Conquer Big Spatial Data: The Stochastic PDE Approach

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Motivation

- The method of approximating a continuous Gaussian field using GMRFs was theoretically good but less practical. The SPDE method represents a Gaussian field with Matern covariance by representing a solution of stochastic partial differential equations as a GMRF using the finite element method.
- The GMRF representation of the Gaussian field, which can be computed explicitly, provides a sparse representation of the spatial effect through a sparse precision matrix. This enables the nice computational properties of the GMRFs which can then be implemented in the INLA package.

The SPDE Model

•**y** | β_0 , **u**, $\sigma_e^2 \sim N(\beta_0 + Au, \sigma_e^2)$

• $\mathbf{u} \sim \textit{GF}(0,\Sigma)$

•y_i are observations at location s_i.

• β_0 is the intercept

•u is a spatial Gaussian random field with mean zero and standard deviation Σ

•A is the projector matrix

Mesh

- Points are often distributed irregularly. Need to construct mesh
- Mesh is used in the Finite Element Method to provide a solution to a SPDE.
- To better approximate the field, need to have more triangles.
- Shiny application within INLA package: meshbuilder()



Understanding the Method

- A Gaussian field with a generalized covariance function obtained in the Matérn correlation function when v>0 is a solution to a SPDE.
- Consider a regular two-dimensional lattice with number of sites tending to infinity.
- The right figure is an example of a GMRF. Another example is AR(1) process
- The GMRF representation is a convolution of processes with precision matrix for distinct values of smoothness v
- As the smoothness parameter v increases, the precision matrix in the GMRF representation becomes denser. This is because the conditional distributions depend on a wider neighborhood.



How Approximation Works

• Illustrate the approximation in two-dimensional space considering piecewise linear basis functions.

 $u(s) = \sum_{j=1}^{m} \psi_k(s) W_k$

- where ψ_k are basis functions and W_k are Gaussian distributed weights, k=1,...,m with m the number of vertices in the triangulation.
- Carefully choose the basis functions to preserve the sparse structure of the resulting precision matrix for the random field at a set of mesh nodes.
- This provides an explicit link between a continuous random field and a GMRF representation, which allows efficient computations.



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To Build a 2-D Mesh INLA package in R

inla.mesh.2d() function

Mandatory arguments:

- *loc, loc.domain, or boundary*
- max.edge: inner domain,

outer extension

Optional arguments:

- cutoff
- offset
- min.angle
- n

* Image is taken from https://becarioprecario.bitbucket.io/spde-gitbook/index.html

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Non-convex Hull Meshes



boundary <- inla.nonconvex.hull(points = , convex = , concave = , resolution = ,...)
D Chinese provide a convex in a convex in the convex is a convex in the convex in the convex is a convex in the convex in the convex is a convex in the convex in the convex is a convex in the convex in the convex is a convex in the convex in the convex is a convex in the convex in the convex is a convex in the convex in the convex is a convex in the convex in the convex is a convex in the convex in the convex in the convex is a convex in the convex in the convex is a convex in the convex in the convex in the convex is a convex in the convex in the convex is a convex in the convex in the convex is a convex in the convex in the convex in the convex in the convex is a convex in the c

R Shiny app: meshbuilder();

* Images are taken from https://becarioprecario.bitbucket.io/spde-gitbook/index.html

J. Wang

Quality of A Mesh

• Goal: Uniform triangle shape and size

mesh <- inla.mesh.2d(loc = coordinates,</pre>

max.edge = c(<inner domain>, <outer extension>),
cutoff = <a numeric value>,
offset = -0.10 <default>)

• To create the projector matrix (A matrix) from the mesh:

A <- inla.spde.make.A(mesh, *loc* = coordinates)

Problem

• Distance is too small:

Range(longitude) = 0.44 degrees, range(latitude) = 0.21 degrees

• Re-scale the coordinates:

(longitude + 96) ×10; (latitude - 41) ×10

• Number of nodes for n = 20,000:

i. node = 3,312: max.edge = c(0.1, 1), cutoff = 0.05

ii. node = 5,962: max.edge = c(0.08, 0.8), cutoff = 0.03

iii. node = 34,861: max.edge = c(0.03, 0.7), cutoff = 0.01

The SPDE Model Construction

- $\mathbf{y} \mid \beta_0, \, \mathbf{u}, \, \sigma_e^2 \sim N(\beta_0 + \mathbf{A}\mathbf{u}, \sigma_e^2)$
- $\mathbf{u} \sim GF(0, \Sigma)$
- Matern covariance with Penalized Complexity prior: inla.spde2.pcmatern()

i.
$$\alpha \in [1, 2]; \alpha = v + d/2 = 2 < default>$$

ii. $P(\sigma > \sigma_0) = p \Rightarrow P(\sigma > 1) = 0.01$:

prior.sigma = c(1, 0.01)

iii.
$$P(r < r_0) = p \Rightarrow P(r < 1.3) = 0.5$$
:

prior.range = c(1.3, 0.5)

MLE & SPDE Run Time



- MLE Estimation passed over one hour on fourth group
- SPDE Models runtime did not increase exponentially
- SPDE Model with most nodes (~33,000) stayed under half hour with all groups

Performance Across Group Size



- Kriging using MLE Estimation & SPDE showed a decline in MSE as size increased
 - SPDE models with higher node counts decreased more significantly than SPDE models with smaller node counts
- Coverage stayed relatively consistent in MLE and SPDE estimation when node count was high (3000, 5500, 33000) but decreased in coverage as size increased when node counts were small

MAD & MSE



- Kriging methods using SPDE estimation with node size >3,000 performed better than MLE models
- Conversely SPDE estimation with node sizes 300, 470 and 600 performed worse than the MLE estimation

Coverage vs. Correlation



- Kriging using MLE Estimation had the highest coverage with ~95%
- As the number of nodes increased, coverage increased for SPDE-based Kriging methods
- SPDE methods using nodes 3,000, 5,500 and 33,000 resulted in higher levels of correlation than the MLE estimation

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0.0

0.2

Methods Overview



Y hat.spde

0.6

0.8

- The predictions made with Kriging using the SPDE estimation better correlate with the test set.
- Additionally, it better matches the test data set in terms of the histogram, with more of the tails captured
- The predictions made with Kriging using the • MLE estimation do not capture the tails of the test set and are more centered around the central value

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