

Hierarchical Spatial Models

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Introduction

Methods for spatial and spatio-temporal modeling are becoming increasingly important in environmental sciences and other sciences where data arise from a process in an inherent spatial setting. Technological advances in remote sensing, monitoring networks, and other methods of collecting spatial data in recent decades have revolutionized scientific endeavor in fields such as agriculture, climatology, ecology, economics, transportation, epidemiology and health management, as well as many other areas. However, such technological advancements require a parallel effort in the development of techniques that enable researchers to make rigorous statistical inference given the wealth of new information at hand. Unfortunately, the application of traditional covariance-based spatial statistical models is either inappropriate or computationally inefficient in many cases. Moreover, conventional methods are often incapable of allowing the researcher to quantify uncertainties corresponding to the model parameters since the parameter space of most complex spatial and spatio-temporal models is very large.

Hierarchical Models

A main goal in the rigorous characterization of natural phenomena is the estimation and prediction of processes as well as the parameters governing processes. Thus a flexible framework capable of accommodating complex relationships between data and process models while incorporating various sources of uncertainty is necessary. Traditional likelihood based approaches to modeling have allowed for scientifically meaningful data structures, though, in complicated situations with heavily parameterized models and limited or missing data, estimation by likelihood maximization is often problematic or infeasible (Hilborn and Mangel 1997). Developments in numerical approximation methods have been useful in many cases, especially for high-dimensional parameter spaces (e.g., Newton-Raphson and E-M methods, Givens and Hoeting 2005), though can still be difficult or impossible to implement and have no provision for accommodating uncertainty at multiple levels.

Hierarchical models, whereby a problem is decomposed into a series of levels linked by simple rules of probability, assume a very flexible framework capable of accommodating uncertainty and potential *a priori* scientific knowledge while retaining many advantages of a strict likelihood approach (e.g., multiple sources of data and scientifically meaningful structure). The years after introduction of the Bayesian hierarchical model and development of MCMC (i.e., Markov Chain Monte Carlo) have brought on an explosion of research, both theoretical and applied, utilizing and (or) developing hierarchical models.

Hierarchical modeling is based on a simple fact from probability that the joint distribution of a collection of random variables can be decomposed into a series of conditional models. For example, if a, b, c are random variables, then basic probability allows the factorization:

$$[a, b, c] = [a|b, c][b|c][c] . \tag{1}$$

where the notation $[.]$ is used to specify a probability distribution. In the case of spatial and spatio-temporal models, the joint distribution describes the behavior of the process at all spatial locations (and, possibly, all times). This joint distribution (left hand side of (1)) is difficult to specify for complicated processes. Typically, it is much easier to specify the distribution of the conditional models (right hand side of (1)). In that case, the product of the series of relatively simple conditional models gives a joint distribution that can be quite complex.

When modeling complicated processes in the presence of data, it is helpful to write the hierarchical model in three basic stages:

Stage 1. Data Model: $[data|process, data\ parameters]$

Stage 2. Process Model: $[process|process\ parameters]$

Stage 3. Parameter Model: $[data\ and\ process\ parameters]$.

The basic idea is to approach the complex problem by breaking it into simpler sub-problems. Although hierarchical modeling is not new to statistics (Lindley and Smith 1972; Gelman et al. 2004), this basic formulation for modeling complicated spatial and spatio-temporal processes in the environmental sciences is a relatively new development (e.g., Berliner 1996; Wikle et al. 1998). The first stage is concerned with the observational process or “data model”, which specifies the distribution of the data given the fundamental process of interest and parameters that describe the data model. The second stage then describes the process, conditional on other process parameters. Finally, the last stage models the uncertainty in the parameters, from both the data and process stages. Note that each of these stages can have many sub-stages (e.g., Wikle et al. 1998, 2001).

Implementation

The goal is to estimate the distribution of the process and parameters given the data. Bayesian methods are naturally suited to estimation in such hierarchical settings, al-

though non-Bayesian methods can sometimes be utilized but often require additional assumptions. Using a Bayesian approach, the “posterior distribution” (i.e., the joint distribution of the process and parameters given the data) is obtained via Bayes’ Theorem:

$$[process, parameters|data] \propto [data|process, parameters] \times [process|parameters][parameters]. \quad (2)$$

Bayesian statistics involves drawing statistical conclusions from the posterior distribution which is proportional to the data model (i.e., the likelihood) times the *a priori* knowledge (i.e., the prior). Bayes’ theorem is thus the mechanism that provides access to the posterior. Although simple in principle, the implementation of Bayes’ theorem for complicated models can be challenging. One challenge concerns the specification of the parameterized component distributions on the right-hand side of (2). Although there has long been a debate in the statistics community concerning the appropriateness of “subjective” specification of such distributions, such choices are a natural part of scientific modeling. In fact, the use of scientific knowledge in the prior distribution allows for the incorporation of uncertainty related to these specifications explicitly in the model. Another, perhaps more important, challenge, from a practical perspective, is the calculation of the posterior distribution. The complex and high-dimensional nature of many scientific models (and indeed, most spatio-temporal models) prohibits the direct evaluation of the posterior. However, MCMC approaches can be utilized to estimate the posterior distribution through iterative sampling. As previously mentioned, the use of MCMC in Bayesian hierarchical models has led to a revolution, in that realistic (i.e., complicated) models can be considered; this is especially evident in the analysis of spatial and spatio-temporal processes. Yet, typically the computational burden must be considered when formulating the conditional models in such problems. Thus, the model building phase requires not only scientific understanding

of the problem, but in what ways that understanding can be modified to fit into the computational framework.

Non-analytical hierarchical models can be fitted to data using high-level programming languages (such as R, S-plus, MATLAB) or low-level languages (such as C, C++, FORTRAN). High-level languages allow for efficient programming, whereas low-level languages often allow for more efficient execution. Alternatively, the freely-distributed Bayesian computation software WinBUGS (<http://www.mrc-bsu.cam.ac.uk/bugs/>) and its spatial package GeoBUGS can be used to carry out Bayesian computations (Banerjee et al. 2004). The developers of the automated Gibbs sampler program, BUGS and WinBUGS (Thomas 1994), have been quick to point out the caveat that comes with misuse of the software (and Bayes methods in general) by cautioning that MCMC methods are not as robust as analytical methods and that the analyst should mindfully utilize such methods, especially when choosing prior distributions.

Spatial Processes

In this section we focus on the process model stage of the hierarchical framework described in the previous section and specifically applied in spatial settings. We consider the two important cases of continuous and areal data, and discuss popular modeling choices and their hierarchical forms. Another class of spatial data and models, are spatial point processes. Point processes refer to cases where points in a spatial domain are considered random and the outcome is often the occurrence of an event at a random location. In cases where covariate information is available, the covariate information marks the random location as belonging to a certain group; in such cases the data represent a marked point process. Point process analyses are often conducted to decide whether points occur independently or whether there is clustering.

General Hierarchical Spatial Model Framework

In order to consider spatial models (*sans* point processes) in a general hierarchical framework, assume $\mathbf{Z} = (Z(r_1), \dots, Z(r_m))'$ as a vector of observations of a spatial process denoted by $\mathbf{y} = (y(s_1), \dots, y(s_n))'$ where the spatial locations of the observations (r_i) do not necessarily correspond to the support of the underlying spatial process of interest (s_i). The general framework for hierarchical spatial modeling is given by the following simple and flexible structure based on the three-stage component models described previously :

$$[\mathbf{Z} \mid \mathbf{y}, \boldsymbol{\theta}_z][\mathbf{y} \mid \boldsymbol{\theta}_y][\boldsymbol{\theta}_z, \boldsymbol{\theta}_y],$$

where $\boldsymbol{\theta}_z$ and $\boldsymbol{\theta}_y$ are parameter vectors. The data model typically specifies that we have measurements from an underlying “true” process in the presence of a measurement error process. A generalized linear framework for the process model can be considered which generalizes the Gaussian model to the exponential family (see Diggle and Tawn 1998). Thus the process model can be written,

$$h(\mathbf{y}) = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\eta},$$

where $h(\cdot)$ is a known link function, $\mathbf{X} = (\mathbf{x}'(s_1), \dots, \mathbf{x}'(s_n))'$ denotes the covariates, and $\boldsymbol{\eta}$ is process model error containing explicit spatial covariance structure. This process specification can easily accommodate most conventional spatial models (e.g., see next section on models for spatially continuous data).

For example, in case of normal data, the data model can be written as,

$$\mathbf{Z} \mid \mathbf{y}, \boldsymbol{\Sigma} \sim N(\mathbf{K}\mathbf{y}, \boldsymbol{\Sigma}),$$

where \mathbf{Z} denotes measurements from an underlying “true” process \mathbf{y} in the presence of a measurement error process $\boldsymbol{\Lambda}$, and \mathbf{K} is a matrix that maps the observations to process locations (allowing for differing observation and process supports, e.g., Wikle

and Berliner 2005). Similarly for count data arising from an appropriate distribution such as the Poisson, the data model can be written as,

$$\mathbf{Z} \mid \boldsymbol{\lambda} \sim \text{Poisson}(\mathbf{K}\boldsymbol{\lambda}),$$

where the observations $\mathbf{Z} = (Z(s_1), \dots, Z(s_n))'$ are assumed to be conditionally independent, and $\boldsymbol{\lambda} = (\lambda(s_1), \dots, \lambda(s_n))'$ is the unknown spatially-varying Poisson intensity. The Poisson intensity process can then be modeled in the process model stage using covariates or latent variables (see e.g., Wikle and Hooten 2006).

Process Models for Spatially Continuous Data

Spatially continuous data (also known as geostatistical data) refer to spatially-indexed data at location s , where s varies continuously over some region R . The modeling of spatially continuous data has long been the dominant theme in spatial statistics. The most common class of models for spatially continuous data are known as Kriging models which are extensions of the method of minimizing the mean squared error in spatial settings. A general Kriging model has the following form:

$$\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\eta}, \text{ where } \boldsymbol{\eta} \sim N(\mathbf{0}, \boldsymbol{\Sigma}), \quad (3)$$

where $\mathbf{y} = \begin{bmatrix} \mathbf{y}_1 \\ \mathbf{y}_2 \end{bmatrix}$, $\mathbf{y}_1 = (y_{s_1}, \dots, y_{s_n})'$ represents the spatial process at locations for which there are data, and \mathbf{y}_2 denotes the process at a new location (e.g., s_0) or locations. Furthermore, the term $\mathbf{X}\boldsymbol{\beta}$ ($= \boldsymbol{\mu}$) represents the mean of the process (possibly explained by covariates \mathbf{X}), and spatially correlated error $\boldsymbol{\eta}$ with covariance,

$$\boldsymbol{\Sigma} = \begin{bmatrix} \boldsymbol{\Sigma}_{11} & \boldsymbol{\Sigma}_{12} \\ \boldsymbol{\Sigma}_{21} & \boldsymbol{\Sigma}_{22} \end{bmatrix}.$$

Kriging, then involves finding the best linear predictor of \mathbf{y}_2 given \mathbf{y}_1 :

$$\mathbf{y}_2 \mid \mathbf{y}_1 \sim N(\boldsymbol{\mu}_2 + \boldsymbol{\Sigma}_{21}\boldsymbol{\Sigma}_{11}^{-1}(\mathbf{y}_1 - \boldsymbol{\mu}_1), \boldsymbol{\Sigma}_{22} - \boldsymbol{\Sigma}_{21}\boldsymbol{\Sigma}_{11}^{-1}\boldsymbol{\Sigma}_{12}),$$

where $\boldsymbol{\mu} = \begin{bmatrix} \boldsymbol{\mu}_1 \\ \boldsymbol{\mu}_2 \end{bmatrix}$, $E(\mathbf{y}_1) = \boldsymbol{\mu}_1$, and $E(\mathbf{y}_2) = \boldsymbol{\mu}_2$.

The model in (3) is known as a universal Kriging model. A hierarchical representation of this model can be written as:

$$\mathbf{Z} \mid \mathbf{y}, \sigma^2 \sim N(\mathbf{K}\mathbf{y}, \sigma^2\mathbf{I}), \quad (4)$$

$$\mathbf{y} \mid \boldsymbol{\beta}, \boldsymbol{\Sigma}_y \sim N(\mathbf{X}\boldsymbol{\beta}, \boldsymbol{\Sigma}_y), \quad (5)$$

$$\boldsymbol{\beta} \mid \boldsymbol{\beta}_0, \boldsymbol{\Sigma}_\beta \sim N(\boldsymbol{\beta}_0, \boldsymbol{\Sigma}_\beta), \quad (6)$$

$$\{\sigma^2, \boldsymbol{\theta}_y, \boldsymbol{\beta}_0, \boldsymbol{\Sigma}_\beta\} \sim [\sigma^2, \boldsymbol{\theta}_y, \boldsymbol{\beta}_0, \boldsymbol{\Sigma}_\beta]. \quad (7)$$

Note that the parameters are assumed to be independent conditioned on the data.

Process Models for Areal Data

Areal data (also known as lattice data) are spatially-indexed data associated with geographic regions or areas such as counties or zip codes, and are often presented as aggregated values over an areal unit with well-defined boundaries. Spatial association among the areal units is specified by defining neighborhood structure for the areas (regular or irregular) of interest. Examples of such data include a wide variety of problems from disease mapping in counties to modeling air-pollution on a grid. Models described in this section are based on Markov random fields (MRFs). MRFs are a special class of spatial models that are suitable for data on discrete (countable) spatial domains in which a joint distribution of y_i (for $i = 1, \dots, n$, where y_i is the spatial process at spatial unit i) is determined by using a set of locally specified conditional distributions for each spatial unit conditioned on its neighbors. MRFs include a wide class of spatial models, such as auto-Gaussian models for spatial Gaussian processes, auto-logistic models for binary spatial random variables, auto-Gamma models for non-negative continuous processes, and auto-Poisson models for spatial count processes. Here we focus on two popular auto-Gaussian models, CAR and SAR models.

Conditionally Autoregressive (CAR) Models

As introduced by Besag (1974), conditionally autoregressive (CAR) models are popular hierarchical spatial models for use with areal data. Here, we consider the Gaussian case. Assume $y_i \equiv y(i^{\text{th}} \text{ spatial area})$ and $\mathbf{y} = (y_1, \dots, y_n)'$, then the CAR model can be defined as the following n conditional distributions:

$$y_i \mid y_j, \tau_i^2 \sim N(\mu_i + \sum_{j \in N_i} c_{ij}(y_j - \mu_j), \tau_i^2), \quad i, j = 1, \dots, n, \quad (8)$$

where N_i is defined as a set of neighbors of area i , $E(y_i) = \mu_i$, τ_i^2 is the conditional variance, and the c_{ij} are constants such that $c_{ii} = 0$ for $i = 1, \dots, n$, and $c_{ij}\tau_j^2 = c_{ji}\tau_i^2$. It can be shown (Cressie 1993) that the joint distribution of \mathbf{y} using the conditional distributions can be written as

$$\mathbf{y} \sim N(\boldsymbol{\mu}, (\mathbf{I} - \mathbf{C})^{-1}\mathbf{M}), \quad (9)$$

where $\boldsymbol{\mu} = (\mu_1, \dots, \mu_n)'$, $\mathbf{C} = [c_{ij}]_{n \times n}$, and $\mathbf{M} = \text{diag}(\tau_1^2, \dots, \tau_n^2)$. Note that $(\mathbf{I} - \mathbf{C})$ has to be invertible and $(\mathbf{I} - \mathbf{C})^{-1}\mathbf{M}$ has to be symmetric and positive-definite.

The implementation of the CAR model is convenient in hierarchical Bayesian settings because of the explicit conditional structure. Perhaps the most popular implementation of the CAR model is the pairwise difference formulation proposed by Besag et al. (1991), where B is decomposed into an adjacency matrix and a diagonal matrix containing information on the number of neighbors for each of the areal units which results in a simple and easy to fit version of the model. Although the convenient specification of CAR models make them attractive for modeling areal data, the usage of these models often involves numerous theoretical and computational difficulties (e.g., singularity of the covariance function of the joint distribution results in the joint distribution being improper; for more details see Banerjee et al. 2004). Several methods to overcome such difficulties have been proposed (e.g., Cressie 1993; Carlin

and Banerjee 2003), however the development of strategies to address the difficulties of CAR models is a topic of ongoing research.

Simultaneous Autoregressive (SAR) Models

Simultaneous autoregressive (SAR) models, introduced by Whittle (1954), are a class of spatial models for areal data. SAR models are a subset of MRFs and are popular in econometrics. Here we consider the Gaussian case,

$$y_i = \mu_i + \sum_j b_{ij}(y_j - \mu_j) + \varepsilon_i$$

or equivalently, in matrix notation,

$$\mathbf{y} = \boldsymbol{\mu} + (\mathbf{I} - \mathbf{B})^{-1}\boldsymbol{\varepsilon},$$

where $\mathbf{B} = [b_{ij}]_{n \times n}$ is a matrix that can be interpreted as the spatial-dependence matrix, $\boldsymbol{\varepsilon} \sim N(\mathbf{0}, \boldsymbol{\Lambda})$, and $\boldsymbol{\Lambda}$ is an $n \times n$ diagonal covariance matrix of $\boldsymbol{\varepsilon}$ (e.g., $\boldsymbol{\Lambda} = \sigma^2 \mathbf{I}$). Thus, $\boldsymbol{\varepsilon}$ induces the following distribution for \mathbf{y} ,

$$\mathbf{y} \sim N(\boldsymbol{\mu}, (\mathbf{I} - \mathbf{B})^{-1}\boldsymbol{\Lambda}(\mathbf{I} - \mathbf{B}')^{-1}),$$

where $(\mathbf{I} - \mathbf{B})$ has to be full rank. There are two common choices for \mathbf{B} ; one is based on a spatial autoregression parameter (ρ) and an adjacency matrix, and the other is based on a spatial autocorrelation parameter (α) and a normalized adjacency matrix. Thus, the following alternative models can be considered:

$$y_i = \rho \sum_{j \in N_i} w_{ij} y_j + \varepsilon_i,$$

or,

$$y_i = \alpha \sum_{j \in N_i} \frac{w_{ij}}{\sum_k w_{ik}} y_j + \varepsilon_i$$

where w_{ij} 's are elements of an adjacency matrix \mathbf{W} , with 0 or 1 entries, describing the neighborhood structure for each unit. CAR and SAR models are equivalent if and only

if their covariance matrices are equal (i.e., $(\mathbf{I} - \mathbf{C})^{-1}\mathbf{M} = (\mathbf{I} - \mathbf{B})^{-1}\mathbf{\Lambda}(\mathbf{I} - \mathbf{B}')^{-1}$). Any SAR model can be represented as a CAR model but the converse is not necessarily true. One main difference between SAR and CAR models is that the spatial-dependence matrix for CAR models (\mathbf{C}) is symmetric, while the spatial-dependence matrix for SAR models (\mathbf{B}) need not be symmetric. Although this might be interpreted as an advantage for SAR models in situations where the spatial dependence of neighboring sites is defined in an asymmetric way, non-identifiability issues related to estimation of model parameters make CAR models more preferable for cases with symmetric dependency structures and a spatio-temporal model can be used when asymmetric dependencies are of interest (for more details on the comparisons between SAR and CAR models see Cressie 1993, pages 408-410).

Spatio-temporal Processes

Spatio-temporal processes are often complex, exhibiting different scales of spatial and temporal variability. Such processes are typically characterized by a large number of observations and prediction locations in space and time, differing spatial and temporal support, orientation and alignment (relative to the process of interest), and complicated underlying dynamics. The complexity of such processes in “real-world” situations is often intensified due to non-existence of simplifying assumptions such as Gaussianity, spatial and temporal stationarity, linearity, and space-time separability of the covariance function. Thus implementation of a joint perspective of modeling for spatio-temporal processes, although relatively easy to formulate, is challenging. On the contrary, a hierarchical formulation allows for modeling of complicated spatial and temporal structures by decomposing an intricate joint spatio-temporal process into relatively simple conditional models. The main advantage of the Bayesian hierarchical model over traditional covariance-based methods is that it allows the complicated structure to be modeled at a lower level in the hierarchy, rather than attempting to

model the complex joint covariance matrix.

General Spatio-Temporal Model

Let $Z(s, t)$ be a spatio-temporal process where $s \in D_s$, and D_s is a continuous or discrete, and potentially time-varying, spatial domain and $t \in D_t$ is a discrete temporal domain. The generality of the definition of the spatial domain allows for the spatio-temporal process to be applicable to both cases of continuous data and areal data. A general decomposition of the process (where $Z(s, t)$ and $Y(s, t)$ have the same spatial support and no missing data) can be written as:

$$Z(s, t) = Y(s, t) + \varepsilon(s, t), \quad (10)$$

where $Y(s, t)$ is the “true” underlying correlated process of interest and $\varepsilon(s, t)$ is a zero-mean measurement error process. The underlying process $Y(s, t)$ can be further decomposed into a mean process, additive error process, and spatial or temporal random effects (e.g., Reinsel 1981, Oehlert 1993, and Stein 1986). Recent approaches to spatio-temporal modeling have focused on the specification of joint space-time covariance structures (e.g., Cressie and Huang 1999, Gneiting 2002, and Stein 2005). However, in high-dimensional settings with complicated non-linear spatio-temporal behavior, such covariance structures are very difficult to formulate. An alternative approach to modeling such complicated processes is using spatio-temporal dynamic models in a hierarchical fashion.

Spatio-Temporal Dynamic Models

Many spatio-temporal processes are dynamic in the sense that the current state of the process is a function of the previous states. There are many examples of spatio-temporal models with dynamic components in the literature (e.g., Guttorp et al. 1994, Huang and Cressie 1996, Waller et al. 1997, Wikle et al. 1998, Wikle and Cressie 1999,

Wikle et al. 2001, Stroud et al. 2001). The joint spatio-temporal process \mathbf{Y} can be factored into conditional models based on a Markovian assumption. That is,

$$[\mathbf{Y} | \boldsymbol{\theta}_t, t = 1, \dots, T] = [\mathbf{y}_0] \prod_{t=1}^T [\mathbf{y}_t | \mathbf{y}_{t-1}, \boldsymbol{\theta}_t], \quad (11)$$

where $\mathbf{y}_t = (y(s_1, t), \dots, y(s_n, t))'$ and the conditional distribution $[\mathbf{y}_t | \mathbf{y}_{t-1}, \boldsymbol{\theta}_t]$ depend on a vector of parameters $\boldsymbol{\theta}_t$ which govern the dynamics of the spatio-temporal process of interest. An example of such spatio-temporal dynamic models is when the process has a first-order Markovian structure:

$$\mathbf{y}_t = \mathbf{H}_{\boldsymbol{\theta}_t} \mathbf{y}_{t-1} + \boldsymbol{\eta}_t, \text{ where } \boldsymbol{\eta}_t \sim N(\mathbf{0}, \boldsymbol{\Sigma}_\eta), \quad (12)$$

where $\boldsymbol{\eta}_t$ is a spatial error process, and $\mathbf{H}_{\boldsymbol{\theta}_t}$ is a “propagator matrix” (sometimes called a transition or evolution matrix) which includes the parameters that govern the dynamics of the process. If these parameters are known or easy to estimate, an implementation of the model through Kalman filtering is possible (e.g., West and Harrison 1989, Huang and Cressie 1996, Wikle and Cressie 1999). If the parameters are unknown, $\mathbf{H}_{\boldsymbol{\theta}_t}$ can be modeled in a hierarchical fashion by specifying prior distributions for $\mathbf{H}_{\boldsymbol{\theta}_t}$ or its parameters $\boldsymbol{\theta}_t$. The hierarchical form for spatio-temporal dynamic models is sometimes motivated by biological processes explained by partial differential equations (PDEs) that describe an approximate behavior of underlying physical processes. For example, Wikle et al. (2001) used the shallow-water PDEs that approximate atmospheric processes in the tropics in order to develop prior distributions for predicting high-resolution wind fields over the tropical ocean.

Conclusion

In this article, a brief overview of hierarchical spatial and spatio-temporal models is presented. In the recent decades, hierarchical models have drawn the attention of scientists in many fields and are especially suited to studying spatial processes. Recent

computational advances and the development of efficient algorithms has provided the tools necessary for performing the extensive computations involved in hierarchical modeling. Advances in hierarchical modeling have created opportunities for scientists to take advantage of massive spatially-referenced databases. The hierarchical approach to spatial modeling is fairly new, but has already attracted researchers and scientists as a versatile modeling tool. Although the literature on hierarchical spatial modeling is rich, there are still many problems and issues yet to be considered. Below we briefly review some of these challenges.

In most spatial and spatio-temporal processes, researchers have to deal with data obtained by different sources and as well as different scales. For example, a combination of Eulerian and Lagrangian data is often collected in sciences such as oceanography. Alignment and change of spatial support often presents a significant challenge for analysts. There is a need for the development of efficient methods to address these issues.

Spatial and spatio-temporal models have recently been extended to accommodate multivariate situations (e.g., popular univariate models such as Kriging and CAR models have been extended to include multivariate cases). The distinction between continuous data and areal data, as described for the univariate case, holds true for the multivariate case. Multivariate approaches have the added advantage of not only being able to rely on covariate and covariance based information, but to “borrow strength” between observation vectors (i.e., response variables) as well. Examples of such multivariate models are cokriging for multivariate continuous data and multivariate CAR models for areal data, multivariate dynamic models.

Spatial and spatio-temporal models are typically high-dimensional. This characteristic complicates the modeling process and necessitates development of efficient computational algorithms on one hand, and implementation of dimension reduction methods

(e.g., recasting the problem in a spectral context), on the other hand.

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