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Filtering remotely sensed chlorophyll concentrations in the Red Sea using a space-time covariance model and a Kalman Filter

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Abstract

A statistical model is proposed to filter satellite-derived chlorophyll concentration from the Red Sea, and to predict future chlorophyll concentrations. The seasonal trend is first estimated after filling missing chlorophyll data using an Empirical Orthogonal Function (EOF)-based algorithm (Data Interpolation EOF). The anomalies are then modeled as a stationary Gaussian process. A method proposed by Gneiting [1] is used to construct positive-definite space-time covariance models for this process. After choosing an appropriate statistical model and identifying its parameters, Kriging is applied in the space-time domain to make a one step ahead prediction of the anomalies. The latter serves as the prediction model of a reduced-order Kalman filter, which is applied to assimilate and predict future chlorophyll concentrations. The proposed method decreases the root mean square (RMS)

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prediction error by about 11% compared with the seasonal average.

1. Introduction

The Red Sea is an elongated basin situated between the Asian and the African shelves, connected to the Mediterranean Sea in the north through the Suez Canal, and to the Gulf of Aden in the south through the Strait of Bab el Mandeb [2]. It is one of the warmest and most saline seas in the world, with a rich ecosystem that has adapted to these extreme conditions [3]. This unique natural resource is however threatened by an abrupt increase of temperature since 1994 [4].

Phytoplankton are small, unicellular, photosynthetic algae. They are the primary producers for marine ecosystems, and at the base of the marine food chain. Phytoplankton concentration is therefore important for fisheries [5]. By fixing atmospheric CO2 and sinking to form sediment at the bottom of the sea, phytoplankton also acts as a biological pump. This phenomenon is crucial to understanding climate change [6]. In the Red Sea, phytoplankton is particularly important to the extensive coral reefs along its shores.

The Red Sea is generally deficient in major inorganic nutrients, and its productivity is relatively low [7, 8]. The high productivity observed in the southern Red Sea is attributed to the intrusion of nutrient-rich waters from the Gulf of Aden [7, 8]. Red Sea chlorophyll concentrations follow seasonal patterns, with a winter bloom following a weak summer productivity. Considerable interannual variability in chlorophyll concentrations has also been
observed [9]. However, the Red Sea ecosystem has not yet been fully explored, and very few in-situ measurements have been collected in its basin [10], increasing the need for remotely sensed data. Satellite observations of chlorophyll concentrations have been shown to be reliable datasets to study the primary productivity of the oceans [11] and, up to this point, they constitute the basis of several studies in the Red Sea [7, 10, 9]. Here we are interested in forecasting chlorophyll concentrations in the Red Sea, and we therefore need a dynamical model to mimic its evolution.

There are two approaches to chlorophyll modeling: deterministic or data-driven (statistical). A broad range of deterministic models has been developed by the marine ecosystem research community, from the very simple NPZ model with only nutrients, phytoplankton and zooplankton as state variables, to much more complex models, such as the European Regional Seas Ecosystem Model (ERSEM) [12]. The latter distinguishes functional phytoplankton and zooplankton groups, and models the complete cycling of different nutrient groups and O$_2$ and CO$_2$, including the effect of higher trophic groups. An example of a 3D ERSEM coupled model has recently been implemented in the Red Sea [8]. Simulating the ecosystem with such a model can be however very challenging, as it requires the coupling to an ocean circulation model, which provides the physical forcing. Configuring these models further requires considerable efforts and expertise [13], because of the high number of parameters (over 50 for ERSEM [14]). This makes such models difficult to calibrate and validate, since there are usually not
enough observations to constrain the parameters [15].

An alternative approach is to follow a statistical framework to model the space-time evolution of chlorophyll concentrations. Space-time statistical methods have not been used yet in this field, which has so far relied on time-series observations. Artificial neural networks were applied to forecasting algal blooms in freshwater and marine systems [16, 17], and generalized additive models have been used for finding explanatory variables for the chlorophyll concentrations in the Pagasitikos Gulf and the subartic North Atlantic [18, 19].

Geostatistical spatio-temporal models are extensions of the spatial classical geostatistical methods [20]. These methods consider space-time data as the realization of a Gaussian process, from which a mean and a covariance function can be estimated. Although, in most applications, such a stochastic modeling approach is not based on a dynamical framework, geostatistical methods may capture some patterns in the data and avoid the difficulties of developing dynamical models [21, 20]. These methods have been widely employed in meteorology to model the surface temperature over land and oceans [22, 23], in an ecological context to study moth populations [24], and in environment applications to characterize soil and pollution [25, 26].

The first proposed spatio-temporal geostatistical model was based on separable covariance functions [20]. Such functions can either be the product, or the sum, of a purely temporal covariance model and a purely spatial covariance model. These covariance models are convenient but have non-physical
properties, limiting their use in many situations [20]. As a result, significant
research has been recently conducted to construct nonseparable families of
covariance functions. [27, 1] proposed methods to construct such families
from known temporal and spatial covariance functions. The method pro-
posed in [1] is appealing because of its modularity and interpretability.

We adopt the aforementioned method to model the anomaly fields of
chlorophyll concentration in the Red Sea. To the best of our knowledge,
this is the first application of a space-time geostatistical modeling approach
to model and predict remotely-sensed chlorophyll data. After filling the
data with an Empirical Orthogonal Function (EOF)-based method (DINEOF
[28, 29]), we model it as the realization of a spatio-temporal Gaussian process
and estimate the mean and covariance functions. Based on these, one may
forecast chlorophyll concentration using Kriging by conditioning on past ob-
servations [20]. However, this approach is not computationally efficient, as it
requires inverting a large matrix at every time step when many observations
are available, which is the case here. We therefore propose a new approach,
in which we reformulate the problem within a state-space framework and
use an efficient reduced-order Kalman filter [30] to filter and forecast chloro-
phyll. The state-space system is further expanded to take into account the
colored model noise [31]. In contrast with the space-time hierarchical models
proposed in [32, 33], the linear dynamical model is derived from a space-time
geostatistical model, avoiding the need to directly estimate the transition
matrix. Furthermore, the dimension reduction is not part of the modeling,
but is implemented at the filtering level through a reduced-order Kalman filter.

The paper is organized as follows. Section 2 describes the satellite data. Section 3 discusses the methodology for the data filling and geostatistical modeling. Section 4 derives the state-space formulation and introduces the space-time Kalman filtering problem and its solution. The experimental setup and numerical results are presented and discussed in section 4. Concluding remarks are provided in section 5.

2. Data and preprocessing

Satellite data provide chlorophyll (CHL) concentrations at high with spatial and temporal resolutions, making them particularly valuable to the Red Sea, where very few in situ data are available.

Level-3 mapped data from the NASA SeaWiFS (Sea-Viewing Wide Field-of-View Sensor) satellite sensor are used in this study. The dataset is publicly available at http://oceancolor.gsfc.nasa.gov. In this study, we use the 9km resolution mapped weekly averages from January 1998 to December 2007 (460 time steps). At each time step, a $133 \times 188$ pixel map is available for a domain extending from longitudes between $33^\circ$E and $44^\circ$E and latitudes between $12^\circ$N and $28^\circ$N, of which 5635 pixels correspond to actual Red Sea surface (see Figure 1(a)). A log-transformation is applied in order to obtain an approximately Gaussian distribution [34]. Pixels with too few observations are discarded, and a control quality check is applied to remove
outliers [35].

Remotely sensed CHL may have missing data because of cloud coverage. The cloud variability in the Red Sea follows a seasonal cycle. Figure 1(c) shows that the cloud coverage is particularly pronounced during summers due to the monsoon and it is sparse during winters. The cloud coverage is, however, not homogeneous over the Red Sea. It is much more pronounced in the south (figure 1(b)). In this region, almost no data are available during summers.

The dataset is separated into a training set, composed of the first seven years of data, and a validation set, containing the remaining three years. Both datasets were preprocessed and log-transformed in the same way. The computations for the data filling and for the covariance model selection and estimation are based only on the training dataset. The testing dataset is used to validate the model predictions outside the training period.

3. Data filling and modeling

3.1. Data filling

The DINEOF (Data Interpolating Empirical Orthogonal Function) is an EOF-based, recursive method for the reconstruction of data matrices with missing values [28, 29]. It estimates the values of the missing data by successive singular values decompositions (SVD) of a given data matrix and truncated reconstructions. The advantage of this method is that it does not require any a priori information about the data. It has been successfully
used reconstructing incomplete chlorophyll datasets in various regions of the ocean [36, 37, 38].

Let $X$ be an $m \times k$ data matrix, where each row is an observation, and each column is a variable. The average of each variable is removed from the corresponding observation in order to center the data matrix. The missing values are initially filled with 0s. Then, until the missing values have converged, the following steps are repeated. An SVD is first applied to the data matrix: $X = U \Sigma V^T$, with $U$ an $m \times m$ unitary matrix, $\Sigma$ an $m \times k$ diagonal matrix and $V$ a $k \times k$ unitary matrix. The missing values are then replaced by the truncated reconstruction order $n$ of the data matrix:

$$\{X\}_{i,j} = \{U^{(n)} \Sigma^{(n)} (V^{(n)})^T\}_{i,j},$$

for $i, j$ indices of the missing values, with $U^{(n)}$ the $m \times n$ matrix composed of the $n$ first columns of $U$, $V^{(n)}$ the $k \times n$ matrix composed of the $n$ first columns of $V$, and $\Sigma^{(n)}$ the $n \times n$ diagonal matrix with the $n$ largest eigenvalues on its diagonal. It is assumed that the eigenvalues and eigenvector are sorted by decreasing order of eigenvalues. In [29], the authors introduced the filtering of the temporal covariance matrix as a way of reducing spurious oscillations that may appear when the data are sparsely sampled in time. This filtering is controlled by the parameter of the Laplacian filter and the number of times the filter is applied.

The values of the DINEOF parameters are determined following the method outlined in [29]. As in the former, the smoothing parameter of the Laplacian filter is set to $0.01d^2$, where $d$ is the laps of time between two successive observations. The number of modes in the truncation and the
The number of times the filter is applied are chosen following a cross-validation technique. A random subset of observed values is taken from $X$ and assumed to be missing before the DINEOF is applied. The algorithm is then run with different numbers of iterations (1, 3, 10, 30, 100) and orders of truncation (from 2 to 50). The set of parameters minimizing the RMS error over the cross-validation data is chosen as the best number of iterations and order of truncation. The approach of [39] is followed to select a cross-validation dataset. The idea is to apply cross-validation over contiguous regions, instead of applying it to randomly selected points. The shape of the cross-validation regions are randomly selected from the shape of missing data regions in the dataset. We set aside the cross-validation data by randomly applying these shapes to the data until 3% of the total number of data points has been extracted.

3.2. Geostatistical modeling

The chlorophyll concentration data are modeled as the realization of a space-time random Gaussian process:

$$Z(s;t), (s;t) \in \mathbb{R}^2 \times \mathbb{R}. \quad (1)$$

Such a process is entirely characterized by the mean function $\mu(s;t) = E(Z(s;t))$, and the covariance function $K(s,r;t,q) = \text{cov}(Z(s;t), Z(r;q))$. The process is further assumed to be stationary, such that the covariance function can be written as $K(s,r;t,q) = C(r - s, q - t)$. 
Covariance functions should be positive-definite, meaning that for any ensemble of space-time coordinates, \( \{(s_i; t_i)\}_{i=1,...,k} \), and real coefficients, \( d_1, ..., d_k \), the following condition should hold:

\[
\sum_{i=1}^{k} \sum_{j=1}^{k} d_id_j C(s_i - s_j; t_i - t_j) \geq 0.
\]  

(2)

One way to enforce positive-definiteness in practice is to assume that the covariance function belongs to a parametric family of covariance functions, denoted by \( C(h; u|\theta) \), where \( h \) is the spatial lag, \( u \) is the temporal lag, and \( \theta \) is the vector of parameters to be estimated [40].

3.3. Mean function estimation and anomalies

As can be seen in Figure 2, the Red Sea chlorophyll data exhibit a strong seasonality. The climatological means explain 50% of the variability in the data. We thus choose to model the mean function \( \mu(s; t) \) as the climatological average. Once the data are filled using the DINEOF algorithm, the seasonal signal is estimated for each week, with the weekly average computed from the training data.

The anomalies are then computed by subtracting the weekly averages from the data. The CHL anomalies of the first 8 weeks of data are displayed in Figure 3. There are large regions of similar colors indicating spatial correlations. Moreover, two maps adjacent in time display similar patterns, suggesting that the dataset is also correlated in time and justifying the use of space-time covariance functions to model CHL anomalies. From the anomaly
time-series at three locations, plotted in Figure 4, one may conclude that the
data have been successfully detrended. Since no increasing pattern appears,
we conclude, as in [27], that the assumption of stationarity for the covariance
model is appropriate.

3.4. Construction and fitting of the covariance model

For space-time processes, families of covariance functions are typically
built by combining known spatial and temporal covariance models. Sep-
arable models are the simplest example of this approach, taking the form
\[ C(h; u|\theta) = C^1(h|\theta_1)C^2(t|\theta_2), \]
where \( C^1 \) and \( C^2 \) are respectively pure spa-
tial and temporal covariance models parametrized by \( \theta_1 \) and \( \theta_2 \). However,
realizations from such families of covariance models were shown to exhibit
non-physical behaviors [1, 27]. Research has been conducted to develop meth-
ods for constructing non-separable covariance functions to overcome these
limitations [27, 1].

In this work, families of space-time covariance functions are constructed
following an approach proposed in [1]. We used this method because of its
flexibility and simplicity, since no transformation of the covariance function
is involved. Given any completely monotonous function, \( \varphi(t), t \geq 0 \), and any
positive function with a completely monotonous derivative, \( \psi(t), t \geq 0 \), the
following space-time covariance function is valid (i.e., positive-definite):
\[
C(h; u) = \sigma^2 \psi(|u|^2)\varphi\left(\frac{||h||^2}{\psi(|u|^2)}\right), \quad (h, u) \in \mathbb{R}^2 \times \mathbb{R}. \tag{3}
\]
In general, \( \varphi \) is defined as a temporal covariance function, and \( \psi \) as a spatial covariance model, e.g., the exponential model or the Matérn model. Tables 1 and 2 present some examples of these functions that were introduced in [1]. To obtain more specialized families, a nugget term can be added.

The typical approach for estimating the parameters of the covariance function is by fitting it to the empirical space-time covariance matrix [40]. The empirical covariance function is first estimated using the formula:

\[
\hat{C}(h(l); u) = \frac{1}{2|N(h(l); u)|^{\times}} \sum_{(i,j,t,t') \in N(h(l); u)} (Z(s_i; t) - \hat{\mu}(s_i; t))(Z(s_j; t') - \hat{\mu}(s_j; t')) ,
\]

where \((s_i, t)\) and \((s_j, t')\) are spatio-temporal locations at which \(Z\) is observed, and \(\hat{\mu}\) is an estimate of the mean function. \(\mathbb{R}^2\) has been divided into a finite number \(N_s\) of bins, \(\text{Bin}(h(l))\), each of which has a representative lag, \(h(l)\). Here, bins are considered of the form \(k \in \text{Bin}(h(l))\) if \((l - 1) \ast \delta h \leq ||k|| \leq l\delta h, l = 0, ..., L\), with \(\delta\) a fixed interval size and \(L\) the number of retained intervals. The parameters are then estimated by fitting the covariance model using the Weighted Least Squares (WLS) approach. This method is popular due to its simplicity and proven efficiency [41]. In practice, the estimator of \(\theta\) minimizes the weighted sum of the squares of errors:

\[
S(\theta) = \sum_{l=1}^{N_s} \sum_{u=0}^{T} w_{l,u} \left( \hat{C}(h(l); u) - C(h(l); u|\theta) \right)^2 ,
\]

(6)
with weights $w_{l,u} = \frac{|N(l(u_l));u|}{(1-C(l(u_l));u)}$. $T$ is the maximum temporal lag considered. $\phi$ and $\psi$ are chosen from the functions presented in Tables 1 and 2, by separately fitting the covariance function in space and time for all possible choices of $\phi$ and $\psi$. The choice of the functions is then made based on the quality of the fit.

4. State-space formulation and filtering

So far, the CHL data are modeled as $Z(s; t) = \mu(s; t) + a(s; t)$, where $\mu$ is a deterministic function representing the seasonality and $a$ is a zero-mean space-time Gaussian process representing the weekly anomalies, with the covariance function estimated as described in the previous section. However, a state-space model needs to be formulated to apply the Kalman filter. This is done by identifying a CHL underlying process, $Y(s; t)$, distinct from the data process $Z(s; t)$. These processes are then discretized in space and time to obtain the state-space formulation, on which a reduced-order variant of the Kalman filter is applied.

4.1. State-space modeling

To derive the state-space model, we start from the following model:

$Y(s; t) = \mu(s; t) + a(s; t)$, where $s, t$ are observable locations, \hspace{1cm} (7)$

$Z(s; t) = Y(s; t) + \varepsilon(s; t)$, where $s, t$ are observed locations. \hspace{1cm} (8)$

\hspace{1cm}
In this formulation, \( \varepsilon(s; t) \) is a white noise process representing the measurement errors, \( a(s; t) \) is the anomaly process with the covariance matrix estimated as described in the previous section, \( \mu(s; t) \) is the seasonal mean function, \( Y(s; t) \) is the underlying CHL log-concentration process, and \( Z(s; t) \) the data process.

To build a state-space model for filtering and forecasting the spatial variability of CHL in time, once the covariance function is estimated, we apply Kriging. In classical geostatistics, Kriging interpolates available observations to provide the best linear unbiased estimate of a spatial process at unobserved locations [40]. This technique generalizes to space-time Gaussian processes, particularly for forecasting. The equations are obtained by conditioning the anomalies at time \( t \) by the anomalies at time \( t - 1 \). Using the vectorial notations \( \mathbf{a}_t = \{a(s_{i,t})\}_{i=1,...,k} \), where \( s_{i,t} \) is the spatial location of the \( i \)-th observation at time \( t \), \( [\mathbf{a}_t^T, \mathbf{a}_{t-1}^T]^T \) is a Gaussian vector. Therefore, by conditioning

\[
\mathbf{a}_t|\mathbf{a}_{t-1} \sim N(\mathbf{M}\mathbf{a}_{t-1}, \mathbf{Q}),
\]

where

\[
\mathbf{M} = \mathbf{C}_1\mathbf{C}_0^{-1},
\]

\[
\mathbf{Q} = \mathbf{C}_0 - \mathbf{C}_1\mathbf{C}_0^{-1}\mathbf{C}_1,
\]
with \( C_0 = E[a_t a_t^T] = \{C(s_{i,t} - s_{j,t}), 0\}_{i,j=1,...,k} \) and \( C_1 = E[a_t a_{t-1}^T] = \{C(s_{i,t} - s_{j,t-1}), 1\}_{i,j=1,...,k} \), the following recursive state-space model can be derived:

\[
y_t = \mu_t + M(y_{t-1} - \mu_{t-1}) + \eta_t, \tag{13}
\]

\[
\tilde{z}_t = H_t y_t + \varepsilon_t, \tag{14}
\]

with \( y_t = a_t + \mu_t \) the CHL process discretized in space, \( \mu_t \) is the vector seasonal component, \( H_t \) is the observation operator that returns the CHL concentration at the observed location, \( \eta_t \) represents the model error, and \( \varepsilon_t \) represents the measurement error. \( y_t \) is a vector of fixed size that represents the whole model domain (Red Sea), whereas \( \tilde{z}_t \) has a variable size equal to the number of available observations at time \( t \).

The model can be reformulated such that only the anomalies are filtered and \( a_t \) is the state vector:

\[
a_t = Ma_{t-1} + \eta_t, \tag{15}
\]

\[
\tilde{z}_t = \tilde{z}_t - H_t \mu_t = H_t a_t + \varepsilon_t. \tag{16}
\]

This system is equivalent to the preceding one, but is more practical to implement. Given \( \eta_t \sim \mathcal{N}(0, Q) \) and assuming \( \varepsilon_t \sim \mathcal{N}(0, \sigma_{\text{obs}}^2 I) \), independent and identically distributed (i.i.d.), with \( \sigma_{\text{obs}}^2 \) a fixed constant that is tuned by minimizing the RMS prediction error.
4.2. Low-rank Kalman filter

Two issues need to be tackled before using the time evolution model for predicting the anomalies. First, the consequent amount of missing data in the CHL satellite observations makes it difficult to obtain frequent initial CHL concentrations to integrate the model forward in time for forecasting, and second, the model is linear with eigenvalues smaller than one\textsuperscript{1} in absolute values, making it inappropriate for long-term predictions. The latter means that CHL model forecasts would decrease exponentially in time, quickly becoming close to 0. The Kalman filter solves both problems by recursively assimilating the observations and providing an optimal estimate, in the mean-square sense, of the anomalies to start a new forecast cycle. The filter operates in two steps to compute the best linear estimate of the state of a linear dynamical model given past observations [42].

\begin{itemize}
  \item \textit{Forecast step:} Starting from the best available estimate of the state, $a_t^{f-1}$, and the associated error covariance matrix, $P_t^{f-1}$, at a given time, $t - 1$, the forecast state, $a_t^f$, and its error covariance matrix, $P_t^f$, are obtained by integrating the model forward to the time of the next available observation:

  \begin{align}
    a_t^f &= M a_t^{f-1}, \quad (17) \\
    P_t^f &= M P_t^{f-1} M^T + Q. \quad (18)
  \end{align}
\end{itemize}

\textsuperscript{1}This was verified numerically in the present work. We are not aware of a general result.
• **Update step:** Once a new observation, \( z_t \), is available, the forecast state, \( a_f^t \), and its error covariance, \( P_f^t \), are updated to their analysis counterparts, \( a_a^t \), and, \( P_a^t \), as:

\[
a_a^t = a_f^t + K_t(z_t - H_t a_f^t),
\]

(19)

\[
P_a^t = (I - K_t H_t) P_f^t.
\]

(20)

\[
K_t = P_f^t H_t^T (H_t P_f^t H_t^T + \sigma_{\text{obs}}^2 I)^{-1}.
\]

(21)

where \( K_t \) is known as the Kalman gain.

The inversion in the calculation of the Kalman gain is computationally quite demanding because the number of observations can be as large as the size of the state. To speed up this computation, the reduced-order Kalman filter [30] is used. This filter approximates the covariance matrices as \( P_f^t = L U_f^t L^T \) and \( P_a^t = L U_a^t L^T \), where \( L \) is a \( n \times r \) matrix whose columns are the \( r \) leading EOFs (computed here with the DINEOF algorithm), and \( U_f^t \) and \( U_a^t \) are \( r \times r \) matrices. The inversion is then applied on the \( r \times r \) matrices \( U_f^t \) and \( U_a^t \), which considerably reduces the computational burden since \( r \ll n \) in practice [30]. Defining

\[
V = L^T Q L,
\]

(22)

\[
W = L^T M L,
\]

(23)
which respectively represent the projection of the model dynamics and the model error on the leading EOFs, the reduced-order Kalman filter equations for the forecast and update of the covariance matrix can be simplified by only updating $U_f^t$ and $U_a^t$ as follows:

\[
U_f^t = WU_a^{t-1}W^T + V, \tag{24}
\]

\[
(U_a^t)^{-1} = (U_f^t)^{-1} + L^T H^T R^{-1} H L. \tag{25}
\]

The Kalman gain can be then computed as

\[
K_t = \sigma_{\text{obs}}^{-2}L U_a^t L^T H^T. \tag{26}
\]

Otherwise, the forecast and update steps are identical to those of the Kalman filter.

4.3. Low-rank Kalman filter with colored noise

The model noise is correlated in time, whereas the Kalman filter assumes it to be white. Indeed, $\eta_t = a_t - M a_{t-1}$ is a space-time Gaussian process. One can show that $[\eta_t^T, \eta_{t-1}^T]^T$ is a Gaussian vector of mean 0 and covariance matrix

\[
\begin{bmatrix}
A_0 & A_1 \\
A_1^T & A_0
\end{bmatrix},
\]

with $A_0 = C_0 - C_1 C_0^{-1} C_1^T$, $A_1 = C_1 C_0^{-1} C_1 C_0^{-1} C_1 - C_2 C_0^{-1} C_1$. We can then predict $\eta_t$ by conditioning on the previous equation and derive the model $\eta_t = N \eta_{t-1} + \xi_t$, with $N = A_1 A_0^{-1}$ and $\xi_t \sim \mathcal{N}(0, A_0 - A_1 A_0^{-1} A_1)$. 

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To take into account this correlation in the Kalman filter, the state-space system is enlarged by including $\eta_t$ in the state vector [31], so the new state-space system becomes:

$$\hat{a}_t = \Gamma \hat{a}_{t-1} + \hat{\eta}_t,$$

$$z_t = \hat{H} \hat{a}_{t-1} + \varepsilon_t,$$

with $\hat{a}_t = [a_t^T, \eta_t^T]^T$, $\Gamma = \begin{bmatrix} M & I \\ 0 & N \end{bmatrix}$, $\hat{\eta}_t = [0, \xi_t^T]^T$ and $\hat{H}_t = [H_t, 0]$, on which the Kalman filter is applied. Of course, $\xi_t$ is again colored, but the procedure can be iterated until no significant correlation is left to exploit in the noise.

5. Results

Figure 5 summarizes the workflow of the experimental setup. The chlorophyll concentration is the sum of a seasonal component and an anomaly. The former is estimated using data filled by the DINEOF algorithm over a learning period (first seven years of CHL data). The anomalies are assumed to be a stationary Gaussian process whose space-time covariance matrix has been estimated. A reduced-order Kalman filter is then applied to perform one-step ahead predictions. These are compared with the observations over the three years of testing data to validate the system’s performance and results. The results of the DINEOF data filling algorithm are presented before a covari-
ance model for the data is chosen and fitted. The results of the filtering are
finally examined and analyzed.

5.1. DINEOF Analysis

Figure 6 plots the RMS error for the cross-validation dataset and for
different values of the number of smoothing iterations and EOFs. The lowest
RMS error is achieved with 24 EOFs and 30 iterations. These parameters
were therefore selected in DINEOF to fill the missing values in the training
dataset.

Figures 7(a) and 7(c) show the first spatial and associated temporal modes
resulting from the DINEOF analysis. Figure 7(c) exhibits a regular peak dur-
ing winters and a secondary peak of varying size during summers. These can
be associated with the winter bloom and the secondary summer bloom, both
described in [9]. The winter bloom has been linked to convective overturning
the northern Red Sea, and enhanced water intrusion from the Gulf of Aden
in the South, whereas the causes of summer bloom are still not completely
understood yet [9]. Figure 7(a) shows that the bloom is relatively homoge-
neous over the Red Sea, except in the southwest corner where the variation
is more pronounced.

Figures 7(b) and 7(d) plot the second spatial and temporal modes re-
spectively. The spatial mode shows a north-south contrast. The time series
displays a large peak around summer 2000, which corresponds to a large
positive anomaly taking place during this period (Figure 7(e)). Except for
the first two modes, all other modes tend to explain a local feature of the data. This makes the interpretation of the EOF analysis difficult and the convergence of the spectrum very slow as shown in Figure 7(f).

5.2. Covariance model estimation

Figure 8a plots the empirical space covariance function $\hat{C}(||h||, 0)$. Among the functions in Table 1, the ones that best fit this curve are selected. A WLS minimization is applied to fit the parameters $c, \gamma$ and $\nu$, as well as the noise variance $\sigma^2_{\text{spatial}}$ while ignoring the observations for $||h|| = 0$. This allows a nugget effect to be taken into account. The results are superimposed on Figure 8a. The Matérn model clearly fits poorly. By examining the residuals, we can obviously eliminate $\varphi_4$. Finally, between the two remaining candidates, both of which appear to be plausible, $\varphi_1$ is chosen since it involves fewer degrees of freedom.

Figure 8b plots the empirical time covariance function $\hat{C}(0; u)$. Among the functions in Table 2, we eliminate $\psi_2$, which defines a temporal covariance function that decreases very slowly, which is not realistic to model anomalies. We also eliminate $\psi_3$ which defines a temporal covariance function that do not tend to 0 for increasing time lags. We therefore use the temporal covariance function defined by $\psi_1$, to which we add a temporal nugget effect. We fit the parameters using WLS and display the results in Figure 8b, where the nugget effect can be clearly identified.

To build the space-time covariance function, we use the reparametrization
in [1] (see example 1), along with the building blocks in time and space that have been derived in Section 3. The reparametrization is used to separate the variables that control the temporal covariance function and those controlling the spatial covariance function. A single parameter $\beta$ controls the space-time interaction strength. A time-only nugget effect is also added, leading to the following covariance model:

$$C(h; u) = \begin{cases} 
\sigma^2 \exp(-c|h|^2\tau), & \text{if } u = 0, \\
\sigma^2 \alpha \tau^{|u|^2+1} \exp\left(-\frac{c|h|^2\tau}{(\alpha|u|^2+1)\beta}\right), & \text{otherwise,}
\end{cases}$$

(29)

with $0 < \tau \leq 1$.

The initial values for the WLS fitting are given by the results of the preceding purely spatial and temporal regressions. The results are shown in Table 3. A contour plot of the resulting covariance function is shown in Figure 9. Our fitted correlation model exhibits level curves very close to those of the empirical covariance model. It does not, however, reproduce their curvature.

5.3. Filtering

$\sigma_{\text{obs}}^2$ is first determined empirically by trial and error, chosen as the value that leads to the minimal averaged RMS error. $\sigma_{\text{obs}}^2 = 1$ is found to be a reasonable choice.

The anomaly model with a space-time covariance function helps decreasing the RMS error by nearly 11% over the test period ($p < 0.01$ using a
permutation test). As Figure 10a indicates, the improvement is most noticeable during periods with large anomalies. The boxplot in Figure 10b shows that the variability of errors is reduced with the proposed filtering approach. Since the modeling is purely statistical and is not based on physical quantities, it was not able to anticipate the start of a bloom. However, it successfully extrapolates the current estimate of the anomaly in space and time and improves the prediction compared to the seasonal component.

In Figures 11 (a to c), the predictions of the model are compared with the observations and the seasonal signal for a fall week in 2006. The model captures some differences with the seasonal regime, such as a larger northern region with a low CHL concentration extending south below 22°N, and a more intense bloom in the south. The usual seasonal dynamic is altered with an extension of the stratified regime in the north and a larger intrusion of the nutrient rich waters of the Gulf of Aden [9].

Figure 11 (d to f) plots a similar comparison for a winter week in 2006. The model captures a weak winter bloom in the northern half of the Red Sea. A similar pattern has been described in [7] for winter 1999, and it seems to be a common feature of El Nina years. The model also successfully captures a high CHL concentration in the south, and a lower than usual CHL concentration in the center [9].

Figure 12 plots the error variance as predicted by the filter and the actual prediction RMS error, computed as the difference between the model forecast and data, averaged over the three year training period at every point of
the grid. The predicted RMS error corresponds to the diagonals of the prediction covariance matrix as estimated by the reduced-order Kalman filter. The prediction RMS error is computed from the error between the model prediction and the observation. One can see that both values are close in the northern half of the Red Sea, but the RMS error is much larger in the south. This is caused by the lack of data and the fact that the dynamics in the south are different from those in the north, making the process non-stationary. Indeed, as shown in Figure 3, the anomalies in the south clearly exhibit smaller spatial and temporal correlation length scales compared with those in the north.

Another way to evaluate the filter’s behavior is to examine the distribution of the innovations and the increments [43]. The innovation corresponds to the difference: $z_t - H_t a_f^t$, whereas the increment corresponds to the difference: $H_t (a^o_t - a_f^t)$. Figure 13(c) shows that the innovation seems to be approximately normally distributed, as expected from a properly tuned Kalman filter. Figure 13(b) shows that the averaged innovation size decreases to a value close to zero as the filter assimilates the data over time, which also suggests that the filter is properly working. However, Figure 13(a) indicates that the increments of the filter are positively biased in some regions. This suggests that the model tends to underestimate the amount of CHL, which may be associated with the statistical model’s inability to forecast CHL blooms.

Figures 14 (a), (b) and (c) plot the spatially averaged correlations between the observations and the seasonal predictions, the Kalman filter forecasts and
the analyses, respectively. Compared with the seasonal correlations, one can see that the model improves the prediction skill over the entire Red Sea, particularly in its central part, with correlations ranging between 0.28 and 0.92. The filter further improves the prediction-data correlation, over the entire domain with correlations up to 0.96.

6. Discussion

We considered the filtering problem of satellite-derived chlorophyll (CHL) concentration in the Red Sea. This is a difficult problem, as ecological systems are highly nonlinear, and difficult to forecast. Moreover, the CHL dataset has a large number of missing data, particularly in the southern Red Sea, during summer. We proposed a data-driven approach in which the CHL spatio-temporal evolution is modeled as a space-time Gaussian process. The DINEOF data-filling algorithm [39] was applied to compute an estimate of the seasonal signal in the data, which was used as the mean function of the process. To model the residual anomalies, the method proposed by Gneiting in [1] was applied to construct an appropriate family of covariance functions.

Based on a space-time Kriging formulation, a linear model was derived to model changes in the chlorophyll concentration. A reduced-order variant of the Kalman filter was then applied to forecast and filter the CHL concentration. The results of our experiments suggest that the proposed system works reasonably well, reducing the RMS error in CHL concentration prediction by about 11% as compared with the seasonal average computed from the
The reduction is more important during large anomalies, in particular in winter (Figure 10a). In general the model reduces the spread of the anomalies (Figure 10b). The RMS error reduction is however not spatially uniform and is more pronounced in the southern Red Sea (Figure 12). This can be caused by the number of missing data in this regions, as well as the fact that the anomaly process is not stationary over the whole domain.

The proposed approach is not difficult to implement, but requires some coding efforts. For the DINEOF, a standalone package is freely available online\(^2\). To the authors’ knowledge, there is currently no R package or library for fitting custom space-time covariance models. The reduced-order Kalman filter is straightforward to implement.

The proposed method requires the estimation of a very few parameters, which may prevent overfitting. Another advantage is its stability, as the anomaly prediction cannot grow over time. One problem with Kriging is that in the case of missing data over large areas, its prediction will be the mean function [44]. In this study, this problem is alleviated using the Kalman filter which always provides a prediction over the whole domain.

Forecasting and filtering CHL concentration in the Red Sea using remotely sensed data is challenging. Because of cloud coverage over the southern Red Sea during the summer, large areas remain unobserved. Moreover, the Red Sea is a heterogenous environment with different ecological and phys-\(^2\)http://modb.oce.ulg.ac.be/mediawiki/index.php/DINEOF
chical dynamics from north to south. We can therefore expect the anomalies
to be non-stationary. Fortunately, Kriging is robust to non-stationarity [44].
A problem with this correlation-based modeling is its inability to forecast
CHL blooms. Our linear model tends to underestimate the amount of CHL
concentration and to miss blooms in some regions, before the Kalman filter
reduces this error when new observations are available.

The proposed method can be further generalized by considering more
sophisticated mean functions. For example, one can use a linear model with
additional covariates, such as sea surface temperature, sea surface height,
thermocline depth or wind speed. Another way to improve the model is to
use non-stationary covariance functions, or to use non-Gaussian models to
predict blooms. These tasks will be considered in future studies.

Acknowledgment

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<thead>
<tr>
<th>Function</th>
<th>Parameters</th>
</tr>
</thead>
<tbody>
<tr>
<td>$\varphi_1(t) = \exp(-ct^\gamma)$</td>
<td>$c &gt; 0, 0 &lt; \gamma \leq 1$</td>
</tr>
<tr>
<td>$\varphi_2(t) = (2^{\nu-1}\Gamma(\nu))^{-1}(ct^{1/2})^\nu K_\nu(ct^{1/2})$</td>
<td>$c &gt; 0, \nu &gt; 0$</td>
</tr>
<tr>
<td>$\varphi_3(t) = (1 + ct^\gamma)^{-\nu}$</td>
<td>$c &gt; 0, 0 &lt; \gamma \leq 1, \nu &gt; 0$</td>
</tr>
<tr>
<td>$\varphi_4(t) = 2^\nu(\exp(ct^{1/2}) + \exp(-ct^{1/2}))^{-\nu}$</td>
<td>$c &gt; 0, \nu &gt; 0$</td>
</tr>
</tbody>
</table>
Table 2: Temporal building blocks proposed by [1]

<table>
<thead>
<tr>
<th>Function</th>
<th>Parameters</th>
</tr>
</thead>
<tbody>
<tr>
<td>( \psi_1(t) = (at^\alpha + 1)^\beta )</td>
<td>( a &gt; 0, 0 &lt; \alpha \leq 1, 0 \leq \beta \leq 1 )</td>
</tr>
<tr>
<td>( \psi_2(t) = \ln(at^\alpha + b) / \ln(b) )</td>
<td>( a &gt; 0, b &gt; 1, 0 &lt; \alpha \leq 1 )</td>
</tr>
<tr>
<td>( \psi_3(t) = (at^\alpha + b) / (b(at^\alpha + 1)) )</td>
<td>( a &gt; 0, 0 &lt; b \leq 1, 0 &lt; \alpha \leq 1 )</td>
</tr>
</tbody>
</table>
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<table>
<thead>
<tr>
<th>$\sigma^2$</th>
<th>$\tau$</th>
<th>$c$</th>
<th>$\gamma$</th>
<th>$a$</th>
<th>$\alpha$</th>
<th>$\beta$</th>
</tr>
</thead>
<tbody>
<tr>
<td>0.08</td>
<td>0.82</td>
<td>0.45</td>
<td>0.10</td>
<td>0.1</td>
<td>1.0</td>
<td>0.92</td>
</tr>
</tbody>
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