

Ecological clustering of the Red Sea and parallel 1D-ecological simulations

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Abstract

Abstract

1. Introduction

3D marine ecological models are useful....

...But expensive and difficult to run.

In this article we look at ways to simulate 3D ecosystems more cheaply by running many parallel 1D regional models.

We are going to test that idea on the Red Sea because....

We will also the hybrid-SEIK data assimilation scheme, because....

We will assimilate Chl data even if it is imperfect because it is the best available data for the Red Sea.

What we are going to do in this paper step by step.

What is new in this paper and why.

Introduce sections.

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13 **2. Data**

14 *2.1. CCI chlorophyll data*

15 *We use CCI chlorophyll data because it has more coverage..*

16 *With a quick look at the data this is what we see....*

17 *2.2. DINEOF*

18 *There are still missing data in CCI, so we use DINEOF for data filling be-*
19 *cause....*

20 *More or less this is the way DINEOF works....*

21 *This is how we applied DINEOF....*

22 *Now we show the results of DINEOF.*

23 *2.3. Clustering*

24 *To do 1D models, we cluster the Red Sea using clustering algorithms. We chose*
25 *GMM because....*

26 *This is more or less the way GMM works....*

27 *This is how we used it....*

28 *This is what we got....*

29 **3. Model and Assimilation**

30 *3.1. 1D-ERSEM model*

31 We use a 1D coupled ERSEM model. The physical forcing comes from a
32 3D circulation simulation of the Red Sea [Yao 2014]. The ecological models
33 are initialized with the results of the 3D Red Sea ecology simulation [Triantfyl-
34 lou2013].

The 1D regional ecological models used for this thesis have been configured and are operational. Three models will be used: for the northern, central and southern Red Sea. The extreme south of the Red Sea is not modeled, as its dynamics is poorly understood and we miss in situ data. The ecology is modeled with ERSEM, and the hydrodynamics is modeled with the MITgcm.

The results of the MITgcm are those from Yao *et al.* (2014a,b), in which a simulation of the Red Sea and part of the Gulf of Aden circulation was run over 50 years. The NCEP data were used for atmospheric forcing, and the ocean ECCO data for the open boundary conditions in the Gulf of Aden. The output of the 50 years run are used for the temperature and vertical circulation at the modeled points.

ERSEM simulates the complete water column with the pelagic and benthic ecosystems, as well as their coupling. The equations model the flow of carbon, nitrogen, phosphorus and silicon in the ecosystem. Living organisms are modeled in terms of population processes (growth and mortality) and physiological processes (ingestion, respiration, excretion, and egestion). The biota is divided into functional groups according to their trophic levels: producers (phytoplankton), consumers (zooplankton) and decomposers (bacteria), and further subdivided according to their sizes (Baretta *et al.*, 1995).

The ecological models are initialized with the results of a 3D ecological simulation of the Red Sea (Triantafyllou *et al.*, 2014). The nutrient concentrations are initialized using values from the World Ocean Atlas 2005 (WOA 2005).

3.2. Data Assimilation

To improve the results of the simulation we use the hybrid-SEIK assimilation scheme, detailed in this subsection.

The assimilation scheme for the ecological models has been implemented and is operational. The chosen scheme is the hybrid-SEIK, described in *Hamill and Snyder (2000)*. It can be seen as a variant of the 3DVAR variational assimilation scheme. 3DVAR assumes that the error forecast covariance is fixed in time. In the case of the hybrid, the covariance is a linear combination of the 3DVAR covariance and the time-evolving SEIK covariance matrix. Figure ?? shows the assimilation scheme improves the fit of the model to the chlorophyll data.

The problem of optimal filtering can be solved exactly by the Kalman Filter for linear systems. For nonlinear models, one can use the Extended Kalman (EK) filter, in which the model is linearized by computing the error covariance function. However, when the state is large, as is often the case for oceanographic applications, the EK is intractable. In that case, SEEK can be used, where the error covariance function is projected into a smaller subspace. This subspace evolves to ensure that most of the error is represented and filtered out. SEIK can be viewed as an ensemble variant of the SEEK, where the error covariance function is represented exactly by an ensemble of states. This avoids the computation of model gradients, and allows the assimilation scheme to perform better when this model is strongly non-linear. SEIK has been shown to be efficient for large-scale 3D ecosystem simulations (*Triantafyllou et al., 2003*).

The Expectation-Maximization scheme to estimate the filter parameters has also been derived. It is similar to that proposed by *Tandeo et al. (2014)*, except that the model is non linear. The scheme will be used to improve the estimates of the observation and model covariance errors.

92 4. Results

93 4.1. Model evaluation

94 Here, we compare the results of the free-run with the assimilated-run. We
95 show that we have a good prediction skill, and that the assimilation improves
96 the model.

97 4.2. Analysis

98 Here we look at the results and interpret them biologically. Do we find
99 comparable results as Acker, Raitsos, Weiker, etc. What can we say about the
100 hypothesis that they made about the process that drive primary productivity in
101 the Red Sea.

102 5. Conclusion

103 Are several 1D parallel 1D models a good alternative to 3D simulations?
104 What did we learn about the Red Sea ecology?
105 Future works?

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109 6. Bibliography

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