Log-Gaussian Cox processes and sampling paths:

towards optimal design

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Abstract

 $Goal\ of\ this\ paper\ (placeholder\ abstract-add\ some\ results\ when\ available).\ \ \text{Eval-}$

uate a wide variety of path designs in terms design-based heuristics and model-

based criteria for spatial prediction using Bayesian LGCP models. Identify

promising path designs. Illuminate any relationships among design characteris-

tics and predictive criteria that will be helpful for constrained optimization.

 ${\it Keywords:} \ \ {\rm log\text{-}Gaussian\ Cox\ process,\ optimal\ sampling,\ model\text{-}based\ design,}$

spatial sampling design

1. Introduction

Spatial point process models have long been considered generally infeasi-

ble because of their computational demands, but recent advances in Bayesian computing have made the Log-Gaussian Cox process an attainable model in

practice (Rue et al., 2009; Lindgren et al., 2011; Illian et al., 2012; Simpson

practice (roac or an, 2000, Emagren or an, 2011, Iman or an, 2012, Simpson

et al., 2016). In some applications, the entire point pattern is not fully observed

due to variable sampling effort. This is referred to as a degraded point pat-

tern (Chakraborty et al., 2011) and it is relatively simple to accommodate variable

sampling effort in these models using modern Bayesian computing tools (Yuan

et al., 2017). However, the literature on optimal sampling for spatial point process models is in its infancy (Liu and Vanhatalo, 2020). In this article, we

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present a variety of sampling path designs and assess their optimality for LGCP models.

Point pattern data are routinely collected in species distribution studies and ordnance response projects. These applications may use quadrat sampling or line-transect sampling, with transect sampling being more common. When the objective is mapping where events occur in space, various spatial mapping procedures have been used. Traditionally these have involved aggregating the data to grid cell counts or computing moving averages. Aggregation has the downside of introducing arbitrary structure into the data by the choice of gridding scheme or averaging window, and requires uneccessary computation effort (Simpson et al., 2016). Software is now available to fit spatial point process models to data acquired via distance sampling and simultaneously estimate the detection function (Johnson et al., 2014; R Core Team, 2019).

In ecological settings, sampling plans are often designed around the goal of estimating total abundance. Ordnance response surveys are typically designed with the objective of detecting (but not necessarily mapping) intensity hotspots (USACE, 2015; Flagg et al., 2020). However, to our knowledge, there has been very little work done in deciding where to collect data when the goal is to map the intensity using a spatial point process model. In this paper, we compare a variety of path design schemes with respect to a suite of model-based and design-based criteria for simulated point pattern data.

1.1. Spatial design

Design-based sampling. Most classical sampling work has been done for points or small quadrats approximated as points, rather than paths. Space-filling criteria may be good starting points (Borkowski and Piepel, 2009). Latin hypercube sampling has space-filling properties McKay et al. (1979); Husslage et al. (2011).

Space-filling curves. Used in design of dense or stretchable circuits (Ogorzałek, 2009; Ma and Zhang, 2016) and high-dimensional data visualization in bioinfor-

matics (Anders, 2009). Peano curve is very flexible for filling irregular shapes (Fan et al., 2014).

Space-filling curves are one-dimensional paths constructed iteratively; as the number of iterations goes to infinity, the limiting path has nonzero area and actually fills the space (Sagan, 1994). For applications we stop after a finite number of iterations. The Hilbert curve is fast and simple to construct.

Model-based spatial design. Regularity is optimal for spatial prediction but randomness and a variety of interpoint distances are best for parameter estimation (Diggle and Lophaven, 2006). Inhibitory plus close pairs is a good compromise (Chipeta et al., 2017).

50 1.2. Paths as sampling designs

While some ideas about the characteristics of a good point design apply to paths, creating an optimal path design is not as simple as connecting the points of a point design with line segments. There are many ways to connect points into a path, so optimal design criteria must apply to the whole path and not only to the waypoints.

Pollard et al. (2002) adaptively zigzagged their line transects in a species abundance survey.

The Visual Sample Plan software includes features to create systamatic transect plans and augment plans with additional transects in regions lacking spatial coverage (Matzke et al., 2014). It helps the user choose the transect spacing to maximize the probability of detecting the presence of a hotspot of specified size and intensity. However, it does not employ criteria to optimize spatial prediction.

Liu and Vanhatalo (2020) used narrow quadrats (swaths along line-transects)
as their sampling units. The transects were short relative to the size of the study
region and not connected into a path.

2. Materials and methods

With an eye toward practical considerations of data collection, we present criteria to compare sampling strategies that impact LGCP estimates. We compare plans with fixed path lengths that avoid sharp turns. Data collection equipment (e.g. metal detectors) may have limited mobility, requiring minimizing the number or angle of turns. The criteria that we evaluate are mean squared prediction error (MSPE) and posterior prediction variance of the Gaussian process.

2.1. Sampling schemes

These are the focus, move them earlier?

2.1.1. Parallel line transects

Parallel straight-line transects are common in ordnance response studies and in ecological studies using distance sampling. Systematic designs are common because they provide good spatial coverage in the sense that any point in the study region has a known maximum distance from the path. For point designs, systematic designs are optimal for prediction, simple random samples are optimal for estimation, and inhibitory with close pair designs are becomming a popular compromise. We adapt all of these to the parallel line transect setting. We use line transects running north-south, with three ways of choosing the horizontal coordinate: simple random sample (SRS), systematic with a random starting point and even spacing, inhibitory plus close pairs. Figure 1 shows an example of each scheme with 25 transects.

We should note that our we will be using an isotropic covariance function in the model. If an anisotropic model is used, parallel transects may need to be augmented with perpendicular transects so that the observed data provide the full range of pairwise distances in other directions.

2.1.2. Parallel serpentine transects

One simple way to observe a greater variety of locations and different directions is to add lateral zigzags to transects. We include alternate right and left turns at right angles to create serpentine transects. This could decrease prediction variance because many points in the study area will be closer to the path than they would be under a similar plan of line transects. (Check the math—is this true? And does similar mean same length or same number of transects?)

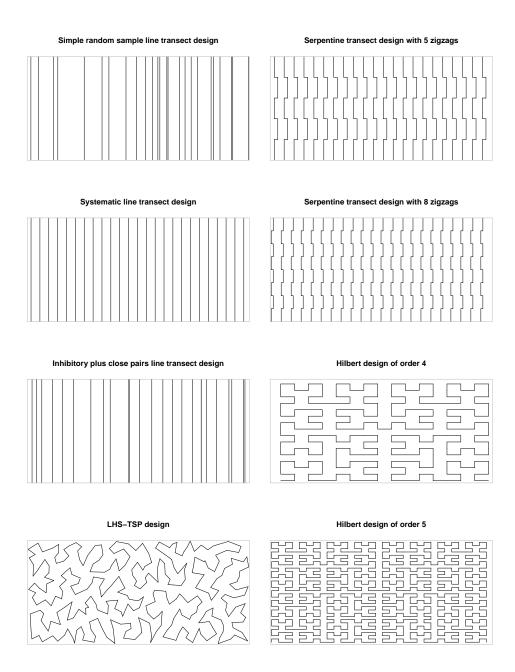


Figure 1: Examples of plans from six design schemes. Left, top to bottom: three different parallel line transect schemes with the same number of transects, and a shortest path through a Latin hypercube sampling design. Right: two serpentine transect plans and two Hilbert curves. Except for the Hilber curve of order 5, all of these plans have approximately the same total length.

They will also improve estimation of covariance parameters in the presence of anisotropy.

2.1.3. Latin hypercube sampling

Random Latin hypercube sampling (LHS) produces a design that spreads discrete points through a (potentially high-dimensional) design space, ensuring that the full range of each dimension is included while remaining balanced and keeping the number of points small. This is done by partitioning each dimension into a specified number k of intervals (thus stratifying the design space into k^d cells), selecting a Latin hypercube design to determine which cells will contain a design point, and then drawing each design point from a uniform distribution over its cell. In two dimensions, this scheme produces point designs with good spatial coverage properties. We use the LHS design as waypoints for a path. Because distance is typically a criterion to be minimized, we treat this as a traveling salesperson problem (TSP) and use the shortest path through the waypoints as our design. This LHS-TSP scheme produces paths that have many sharp corners but leaves few large voids (example in Figure 1). A downside of this design scheme is that the length cannot be specified directly, and only certain distances are possible depending on the number of bins used.

Waypoints are generated by the 1hs R package Carnell (2020) and connected into a the shortest path by the TSP package (Hahsler and Hornik, 2020).

2.1.4. Space-filling curves

Hilbert curve generated by HilbertVis package (Anders, 2009). This is a deterministic design, so a random offset is added.

2.2. Model fitting

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INLA (Rue et al., 2009), SPDE (Lindgren et al., 2011), off-grid (Simpson et al., 2016), Flagg and Hoegh (2020) knock on wood

5 3. Simulation Study

We consider a fictitious site \mathcal{R} with the simple shape of a 1500 unit by 700 unit rectangle. In this site, we will simulate two data generating models meant to produce random intensity functions with with hotspots. First, a LGCP with latent GP mean $\mu = \log(250/|\mathcal{R}|)$ and a Matérn covariance with $\nu = 1$, $\sigma = 2$, and range = 200. This model produces relatively unstructured hotspots due to large variability in the GP.

Second, the superposition of a two-stage cluster process superposed and a LGCP. The cluster process (a Neyman-Scott or, more specifically, a Thomas process) is constructed as follows. The number of clusters is Poisson with mean 3. The number of events per cluster is Poisson with mean 200. The cluster centers distributed uniformly over \mathcal{R} . Events come from a bivariate normal distribution with mean equal to the cluster center and variance $\Sigma = \tau^2 \mathbf{I}$, $\tau = 50$. The LGCP has $\mu = \log(250/|\mathcal{R}|)$ and Matérn covariance with $\nu = 1$, $\sigma = 1$, and range = 200. This model is based upon the typical conceptual model of a firing range, with a background process (represented by the LGCP) and a small number of higher-intensity foreground clusters containing the events of interest.

Path design schemes:

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- Simple random sample of north-south line transects
 - Number of transects = 10, 25, 50, 70
 - Expect high variance, large prediction error in big gaps.
- Systematic sample of north-south line transects
 - Number of transects = 10, 25, 50, 70
 - Uniformly distributed starting point
 - Constant spacing
 - Expect low bias and ok variance, can miss structures at certain sizes,
 may not have best space-filling properties.

- Inhibitory plus close pairs sample of north-south line transects
 - Total number of transects (including pairs) = 10, 25, 50, 70
 - Number of pairs = 0.1, 0.2 times the total number of transects (rounded up or down to nearest whole number)
 - Pairs uniformly distributed within radius of primaries, max pair radius =
 1500/total number of transects
 - Position of primaries generated from a 1-dimensional Strauss process with $\gamma = 0.05$
 - A compromise between SRS and systematic in every way.
 - For the inhibitory plus close pairs designs, we vary the numbers of paired and unpaired transects. The total number of transects is 10, 25, 50, or 70, with 10% and 20% of the transects (rounded to the nearest integer) as redundant members of a pair. The remaining primary transects were placed according to a one-dimensional Strauss process with $\gamma = 0.05$ and a radius of 80. Then each redundant transect was randomly paired to a primary transect, an placed within an 80 unit radius of the primary transect according to a uniform distribution.
- Systematic sample of north-south serpentine transects
 - Number of transects = 7, 22, 47, 67
 - Uniformly distributed starting point
 - Constant spacing

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- Number of zigzags = 5.8
- Horizontal zigzag distance set so that the total horizontal distance traveled equals 2100 units (the length of 3 non-zigzag line transects)
- Expect better space-filling properties than line-transect designs, lower bias/variance farther from path, would be better at estimating anisotropic covariance than line-transects.

- We generate designs with 7, 22, 47, and 67 serpentine transects. We vary the complexity of the serpentines by using versions with 5 and 8 zigzags. We define a zigzag as a single north-south segment or a pair of connected north-south and east-west segments. The lengths of the east-west segments are set so that the total distance equals the total distance of a line transect design with three more transects.

• Latin Hypercube Sampling waypoints

- Number of bins = 50, 300, 1200, 2400
- Expect low bias/variance per unit distance traveled, many sharp corners, some big open areas.

• Hilbert curve

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- Order = 3, 4, 5, 6
- Created in square and then scaled to fit in \mathcal{R}
- A uniform random offset added equal to spacing between segments
- Expect good space filling, good bias and variance, lots of short segments.

(Probably should move explanations of schemes to appendix.)

100 designs from each scheme. All events within a 2 unit radius of the path are observed. Whole experiment repeated for 5 realizations from each data generating model.

Model:

- **X** is a Poisson process on \mathcal{R} with intensity $\lambda(u)$
- $\log[\lambda(u)] = \mu + \mathbf{e}(u)$
- $\mu \sim \text{Unif}(-\infty, \infty)$
- e is a Gaussian process with mean 0 and a Matérn covariance function with fixed $\nu=1$

Mesh with dual colored by node weight

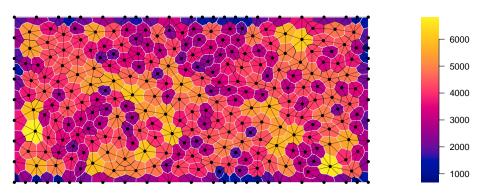


Figure 2: Illustration of the mesh and associated numerical integration scheme used to approximate the latent GP.

- PC prior on σ and ρ with $Pr(\sigma > 3) = 0.1$ and $Pr(\rho < 100) = 0.1$ (Fuglstad et al., 2019; Simpson et al., 2017)
- SPDE approach of Lindgren et al. (2011) using mesh in Figure 2
- Likelihood factorization of Simpson et al. (2016)

210 4. Results

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look at examples of designs that minimize each criterion look at examples of designs along the Pareto front

5. Discussion

discuss starting points for optimization and sequential design

practical issue: path will be smoothed, no instantaneous direction changes at corners, equipment may have limitations which is why we looked at number and distribution or turn angles

could incorporate turns into loss function or use multi-objective optimization (Lark, 2016)

6. Conclusions

Appendix A. Notation and Terminology

- process defined on $\mathcal{D} \subset \mathbb{R}^d$, domain of the intensity function, in this manuscript d=2
- observation window $S \subset \mathcal{D}$
- define three regions:

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- the domain \mathcal{D} over which the process mathematically operates
- the study region \mathcal{R} over which inferences are desired
- the observed/sampled observation window \mathcal{S}
- 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"
- \mathcal{D} can be bounded or unbounded (often equal to \mathbb{R}^d), \mathcal{S} practically always bounded, \mathcal{R} bounded or unbounded depending on application and inferential goals
- the "fully surveyed" (censused) situation is S = R
- survey path \mathcal{P} is a one-dimensional subset of \mathcal{R}
 - set of one or more sequences of waypoints connected by line segments
 - ${\mathcal S}$ is the set of all points within a fixed (and assumed known) radius of ${\mathcal P}$
 - **X** point process on \mathcal{R} , $\mathbf{x} = \{x_1, \dots, x_n\}$ realized point pattern
 - $-\mathbf{X}_{\mathcal{S}} = \mathbf{X} \cap \mathcal{S}$ the restriction of \mathbf{X} to \mathcal{S} , $\mathbf{x} = \mathbf{X} \cap \mathcal{S}$ the realized observeable point pattern
 - point $x \in \mathbf{x}$ called an event
 - intensity function $\lambda(u)$

- types of "points" in space:
 - -x event in the point pattern
 - s numerical integration node
 - u arbitrary location in \mathcal{D} used to index intensity function and predictors
- z(u) a column vector of covariates/predictors at u (not used in this manuscript)
- "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. marks, pseudodata
- distance sampling fits into the framework with expansion of notation to include a (nontrivial) detection function and differentiate between the observed and observable point patterns

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