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Previous word count: 2342
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Current page count:

REVIEW ARTICLE

The Integrated Nested Laplace Approximation applied to Spatial Point Process Models

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ARTICLE HISTORY

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ABSTRACT

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KEYWORDS

INLA, spatial prediction, log-Gaussian Cox process, spatial point process

1. Introduction

Spatial prediction is a high-dimensional inference problem. When the goal of statistical modeling is to produce a graphical map of a random variable over space, the model ultimately must be able to predict that random variable at every pixel of the image. A map image will typically be at least several hundred by several hundred pixels, so in total there can easily be hundreds of thousands of pixels requiring predictions. Thus, even when a model has only half a dozen parameters, it may include hundreds of thousands of latent variables.

Spatial point process models further complicate the situation with difficult likelihoods. (*Cite some computational papers — Baddely?*) Both maximum likelihood and Bayesian model fitting require integrating the intensity function over space, but the integral is generally not available in closed form. Many methods have been introduced including quadrature-based approximations (*cite Baddeley*), pseudodata approaches (*cite Baddeley/Berman/Turner etc*), and Markov chain Monte Carlo [11].

Development of the integrated nested Laplace approximation (INLA) has made accurate approximate model fitting considerably more feasible for a particular class of log-Gaussian Cox process (LGCP) models. INLA was developed to fit Bayesian hierarchical models with many latent Gaussian variables [13]. A key part of INLA's computational simplicity is that it calculates the posterior distribution of each latent Gaussian variable one at a time; that is, it provides only the posterior marginal distributions rather than the full joint distribution.

When using a LGCP for spatial mapping, two aspects make INLA a suitable approach. First, the LGCP is driven by a spatial Gaussian process (GP), so the latent variables are Gaussian. Second, even though the latent variables are expected to ex-

hibit spatial dependence, their full joint distribution is not needed. In most situations it suffices to map their predicted values, variance, and upper and lower interval bounds pointwise across space.

We focus on spatial mapping using a hierarchical construction of the LGCP, and do so without discussing many classic spatial point process concepts such as Gibbs processes, Markov processes, point process densities, or the Papangelou conditional intensity function. For readers interested in the general spatial point process context in which the LGCP originates, we recommend other references [6, 7, 11].

Much recent work in point process modeling involves spatiotemporal models, which are easily implemented in INLA by allowing the parameters of a spatial point process model to vary over time. To keep the focus on the spatial aspects, we do not discuss spatiotemporal models. This article provides a review of recent advances in the fitting of spatial LGCP models via INLA, including dimension reduction by triangulation, a likelihood factorization that avoids gridding, and incorporation of sampling or false negatives.

1.1. *Log-Gaussian Cox Process*

Notes from [11]:

- applications mentioned: seismology, ecology, forestry, geography, spatial epidemiology and materials science
- modern computing (e.g. MCMC) allows for model-based inference without a stationarity assumption
- Minke whale distance sampling data used as an example of incomplete observation (they used a shot noise Cox process model)
- distribution of point process is the joint distribution of n and the n events, or equivalently the distribution of the number of events in any arbitrary region
- in their presentation of the two Poisson process postulates, the second one is given the number of events in a region they are iid proportional to the intensity function
- Cox process is driven by a non-negative random process such that conditional on the driving process the Cox process is a Poisson process
- LGCP has no edge effects because the point process in the window is fully specified by the GP restricted to the window
- quite a bit about Gibbs processes, densities wrt stationary Poisson processes, and Papangelou conditional intensities (comments: The density was traditionally used to facilitate MCMC simulation and ML fitting when the likelihood was not explicitly available. The conditional intensity of a Poisson process is the intensity function because of independence among the points. The hierarchical setup of the LGCP lets us work directly with a Poisson process so the density and conditional intensity are not needed. LGCP models with constructed covariate can model Gibbs processes.)
- more things about infinite Gibbs and Markov processes that go over my head but we don't need because we deal with bounded and finite processes

Notation and terminology:

- process defined on $\mathcal{D} \subset \mathbb{R}^2$, domain of the intensity function, define $d = \dim(\mathcal{D})$
- observation window $\mathcal{S} \subset \mathcal{D}$

- (Note: When thinking about sampling, we need to define three regions: the domain \mathcal{D} over which the process mathematically operates, the region \mathcal{R} over which inferences are desired, and the observed/sampled observation window \mathcal{S} . The general relationship is $\mathcal{S} \subset \mathcal{R} \subset \mathcal{D} \subset \mathbb{R}^d$, where all of the subset symbols taken to mean “subset or equal”. The “fully surveyed” situation is $\mathcal{S} = \mathcal{R}$.)
- \mathbf{X} point process on \mathcal{R} , $\mathbf{x} = \{x_1, \dots, x_n\}$ realized point pattern
- point $x \in \mathbf{x}$ called an event
- intensity function $\lambda(u)$
- x event, s numerical integration node, u arbitrary location in \mathcal{D}
- $z(u)$ a column vector of covariates/predictors at u
- “point” refers to a u unless clearly stated otherwise
- bold for sets and spatial processes, normal italics for spatial vectors
- y and variations will be used for objects derived from the point pattern, e.g. pseudodata

The LGCP is a Poisson process driven by a latent Gaussian process [12]. A Poisson process is characterized entirely by its intensity function $\lambda(u)$, which gives the mean number of events per unit area, and the process satisfies the following two properties (*find a good citation*).

- (1) The number of events in any region \mathcal{B} follows a Poisson distribution with mean $\int_{\mathcal{B}} \lambda(u) du$.
- (2) The numbers of events in disjoint regions are independent.

Commonly, covariates are incorporated via a log-linear model for the intensity,

$$\log \lambda(u) = \mathbf{z}'(u)\boldsymbol{\beta}.$$

The LGCP adds another stochastic layer,

$$\log \lambda(u) = \mathbf{z}(u)'\boldsymbol{\beta} + \boldsymbol{\Psi}(u),$$

where $\boldsymbol{\Psi}$ is a Gaussian process.

Where the spatial Poisson process has a (stochastically) fixed intensity to be estimated from data, the spatial LGCP induces a hierarchical model where the intensity function is itself a random process to be predicted.

Möller and Waagepetersen (2007) present model checking tools for LGCP models [11].

1.2. Integrated Nested Laplace Approximation

INLA was developed with the goal of providing fast, accurate, deterministic approximations for posterior marginal distributions of the parameters in latent Gaussian models [13]. The setting is Bayesian generalized additive models but is kept very general; this includes linear models, models with nonlinear (spline, random walk, etc.) functions of predictors, and models with temporal and/or spatial dependence. Such models commonly have a large number of Gaussian latent variables or parameters with Gaussian priors and relatively few non-Gaussian parameters. The INLA approach makes repeated use of Laplace expansion, numerical integration, and numerical search. The method has established usefulness for LGCP [8]. It is readily implemented using the standalone INLA software or in R via the R-INLA package [9].

2. Methodology

INLA provides an efficient computational framework for fitting Bayesian models with latent Gaussian variables. On top of this framework are built several tools to further simplify the fitting of spatial LGCP models. The stochastic partial differential equation (SPDE) approach provides dimension reduction for the spatial GP [10]. The SPDE approach employs a numerical integration scheme which can also be used to approximate the LGCP likelihood and negate the need to grid the events into Poisson counts [14]. With these computational improvements, researchers are now able to efficiently fit LGCP models to incompletely-observed point patterns [15].

2.1. The SPDE Approach

Because the LGCP includes a Gaussian process (GP), efficient computation for Gaussian processes is critical when working with LGCP models.

The GP imposes a dense covariance matrix on the latent variables [3]. For GPs with a Matérn covariance function, a Gaussian Markov random field (GMRF) approximation can simplify computation, requiring only a sparse covariance structure.

The GMRF approximation is motivated by the fact that Gaussian fields with Matérn covariances are solutions to the below stochastic partial differential equation (SPDE) [10].

$$(\kappa^2 - \Delta)^{\alpha/2} \Psi(u) = \mathbf{W}(u), \quad u \in \mathbb{R}^d, \quad \kappa > 0, \quad \alpha = \nu + d/2, \quad \nu > 0$$

Here, \mathbf{W} is a Gaussian white noise process with variance 1, and Δ is the Laplacian operator. The stationary solution Ψ is a Gaussian field with a Matérn covariance function with scaling parameter κ (approximately inversely proportional to the range) and smoothness parameter ν . The variance is a function of κ , ν , and d .

Lindgren, Rue, and Lindström investigate the limit as $\nu \rightarrow 0$ for $d = 2$, finding that the solution is a GMRF on a unit lattice [10]. They then construct approximations for positive integer values of ν by ν -fold convolution of Ψ with itself. Finally, they use a finite element method to generalize the approximation to arbitrary triangulations of the support. This approximation has considerable computational benefits because the GMRF has a sparse covariance structure; the only nodes with nonzero covariances are those directly connected by edges in the triangulation.

In practice, the SPDE approach goes as follows. Choose nodes s_i at which to model $\Psi(s_i)$, then build a triangular mesh using these nodes. Typically the nodes will include locations where data or covariates are available, then the rest will be filled in with a Delaunay triangulation under some edge length constraints. The $\Psi(s_i)$ are modeled as a GMRF where the distribution of each $\Psi(s_i)$ depends only on the $\Psi(s_j)$ where s_i and s_j are connected by an edge. The GMRF representation is assumed to be a piecewise linear approximation of the continuous Gaussian field; values of $\Psi(s)$ for s not in the set of nodes are predicted by linear interpolation using the barycentric coordinates of s .

The SPDE approach is implemented in the R-INLA package along with tools for constructing meshes. A wrapper for easily fitting LGCP models is provided in the `inlabru` package [1].

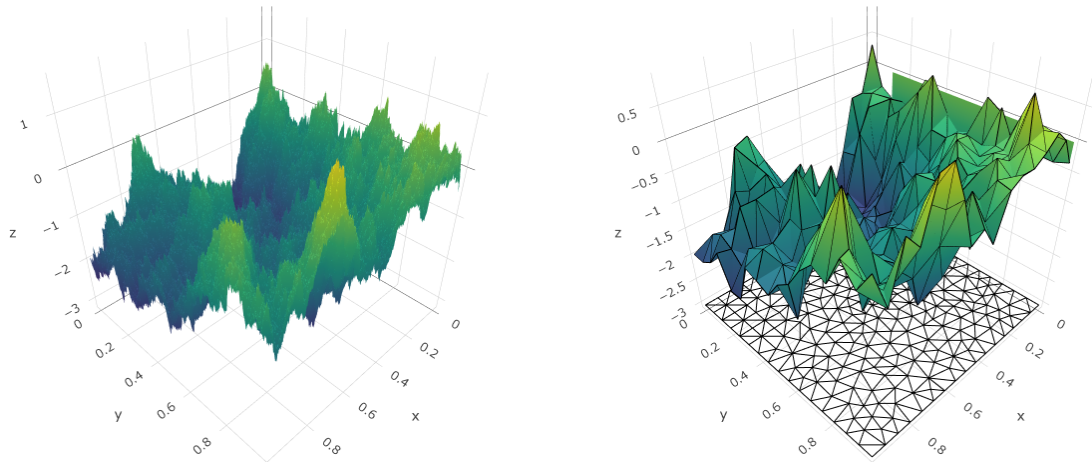


Figure 1. A realization of a spatial Gaussian process (left) and an approximation of that realization over a triangular mesh (right).

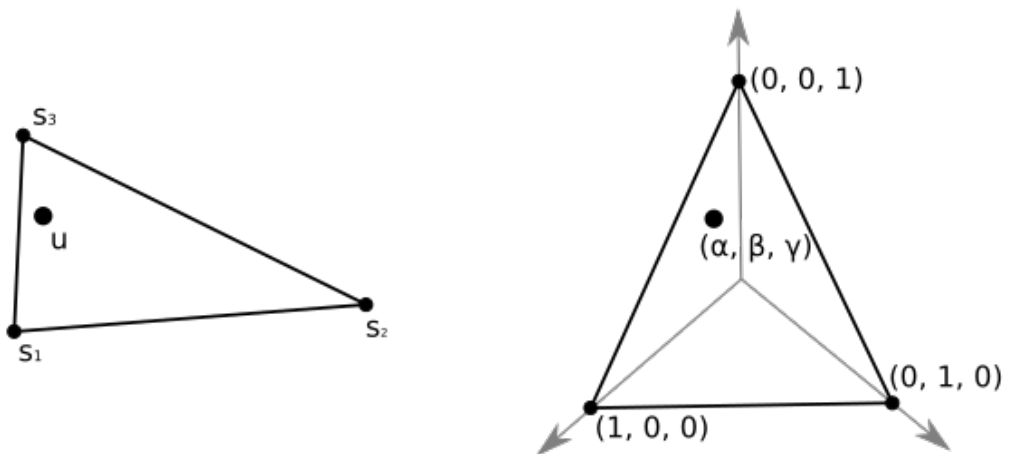


Figure 2. An illustration of the linear transformation from a mesh triangle to the simplex. (α, β, γ) are the barycentric coordinates of u .

Example Mesh

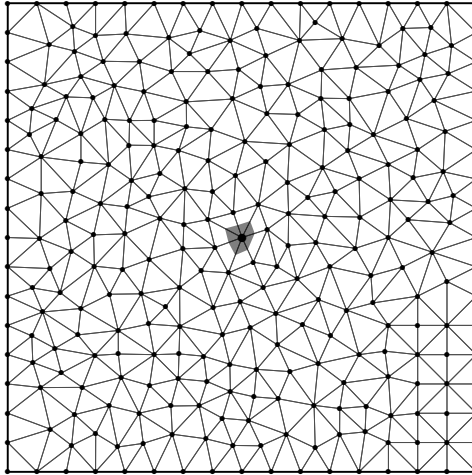


Figure 3. An illustration of the nodes (dots) and the weighting scheme for one node (large dot). This node represents the shaded region, so its weight is proportional to the shaded area. The shaded region is constructed by connecting the midpoints of the node’s edges.

2.2. Going Off the Grid

Point pattern data are sometimes known as presence-only data, underscoring the fact that information about where events did *not* occur is both important and often overlooked. There have been many proposed methods to account for regions that were observed to contain no events (in contrast to unobserved regions where it is unknown if any events are present). Many of these methods involve imputation of dummy points or discretization. In the world of maximum likelihood, perhaps the most well-developed of these use approximations based on logistic regression on presence/absence information in small disjoint regions (*cite a bunch of Baddely etc papers*).

Another alternative, which is probably the most popular approach to Bayesian fitting of LGCP models but is also common in frequentist analyses, is to grid the domain and model the induced Poisson counts. It has long been understood that results are sensitive to the discretization scheme [4]. Simpson et. al. (2016) explain that this is also computationally wasteful [14].

Simpson et. al. took LGCP inference “off grid” by introducing a computationally-efficient approximation to the Poisson process likelihood that requires the intensity function only to be evaluated at the locations of observed events and at the nodes of a mesh. Thus, the SPDE approach can be employed to model the intensity surface and the same nodes reused in evaluation of the Poisson process likelihood. The result is a substantial improvement in both computing time and accuracy of the approximation compared to gridding.

The approximation arises from a factorization of the Poisson processes likelihood. The exact log-likelihood is

$$\ell(\lambda) = C - \int \lambda(u) du + \sum_{i=1}^n \log [\lambda(x_i)]$$

where C is a normalizing constant. The log-intensity is projected into the space

spanned by a finite set of basis function representation is used for $\log[\lambda(u)]$, namely

$$\log[\lambda(u)] \approx \sum_{j=1}^m \psi_j \phi_j(u)$$

with $\boldsymbol{\psi} = (\psi_1, \dots, \psi_m)'$ a multivariate normal random vector and $\{\phi_1, \dots, \phi_m\}$ a set of linearly independent basis functions. This defines a numerical integration scheme at nodes s_i with weights α_i and yields the approximation,

$$\begin{aligned} \ell(\lambda) &\approx C - \sum_{i=1}^m \tilde{\alpha}_i \exp \left[\sum_{j=1}^m \psi_j \phi_j(s_i) \right] + \sum_{i=1}^n \sum_{j=1}^m \psi_j \phi_j(x_i) \\ &= C - \boldsymbol{\alpha}' \exp[\mathbf{A}_1 \boldsymbol{\psi}] + \mathbf{1}' \mathbf{A}_2 \boldsymbol{\psi}. \end{aligned}$$

This is equivalent to the likelihood of independent Poisson random variables with means (*translate the confusing algebra from the appendix*). Thus, the SPDE approach, which can be easily fit in R-INLA, can be combined with a Poisson GLM to rapidly fit a LGCP model. There is no need to compute grid counts, only to define dummy points at the mesh nodes.

2.3. Variable Sampling Effort

There has long been a gap between point process modeling theory and practice, where point process models are only fit to data from completely-observed domains under the assumption that every event was detected perfectly. In practice, this is not the case as it may be impossible or impractical to census the entire region of interest. For example, line-transect surveys routinely generate point pattern data but are analyzed after aggregation rather than using a point process model. Another issue is that of false negatives, in other words events which exist but are not detected during the survey. False negatives are an accepted part of species abundance surveys, where the species of interest may be camouflaged or hidden in thick plant cover.

The idea of the incompletely-observed domain has been around for some time. Brix and Möller (2001) fit an LGCP model to weed data observed in rectangular frames and used a Metropolis-adjusted Langevin algorithm to predict the intensity outside of the observed frames [4]. Chakraborty et. al (2011) discuss nonhomogeneous Poisson process modeling as a richer alternative to ecological presence/absence models and describe their data as “degraded” in the sense that sampling bias prevented the entire region from being fully observed; they fit their model by aggregating the point pattern to counts in grid cells and using MCMC [5].

With INLA, the SPDE approach, and the “off grid” approximation facilitating the routine fitting of LGCP models, there is now renewed interest in accounting for variable sampling effort in spatial point process models [14, 15].

Sampling effort is accounted for using the theory of thinned point processes. Thinning refers to the events of one point process being kept or discarded probabilistically. Let $\lambda(u)$ be the intensity of the parent process, and let an event at u (if it exists) be observed with probability $p(u)$. The observed point process is a thinned point process with intensity

$$\lambda_p(u) = p(u)\lambda(u),$$

and if the parent process is a Poisson process then the observed process is also a Poisson process [11].

We seek to make posterior inferences about the parent intensity, $\lambda(u)$. If $p(u)$ is known, its value at each node is used to adjust the SPDE integration weights. Most usefully, if it is known that $p(s) = 0$ at certain nodes s because they were outside the surveyed domain, the weights become zero so those nodes do not contribute to the integral. If $p(u)$ is unknown, it can be modeled. Taking the logarithm of the above and substituting in the basis function representation, we have the log-linear model

$$\log [\lambda_p(u)] = \log [p(u)] + \sum_{j=1}^m \psi_j \phi_j(u).$$

Thus, any log-linear model for $p(u)$ that can be fit by INLA and the SPDE approach can be incorporated into the LGCP model. For example, Yuan et. al (2017) fit such a model to data from a line-transect survey using a spline model to account for the unknown detection function [15].

3. Applications

3.1. Simulation Study

ideas:

- simulate LGCPs
- fit via Simpson method
- vary mesh coarseness
- explore complexity of predictors vs latent GP
- explore bias/variability in parameter posteriors
- explore spatial prediction error in latent GP

3.2. Data Application

Examples with data, maybe `bei` dataset or Victorville. Should read like a tutorial because there aren't many good resources yet.

3.2.1. Data and Model

For an example analysis of real-world data, we consider the locations of 3605 *Beilschmiedia pendula* Lauraceae trees in a tropical rainforest [11]. The data are available as the `bei` dataset in the `spatstat` R package [2]. The point pattern exhibits inhomogeneity (Figure 4) which appears to be associated with elevation (Figure 5) and the magnitude of the gradient (Figure 6).

3.2.2. Fitting in R-INLA

- explain the mesh
- explain the weight calculations
- walk through the spatial predictions

bei

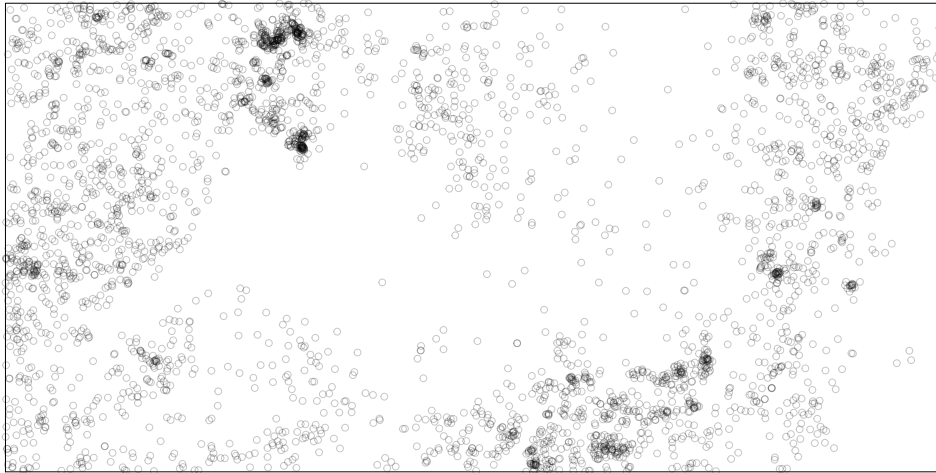


Figure 4. Locations of trees (events).

3.2.3. Results

3.2.4. Model Checking

Pearson residuals: small magnitude in the region with no observed events and low posterior intensity (not possible to see much variation), the model overestimates the intensity in the canyon (too small to be accurately represented on the mesh), random with no obvious spatial patterns elsewhere, a few large-magnitude positive residuals (maybe there is some clustering?), tried a few different grids (not shown) and saw the same patterns

4. Conclusion and Discussion

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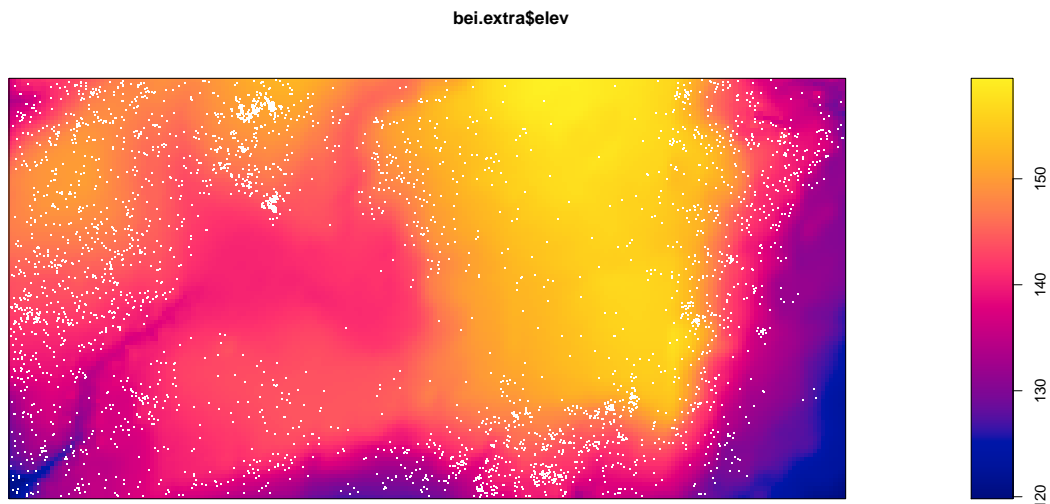


Figure 5. Elevation data.

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bei.extra\$grad

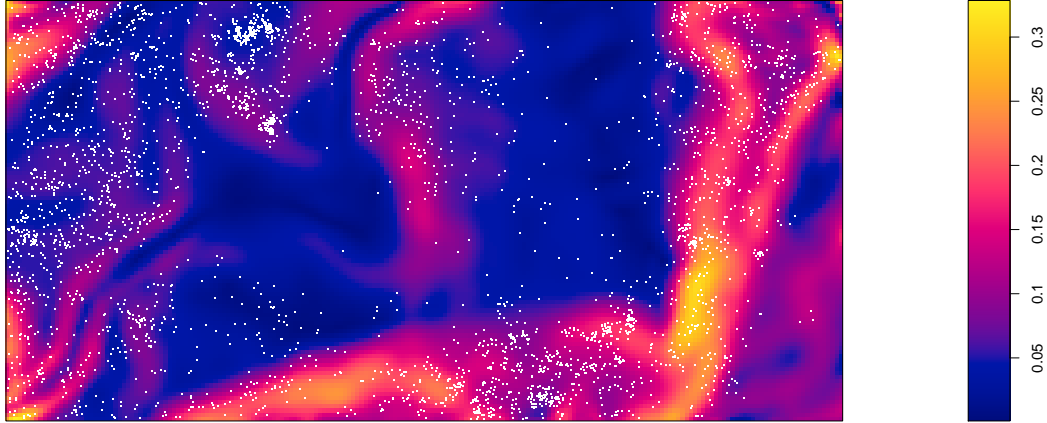


Figure 6. Gradient data.

Mesh Over Bei Data

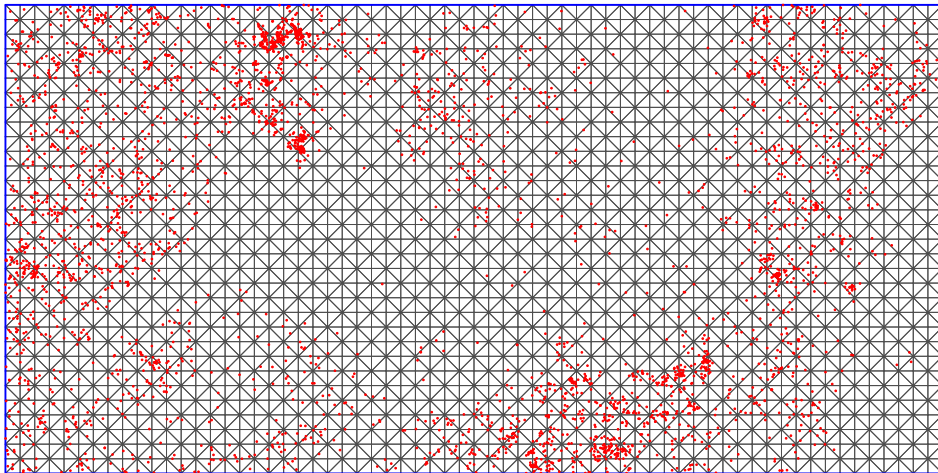


Figure 7. Triangular mesh with events overlaid.

Piecewise Linear Approximation of Elevation

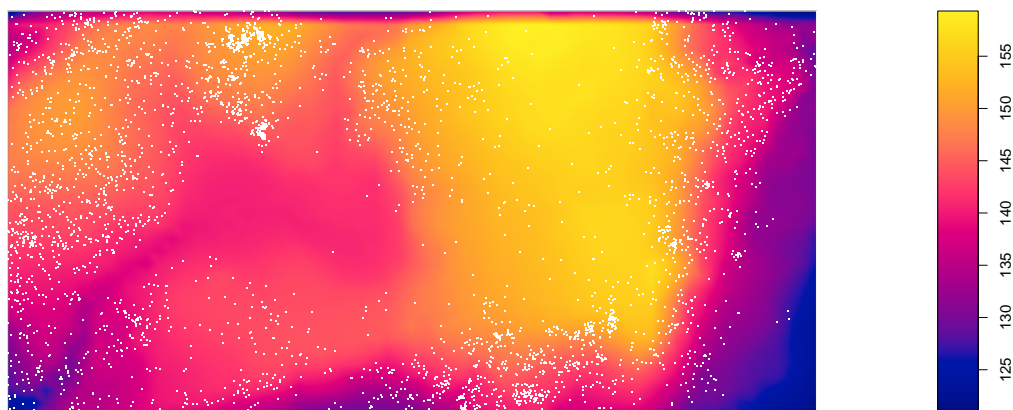


Figure 8. Piecewise linear approximation of the elevation surface.

Piecewise Linear Approximation of Gradient

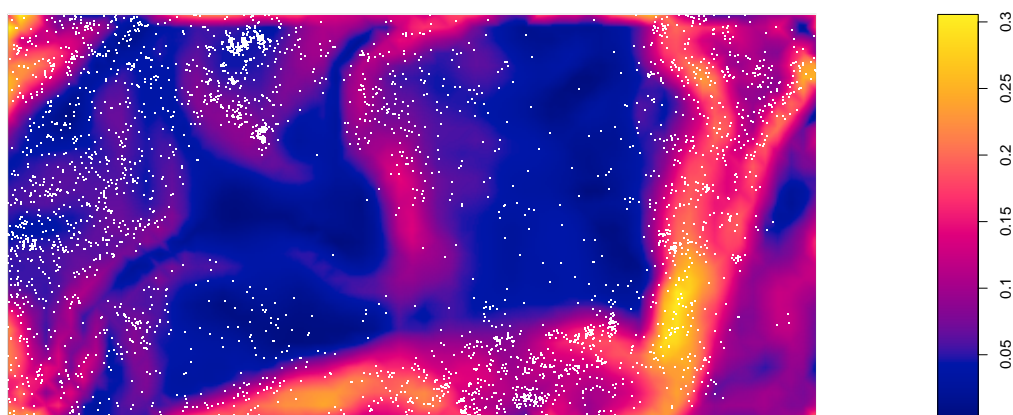


Figure 9. Piecewise linear approximation of the gradient surface.

Posterior Mean of Latent GP

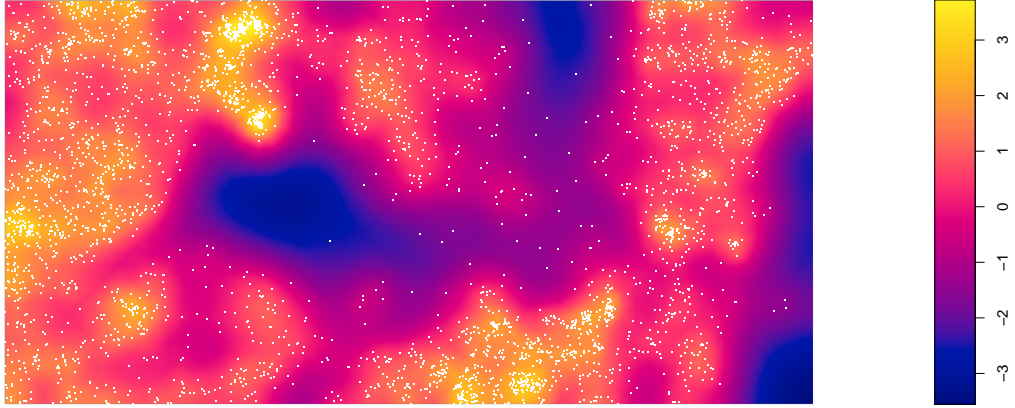


Figure 10. Posterior mean prediction surface.

Posterior SD of Latent GP

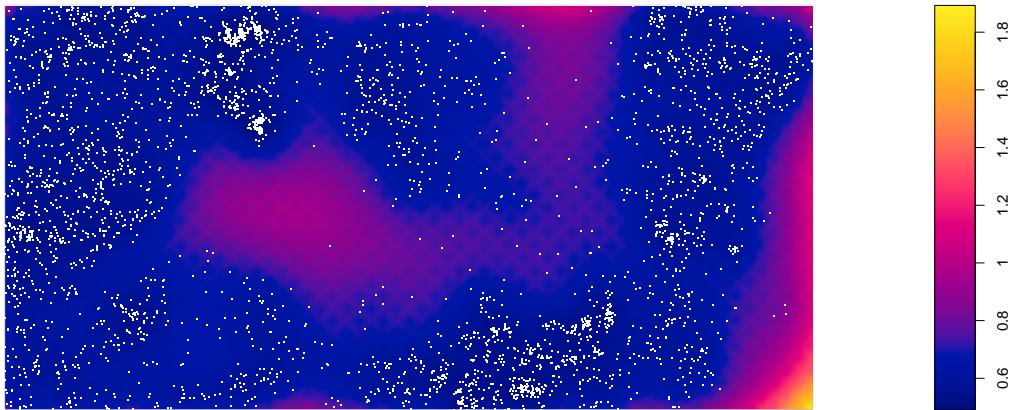


Figure 11. Pointwise posterior standard deviation of predictions.

Posterior Intensity Function

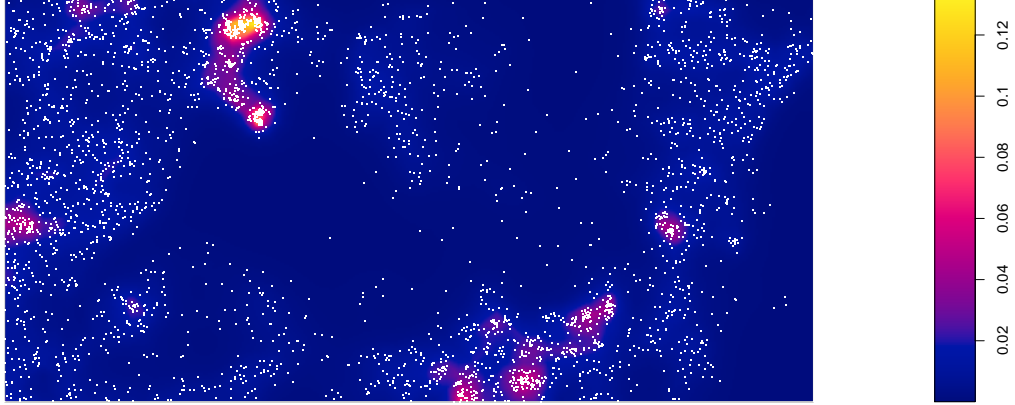


Figure 12. Posterior intensity surface in events per square meter, calculated using the piecewise linear approximate covariate surfaces and the posterior means of the intercept, coefficients, and GP.

Gridded Pearson Residuals

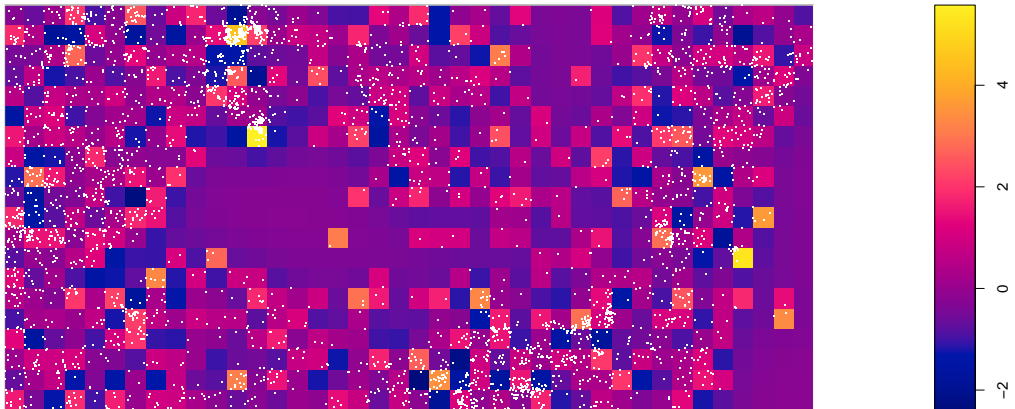


Figure 13. Pearson residuals calculated on a grid.