Gaussian Copulas for Large Spatial Fields

Modeling Data-Level Spatial Dependence in Multivariate Generalized Extreme Value Distributions

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Introduction

- ► UKCP Local Projections on a 5km grid over the UK (1980-2080)
- Challenge: Modeling maximum daily precipitation in yearly blocks
 - ▶ 43,920 spatial locations on a 180 x 244 grid
- Two aspects of spatial dependence:
 - 1. GEV parameters (ICAR models)
 - 2. Data-level dependence (Copulas)



Calculating Multivariate Normal Densities

Log Density Formula

$$\log f(\mathbf{x}) \propto \frac{1}{2} \left(\log |\mathbf{Q}| - \mathbf{x}^T \mathbf{Q} \mathbf{x} \right)$$

Key Components

- 1. Log Determinant: $\log |\mathbf{Q}|$
 - Constant for a given precision matrix
- 2. Quadratic Form: $\mathbf{x}^T \mathbf{Q} \mathbf{x}$
 - Needs calculation for each density evaluation

Computational Challenges

- ▶ Log determinant calculation
 - Time complexity: $O(n^3)$ for naive methods
 - ightharpoonup Memory complexity: $O(n^2)$
- Quadratic form calculation
 - ightharpoonup Time complexity: $O(n^2)$
 - Critical for performance in large spatial fields

Spatial Model Considerations

- Some models (e.g., ICAR) avoid log determinant calculation
- Efficient computation crucial for large-scale applications

Spatial Models

Conditional Autoregression (CAR)

- **D** is a diagonal matrix with $D_{ii} = n_i$, the number of neighbours of i
- A is the adjacency matrix with
 - $A_{ij} = A_{ji} = 1$ if $i \sim j$
- ightharpoonup au models overall precision

Besag's Intrinsic Conditional Autoregression (ICAR)

- ightharpoonup lpha = 1, so ${f Q}$ is singular, but constant
- lacksquare Don't have to calculate $\log |\mathbf{Q}|$
- ightharpoonup au is a precision parameter

 $\mathbf{x} \sim N(\mathbf{0}, \tau \mathbf{Q})$

 $\mathbf{Q} = \mathbf{D} \left(\mathbf{I} - \alpha \mathbf{A} \right)$

 $\mathbf{x} \sim N(\mathbf{0}, \tau \mathbf{Q})$

Q = D - W

Spatial Models

BYM (Besag-York-Mollié) Model

- u is the structured spatial component (Besag model)
- v is the unstructured component (i.i.d. normal)
- $\blacktriangleright \ \tau_u$ and τ_v are precision parameters for each component

BYM2 Model

- Rewrite the combination to get proper scaling
- ho models how much of variance is spatial
- lacksquare s is a scaling factor chosen to make $Var(\mathbf{u}_s) \approx 1$

$$x = u + v$$

 $\mathbf{u} \sim \mathrm{ICAR}(\tau_u)$ $\mathbf{v} \sim N(\mathbf{0}, \tau_v^{-1})$

$$\mathbf{x} = \left(\left(\sqrt{\rho/s} \right) \mathbf{u} + \left(\sqrt{1 - \rho} \right) \mathbf{v} \right) \sigma$$
$$\mathbf{u} \sim ICAR(1)$$

 $\mathbf{v} \sim N(\mathbf{0}, n)$

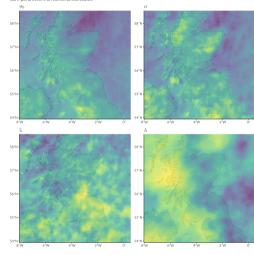
Spatial Modeling on Parameter-level

- $\blacktriangleright \mu = \mu_0 \left(1 + \Delta \left(t t_0\right)\right)$, location
- σ : scale
- ξ : shape

$$\begin{split} \log(\mu_0) &= \psi \sim \text{BYM2}(\mu_{\psi}, \rho_{\psi}, \sigma_{\psi}) \\ \log(\mu_0) &- \log(\sigma) = \tau \sim \text{BYM2}(\mu_{\tau}, \rho_{\tau}, \sigma_{\tau}) \\ f_{\xi}(\xi) &= \phi \sim \text{BYM2}(\mu_{\phi}, \rho_{\phi}, \sigma_{\phi}) \\ f_{\Delta}(\Delta) &= \gamma \sim \text{BYM2}(\mu_{\gamma}, \rho_{\gamma}, \sigma_{\gamma}) \end{split}$$

Our BYM2 hyperparameters point to a large degree of spatial variation									
variable	mean	median	sd	mad	q5	q95	rhat	ess_bulk	ess_tail
σψ	0.072	0.072	0.001	0.001	0.070	0.074	1.012	137	387
μφ	2.133	2.133	0.001	0.001	2.131	2.135	1.001	5,029	3,417
Pe	0.998	0.998	0.001	0.001	0.997	0.999	1.000	3,161	3,218
σ_t	0.102	0.102	0.002	0.002	0.098	0.106	1.015	381	973
με	-0.923	-0.923	0.001	0.001	-0.926	-0.921	1.000	4,130	3,507
ρι	0.997	0.998	0.001	0.001	0.996	0.999	1.001	3,192	2,903
σ_{Φ}	0.358	0.358	0.009	0.009	0.343	0.372	1.008	401	730
μφ	0.341	0.341	0.004	0.004	0.335	0.347	1.001	3,728	3,049
Po	0.996	0.997	0.001	0.001	0.994	0.998	1.002	1,995	2,677
σ_{y}	0.332	0.333	0.011	0.011	0.313	0.351	1.029	105	208
μ,	1.438	1.438	0.012	0.012	1.419	1.458	1.001	2,517	3,053
ρ,	0.996	0.996	0.002	0.001	0.993	0.998	1.000	3,823	2,875

Spatial distribution of posterior means



From Parameter-level to Data-level Dependence

Parameter-level Dependence

- Assumes conditional independence
- ▶ Biased joint probability estimates
- Underestimates parameter variance

Copula

- Improves joint probabilities
- ► Enhances spatial risk assessment
- Better variance estimates

Sklar's Theorem: For any multivariate distribution H, there exists a unique copula C such that:

$$H(\mathbf{x}) = C(F_1(x_1), \dots, F_d(x_d))$$

where F_i are marginal distributions. We can also write this as a log-density

$$\log h(x) = \log c(F_1(x_1), \dots, F_d(x_d)) \sum_{i=1}^d \log f_i(x_i)$$

Our Approach: Matérn-like Gaussian Copula

Marginal CDFs, $F_i(x_i)$, is $\operatorname{GEV}(\mu_i, \sigma_i, \xi_i)$

$$\begin{split} \log h(\mathbf{x}) &= \log c(u_1, \dots, u_d) + \sum_{i=1}^d f_{\text{GEV}}(x_i | \mu_i, \sigma_i, \xi_i) \\ u_i &= F_{\text{GEV}}(x_i | \mu_i, \sigma_i, \xi_i) \end{split}$$

Gaussian Copula

$$\log c(\mathbf{u}) \propto \frac{1}{2} \left(\log |\mathbf{Q}| - \mathbf{z}^T \mathbf{Q} \mathbf{z} + \mathbf{z}^T \mathbf{z} \right)$$
$$\mathbf{z} = \Phi^{-1}(\mathbf{u})$$

The Precision Matrix

 \mathbf{Q} defined as Kronecker sum of two AR(1) precision matrices

$$\mathbf{Q} = \left(\mathbf{Q}_{\rho_1} \otimes \mathbf{I_{n_2}} + \mathbf{I_{n_1}} \otimes \mathbf{Q}_{\rho_2}\right)^{\nu+1}, \quad \nu \in \{0,1,2\}$$

$$\mathbf{Q}_{\rho} = \frac{1}{1 - \rho^2} \begin{bmatrix} 1 & -\rho & 0 & \cdots & 0 \\ -\rho & 1 + \rho^2 & -\rho & \cdots & 0 \\ 0 & -\rho & 1 + \rho^2 & \cdots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & 0 & \cdots & 1 \end{bmatrix}$$

Eigendecomposition

Because of how Q is defined, we know that

$$\begin{split} \mathbf{Q} &= \mathbf{V} \boldsymbol{\Lambda} \mathbf{V} \\ &= (\mathbf{V}_{_{\mathbf{1}}} \otimes \mathbf{V}_{_{\mathbf{2}}}) (\boldsymbol{\Lambda}_{\rho_{_{\mathbf{1}}}} \otimes \mathbf{I} + \mathbf{I} \otimes \boldsymbol{\Lambda}_{\rho_{_{\mathbf{2}}}})^{\nu+1} (\mathbf{V}_{_{\mathbf{1}}} \otimes \mathbf{V}_{_{\mathbf{2}}})^{T} \\ \mathbf{Q}_{\rho_{_{\mathbf{1}}}} &= \mathbf{V}_{_{\mathbf{1}}} \boldsymbol{\Lambda}_{\rho_{_{\mathbf{1}}}} \mathbf{V}_{_{\mathbf{1}}}^{T} & \& \quad \mathbf{Q}_{\rho_{_{\mathbf{2}}}} &= \mathbf{V}_{_{\mathbf{2}}} \boldsymbol{\Lambda}_{\rho_{_{\mathbf{2}}}} \mathbf{V}_{_{\mathbf{2}}}^{T} \end{split}$$

Spectral decomposition defined by value/vector pairs of smaller matrices

$$\left\{\lambda_{\rho_{1}}\right\}_{i}+\left\{\lambda_{\rho_{2}}\right\}_{j} \qquad \qquad \left\{\mathbf{v}_{\rho_{1}}\right\}_{i}\otimes\left\{\mathbf{v}_{\rho_{2}}\right\}_{j}$$

- Problem: $\Sigma_{ii} = (\mathbf{Q}^{-1})_{ii} \neq 1$
- \blacktriangleright Solution: $\mathbf{\tilde{Q}}=\mathbf{DQD},$ where $\mathbf{D}_{ii}=\sqrt{\Sigma_{ii}}$

Marginal Standard Deviations

$$\Sigma = \mathbf{Q}^{-1} = (\mathbf{V}\Lambda\mathbf{V}^T)^{-1} = \mathbf{V}\Lambda^{-1}\mathbf{V}$$

We know that if A=BC then $A_{ii}=B_{i..}C_{..i}$, so

$$\Sigma_{ii} = \sum_{k=1}^{n} v_{ik} \frac{1}{\lambda_k} (v^T)_{ki} = \sum_{k=1}^{n} v_{ik} \frac{1}{\lambda_k} v_{ik} = \sum_{k=1}^{n} v_{ik}^2 \frac{1}{\lambda_k}$$

Compute vector σ^2 containing all marginal variances

$$\sigma^2 = \sum_{i=1}^{n_1} \sum_{j=1}^{n_2} \frac{\left(\left\{\mathbf{v}_{\rho_1}\right\}_i \otimes \left\{\mathbf{v}_{\rho_2}\right\}_j\right)^2}{\left(\left\{\lambda_{\rho_1}\right\}_i + \left\{\lambda_{\rho_2}\right\}_j\right)^{\nu+1}}$$

Marginal Standard Deviations

dim1 <- 50: dim2 <- 50

```
rho1 <- 0.5; rho2 <- 0.3
                                                                   E1 <- eigen(Q1)
nii <- 2
                                                                   E2 <- eigen(Q2)
Q1 <- make AR prec matrix(dim1, rho1)
                                                                   marginal_sd_eigen(
Q2 <- make AR prec matrix(dim2, rho2)
                                                                     E1$values, E1$vectors, dim1.
I1 <- Matrix::Diagonal(dim1)</pre>
                                                                     E2$values, E2$vectors, dim2,
I2 <- Matrix::Diagonal(dim2)
Q <- temp <- kronecker(Q1, I2) + kronecker(I1, Q2)
                                                                   ) |> sort()
for (i in seg_len(nu)) Q <- Q %*% temp
  bench::mark(
    "solve" = solve(Q) \mid > diag() \mid > sort() \mid > sort().
    "inla.giny" = inla.giny(Q) |> diag() |> sgrt() |> sort().
    "marginal_sd_eigen" = msd(Q1, Q2),
    iterations = 10, filter gc = FALSE
  # A tibble: 3 \times 6
                                                median `itr/sec` mem alloc `gc/sec`
     expression
                                        min
                                <bch:tm> <bch:tm>
                                                                                              <dbl>
     <bch:expr>
                                                                 <dbl> <bch:bvt>
                                                                                              0.757
  1 solve
                                     1.27s
                                                  1.29s
                                                                 0.757
                                                                             78.17MB
                                                                 2.72
                                                                               4.33MB
  2 inla.ginv
                                  358.2ms 367.86ms
  3 marginal sd eigen
                                   3.29ms
                                                 3.35 ms
                                                              279.
                                                                            649.35KB
```

msd <- function(Q1, Q2) {