SPATIAL FACTOR MODELS FOR HIGH-DIMENSIONAL AND LARGE SPATIAL DATA: AN APPLICATION IN FOREST VARIABLE MAPPING

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Abstract: Gathering information about forest variables is an expensive and arduous activity. Therefore, directly collecting the data required to produce high-resolution maps over large spatial domains is infeasible. Next-generation collection initiatives for remotely sensed light detection and ranging (LiDAR) data are specifically aimed at producing complete-coverage maps over large spatial domains. Given that Li-DAR data and forest characteristics are often strongly correlated, it is possible to use the former to model, predict, and map forest variables over regions of interest. This entails dealing with high-dimensional ($\sim 10^2$) spatially dependent LiDAR outcomes over a large number of locations ($\sim 10^5 - 10^6$). With this in mind, we develop the spatial factor nearest neighbor Gaussian process (SF-NNGP) model, which we embed in a two-stage approach that connects the spatial structure found in LiDAR signals with forest variables. We provide a simulation experiment that demonstrates the inferential and predictive performance of the SF-NNGP, and use the two-stage modeling strategy to generate complete-coverage maps of the forest variables, with associated uncertainty, over a large region of boreal forests in interior Alaska.

Key words and phrases: Forest outcomes, LiDAR data, nearest neighbor Gaussian processes, spatial prediction.

1. Introduction

Strong relationships between remotely sensed light detection and ranging (LiDAR) data and forest variables have been documented in the literature (Asner et al. (2009); Babcock et al. (2013); Næsset (2011)). When used in forested settings, LiDAR data provide a high-dimensional signal that characterizes the

vertical structure of the forest canopy at point-referenced locations. Traditionally, LiDAR data acquisition campaigns have sought complete-coverage at a high spatial resolution over relatively small spatial domains, resulting in a fine grid of point-referenced LiDAR signals. In such settings, the link between the LiDAR data and the forest variable measurements on sparsely sampled forest inventory plots has been exploited to create high-resolution complete-coverage predictive maps of the forest variables. Commonly, this link is established by first extracting the relevant features of the high-dimensional LiDAR signals using a dimension-reduction step (Babcock et al. (2015); Junttila and Laine (2017)). Then the LiDAR features are used as predictors in a regression model to explain the variability in the spatially coinciding forest variable outcomes. Lastly, the model is applied to predict the forest outcomes at all locations across the domain where LiDAR signals have been observed.

Considerably more ambitious next-generation LiDAR collection initiatives, such as ICESAT-2 (ICESat-2 (2015)), Global Ecosystem Dynamics Investigation LiDAR (GEDI) (GEDI (2014)), and NASA Goddard's LiDAR, Hyper-Spectral, and Thermal imager (G-LiHT) (G-LiHT (2016)), seek to quantify and map forest variables over vast spatial extents. To fulfill their goals in a cost-effective manner, these data-gathering programs do not collect LiDAR data over the entire domain. Instead they sparsely sample locations across the domain extent and over forest inventory plots (i.e., where forest variables have been measured). While generating complete-coverage high-resolution maps of forest outcomes remains the primary intended use of these data, there is also interest in creating maps of LiDAR data over nonsampled locations and assessing the spatial dependence within and among LiDAR signals.

Our motivating application focuses on forest variable prediction and mapping in the boreal forests of interior Alaska using sparsely sampled LiDAR and forest variable measurements. Within these regions, acquiring complete-coverage LiDAR data is prohibitive from a cost perspective (Andersen, Strunk and Temesgen (2011); Bolton, Coops and Wulder (2013); Nelson et al. (2012)). Because generating complete-coverage maps of forest variables (and perhaps LiDAR signals) is still the goal, the sparsely sampled LiDAR must be leveraged to inform the forest variable predictions. One attractive solution is to move the LiDAR predictor variables to the left-hand side of the regression and then to model them jointly with the forest outcomes. When the number of LiDAR and forest variables is small, such joint models are possible via linear models of coregionalization; for example, see, Babcock et al. (2018) and Finley, Banerjee and Cook

(2014). Alternatively, if the LiDAR signal is high-dimensional, but observed at a small number of locations, reduced-rank models can be employed. For example, Banerjee et al. (2008), Ren and Banerjee (2013), and Finley et al. (2017) applied a reduced-rank *predictive process* modeling strategy to analyze similar high-dimensional data. However, such approaches cannot scale to data sets with tens of thousands of locations and can yield poor predictive performance (Stein (2014)).

Models able to handle high-dimensional signals observed over a large number of locations and capable of estimating within and among location dependence structures are needed. Recent modeling developments reviewed in Heaton et al. (2017) and Banerjee (2017) highlight several options for robust and practical approximation of univariate Gaussian process (GP) models. A subset of these models can be easily extended to accommodate relatively small multivariate response vectors (five or less) for example, see (Datta et al. (2016a)). Nevertheless, for our particular application, we require an approach that can cope with both the high-dimensional LiDAR measurements, ~50 outcomes at a location, and use the large collection of observed locations.

The nearest neighbor Gaussian process (NNGP) developed in Datta et al. (2016a), Datta et al. (2016b), and Datta et al. (2016c) can be used with a large number of locations, because its scalability is not mediated by the number of observed locations, but rather by the size of the nearest neighbor sets considered—a quality that yields minimal storage and computational requirements. These models belong to the class of methods that induce sparsity on the spatial precision matrix, and exploit the natural representation of sparsity provided by graphical models (Lauritzen (1996); Murphy (2012)) to build a sparse GP that accurately approximates the original dense GP.

To tackle the high-dimensional LiDAR data set, we develop a Bayesian NNGP spatial factor model (SFM), referred to as the SF-NNGP. Following Christensen and Amemiya (2002), Hogan and Tchernis (2004), and Ren and Banerjee (2013), the SFM structure enables approximating the dependence between multivariate (spatially dependent) outcomes through a lower-dimensional set of spatial factors, alleviating the difficulty of dealing directly with high-dimensional outcomes. The SF-NNGP allows us to model and map the LiDAR signals on both observed and unobserved locations, and, conditioning on the LiDAR spatial signatures, we can similarly map the forest variables over the entire spatial domain of interest. Furthermore, using a Bayesian approach for model fitting enables us to equip the derived estimates and predictions with associated measures of uncer-

tainty, an essential requirement of many high-profile initiatives. Our methods are fully implemented in C++, using BLAS (Blackford et al. (2001); Zhang (2016)) to leverage efficient multiprocessor matrix operations and openMP (Dagum and Menon (1998)) to improve key steps of the algorithm through parallelization.

The structure of the remainder of this paper is as follows. Section 2 introduces the Bonanza Creek data set. In Section 3, we formulate the proposed hierarchical Bayesian modeling strategy. Section 4 presents an analysis of a synthetic data set to validate the performance of the SF-NNGP model. Using the available LiDAR and forest inventory data, in Section 5, we develop and validate a predictive model for the forest variables. We close by providing insights, recommendations, and directions for future in Section 6.

2. Data Description

The Bonanza Creek Experimental Forest (BCEF) is a Long-Term Ecological Research (LTER) site consisting of vegetation and landforms typical of interior Alaska. The BCEF is 21,000 ha and includes a section of the Tanana River floodplain along the southeastern borders (Bonanza Creek LTER (2016)). Figure 1 shows the location and extent of the BCEF data detailed in this section.

Forest variables were collected on 197 plots in 2014 using the USDA Forest Service Forest Inventory and Analysis Program protocol (Bechtold and Patterson (2005)). We consider three forest variables commonly used by forest professionals to make management decisions: above-ground biomass (AGB); tree density (TD); and basal area (BA). The AGB for individual trees was estimated using the Component Ratio Method described in Woodall et al. (2015). The TD for a plot is expressed in thousands of trees per hectare. The BA for a plot is the sum of the individual trees' cross-sectional areas in m^2 at breast height, scaled to a per hectare basis.

In the summer of 2014, LiDAR data were collected using a flight-line strip sampling approach using NASA Goddard's G-LiHT sensor (Cook et al. (2013)), which is a portable multisensor system that accurately characterizes complex terrain and the vertical distribution of canopy elements (Jakubowski, Guo and Kelly (2013); White et al. (2013)). Point cloud information was summarized to a 13×13 m grid cell size to approximate field plot areas. Over each grid cell, psuedowaveforms were generated by calculating the LiDAR return count densities for .5 m height bins between 0 and 28.5 m (i.e., 57 LiDAR outcomes per location). The LiDAR return count density for height bin l is defined as the number of returns

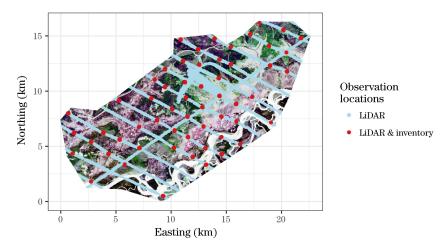


Figure 1. Bonanza Creek Experimental Forest extent with color enhanced Landsat image and locations where the LiDAR signals were measured (LiDAR in the legend) and locations where both LiDAR signals and forest variables were measured (LiDAR & inventory in the legend).

in height bin l divided by the total number of LiDAR returns over the grid cell. Identical LiDAR psuedo-waveforms were obtained using point clouds extracted over each field plot. G-LiHT data for the study area are available online at https://gliht.gsfc.nasa.gov. For this analysis, 50,197 LiDAR observations were used for model-fitting.

A Landsat 8 top of atmosphere (TOA) reflectance product was procured for the BCEF area for June of 2015. The June 2015 image was preferred to the June 2014 image owing to the excessive cloud cover in the 2014 image. A tasseled cap transformation was applied to the raw Landsat 8 TOA reflectance bands to obtain *brightness*, *greenness*, and *wetness* tasseled cap indices (Baig et al. (2014)). These indices are used as covariates in the subsequent analysis.

Further details on the data set and our analysis are provided in Section 5.

3. Modeling Strategy

Our goal is to model and generate uncertainty-equipped predictions of forest variables using information contained in LiDAR signals. Consider a LiDAR signal, $\mathbf{z}(\cdot)$, observed at a finite collection of locations, $\mathcal{T}_z = \{\mathbf{s}_1, \dots, \mathbf{s}_{n_z}\}$, and a set of forest outcomes, $\mathbf{y}(\cdot)$, observed at locations in the set $\mathcal{T}_y = \{\mathbf{r}_1, \dots, \mathbf{r}_{n_y}\} \subset \mathcal{T}_z$. Furthermore, let $\mathcal{T}_\emptyset = \{\mathbf{t}_1, \dots, \mathbf{t}_{n_\emptyset}\}$ denote a set of locations where neither LiDAR signals nor forest outcomes are available, but where predictions are of interest.

Thus, the set of locations where both LiDAR and forest outcomes are mapped corresponds to $\mathcal{T} = (\mathcal{T}_z \cup \mathcal{T}_{\emptyset})$, with $\mathcal{T} \subset \mathcal{D} \subset \mathbb{R}^2$, where \mathcal{D} is the spatial domain of interest. Note that although $\mathbf{z}(\cdot)$ and $\mathbf{y}(\cdot)$ are "observed" at locations in \mathcal{T}_z and \mathcal{T}_y , respectively, we allow for missing values that are to be imputed in these sets. We make this distinction because locations where imputation is performed are part of the model fitting, whereas for locations in \mathcal{T}_{\emptyset} , predictions are drawn ex post facto from the posterior predictive distribution; see Section 3.4.

The LiDAR signals are high-dimensional vectors of measurements in \mathbb{R}^{h_z} , whereas the forest outcomes are relatively small-dimensional vectors (i.e., $h_y << h_z$), assumed to have support on \mathbb{R}^{h_y} . The forest outcomes and LiDAR signals are strongly dependent on each other; LiDAR signals vary with the composition of a forest, and, as a plethora of examples in the literature have demonstrated (Ene et al. (2018); Finley et al. (2014); Nelson et al. (2017)), variability in forest outcome variables can be partially explained by LiDAR characteristics.

3.1. Linking the LiDAR and forest inventory data

We seek to connect the forest outcomes and LiDAR signals as a two-step process. First, we formulate a generative model to extract the spatial signature from the LiDAR data at locations in \mathcal{T}_z , which can also be used to interpolate LiDAR signals in \mathcal{T}_{\emptyset} . Along with other spatially referenced predictors, the LiDAR spatial signatures for locations in \mathcal{T}_y are used as predictors to build the model for the forest outcomes. Moreover, a component that captures the spatial variation exclusive to the forest outcomes can also be specified, if required. For $\mathbf{s} \in \mathcal{D}$, this two-stage model is given by

Stage 1:
$$\mathbf{z}(\mathbf{s}) = \mathbf{X}_z(\mathbf{s})'\beta_z + \mathbf{w}^*(\mathbf{s}) + \varepsilon_z(\mathbf{s}),$$
 (3.1)

Stage 2:
$$\mathbf{y}(\mathbf{s}) = \mathbf{X}_{y}(\mathbf{s})'\beta_{y} + \Upsilon \mathbf{w}^{*}(\mathbf{s}) + \mathbf{v}^{*}(\mathbf{s}) + \varepsilon_{y}(\mathbf{s}).$$
 (3.2)

Note that the influence of $\mathbf{z}(\mathbf{s})$ over $\mathbf{y}(\mathbf{s})$ in (3.2) is exerted solely through its spatial component, $\mathbf{w}^{\star}(\mathbf{s})$. There are several arguments in favor of this approach, as opposed to substituting $\mathbf{z}(\mathbf{s})$ or $\boldsymbol{\mu}_z(\mathbf{s}) = \mathbf{X}_z(\mathbf{s})'\boldsymbol{\beta}_z + \mathbf{w}^{\star}(\mathbf{s})$ as covariates directly into (3.2). Among these, and most importantly for our setting, $\mathbf{z}(\mathbf{s})$, $\boldsymbol{\mu}_z(\mathbf{s})$ and $\mathbf{w}^{\star}(\mathbf{s})$ are all high-dimensional objects. Using $\mathbf{w}^{\star}(\mathbf{s})$ reduces the dimensionality of the problem by casting it under the factor model structure, as shown in Section 3.2. In addition, the elements within $\mathbf{z}(\mathbf{s})$ are strongly correlated, hence, multicollinearity issues would arise if it was included directly in (3.2).

In (3.1) and (3.2), the terms $\mathbf{X}_z(\mathbf{s})'\boldsymbol{\beta}_z$ and $\mathbf{X}_y(\mathbf{s})'\boldsymbol{\beta}_y$ capture large-scale variation. For $\kappa \in \{z, y\}$, $\mathbf{X}_{\kappa}(\mathbf{s})'$ represents a fixed $h_{\kappa} \times p_{\kappa}$ block-diagonal matrix of

spatially referenced predictors, where $p_{\kappa} = \sum_{j=1}^{h_{\kappa}} p_{\kappa,j}$, having as its jth diagonal block the length- $p_{\kappa,j}$ vector $\mathbf{x}_{j}^{\kappa}(\mathbf{s})'$. The length- p_{κ} vector $\boldsymbol{\beta}_{\kappa}$ corresponds to the regression coefficients associated with $\mathbf{X}_{\kappa}(\mathbf{s})'$. The vectors $\mathbf{w}^{\star}(\mathbf{s})$ and $\mathbf{v}^{\star}(\mathbf{s})$ are h_{z} - and h_{y} -dimensional zero-centered stochastic processes over \mathcal{D} , respectively. The process $\mathbf{w}^{\star}(\mathbf{s})$ captures the spatial variation of $\mathbf{z}(\mathbf{s})$, and $\mathbf{v}^{\star}(\mathbf{s})$ synthesizes additional spatial variation in the forest outcomes. The $h_{y} \times h_{z}$ matrix Υ connects the spatial information extracted from the LiDAR model into the forest outcomes model. The vectors $\boldsymbol{\varepsilon}_{z}(\mathbf{s}) \sim N_{h_{z}}(\mathbf{0}, \Psi_{z})$ and $\boldsymbol{\varepsilon}_{y}(\mathbf{s}) \sim N_{h_{y}}(\mathbf{0}, \Psi_{y})$ represent uncorrelated random errors (i.e., Ψ_{z} and Ψ_{y} are diagonal) at finer scales.

Implementing this modeling strategy directly is challenging owing to the high-dimensionality of the LiDAR signals ($h_z \sim 50$) and the massive number of spatially dependent observations ($n \sim 10^5$). Thus, it is impossible to attempt using common computing resources. In the following section, we formulate a viable alternative to models (3.1) and (3.2).

3.2. The spatial factor NNGP model

To make models (3.1) and (3.2) tractable with limited computing power, we combine a dimension-reduction approach and a sparsity-inducing technique. In particular, we introduce the SF-NNGP model, which brings together the SFM structure (Schmidt and Gelfand (2003); Finley et al. (2008); Zhang (2007); Ren and Banerjee (2013)) with NNGPs (Datta et al. (2016b,c,a)).

While the SFM structure enables the analysis of high-dimensional response vectors by using linear combinations of a relatively small number of independent stochastic processes, NNGPs make it possible to fit spatial process models when the number of spatial observations is particularly large. NNGPs approximate the parent (dense) GP using the natural representation of sparsity provided by graphical models (Lauritzen (1996); Murphy (2012)), by assuming conditional independence—where conditioning is on the nearest neighbors—with locations outside of the neighbor set. The result is a proper (but sparse) GP that accurately approximates the original dense GP. In contrast to other sparsity-inducing approaches, NNGPs allow for interpolation at unobserved locations and can be used to make full inference on model parameters, including the latent processes. Combining the SFM structure with NNGPs provides a methodology capable of coping simultaneously with high-dimensional response vectors and a large number of spatially dependent observations.

Under the traditional SFM structure, spatial dependence is introduced by defining the spatial process as $\mathbf{w}^*(\mathbf{s}) = \mathbf{\Lambda}\mathbf{w}(\mathbf{s}) \sim \mathrm{GP}(\mathbf{0}, \mathcal{H}(\cdot \mid \boldsymbol{\phi}))$, where $\mathbf{\Lambda}$

is a factor loadings matrix (commonly tall and skinny) and $\mathbf{w}(\mathbf{s})$ is a small-dimensional vector of independent spatial GPs, providing the nonseparable multivariate cross-covariance function given by

$$\mathcal{H}(\mathbf{h} \mid \boldsymbol{\phi}) = \operatorname{cov}(\boldsymbol{\Lambda} \mathbf{w}(\mathbf{s}), \boldsymbol{\Lambda} \mathbf{w}(\mathbf{s} + \mathbf{h}))$$
$$= \sum_{k=1}^{q_w} C_k(\mathbf{h} \mid \phi_k) \boldsymbol{\lambda}_k \boldsymbol{\lambda}'_k, \tag{3.3}$$

for locations $\mathbf{s}, \mathbf{s} + \mathbf{h} \in \mathcal{D}$. Here, $\mathcal{C}_k(\mathbf{h}|\phi_k)$ denotes a univariate parametric correlation function, and λ_k is the kth column of Λ . This cross-covariance matrix is induced by q-variate ($q \leq l$) spatial factors $\mathbf{w}(\mathbf{s})$ with independent components $w_k(\mathbf{s}) \sim \mathrm{GP}(0, \mathcal{C}_k(\cdot | \phi_k))$.

As such, models (3.1) and (3.2) can be reformulated as SF-NNGPs by characterizing the spatial processes $\mathbf{w}^{\star}(\mathbf{s})$ and $\mathbf{v}^{\star}(\mathbf{s})$ as

$$\mathbf{w}^{\star}(\mathbf{s}) = \mathbf{\Lambda}_z \mathbf{w}(\mathbf{s}) \text{ and } \mathbf{v}^{\star}(\mathbf{s}) = \mathbf{\Gamma} \mathbf{v}(\mathbf{s}),$$
 (3.4)

where the matrices $\mathbf{\Lambda}_z = ((\lambda_{hk}^{(z)}))_{h_z \times q_w}$ and $\mathbf{\Gamma} = ((\gamma_{lr}))_{h_y \times q_v}$ correspond to the factor loadings matrices, and the new spatial factors for $\mathbf{s} \in \mathcal{D}$ are given by

$$\mathbf{w}(\mathbf{s}) \sim \prod_{k=1}^{q_w} \text{NNGP}\left(0, \tilde{\mathcal{C}}(\cdot \mid \phi_k^w)\right), \text{ and}$$

$$\mathbf{v}(\mathbf{s}) \sim \prod_{r=1}^{q_v} \text{NNGP}\left(0, \tilde{\mathcal{C}}(\cdot \mid \phi_r^v)\right).$$

The expressions NNGP $\left(0, \tilde{\mathcal{C}}(\cdot \mid \phi_k^w)\right)$ and NNGP $\left(0, \tilde{\mathcal{C}}(\cdot \mid \phi_r^v)\right)$ denote the NNGPs derived from the parent processes $\operatorname{GP}(0, \mathcal{C}(\cdot \mid \phi_k^w))$ and $\operatorname{GP}(0, \mathcal{C}(\cdot \mid \phi_r^v))$, respectively. Here, $\mathcal{C}(\cdot \mid \phi)$ represents the spatial correlation function with spatial decay parameter ϕ . The factor model representation in (3.4) leads to a significant reduction in the dimensionality of the problem because the spatial factors $\mathbf{w}(\mathbf{s}) = (w_k(\mathbf{s}) : 1 \leq k \leq q_w)$ and $\mathbf{v}(\mathbf{s}) = (v_r(\mathbf{s}) : 1 \leq r \leq q_v)$ have dimensions $q_w << h_z$ and $q_v \leq h_y$, respectively.

Combining these elements, and letting $\mathbf{\Lambda}_y = \mathbf{\Upsilon} \mathbf{\Lambda}_z = ((\lambda_{lk}^{(y)}))_{h_y \times q_w}$, a computationally viable version of (3.1) and (3.2) is

Stage 1:
$$\mathbf{z}(\mathbf{s}) = \mathbf{X}_z(\mathbf{s})' \boldsymbol{\beta}_z + \boldsymbol{\Lambda}_z \mathbf{w}(\mathbf{s}) + \boldsymbol{\varepsilon}_z(\mathbf{s})$$
 (3.5)

Stage 2:
$$\mathbf{y}(\mathbf{s}) = \mathbf{X}_{u}(\mathbf{s})'\boldsymbol{\beta}_{u} + \boldsymbol{\Lambda}_{u}\mathbf{w}(\mathbf{s}) + \boldsymbol{\Gamma}\mathbf{v}(\mathbf{s}) + \boldsymbol{\varepsilon}_{u}(\mathbf{s}).$$
 (3.6)

In general, additional constraints are required for factor models to be identifiable (Anderson (2003)). Identifiability for SFMs can be achieved either by making the upper triangle of the loadings matrix equal to zero and its diagonal

elements all equal to one (Geweke and Zhou (1996); Lopes and West (2004); Aguilar and West (2010)), or, as in Ren and Banerjee (2013), by fixing the sign of one element in each column of the factor loadings matrix, while enforcing an ordering constraint among the spatial decay parameters of the univariate correlation functions. We choose to ensure rotation and scale identifiability by using the former approach.

With the SFM structure in place, introducing the NNGP reduces the expensive ($\sim n_z^3 q_w$ and $\sim n_y^3 q_v$) calculation required to invert the dense covariance matrices from the parent GPs by $n_z q_w$ and $n_y q_v$ parallel operations, each of order m^3 . Here, m is the number of neighbors considered for the NNGP, with $m << n_y \le n_z$. In simulations, Datta et al. (2016b) found that, in most cases, $10 \le m \le 20$ provides an excellent approximation to the parent process; thus, the number of operations required is nearly linear in n.

For completeness, additional details on SFMs, NNGPs, and the sampling algorithm are included in the online supplement. For a more thorough treatment of SFM's, refer to Ren and Banerjee (2013) and Genton and Kleiber (2015), and for NNGPs, refer to Datta et al. (2016c).

3.3. Prior specification and hierarchical formulation

Importantly, models (3.5) and (3.6) are fitted separately such that $\mathbf{w}(\mathbf{s})$ exclusively captures the spatial signal present in the LiDAR signals. However, using plug-in estimates for $\mathbf{w}(\mathbf{s})$ (e.g., the posterior means) in (3.6) disregards the uncertainty present in the LiDAR spatial signal. Thus, to propagate this uncertainty through the forest outcome predictions, at each iteration of the Markov Chain Monte Carlo (MCMC) algorithm for $\mathbf{y}(\mathbf{s})$, we draw a sample for $\mathbf{w}(\mathbf{s})$ ($\mathbf{s} \in \mathcal{T}_u$) MCMC samples obtained when fitting model (3.5).

As mentioned in the previous section, the stochastic processes that capture the spatial structure are assumed to follow NNGPs. Given that an NNGP is a proper GP, at a finite collection of locations, the NNGPs induce zero-centered multivariate normal priors, with covariance matrices given by $\tilde{\mathbf{C}}^{(w)}$ and $\tilde{\mathbf{C}}^{(v)}$, respectively. Additionally, we use suitably noninformative priors for all other parameters, thus providing a direct sampling strategy.

In particular, we assume that β is either flat or conjugate normal. The matrices Γ and Λ_z are constrained as described above, with elements below the diagonal assumed to be standard normal. All elements in Λ_y are also assumed to follow a standard normal distribution. The diagonal entries in Ψ_z and Ψ_y are assigned half-t priors. Lastly, we assume uniform priors for the elements

of the spatial decay vectors $\phi_w = (\phi_{w,k} : 1 \le k \le q_w)$ and $\phi_v = (\phi_{v,r} : 1 \le r \le q_v)$ in the interval $(-\log 0.05/\zeta_{\text{max}}, -\log 0.01/\zeta_{\text{min}})$, where ζ_{min} and ζ_{max} are the minimum and maximum distances, respectively, across all locations. Given that ϕ_z and ϕ_y are not conjugate with their corresponding likelihood, these are sampled using random walk Metropolis steps.

The joint posterior densities for the first and second stages of the algorithm are proportional to

Stage 1:

$$\pi(\boldsymbol{\phi}_{w}) \operatorname{N}_{n_{z}q_{w}}(\mathbf{w}_{\mathcal{T}_{z}}|\mathbf{0}, \tilde{\mathbf{C}}^{(w)}) \left(\prod_{k=1}^{q_{w}} \prod_{j>k}^{h_{z}} \operatorname{N}(\lambda_{jk}^{(z)}|0, 1) \right)$$

$$\times \pi(\boldsymbol{\beta}_{z}) \left(\prod_{j=1}^{h_{z}} \mathcal{IG}\left(\psi_{j}^{z} \middle| \frac{\nu}{2}, \frac{\nu}{a_{z,j}}\right) \mathcal{IG}\left(a_{z,j} \middle| \frac{1}{2}, \frac{1}{A^{2}}\right) \right)$$

$$\times \left(\prod_{\mathbf{s}_{i} \in \mathcal{T}_{z}} \operatorname{N}_{h_{z}}(\mathbf{z}(\mathbf{s}_{i})|\mathbf{X}_{z}(\mathbf{s}_{i})'\boldsymbol{\beta}_{z} + \boldsymbol{\Lambda}_{z}\mathbf{w}(\mathbf{s}_{i}), \boldsymbol{\Psi}_{z}) \right),$$

$$(3.7)$$

Stage 2:

$$\pi(\boldsymbol{\phi}_{v}) \operatorname{N}_{n_{y}q_{v}}(\mathbf{v}_{\mathcal{T}_{y}}|\mathbf{0}, \tilde{\mathbf{C}}^{(v)}) \left(\prod_{k=1}^{q_{w}} \prod_{j=1}^{h_{y}} \operatorname{N}(\lambda_{jk}^{(y)}|0, 1) \right) \left(\prod_{r=1}^{q_{v}} \prod_{j>r}^{h_{y}} \operatorname{N}(\gamma_{jr}|0, 1) \right)$$

$$\times \pi(\boldsymbol{\beta}_{y}) \left(\prod_{j=1}^{h_{y}} \mathcal{I}\mathcal{G}\left(\psi_{j}^{y} \middle| \frac{\nu}{2}, \frac{\nu}{a_{y,j}}\right) \mathcal{I}\mathcal{G}\left(a_{y,j} \middle| \frac{1}{2}, \frac{1}{A^{2}}\right) \right)$$

$$\times \left(\prod_{\mathbf{s}_{i} \in \mathcal{T}_{y}} \operatorname{N}_{h_{y}}(\mathbf{y}(\mathbf{s}_{i})|\mathbf{X}_{y}(\mathbf{s}_{i})'\boldsymbol{\beta}_{y} + \boldsymbol{\Lambda}_{y}\mathbf{w}(\mathbf{s}_{i}) + \boldsymbol{\Gamma}\mathbf{v}(\mathbf{s}_{i}), \boldsymbol{\Psi}_{y}) \right), \quad (3.8)$$
where $\mathbf{w}_{\mathcal{T}_{z}} = (\mathbf{w}(\mathbf{s}_{i})' : \mathbf{s}_{i} \in \mathcal{T}_{z})' \text{ and } \mathbf{v}_{\mathcal{T}} = (\mathbf{v}(\mathbf{s}_{i})' : \mathbf{s}_{i} \in \mathcal{T}_{y})', \text{ such that}$

where
$$\mathbf{w}_{\mathcal{T}_{z}} = (\mathbf{w}(\mathbf{s}_{i})' : \mathbf{s}_{i} \in \mathcal{T}_{z})'$$
 and $\mathbf{v}_{\mathcal{T}} = (\mathbf{v}(\mathbf{s}_{i})' : \mathbf{s}_{i} \in \mathcal{T}_{y})'$, such that
$$N_{n_{z}q_{w}}(\mathbf{w}_{\mathcal{T}}|\mathbf{0}, \tilde{\mathbf{C}}^{(w)}) = \prod_{\mathbf{s}_{i} \in \mathcal{T}_{z}} N_{q_{w}}(\mathbf{w}(\mathbf{s}_{i}) | \mathbf{B}_{i}^{(w)} \mathbf{w}_{N(i)}, \mathbf{F}_{i}^{(w)}), \text{ and}$$

$$N_{n_{y}q_{v}}(\mathbf{v}_{\mathcal{T}}|\mathbf{0}, \tilde{\mathbf{C}}^{(v)}) = \prod_{\mathbf{s}_{i} \in \mathcal{T}_{y}} N_{q_{v}}(\mathbf{v}(\mathbf{s}_{i}) | \mathbf{B}_{i}^{(v)} \mathbf{v}_{N(i)}, \mathbf{F}_{i}^{(v)}). \tag{3.9}$$

The expressions on the right-hand side of (3.9) result from the construction of the NNGP (see online supplement). For an m-neighbor NNGP, let $m_i = \min\{m, i-1\}$ denote the number of neighbors for location \mathbf{s}_i . The index set N(i) for location $\mathbf{s}_i \in \mathcal{T}_z$ contains its m_i nearest neighbors; thus, $\mathbf{w}_{N(i)}$ corresponds to the vector $(\mathbf{w}(\mathbf{s}_j)': \mathbf{s}_j \in N(i) \subset \mathcal{T}_z)'$. The neighbor set for $\mathbf{v}(\mathbf{s}_i)$ is defined

analogously. Letting $u \in \{w, v\}$, $\mathbf{B}_i^{(u)}$ denotes a $q_u \times m_i q_u$ block matrix, with the $q_u \times q_u$ diagonal blocks containing the kriging weights for the q_u spatial factors for each neighbor. In addition, $\mathbf{F}_i^{(u)}$ corresponds to the $q_u \times q_u$ diagonal matrix with the variances for the q_u spatial factors conditioned on the neighbor set N(i) (see Section S2 in the supplement for details on $\mathbf{B}_i^{(u)}$ and $\mathbf{F}_i^{(u)}$). Lastly, the parameters $\{a_{y,j}\}_{j=1}^{h_y}$ and $\{a_{z,k}\}_{k=1}^{h_z}$ complete the hierarchical representation of the half-t prior distribution assumed for ψ_j^y and ψ_k^z , respectively, and the hyperparameter A is simply chosen to be some large value (say, 100).

Owing to prior conjugacy, the full conditional densities for all parameters except ϕ_w and ϕ_v can be sampled using simple Gibbs steps. Further details on the sampling algorithm are deferred to the online supplement.

3.4. Imputation and prediction

As mentioned before, LiDAR signals are collected over the large spatial region \mathcal{T}_z , whereas forest outcome observations are confined to the smaller subset of locations in \mathcal{T}_y . Additionally, there are relevant out-of-sample locations where neither LiDAR nor forest outcomes are observed, \mathcal{T}_{\emptyset} . Finally, there are some locations within the corresponding reference sets \mathcal{T}_z and \mathcal{T}_y that have some or all missing outcomes. It is thus essential for this modeling effort to provide the means to accurately impute the missing values in \mathcal{T}_z or \mathcal{T}_y . Having the imputed data ultimately enables us to generate LiDAR predictions in \mathcal{T}_{\emptyset} and forest outcome predictions within $\mathcal{T}_{\emptyset} \cup (\mathcal{T}_z \setminus \mathcal{T}_y)$. Given the NNGP formulation, both the imputation and the out-of-sample prediction are remarkably inexpensive.

Imputation is straightforward. Let $\mathbf{s}_{\bullet} \in \mathcal{T}_z$ be a location where $\mathbf{z}(\mathbf{s}_{\bullet})$ is missing. Then, $\mathbf{z}(\mathbf{s}_{\bullet})$ is drawn as part of the sampling algorithm from $N_{h_z}(\mathbf{X}_z(\mathbf{s}_{\bullet})'\beta_z + \mathbf{\Lambda}_z\mathbf{w}(\mathbf{s}_{\bullet}), \mathbf{\Psi}_z)$, where $\mathbf{w}(\mathbf{s}_{\bullet})$ is sampled from the full conditional posterior density in Equation (S3.1) of the online supplement. For a missing value $\mathbf{y}(\mathbf{s}_{\bullet})$, where $\mathbf{s}_{\bullet} \in \mathcal{T}_y$, the procedure is analogous using the full conditional posterior for $\mathbf{v}(\mathbf{s}_{\bullet})$ and the likelihood for $\mathbf{y}(\mathbf{s}_{\bullet})$.

The procedure to predict a new LiDAR observation $\mathbf{z}(\mathbf{s}_{\circ})$, $\mathbf{s}_{\circ} \in \mathcal{T}_{\emptyset}$, begins by sampling the spatial factor $\mathbf{w}(\mathbf{s}_{\circ})$ from $N_{q_{w}}(\mathbf{B}_{\circ}^{(w)}\mathbf{w}_{N(\mathbf{s}_{\circ})}, \mathbf{F}_{\circ}^{(w)})$, with $\mathbf{B}_{\circ}^{(w)}$ and $\mathbf{F}_{\circ}^{(w)}$ defined as before. Note that the nearest neighbor set $N(\mathbf{s}_{\circ})$ is assumed to be in \mathcal{T}_{z} . Then, we draw $\mathbf{z}(\mathbf{s}_{\circ}) \mid \mathbf{z}_{\mathcal{T}_{z}}$ from $N_{h_{z}}(\mathbf{X}_{z}(\mathbf{s}_{\circ})'\boldsymbol{\beta}_{z} + \boldsymbol{\Lambda}_{z}\mathbf{w}(\mathbf{s}_{\circ}), \boldsymbol{\Psi}_{z})$. This is done by conditioning on the posterior samples of $\{\boldsymbol{\beta}_{z}, \boldsymbol{\Lambda}_{z}, \boldsymbol{\Psi}_{z}, \boldsymbol{\phi}_{w}\}$ obtained from the fitting algorithm.

To predict the forest outcomes $\mathbf{y}(\mathbf{s}_{\circ})$ at $\mathbf{s}_{\circ} \in \mathcal{T}_{\emptyset} \cup (\mathcal{T}_z \setminus \mathcal{T}_y)$, we first generate samples of $\mathbf{v}(\mathbf{s}_{\circ}) \sim \mathrm{N}_{q_v}(\mathbf{B}_{\circ}^{(v)}\mathbf{v}_{N(\mathbf{s}_{\circ})}, \mathbf{F}_{\circ}^{(v)})$. Given that $\mathbf{y}(\mathbf{s}_{\circ})$ depends on $\mathbf{w}(\mathbf{s}_{\circ})$, we

combine the posterior draws of $\{\beta_y, \Lambda_y, \Gamma, \Psi_y, \phi_v\}$ with those of $\mathbf{w}(\mathbf{s}_{\circ})$, obtained when fitting or predicting $\mathbf{z}(\mathbf{s}_{\circ})$, and draw predicted values for $\mathbf{y}(\mathbf{s}_{\circ}) \mid \mathbf{y}_{\mathcal{T}_y}$ from $N_{h_y}(\mathbf{X}_y(\mathbf{s}_{\circ})'\beta_y + \Lambda_y\mathbf{w}(\mathbf{s}_{\circ}) + \Gamma\mathbf{v}(\mathbf{s}_{\circ}), \Psi_y)$.

4. Simulation: Recovering Low-dimensional Structure

In the following simulation exercise we focus exclusively on the high-dimensional component (i.e., the first stage) of the model described above. The simulation below was devised to illustrate the ability of our approach to recover the true low-dimensional structure when the data are generated from a low-dimensional SFM with dense spatial factors.

We generate a synthetic data set for $h_z = 50$ outcomes in $n_z = 10,000$ locations from the spatial factor model $\mathbf{z}(\mathbf{s}) = \mathbf{X}_z(\mathbf{s})' \tilde{\boldsymbol{\beta}}_z + \tilde{\boldsymbol{\Lambda}}_z \tilde{\mathbf{w}}(\mathbf{s}) + \tilde{\boldsymbol{\varepsilon}}_z(\mathbf{s})$. Here, $\mathbf{X}_z(\mathbf{s})'$ is a 50×150 block-diagonal matrix of predictors, and $\tilde{\boldsymbol{\beta}}_z$ is the vector of regression coefficients, both defined as before. We consider the same three predictors for all outcomes. The spatial factors $\tilde{\mathbf{w}}(\mathbf{s}) \sim \prod_{k=1}^8 \mathrm{GP}(0, \mathcal{C}(\cdot | \tilde{\phi}_k^z),$ where $\mathcal{C}(\cdot | \tilde{\phi}_k^z)$ is an exponential correlation function with decay parameter $\tilde{\phi}_k^z$. Additionally, for identifiability, we assume that the 50×8 factor loadings matrix $\tilde{\boldsymbol{\Lambda}}_z$ has zeros in the upper triangle and ones along the diagonal. Finally, $\tilde{\boldsymbol{\varepsilon}}_z \sim N_{h_z} \left(\mathbf{0}, \tilde{\boldsymbol{\Psi}}_z\right)$, with $\tilde{\boldsymbol{\Psi}}_z = \mathrm{diag}(\tilde{\psi}_k^z : k = 1, \dots, 8)$.

We assess the ability of model (3.5) to recover the model parameters from the true data-generating process, impute missing outcomes, and predict at out-of-sample locations. The SF-NNGP model was fitted for $q_w \in \{3, 5, 8, 10\}$ spatial factors and m = 10 neighbors. Of the 10,000 locations, we assume all 50 outcomes to be missing in 200 locations chosen at random. These outcomes are to be imputed. Additionally, we use $n_0 = 500$ locations for out-of-sample prediction and model validation.

The first result worth highlighting is the gains in computational efficiency provided by the SF-NNGP. For this simulation exercise—a relatively computationally challenging problem—fitting the largest model considered (i.e., $q_w = 10$) with 50,000 MCMC iterations on a Linux server with an Intel i7 processor (two eight-core) and 16 GB of memory, the runtime was 4.88 hours. As shown below, the proposed approach is able to recover the true model parameters, accurately impute missing data, and generate precise predictions, all with suitable uncertainty estimates.

For all values of q_w , the SF-NNGP accurately recovered the regression coefficients $\tilde{\beta}_z$ for all predictors and responses (Figure 1 in the online supplement).

In contrast, the quality of the estimates for the small-scale variance components $\tilde{\psi}_k^z$'s was compromised when q_w was lower than the true number of spatial factors. This behavior is expected. For lower values of q_w , the ψ_k^z 's attempt to compensate for the additional signal that the spatial component with too few spatial factors is unable to capture (Figure 2 in the supplementary material). For $q_w = 8$ and $q_w = 10$, the coverage for $\tilde{\psi}_z$ was 88% and 84%, respectively, with all ψ_k^z close to $\tilde{\psi}_k^z$ with tight 95% credible sets.

When $q_w \neq 8$, the dimensions of the fitted Λ_z , ϕ_w , and $\mathbf{w}(\mathbf{s})$ do not match those of their analogs in the true model. Therefore, to assess the quality of the fit for the spatial signal for all values of q_w considered, we instead compare the fitted spatial component $\mathbf{w}^*(\mathbf{s}) = \Lambda_z \mathbf{w}(\mathbf{s})$, for $\mathbf{s} \in \mathcal{T}_z$, to that of the true model, given by $\tilde{\mathbf{w}}^*(\mathbf{s}) = \tilde{\Lambda}_z \tilde{\mathbf{w}}(\mathbf{s})$.

For all locations in \mathcal{T}_z , we calculate $\Delta(\mathbf{s}) = \mathbf{w}^*(\mathbf{s}) - \tilde{\mathbf{w}}^*(\mathbf{s})$ (fitted minus true spatial signal) for each MCMC draw of the parameters. For all $\mathbf{s} \in \mathcal{T}_z$, we obtained the median and 95% credible set for $\Delta(\mathbf{s})$. To facilitate visualization, in Figure 2, we show the results for only three responses, selected at random, from the 50 considered. The columns of each panel map the quantiles 2.5, 50, and 97.5 for $\Delta(\mathbf{s})$, with three locations (13, 23, and 48) plotted by row. The fitted spatial signal when $q_w \in \{3, 5\}$ only partially recovers the true signal, with coverages of 26.13% and 42.06%, respectively, for $q_w = 3$ and $q_w = 5$. When $q_w \in \{8, 10\}$, the recovery of the spatial signal is extremely accurate: over all responses, the coverage is 94.78% with $q_w = 8$, and 94.18% with $q_w = 10$.

In addition to the previous results, it is also encouraging to find that when the dimension of the SF-NNGP model matches that of the true model, both the factor loadings $(\tilde{\Lambda}_z)$ and the spatial decay parameters $(\tilde{\phi}_z)$ from the true spatial process can be recovered accurately (Figures 3 and 4).

Model performance in terms of the accuracy of imputation and prediction improves drastically as the number of factors approaches that of the true model; see Figures 5 and 6 in the online supplement.

Table 1 compares q_w in the SF-NNGP using different measures of out-of-sample predictive performance. In particular, the continuous rank probability score (CRPS) (Equation (21) in Gneiting and Raftery (2007)) and the root mean squared prediction error (RMSPE) (Yeniay and Goktas (2002)) favor the model with $q_w = 8$. The coverage of the 95% credible intervals of the predictions was close to the nominal value for all q_w ; however, the width of the interval rapidly decreases as q_w approaches the true number of spatial factors.

Both the fitted values for the spatial signals and the out-of-sample predictions

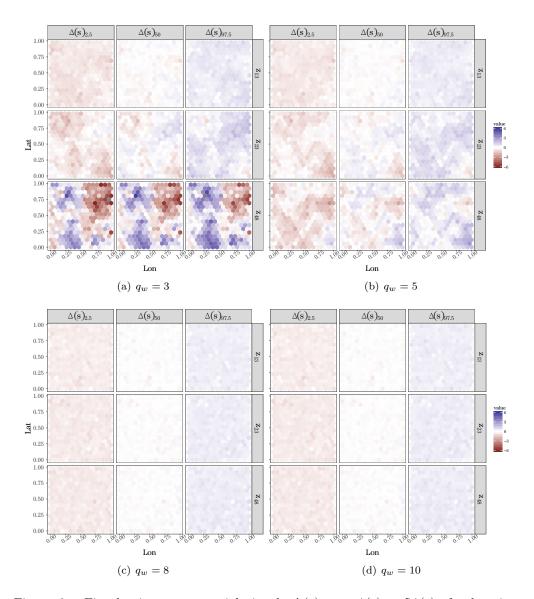


Figure 2. Fitted minus true spatial signal, $\Delta(\mathbf{s}) = \mathbf{w}^*(\mathbf{s}) - \tilde{\mathbf{w}}^*(\mathbf{s})$, for locations $\mathbf{s}_{13}, \mathbf{s}_{23}, \mathbf{s}_{48}$. From left to right, the columns in each panel show percentiles 2.5, 50, and 97.5 for $\Delta(\mathbf{s})$, respectively.

with $q_w=8$ and $q_w=10$ are practically indistinguishable from each other. Furthermore, the model with $q_w=8$ accurately recovers all of the true factor loadings (Figure 3). Interestingly, with $q_w=10$, a visual inspection of the estimates for columns 1 through 6 in Λ_z indicates that this model accurately

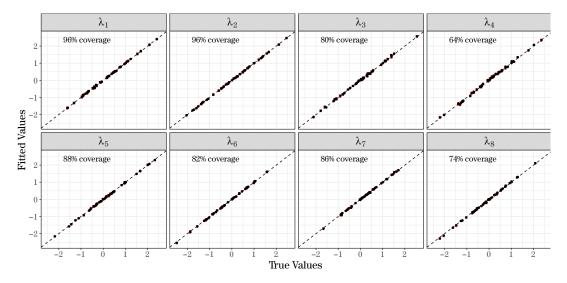


Figure 3. Fitted vs. true factor loadings matrix parameters (95% credible sets and medians) for $q_w = 8$.

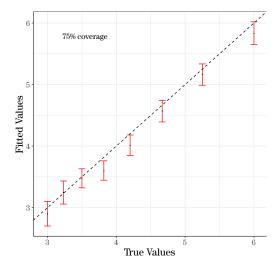


Figure 4. Fitted vs. true spatial decay parameters (95% credible sets and medians) for $q_w = 8$.

estimates the corresponding true parameter values (see Figure 4 in the online supplement). However, in this same model, the estimated parameter values in columns 7 and 8 of Λ_z display departures from their true values. Furthermore, the 95% credible sets for all unconstrained elements in the 9th and 10th columns of Λ_z contain zero (see Figure 3 in the online supplement). These results provide

Table 1. Out-of-sample prediction comparison across models with different numbers of spatial factors.

q_w	CRPS	RMSPE	95% Coverage	95% CI Width
3	0.85	1.61	95.82	6.14
5	0.67	1.28	95.43	4.79
8	0.45	0.83	94.78	3.10
10	0.45	0.83	94.84	3.10

guidance on the number of factors q_w to use. Because there is no gain in using the model with $q_w = 10$ over that with $q_w = 8$ in terms of predictive accuracy or parameter fit, the results favor the more parsimonious model of the two.

5. Modeling LiDAR Signals and Forest Structure

Our focus in the subsequent analysis is to assess and interpret the utility of SF-NNGP spatial factors to explain the variability in the three forest outcomes defined in Section 2, measured on the BCEF. Following the two-stage model developed in Section 3.2, we fit (3.5) using $q_w \in \{1, 2, 3, 4, 5, 6, 7, 8\}$ spatial factors and m = 10 neighbors to the BCEF LiDAR data comprising $n_z = 50,197$ signals, each of length $h_z = 57$. The model mean included only an intercept. The specification for the priors follows Section 3.3, with the support for elements in ϕ_w adjusted to match the BCEF spatial extent.

The $n_y = 197$ locations with $h_y = 3$ forest outcomes were used in the secondstage model (3.6). To more clearly interpret the spatial factors' ability to explain the variability in forest outcomes, we decided to avoid potential issues with spatial confounding (Hanks et al. (2015)) and set $\mathbf{v}(\mathbf{s})$ to zero. In practice, however, if our main objective is to maximize predictive performance, then this residual spatial random effect should likely be included in the model. In addition to the spatial factors, the second-stage model was informed by the three Landsat 8 tasseled cap predictor variables defined in Section 2, which, along with an intercept, were included in $\mathbf{X}_y(\mathbf{s})$. Importantly, these predictor variables are available across the entire BCEF; hence, given the predicted values of the spatial factors at unobserved locations, we can create complete-coverage forest outcome maps.

Posterior inference for all candidate models was based on three chains of 50,000 post-burn-in MCMC samples. Chains converged by 20,000 MCMC iterations. Using the same computer configuration detailed in Section 4, the total runtime for the most demanding model, $q_w = 8$, was ~ 36 hours.

Table 2. Cross-validation prediction summary for forest outcomes given increasing number of spatial factors q_w . Bold values identify lowest CRPS and RMSPE.

	q_w	CRPS	RMSPE	95% Coverage	95% CI Width
AGB	1	26.21	51.37	91.88	161.24
	2	26.36	52.02	92.39	162.14
AGD	3	23.64	46.95	95.94	155.71
	4	23.53	46.93	93.91	155.66
	5	24	47.54	96.45	157.75
	6	24.47	47.8	94.92	172.64
	7	24.75	47.84	95.43	174.44
	8	24.76	48.02	96.45	182.12
	1	1,017.7	1,980.62	92.39	6,010.6
TD	2	1,006.02	1,957.54	93.4	5,944.81
	3	1,007.72	1,954.87	93.4	6,068.29
	4	997.32	1,955.2	93.4	6,040.06
	5	989.31	1,930.76	94.92	$6,\!182.2$
	6	998.3	1,944.22	94.42	$6,\!223.73$
	7	1,005.26	1,965.81	95.43	$6,\!450.5$
	8	1,004.36	1,955.08	96.95	6,503.17
	1	5.53	10.29	91.88	36.34
BA	2	5.4	10.01	94.42	36.85
DA	3	5.13	$\boldsymbol{9.54}$	93.91	35.16
	4	5.17	9.62	93.4	36.21
	5	5.16	9.58	93.4	36.51
	6	5.2	9.59	96.45	38.62
	7	5.24	9.73	95.43	38.34
	8	5.27	9.72	94.42	37.93

The eight candidate models, specified by q_w , were assessed based on their ability to inform the forest outcome predictions. This was done by fitting each of the first-stage models, then fitting their corresponding second-stage models using data from 99 of the 197 available locations in \mathcal{T}_y . The three forest outcomes were then predicted for the remaining 98 out-of-sample locations. The scoring rules and other summaries of the posterior predictive distributions for the 98 out-of-sample locations are presented in Table 2.

Increasing the number of spatial factors improves the CRPS and RMSPE for each forest outcome shown in Table 2. Exploratory analysis showed that the gains in predictive performance were negligible beyond $q_w = 4$ for AGB, $q_w = 5$ for TD, and $q_w = 3$ for BA. As such, the model with $q_w = 5$ was selected for exposition below.

Table 3 provides estimates for the second-stage model's spatial factor regres-

Table 3. Elements of Λ_y median and 95% credible intervals for the $q_w = 5$ model. Bold entries indicate where the 95% credible interval excludes zero.

Parameter	50%	(2.5%, 97.5)	%)
$\lambda_{AGB,1}^{(y)}$	-6.65	(-8.89,	-4.23)
$\lambda_{AGB,2}^{(y)}$	27.20	(-14.11,	65.14)
$\lambda_{AGB,3}^{(y)}$	-278.29	(-324.52,	-232.28)
$\lambda_{AGB,4}^{(y)}$	-46.15	(-162.56,	75.91)
$\lambda_{AGB,5}^{(y)}$	-308.81	(-524.42,	-90.45)
$\lambda_{TD,1}^{(y)}$	-1.77	(-21.35,	17.60)
$\lambda_{TD,2}^{(y)}$	-357.49	(-718.82,	-7.86)
$\lambda_{TD,3}^{(y)}$	269.03	(-137.51,	667.62)
$\lambda_{TD,4}^{(y)}$	-1,777.21 ((-2,696.67,	-708.08)
$\lambda_{TD,5}^{(y)}$	$2,\!457.52$	(681.18,	4,337.97)
$\lambda_{BA,1}^{(y)}$	-2.93	(-3.94,	-1.75)
$\lambda_{BA,2}^{(y)}$	-2.07	(-19.02,	15.79)
$\lambda_{BA,3}^{(y)}$	-98.64	(-119.79,	-76.24)
$\lambda_{BA,4}^{(y)'}$	-72.00	(-120.60,	-23.00)
$\lambda_{BA,5}^{(y)'}$	-80.55	(-177.44,	20.51)

sion coefficients, that is, the elements in Λ_y . These results show that several of the spatial factors explain a substantial portion of the variability in the forest outcomes. It is, however, difficult to interpret the different $\lambda^{(y)}$ without a sense of what characteristic of $\mathbf{z}(\mathbf{s})$ the spatial factors are capturing. When considered with the estimates in Table 3, Figure 5 provides a biological interpretation of the spatial factors. Specifically, each panel in Figure 5 represents a spatial factor. The 50 lines in each panel are observed LiDAR signals, with the lines corresponding to the 25 largest (lighter colored lines) and 25 smallest (darker lines) estimated spatial factor values.

There are some general biological relationships between the forest canopy structure and AGB, TD, and BA. A very low maximum canopy height is indicative of a young regenerating forest (e.g., regrowth after a fire), which would be characterized by low AGB, high TD, and low BA. If the majority of trees in a forest have a high canopy height, then we expect high AGB, low TD, and high BA (i.e., a few large-diameter mature trees dominate the area). When the forest is characterized by trees of many different heights (i.e., tree crowns in several vertical strata), then we might expect moderate/high AGB, moderate TD, and moderate/high BA. Some of these expected relationships are observed when

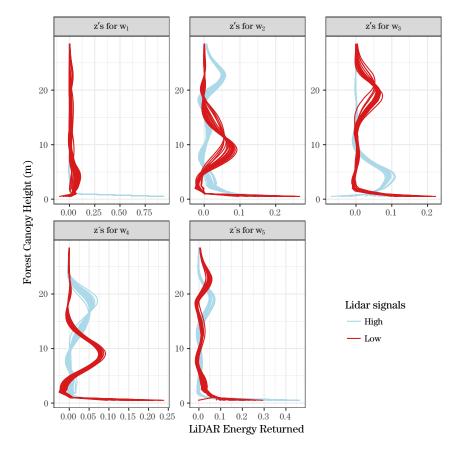


Figure 5. Observed LiDAR signals with the 25 largest (*High* in the legend) and 25 smallest (*Low* in the legend) values of $\mathbf{w}(\mathbf{s})$ from the $q_w = 5$ model.

comparing Table 3 and Figure 5. For example, the top left panel in Figure 5 differentiates between non-forested areas and all forest structures, that is, the lighter lines show a spike of energy returned at or near ground level versus red lines which show the majority of the energy is returned at or above a height of several meters. Hence, we have negative regression coefficients $\lambda_{AGB,1}^{(y)}$ and $\lambda_{BA,1}^{(y)}$ in Table 3. The LiDAR signals shown in the top right panel in Figure 5 differentiate between young and old single-cohort forests (i.e., all trees were regenerated around the same time and there is little vertical variation in canopy height); hence, we have negative $\lambda_{AGB,3}^{(y)}$ and $\lambda_{BA,3}^{(y)}$ in Table 3. The top middle and bottom left panels in Figure 5 generally separate the signal for mature 20+ and \sim 20 meter canopy heights (lighter lines), respectively, from the lower stature \sim 10 meter canopy height forest (darker lines). Consistent with the biological ex-

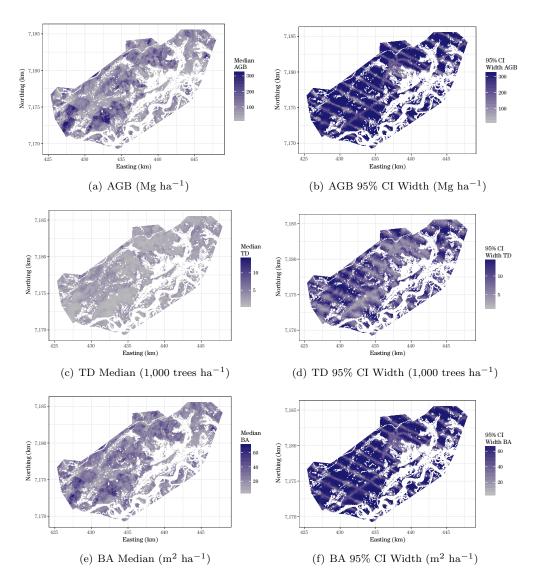


Figure 6. Model $q_w = 5$ posterior predictive distribution median and 95% CI width for AGB, TD, and BA forest variables over Bonanza Creek Experimental Forest.

pectation, the negative $\lambda_{TD,2}^{(y)}$ and $\lambda_{TD,4}^{(y)}$ suggest that forests associated with red LiDAR signals have higher tree density relative to the older taller forests.

As detailed in Section 1, complete-coverage maps of the forest outcomes with associated uncertainty estimates are important data products that can be delivered by the proposed two-stage model. Following Section 3.4 and using

the full data set depicted in Figure 1, we predicted the forest outcomes on a 30×30 m grid over the BCEF. Figure 6 provides the median and 95% credible interval width maps for each outcome. Nonforested areas are omitted (white regions on the maps). The posterior predictive point estimates match well with the distribution of the forest outcomes across the BCEF and are clearly informed by the LiDAR factors, which are capturing key forest structure characteristics. Most importantly, the prediction uncertainty maps, displayed in the right column of Figure 6, accurately reflect our lack of information for prediction units that are far from the flight lines where LiDAR data are available, that is, we achieve more precise posterior predictive distributions along and adjacent to locations where LiDAR data are available. Far from the LiDAR flight lines, prediction is only informed by the Landsat 8 tassel cap predictor variables, which in this study explained very little variability in the forest outcomes.

6. Concluding Remarks

We formulated an approach to model high-dimensional spatial data over a large set of locations, and developed an efficient implementation in C++. The SF-NNGP enables the analysis of multivariate spatially referenced data sets that, due to their magnitude, could not be rigorously explored before. It does so by combining the ability of SFMs to compress the signal from high-dimensional structures into a few dimensions with the computational scalability of NNGPs.

The algorithm was used to exploit the information from the high-dimensional LiDAR signals to jointly model and generate LiDAR-based maps of multiple forest variables. Importantly, the proposed two-stage model provides a viable approach to producing spatially continuous maps from sparsely sampled LiDAR and forest measurements. Furthermore, the model delivers spatially explicit uncertainty quantification that captures the irregular distribution of information across the domain of interest. Such frameworks will become increasingly important as sampling LiDAR systems, such as GEDI, come on-line in the near future. These approaches can also be extended to help guide LiDAR and field data acquisition to minimize prediction uncertainty.

Importantly, when fitting a spatial factor model, one must choose the number of factors q_w to use in the model; there are different strategies to address this issue. Here, we consider out-of-sample evaluation metrics for different choices of q_w and select the one where the curves flatten out. This is a pragmatic solution, similar in spirit to cross-validation approaches commonly used to tune

hyper-parameters in richly parametrized models. Like any other cross-validation approach, this leads to additional computation, but parallel computing opens the possibility of conducting simultaneous MCMC runs for different values of q_w . As shown, both in the simulation experiment and in the BCEF data analysis, this heuristic provides sufficiently good results. Other automated rank selection schemes are available in the literature, such as those proposed in Lopes and West (2004) and in Ren and Banerjee (2013); however, these drastically increase the computational burden of an already computationally costly problem.

In future research, we would like to explore an extension for spatio-temporal data. For this type of data, it is necessary to posit a strategy to select the neighbors in the spatio-temporal domain, following the discussion presented in Datta et al. (2016a).

Although our method presents a substantial improvement in terms of scalability over existing approaches, further efforts are required to scale multivariate spatial methods to truly massive data sets. For instance, the ultimate goal for forest variable mapping assisted by sampled LiDAR in interior Alaska is a complete-coverage map of the entire domain (e.g., 46 million ha), which could easily require models capable of assimilating LiDAR signals in more than 10⁸ locations.

Supplementary Materials

The supplementary materials include (1) background information on NNGPs and spatial factor models, (2) the sampling algorithm for the SF-NNGP, and (3) additional simulation results.

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