

# Parallel Hierarchical Matrix Technique to Approximate Large Covariance Matrices, Likelihood Functions and Parameter Identification

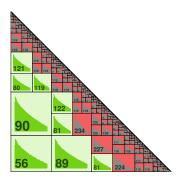
Alexander Litvinenko<sup>1</sup>, (joint work with V. Berikov<sup>2</sup>(Novosibirsk State University), M. Genton<sup>3</sup>, D. Keyes<sup>3</sup>, R. Kriemann<sup>4</sup>, and Y. Sun<sup>3</sup>) SIAM CSE 2021

<sup>1</sup>RWTH Aachen, Germany, <sup>2</sup>Novosibirsk Staate University, <sup>3</sup>KAUST, Saudi Arabia, <sup>4</sup>MPI for Mathematics in the Sciences in Leipzig, Germany

## Analysis of large datasets



Given: Dense (covariance) matrix of size  $2,000,000 \times 2,000,000$ 



#### We show how to:

- 1. reduce storage cost from 32TB to 16 GB
- 2. approximate its Cholesky factorisation, determinant, inverse in 8 minutes on modern desktop computer.
- 3. use it for prediction

#### The structure of the talk



- 1. Motivation: improve statistical model and data analysis
- 2. Identification of unknown parameters via maximizing Gaussian log-likelihood (MLE)
- 3. Tools: Hierarchical matrices [Hackbusch 1999]
- 4. Matérn covariance function, joint Gaussian log-likelihood
- 5. Error analysis
- 6. Prediction at new locations
- 7. Comparison with machine learning methods



Given:  $Z = \{Z(s_1), \dots, Z(s_n)\}^{\top}$ , where Z(s) is a Gaussian random field indexed by a spatial location  $s \in \mathbb{R}^d$ ,  $d \ge 1$ .

Assumption: Z has mean zero and stationary parametric covariance function, e.g. Matérn:

$$\mathsf{C}(oldsymbol{ heta}) = rac{2\sigma^2}{\mathsf{\Gamma}(
u)} \left(rac{r}{2\ell}
ight)^
u \mathsf{K}_
u \left(rac{r}{\ell}
ight) + au^2 \mathsf{I}, \quad oldsymbol{ heta} = (\sigma^2, 
u, \ell, au^2).$$

To identify: unknown parameters  $\theta := (\sigma^2, \nu, \ell, \tau^2)$ .

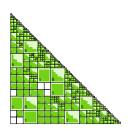


Statistical inference about heta is then based on the Gaussian log-likelihood function:

$$\mathcal{L}(\boldsymbol{\theta}) = -\frac{n}{2}\log(2\pi) - \frac{1}{2}\log|\mathsf{C}(\boldsymbol{\theta})| - \frac{1}{2}\mathsf{Z}^{\top}\mathsf{C}(\boldsymbol{\theta})^{-1}\mathsf{Z},\tag{1}$$

where the covariance matrix  $C(\theta)$  has entries  $C(s_i - s_j; \theta)$ ,  $i, j = 1, \ldots, n$ . The maximum likelihood estimator of  $\theta$  is the value  $\widehat{\theta}$  that maximizes (1).









 ${\cal H}$ -matrix approximations of the exponential covariance matrix (left), its hierarchical Cholesky factor  $\tilde{\rm L}$  (middle), and the zoomed upper-left corner of the matrix (right), n=4000,  $\ell=0.09$ ,  $\nu=0.5$ ,  $\sigma^2=1$ .

6

## What will change after $\mathcal{H}$ -matrix approximation?



#### Approximate C by $C^{\mathcal{H}}$

- 1. How the eigenvalues of C and  $C^{\mathcal{H}}$  differ ?
- 2. How det(C) differs from  $det(C^{\mathcal{H}})$ ? [Below]
- 3. How L differs from  $L^{\mathcal{H}}$  ? [Mario Bebendorf et al]
- 4. How  $C^{-1}$  differs from  $(C^{\mathcal{H}})^{-1}$ ? [Mario Bebendorf et al]
- 5. How  $\tilde{\mathcal{L}}(\theta, k)$  differs from  $\mathcal{L}(\theta)$ ? [Below]
- 6. What is optimal H-matrix rank? [Below]
- 7. How  $\theta^{\mathcal{H}}$  differs from  $\theta$ ? [Below]

For theory, estimates for the rank and accuracy see works of Bebendorf, Grasedyck, Le Borne, Hackbusch,...

#### Details of the identification

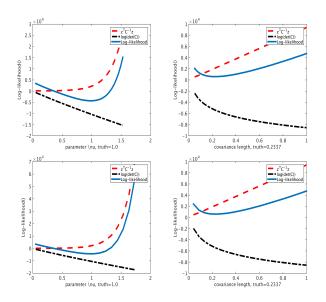


To maximize the log-likelihood function we use the Brent's method (combining bisection method, secant method and inverse quadratic interpolation) or any other.

- 1.  $C(\theta) \approx \widetilde{C}(\theta, \varepsilon)$ .
- 2.  $\widetilde{C}(\boldsymbol{\theta}, k) = \widetilde{L}(\boldsymbol{\theta}, k)\widetilde{L}(\boldsymbol{\theta}, k)^T$
- 3.  $Z^T \widetilde{C}^{-1} Z = Z^T (\widetilde{L} \widetilde{L}^T)^{-1} Z = v^T \cdot v$ , where v is a solution of  $\widetilde{L}(\theta, k) v(\theta) := Z$ .

$$\log \det\{\widetilde{\mathsf{C}}\} = \log \det\{\widetilde{\mathsf{LL}}^T\} = \log \det\{\prod_{i=1}^n \lambda_i^2\} = 2\sum_{i=1}^n \log \lambda_i,$$

$$\widetilde{\mathcal{L}}(\boldsymbol{\theta}, k) = -\frac{N}{2} \log(2\pi) - \sum_{i=1}^{N} \log\{\widetilde{\mathsf{L}}_{ii}(\boldsymbol{\theta}, k)\} - \frac{1}{2} (\mathsf{v}(\boldsymbol{\theta})^T \cdot \mathsf{v}(\boldsymbol{\theta})). \tag{2}$$

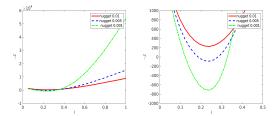


Dependence of log-likelihood ingredients on parameters, n = 4225.

#### Remark: stabilization with nugget



To avoid instability in computing Cholesky, we add:  $\widetilde{C}_m = \widetilde{C} + \delta^2 I$ . Let  $\lambda_i$  be eigenvalues of  $\widetilde{C}$ , then eigenvalues of  $\widetilde{C}_m$  will be  $\lambda_i + \delta^2$ ,  $\log \det(\widetilde{C}_m) = \log \prod_{i=1}^n (\lambda_i + \delta^2) = \sum_{i=1}^n \log(\lambda_i + \delta^2)$ .



(left) Dependence of the negative log-likelihood on parameter  $\ell$  with nuggets  $\{0.01, 0.005, 0.001\}$  for the Gaussian covariance. (right) Zoom of the left figure near minimum; n=2000 random points from moisture example, rank k=14,  $\sigma^2=1$ ,  $\nu=0.5$ .



#### Theorem (1)

Let  $\widetilde{C}$  be an  $\mathcal{H}$ -matrix approximation of matrix  $C \in \mathbb{R}^{n \times n}$  such that

$$\rho(\widetilde{\mathsf{C}}^{-1}\mathsf{C}-\mathsf{I})\leq \varepsilon<1.$$

Then

$$|log|C| - log|\widetilde{C}|| \le -nlog(1 - \varepsilon),$$
 (3)

Proof: See [Ballani, Kressner 14] and [Ipsen 05]. Remark: factor *n* is pessimistic and is not really observed numerically.



#### Theorem (2)

Let  $\widetilde{C} \approx C \in \mathbb{R}^{n \times n}$  and Z be a vector,  $\|Z\| \le c_0$  and  $\|C^{-1}\| \le c_1$ . Let  $\rho(\widetilde{C}^{-1}C - I) \le \varepsilon < 1$ . Then it holds

$$\begin{split} |\widetilde{\mathcal{L}}(\boldsymbol{\theta}) - \mathcal{L}(\boldsymbol{\theta})| &= \frac{1}{2}(log|\mathsf{C}| - log|\widetilde{\mathsf{C}}|) + \frac{1}{2}|\mathsf{Z}^{\mathrm{T}}\left(\mathsf{C}^{-1} - \widetilde{\mathsf{C}}^{-1}\right)\mathsf{Z}| \\ &\leq -\frac{1}{2} \cdot nlog(1 - \varepsilon) + \frac{1}{2}|\mathsf{Z}^{\mathrm{T}}\left(\mathsf{C}^{-1}\mathsf{C} - \widetilde{\mathsf{C}}^{-1}\mathsf{C}\right)\mathsf{C}^{-1}\mathsf{Z}| \\ &\leq -\frac{1}{2} \cdot nlog(1 - \varepsilon) + \frac{1}{2}c_0^2 \cdot c_1 \cdot \varepsilon. \end{split}$$

#### $\mathcal{H}$ -matrix approximation

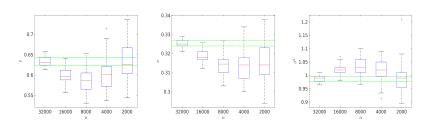


Denote accuracy in each sub-block by  $\varepsilon$ , n=16641,  $\nu=0.5$ , c.r.=compression ratio.  $\log |C|=2.63$  for  $\ell=0.0334$  and  $\log |C|=6.36$  for  $\ell=0.2337$ .

ε	$ log C  - log \widetilde{C}  $	$\left  \frac{\log C  - \log \widetilde{C} }{\log \widetilde{C} } \right $	$\ C-\widetilde{C}\ _{F}$	$\frac{\ C-\widetilde{C}\ _2}{\ C\ _2}$	$\ \mathbf{I} - (\mathbf{L}\widetilde{\mathbf{L}})^{-1}\mathbf{C}\ _2$	c.r. in %
$\ell = 0.0334$				<del>-</del>		
1e-1	3.2e-4	1.2e-4	7.0e-3	7.6e-3	2.9	9.16
1e-2	1.6e-6	6.0e-7	1.0e-3	6.7e-4	9.9e-2	9.4
1e-4	1.8e-9	7.0e-10	1.0e-5	7.3e-6	2.0e-3	10.2
1e-8	4.7e-13	1.8e-13	1.3e-9	6e-10	2.1e-7	12.7
$\ell = 0.2337$						
1e-4	9.8e-5	1.5e-5	8.1e-5	1.4e-5	2.5e-1	9.5
1e-8	1.45e-9	2.3e-10	1.1e-8	1.5e-9	4e-5	11.3

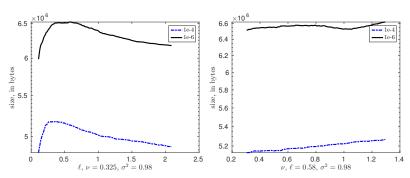
## Boxplots for unknown parameters





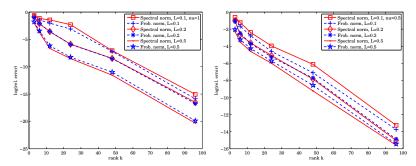
Moisture data example. Boxplots for the 100 estimates of  $(\ell, \nu, \sigma_2)$ , respectively, when n=32K, 16K, 8K, 4K, 2K.  $\mathcal{H}$ -matrix with a fixed rank k=11. Green horizontal lines denote 25% and 75% quantiles for n=32K.





(left) Dependence of the matrix size on the covariance length  $\ell$ , and (right) the smoothness  $\nu$  for two different  $\mathcal{H}$ -accuracies  $\varepsilon = \{10^{-4}, 10^{-6}\}$ 





Convergence of the  $\mathcal{H}$ -matrix approximation errors for covariance lengths  $\{0.1,0.2,0.5\}$ ; (left)  $\nu=1$  and (right)  $\nu=0.5$ , computational domain  $[0,1]^2$ .



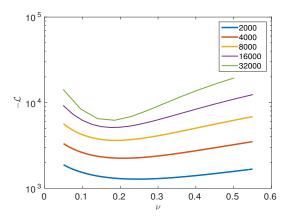


Figure: Dependence of negative log-likelihood function on different number of locations  $n = \{2000, 4000, 8000, 16000, 32000\}$  in log-scale.

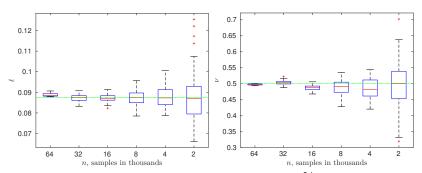
## Time and memory for parallel ${\mathcal H}$ -matrix approximation



Maximal # cores is 40,  $\nu = 0.325$ ,  $\ell = 0.64$ ,  $\sigma^2 = 0.98$ 

n	Ĉ			$\widetilde{L}\widetilde{L}^{ op}$		
	time	size	kB/dof	time	size	$\ I - (\tilde{L} \tilde{L}^{ op})^{-1} \tilde{C} \ _2$
	sec.	MB		sec.	MB	
32.000	3.3	162	5.1	2.4	172.7	$2.4 \cdot 10^{-3}$
128.000	13.3	776	6.1	13.9	881.2	$1.1\cdot 10^{-2}$
512.000	52.8	3420	6.7	77.6	4150	$3.5\cdot 10^{-2}$
2.000.000	229	14790	7.4	473	18970	$1.4\cdot 10^{-1}$





Synthetic data with known parameters  $(\ell^*, \nu^*, \sigma^{2^*}) = (0.5, 1, 0.5)$ . Boxplots for  $\ell$  and  $\nu$  for  $n=1,000\times\{64,32,...,4,2\}$ ; 100 replicates.



Let  $(Z_1, Z_2)$  has mean zero and a stationary covariance

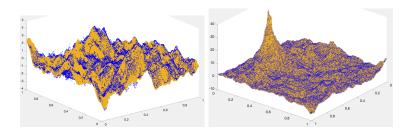
$$C = \begin{bmatrix} C_{11} & C_{12} \\ C_{21} & C_{22}, \end{bmatrix}$$

We compute predicted values

$$Z_2 = C_{21}C_{11}^{-1}Z_1$$

 $Z_2$  has the conditional distribution with the mean value  $C_{21}C_{11}^{-1}Z_1$  and the covariance matrix  $C_{22}-C_{21}C_{11}^{-1}C_{12}$ .





Prediction for two datasets. The yellow points at 900.000 locations were used for training and the blue points were predicted at 100.000 new locations.

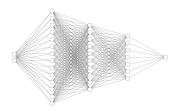
## Machine learning methods to make prediction



**k-nearest neighbours (kNN):** For each point find its k nearest neighbors  $x_1, \ldots, x_k$ , and set:  $\hat{y} = \frac{1}{k} \sum_{i=1}^k y_i$ .

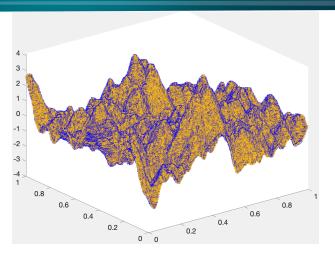
Random Forest (RF): a large number of decision (or regression) trees are generated independently on random sub-samples of data. The final decision for x is calculated over the ensemble of trees by averaging the predicted outcomes.

Deep Neural Network (DNN): fully connected neural network



Input layer consists of two neurons (input feature dimensionality), and output layer consists of one neuron (predicted feature dimensionality).





Prediction obtained by the kNN method. The yellow points at 900.000 locations were used for training and the blue points were predicted at 100.000 new locations. One can see a very well alignment of both.



- ▶ With H-matrices you can approximate Matérn covariance matrices, Gaussian log-likelihoods, identify unknown parameters and make predictions
- ▶ MLE estimate and predictions depend on  $\mathcal{H}$ -matrix accuracy
- parameter identification problem has multiple solutions
- ► Investigated dependence of *H*-matrix approximation error on the estimated parameters
- ► Each of ML methods needs fine-tuning stage to optimize its hyperparameters or architecture.

## Open questions and TODOs



- A good starting point for optimization is needed
- ▶ a "preconditioner" is needed
- → H-matrices become expensive for large number of parameters to be identified
- error estimates are needed



## All test are reproducible

#### https://github.com/litvinen/large\_random\_fields.git

- A. Litvinenko, R. Kriemann, M.G. Genton, Y. Sun, D.E. Keyes, HLIBCov: Parallel hierarchical matrix approximation of large covariance matrices and likelihoods with applications in parameter identification, MethodsX 7, 100600, 2020
- A. Litvinenko, Y. Sun, M.G. Genton, D.E. Keyes, Likelihood approximation with hierarchical matrices for large spatial datasets, Computational Statistics & Data Analysis 137, 115-132, 2019
- M. Espig, W. Hackbusch, A. Litvinenko, H.G. Matthies, E. Zander, Iterative algorithms for the post-processing of high-dimensional data, Journal of Computational Physics 410, 109396, 2020

#### Literature



- A. Litvinenko, D. Keyes, V. Khoromskaia, B.N. Khoromskij, H.G. Matthies, Tucker tensor analysis of Matérn functions in spatial statistics, Computational Methods in Applied Mathematics 19 (1), 101-122, 2019
- B. N. Khoromskij, A. Litvinenko, H.G. Matthies, Application of hierarchical matrices for computing the Karhunen-Loéve expansion, Computing 84 (1-2), 49-67, 31, 2009
- W. Nowak, A. Litvinenko, Kriging and spatial design accelerated by orders of magnitude: Combining low-rank covariance approximations with FFT-techniques, Mathematical Geosciences 45 (4), 411-435, 2013
- M. Espig, W. Hackbusch, A. Litvinenko, H.G. Matthies, E. Zander, Efficient analysis of high dimensional data in tensor formats, Sparse Grids and Applications, 31-56, Springer, Berlin, 2013