

Modelling multivariate counts varying continuously in space

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Outline of the talk

- ▶ Motivation and aims
- ▶ Distributions for multivariate counts: a brief review
- ▶ Our proposed model
- ▶ Covariates in covariance structure
- ▶ Model comparison and results
- ▶ Discussion

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

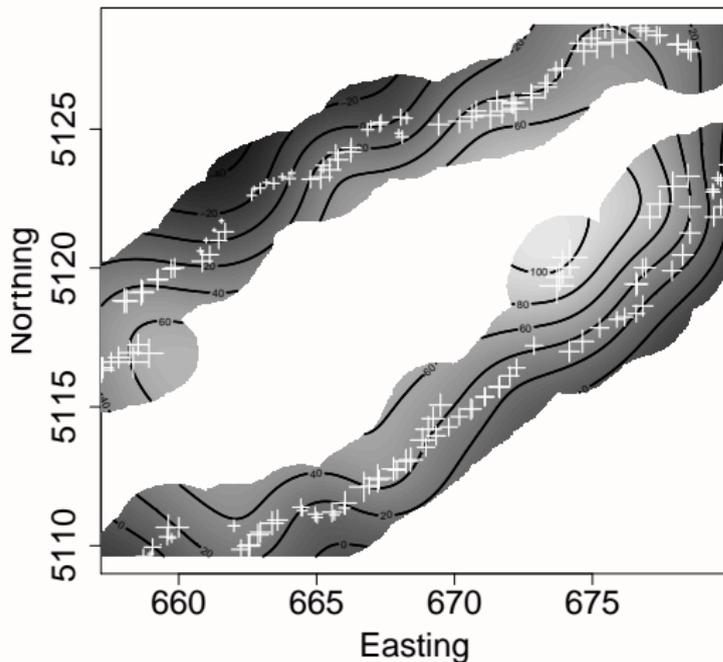
Inference procedure

Results

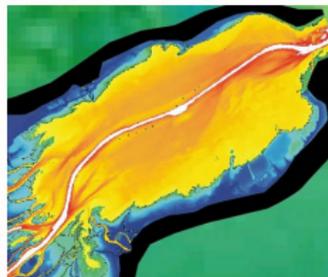
Model comparison
Results

Conclusions

Main references



Sampling locations in Lake St. Pierre (+) and geodetic depth contours (curves)



Central navigation channel (deep; strong current)

- ▶ On each sampling date, measurements were made at a cluster of locations on a shore, selected randomly among all clusters on that shore
- ▶ Sampling dates were unevenly spaced in time over a period of 70 days; the North and South shores were visited in alternation on consecutive sampling dates

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Fish were collected by electrofishing: fish are attracted to anodes hanging from the booms



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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Most abundant fish species observed in Lake St. Pierre



(a) Yellow perch



(b) Brown bullhead



(c) Golden shiner



(d) Pumpkinseed

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Main aims

- ▶ Assess the influence of local habitat (characterized by environmental covariates [water depth](#), [water transparency](#), [substrate](#), [vegetation](#)) on the abundance of fish species
- ▶ Determine whether species abundances are correlated across space and among themselves
- ▶ Understand the spatial distribution of each species

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Distributions for multivariate counts

Probability distributions for multivariate counts can be defined through:

the sum of independent Poisson distributions → **does not account** for overdispersion or negative covariance;

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Distributions for multivariate counts

Probability distributions for multivariate counts can be defined through:

continuous mixtures of independent Poisson distributions. Let $\mathbf{Y} = (Y_1, \dots, Y_K)$, with $Y_k | \delta_k \sim Poi(\delta_k)$, $k = 1, \dots, K$, conditionally independent, then

$$f(\mathbf{y} | \theta) = \int \prod_{k=1}^K p(y_k | \delta_k) g(\delta | \theta) d\delta,$$

where $\delta = (\delta_1, \dots, \delta_K)$.

- ▶ $g(\cdot)$ can be the pdf of a multivariate gamma \rightarrow **accounts** for overdispersion **but not** for negative covariance
- ▶ $g(\cdot)$ can be the pdf of a multivariate log-normal distribution

Multivariate Poisson as a Sum of Independent Poisson Random Variables

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Let

$$Y_1 = W_1 + W_{12} + W_{13} + W_{123}$$

$$Y_2 = W_2 + W_{12} + W_{23} + W_{123}$$

$$Y_3 = W_3 + W_{23} + W_{13} + W_{123},$$

with $W_i \sim Poi(\lambda_i)$, $W_{ij} \sim Poi(\lambda_{ij})$, $W_{ijl} \sim Poi(\lambda_{ijl})$,
 $i, j, l = \{1, 2, 3\}$, $i < j < l$.

More generally, $\mathbf{Y} = \mathbf{B}\mathbf{W}$, such that, $E(\mathbf{Y}) = \mathbf{B}\boldsymbol{\lambda}$ and
 $V(\mathbf{Y}) = \mathbf{B}\boldsymbol{\Sigma}\mathbf{B}^T$, where $\boldsymbol{\Sigma} = \text{diag}(\lambda_1, \dots, \lambda_q)$.

- ▶ Mean and variance of Y_i are assumed identical.
- ▶ Model only accounts for positive covariance structures.

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Multivariate Count Distributions Based on Mixtures

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Let $\mathbf{Y} = (Y_1, \dots, Y_K)$, with $Y_k \mid \delta_k \sim Poi(\delta_k)$,
 $k = 1, \dots, K$, conditionally independent, then

$$f(\mathbf{y} \mid \theta) = \int \prod_{k=1}^K p(y_k \mid \delta_k) g(\delta \mid \theta) d\delta,$$

where $\delta = (\delta_1, \dots, \delta_K)$.

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Multivariate Poisson-Gamma Mixture

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A multivariate gamma distribution for a K -dimensional random vector δ can be obtained by defining

$$\delta = \mathbf{B} \mathbf{W} = \begin{pmatrix} \frac{b_0}{b_1} & 1 & 0 & \cdots & 0 \\ \frac{b_0}{b_2} & 0 & 1 & \cdots & 0 \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ \frac{b_0}{b_K} & 0 & 0 & \cdots & 1 \end{pmatrix} \begin{pmatrix} W_0 \\ W_1 \\ \vdots \\ W_K \end{pmatrix}, \quad (1)$$

with $W_k \sim Ga(a_k, b_k)$ for $k = 0, 1, 2, \dots, K$; W_k independent among themselves (Mathai & Moschopoulos, 1991).

It is easy to show that $\text{Cov}(\delta_k, \delta_l) = \frac{a_0}{b_k b_l}$,
 $k, l = 1, 2, \dots, K$.

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

If $\mathbf{Y} = (Y_1, \dots, Y_K)$, such that $Y_i | \delta_i \sim Poi(\delta_i)$, independently, and $\delta = \mathbf{BW}$ it can be shown that, marginally,

$$E(Y_i) = \frac{a_i}{b_i} + \frac{a_0}{b_i} = \mu_i \quad i, j = 1, \dots, K$$

$$V(Y_i) = \mu_i + \frac{a_0}{b_i^2} + \frac{a_i}{b_i}$$

$$\text{Cov}(Y_i, Y_j) = \frac{a_0}{b_i b_j}$$

- ▶ It accounts for overdispersion,
- ▶ however, it only captures positive covariance structures.

[Outline](#)[Motivation and aims](#)[Distr. multiv. counts](#)[Proposed model](#)[Inference procedure](#)[Results](#)Model comparison
Results[Conclusions](#)[Main references](#)

Multivariate Poisson-lognormal mixture (Aitchison & Ho 1989)

Assume that

$$\log \boldsymbol{\delta} \sim N_K(\boldsymbol{\mu}, \boldsymbol{\Sigma}),$$

where $\boldsymbol{\Sigma}$ has elements σ_{kl} . Assuming $Y_k \mid \delta_k \sim Poi(\delta_k)$ independently.

Marginally,

$$E(Y_k) = \exp\left(\mu_k + \frac{1}{2}\sigma_{kk}\right) = \alpha_k$$

$$V(Y_k) = \alpha_k + \alpha_k^2\{\exp(\sigma_{kk}) - 1\}$$

$$\text{Cov}(Y_k, Y_l) = \alpha_k \alpha_l \{\exp(\sigma_{kl}) - 1\}, \quad k, l = 1, \dots, K.$$

- ▶ This model **accounts** for overdispersion
- ▶ Allows for **both** positive and negative covariance structures

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

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$$\log \boldsymbol{\delta} \sim N_K(\boldsymbol{\mu}, \boldsymbol{\Sigma}),$$

where $\boldsymbol{\Sigma}$ has elements σ_{kl} . Assuming $Y_k \mid \delta_k \sim Poi(\delta_k)$ independently.

This approach is appealing, as we can write

$$\begin{aligned} \log \delta_k &= \beta X_k + \epsilon_k, \quad k = 1, \dots, K \\ \epsilon &\sim N_K(0, \boldsymbol{\Sigma}) \end{aligned}$$

- ▶ Chib and Winkelmann (2001) were the first to provide a full Bayesian treatment of this model
- ▶ A similar approach has been widely used for multiple disease mapping (e.g., Carlin and Banerjee 2003, Gelfand and Vounatsou 2003, Jin et al. 2007; good overview in Lawson 2009)

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Proposed model

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Let $Y_k(s_{t_j})$ represent the number of individuals (counts) of species k , $k = 1, \dots, K$, observed at location s_{t_j} , $j = 1, \dots, n_t$ and time $t = 1, \dots, T$. We assume

$$Y_k(s_{t_j}) \mid \theta_k(s_{t_j}), \delta_k(s_{t_j}) \sim Poi(\theta_k(s_{t_j})\delta_k(s_{t_j})),$$

where

$$\log \theta_k(s_{t_j}) = \mathbf{X}_k(s_{t_j})\boldsymbol{\beta}_k$$

which captures local habitat structures.

The parameter vector $\boldsymbol{\delta}(s_{t_j}) = (\delta_1(s_{t_j}), \dots, \delta_K(s_{t_j}))'$ plays the role of the mixing component.

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Proposed model - Mixing component

$$\log \delta_k(s_{t_j}) = \gamma_k(s_t) + \nu_k(s_{t_j}),$$

$\log(\text{mixing component}) = \text{temporal effect} + \text{local effect}$

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Proposed model - Mixing component

$$\log \delta_k(s_{t_j}) = \gamma_k(s_t) + \nu_k(s_{t_j}),$$

Modelling the temporal effect:

Let $\gamma(s_t) = (\gamma_1(s_t) \cdots \gamma_K(s_t))'$.

We assume

$$\gamma(s_t) \sim N_K(0, \mathbf{\Omega}), \quad \forall t = 1, \dots, T,$$

where $\mathbf{\Omega}$ captures the covariance among species at each time t .

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Proposed model - Mixing component

$$\log \delta_k(s_{t_j}) = \gamma_k(s_t) + \nu_k(s_{t_j}),$$

Modelling the local effect:

Let $\boldsymbol{\nu}(s_{t_j}) = (\nu_1(s_{t_j}), \dots, \nu_K(s_{t_j}))$, following the LMC (Gelfand et al 2004)

$$\boldsymbol{\nu}(s_{t_j}) = \mathbf{A}\boldsymbol{\omega}(s_{t_j}).$$

\mathbf{A} is lower triangular and $\boldsymbol{\omega}(s_{t_j}) = (\omega_1(s_{t_j}) \cdots \omega_K(s_{t_j}))'$ such that, independently,

$$\omega_k(\cdot) \sim GP(0, \rho(\boldsymbol{\vartheta}_k; d)), \quad k = 1, \dots, K.$$

Then

$$\text{Cov}(\boldsymbol{\nu}(s), \boldsymbol{\nu}(s')) = \sum_{k=1}^K \rho(\boldsymbol{\vartheta}_k; d) \mathbf{M}_k, \quad \text{with } \mathbf{M}_k = \mathbf{a}_k \mathbf{a}_k^T$$

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Joint distribution of $\log \delta$

Let δ be the nK -dimensional vector containing the elements of the mixing component,

$$\delta = (\delta(s_{1_1}), \dots, \delta(s_{T_{n_T}}))'$$

- ▶ **Separable model** ($\rho(\vartheta; d)$, for $k = 1, \dots, K$)

$$\log \delta \mid \gamma \sim N((I_K \otimes C)\gamma, \mathbf{R} \otimes \mathbf{M}),$$

where $\mathbf{M} = \mathbf{A}\mathbf{A}^T$.

- ▶ **Non-separable model** ($\rho(\vartheta_k; d)$)

$$\log \delta \mid \gamma \sim N\left((I_K \otimes C)\gamma, \sum_{j=1}^K \mathbf{R}_j \otimes \mathbf{M}_k\right).$$

We now discuss the specification of the spatial correlation function (elements of \mathbf{R}).

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- ▶ **Non-separable model** ($\rho(\vartheta_k; d)$)

$$\log \delta \mid \gamma \sim N\left((I_K \otimes C)\gamma, \sum_{j=1}^K \mathbf{R}_j \otimes \mathbf{M}_k\right).$$

We now discuss the specification of the spatial correlation function (elements of \mathbf{R}).

The correlation structure of $\omega_k(\cdot)$ is given by

$$\rho(s, s'; \boldsymbol{\vartheta}_k) = \exp(-\phi_k \|s - s'\|).$$

with parameter $\vartheta_k = \phi_k$.

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

The correlation structure of $\omega_k(\cdot)$ is given by

$$\rho(s, s'; \boldsymbol{\vartheta}_k) = \exp(-\phi_k \|d_k(s) - d_k(s')\|),$$

with $d_k(s) = s D_k$, and

$$D_k = \begin{bmatrix} \cos \psi_{A_k} & -\sin \psi_{A_k} \\ \sin \psi_{A_k} & \cos \psi_{A_k} \end{bmatrix} \begin{bmatrix} 1 & 0 \\ 0 & \psi_{R_k}^{-1} \end{bmatrix},$$

where ψ_{A_k} is the *anisotropy angle* and $\psi_{R_k} > 1$ is the *anisotropy ratio*, with parameters $\boldsymbol{\vartheta}_k = (\phi_k, \psi_{A_k}, \psi_{R_k})$.

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Covariates in the covariance structure (Schmidt et al. 2011)

Now location has a broader interpretation.

Let $\mathbf{w} = (\text{long}, \text{lat}, z_3, \dots, z_C)$, and

$$\rho(\mathbf{w}, \mathbf{w}'; \boldsymbol{\vartheta}_k) = \exp \left(-\sqrt{(\mathbf{w} - \mathbf{w}')^T \Phi^{-1} (\mathbf{w} - \mathbf{w}')} \right).$$

Similar to assuming a GP in R^C , generating an anisotropic correlation function in R^2 (“geographical” space).

Our model assumes

$$\rho(\mathbf{w}, \mathbf{w}'; \boldsymbol{\vartheta}_k) = \exp \left(-\phi_1 \|s - s'\| - \phi_2 |z - z'| \right),$$

where z is *geodetic depth* (because lake level fluctuates, local habitat [travels](#), but geodetic depth is [fixed](#)).

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

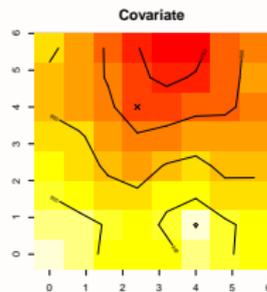
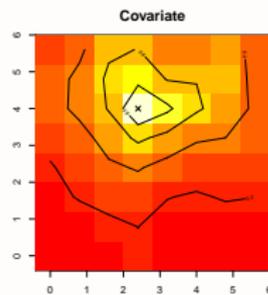
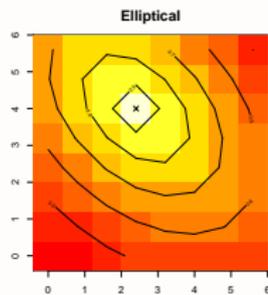
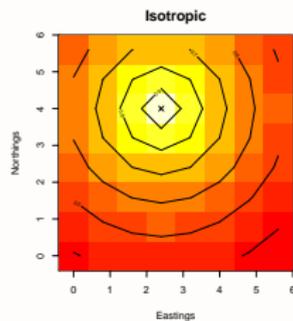
Main references

Comparing correlation structures

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Likelihood function

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Let

$\mathbf{y} = (\mathbf{y}(s_{1_1}), \mathbf{y}(s_{1_2}), \dots, \mathbf{y}(s_{1_{n_1}}), \dots, \mathbf{y}(s_{T_1}), \dots, \mathbf{y}(s_{T_{n_T}}))'$
be the observed counts over the sampling period at
each location s_{t_j} , $t = 1, \dots, T$, $j = 1, \dots, n_t$.

The likelihood function is

$$l(\mathbf{y} \mid \boldsymbol{\theta}, \boldsymbol{\delta}) \propto \prod_{t=1}^T \prod_{j=1}^{n_t} \prod_{k=1}^K \exp \{ -\theta_k(s_{t_j}) \delta_k(s_{t_j}) \} [\theta_k(s_{t_j}) \delta_k(s_{t_j})]^{y_k(s_{t_j})} .$$

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

- ▶ $\beta \sim N(0, \sigma_\beta^2 I_p)$
- ▶ $\phi_{ij} \sim IG(2, b)$, $i = 1, 2, j = 1, \dots, K$, and b is fixed such that the practical range (correlation = 0.05) is reached at half of the maximum distance between observations
- ▶ $\Omega \sim IW(K + 1, c I_k)$
- ▶ $\psi_R \sim \text{Pareto}(1, 2)$, $\psi_A \sim U(0, \pi)$
- ▶ $a_{ij} \sim N(0, 5)$ and $\log a_{ii} \sim N(0, 5)$

MCMC sampling scheme

We reparametrize the model such that

$\varphi_k(s_{t_j}) = \theta_k(s_{t_j}) \delta(s_{t_j})$, and

$$\log \varphi_k(s_{t_j}) = X_k^*(s_{t_j}) \beta_k^* + W_k(s_{t_j})$$

and $W_k(s_{t_j}) = \beta_{1k} + \gamma_k(s_{t_j}) + \nu_k(s_{t_j})$, where

$X_k^*(\cdot)$ does not have a column of ones, and

$\beta_k^* = (\beta_{2k}, \dots, \beta_{p_k k})^T$.

- ▶ β_k and $W_k(\cdot)$ M-H algorithm proposed by Gamerman (1997)
- ▶ $\beta_{1\cdot} = (\beta_{11}, \dots, \beta_{1K})'$ and γ normal posterior full conditionals
- ▶ Ω follows an inverse-Wishart posterior full conditional
- ▶ $\phi_1, \phi_2, \psi_R, \psi_A$, and elements of \mathbf{A} are sampled through M-H steps

Fitted models

M1: Separable isotropic covariance structure

M2: Separable elliptical anisotropy covariance structure

M3: Separable covariate-dependent ($z =$ geodetic depth) covariance structure

M4: Non-separable isotropic covariance structure

M5: Non-separable elliptical anisotropy covariance structure

M6: Non-separable, covariate-dependent ($z =$ geodetic depth) covariance structure

- ▶ $\gamma(s_t)$ assumed to be independent across time and common to both shores
- ▶ We allow for different local random effects for the North and South shores (Glémet and Rodríguez 2007; Schmidt et al. 2010b)

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Stats
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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Fitted models

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in space

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Stats
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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Fitted models

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Búzios, Brazil, 2014

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Fitted models

M1: Separable isotropic covariance structure

M2: Separable elliptical anisotropy covariance structure

M3: Separable covariate-dependent ($z =$ geodetic depth) covariance structure

M4: Non-separable isotropic covariance structure

M5: Non-separable elliptical anisotropy covariance structure

M6: Non-separable, covariate-dependent ($z =$ geodetic depth) covariance structure

- ▶ $\gamma(s_t)$ assumed to be independent across time and common to both shores
- ▶ We allow for different local random effects for the North and South shores (Glémet and Rodríguez 2007; Schmidt et al. 2010b)

Multivariate counts
in space

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PASI 2014
Multivariate Spatial
Stats
Búzios, Brazil, 2014

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Fitted models

M1: Separable isotropic covariance structure

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Multivariate counts
in space

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PASI 2014
Multivariate Spatial
Stats
Búzios, Brazil, 2014

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Fitted models

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Multivariate counts
in space

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PASI 2014
Multivariate Spatial
Stats
Búzios, Brazil, 2014

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Fitted models

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Multivariate counts
in space

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PASI 2014
Multivariate Spatial
Stats
Búzios, Brazil, 2014

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Model comparison

Yellow perch

Model	p_D	DIC	EPD
M1	144.8	1039.0	2019063.4
M2	145.2	1038.6	2019061.1
M3	137.1	1031.3	2019023.1
M4	145.1	1040.0	2019030.6
M5	144.0	1036.4	2019067.2
M6	136.5	1032.1	2018988.0

Brown bullhead

Model	p_D	DIC	EPD
M1	109.4	626.2	133777.8
M2	110.1	628.3	133778.2
M3	101.1	621.3	133755.5
M4	111.9	633.7	133768.1
M5	112.6	635.1	133770.4
M6	101.1	622.9	133754.1

Golden shiner

Model	p_D	DIC	EPD
M1	106.0	590.1	246747.8
M2	106.0	591.7	246729.9
M3	98.6	585.4	246714.3
M4	103.2	592.1	246734.8
M5	105.5	596.2	246741.1
M6	99.6	590.6	246699.6

Pumpkinseed

Model	p_D	DIC	EPD
M1	81.9	432.4	54044.9
M2	81.4	429.8	54056.2
M3	75.6	428.9	54037.7
M4	82.9	434.0	54050.2
M5	82.8	434.7	54046.9
M6	75.7	432.7	54038.1

Table : Values of the effective number of parameters, p_D , DIC (Spiegelhalter et al. 2001), and EPD (Gelfand and Ghosh, 1998) for the six fitted models, by fish species.

Models including geodetic depth as a covariate in the correlation structure of the spatial effects generally fit better than those assuming isotropy or geometrical anisotropy

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison

Results

Conclusions

Main references

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

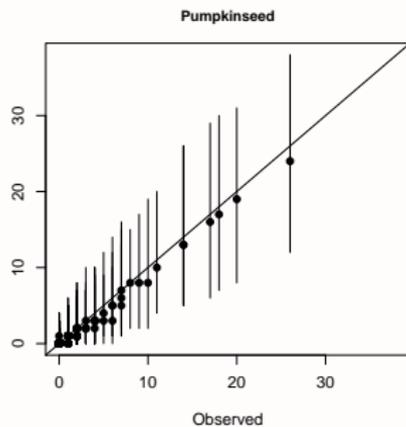
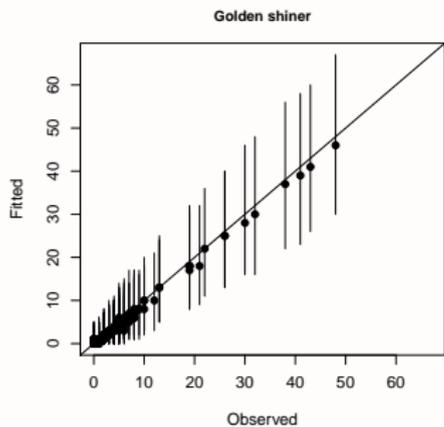
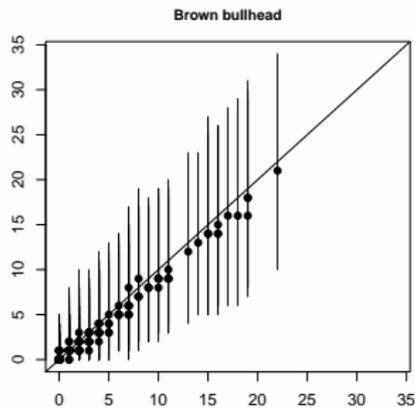
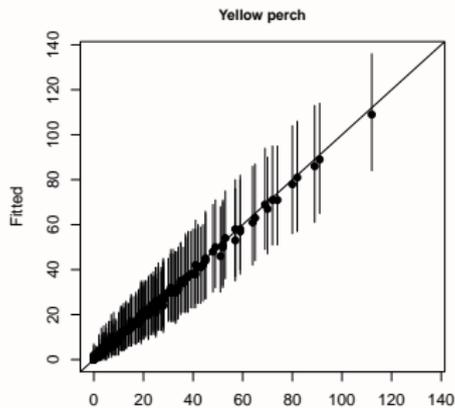
Results

Model comparison

Results

Conclusions

Main references



Posterior predictive distributions for counts of the four species
under M6 showed agreement with observed counts

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

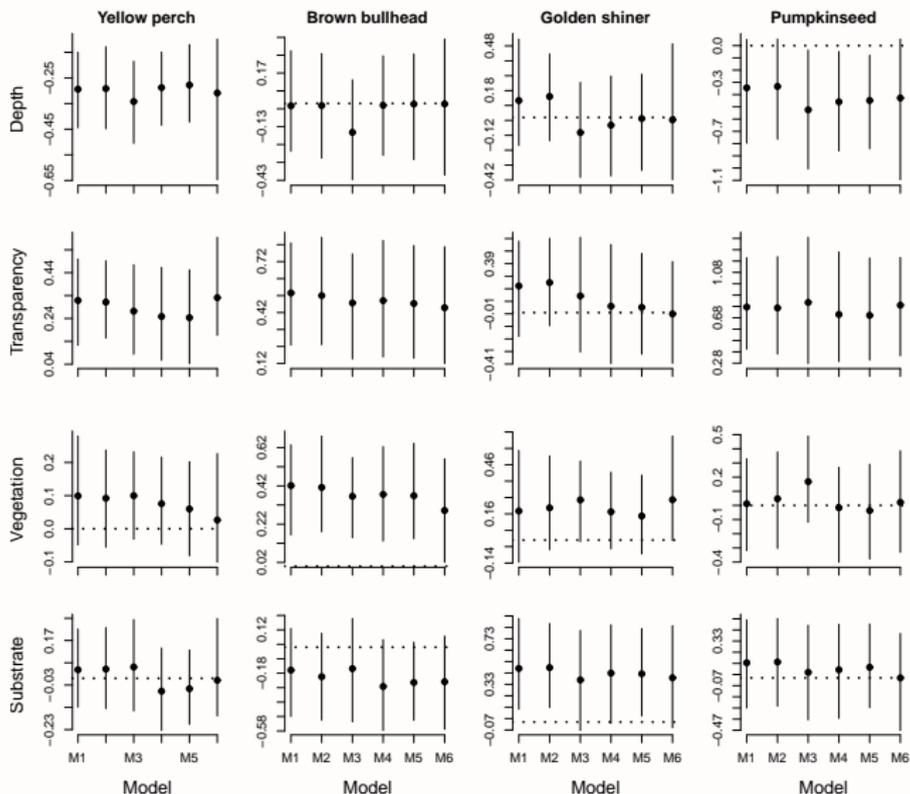
Results

Model comparison

Results

Conclusions

Main references



Water transparency had positive influence on the abundance of three of the four fish species, whereas water depth and vegetation each influenced the abundance of one fish species. Substrate composition had no apparent effect on fish abundances

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

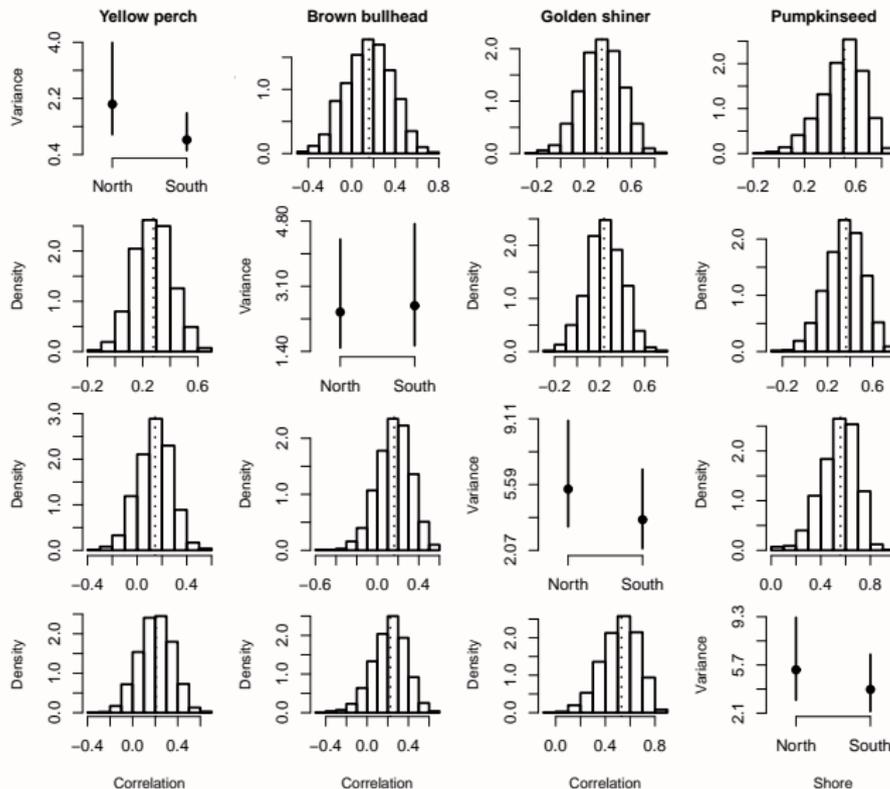
Results

Model comparison

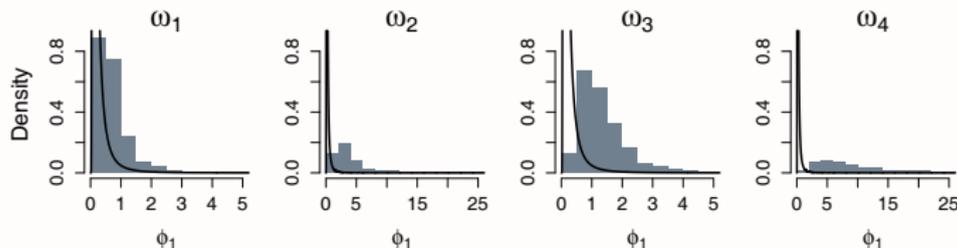
Results

Conclusions

Main references



Variations and correlations tended to be greater in the North shore than in the South shore. The two strongest correlations, both in the North shore, were positive. Negative correlations, possibly indicative of competition between species, were not apparent.



For all non-separable models, the decay parameter ϕ_1 showed well-defined information gain and marked differences among component spatial processes $\omega_k(\cdot)$.

[Outline](#)

[Motivation and aims](#)

[Distr. multiv. counts](#)

[Proposed model](#)

[Inference procedure](#)

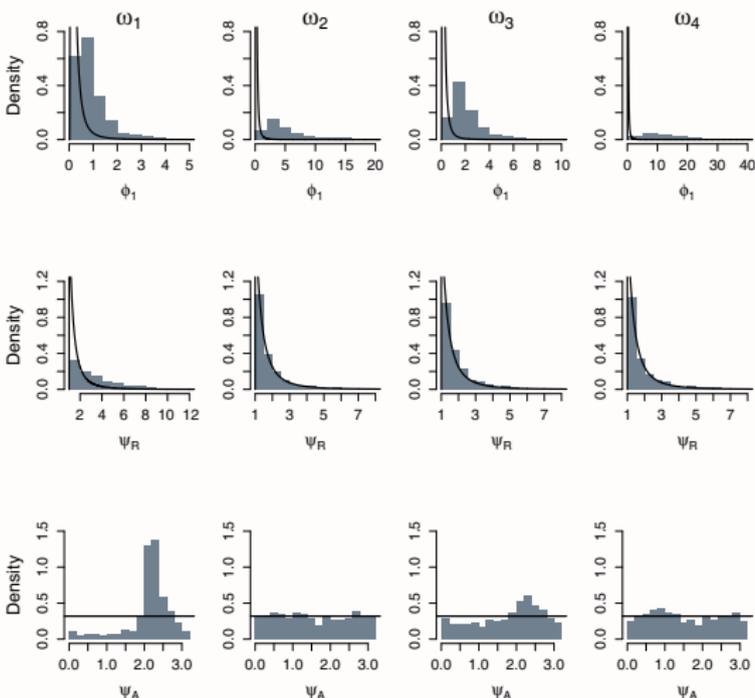
[Results](#)

Model comparison

Results

[Conclusions](#)

[Main references](#)



M5 provided strong evidence of anisotropy associated with spatial process $\omega_1(\cdot)$, but little deviation from the priors for components $\omega_2(\cdot)$, $\omega_3(\cdot)$, and $\omega_4(\cdot)$, indicating a single anisotropic pattern shared across species and suggesting that the spatial component of M5 may be overparametrized.

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

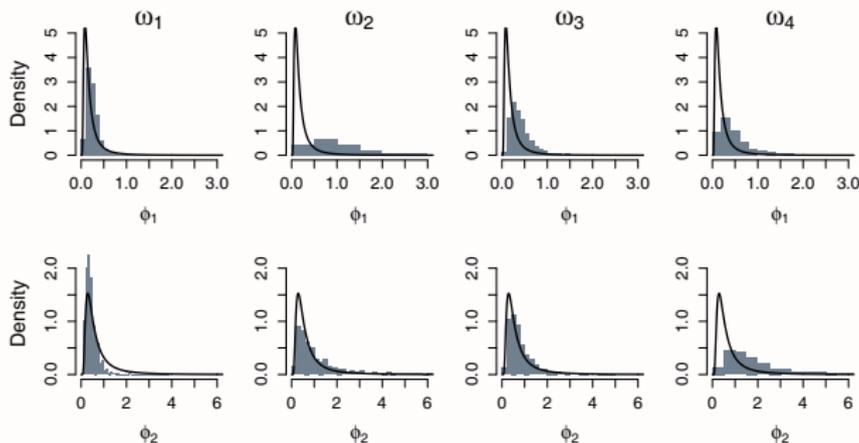
Results

Model comparison

Results

Conclusions

Main references



Similar to M5, the posterior distribution of ϕ_2 under M6 provides strong evidence of anisotropy. However, in contrast to M5, the gain in information reflected in differences between priors and posteriors is distributed across components, indicating that species do not share a common anisotropic pattern.

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

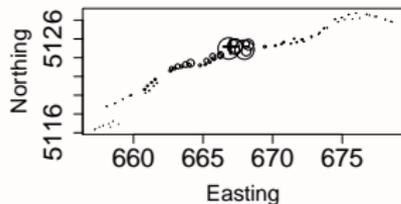
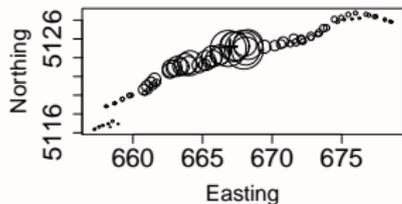
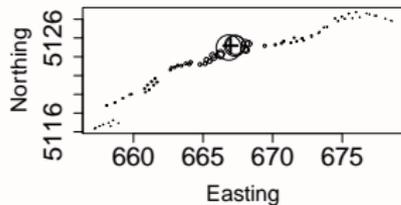
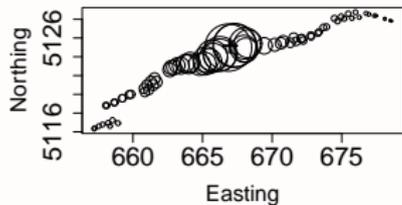
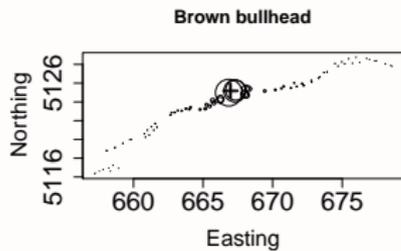
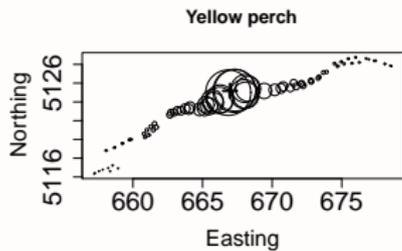
Results

Model comparison

Results

Conclusions

Main references



Decay of correlation under M6 had directionality generally similar to that of M5, but was further modified by geodetic lake depth.

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

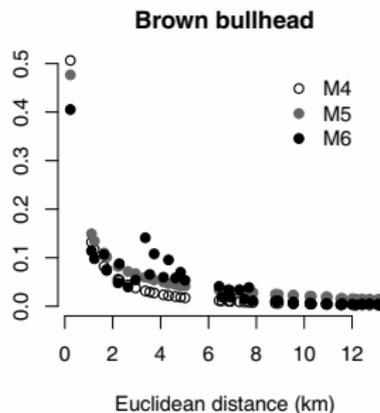
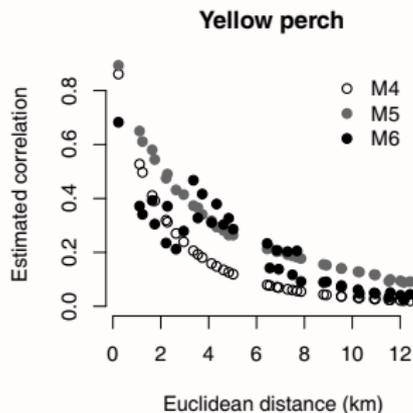
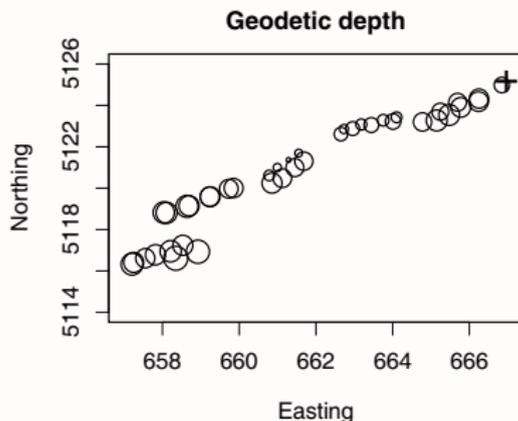
Results

Model comparison

Results

Conclusions

Main references



[Outline](#)

[Motivation and aims](#)

[Distr. multiv. counts](#)

[Proposed model](#)

[Inference procedure](#)

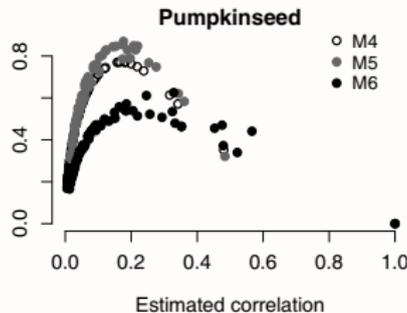
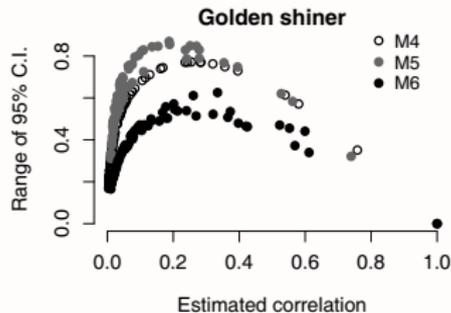
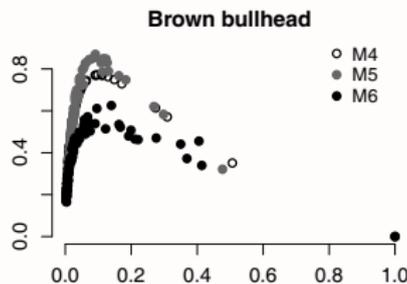
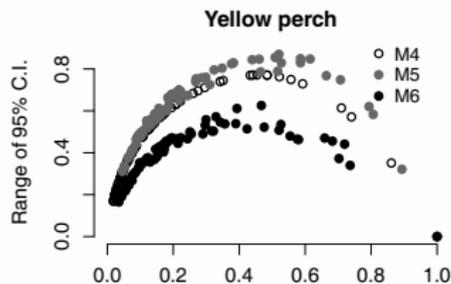
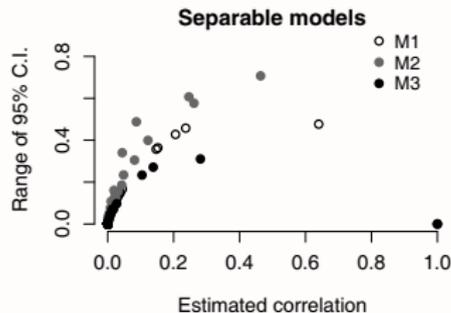
[Results](#)

Model comparison

Results

[Conclusions](#)

[Main references](#)



Conclusions

- ▶ Our model **accounts for overdispersion** and **both** positive and negative covariances (among species and across space)
- ▶ The LMC provided **flexible** covariance structures for the mixing component
- ▶ Anisotropic spatial effects improved fit (DIC and EPD)
- ▶ Including information on **geodetic lake depth** in the spatial covariance structure of the spatial process provided a flexible, yet relatively simple, means of capturing **anisotropy** along the shorelines of the lake

Multivariate counts
in space

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Multivariate Spatial
Stats
Búzios, Brazil, 2014

Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references

Main references

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Outline

Motivation and aims

Distr. multiv. counts

Proposed model

Inference procedure

Results

Model comparison
Results

Conclusions

Main references