



1 **Assessing the value of BGC Argo profiles versus ocean colour observations for** 2 **biogeochemical model optimization in the Gulf of Mexico**

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8 **Abstract.** Biogeochemical ocean models are useful tools subject to uncertainties arising from
9 simplifications, inaccurate parameterization of processes, and poorly known model parameters. Parameter
10 optimization is a standard method for addressing the latter but typically cannot constrain all
11 biogeochemical parameters because of insufficient observations. Here we assess the trade-offs between
12 satellite observations of ocean colour and biogeochemical (BGC) Argo profiles, and the benefits of
13 combining both observation types, for optimizing biogeochemical parameters in a model of the Gulf of
14 Mexico. A suite of optimization experiments is carried out using different combinations of satellite
15 chlorophyll and profile measurements of chlorophyll, phytoplankton biomass, and particulate organic
16 carbon (POC) from autonomous floats. As parameter optimization in 3D models is computationally
17 expensive, we optimize the parameters in a 1D model version, and then perform 3D simulations using
18 these parameters. We show first that the use of parameters optimized in 1D improves the skill of the 3D
19 model. Parameters that are only optimized with respect to surface chlorophyll cannot reproduce
20 subsurface distributions of biological fields. Adding profiles of chlorophyll in the parameter optimization
21 yields significant improvements for surface and subsurface chlorophyll but does not accurately capture
22 subsurface phytoplankton and POC distributions because the parameter for the maximum ratio of
23 chlorophyll to phytoplankton carbon is not well constrained in that case. Using all available observations
24 leads to significant improvements of both observed (chlorophyll, phytoplankton, and POC) and
25 unobserved variables (e.g. primary production). Our results highlight the significant benefits of BGC
26 Argo measurements for biogeochemical parameter optimization and model calibration.



27 **1. Introduction**

28 Oceanic primary production forms the basis of the marine food web and fuels the biological pump,
29 which contributes to the sequestration of atmospheric CO₂ in the ocean's interior thus mitigating global
30 warming. An accurate quantification of primary production and biological carbon export is therefore
31 important for our understanding of the marine carbon cycle and for predicting how carbon cycling and
32 marine ecosystems will interact with climate change.

33 Direct observations of primary production and export flux are relatively sparse because of the cost
34 and effort involved in measuring these fluxes. Numerical models can complement sparse observations.
35 Well validated and calibrated models are useful tools for hindcasting and nowcasting past and present
36 biogeochemical fluxes and are the most common tool for projecting future changes.

37 In recent years, many biogeochemical models with different complexities, ranging from three to more
38 than thirty biological state variables, have been developed to study ocean biogeochemical processes.
39 Regardless of their complexities, the performance of these models is highly dependent on the appropriate
40 choice of model parameter values (e.g. maximum growth, grazing and mortality rates), most of which are
41 poorly known. Some parameter choices are informed by laboratory experiments (e.g. light and nutrient
42 dependence of phytoplankton growth), although isolated cultures in the lab may not be representative of
43 the behavior of natural communities. Other parameters cannot be directly determined (e.g. mortality
44 rates). Choosing appropriate parameter values becomes more challenging as model complexity grows
45 because the number of parameters increases exponentially with the number of state variables (Denman,
46 2003).

47 A standard method for addressing the problem of poorly known model parameters is parameter
48 optimization, a process by which the misfit between model results and available observations is
49 minimized by iteratively varying parameters (Matear, 1995; Prunet et al., 1996a, 1996b; Fennel et al.,
50 2001; Friedrichs et al., 2007; Kuhn et al., 2015, 2018). However, even formal optimization typically
51 cannot constrain all biogeochemical parameters (i.e. provide optimal parameter estimates with relatively
52 small uncertainties) because of insufficient information in the available observations (Matear, 1995;
53 Fennel et al., 2001; Ward et al., 2010; Bagniewski et al., 2011). For example, Matear (1995) used a so-



54 called simulated annealing algorithm to optimize three different ecosystem models and found that even
55 for the simplest nutrient-phytoplankton-zooplankton model, not all independent parameters could be
56 constrained well, leaving the others with large uncertainty ranges. A more recent study reported that the
57 lack of zooplankton observations led to poor accuracy of the optimized zooplankton-related parameters
58 when using a suite of Lagrangian-based observations during the North Atlantic spring bloom (Bagniewski
59 et al., 2011). A broader suite of observation types should be favourable to parameter optimization although
60 complications can arise. For example, when optimizing a suite of 1D models for the Mid-Atlantic Bight,
61 the use of satellite POC observations in addition to satellite chlorophyll did not yield further
62 improvements in model-data fit but degraded the representation of chlorophyll (Xiao and Friedrichs,
63 2014a).

64 Typically surface ocean chlorophyll from satellite is the main source of observations for model
65 validation (e.g. Doney et al., 2009; Gomez et al., 2018; Lehmann et al., 2009) and parameter optimization
66 (Prunet et al., 1996a; Xiao and Friedrichs, 2014a, 2014b), supplemented by other observation types as
67 available. However, satellites only see the ocean surface and do not resolve the vertical distribution of
68 chlorophyll. This is especially problematic in oligotrophic regions where the maximum chlorophyll
69 concentration (referred as the deep chlorophyll maximum, DCM) is pronounced near the base of the
70 euphotic zone because of photoacclimation (Cullen, 2015; Fennel and Boss, 2003). In addition, although
71 chlorophyll has long been used as a proxy of phytoplankton biomass and to estimate primary production
72 based on some assumptions (Behrenfeld and Falkowski, 1997), it is not a direct measure of carbon-based
73 phytoplankton biomass. The ratio of chlorophyll-to-phytoplankton carbon varies by at least an order of
74 magnitude due to physiological responses of phytoplankton to their ambient environment (e.g. nutrients,
75 light, and temperature) (Cullen, 2015; Fennel and Boss, 2003; Geider, 1987). Thus, changes in
76 chlorophyll may result from physiologically induced modifications of the chlorophyll-to-phytoplankton
77 ratio rather than actual changes of phytoplankton biomass (Fommervault et al., 2017; Mignot et al., 2014).
78 Satellite surface chlorophyll alone is therefore likely insufficient for model validation and for constraining
79 biogeochemical models via parameter optimization.

80 Recent advances in autonomous platforms and sensors have opened opportunities for simultaneous



81 measurement of several biological and chemical properties throughout the upper ocean with high
82 resolution, over broad spatial scales and for sustained periods (Roemmich et al., 2019). In particular, the
83 biogeochemical (BGC) Argo program (Johnson and Claustre, 2016; Roemmich et al., 2019) will provide
84 temporally evolving 3D information on biogeochemical variability at previously unobserved scales. Here
85 we assess to what degree observations of chlorophyll fluorescence and particle backscatter from Argo
86 profiles improve the prospects of optimizing a biogeochemical model for the Gulf of Mexico.

87 Since the high computational cost and storage demands of 3D models make direct application of
88 most parameter optimization techniques difficult (but see Mattern et al., 2012; Mattern and Edwards,
89 2017; Tjiputra et al., 2007 for exceptions), they are typically applied in computationally efficient 1D
90 models before using the resulting parameters in 3D version (e.g. Hoshiba et al., 2018; Kane et al., 2011;
91 Kuhn and Fennel, 2019; Schartau and Oschlies, 2003). We follow the latter approach here.

92 The main objective of this study is to assess the added value of bio-optical profile information from
93 Argo floats for biogeochemical model optimization in the Gulf of Mexico. We first examine the feasibility
94 of improving the 3D model by applying the optimal parameters from 1D model optimizations. We find
95 that the gains from the 1D optimizations transfer to the 3D version. Then, by using different combinations
96 of satellite and float observations we show that parameters optimized with respect to satellite data cannot
97 reproduce subsurface distributions unless the float observations (i.e. chlorophyll, phytoplankton, and POC)
98 are also used.

99 **2. Study region**

00 The Gulf of Mexico (GOM) is a semi-enclosed marginal sea (Figure 1) which is characterized by
01 eutrophic coastal waters on the northern shelf and an oligotrophic deep ocean. The high productivity in
02 the northern coastal region is fueled by large nutrient and freshwater inputs from the Mississippi and
03 Atchafalaya Rivers. The large nutrient load and strong stratification driven by Mississippi and
04 Atchafalaya River inputs lead to summer hypoxia and ocean acidification in bottom waters on the
05 northern shelf (Laurent et al., 2017; Yu et al., 2015), but nutrient export across the shelf break into the
06 open Gulf is minor (Xue et al., 2013).



07 The deep ocean of the GOM is oligotrophic. Previous satellite-based studies have revealed a clear
08 seasonal cycle in surface chlorophyll with highest concentrations in winter and lowest in summer
09 (Martínez-López and Zavala-Hidalgo, 2009; Muller-Karger et al., 1991, 2015). Thanks to advances in
10 autonomous profiling technology, recent studies based on simultaneous measurements of subsurface
11 chlorophyll and backscatter have demonstrated that the seasonal variability of surface chlorophyll might
12 be a result of the vertical redistribution of subsurface chlorophyll and/or physiological response to solar
13 radiation of phytoplankton (Fommervault et al., 2017; Green et al., 2014).

14 3. Methods

15 3.1. Biological observations

16 Satellite-derived chlorophyll from the Ocean-Colour Climate Change Initiative project (OC-CCI,
17 <https://www.oceancolour.org>) with a spatial resolution of 4 km from 2010 to 2015 is used for model
18 validation and parameter optimization. These data were provided by the European Space Agency (ESA),
19 which produced a set of validated and error-characterised global ocean-color products by merging
20 SeaWiFS (Sea-viewing Wide Field-of-view Sensor), MODIS (Moderate-resolution Imaging
21 Spectroradiometer), and MERIS (medium-spectral resolution imaging spectrometer) products.

22 In addition to the satellite-based measurements, bio-optical measurements from six autonomous
23 profiling floats are used (Figure 1), which were deployed by the Bureau of Ocean Energy Management
24 (BOEM) and operated in the deep GOM from 2011 to 2015. These floats were equipped with a CTD and
25 bio-optical sensors to collect biweekly profiles of temperature, salinity, chlorophyll fluorescence, and
26 backscatter at 700 nm ($bbp700$ (m^{-1})) from the surface to 1000 m depth (see Fommervault et al., 2017
27 and Green et al., 2014 for more details). Chlorophyll was derived from fluorescence based on the sensor
28 manufacturer's calibrations and cross-validated with the satellite estimates of surface chlorophyll. While
29 the surface chlorophyll measurements from the floats and the satellite estimates both show a typical
30 seasonal cycle and are highly correlated ($R^2=0.74$; see Figures S1 and S2a in the Supplement), the satellite
31 underestimates the float-measured chlorophyll concentrations in winter (Figure S1c). Satellite estimates
32 were therefore corrected following the regression equation shown in Figure S2a (Figure S1c).

33 The backscatter sensor carried by the floats provides the volume scattering function at a centroid



34 angle of 140° and a wave length of 700 nm ($\beta(140^\circ, 700\text{nm}) \text{ m}^{-1} \text{ sr}^{-1}$) (Green et al., 2014). The profiles
35 were filtered following Briggs et al. (2011) to remove spikes. To cross-validate the float-measured *bbp700*
36 with the satellite estimates, the *bbp670* from OC-CCI was firstly converted to *bbp700* through a power
37 law (Boss and Haëntjens, 2016):

38

$$39 \quad bbp(\lambda_1) = \left(\frac{\lambda_1}{\lambda_2}\right)^{-\gamma} bbp(\lambda_2), \quad (1)$$

40

41 where λ_1 and λ_2 represent the measured wavelength, and γ was estimated as 0.78 based on the global
42 measurements. Compared to surface chlorophyll, surface *bbp700* has a less distinct seasonal cycle (Figure
43 S3). For example, the coefficient of variation, defined as the ratio between standard deviation and mean
44 to show the extend of variability, is much lower for *bbp700* (0.09 and 0.07 for floats and satellite,
45 respectively) than for chlorophyll (0.31 and 0.26 for floats and satellite, respectively). The *bbp700* from
46 the floats is weakly correlated with the satellite estimates ($R^2=0.11$) and generally lower by a factor of
47 ~ 0.45 than the satellite estimates (Figure S2b). The *bbp700* profiles were therefore multiplied by 2.2
48 before being converted to *bbp470* following the equ. 1.

49 Profiles of phytoplankton and POC were derived from the validated *bbp470* profiles based on the
50 following empirical relationships

51

$$52 \quad Phy = 30100 \times (bbp470 - 76 \times 10^{-5}) \frac{1}{12 \times 6.625}, \quad (2)$$

53

$$54 \quad \log_{10}(POC) = 1.22 \times \log_{10}(bbp470) + 5.15. \quad (3)$$

55

56 The relationships for phytoplankton (Martinez-Vicente et al., 2013; equ. 2) and POC (Rasse et al., 2017;
57 equ. 3) were obtained from a data set for the Atlantic Ocean that covers a wide range of oceanographic
58 regimes from eutrophic to oligotrophic ecosystems. The scale factors of 12 and 6.625 in equ. 2 represent



59 the molecular weight of carbon and the Redfield ratio to convert phytoplankton concentrations from mg
60 C m⁻³ to mmol N m⁻³. According to Behrenfeld et al. (2005), the intercept 76×10^{-5} in equ. 2 represents
61 the background backscatter of nonalgal detritus. In this study, both chlorophyll and phytoplankton
62 approach zero when *bbp470* is $76 \times 10^{-5} \text{ m}^{-1}$, implying that the use of equ. 2 is appropriate.

63 **3.2. 3D model description**

64 The physical model is configured based on Regional Ocean Modeling System (Haidvogel et al.,
65 2008; ROMS, <https://www.myroms.org>) for the Gulf of Mexico (Figure 1). The model has a horizontal
66 resolution of 6~7 km and 36 terrain-following sigma layers with refined resolution near the surface and
67 bottom. The model solves for the horizontal and vertical advection of tracers using the Multidimensional
68 positive definitive advection transport algorithm (MPDATA, Smolarkiewicz and Margolin 1998).
69 Horizontal viscosity and diffusivity are parameterized by a Smagorinsky-type formula (Smagorinsky,
70 1963), and vertical turbulent mixing is calculated by the Mellor-Yamada 2.5-level closure scheme (Mellor
71 and Yamada, 1982). Bottom friction is specified by a logarithmic drag formulation with a bottom
72 roughness of 0.02 m. The model is forced by 3-hourly surface heat and freshwater fluxes, 6-hourly air
73 temperature, sea level pressure and relative humidity, and 10-m winds from the European Centre for
74 Medium-Range Weather Forecast ERA-Interim product with a horizontal resolution of 0.125° (ECMWF
75 reanalysis, <https://www.ecmwf.int/en/forecasts/datasets/reanalysis-datasets/era-interim>). A bulk
76 parameterization is applied to calculate the surface net heat fluxes and wind stress. The model is one-way
77 nested inside the 1/12° data-assimilative global HYCOM/NCODA (<https://www.hycom.org>). Tidal
78 constituents are neglected in the model.

79 The biogeochemical model uses a 7-component model (Fennel et al., 2006) to simulate the nitrogen
80 cycle in the water column. The model describes the dynamics of two species of dissolved inorganic
81 nitrogen (nitrate, NO₃, and ammonium, NH₄), one function of phytoplankton (Phy), chlorophyll (Chl) as
82 a separate state variable which allows photo-acclimation based on the model of (Geider et al., 1997), one
83 function of zooplankton (Zoo), and two pools of detritus (i.e. small suspended detritus, SDeN, and large
84 fast-sinking detritus, LDeN). Water-sediment interactions are simplified by an instantaneous
85 remineralization parameterization, where detritus sinking out of water column immediately results in a



86 corresponding influx of ammonium into the bottom layer. Detailed descriptions of the model equations
87 can be found in Fennel et al. (2006). The biological model parameters are listed in Table 1.

88 The model receives freshwater, nutrients (NO₃ and NH₄) and organic matter inputs from major rivers
89 along the Gulf coast. Freshwater and nutrients from the Mississippi and Atchafalaya rivers are prescribed
90 based on the daily measurements by the US Geological Survey river gauges. River particulate organic
91 nitrogen (PON) is assigned to the small detritus pool and determined as the difference between total
92 Kjeldahl nitrogen and ammonium (Fennel et al., 2011). Other rivers utilize the climatological estimates
93 of freshwater, nutrients, and PON as in Xue et al. (2013).

94 Initial and open boundary conditions for NO₃ are specified by applying an empirical relationship
95 between NO₃ and temperature, derived from the World Ocean Atlas (WOA; Figure S4a), that is applied
96 to the temperature fields from HYCOM/NCODA. Analogously, empirical relationships between
97 chlorophyll and density (Figure S4b), phytoplankton and density (Figure S4c), and POC and density
98 (Figure S4d) were obtained from the median profiles of the bio-optical floats and used to derive initial
99 and boundary conditions for these variables. Zooplankton and small detritus were assumed to amount to
00 10% of phytoplankton biomass and the remaining fractions of POC attributed to large detritus.

01 A 6-year (5 January 2010 – 31 December 2015) hindcast was performed that includes the period of
02 operation of the bio-optical floats. The first year is considered model spin-up and the next five years will
03 be discussed.

04 **3.3. 1D model description**

05 As optimizing a 3D biogeochemical model is computationally expensive, it is more practical to
06 perform the optimization using a reduced-order model surrogate. A surrogate can be a coarser resolution
07 model, a simplified model, or a reduced-dimension model. In this study, a 1D model is used to optimize
08 the biological parameters of the 3D model. This approach has been successfully used previously (Hoshiba
09 et al., 2018; Kane et al., 2011; Oschlies and Schartau, 2005).

10 The 1D model, which is similar to that used by Lagman et al. (2014) and Kuhn et al. (2015), covers
11 the upper 200 m of the ocean with a vertical resolution of 5 m and is configured at one location in the
12 central Gulf (see Figure 1). In the vertical direction, the water column is divided into two layers: the



13 turbulent surface layer and a quiescent layer below. A higher diffusion coefficient ($K_{Z1} =$
14 $\max(H_{MLD}^2/400, 10)$) is applied in the turbulent surface layer and a lower diffusion coefficient ($K_{Z2} =$
15 $K_{Z1}/2$) is assigned to the quiescent bottom layer. The interface between these two layers is determined
16 by the mixed layer depth (H_{MLD}), defined as the depth where the temperature is 5°C lower than at the
17 surface, and was obtained from a simulation of the 3D model. The model is integrated in time using the
18 Crank-Nicolson scheme for vertical turbulent mixing and an implicit time-stepping scheme for the
19 biogeochemical tracers, which are treated identically to the 3D model. Some of the biogeochemical
20 parameterizations require input of temperature and solar radiation, which are also taken from the 3D
21 model. As the 1D model does not consider horizontal and vertical advection, NO₃ below 100 m is nudged
22 to that from the 3D base simulation with a nudging time scale of 20 days. The model is run for the year
23 2010 repeatedly for three cycles, with the first two are model spin-up and the last annual cycle used to
24 calculate the misfit between model and observations.

25 3.4. Parameter optimization method

26 The evolutionary algorithm described by Kuhn et al. (2015, 2018) is used to search for optimal
27 model parameters by minimizing the misfit between model and observations. The misfit is measured by
28 the following cost function:

29

30

$$F(\vec{p}) = \sum_{v=1}^V F_v(\vec{p}), \quad (4)$$

31

32

$$F_v(\vec{p}) = \frac{1}{N_v \sigma_v^2} \sum_{i=1}^{N_v} (\hat{y}_{i,v} - y_{i,v}(\vec{p}))^2, \quad (5)$$

33

34 where \vec{p} represents the parameters vector, V is the number of different observation types, N_v is the
35 number of observations for each variable, $F_v(\vec{p})$ is the misfit for observation type v measured as the
36 mean-square difference between observations (\hat{y}) and corresponding model estimates ($y(\vec{p})$). The cost



37 function $F_v(\vec{p})$ is normalized by the standard deviation of each variable type (σ_v) in order to remove the
38 effect of different units.

39 The algorithm is inspired by the rules of natural selection. Following Kuhn et al. (2015), an initial
40 parameter population of 30 parameter vectors is randomly generated within a predefined range of
41 parameters (see Table 1). The model is evaluated for each parameter vector and the resulting cost function
42 is calculated. For this initial generation and each of the following generations, the half of the population
43 with the lower misfit survives into the next generation. The other half is regenerated through a
44 recombination of survivors in a process analogous to genetic crossover. In addition, each newly generated
45 population is subject to random mutations by multiplying the parameter values by a random value
46 between 0 and 2. Parameter values exceeding the predefined range are replaced by their corresponding
47 minimum or maximum limits to avoid unrealistic values. The above procedure is performed iteratively
48 for 300 generations to reach the minimum of the cost function, which corresponds to the optimal
49 parameter set.

50 Previous parameter optimization studies have shown that it is difficult to constrain all model
51 parameters even for very simple ecosystem models because the information content of available
52 observations is typically insufficient (Matear, 1995; Fennel et al., 2001; Ward et al., 2010). Here we
53 conducted sensitivity tests to identify the parameters that are most sensitive to the available observations
54 and chose a subset of these to be optimized. In the **base case**, all parameters were at their initial guess
55 values obtained from the previous literature and some initial tuning. Then the **test cases** were run multiple
56 times by incrementally changing one parameter at a time to be the minimum, the first, second and third
57 quartile, and the maximum of its corresponding range while setting the other parameters to their initial
58 guess value (Table 1). The sensitivity was measured as the sum of a normalized absolute difference
59 between the base case (y_{Base}) and the test case (y_{Test})

60

61

62

$$Q(y, \vec{p}) = \frac{1}{m} \sum_{i=1}^m \frac{1}{n} \sum_{j=1}^n \frac{|y_{Base} - y_{Test}|}{y_{Base}} \quad (6)$$



63 where m is the number of parameter increments (here 5) and n is the number of base-test pairs including
64 all 1D model grid cells throughout the whole simulation period for all variables to be compared.

65 Results of the sensitivity analysis are shown in Figure 2, where parameters are ranked by sensitivity
66 with respect to chlorophyll (Figure 2a) and the sum of chlorophyll, phytoplankton, and POC (Figure 2b).
67 POC is the sum of phytoplankton, zooplankton, and small and large detritus.

68 **3.5. Parameter optimization experiments**

69 For the parameter optimization of the 1D model, satellite chlorophyll within a 3×3 pixel ($12\text{ km}\times 12$
70 km area) around the 1D station and climatological monthly averages of the profiles from the bio-optical
71 floats were used.

72 To assess the effects of the optimization with respect to the different observation types, we conducted
73 three groups of experiments in which (A) surface satellite chlorophyll only, (B) surface satellite
74 chlorophyll and float profiles of chlorophyll, and (C) surface satellite chlorophyll and float profiles of
75 chlorophyll, phytoplankton, and POC were used. For each of these three groups, four to five optimizations
76 were conducted starting with the three most sensitive parameters and then adding one more parameter at
77 a time (Table 2) guided by the sensitivity analysis with respect to observed variables they used.
78 Specifically, groups A and B were based on the sensitivity analysis with respect to chlorophyll, while
79 group C was based on sensitivity analysis with respect to the sum of chlorophyll, phytoplankton, and
80 POC. Each optimization was replicated four times. The optimization with smallest model-data misfit
81 within each group was then used. Prior tests have shown that the available observations cannot
82 simultaneously constrain the sinking rates of small and large detritus (w_{SDet} and w_{LDet}). Therefore, a
83 constant ratio of 0.1 between these two parameters ($w_{SDet} = 0.1 \times w_{LDet}$) was imposed and only one of the
84 two was optimized. In groups A and B, the aggregation parameter τ was fixed at 0.05 because prior tests
85 generated unreasonably high values for this parameter.

86 We report two different metrics of misfits for these groups of experiments. The first metric, which
87 we refer to as the case-specific cost function value, is based on the optimized observations in a given
88 experiment and is minimized by the optimization algorithm, i.e.

89



90
$$F_A(\vec{p}) = F_{SurfCHL}(\vec{p}), \quad (8)$$

91
92
$$F_B(\vec{p}) = F_{SurfCHL}(\vec{p}) + F_{CHL}(\vec{p}), \text{ and} \quad (9)$$

93
94
$$F_C(\vec{p}) = F_{SurfCHL}(\vec{p}) + F_{CHL}(\vec{p}) + F_{Phy}(\vec{p}) + F_{POC}(\vec{p}). \quad (10)$$

95
96 However, the models with lower case-specific misfit does not necessarily have better predictive skill in
97 reproducing the unoptimized observations because of the so-called overfitting problem, e.g. the model
98 may be tuned to reproduce optimized observations through wrong mechanisms (Friedrichs et al., 2006).
99 To account for this, a second metric referred to as the total misfit is given by equ. 10. For group C, the
00 second metric is the same as the case-specific cost function. For groups A and B, the total misfit metric
01 allows us to assess improvements in the model's predictive skill to represent unoptimized fields.

02 **4. Optimization of 1D models**

03 **4.1. Observations and base case**

04 To provide context for the evaluation of our optimization experiments, the observations and the
05 base case will be described first. As shown in Figure 3a, the observed surface chlorophyll shows a clear
06 seasonality with the high concentrations in winter and low concentrations in summer. In the base case,
07 the simulated surface chlorophyll fits observations well. Unlike the surface chlorophyll, the vertically
08 integrated chlorophyll as well as the phytoplankton and POC over the upper 200 m tend to be more
09 constant with much less seasonality (Figure 3b-d). This has been reported by Fommervault et al. (2017)
10 who attributed the seasonality of surface chlorophyll to the vertical redistributions of subsurface
11 chlorophyll and/or photoacclimation, rather than real changes in biomass.

12 The DCM is a ubiquitous phenomenon in the oligotrophic regions (Cullen, 2015). Observations
13 detect a predominant DCM at around 60-100m depth throughout the whole year, with a sharp deepening
14 in June and gradual shoaling after July (Figure 3e), reflecting the seasonality of the solar radiation. Unlike
15 the large variability in the depth of the DCM, its magnitude is relatively constant at around 0.62 mg m^{-3}
16 (Figure 3f). In the annually averaged profiles, the observed DCM is located at about 80 m depth with a



17 concentration of 0.52 mg m^{-3} (Figure 4a). The base case succeeds in reproducing the DCM at $65 \pm 7 \text{ m}$
18 depth. However, it fails to reproduce the deepening of the DCM in June and the simulated annually
19 averaged depth of DCM is shallower by about 15 m than the observed. The simulated magnitude of the
20 DCM is about 2-fold larger than the observed (Figure 3f and Figure 4a) and hence the base case generally
21 overestimates vertically integrated chlorophyll (Figure 3b).

22 With respect to phytoplankton and POC, the observed maximum concentration occurs at about 60
23 m depth, which is 20 m above the DCM (Figure 4b-c). The observed vertical distributions of
24 phytoplankton and POC are not well captured by the base case. For example, phytoplankton and POC in
25 the upper layer are overestimated with the model-data discrepancies exceeding the variability of the
26 observations (Figure 4b-c). As a result, the base case yields an overall overestimation of the vertically
27 integrated phytoplankton and POC (Figure 3c-d).

28 Figure 4b also shows that both observed and simulated phytoplankton approach zero at about 160
29 m depth. Unlike phytoplankton, the observations show that the POC concentrations are 19 mg C m^{-3} at
30 about 200 m depth because of the existence of detritus (Figure 4c). However, the base case fails to
31 reproduce this non-zero POC concentrations, indicating that the model might be underestimating the
32 carbon export fluxes at 200 m.

33 **4.2. Results of the optimizations**

34 **4.2.1. Model-data misfits**

35 The case-specific cost function values and total misfits for the different 1D optimizations are shown
36 in Figure 5. Not surprisingly, all optimizations result in a reduction of the case-specific cost function
37 values. The extent of the reductions depends on the specific subset of parameters that were optimized.
38 However, the total misfits are not reduced in all optimizations. Optimizations A1 and A2 lead to slightly
39 larger total misfits than the base case and optimization B2 leads to a significantly larger total misfit. The
40 smallest total cost function values are achieved in A4, B4, and C4, i.e. in the experiments where a larger
41 subset of parameters was optimized (6 parameters). The optimal parameter sets (A4, B2, and C4), which
42 are selected based on case-specific misfit from these three groups, will be used in subsequent analyses



43 and hereafter are denoted simply as experiment A, experiment B, and experiment C. Further comparisons
44 are presented below to assess the impact of the different combinations of observations.

45 **4.2.2. Experiment A**

46 The optimal parameters from experiment A yield a 58% reduction in the misfit for surface
47 chlorophyll (Figure 5d). However, the vertical structure of chlorophyll deteriorates relative to the base
48 case (Figure 4a) because of inappropriate estimates of the initial slope ($\alpha=0.0101$; see table 2) and the
49 maximum ratio of chlorophyll to carbon ($\theta_{max}=0.0191$; see table 2). The annually averaged depth of the
50 DCM is lifted up to around 30 ± 10 m and the magnitude of DCM strongly decreases (Figure 3a, 4b).
51 Similar to chlorophyll, these deteriorations also manifest in the vertical phytoplankton and POC
52 distributions (Figure 4b-c). As a result, experiment A underestimates vertically integrated chlorophyll,
53 phytoplankton, and POC (Fig. 3b-d).

54 **4.2.3. Experiment B**

55 Due to the addition of observed chlorophyll profiles to the optimization in experiment B, the misfits
56 for surface and subsurface chlorophyll decrease relative to the base case (Figure 5d), but the reduction in
57 the misfit for surface chlorophyll (38%) is smaller than that in experiment A (58%). A straightforward
58 interpretation is that the addition of subsurface observations reduces the model's degrees of freedom to
59 fit one single observation type (surface chlorophyll). The vertical profile of chlorophyll is reproduced
60 better in experiment B than in the base case and experiment A in that the magnitude of the DCM is closer
61 to the observations, although the DCM depth is still too shallow, on average by about 20 m (Figure 4a).
62 The improvement in the vertical chlorophyll structure results in a better model-data fit of vertically
63 integrated chlorophyll (Figure 3b).

64 Despite the improvements in chlorophyll, the vertical profiles of phytoplankton and POC exhibit a
65 marked deterioration relative to the base case and experiment A (Figure 4b-c) because the parameter
66 optimization underestimates the maximum chlorophyll-to-carbon ratio ($\theta_{max}=0.0158$; see table 2).
67 Experiment B leads to an overestimation of phytoplankton and POC relative to the base case with misfits
68 roughly 2.7 and 1.6 times larger than those of the base case, respectively (Figure 5d). Although
69 experiment B reproduces the non-zero POC concentrations at about 200 m depth, the proportion of



70 phytoplankton in the POC pool is incorrect. In contrast to the observations where detritus dominates POC,
71 the simulated POC at 200 m is dominated by phytoplankton (49%) followed by zooplankton (39%).

72 **4.2.4. Experiment C**

73 Incorporating all observations (i.e. surface chlorophyll and profiles of chlorophyll, phytoplankton,
74 and POC) in experiment C improves the model-data misfits for almost all aspects except for surface
75 chlorophyll (Figure 3). Although a slight increase in the misfit occurs for the surface chlorophyll (~5%),
76 the total misfit is reduced by 75% compared to the base case. As shown in Figure 4a, the annually
77 averaged depth of DCM of 80 m coincides with the observed DCM, primarily because experiment C
78 reproduces the deepening of the DCM in summer. The magnitude of the DCM is also decreased relative
79 to the base case but remains higher than the observed. Phytoplankton and POC profiles exhibit only minor
80 deviations from the observations (Figure 4b-c). Importantly, experiment C reproduces the non-zero POC
81 concentrations at 200 m. In contrast to experiment B, phytoplankton in experiment C drops to zero at
82 about 160 m and POC is dominated by detritus (85%), which is more consistent with the observations.

83 **4.3. Simulated carbon fluxes**

84 Annually averaged carbon fluxes within the upper 200 m are shown for each experiment in Figure
85 6. The primary production in the base case amounts to $0.78 \text{ g C m}^{-2} \text{ day}^{-1}$, of which 37% is consumed by
86 zooplankton, and the remaining 63% flows into detritus pools through sloppy feeding, mortality, and
87 aggregation of phytoplankton. As for the production of detritus, contributions from the phytoplankton
88 and zooplankton pools account for 70% and 30%, respectively. Most of the produced detritus is recycled
89 into the nutrient pool fueling recycled primary production, and only a small fraction is removed from the
90 upper layer through gravitational sinking. As a result, carbon export, which is estimated as the sum of
91 sinking fluxes by phytoplankton and detritus, is only $0.00032 \text{ g C m}^{-2} \text{ day}^{-1}$ and accounts for 0.04% of
92 primary production.

93 Due to the underestimation of phytoplankton in experiment A, primary production is reduced to 0.21
94 $\text{g C m}^{-2} \text{ day}^{-1}$ in that case. All other fluxes in the top 200 m decrease relative to the base case as well,
95 except for the export flux which increases to about 0.8% of primary production. This relative increase in
96 export is the result of larger sinking rates of phytoplankton and detritus ($w_{Phy}=0.95$, $w_{LDet}=4.97$; see table



97 2) than those used in the base case.

98 In contrast to experiment A, experiment B yields an increase of primary production relative to the
99 base case. The proportion of the grazing flux to primary production and the contribution of zooplankton
100 to the production of detritus also increase to about 59% and 52%, respectively. Unlike in the other three
101 experiments, carbon export in experiment B is dominated by the sinking of phytoplankton, reflecting its
102 large contribution to POC at 200 m. Although the simulated POC concentration at 200 m is very close to
103 the observations, the relative contributions of phytoplankton, zooplankton, and detritus are problematic
104 and likely do not result in a better estimation of carbon export (in this case 0.3% of primary production).

105 In experiment C, primary production is $0.30 \text{ g C m}^{-2} \text{ day}^{-1}$ with 24% flowing to zooplankton. The
106 mortality of zooplankton causes a flux of $0.047 \text{ g C m}^{-2} \text{ day}^{-1}$ to detritus, which accounts for 17% of the
107 production of detritus. Finally, about 24% of primary production is removed from the upper 200 m
108 through gravitational sinking. The simulated export ratio of 24% is within the wide range of reported
109 export ratios, from 6% to 43%, at 120 m depth in the Gulf of Mexico (see Table 3 of Hung et al., 2010).
110 Despite the high degree of uncertainty that exists when estimating export ratios (e.g., the global mean
111 export ratio varies from ~10% (Henson et al., 2012; Lima et al., 2014; Siegel et al., 2014) to ~20%
112 (Henson et al., 2015; Laws et al., 2000)), it is obvious that only experiment C reproduced an export ratio
113 of a reasonable magnitude. A more detailed validation of primary production and export fluxes will be
114 presented in the following sections.

15 **5. 3D biogeochemical model**

16 The optimal parameter sets from the 1D optimizations of A, B, and C were applied in the 3D model
17 for the whole GOM for five years (2011-2015). The performance of the 3D model can be regarded as a
18 cross-validation of the parameters optimized in 1D at different times and locations. It is possible that the
19 optimization algorithm has modified parameters to compensate for biases in the 1D simulations, e.g. the
20 absence of horizontal and vertical advection or the simplification of vertical diffusion, that degrades the
21 3D model performance. Indeed, directly applying the optimal parameter sets from 1D version to the 3D
22 model yields lower model-data agreement than the 1D counterpart and the following modifications to the
23 optimized parameters were made manually to bring the model-data agreement of 3D model in better



24 alignment with that of 1D version: the half-saturation for NH_4 uptake (k_{NH_4}) was decreased to 0.01 in
25 experiment B and C, and the aggregation parameter (τ) was decreased to 0.05 in the experiment C.

26 **5.1. Spatial patterns of surface chlorophyll**

27 First, the annual climatological surface chlorophyll from satellite and model are compared from 2011
28 to 2015. The satellite estimates show high chlorophyll in the coastal regions and low chlorophyll in the
29 deep ocean (Figure 7a). This spatial pattern of surface chlorophyll is well reproduced in all simulations
30 except in the experiment A which even fails to reproduce the relatively high chlorophyll near the
31 Mississippi-Atchafalaya river systems because of the high sinking rate of phytoplankton ($w_{\text{Phy}}=0.95$; see
32 Table 2). The largest model-data differences occur in the coastal regions, where all simulations
33 underestimate the observed surface chlorophyll because parameter optimization is only performed at one
34 station located in the deep ocean without considering the coastal environments. Based on this and the fact
35 that the floats operated in the deep ocean (Figure 1), only the model results in the deep ocean (depth >
36 1000 m) will be considered in the following discussion.

37 **5.2. Subsurface distributions**

38 Figure 8 shows the seasonal cycles of surface chlorophyll as well as the vertically integrated
39 chlorophyll, phytoplankton, and POC within the deep ocean (depth>1000 m). Analogous to the 1D
40 models, chlorophyll, phytoplankton, and POC are integrated over the upper 200 m. Comparisons of
41 vertical profiles between observations and model results are given in Figure 9. In general, the main
42 features in the 3D models are very similar to those in 1D. Experiment A cannot constrain the vertical
43 profiles of chlorophyll because of the inappropriate estimation of initial slope (α), experiment B
44 overestimates phytoplankton and its contribution to POC since the maximum ratio of chlorophyll to
45 carbon (θ_{max}) is weakly constrained, and experiment C shows significant improvements in the model-data
46 agreement. However, there are some differences between the 1D and 3D models. For example, the base
47 case of the 1D model overestimates the magnitude while underestimating the depth of the observed DCM.
48 Experiment B and C best improve the magnitude and depth of DCM, respectively. In contrast, in the 3D
49 model the vertical profile of chlorophyll and the magnitude of the DCM in the base case are already very
50 close to the observations and neither of the optimizations yield further improvement. These differences



51 between the 1D and 3D models might be a result of different spatio-temporal scales between the two
52 model versions, or the simplifications of physical processes in the 1D model.

53 We have also compared the chlorophyll-to-carbon ratio, primary production, and carbon export
54 fluxes from 1D and 3D models with observations (Figure 10). The chlorophyll-to-carbon ratio is
55 estimated as the vertically integrated chlorophyll divided by the phytoplankton in the upper 200 m (Figure
56 10a). As an important indicator of phytoplankton physiological status (Geider, 1987), the observed
57 chlorophyll-to-carbon ratio varies considerably in response to the ambient environment. In general, the
58 ratios derived from the 3D models are lower than their corresponding 1D values, but the differences are
59 still within the range of variability. Without utilizing the observations of phytoplankton and POC,
60 experiments A and B in both 1D and 3D versions underestimate the chlorophyll-to-carbon ratio. In
61 experiment C, the simulated chlorophyll-to-carbon ratios from 1D and 3D are in good agreement with the
62 observations although the observed variability is underestimated.

63 For reference, satellite-based primary production (PP) is provided by two algorithms, the Vertically
64 Generalized Production Model (VGPM, Behrenfeld and Falkowski 1997) and the Carbon-based
65 Productivity Model (CbPM, Westberry et al. 2008). As shown in Figure 10b, satellite-based PP differs
66 depending on the algorithm applied. PP results from all 3D simulations are qualitatively similar to the 1D
67 simulations. Experiment C provides the best estimates of PP when compared to satellite-based estimates
68 from VGPM and CbPM, both in 1D and 3D.

69 The base case and experiments A and B yield carbon export fluxes smaller by one to two orders of
70 magnitude than experiment C. Thus, only experiment C from the 1D and 3D models are shown in Figure
71 10b in comparison to observations from sediment traps (see supplementary material). The carbon export
72 fluxes at 200 m from the 1D and 3D are similar in magnitude although the 1D model yields higher fluxes
73 and larger variability. Despite the scarcity of carbon export observations in the GOM, the model estimates
74 are within the range of observations down to ~1,600 m and capture the observed declining trend of carbon
75 export with depth.

76 In summary, all the results above demonstrate the feasibilities of using the locally optimized
77 parameters from the 1D model to improve the 3D simulation. In addition, by incorporating all available



78 observations (i.e. surface chlorophyll from satellite estimates, profiles of chlorophyll, phytoplankton, and
79 POC from bio-optical floats), experiment C cannot only simulate the biogeochemical processes well in
80 the upper 200 m, but also reproduce the carbon export flux and its associated attenuation in the deep ocean
81 (200-1600m) of the GOM.

82 **6. Discussion**

83 **6.1. Trade-offs between different observations for parameter optimization**

84 The results of the optimization experiments vary dramatically depending on how many observation
85 types are used. Using only satellite surface chlorophyll in experiment A succeeds in reducing the misfits
86 of surface chlorophyll, but at the expense of the vertical structure since the predominant DCM disappears
87 in experiment A. Satellite surface chlorophyll alone cannot constrain several vital parameters, including
88 the initial slope of the productivity-irradiance curve (α) and the maximum ratio of chlorophyll to carbon
89 (θ_{max}), with confidence. This result highlights the importance of subsurface observations for parameter
90 optimization and similarly for model validation.

91 The floats provide valuable subsurface observations, but chlorophyll profiles alone are not sufficient
92 for parameter optimization. In experiment B, the addition of chlorophyll profiles leads to significant
93 improvements in vertical chlorophyll distributions; however, the profiles of phytoplankton and POC
94 deteriorate largely because the maximum ratio of chlorophyll to carbon (θ_{max}) is weakly constrained.
95 Using estimates of phytoplankton biomass and POC derived from backscatter measurements in
96 experiment C yields the best estimation of plankton-related state variables and carbon fluxes (i.e. primary
97 production and carbon export). Only in this experiment do the improvements obtained from observations
98 in the upper 200 m extend to the deep ocean as reflected in the improved carbon export estimates below
99 1,000 m.

00 It should be noted, however, that degradation of unoptimized variables did not occur in all
01 optimizations within experiments A and B. In some cases, the unoptimized fields were improved. For
02 example, the A2 optimization yields a 69% reduction in the misfit for subsurface chlorophyll (Figure 5d)
03 and large improvements of chlorophyll profiles (Figure S5a) even though no observations of subsurface
04 chlorophyll are used. Another example is that B1 optimization improves simulations of phytoplankton



05 and POC (Figure 5d and Figure S5b-c) through the correlations between the observed chlorophyll and
06 phytoplankton ($r^2 = 0.69$) and POC ($r^2 = 0.69$). Similar findings have been reported in Prunet et al. (1996b)
07 where the improvements of chlorophyll profiles within the mixed layer were obtained by using surface
08 chlorophyll in a 1D model. In a more recent study by Xiao and Friedrichs (2014a) where satellite data
09 was used subsurface fields were improved in addition to surface fields.

10 In optimizations A2 and B1, the improvement in unoptimized fields occurred because the poorly
11 constrained parameters were not optimized but well defined ($\alpha = 0.125$ in the optimization A2 and $\theta_{max} =$
12 0.0535 in the optimization B1; see table 2). Including the unconstrained parameters into the parameter
13 optimization can return a lower misfit with respect to the observations used in optimization but increases
14 the risk of overfitting and reduces the model's predictive skill, i.e. the ability to simulate unoptimized
15 observations and those collected at different locations and times. This is consistent with previous studies
16 (Friedrichs et al., 2006, 2007; Ward et al., 2010). For example, Friedrichs et al. (2006) optimized three
17 ecosystem models of different complexities against three seasons of observations and the resulting
18 parameters were used to quantify the predictive skill for the fourth season. Cross-validation showed that
19 exclusion of the poorly constrained parameters from the optimization increased the predictive skill.

20 Although prior knowledge of the parameters allows one to exclude those poorly constrained ones
21 from the optimization and thus can prevent degradation in unoptimized variables, most parameters are
22 poorly known. Thus, the ultimate resolution of this issue should be to increase availability of observations
23 so that more parameters can be constrained with confidence. In addition, even if the unconstrained
24 parameters are well-known, a lack of observations hampers our ability to recognize improvements in the
25 model's predictive skill and hence may prevent us from identifying the optimal solutions. For example,
26 without the observations of phytoplankton and POC, we could not have known that optimization B1
27 improved simulations of phytoplankton and POC, let alone that the optimization B1 was a better solution
28 than the optimization B2 (the experiment B) in terms of the lower total misfit as shown in Figure 5d.

29 It has been suggested that when performing a parameter optimization, not only parameter values but
30 also parameter uncertainties should be taken into account (Fennel et al., 2001; Ward et al., 2010;
31 Bagniewski et al., 2011). The parameter uncertainties can be assessed by performing an uncertainty



32 analysis (Fennel et al., 2001; Prunet et al., 1996a, 1996b), replicating the parameter optimization (Ward
33 et al., 2010), and cross-validating the resulting parameters (Xiao and Friedrichs, 2014a). In this study, a
34 cross-validation of the parameters was conducted by evaluating the model's predictive skill with respect
35 to different variables, times, and locations. Although this cross-validation at different times and locations
36 may give some indication of overfitting, it cannot determine whether the model reproduces observations
37 through wrong mechanisms because a small misfit of cross-validation can be caused by missing
38 validations of key variables or fluxes, e.g. ignorance of phytoplankton and PP in the experiment B, while
39 a large misfit can be a result of the intrinsic heterogeneity of biological parameters in different times
40 (Mattern et al., 2012) and locations (Kidston et al., 2011), e.g. underestimation of coastal surface
41 chlorophyll in the experiment C. Therefore, it is important to evaluate the predictive skill of unoptimized
42 variables.

43 Collectively, the discussion above highlights the values of BGC float data for parameter optimization
44 and model validation, not only because of their high spatio-temporal coverage but also their ability to
45 measure multiple properties simultaneously.

46 **6.2. Feasibilities of applying the local optimized parameters to 3D models**

47 As the high computational cost makes direct optimization for a 3D biogeochemical model
48 impractical, we performed parameter optimizations first in a 1D surrogate model with the same
49 biogeochemical component as the 3D model. However, there are some difficulties in porting the locally
50 optimized parameters to the basin-scale model. Firstly, the 1D model necessarily neglects horizontal
51 advection, which can result in differences between the 1D and 3D models. On the one hand, the optimized
52 parameters from the 1D model may have been adjusted to compensate for biases in the biological
53 properties caused by neglecting advection and, as a result, this may degrade the 3D simulations (Kane et
54 al., 2011). On the other hand, counter examples exist where the 3D simulations outperform the 1D models
55 (Hoshiba et al., 2018). Secondly, the spatial heterogeneity of parameters (e.g., Kuhn and Fennel 2019) is
56 another issue that influences the portability of parameters from 1D to 3D models. In some studies, the
57 parameter optimization has been performed at several contrasting stations with the goal of using different
58 parameter sets in different regions of the 3D model (Hoshiba et al., 2018). In other studies different



59 stations were optimized simultaneously to obtain one single optimized parameter set (Kane et al., 2011;
60 Oschlies and Schartau, 2005; Schartau and Oschlies, 2003). Such parameters compromise the misfit in
61 each single station but take account into all stations and can often yield an overall better simulation of all
62 these stations as shown by Kuhn and Fennel (2019).

63 In our study, the similarities in general features between the 1D and 3D models confirm the
64 portability of the resulting parameters in the deep ocean of the GOM while the underestimation of surface
65 chlorophyll in the coastal regions may result from the contrasting ecosystem functioning between coastal
66 regions and deep ocean. For example, the highly productive continental shelf in the northern GOM is
67 fueled by the large nutrient load from the Mississippi and Atchafalaya river systems with primary
68 production being as high as $4 \text{ g C m}^{-2} \text{ day}^{-1}$ near the Mississippi river delta (Fennel et al., 2011), while
69 the deep ocean is oligotrophic and nutrient limited with the primary production ranging from 0.2 to 0.5 g
70 $\text{C m}^{-2} \text{ day}^{-1}$ (see Figure 10).

71 7. Conclusions

72 In this study, we have performed parameter optimization for a 1D biogeochemical model and then
73 used the resulting parameters to generate simulations with a corresponding 3D model in the GOM. Three
74 experiments have been conducted by using different combinations of observations (surface chlorophyll
75 from satellite estimates, vertical profiles of chlorophyll, phytoplankton biomass and POC from BGC Argo
76 floats) in order to examine the trade-offs between the different observations for parameter optimization.
77 Two misfit metrics have been defined using the case-specific misfit and the total misfit to measure the
78 models' abilities to reproduce the optimized and unoptimized observations.

79 Model results show that satellite surface chlorophyll alone cannot reproduce well the vertical
80 structures in a biogeochemical model unless profile observations are used in addition. BGC Argo floats
81 are an excellent platform for obtaining such observations at high spatio-temporal coverage and for a
82 relatively broad suite of parameters. Only adding chlorophyll profiles is not sufficient because it fails to
83 constrain the ratio of chlorophyll to phytoplankton, but the addition of backscatter, which allows
84 estimation of phytoplankton biomass and POC, enables us to constrain the subsurface carbon state
85 variables and reproduce well PP and carbon export fluxes to below 1000 m depth. Finally, our 3D model



86 was improved and reproduced similar results as the 1D version, which is promising for the application of
87 parameter optimization.

88

89 *Code and data availability:* The ROMS model code can be accessed at <http://www.myroms.com> (last
90 access: 16 June 2016). HYCOM data can be downloaded at <http://tds.hycom.org/thredds/dodsC/datasets>
91 (last access: 16 August 2018). Profiling data from the BGC-Argo floats are available at the National
92 Oceanographic Data Center (NOAA), <https://data.nodc.noaa.gov/cgi-bin/iso?id=gov.noaa.nodc:159562>
93 (Hamilton and Leidos, 2017)

94

95 *Author contributions.* BW and KF conceived the study. BW carried out optimization experiments, model
96 simulations and analyses. LY assisted with set-up and validation of the 3D model. CG assisted with
97 processing of the BGC float data. BW and KF discussed the results and wrote the paper with contributions
98 from the coauthors.

99

00 *Competing interests.* The authors declare that they have no conflict of interest.

01

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29 **Table list**

30 Table 1. Initial values and ranges of primary parameters used in the biogeochemical model

Descriptions (unit)	Symbol	Value	Range
Radiation threshold for nitrification (W m^{-2})	I_0	0.0095 ^a	0.005 ^b -0.01 ^b
Half-saturation radiation for nitrification (W m^{-2})	k_I	0.1 ^a	0.01 ^b -0.5 ^b
Maximum nitrification rate (day^{-1})	n_{max}	0.2 ^c	0.01 ^b -0.35 ^b
Phytoplankton growth at 0 °C (Dimensionless)	μ_0	0.69 ^a	0.1 ^b -3.0 ^b
Initial slope of P-I curve ($\text{mg_C (mg_Chl W m}^{-2} \text{ day)}^{-1}$)	α	0.125 ^a	0.007 ^a -0.13 ^a
Half-saturation for NO_3 uptake (mmol_N m^{-3})	k_{NO3}	0.5 ^a	0.007 ^a -1.5 ^a
Half-saturation for NH_4 uptake (mmol_N m^{-3})	k_{NH4}	0.5 ^a	0.007 ^a -1.5 ^a
Phytoplankton mortality (day^{-1})	m_P	0.075	0.01 ^b -0.2 ^b
Aggregation parameter (day^{-1})	τ	0.1	0.01 ^b -25 ^b
Maximum chlorophyll to carbon ratio (mg_Chl mg_C^{-1})	θ_{max}	0.0535 ^c	0.005 ^a -0.15 ^b
Phytoplankton sinking velocity (m day^{-1})	w_{Phy}	0.1 ^a	0.009 ^a -25 ^a
Maximum grazing rate (day^{-1})	g_{max}	0.6 ^a	0.1 ^b -4 ^b
Half-saturation for phytoplankton ingestion ($(\text{mmol_N m}^{-3})^2$)	k_P	0.5	0.01 ^b -3.5 ^a
Zooplankton assimilation efficiency (Dimensionless)	β	0.75 ^a	0.25 ^b -0.75 ^b
Zooplankton basal metabolism (day^{-1})	l_{BM}	0.01	0.01 ^b -0.15 ^b
Zooplankton specific excretion (day^{-1})	l_E	0.1 ^a	0.05 ^b -0.35 ^b
Zooplankton mortality (day^{-1})	m_Z	0.2	0.02 ^b -0.35 ^b
Small detritus remineralization (day^{-1})	r_{SD}	0.3 ^c	0.005 ^b -0.25 ^a
Large detritus remineralization (day^{-1})	r_{LD}	0.1	0.005 ^b -0.25 ^a
Small detritus sinking velocity (m day^{-1})	w_{SDet}	0.1 ^a	0.009 ^a -25 ^a
Large detritus sinking velocity (m day^{-1})	w_{LDet}	1 ^a	0.009 ^a -25 ^a

31 a Fennel et al. (2006); b Kuhn et al. (2018); c Yu et al. (2015)



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Table 2. The best fit of parameter set for each experiment

	w_{Phy}	m_P	k_{NH4}	τ	θ_{max}	α	w_{LDet}
Base	0.1000	0.0750	0.5000	0.1000	0.0535	0.1250	1.000
A1	0.0608	0.0100	1.5000	--	--	--	--
A2	0.6863	0.0100	0.0195	--	0.0169	--	--
A3	1.6567	0.1978	0.1004	--	0.0250	0.0219	--
A4	0.9468	0.0737	0.2454	--	0.0191	0.0101	4.9694
B1	0.2863	0.0983	1.5000	--	--	--	--
B2	0.4217	0.0130	0.0300	--	0.0158	--	--
B3	2.1016	0.0176	1.5000	--	0.0346	0.0079	--
B4	0.0090	0.0100	1.5000	--	0.0361	0.0405	8.3514
	w_{Phy}	r_{LD}	m_P	τ	k_{NH4}	w_{LDet}	θ_{max}
Base	0.1000	0.1000	0.0750	0.1000	0.5000	1.0000	0.0535
C1	1.9231	0.2500	0.1805	--	--	--	--
C2	0.9755	0.2500	0.0100	1.1402	--	--	--
C3	0.4071	0.0630	0.0100	1.8531	0.0070	--	--
C4	0.0090	0.0050	0.0634	0.0995	0.0431	5.6623	--
C5	0.0090	0.2245	0.0100	0.6451	1.5000	2.5202	0.0614

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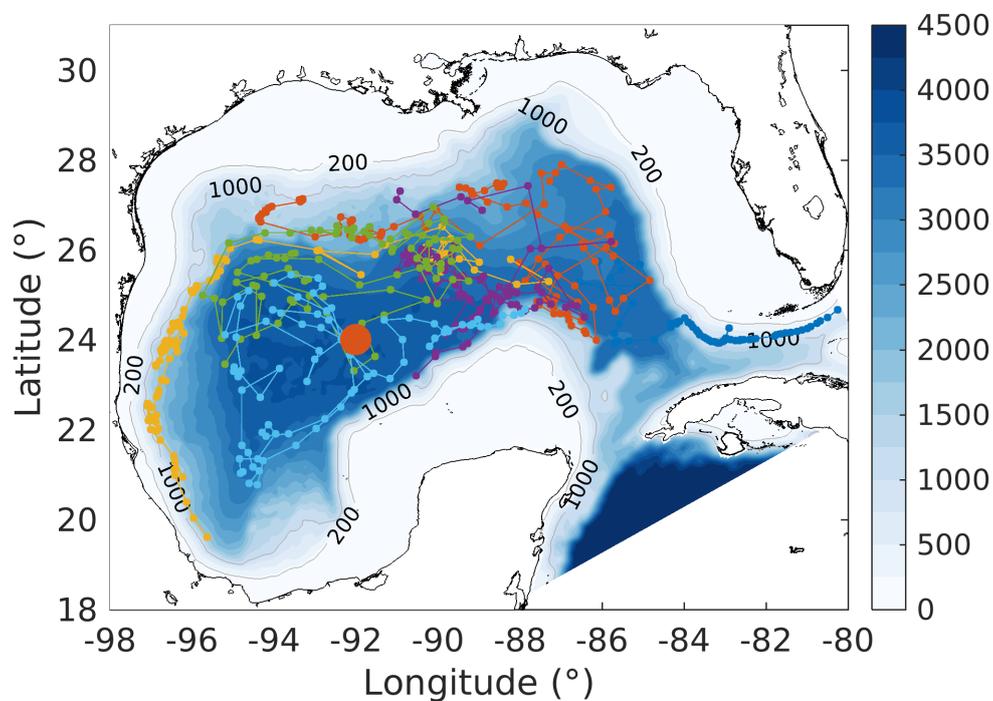


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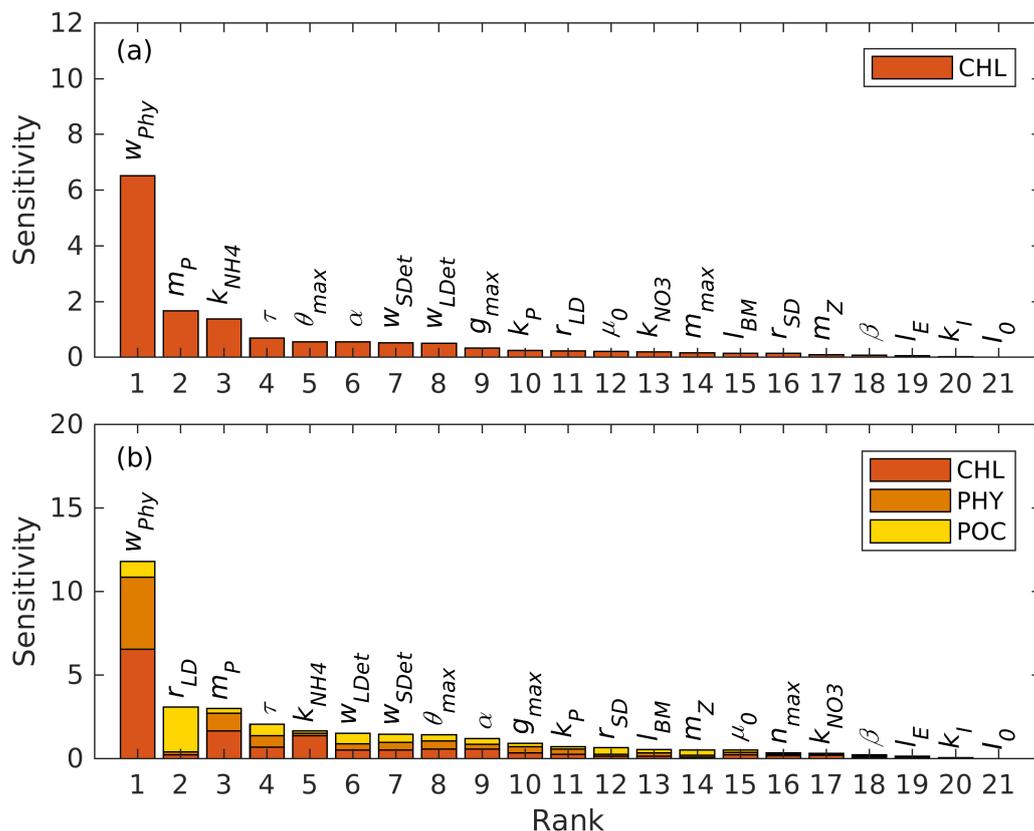
44 **Figure captions**

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47 Figure 1. Model bathymetry (unit: m) with trajectories of six bio-optical floats (small colored dots and
48 lines) which operated in the Gulf of Mexico from 2011 to 2015. The location of the 1D model is denoted
49 by the large orange dot.

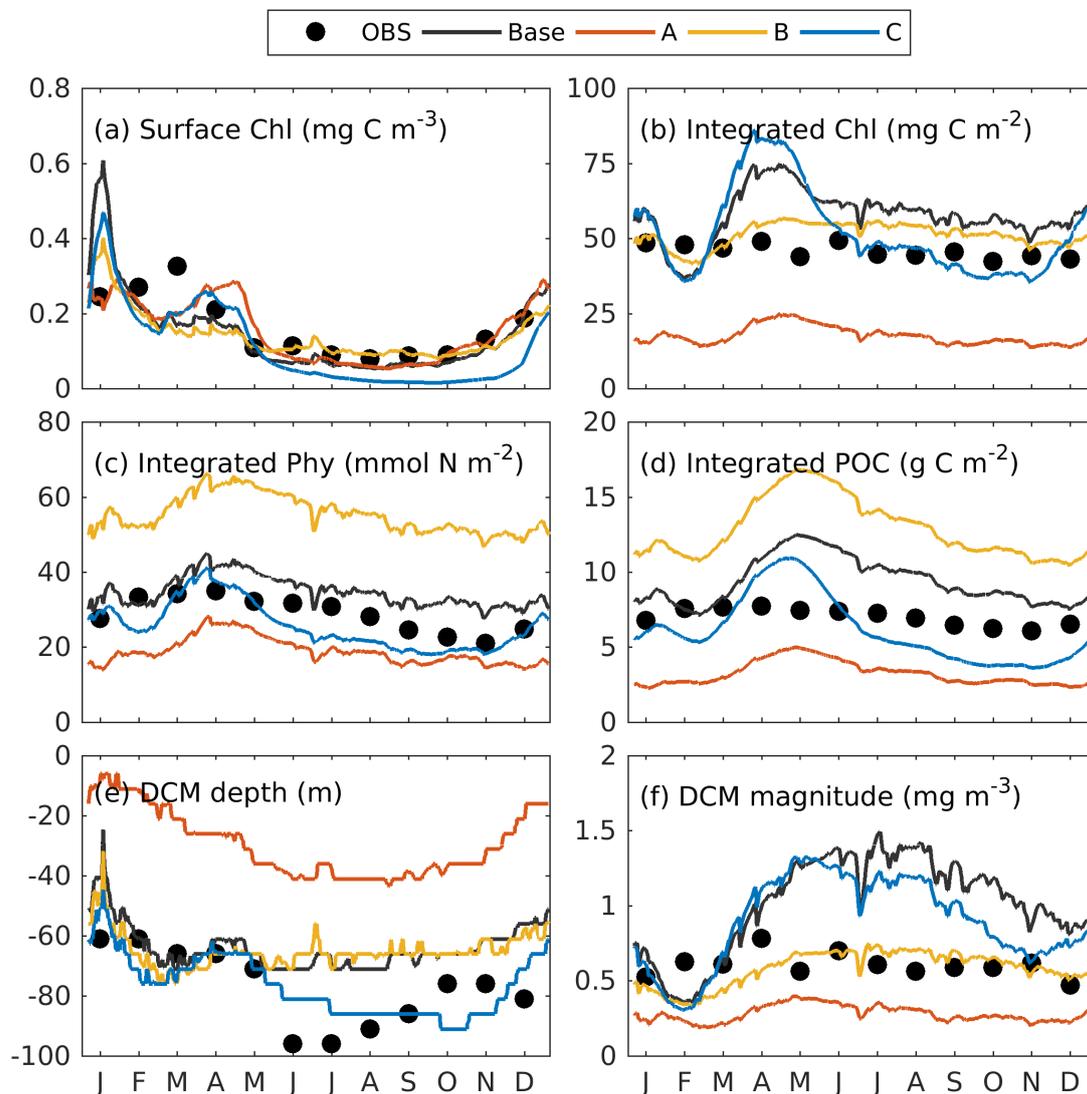


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51 Figure 2. Parameter sensitivities (unit: dimensionless) with respect to (a) chlorophyll and (b) the sum of
 52 chlorophyll, phytoplankton, and POC.

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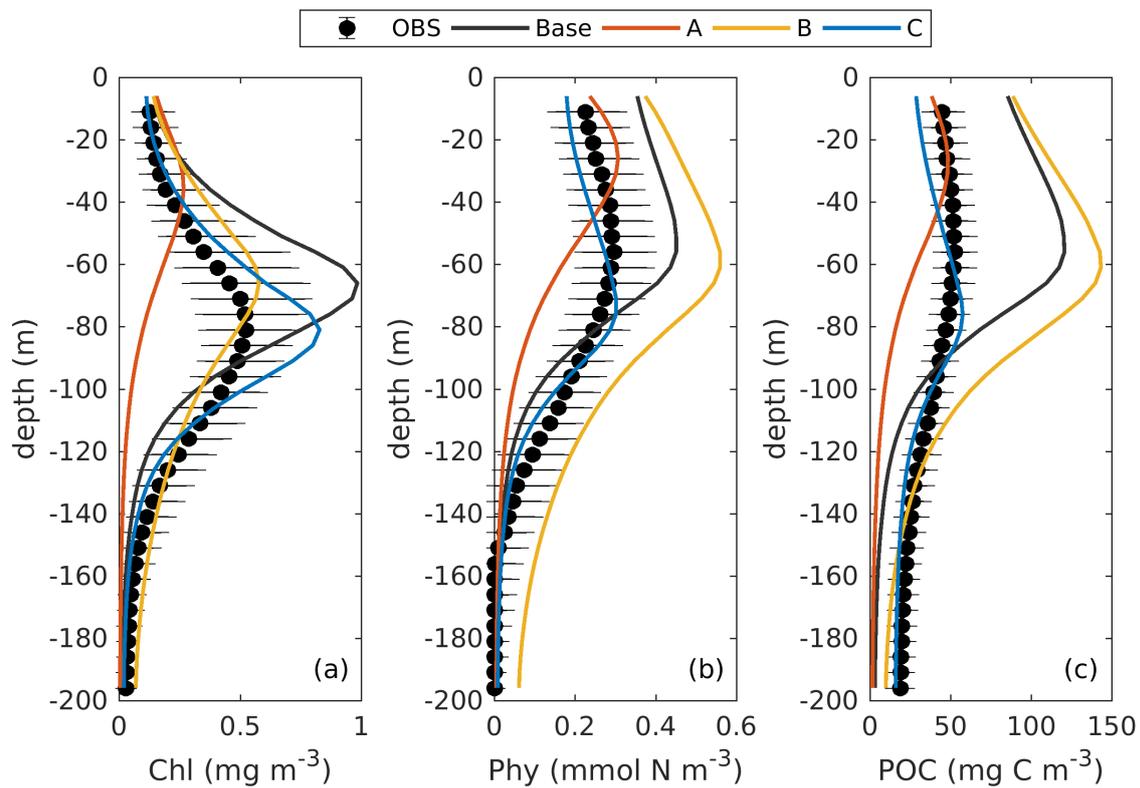


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56 Figure 3. Observed (black dots) and simulated (colored lines) annual cycle of surface chlorophyll (a),
57 vertically integrated chlorophyll (b), vertically integrated phytoplankton (c), vertically integrated POC
58 (d), and the depth (e) and magnitude (f) of the DCM. Chlorophyll, phytoplankton, zooplankton, and POC
59 are integrated over the top 200 m.

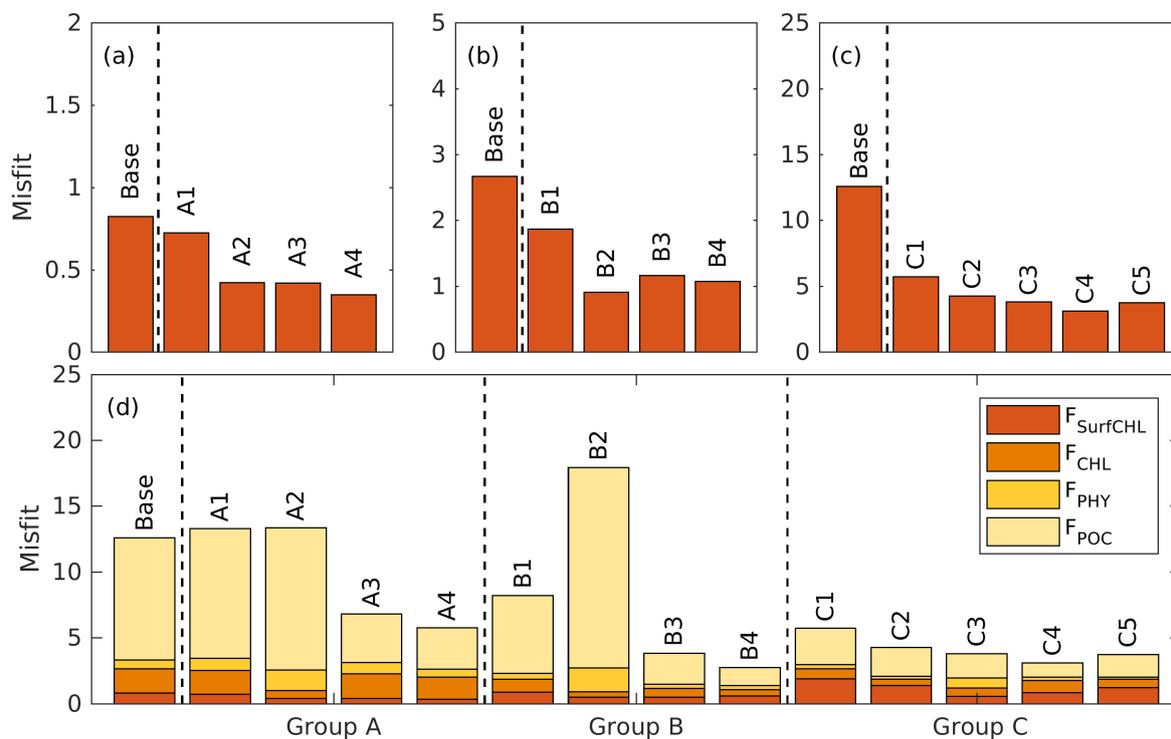
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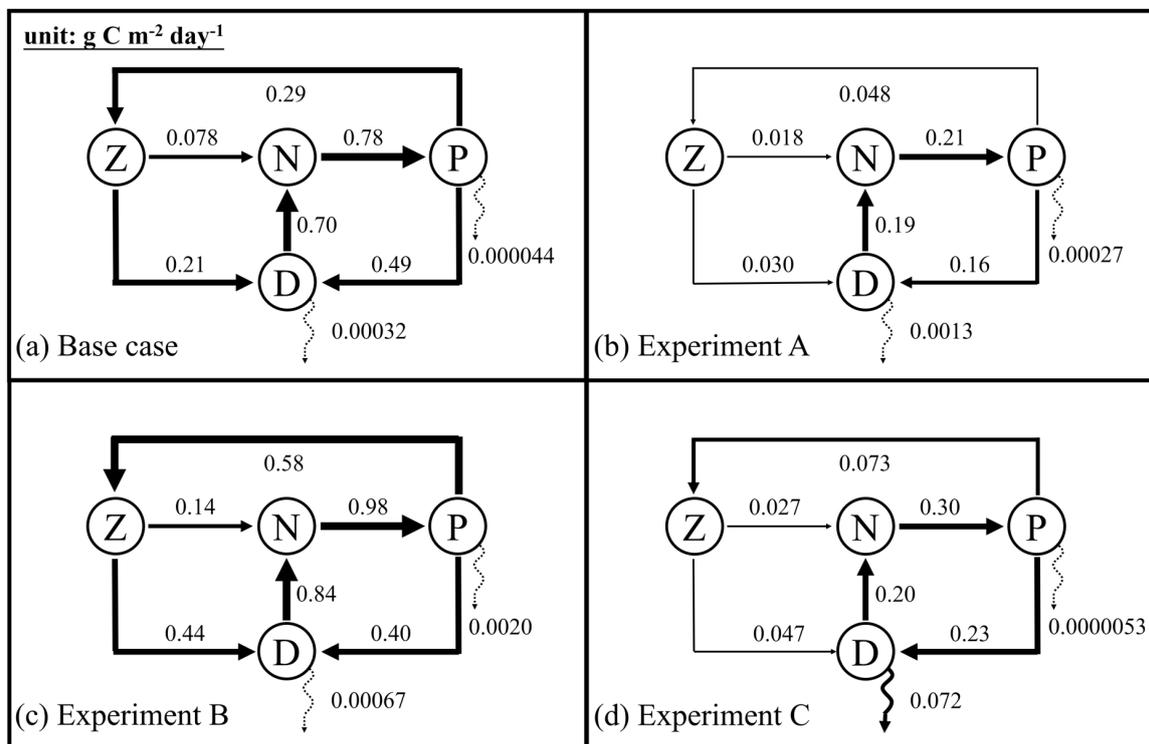


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63 Figure 4. Observed (black dots with error bars) and simulated (colored lines) vertical profiles of
64 chlorophyll, phytoplankton, and POC.

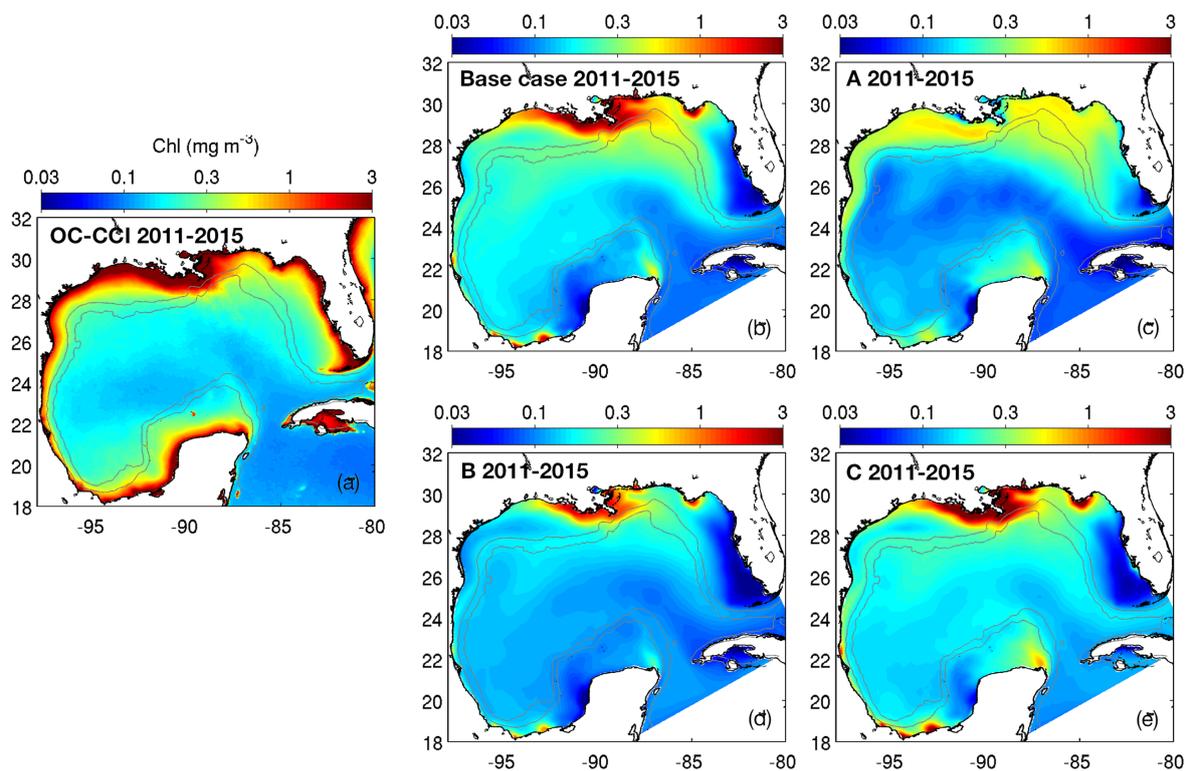


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 66 Figure 5. The case-specific cost function values (a-c) and total misfits (d) of the base case and the
 67 different optimizations.



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 69 Figure 6. Annually averaged carbon fluxes integrated over the upper 200 m (unit: $\text{g C m}^{-2} \text{ day}^{-1}$) for the
 70 base case (a) and optimized experiments A, B, and C. The N, P, Z, and D stand for the pools of nutrient,
 71 phytoplankton, zooplankton, and the sum of small and large detritus, respectively. The thickness of arrows
 72 scales with the magnitude of fluxes. Dashed arrows represent fluxes lower than $0.01 \text{ g C m}^{-2} \text{ day}^{-1}$.

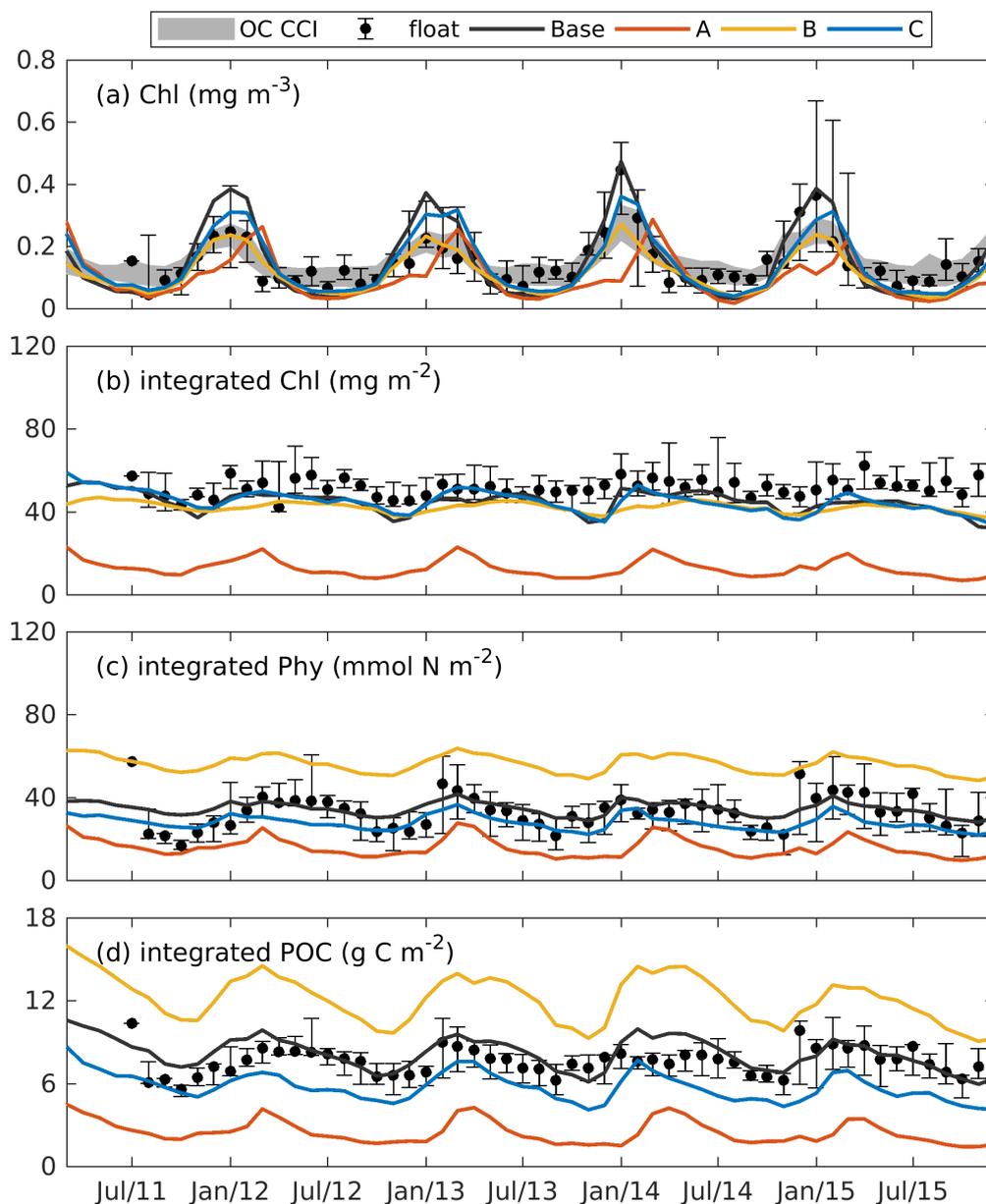
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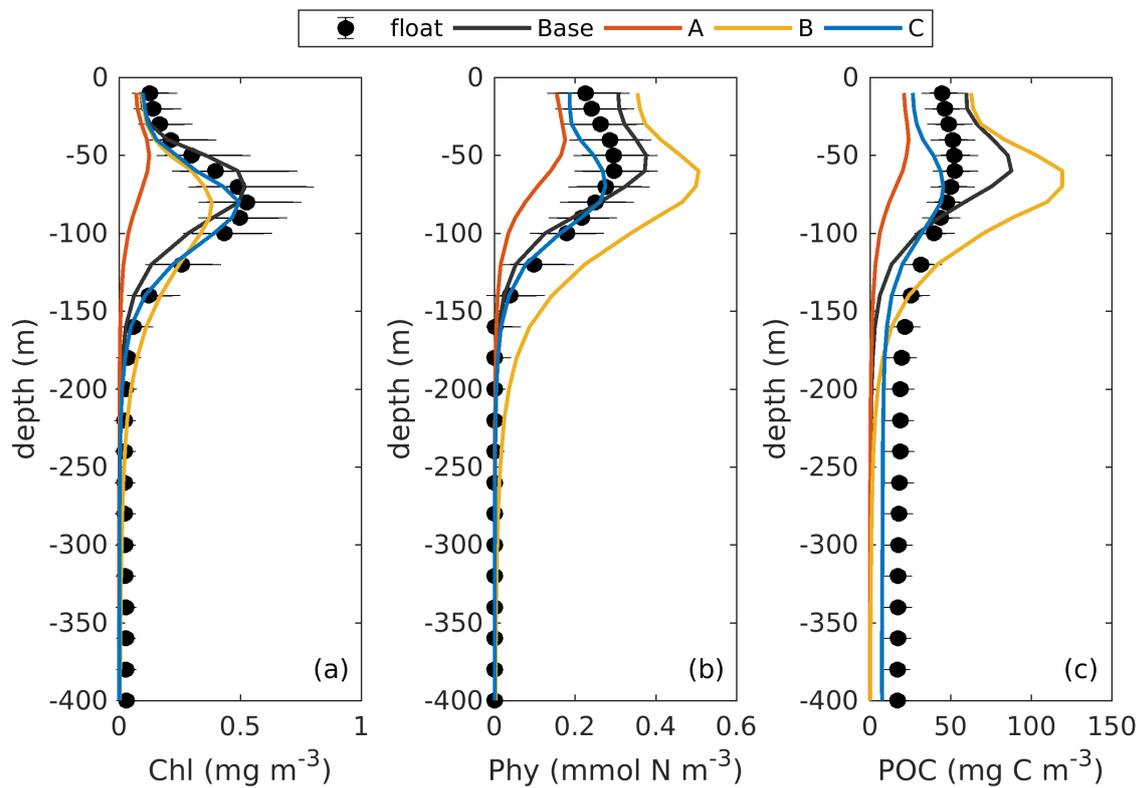
75 Figure 7. Spatial distributions of the annual mean chlorophyll in the surface layer from the satellite (OC-
76 CCI) climatology (2011-2015) and the different model versions. The gray contours mark the bathymetric
77 depths of 200 and 1000 m.

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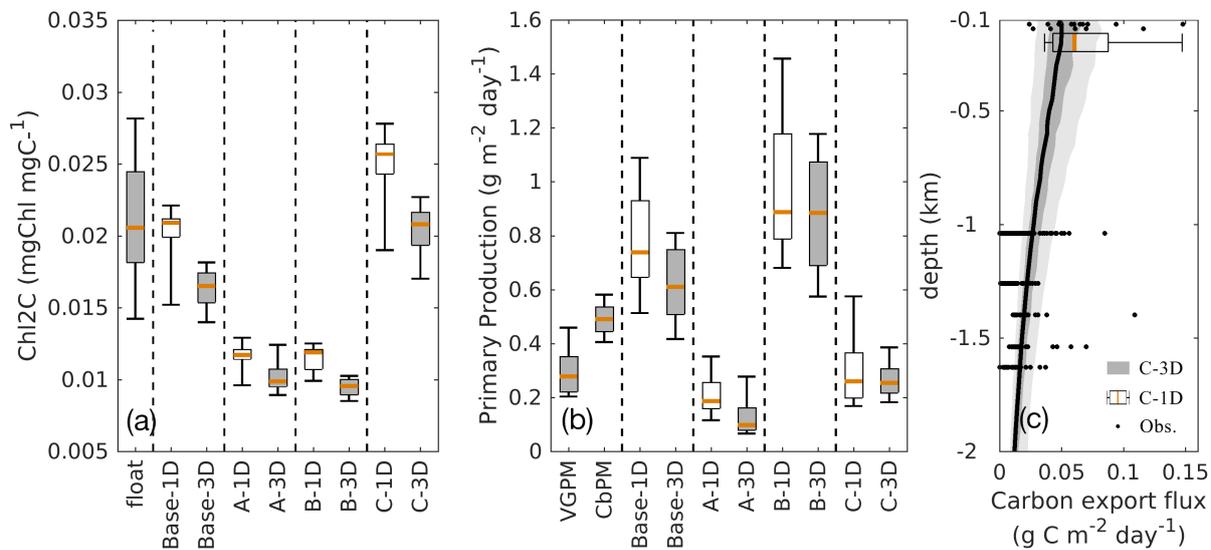
80 Figure 8. Observed and simulated seasonal cycles of surface chlorophyll (a), vertically
81 chlorophyll (b), vertically integrated phytoplankton (c), and vertically integrated POC (d) from each 3D
82 models. Solid lines represent the median values over the deep ocean of GOM (depth>1000m). Error bars
83 and shades show the 25% and 75% percentiles. Chlorophyll, phytoplankton, and POC are integrated over
84 the top 200m.



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86 Figure 9. Observed and simulated vertical profiles of chlorophyll, phytoplankton, and POC from each
87 3D models.

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90 Figure 10. Comparisons of the chlorophyll to carbon ratio (a), primary production (b), and carbon export
91 fluxes (c) between the 1D and 3D models.

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