fire Frequency with Negative Binomial Spatial Regression Models** YOLANDA LI<sup>\*</sup> (VICTORIA) STEVE TAYLOR (PACIFIC FORESTRY) AND FAROUK NATHOO<sup>\*</sup> (VICTORIA) YOLANDA LI\* (VICTORIA), STEVE TAYLOR (PACIFIC FORESTRY), AND FAROUK NATHOO\* (VICTORIA) \* Department of Mathematics and Statistics, University of Victoria, Victoria, B.C.

# 4. Overdispersed Spatial Count Model

### $N_i \mid \lambda_i, a \sim \operatorname{Negbin}(\lambda_i, a) \qquad i = 1, ..., I$

where  $N_i$  is the total fire count in region i,  $\lambda_i$  describes the mean and  $a \ge 0$  is the dispersion parameter  $(a \rightarrow 0 \text{ yields the Poisson}(\lambda_i))$  (Lawless, 1987).

Under a Bayesian hierarchical framework, we can use an equivalent Poisson-Gamma mixture representation:

- $N_i \mid \nu_i, \lambda_i \sim \text{Poisson}(\nu_i \lambda_i)$  $\log(\lambda_i) = \boldsymbol{\beta}^T \mathbf{X}_i + b_i$  $\mathbf{b} = (b_1, ..., b_n)$  are spatial random effects,  $\mathbf{b} \mid \sigma_{\mathbf{b}}^2 \sim \text{CAR}(\sigma_{\mathbf{b}}^2)$
- $\nu_i \mid a \stackrel{i.i.d}{\sim} \operatorname{Gamma}(\frac{1}{a}, a)$  accommodate extra Poisson variation

where  $\beta$  is a vector of regression coefficients and  $\mathbf{X}_i$  is a vector of covariates for region *i*. The conditional autoregressive model (CAR) (Besag, 1974) employed for  $\mathbf{b}$  accounts the spatial effect of region i conditionally on its neighboring regions based on the adjacency matrix W.

Identifiability: Note the random effects  $b_i$  and  $\nu_i$  are not uniquely identified; however the sum  $\alpha_i = b_i + \log(\nu_i)$  is identified. In order to deal with the weak identifiability issue among  $\nu_i$  and  $b_i$ , we monitor  $\alpha_i$  instead.

Model estimation of  $\theta = (\beta, \alpha, a, \sigma_b^2)$  given the data is carried out using a Markov Chain Monte Carlo Algorithms programed in Matlab.

### Posterior Summaries (NegBin with CAR)

	Mean	Std.	Deviation	2.50%	97.5%
$\beta_0$ (Intercept)	2.6879		0.0145	2.6587	2.7157
$eta_1~(\mathrm{MPB})$ -	-0.0335		0.0553	-0.1446	0.0746
$\beta_2$ (DCmean)	0.8457		0.0811	0.6941	1.0119
$\beta_3 \text{ (forest)}$	0.8316		0.0377	0.7577	0.9058
$eta_4~({ m pine})$ -	-0.3491		0.0418	-0.4313	-0.2655
$\beta_5 \pmod{2}$	0.4220		0.0425	0.3376	0.5058
$\tau = 1/\sigma_{\mathbf{b}}^2$ (CAR precision)	0.3221		0.0143	0.2949	0.3508
a (overdispersion)	0.0002		0.0010	0.0000	0.0020

### $p_D$ and DIC

model  $Poisson(\lambda_i), log(\lambda_i) = \beta^T X_i$ Poisson $(\lambda_i), log(\lambda_i) = \boldsymbol{\beta}^T X_i + b_i, b_i \overset{i.i.d}{\sim} \mathbb{N}$ Poisson $(\lambda_i), log(\lambda_i) = \boldsymbol{\beta}^T X_i + b_i, \mathbf{b} \sim CA$ NegBin $(\lambda_i, a), log(\lambda_i) = \boldsymbol{\beta}^T X_i$ NegBin $(\lambda_i, a), log(\lambda_i) = \boldsymbol{\beta}^T X_i + b_i, \mathbf{b} \sim CA$ 

Comment: The Negative Binomial model with spatial random effects is preferred since this model has the smallest DIC.

	$p_D$	DIC
	6	95292
$\mathbb{N}(0,1/ au)$	1476	10849
$\mathrm{AR}(1/ au)$	1315	10697
	1403	10710
$AR(1/\tau)$	1295	10678

Posterior Predictive Checking: (Gelman et al., 2004)

- predictive distribution
- Posterior Predictive P-value is defined as

where T(.) is a test quantity that is a scalar summary of parameters and data



# Heavy Zeros:

 $T = \sum_{i=1}^{I} I(N_{ij}^{rep} = 0)$ 

I(.) is the indication function



Comment: No evidence of lack of fit for extremes and overdispersion aspects; however, not adequately detail with zeros.

# 6. Future Work

• Use zero heavy Poisson Mixture (Hougaard et al., 1997) to deal with the heavy tail property of the data.  $f^*(y \mid \lambda, \rho) = \rho f(y \mid \lambda) + (1 - \rho)I\{y = 0\}$  where  $\rho \in [0, 1], f(y \mid \lambda)$  is the Poisson pmf

• Move towards both spatial and temporal modeling. References:

- the negative binomial distribution. *Biostatistics*  $\mathbf{1}$ , 453-463.
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5. Goodness of Fit

• J=4000 replicated data sets,  $N^{rep}$ , generated from posterior

 $P_B = \Pr(T(\mathbf{N}^{rep}, \boldsymbol{\theta}) \ge T(\mathbf{N}, \boldsymbol{\theta}) \mid \mathbf{N})$ 

Extremes: T =the largest value of fire counts in each data set

°<sub>R</sub>=0.003

Overdispersion:  $T = \frac{\operatorname{mean}(\mathbf{N}_{j}^{rep})}{\operatorname{variance}(\mathbf{N}_{j}^{rep})}$ 

number of fire count = 0

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• Hougaard, P., Lee, T. M. and Whitmore, A. G. (1997). Analysis of overdispersed count data by mixtures of