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Version Date: Sun Jun 26 16:40:23 2016 -0500

3 Abstract

Bio-geo-chemical models are useful tools in environmental sciences that can guide management and policy-making. Consequently, significant time and resources are spent developing these models in system-specific contexts. The optimization of model parameters to maximize precision, including transferability of these models to different systems, are fundamental concerns in the development and application of these tools. This study describes quantitative limitations of coupled hydrodynamic-ecological modelling by contrasting numeric and ecological certainty with a systematic framework for characterizing parameter sensitivity and identifability. We evaluate a simple bio-geo-chemical model that is the one-dimensional (1-D) unit of a larger spatio-temporal model of hypoxia on the Louisiana continental shelf of Gulf of Mexico as an example. Results from analysis of the 1-D model are used to infer larger trends in dissolved oxygen dynamics over time, having implications for understanding factors that contribute to environmental conditions that are detrimental to aquatic resources. In particular, we focus on issues of parameter identifiability using local sensitivity analyses to provide quantitative descriptions of numerical constraints on model precision. We argue that quantitative and ecological certainty in model calibration are often at odds and the practitioner must explicitly choose model components to optimize given tradeoffs between the two. We further conclude that numerically optimal parameter sets for models of hypoxia are often small subsets of the complete parameter set because of redundancies in the unique effects of paramater perturbations on model output. As a result, we demonstrate that use of a model for inference into ecological mechanisms of observed or predicted changes in hypoxic condition can be potentially misguided in the absence of quantitative descriptions of identifiability. Although these concerns have been expressed in the literature, they are rarely explicitly addressed or included in model evaluations. In addition to immediate implications for regional models, we provide a framework for describing the effects of parameter uncertainty and identifiability that can be applied to similar models to better inform environmental management.

1 Introduction

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- 1. Simulation/biogeochemical/process-based models overview, contrast with statistical models
- 2. What models seek to provide generality, precision, realism Levins (1966), there is a

tradeoff so models are 1) developed in partial independence and dependence on the world
and theory, 2) function autonomously from both, or 3) represent both at the same time, from
Morrison and Morgan (1999), cited in Ganju et al. (2016). This is similar to the
bias-variance tradeoff for statistical models, e.g., overparameterization of a model makes it
very biased as it fits the data (the world) exactly, tradeoff between sensitity and error with
changes in model complexity (more complexity is less error but increasing sensitivity)
described in Snowling and Kramer (2001)

- 3. How is model performance/uncertainty evaluated regarding what they should provide structural, observational, parameter Beck (1987)?
- 4. Parameter uncertainy as low-hanging fruit can do post-hoc and from inner to outer level of
 complexity, parameter uncertainty is the most common, e.g., marine ecological model
 Mateus and Franz (2015), but stopped short, global sensitivity analysis of eutrophication
 model Estrada and Diaz (2010)
- 5. Challenges related to uncertainty similar to degrees of freedom, identifiability definition 46 from Brun et al. (2001) and need to evaluate identifiability Fasham et al. (2006), Omlin 47 et al. (2001) did a similar analysis with freshwater biogeochem model. Identifiability 48 describes the ability to estimate a parameter in relation to variation among the remaining 49 parameters. A parameter is identifiable if all parameters within the set can be uniquely 50 estimated based on the observed data. Parameters that are unidentifiable typically produce 51 similar model outputs for a given relative perturbation, i.e., the effect of altering one 52 parameter can be undone by altering one or more other parameters. Model calibration will 53 not converge for parameters sets that are unidentifiable.

This study describes a parameter sensitivity analysis to evaluate identifiability for a bio-geo-chemical model of hypoxia for the northern Gulf of Mexico (GOM). We evaluate a simple one-dimensional (1-D) unit of a larger spatial-temporal model to explore relationships between multiple parameter sets and hypoxia dynamics on the Louisiana continental shelf (LCS). The study also provides a general framework for sensitivity analysis and parameter identifiability that can be used on similar mechanistic models. Specifically, an assumption is that models are

generally over-parameterized and only a finite and smaller subset of the larger parameter set can
be optimized for a given research question or dataset. We provide explicit guidance for choosing
such subsets of the parameter space given constraints on identifiability as directly related to
sensitivity analyses. The specific objectives are to 1) identify the parameters that have the
greatest influence on dissolved oxygen (O₂) using local sensitivity analysis, 2) quantify the
identifiability of subsets of the total parameter space based on sensitivity, 3) provide a set of
heuristics for choosing parameters based on sensitivity and parameter categories with the larger
mechanistic model, including extension to other state variables, and 4) discuss implications for
hypoxia formation in coastal regions, including management strategies for nutrient reduction and
use of mechanistic models to inform decision-making. The 'optimum' parameter space is defined
as the chosen subset that represents the maximum number of identifiable parameters. Here,
'optimum' is both a qualitative description based on a research question or management goal and
a quantitative objective based on numerical optimization criteria for fitting model output to a
calibration dataset. These results can be used to refine existing models or guide application of
models to novel contexts, such as downscaling or application to new environments.

6 2 Methods

7 2.1 Model description

Low concentrations of dissolved oxygen occur seasonally on the LCS in the northern
GOM. These hypoxic events, defined as <2 mg L⁻¹ (< 64 mmol m⁻³ of O₂), occur in bottom
waters and are caused by nutrient inputs from the Mississippi-Atchafalaya River Basin (MARB)
that drains a significant portion of the continental United States. Nutrient-stimulated primary
production in surface waters increases biological oxygen demand in bottom waters as sinking
organic matter is decomposed. The hypoxic area averages 15,540 km² annually (1993-2015) with
minimum concentrations observed from late spring to early fall. Seasonal variation is strongly
related to carbon and nutrient export from the MARB (Lohrenz et al. 2008, Bianchi et al. 2010),
whereas factors related to hydrologic variation and wind patterns can affect vertical salinity
gradients that contribute to the formation of hypoxia (Wiseman et al. 1997).

Three-dimensional numerical simulation models have been developed to describe factors
contributing to hypoxia and to predict the effects of management actions or climate scenarios on

future patterns (Fennel et al. 2013, Pauer et al. 2016, Lehrter et al. in review). This study evaluates a recently developed hydrodynamic and ecological model that describes horizontal and vertical transport and mixing of state variables relevant for hypoxia. The Coastal General Ecosystem Model (CGEM) includes elements from the Navy Coastal Ocean Model (Martin 2000) to describe hydrodynamics on the LCS and a biogeochemical model with multiple plankton groups, water-column metabolism, and sediment diagenesis (Eldridge and Roelke 2010). The hydrodynamic component of CGEM provides a spatially-explicit description of hypoxia dynamics using an orthogonal grid with an approximate horizontal resolution of 1.9 km² and twenty equally-spaced vertical sigma layers on the shelf (depth < 100 m, with additional hybrid layers at deeper depths). The biogeochemical component includes equations for 36 state variables including six phytoplankton groups (with nitrogen and phosophorus quotas for each), two zooplankton groups, nitrate, ammonium, phosphate, dissolved inorganic carbon, oxygen, silica, 101 and multiple variables for dissolved and particulate organic matter from different sources. The 102 model can be run for a set period of time at a given time step using atmospheric and hydrological 103 boundary conditions described in Hodur (1997) and Lehrter et al. (2013).

The core unit of CGEM is a 1-D model called FishTank that implements the 105 biogeochemical equations in Eldridge and Roelke (2010). This model acts as a standalone 106 operational unit from the 3-D model that does not include any form of transport (i.e., advection, 107 mixing, or surface flux) such that it can easily be applied to other hydrodynamic grids. 108 Accordingly, the sensitivity and identifiability analysis described below are informative for both the LCS gridded model as well as potential applications to different systems. The FishTank model 110 provides estimates for the 36 state variables described above using a 1-D parcel that is uniformly 111 mixed. A set of initial conditions is provided as input to the model that was based on observations 112 of relevant variables obtained from research cruises in April, June, and September 2006 (Table 1, 113 Murrell et al. (2014)). The FishTank model was executed from January 1st to December st, 2006 at a timestep of five minutes. 115

Results from FishTank are based on time-dependent differential equations that describe energy flow between phytoplankton (up to six groups) and zooplankton (two groups) as affected by nutrient uptake rates, organic matter inputs and losses, inherent optical properties, sediment diagenesis, and temperature (Penta et al. 2008, Eldridge and Roelke 2010, see appendix in Lehrter

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et al. in review). A total of 108 equations are estimated at each time step to return a value for each of the 36 state variables described by the model. In addition to the initial conditions, a set of 121 parameter values for each of the equations is also supplied at model execution. These parameters 122 define relationships among fixed effects in the equations and represent ecological properties 123 described by the model that influence hypoxia formation. Values for each of the parameters were 124 based on estimates from the literature, field or laboratory-based measurements, or expert-based knowledge in absence of the former. The sensitivity of O_2 was estimated at each timestep from 126 FishTank was evaluated in relation to 249 parameters that are included in the 108 equations. For 127 simplicity in the discussion below, the parameters were grouped into one of six categories in 128 relation to applicable equation: optics, phytoplankton, zooplankton, temperature, and organic matter. A full description of the equations and parameters is available as an appendix in (Lehrter 130 et al. in review). 131

2.2 Local sensitivity analysis

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The analysis focused on sensitivity of O_2 in the 1-D model to identify parameters that may affect spatial and temporal variation of hypoxia in the larger model. A local sensitivity analysis was performed for each of the 249 parameters using a simple perturbation approach to evaluate the change in dissolved oyxgten from the original parameter values. The analyses relied exlusively on concepts used in the FME package developed for the R statistical programming language (Soetaert and Petzoldt 2010). Each parameter was perturbed by 50% of its original value and the model was executed to obtain an estimate of the effect on O_2 . For each perturbation, a sensitivity value S was estimated for each time step i given a set value for parameter j as:

$$S_{ij} = \frac{\partial y_i}{\partial \Theta_j} \cdot \frac{w_{\Theta_j}}{w_{u_i}} \tag{1}$$

where the estimate depended on the change in the predicted value for response variable y divided by the change in the parameter Θ_j multiplied by the quotient of scaling factors w for each. The scaling factors, w_{Θ_j} for the parameter Θ_j and w_{y_i} for response variable y, were set as the default value of the unperturbed parameter and the predicted value of y_i after perturbation (Soetaert and Petzoldt 2010). The scaling ensures the estimates are unitless such that the relative magnitudes provide a comparison for model sensitivity to parameter changes that may vary in scale. Individual estimates for S_{ij} were as L1 and L2 across the time series to obtained a single sensitivity value of O_2 in response to a change in parameter j:

$$L1 = \sum |S_{ij}|/n \tag{2}$$

$$L2 = \sqrt{\sum \left(S_{ij}^2\right)/n} \tag{3}$$

The mean, minimum, and maximum S_{ij} for each parameter were also evaluated. In 149 general, positive mean sensitivity estimates suggested a parameter had a positive effect on O₂ for 150 a given increase in the parameter, whereas the converse was true for negative sensitivity estimates. 151 However, the effect of a parameter change may not be uniform over time such that S_{ij} can change 152 in magnitude and sign depending on the time step i. Time series of O_2 estimates before and after 153 perturbation were also evaluated to identify patterns not captured the summary statistics. All 154 parameters that are considered 'sensitive' should be further evaluated by plotting the predictions before and after perturbation and across the total range of the parameter. All parameters for each 156 of the six equation categories (optics, phytoplankton, zooplankton, temperature, and organic 157 matter) that had non-zero L1 or L2 were retained for identifiability analysis. 158

2.3 Identifiability

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Identifiability of parameter subsets was estimated from the minimum eigenvector of the cross-product of a selected sensitivity matrix (Brun et al. 2001, Omlin et al. 2001):

$$\gamma = \frac{1}{\sqrt{\min\left(\text{EV}[\hat{S}^{\dagger}\hat{S}]\right)}} \tag{4}$$

where γ ranges from one to infinity for perfectly identifiable (orthogonal) or unidentifiable (perfectly collinear) results for a set of parameters in the sensitivity matrix S. Identifiability can be estimated for any combination of parameters for a model. Values less than 10-15 indicate parameters are generally identifiable. The FME package provides the 'collin' function to estimate identifiability of output from the 'sensFun'. For reasons described above, a custom function was created to determine identifiability from model residuals, where the residauls were based on a relative perturbation of the model parameters. A comparison of results of identifiability using the

custom functions and those from the FME package were generally in agreement. Standard output from the identifiability function is shown below, where identifiability is displayed as a matrix with increasing combinations of subsets of the parameters.

The identifiability functions evaluate the ability to identify all subsets from pairwise to all parameters in the subset. As an example, the identifiability of pairwise combinations for each of the categories are shown below.

Pairwise identifiability of the top two most sensitive parameters in each category is shown, first as a table showing all unique combinations from two to all parameters and second as a figure showing pairwise combinations.

2.4 Identifying subsets and testing calibration

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Model calibration must consider the competing objectives of parameter inclusion and 179 identifability. Estimating parameters for novel datasets increases model complexity and the 180 ability to identify parameters decreases with the inclusion of parameters to estimate. The above analyses indicated that estimating all of the parameters was impossible because of redundancies in the the model output. Calibration to novel datasets should balance the competing objectives of 183 identifiability while including parameters for which model output is most sensitive. The plots 184 below show different scenarios of parameter inclusion for model calibration. Reading each plot 185 from left to right can be interpreted as including additional parameters, where each parameter is ranked by relative sensitivity. The inset in each plot shows the identifiability of including 187 parameters, up to a maximum where additional inclusion exceeds the identifability threshold of 188 fifteen. The scenarios for including parameters all begin with the parameters that have the greatest 189 effect on the model output. The inclusion of additional parameters depends on the scenario. The first scenario selected parameters by decreasing sensitivity within each category (i.e., four 191 separate models calibrated for optics, organics, phytoplankton, or zooplankton), the second 192 scenario selected parameters by sensitivity regardless of category, and the third scenario selects 193 parameters by sensitivity with equal representation between categories. Boxes in each plot 194 represent a unique parameter and the numbers in each box represent the relative rank of a 195 parameter's sensitivity to the model output within each category.

The following plots show identifiability with the addition of parameters.

3 Results

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3.1 Local sensitivity analysis

The parameters that have the greatest effect on the model by category (optics, organics, phytoplankton, zooplankton) are as follows.

Plotting the raw values from the sensitivity analysis provides a visual assessment of changes.

4 Discussion

Questions specific to GOM - what initial conditions are important? How many phytoplankton groups do we need (e.g., related to structural uncertainty)?

How does the assimilation of additional parameters (e.g., other state variables) during calibration influence the conclusions?

How does uncertainty translate to what a model should provide (generality v precision)? The first step - find out what can be optimized but then do not overfit....

What about structural uncertainty - does sensitivity of a model to variation in a parameter imply parameter uncertainty and/or structural uncertainty?

A final point about optimization with identifiable parameter sets - optimization to fit the data still does not ensure a correct model. Failing in one way can be over-compensated by another feature, e.g., the parameter set that is optimized (see Flynn (2005), p. 1207, third paragraph)

Omlin et al. (2001) state that the sensitivity, identifiability, estimation process is iterative (p. 113), need to rinse and repeat for proper calibration.

How to improve identifiability - get more/better observed data, include obs from other state variables in RSS minimization (eqn q in Omlin et al. (2001))

Alternative methods for uncertainty analysis - bayesian, MCMC, nonlinear calibration-constrained optimization (Gallagher and Doherty 2007)

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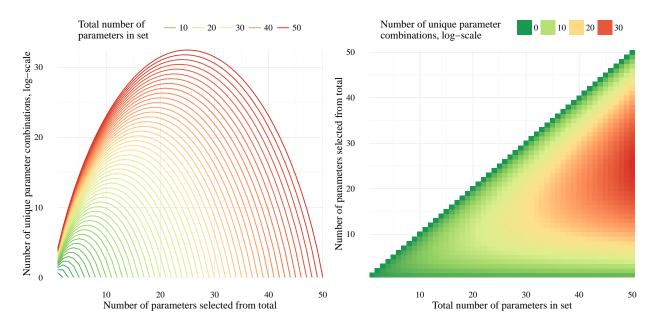


Fig. 1: Number of unique combinations that can occur by selecting parameters for optimization from different parameter sets. The number of combinations are shown for increasing numbers of selected parameters from the total in the set, where 100 parameter sets are shown each with one through 100 total parameters. For example, there are six unique parameter combinations that result from choosing two parameters from a set. Note that the number of unique combinations is shown as the natural-log. The same information is viewed differently in each plot.