

Improved consistency between the modelling of ocean optics, biogeochemistry and physics, and its impact on the North-West European Shelf seas

Jozef Skakala^{1,1}, Jorn Bruggeman^{2,2}, David Andrew Ford^{3,3}, Sarah L Wakelin^{4,4}, Anıl Akpınar^{4,4}, Tom Hull^{5,5}, Jan Kaiser^{6,6}, Benjamin Roger Loveday^{1,1}, Charlotte Anne June Williams^{4,4}, and Stefano Ciavatta^{7,7}

¹Plymouth Marine Laboratory

²PML

³Met Office

⁴National Oceanography Centre

⁵Centre for Environment, Fisheries and Aquaculture Science

⁶University of East Anglia

⁷Plymouth Marine Laboratory/National Centre for Earth Observation

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Abstract

We use a recently developed spectrally resolved bio-optical module to better represent the interaction between the incoming irradiance and the heat fluxes in the upper ocean within the (pre-)operational physical-biogeochemical model on the North-West European (NWE) Shelf. The module attenuates light based on the simulated biogeochemical tracer concentrations, and thus introduces a two-way coupling between the biogeochemistry and physics. We demonstrate that in the late spring-summer the two-way coupled model heats up the upper oceanic layer, shallows the mixed layer depth and influences the mixing in the upper ocean. The increased heating in the upper oceanic layer reduces the convective mixing and improves by ~5 days the timing of the late phytoplankton bloom of the ecosystem model. This improvement is relatively small compared with the existing model bias in bloom timing, but sufficient to have a visible impact on model skill. We show that the changes to the model temperature and salinity introduced by the module have mixed impact on the physical model skill, but the skill can be improved by assimilating the observations of temperature, salinity and chlorophyll into the model. However, in the situations where we improved the simulation of temperature, either via the bio-optical module, or via assimilation of temperature and salinity, we have shown that we also improved the simulated oxygen concentration as a result of the changes in the simulated air-sea gas flux. Overall, comparing different 1-year experiments showed that the best model skill is achieved with joint physical-biogeochemical assimilation into the two-way coupled model.

1 Highlights

2 **Improved consistency between the modelling of ocean optics, bio-** 3 **geochemistry and physics, and its impact on the North-West Eu-** 4 **ropean Shelf seas**

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- 8 • We established two-way coupling between biogeochemistry and physics.
- 9 • We assessed the impact of the coupling in free and assimilative exper-
10 iments.
- 11 • The two-way coupling improves the simulated biogeochemistry on the
12 NWE Shelf.
- 13 • We recommend to implement this development into the operational
14 system.

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Jozef Skákala^{a,b}, Jorn Bruggeman^a, David Ford^c, Sarah Wakelin^d, Anıl
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Williams^d, Stefano Ciavatta^{a,b}

^a*Plymouth Marine Laboratory, Prospect Place, The Hoe, Plymouth, PL1 3DH, United Kingdom*

^b*National Centre for Earth Observation, Prospect Place, The Hoe, Plymouth, PL1 3DH, United Kingdom*

^c*Met Office, FitzRoy Road, Exeter, EX1 3PB, United Kingdom*

^d*National Oceanography Centre, Joseph Proudman Building, 6 Brownlow Street, Liverpool, L3 5DA, United Kingdom*

^e*Centre for Environment, Fisheries and Aquaculture Science, Lowesoft, NR33 0HT, United Kingdom*

^f*Centre for Ocean and Atmospheric Science, University of East Anglia, Norwich, NR4 7TJ, United Kingdom*

^g*Innoflair UG, Richard-Wagner-Weg 35, Darmstadt, 64287, Germany*

Abstract

We use a recently developed spectrally resolved bio-optical module to better represent the interaction between the incoming irradiance and the heat fluxes in the upper ocean within the (pre-)operational physical-biogeochemical model on the North-West European (NWE) Shelf. The module attenuates light based on the simulated biogeochemical tracer concentrations, and thus introduces a two-way coupling between the biogeochemistry and physics. We demonstrate that in the late spring-summer the two-way coupled model heats up the upper oceanic layer, shallows the mixed layer depth and influences the mixing in the upper ocean. The increased heating in the upper oceanic layer reduces the convective mixing and improves by ~ 5 days the timing of the late phytoplankton bloom of the ecosystem model. This improvement is relatively small compared with the existing model bias in bloom timing, but sufficient to have a visible impact on model skill. We show that the changes to the model temperature and salinity introduced by the module have mixed

impact on the physical model skill, but the skill can be improved by assimilating the observations of temperature, salinity and chlorophyll concentrations into the model. However, in the situations where we improved the simulation of temperature, either via the bio-optical module, or via assimilation of temperature and salinity, we have shown that we also improved the simulated oxygen concentration as a result of the changes in the simulated air-sea gas flux. Overall, comparing different 1-year experiments showed that the best model skill is achieved with joint physical-biogeochemical assimilation into the two-way coupled model.

22 *Keywords:* two-way coupled physical-biogeochemical model, ocean
 23 chlorophyll concentration, sea surface temperature, phytoplankton spring
 24 bloom, North-West European Shelf (10E-10W, 40N-68N), data assimilation

25 1. Introduction

26 Physical-biogeochemical ocean models are an essential element in mon-
 27 itoring and forecasting of global and shelf-sea ecosystem indicators ([1, 2]).
 28 However, coupled physical-biogeochemical marine modelling is a complex un-
 29 dertaking and a common way to simplify coupled models is to neglect the
 30 impact of the biogeochemical model state on physics ([3, 2]). Although ma-
 31 rine ecosystem models often neglect the coupling from the biogeochemical
 32 model state to physics, there are number of established mechanisms through
 33 which biogeochemistry influences physics and climate ([4, 1, 2]): (i) marine
 34 ecosystems play an essential part in the carbon cycle through biological and
 35 microbial carbon pump, influencing atmospheric carbon concentrations and
 36 the Earth surface temperature, (ii) phytoplankton influences oceanic albedo
 37 (e.g. [5]) having an overall impact on the radiative terms and Earth energy
 38 budget, (iii) some biogeochemical tracers influence light attenuation, mod-
 39 ifying the short-wave heat fluxes in the water column and therefore ocean
 40 stratification ([6, 7, 8, 9, 10, 11, 12, 13, 14]), and (iv) marine ecosystems have
 41 an impact on cloud condensation nuclei through the production of dimethyl
 42 sulfide (DMS, [15, 16, 17, 18]), or more directly via bubble formation ([19]).
 43 The size of life’s impact on Earth’s physics has been subject to much de-
 44 bate ([2]), often in connection with “the Gaia hypothesis” ([20, 21]), which
 45 proposes that life plays a central role in regulating climate.

46 For coupled physical-biogeochemical marine models the main source of
 47 impact of ecosystems on physics is through the absorption and backscatter-

ing of short-wave radiation by some biogeochemical substances in the sea water (e.g. [10]). The presence of optically active tracers, such as chlorophyll, suspended particulate matter (SPM), or colored dissolved organic matter (CDOM), in the oceanic upper layer increases light attenuation near the oceanic surface, warms the sea temperature in the upper ocean, which typically influences the mixing in the upper oceanic layer (e.g. [6]), e.g. shallowing the thermocline and the mixed layer depth (MLD). The changes to the vertical mixing can in turn impact the biogeochemical model, by influencing the nutrient concentrations and growth conditions in the upper ocean.

In this work we focus on the Copernicus Marine Environmental Monitoring Service (CMEMS) operational system for the North-West European (NWE) Shelf biogeochemistry, which is of a substantial societal benefit, as the NWE Shelf is a key region for fisheries, and an important contributor to the global carbon cycle ([22, 23, 24]). The presently used physical-biogeochemical operational model for the NWE Shelf is the marine physical model Nucleus for European Modelling of the Ocean (NEMO, [25]) coupled through the Framework for Aquatic Biogeochemical Models (FABM, [26, 27]) to the European Regional Seas Ecosystem Model (ERSEM, [28, 29, 30]). NEMO-FABM-ERSEM drives its physics and biogeochemistry by two separate irradiance modules: (i) the physical model calculates heat fluxes from the incoming net short-wave radiation (SWR) split into two wavebands, the 400-700nm visible band reduced through attenuation obtained from a monthly climatology of a satellite surface K_d product at 490nm wavelength (European Space Agency product version 2.0, <https://www.esa-oceancolour-cci.org/>), and the UV/infrared band reduced with a preset attenuation with an e-folding scale of 0.35m, (ii) the biogeochemical model reduces incoming photosynthetic active radiation (PAR) by taking into account both absorption and backscattering by the sea water and the simulated Phytoplankton Functional Types (PFTs), and also by including absorption by Particulate Organic Matter (POM), CDOM and sediment represented by an external satellite product (for details see [29, 31]). The presently used scheme means that, although some impact of biogeochemical tracers on the physical model is implicitly included in the 490nm K_d satellite climatology, there is no feedback from the biogeochemical model state to the simulated physics.

In [31] we implemented into ERSEM a stand-alone bio-optical module (based on OASIM, [32, 33, 34]), that resolves irradiance spectrally and splits the irradiance into diffuse and direct streams ([35]). The module then propagates irradiance through the water column, based on attenuation by sea water

86 and the biogeochemical substances in the water. The new module drove only
87 the biogeochemical part of the coupled NEMO-FABM-ERSEM model, sub-
88 stantially improving the underwater irradiance, but without a major impact
89 on the ERSEM model skill on the NWE Shelf ([31]). This version of NEMO-
90 FABM-ERSEM model will be used in the present study as a reference run
91 and will be called a “one-way coupled model”. In this work we expand the
92 development implemented in [31] by using the bio-optical module to drive
93 both the biogeochemistry and the physics (i.e. heating by light absorption).
94 Since the physical heat fluxes will be driven by the underwater irradiance
95 that is attenuated by biogeochemical substances, the module establishes an
96 important feedback from the biogeochemical model to physics. We will fur-
97 ther call this new implementation a “two-way coupled model”, to distinguish
98 it from the “one-way coupled” reference run.

99 This work aims at answering two main questions: (i) What is the size
100 of the biogeochemical impact on the marine physics within the NWE Shelf?
101 (ii) Does the impact of the spectrally resolved bio-optical module on physics
102 lead to more internally consistent ecosystem dynamics on the NWE Shelf,
103 and hence, an improvement in the operational biogeochemical model skill?
104 Those two questions are answered both in the context of free simulations
105 and also in the context of (physical, biogeochemical, coupled) assimilative
106 runs. The second question is particularly relevant: It has been established
107 that NEMO-FABM-ERSEM displays on the NWE Shelf late and intense
108 spring blooms ([31, 36]). Since a spring bloom is a major ecosystem driver
109 ([37, 38]), the simulated late blooms severely limit the ecosystem model skill
110 ([31, 36]). Although many factors can influence the bloom timing (including
111 biological drivers, such as zooplankton grazing, e.g. [39]), one of the leading
112 hypotheses for how phytoplankton blooms are triggered in the North Atlantic
113 is based on the interplay between PAR and an effective mixing depth (the
114 critical turbulence hypothesis, [40, 41]), i.e. the bloom sets in when the
115 effective mixing depth becomes fully contained within the euphotic layer
116 ([42]). Within the scope of the critical turbulence hypothesis, the delay in
117 bloom timing could then be explained by multiple components within the
118 physical-biogeochemical coupled model: (a) atmospheric wind stress forcing,
119 (b) model upper-ocean mixing scheme, (c) vertical stratification (thermocline
120 and pycnocline), (d) incoming surface PAR, (e) underwater light attenuation,
121 (f) the phytoplankton growth response to light (e.g. model parameters, such
122 as P-I curves, maximum PFT chlorophyll-to-carbon ratios). In [31] we have
123 addressed to a varying degree the points (d) and (e) without a significant

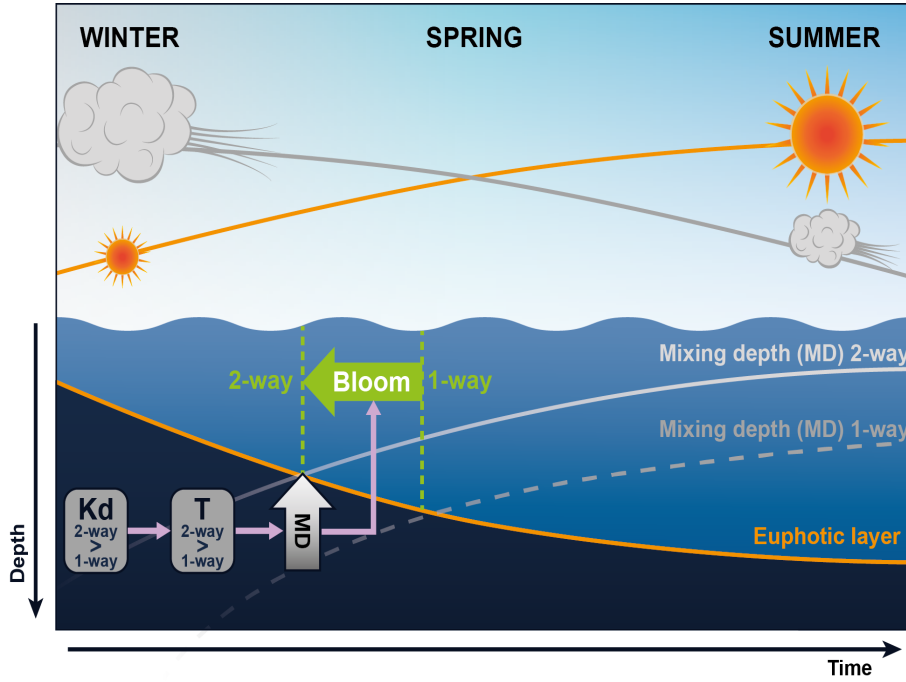


Figure 1: A schematic representation of the hypothesis about the impact of the two-way coupled model on the timing of the simulated bloom.

124 impact on the bloom timing. However, [31] observed that attenuation of
 125 light based on the satellite K_d product for the 490 nm wavelength is most
 126 likely an underestimate of the total PAR absorbed in the upper oceanic layer.
 127 Calculating heat fluxes using the bio-optical module is therefore expected to
 128 produce extra heat in the upper oceanic layer (Fig.5 of [31]), which is thought
 129 to shallow the MLD, but it can also reduce turbulent convective mixing near
 130 the oceanic surface ([43, 44]). The hypothesis tested in this work (see Fig.1) is
 131 that the reduced convective mixing can lead to a shallower turbulent mixing
 132 depth and help trigger an earlier phytoplankton bloom, as suggested by the
 133 critical turbulence hypothesis ([40, 44]). The biogeochemical feedback to the
 134 simulated physics could therefore improve the ERSEM skill on the NWE
 135 Shelf.

2. Methods

2.1. The physical model: NEMO

The NEMO ocean physics component (OPA) is a finite difference, hydrostatic, primitive equation ocean general circulation model ([25]). The NEMO configuration used in this study is similar to the one used by [45, 46, 31], and identical to the configuration used in [36]: we use the CO6 NEMO version, based on NEMOv3.6, a development of the CO5 configuration explained in detail by [47]. The model has 7 km spatial resolution on the Atlantic Margin Model (AMM7) domain using a terrain-following $z^* - \sigma$ coordinate system with 51 vertical levels ([48]). The lateral boundary conditions for physical variables at the Atlantic boundary were taken from the outputs of the Met Office operational 1/12° North Atlantic model (NATL12, [49]); the Baltic boundary values were derived from a reanalysis produced by the Danish Meteorological Institute for CMEMS. We use annually varying river discharge based on data from [50].

The model was forced at the surface by atmospheric fluxes provided by an hourly and 31km resolution realisation (HRES) of the ERA5 data-set (<https://www.ecmwf.int/>). In case of the one-way coupled model the ERA5 fluxes provide also the total incoming net shortwave radiation whose visible fraction is attenuated inside the water column based on the K_d for 490nm wavelength supplied by a monthly climatology from an Ocean Color - Climate Change Initiative (OC-CCI) product of European Space Agency (ESA), version 4.1 (<https://www.esa-oceancolour-cci.org/>). For the two-way coupled model the incoming net short-wave radiation is decomposed into direct and diffuse streams and spectrally resolved, and is provided by the bio-optical module ([31]) that will be described later in the ecosystem model section. The direct and diffuse streams are attenuated throughout the water column by the bio-optical module, and subsequently integrated by NEMO to calculate the heating within each vertical layer.

2.2. The ecosystem model: ERSEM

ERSEM ([28, 29]) is a lower trophic level ecosystem model for marine biogeochemistry, pelagic plankton, and benthic fauna ([51]). The model splits phytoplankton into four functional types largely based on their size ([28]): picophytoplankton, nanophytoplankton, diatoms and dinoflagellates. ERSEM uses variable stoichiometry for the simulated plankton groups ([52, 53]) and each Phytoplankton Functional Type (PFT) biomass is represented in terms

172 of chlorophyll, carbon, nitrogen and phosphorus, with diatoms also repre-
 173 sented by silicon. ERSEM predators are composed of three zooplankton
 174 types (mesozooplankton, microzooplankton and heterotrophic nanoflagel-
 175 lates), with organic material being decomposed by one functional type of
 176 heterotrophic bacteria ([29]). The ERSEM inorganic component consists of
 177 nutrients (nitrate, phosphate, silicate, ammonium and carbon) and dissolved
 178 oxygen. The carbonate system is also included in the model ([54]).

179 We applied in this study the ERSEM configuration from [36], based on a
 180 new ERSEM version 20.10, which has an updated benthic component with
 181 respect to [29]. The ERSEM parametrization is identical to the one described
 182 in [29]. The Atlantic boundary values for nitrate, phosphate, silicate and
 183 oxygen were taken from World Ocean Atlas ([55]) and dissolved inorganic
 184 carbon from the GLODAP gridded dataset ([56, 57]), while plankton and
 185 detritus variables were set to have zero fluxes at the Atlantic boundary.

186 The irradiance at the ocean surface was calculated using the bio-optical
 187 module implemented into the NEMO-FABM-ERSEM AMM7 configuration
 188 by [31]. The bio-optical module resolves irradiance spectrally and distin-
 189 guishes between downwelling direct and diffuse streams. The module is
 190 forced by the ERA5 atmospheric inputs (<https://www.ecmwf.int/>) for to-
 191 tal vertically integrated ozone, water vapour, cloud cover, cloud liquid water
 192 and sea-level air pressure, as well as by a satellite product for aerosol op-
 193 tical thickness (MODerate resolution Imaging Spectroradiometer, MODIS,
 194 <https://modis.gsfc.nasa.gov/data/dataproduct/>), and also by data for surface
 195 wind speed, air humidity, and air temperature, all provided by the NEMO at-
 196 mospheric (ERA5) forcing. The attenuation of the irradiance was described
 197 in detail by [31], here it is briefly summarized: The module distinguishes
 198 between the absorption and backscattering by the sea water and the 4 PFTs
 199 based on the model of [58]. The scheme for the underwater irradiance was
 200 based on [33], i.e. the irradiance was resolved at 33 wavelengths in the
 201 250 - 3700nm range, and so were the wavelength-dependent absorption and
 202 backscattering coefficients for clear water and PFTs. Although we included
 203 the impact of backscattering on the light attenuation, similarly to [31], we
 204 did not explicitly track the upwelling stream. Besides the clear sea water
 205 and PFTs, we included into the light attenuation also the absorption by
 206 POM, CDOM and sediment, which was (the same as in [31]) forced by an
 207 external product extrapolated from the 443nm data of [59]. The bio-optical
 208 module was extensively validated in [31], and was shown to be skilled in its
 209 representation of SWR, PAR and the underwater irradiances.

210 2.3. Observations: assimilated and validation data

211 2.3.1. Assimilated data

212 In the physical data assimilation component we have included: a) sea
213 surface temperature data from the GCOM-W1/AMSR-2, NOAA/AVHRR,
214 MetOp/AVHRR, MSG/SEVIRI, Sentinel-3/SLSTR, Suomi-NPP/VIIRS satel-
215 lite products and in situ SST observations from ships, surface drifters and
216 moorings, distributed over the Global Telecommunication System (GTS) in
217 near-real time, b) temperature and salinity from the EN4 dataset ([60]),
218 which includes in situ profiles from Argo floats, fixed moored arrays, XBTs,
219 CTDs, gliders, marine mammals, and c) temperature and salinity data from
220 a specific Slocum glider Cabot (Unit 345, see [36]) that has been deployed
221 in the central North Sea during 08/05/2018 - 15/08/2018 as a part of the
222 Alternative Framework to Assess Marine Ecosystem Functioning in Shelf
223 Seas (AlterECO) programme (<https://altereco.ac.uk/>). The satellite SST
224 was bias-corrected following the scheme from [61], using the VIIRS and in
225 situ SST data as the reference.

226 In the biogeochemical data assimilation we have included total log-chlorophyll
227 derived from the ocean color based satellite product of ESA (version 2.0, [62])
228 and also log-chlorophyll derived from the fluorescence measurements by the
229 same AlterEco glider Cabot, that was used in the physical data assimilation.
230 The assimilation is performed for log-chlorophyll, rather than chlorophyll, as
231 chlorophyll is widely known to be log-normally distributed ([63]).

232 The assimilated in situ (EN4, glider) observations were thinned to a res-
233 olution of 0.08° (EN4), or up-scaled to the AMM7 grid (glider), with addi-
234 tional temporal averaging applied to the same-day glider observations. The
235 thinning/up-scaling is performed to avoid assimilating many observations
236 at higher resolution than the model can represent. After the thinning/up-
237 scaling there were $O(10^5)$ EN4 and $O(10^4)$ Cabot glider data-points to as-
238 simulate throughout the year 2018.

239 2.3.2. Validation data

240 The assimilated observations were used for the validation of those exper-
241 iments in which they were excluded from the assimilation (e.g. chlorophyll
242 data for the physical data assimilative run). However, we excluded the bias-
243 corrected satellite SST from the temperature validation, so that the only
244 assimilated SST data used for validation were a) the high quality SST data
245 from the VIIRS satellite product and from ships, drifters and moorings (we
246 will call this “VIIRS/in situ SST data”), and the SST that was part of b) EN4

247 and c) Cabot glider data. Besides the assimilated observations, all the exper-
 248 iments were validated with other (non-assimilated) AlterEco glider data for
 249 temperature, salinity, chlorophyll, oxygen and the sum of nitrate and nitrite
 250 (all the gliders included in the validation are listed in Tab.1). The processing
 251 of the physical, chlorophyll and oxygen data was described in [36]. The sum
 252 of nitrate and nitrite concentrations (abbreviated as $\text{NO}_x^- = \text{NO}_3^- + \text{NO}_2^-$)
 253 were determined using a Lab-on-Chip (LoC) analyser designed and fabricated
 254 at the National Oceanography Centre ([64]), which were implemented by the
 255 AlterEco team into Seagliders following a similar protocol as used by [65].
 256 The combined uncertainty (random and systematic errors) of measurements
 257 made using these LoC analysers has been calculated as 5% (coverage interval
 258 $k = 1$) ([66]). The nitrite concentrations were relatively negligible compared
 259 to the nitrate concentrations, so the NO_x^- data were used to validate model
 260 nitrate outputs. All of the used AlterEco gliders operated during 2018 in
 261 the central North Sea (for both the glider and the EN4 data locations see
 262 Fig.S1 of the Supporting Information (SI)), moving throughout the whole
 263 water column. Similar to the assimilated Cabot glider, the remaining glider
 264 data were up-scaled onto the model grid (on a daily basis) and after the up-
 265 scaling there remained $\text{O}(10^4)$ AlterEco glider observations for each variable
 266 in 2018.

267 The EN4 data-set contained subsurface observations that were approxi-
 268 mately homogeneously distributed both with depth and in time, with slightly
 269 lower number of observations towards the end of the year (November-December
 270 2018). Beyond the assimilated data and the AlterEco data, we used for vali-
 271 dation a 1960-2014 monthly climatological dataset for total chlorophyll, oxy-
 272 gen, nitrate, phosphate and silicate concentrations, compiled in the North
 273 Sea Biogeochemical Climatology (NSBC) project ([67]). The NSBC dataset
 274 covers most of the NWE Shelf and the full range of depths. Finally, we also
 275 included validation of surface CO_2 fugacity using 2018 SOCAT (v2019) data
 276 (<https://www.socat.info/index.php/about/>).

277 *2.4. The assimilative system: NEMOVAR*

278 NEMOVAR is a variational (in this study a 3DVar) DA system ([68, 69,
 279 70]) used at the Met Office for operational reanalyses and forecasting on
 280 the NWE Shelf. The assimilation of ocean color-derived chlorophyll using
 281 NEMOVAR is highly successful in improving the NWE Shelf phytoplankton
 282 phenology, PFT community structure (using PFT chlorophyll assimilation),
 283 underwater irradiance and to a more limited degree also carbon cycle ([46, 31,

Table 1: The AlterEco gliders and the variables measured by the gliders used for assimilation (6-th column), or validation (7-th column). The table uses the following abbreviations: deployment: “dpl”, data assimilation: “DA”, temperature: “T”, salinity: “S”, oxygen concentrations: “O₂”, chlorophyll *a* concentrations: “Chl *a*” and sum of nitrate and nitrite concentrations: “NO_x-”.

Campaign	platform	dpl	serial	mission period	DA	validation
AlterEco 1	Stella	440	unit_436	02/02/2018 - 08/05/2018	none	T,S,O ₂ ,Chl <i>a</i>
AlterEco 1	Cook	441	unit_194	15/11/2017 - 07/02/2018	none	T,S,O ₂ ,Chl <i>a</i> ,NO _x -
AlterEco 2	Orca	493	SG510	07/03/2018 - 27/03/2018	none	Chl <i>a</i> ,NO _x -
AlterEco 2	Melonhead	496	SG620	07/02/2018 - 02/04/2018	none	Chl <i>a</i>
AlterEco 3	Cabot	454	unit_345	08/05/2018 - 15/08/2018	T,S,Chl <i>a</i>	T,S,O ₂ ,Chl <i>a</i>
AlterEco 3	Orca	455	SG510	16/03/2018 - 24/07/2018	none	Chl <i>a</i> ,NO _x -
AlterEco 3	Humpback	497	SG579	09/05/2018 - 25/06/2018	none	Chl <i>a</i>
AlterEco 4	Dolomite	477	unit_305	13/08/2018 - 10/10/2018	none	T,S,Chl <i>a</i> ,NO _x -
AlterEco 4	Eltanin	478	SG550	15/08/2018 - 28/09/2018	none	Chl <i>a</i>
Altereco 5	Kelvin	481	unit_444	26/09/2018 - 02/12/2018	none	T,S,Chl <i>a</i>
AlterEco 6	Dolomite	499	unit_305	02/12/2018 - 12/03/2018	none	T,S,O ₂ ,Chl <i>a</i>
AlterEco 6	Coprolite	500	unit_331	02/12/2018 - 12/03/2018	none	T,S,O ₂ ,Chl <i>a</i>

71]). NEMOVAR includes capability to assimilate multi-platform (satellite, in situ) data, which has been established first for physics (e.g. [70, 72]) and subsequently for biogeochemistry ([73]), including validating the multi-platform DA system for the NWE Shelf ([36]).

The NEMOVAR set-up used in this study for the multi-platform physical-biogeochemical assimilation is the same as the one described in detail by [36]. Here we offer only a short summary: The 3DVar version of NEMOVAR uses a First Guess at Appropriate Time (FGAT) to calculate a daily set of increments for the directly updated variables ([70, 72]). In the physical DA application NEMOVAR applies balancing relationships within the assimilation step and delivers a set of increments for temperature, salinity, sea surface height (SSH) and the horizontal velocity components. For the total chlorophyll assimilation NEMOVAR calculates a set of log-chlorophyll increments and then a balancing scheme is used to distribute those increments into the PFT components (chlorophyll, carbon, nitrogen, phosphorus and for diatoms also silicon), all of which are being updated based on the background community structure and stoichiometric ratios (e.g. [46, 31, 36]). After the assimilation step, the model is re-run with the increments applied to the model variables gradually at each model time-step using incremental analysis updates (IAU, [74]).

304 NEMOVAR uses externally supplied spatio-temporally varying observa-
 305 tion and background error variances, with the background error variances
 306 typically 1-3 times larger than the observational error variances ([36]). The
 307 system combines two horizontal correlation length-scales, one fixed 100km
 308 length-scale with another length-scale based on the baroclinic Rossby radius
 309 of deformation ([72]). The vertical length-scales follow the scheme from [72],
 310 where NEMOVAR calculates directly the set of 3D increments using flow-
 311 dependent vertical length-scales (ℓ), which are at the surface equal to half
 312 of the MLD, decreasing in the mixed layer to become two-times the vertical
 313 model grid spacing at, and beneath the MLD.

314 2.5. The experiments

315 In this study we compared the performance of both one-way and two-way
 316 coupled versions of the NEMO-FABM-ERSEM model. We also tested the
 317 impact of assimilating different types of data (physical-only, biogeochemical-
 318 only and physical and biogeochemical jointly) on the skill of both the one-
 319 way and two-way coupled models. The various experiments used exactly the
 320 same model configuration, apart from the difference in the coupling between
 321 physics and biogeochemistry. The experiments all started from the same
 322 initial value conditions on the 01/09/2017 to allow a 4 month spin-up time for
 323 the final 2018 simulation. The initial values were provided by the 2016-2018
 324 free simulation (using bio-optical module) from the study of [31]. Finally,
 325 Tab.2 provides a list of all experiments with their abbreviated names that
 326 we will use in the paper.

327 2.6. Skill metrics

328 The performance of the different simulations will be evaluated using two
 329 skill metrics. The first metric is the model bias (ΔQ_{mo}):

$$\Delta Q_{mo} = \langle Q_m - Q_o \rangle \quad (1)$$

330 where Q_o are the observations mapped into the model grid and the Q_m are
 331 the corresponding model outputs. The second metric is the bias-corrected
 332 root mean square difference (BC RMSD, $\Delta_{RD}Q_{mo}$):

$$\Delta_{RD}Q_{mo} = \sqrt{\langle (Q_m - Q_o - \Delta Q_{mo})^2 \rangle}. \quad (2)$$

Table 2: The different experiments compared in this study. The first column shows the abbreviated experiment name, the second column indicates whether the two-way coupling is used and the following columns list the assimilated data. The table uses the following abbreviations: satellite: “sat”, Cabot glider: “Cabot”, EN4 dataset: “EN4”, temperature: “T”, sea surface temperature: “SST”, salinity: “S”, chlorophyll *a*: “Chl *a*”.

abbreviation	two-way	SST (sat./in situ)	T & S (EN4)	T & S (Cabot)	Chl <i>a</i> (sat.)	Chl <i>a</i> (Cabot)
free 1-way	no	no	no	no	no	no
free 2-way	yes	no	no	no	no	no
phys DA 1-way	no	yes	yes	yes	no	no
phys DA 2-way	yes	yes	yes	yes	no	no
chl DA 1-way	no	no	no	no	yes	yes
chl DA 2-way	yes	no	no	no	yes	yes
phys+chl DA 1-way	no	yes	yes	yes	yes	yes
phys+chl DA 2-way	yes	yes	yes	yes	yes	yes

3. Results and Discussion

3.1. The impact of the two-way coupling and assimilation on the simulated physics

The reference one-way coupled model simulates well the seasonal increase of temperature in the surface ocean in late-spring / summer (Fig.2:A, Fig.3). The novel two-way coupling further increased the temperature in the upper 20m by around 1°C (Fig.2:B, Fig.3). This is a relatively major change with respect to the reference run, when compared to the changes introduced to the simulated temperature by the physical data assimilation during the same period of the year (Fig.2:D, for all the assimilative runs see Fig.S2-S3 in the Supporting Information (SI)). The increase in the upper ocean temperature in the two-way coupled model cannot be explained by the enhanced shortwave radiation flux in the water column, since the bio-optical module and the ERA5 short-wave radiation product, which forces the one-way coupled run, have a negligible mutual bias ([31]). Therefore, the temperature increase is likely a consequence of an increased rate of absorption inside the upper oceanic layer. The increased absorption in the two-way coupled run was anticipated since: a) the bio-optical module appears to have higher level of light attenuation near the water surface than the satellite observations used to force the physics in the one-way coupled run (this was observed for 490nm wavelength in Fig.5:A of [31]), b) the “broadband” visible light attenuation in the one-way coupled run was represented by the satellite K_d for 490nm

355 wavelength, but K_d at 490nm wavelength is clearly an underestimate of the
356 K_d for the 400-700nm waveband (see Fig.5:B of [31]).

357 The impact of phytoplankton biomass on the simulated temperature can
358 be analysed by comparing the chlorophyll-assimilative run (chl DA 2-way)
359 with its corresponding two-way coupled free run (free 2-way): In the late
360 spring - summer, the assimilation of chlorophyll into the two-way coupled
361 model removes a large amount of phytoplankton biomass from the mixed
362 layer (see Fig.S4:B of SI), increases the light penetration into the water col-
363 umn and heats up a deeper oceanic layer than the free run (Fig.2:B-C). The
364 temperature is then raised in the 20-60m depth range by 0.1-0.2°C in the
365 summer and by less than that in the late spring (see Fig.S5 of SI). The extra
366 heat captured by the two-way coupled model near the ocean surface shallows
367 the MLD (Fig.4:B, Fig.S6 of SI), which is indicative of important changes to
368 mixing of biogeochemical tracers in the upper ocean.

369 Outside of the late spring - summer, both two-way coupling (Fig.2:B) and
370 chlorophyll assimilation (Fig.2:C) have comparably smaller impact on the
371 simulated oceanic temperature than the physical data assimilation (Fig.2:D,
372 see also Fig.S2-S4 of SI). The impact of physical data assimilation is most
373 important around the winter, when it corrects a negative temperature bias
374 ($\sim -0.5^\circ\text{C}$) of the physical model (Fig.2-3, Fig.S3-S4 of SI). The physical
375 data assimilation influences the simulated temperature more evenly across
376 the water column than the bio-optical module (Fig.2), which is likely a com-
377 bination of model dynamical response to the temperature increments in the
378 mixed layer and some assimilated sub-surface data (EN4 and Cabot glider).
379 If the reanalysis state is sufficiently stable with respect to the model dynam-
380 ics, it is known ([46, 31, 36]) that, within NEMOVAR on the NWE Shelf,
381 the assimilated variables in the reanalysis tend to converge to the assimilated
382 data. This is evident in the Fig.5:D, Fig.S3,S7 of SI, comparing the
383 SST of the physical data assimilation runs with the assimilated satellite SST
384 observations.

385 We evaluated (Fig.6 and Fig.7) the skill of both the two-way coupled
386 model and the different assimilative experiments to represent temperature
387 and salinity on the NWE Shelf. Fig.6 compares the two-way coupled free and
388 chlorophyll-assimilative runs with the temperature and salinity measured by
389 the Cabot glider mission in the central North Sea during late spring - summer
390 of 2018 (for more details about the mission see [36], Fig.S1 of SI and Tab.1).
391 Glider-observed temperature is warmer in the upper 30-40m of the water col-
392 umn than the temperature simulated by the one-way coupled model, whereas

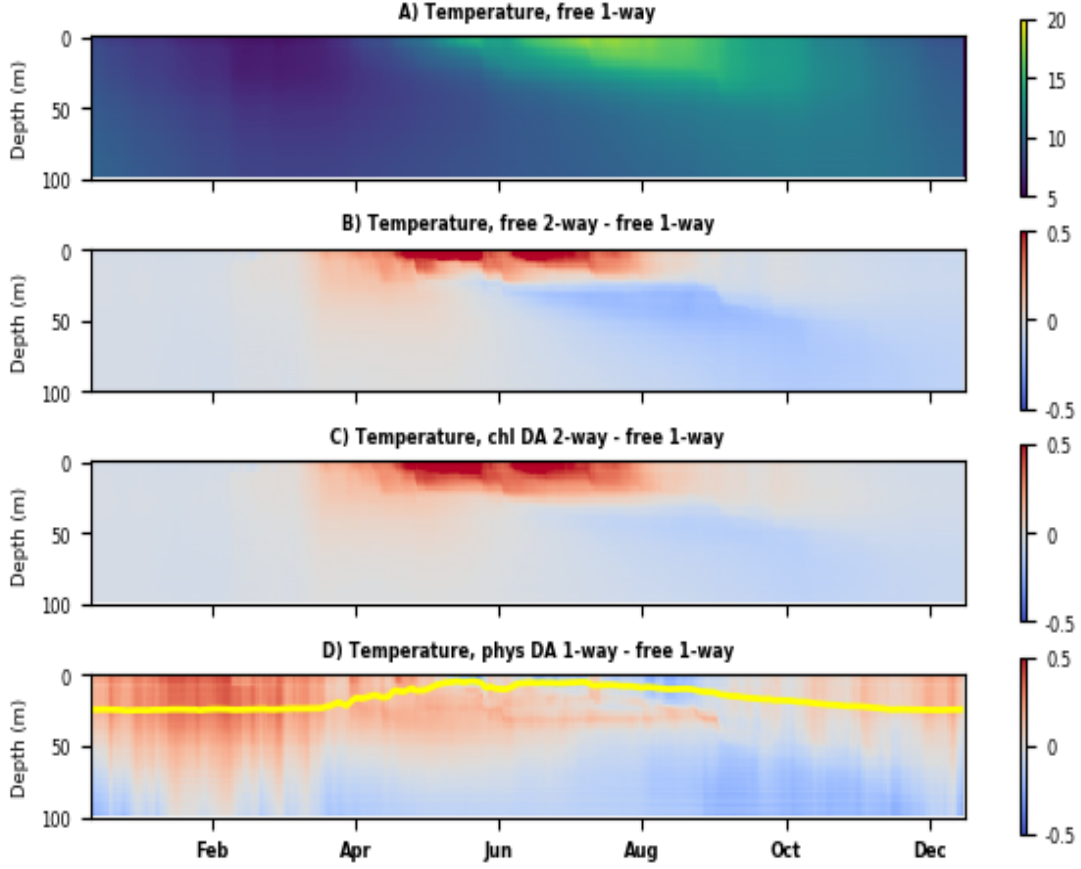


Figure 2: Panel A shows Hovmöller diagram (time on the x-axis vs depth on the y-axis) for the temperature ($^{\circ}\text{C}$) of the one-way coupled free run, where the values for each day and depth represent the horizontal spatial averages throughout the NWE Shelf (bathymetry $< 200\text{m}$). Panels B-D show the same Hovmöller diagrams as panel A, but for the temperature differences between the two-way coupled, or assimilative runs and the reference, free one-way coupled model run from the panel A (for the abbreviations used in the titles see Tab.2). In particular, panels B-D compare the impact of two-way coupling on the simulated temperature (panel B), joint impact of chlorophyll-assimilation and two-way coupling on the simulated temperature (panel C) and the impact of physical data assimilation on the simulated temperature (panel D). The yellow line in the panel D shows the MLD of the physical data assimilative run to indicate the vertical scale of impact of the SST assimilation.

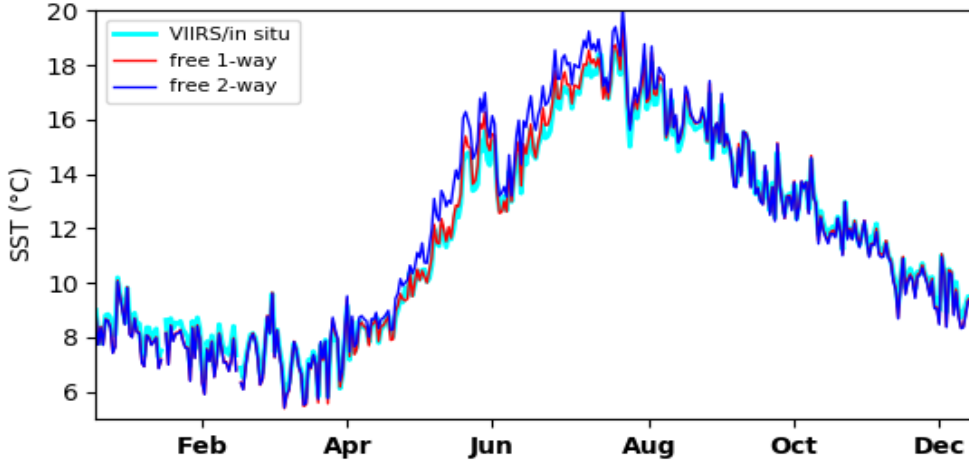


Figure 3: The 2018 time-series of SST averaged throughout the NWE Shelf compared between the one-way and two-way coupled free simulations, and the VIIRS satellite/in situ data. To consistently compare the model simulations with the observed SST, the model outputs were masked wherever there were missing observations. The missing satellite observations are due to the movements of clouds and atmospheric disturbances and the missing values are responsible for the small time-scale fluctuations in the different curves shown in the plot.

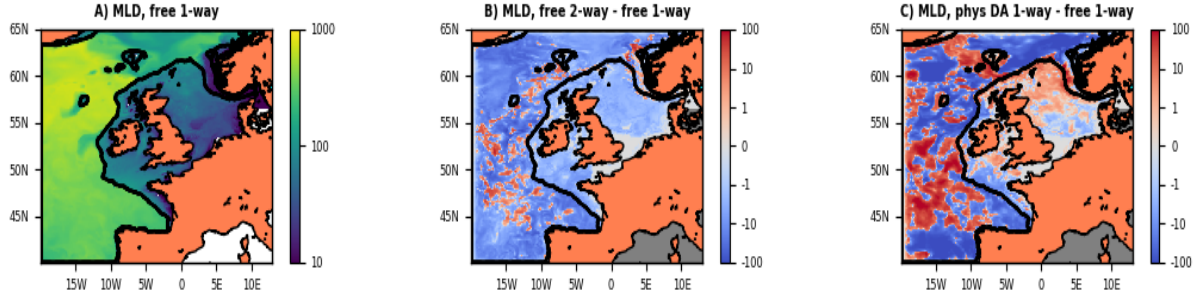


Figure 4: Panel A shows the mixed layer depth (MLD, in m) of the one-way coupled reference free run. The MLD values are averaged for the spring bloom period between March-May 2018. Panels B-C show the relative changes in MLD carried by the two-way coupled free run (panel B) and physical data assimilation into the one-way coupled model (panel C). Both panels B,C show the difference (in m) between the MLD of the two-way coupled, or physical data assimilative run and the one-way coupled model free run (panel A). The blue color in panels B-C (negative values) indicates shallowing of MLD, whilst the red color (positive values) indicates deepening of MLD. The black line shows the boundary of the continental shelf (bathymetry < 200m).

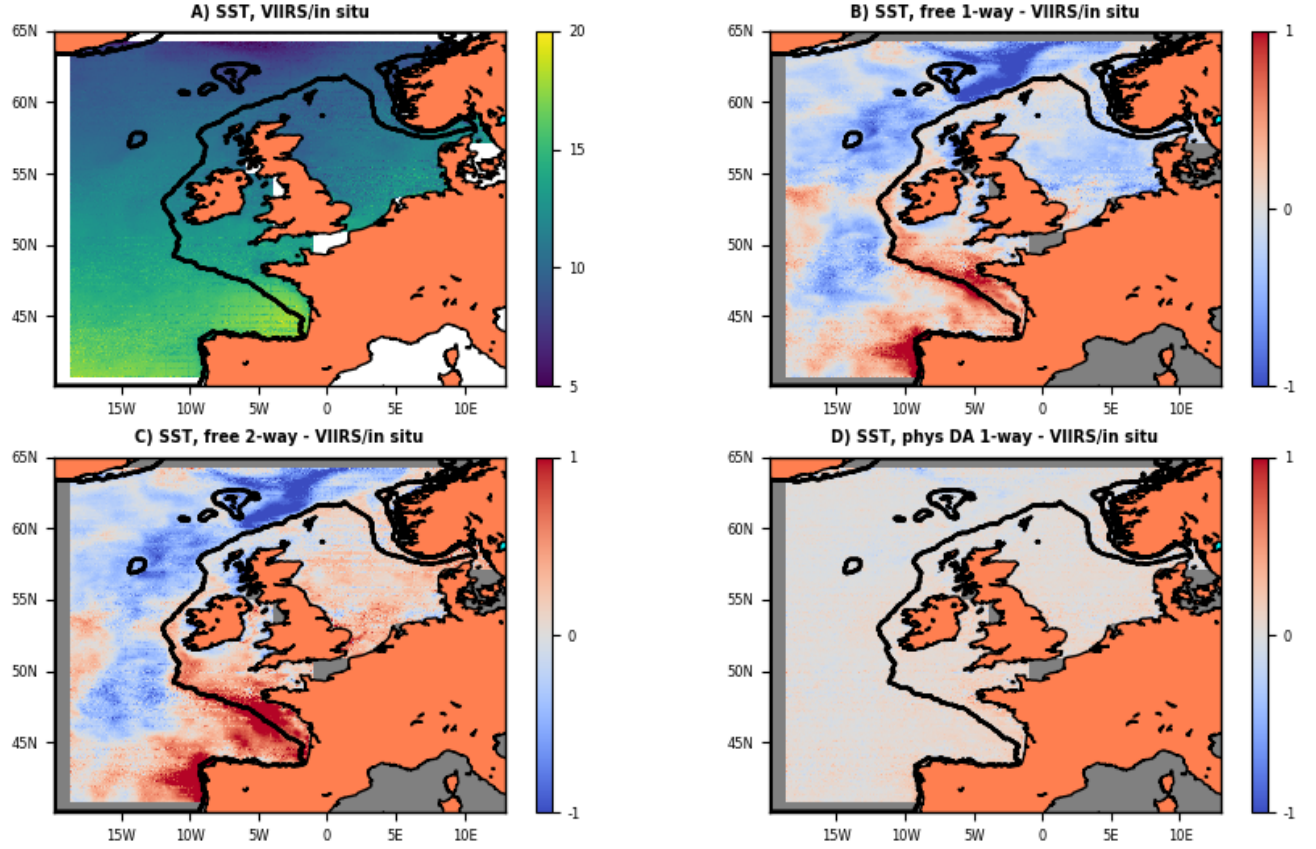


Figure 5: The assimilated 2018 median satellite data for SST (panel A, in $^{\circ}\text{C}$) and the corresponding model to VIIRS/in situ SST differences (panels B-D, in $^{\circ}\text{C}$) for one-way coupled model free run (panel B), two-way coupled model free run (panel C) and physical data assimilation into the one-way coupled free run (panel D). The masked values indicate the regions where there was no assimilation of VIIRS/in situ data into the model.

393 the opposite is true beneath 40m depth (Fig.6:A). This means the observed
 394 thermocline represents a larger gradient in temperature than the simulated
 395 thermocline. The bio-optical module substantially (by $> 1^{\circ}\text{C}$) heats up the
 396 upper 20-30m layer, increasing the vertical temperature gradient (Fig.6:C),
 397 however the near-surface temperature of the two-way coupled run rises well
 398 above the levels observed by the glider (Fig.6:D). The thermocline of the
 399 two-way coupled model free run appears to be located above the glider ther-
 400 mocline (e.g. Fig.6:D) and the impact of the two-way coupling on the model
 401 skill in representing glider temperature is somewhat mixed (it improves bias,
 402 but degrades BC RMSD, Fig.7:A). The skill validation presented in Fig.7
 403 shows similarly mixed results: the summer temperature bias is improved
 404 across the EN4 and AlterEco glider data, but degraded relative to the VI-
 405 IRS/in situ data (see also Fig.3), with the BC RMSD consistently degraded
 406 across the different validation data. The Fig.7:A indicates that the two-
 407 way coupling produces better results for sub-surface summer temperature,
 408 than for SST (VIIRS/in situ data). The two-way coupling has a similarly
 409 mixed impact on the free run skill to represent summer salinity (Fig.7:C),
 410 and both small ($< 0.05^{\circ}\text{C}$) and mixed impact on winter temperature and
 411 salinity (Fig.7:B,D, for temperature see also Fig.2-3). However, it should be
 412 noted that chlorophyll assimilation into the two-way coupled model slightly
 413 improves the skill of the free run in representing temperature and salinity
 414 across most of the data and throughout the whole year 2018 (Fig.7). Finally,
 415 the comparison with the non-assimilated temperature validation data clearly
 416 demonstrates that the physical data assimilation improves the model skill in
 417 temperature both in summer and winter half-year (Fig.7:A-B) and also the
 418 model skill in salinity in the winter half-year (Fig.7:D).

419 3.2. *The impact of the two-way coupling and assimilation on biogeochemistry*

420 As the days in spring become longer, the layer that is effectively lit by
 421 the sunlight expands into the water column, whilst the effective mixing depth
 422 shrinks. It is often assumed, that the effective mixing depth reaching a critical
 423 threshold marks the onset of the spring bloom (Fig.1). This process might be
 424 misrepresented by the one-way coupled reference free simulation, which could
 425 be why the model shows on the NWE Shelf late (by ~ 1 month) and intense
 426 blooms (Fig.8, see also [31, 36]). The effective mixing depth has often been
 427 interpreted as the seasonal MLD (this is the frequent understanding of the
 428 critical depth hypothesis of [75]), but it is assumed that on the NWE Shelf
 429 the onset of the bloom might be better described by the critical turbulence

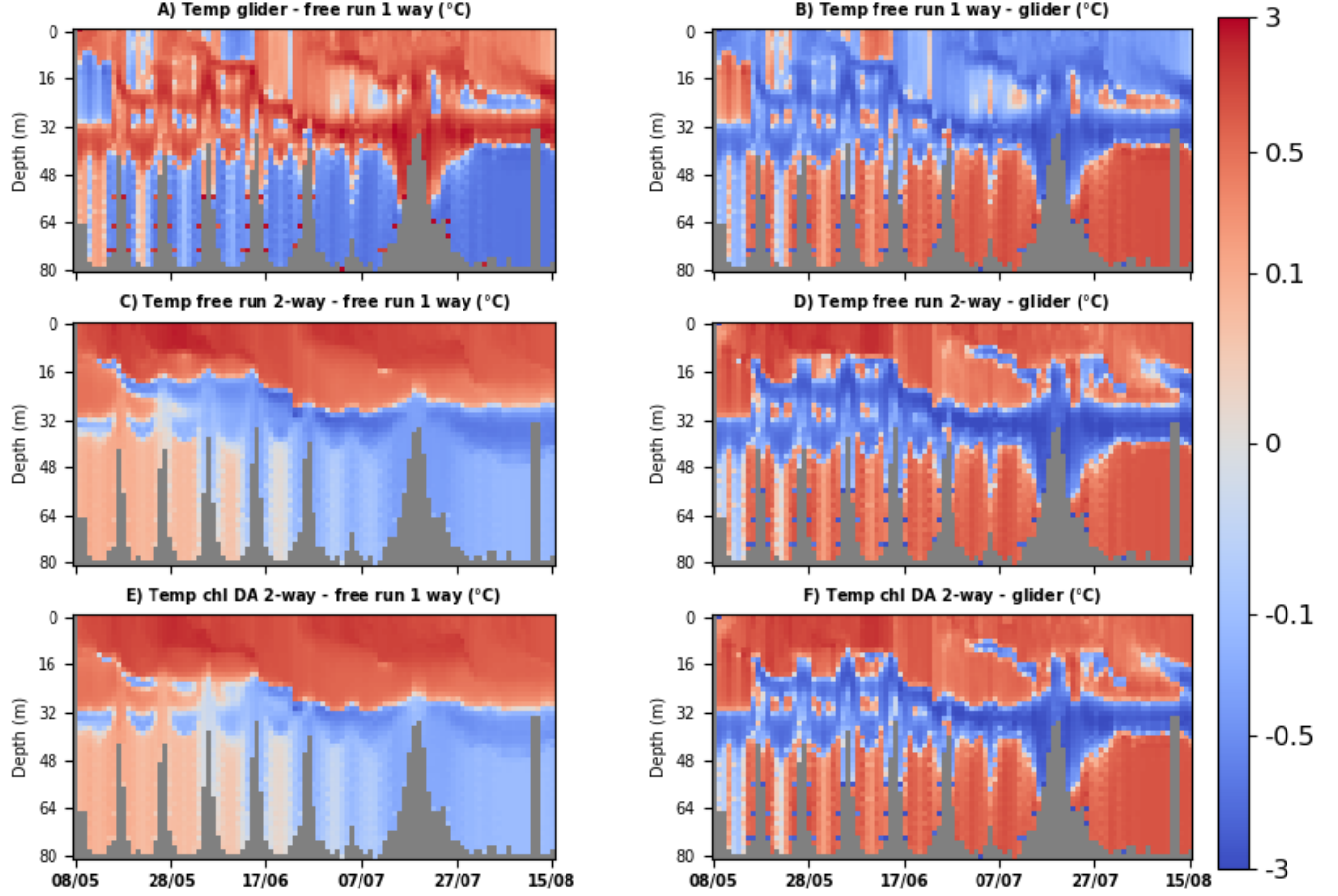


Figure 6: Hovmöller diagram for temperature ($^{\circ}\text{C}$) along the trajectory covered by the Cabot glider in the central North Sea during an early May to mid-August 2018 mission. The right-hand panels (B,D,F) show the temperature differences between the free one-way coupled run (panel B), free two-way coupled run (panel D), the chlorophyll assimilation into the two-way coupled model (panel F) and the Cabot glider observations (model minus glider). The left hand panels (A,C,E) show the differences between the observations, or model simulations and the reference, free one-way coupled model run. The purpose of the left-hand panels is to show the desired changes to the one-way coupled model (panel A) and how these changes are realized by the biogeochemical feedback in the free run (panel C) and in the chlorophyll-assimilative run (panel E). The main advantage of those left-hand (A,C,E) panels is that they allow relatively easy interpretation of the dynamical changes introduced to the reference run by the biogeochemical feedback to physics and/or data assimilation.

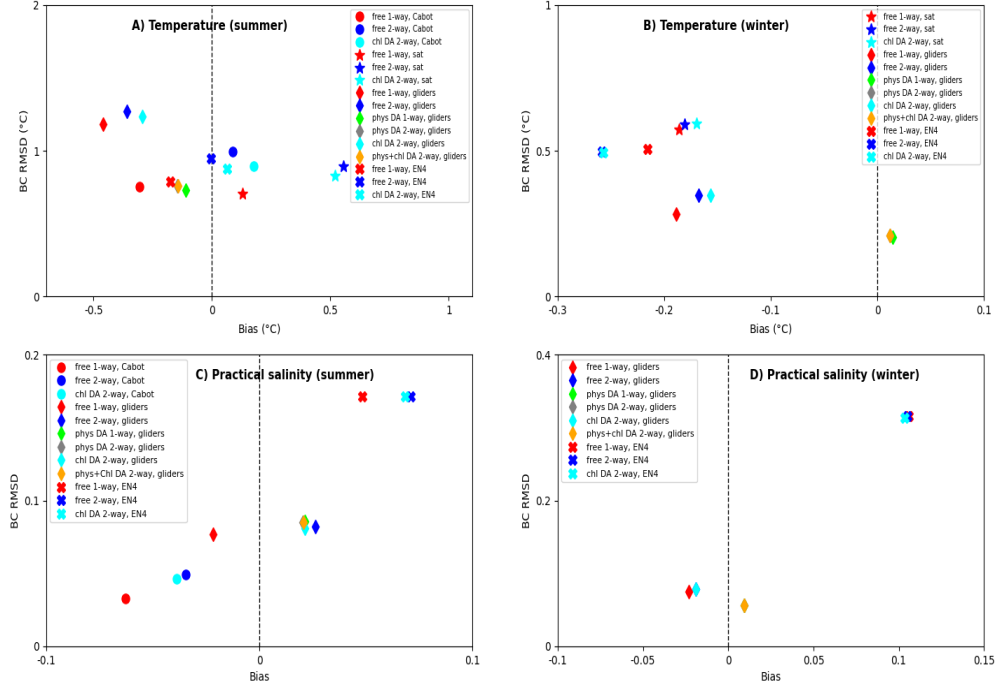


Figure 7: Skill of the different model simulations to represent temperature ($^{\circ}\text{C}$, panels A-B) and practical salinity (panels C-D). The skill is measured by bias (x-axis, Eq.1) and BC RMSD (y-axis, Eq.2). The skill is evaluated for two half-year periods of 2018, the “summer” (panels A,C) defined as May-October and the “winter” (panels B,D) defined as November-April (data averaged through January-April 2018 and November-December 2018). The different simulations are represented by different colors: free run of the one-way coupled model (red), free run of the two-way coupled model (blue), assimilation of chlorophyll into the two-way coupled model (cyan), physical data assimilation into the one-way coupled model (lime), physical data assimilation into the two-way coupled model (grey) and joint physical data-chlorophyll assimilation into the two-way coupled model (orange). The different markers show comparison with different data-sets: the star stands for the VIIRS/in situ SST, the circle for the Cabot glider observations, the diamond for the remaining available glider observations (the 2018 AlterEco mission without Cabot) and the cross for the EN4 data-set. The data (SST, Cabot, EN4) which were assimilated in some of the simulations were used to validate only the simulations that avoided their assimilation.

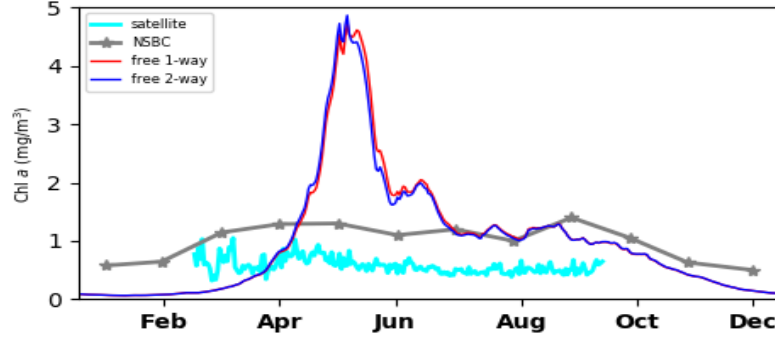


Figure 8: The 2018 time-series of surface chlorophyll a concentrations (mg/m^3) averaged throughout the NWE Shelf compared between the one-way and two-way coupled free simulations, the satellite data, as well as with the NSBC climatological data-set. The satellite data were considered only in the March-September period as the data outside this period are scarce and limited only to the southern part of the NWE domain. The small time-scale fluctuations in the satellite data are due to the missing values caused by the movement of clouds and atmospheric disturbances.

430 hypothesis ([40]). In the critical turbulence hypothesis the bloom starts when
 431 the turbulent mixing in the upper ocean drops beneath a critical level, whilst
 432 the effective rate of turbulent mixing is largely decoupled from the seasonal
 433 MLD ([40, 41, 42]).

434 The implementation of the bio-optical module was shown to shallow the
 435 MLD (Fig.4), but it can also reduce convection within the mixed layer and
 436 the turbulent mixing. The starting hypothesis of this work was that the ex-
 437 tra heat captured in the upper oceanic layer could trigger an earlier bloom
 438 and improve the ERSEM skill. Fig.8, Fig.9:B and Fig.10:C-D show that the
 439 changes to the simulated physics introduced through the two-way coupled
 440 model indeed trigger an earlier phytoplankton bloom, but the difference in
 441 the bloom timing is only on the scale of several days, rather than weeks.
 442 However, the shift to the bloom timing has an impact on many subsequent
 443 features, such as the deep chlorophyll maxima (e.g. [36]), so the changes
 444 to the bloom onset can gradually propagate to the subsurface chlorophyll
 445 (Fig.9:C). The model skill to simulate chlorophyll is improved by the two-
 446 way coupling quite notably in the central North Sea and the period covered
 447 by the Cabot glider (Fig.11:A), however comparisons with other data spread
 448 throughout the year 2018 (satellite ocean color, remaining AlterEco gliders
 449 and the NSBC climatology) show only small improvement (Fig.11:A). The

450 modest improvement to the timing of the (delayed) spring bloom through
 451 the changed mixing is certainly a disappointment, and we suspect that to
 452 introduce a larger correction to the timing of the bloom it would be nec-
 453 essary to either improve the physical model mixing scheme, or to improve
 454 some key ERSEM parameters and processes, such as P-I curves, the max-
 455 imum chlorophyll-to-carbon ratios, zooplankton grazing and representation
 456 of plankton mixotrophy ([29]).

457 Although the (modest) improvements to the simulated chlorophyll by the
 458 two-way coupled model originate from its changes to the simulated physics
 459 (i.e. vertical mixing), the physical data assimilation, which substantially im-
 460 proves the simulated physics (Fig.7) does not improve (even slightly degrades)
 461 the model skill in chlorophyll (Fig.11:A). This is likely because the physical
 462 data assimilation is for large part the assimilation of SST. The improvement
 463 in the ecosystem model skill depends mostly on the vertical mixing and lim-
 464 ited changes to vertical mixing are expected by assimilating SST. Assimilated
 465 subsurface temperature and salinity data are quite sparse, and have only a
 466 limited impact on the modelled biogeochemistry. In the case of the Cabot
 467 glider “case-study” presented in Fig.10 (for a more complete view see Fig.S8
 468 of SI), the glider temperature and salinity assimilation did not improve the
 469 simulated chlorophyll at the glider locations (Fig.11:A) mostly because the
 470 impact of physics on biogeochemistry needs some spin-up time. In fact in
 471 the last part of the glider mission period (late July-August in Fig.10:E) the
 472 physical assimilation has some potential to improve the chlorophyll concen-
 473 trations, as was demonstrated by the assimilation of the same Cabot glider
 474 data in Fig.6E of [36]. Finally, the chlorophyll assimilation dominates over
 475 both physical assimilation and two-way coupling in its impact on the simu-
 476 lated chlorophyll concentrations across the whole water column and the whole
 477 simulation year (Fig.9:D and Fig.S9 of SI). Since the chlorophyll assimi-
 478 lation is almost entirely based on the satellite ocean color, chlorophyll beneath
 479 the mixed layer is updated through the model dynamical response to the
 480 assimilation (e.g. vertical mixing). Similarly to temperature, the chlorophyll
 481 reanalyses look very similar to the assimilated data (Fig.12:B-C, Fig.S5 and
 482 Fig.S10 of SI) and also validate much better than the free runs relative to
 483 the non-assimilated AlterEco glider data (Fig.11:A).

484 We validated the model simulation of additional biogeochemical variables
 485 with available observational data: oxygen, nitrate, phosphate, silicate and
 486 CO₂ fugacity. The oxygen concentrations are mostly driven by the primary
 487 productivity, respiration and outgassing, which largely depends on the sea

488 temperature. The two-way coupled model improves the model skill in repre-
 489 senting Cabot oxygen (Fig.11:B), which is likely triggered by the fact that the
 490 same simulation improves both Cabot chlorophyll (Fig.11:A) and the temper-
 491 ature bias (Fig.7:A). Equivalently, model skill in representing Cabot glider
 492 oxygen can be improved by assimilating physical data into the model (phys
 493 DA 1-way), and it is to some degree also improved by assimilating chloro-
 494 phyll (chl DA 1-way, chl DA 2-way), with the best performance achieved
 495 when both the physical data and chlorophyll are assimilated into the model
 496 (Fig.11:B). However, the Cabot glider study is specific, since the glider mis-
 497 sion took place in the period of the largest discrepancy in the simulated
 498 and observed productivity (Fig.8) and the oxygen concentrations were mea-
 499 sured by the same glider that provided temperature, salinity and chlorophyll
 500 data for assimilation. For the remaining non-assimilated AlterEco gliders
 501 the impact of two-way coupling and assimilation on simulated oxygen is less
 502 clear (Fig.11:B), i.e. even though AlterEco chlorophyll is improved by the
 503 chlorophyll-only assimilative runs (Fig.11:A) they mostly degrade simulated
 504 oxygen (Fig.11:B). This is likely due to the complex relationship between phy-
 505 toplankton chlorophyll and oxygen (see [36]), which includes respiration of
 506 oxygen by the higher trophic-level species (in ERSEM it is zooplankton and
 507 heterotrophic bacteria). However, improved representation of temperature
 508 consistently improves model oxygen bias across all the used data (Fig.11:A),
 509 which indicates that an important part of oxygen bias is due to model biases
 510 in temperature and not due to errors in the simulated biogeochemistry.

511 Besides oxygen, we looked at the model skill in how it represents the
 512 surface CO_2 fugacity, which is influenced by the model skill in simulating
 513 primary productivity and sea temperature (gas solubility). Fig.11:C shows
 514 that CO_2 fugacity is substantially improved by all the runs that included
 515 chlorophyll assimilation, which indicates that the assimilation of chlorophyll
 516 improved the phytoplankton carbon biomass and therefore the simulated
 517 carbon cycle (see also [46]). The physical data-only assimilative runs, and
 518 the two-way coupled free run, had more limited impact on the model skill
 519 to represent surface CO_2 fugacity, but they sometimes reduced the model
 520 bias in CO_2 fugacity. Both the two-way coupling and the physical assimi-
 521 lation, have a relatively small impact on the nitrate and phosphate concen-
 522 trations (Fig.11:D-E), however the changed phytoplankton biomass through
 523 the chlorophyll assimilation lowers the nitrate and phosphate concentrations
 524 at the NSBC data-set locations. This has a positive impact on the nitrate
 525 bias and a negative impact on the phosphate bias (Fig.11:D-E). Silicate is

526 impacted more by the physical data assimilation than nitrate and phosphate,
527 but it is mostly degraded by all the assimilative runs (Fig.11:F).

528 4. Summary

529 In this work we used a recently developed bio-optical module to improve
530 the representation of oceanic heat fluxes and to introduce a biogeochemical
531 feedback to the physical marine model (we call the model with such feedback
532 “a two-way coupled model”). We have estimated the scale of the biogeochem-
533 ical impact on the simulated physics and we have shown that in the upper
534 oceanic layer, in the late spring - summer period, the feedback is comparable
535 to the physical data assimilation in its impact on the simulated tempera-
536 ture. The bio-optical module increases the heat captured in the upper part
537 of the water column, steepens the vertical temperature gradient and shal-
538 lows the mixed layer depth. We have shown that the changes introduced by
539 the bio-optical module into the physical marine model have a mixed impact
540 on the physical model skill. The skill is however (slightly) improved by the
541 chlorophyll assimilation into the two-way coupled model and substantially
542 improved by the physical data assimilation.

543 The increased stratification of the water column and the shallowed mixed
544 layer depth have a modest positive impact on the timing of the late bloom
545 displayed by the biogeochemical model. The shift in the timing of the bloom
546 in the two-way coupled model improves the model skill in representing chloro-
547 phyll. We conclude that, for a more substantial improvement of the timing
548 of the bloom, it will be necessary to either improve the physical model mix-
549 ing scheme, or to improve the process description, or parametrization of the
550 biogeochemical model. We have expanded our analysis to include other bio-
551 geochemical tracers, and we have found that the two-way coupled model and
552 the physical data assimilation may sometimes help improve the agreement of
553 simulated oxygen concentrations and CO_2 fugacity with observations, both
554 due to improved simulation of the sea water temperature (saturation levels)
555 and productivity.

556 This study provides important evidence to support the inclusion of two-
557 way coupling into future operational models of the NWE Shelf. Furthermore,
558 the physical-biogeochemical assimilative runs on the NWE Shelf, including
559 this work, are typically only weakly coupled (for one recent exception see
560 [76]), in the sense that the physical and the biogeochemical variables are
561 updated independently and interact only through the model dynamics. The

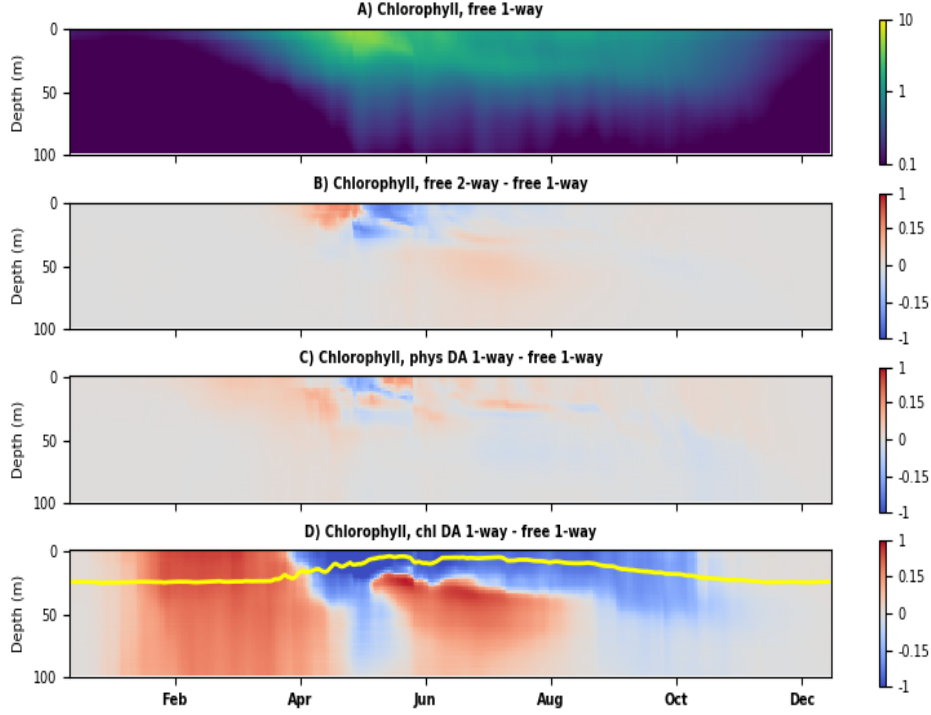


Figure 9: Impact of two-way coupling and assimilation on the simulated chlorophyll concentrations (mg/m^3). Panel A shows Hovmöller diagram (time on the x-axis vs depth on the y-axis) for the one-way coupled model free run, where the values for each day and depth represent the horizontal spatial averages throughout the NWE Shelf (bathymetry $< 200\text{m}$). Panels B-D show the same Hovmöller diagrams, but for the difference between the specific simulation and the reference, free one-way coupled run. The purpose of the panels B-D is to provide an understanding of how the two-way coupling (panel B), the biogeochemical feedback (panel C) and the chlorophyll-assimilation (panel D) influence the chlorophyll concentrations of the reference free one-way coupled run. The yellow line in the panel D shows the mixed layer depth, providing the boundary of the region in which the ocean color assimilation directly updates the simulated chlorophyll.

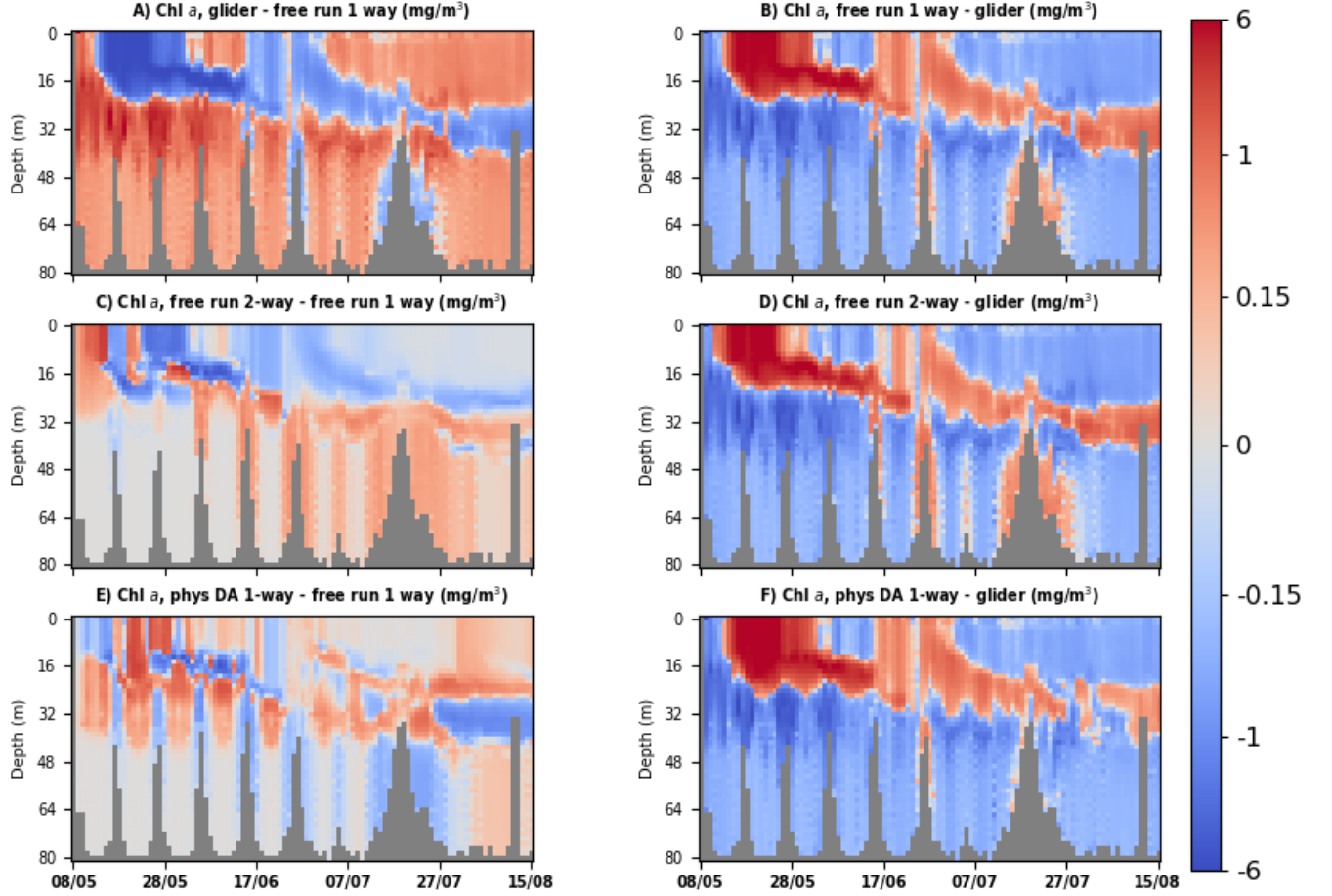


Figure 10: Hovmöller diagram for chlorophyll concentrations (mg/m^3) along the Cabot glider trajectory in the central North Sea during an early May to mid-August 2018 mission. The right-hand panels (B,D,F) show the chlorophyll differences between the free one-way coupled model run (panel B), free two-way coupled model run (panel D), the physical data assimilation into the one-way coupled model (panel F), and the Cabot glider observations (model minus glider). The left hand panels (A,C,E) show the differences between the observations, or model simulations and the reference, free one-way coupled model run. The purpose of the left-hand panels is to show the desired changes to the one-way coupled model (panel A) and how these changes are realized by the biogeochemical feedback in the free run (panel C) and in the physical data-assimilative run (panel E). The main advantage of those left-hand panels is that they allow relatively easy interpretation of the dynamical changes introduced to the reference run by the biogeochemical feedback to physics and/or data assimilation.

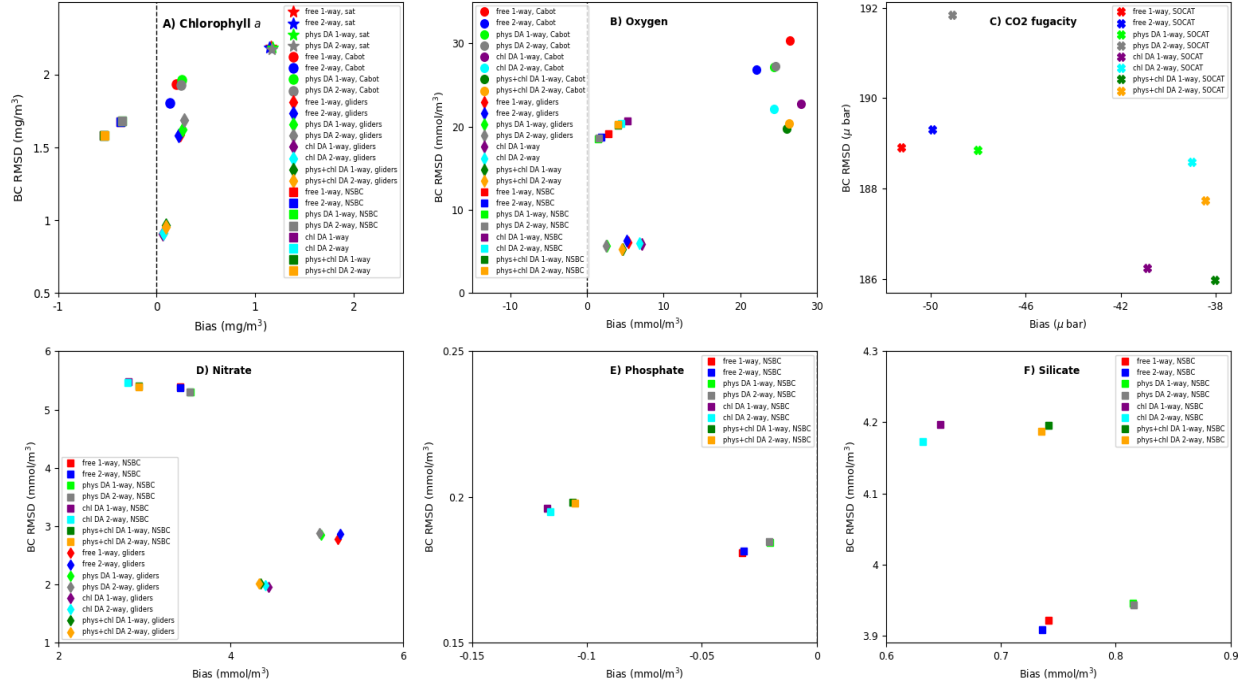


Figure 11: Skill of the different model simulations to represent chlorophyll *a* (mg/m³, panel A), oxygen (mmol/m³, panel B), CO₂ fugacity (μ bar, panel C), nitrate (mmol/m³, panel D), phosphate (mmol/m³, panel E) and silicate (mmol/m³, panel F) concentrations. The skill is measured by bias (x-axis, Eq.1) and BC RMSD (y-axis, Eq.2). The skill is evaluated for the full year 2018. The different simulations are represented by different colors: free run of the one-way coupled model (red), free run of the two-way coupled model (blue), assimilation of chlorophyll into the one-way coupled model (purple), assimilation of chlorophyll into the two-way coupled model (cyan), physical data assimilation into the one-way coupled model (lime), physical data assimilation into the two-way coupled model (grey), joint physical data-chlorophyll assimilation into the one-way coupled model (green) and joint physical data-chlorophyll assimilation into the two-way coupled model (orange). The different markers show comparison with different data-sets: the star stands for the satellite ocean color data, the circle for the Cabot glider observations, the diamond for the remaining available glider observations (the 2018 AlterEco mission without Cabot), the cross for the SOCAT data and the square for the NSBC climatological data-set.

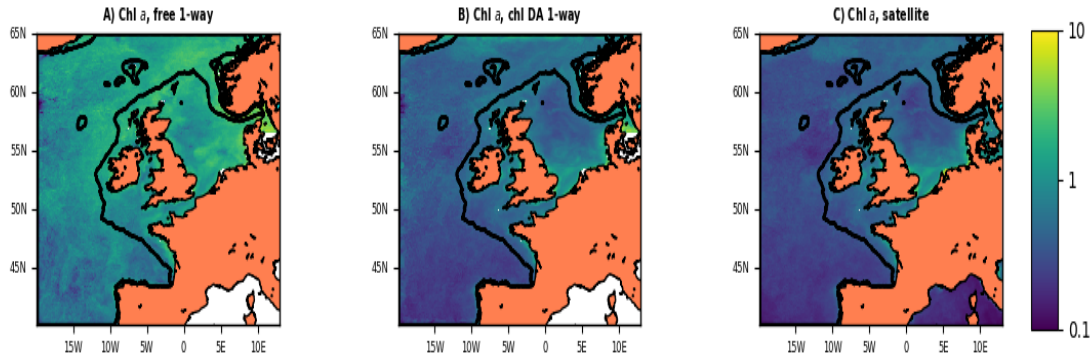


Figure 12: The 2018 mean surface chlorophyll concentrations (in mg/m^3). The different panels compare: the one-way coupled model free run (panel A), the chlorophyll assimilation into the one-way coupled model free run (panel B), and the assimilated satellite ocean color observations (panel C). In the annual averaging we masked the model outputs wherever the satellite data were missing. The black line shows the continental shelf boundary (bathymetry $< 200\text{m}$).

562 interaction between physics and biogeochemistry via the coupled model dy-
 563 namics has been strengthened through the two-way coupling, but it would
 564 be much more efficient if the assimilative updates to the physics and bio-
 565 geochemistry interacted directly through their cross-covariances, or a bal-
 566 ancing component within a data assimilation system. Such scheme is called
 567 “strongly coupled”, and would provide the physical assimilation with both
 568 faster and greater impact on the biogeochemical model skill, and vice versa.
 569 Future work will use the two-way coupled model and expand the data assim-
 570 ilation scheme to include such strong coupling into our operational system.

571 Acknowledgments

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Supporting Information for ”Improved consistency between the modelling of ocean optics, biogeochemistry and physics, and its impact on the North-West European Shelf seas”

Jozef Skákala^{1,2}, Jorn Bruggeman¹, David Ford³, Sarah Wakelin⁴, Anıl Akpınar⁴, Tom Hull^{5,6}, Jan Kaiser⁶, Benjamin R. Loveday⁷, Charlotte A.J. Williams⁴ and Stefano Ciavatta^{1,2}

¹Plymouth Marine Laboratory, The Hoe, Plymouth, PL1 3DH United Kingdom.

²National Centre for Earth Observation, Plymouth, PL1 3DH, UK.

³Met Office, FitzRoy Road, Exeter, EX1 3PB UK.

⁴National Oceanography Centre, Joseph Proudman Building, 6 Brownlow Street, Liverpool, L3 5DA UK.

⁵Centre for Environment, Fisheries and Aquaculture Science, Lowestoft, NR33 0HT UK.

⁶Centre for Ocean and Atmospheric Science, University of East Anglia, Norwich, NR4 7TJ, UK.

⁷Innoflair UG, Richard-Wagner-Weg 35, 64287, Darmstadt, Germany.

Contents of this file

1. Figures S1 to S10

Figures

The panels in the Fig.S1-S10 use the following abbreviations: “free 1-way”: free run of the one-way coupled model, “free 2-way”: free run of the two-way coupled model, “phys DA 1-way”: physical data assimilation into the one-way coupled model, “phys DA 2-way”: physical data assimilation into the two-way coupled model, “chl DA 1-way”: chlorophyll assimilation into the one-way coupled model, “chl DA 2-way”: chlorophyll assimilation

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into the two-way coupled model, “phys+chl DA 1-way”: joint physical data - chlorophyll assimilation into the one-way coupled model, “phys+chl DA 2-way”: joint physical data - chlorophyll assimilation into the two-way coupled model.

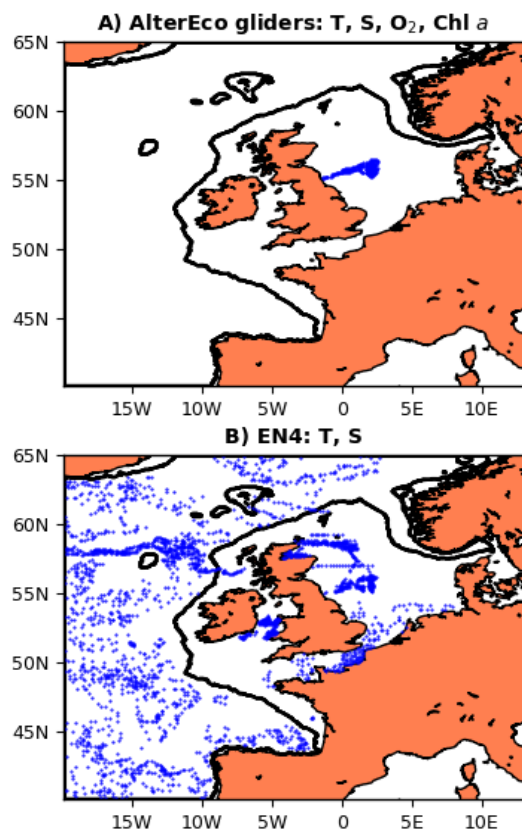


Figure S1. The locations of the 2018 in situ data used both for the assimilation and the validation. The panel A shows the locations of the AlterEco glider measurements and the bottom panel B shows the locations of the EN4 data for temperature and salinity. The EN4 data located outside of the NWE Shelf (bounded by the black line) were used only for assimilation, not for validation.

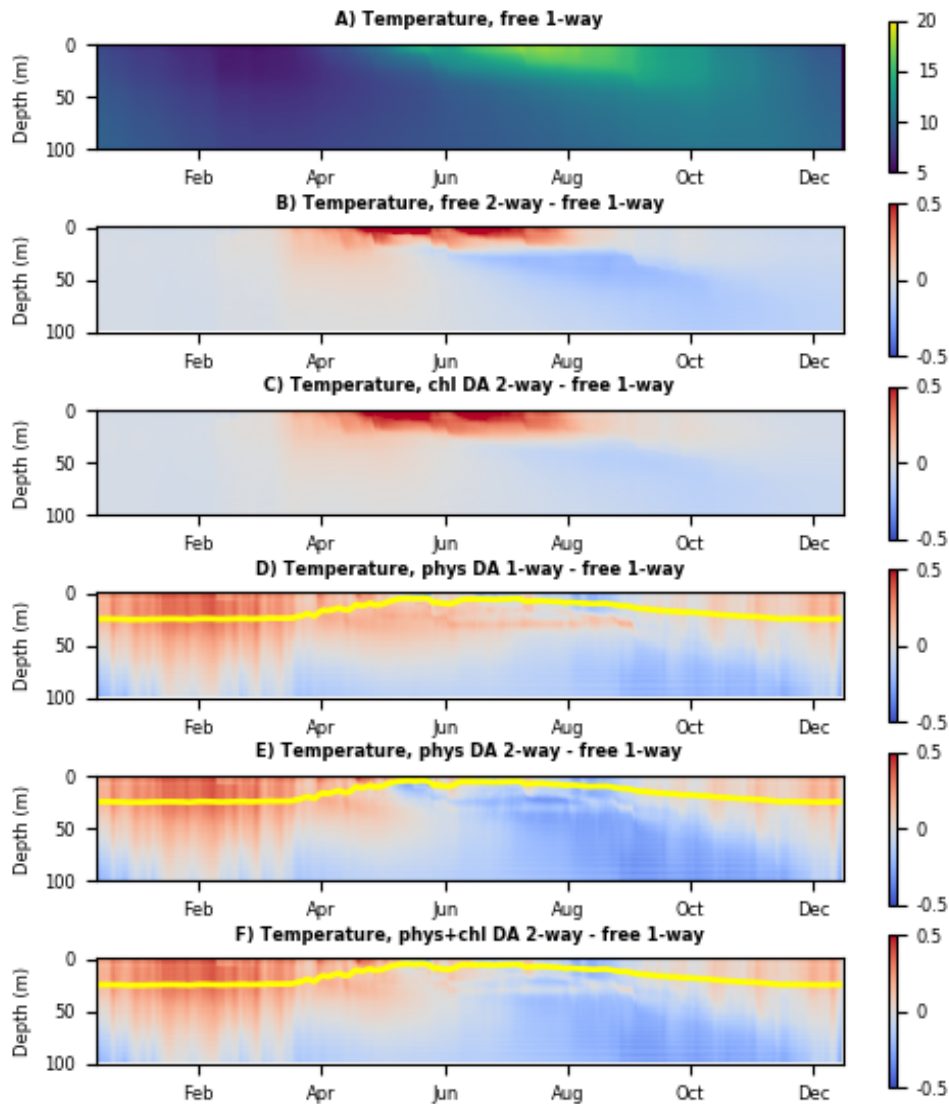


Figure S2. The panel A shows Hovmöller diagram (time on the x-axis vs depth on the y-axis) for the temperature (C) of the one-way coupled free run ("free 1-way"), where the values for each day and depth represent the horizontal spatial averages throughout the NWE Shelf (bathymetry < 200m). Panels B-F show the same Hovmöller diagrams, but for the differences between the two-way coupled, or assimilative runs and the reference, free one-way coupled model run. The purpose of the panels B-F is to provide an understanding of how the bio-optical module and the assimilative model components influence the temperature of the reference free one-way coupled run. The yellow lines in the panels D-F show the MLD of the physical data assimilative runs to indicate the vertical scale of impact of the SST assimilation.

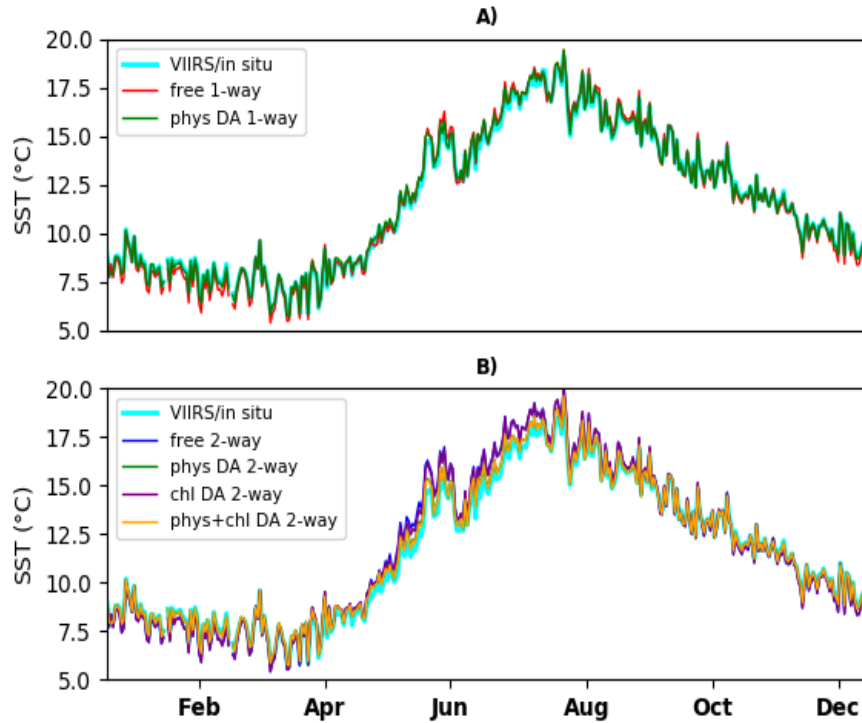


Figure S3. The 2018 time-series of SST averaged throughout the NWE Shelf compared between the different one-way, two-way coupled, free, or assimilative simulations and the VIIRS/in situ data. Panel A compares the different one-way coupled runs, i.e. the one-way coupled free run with the physical data assimilative run, panel B compares the different two-way coupled runs, i.e. the two-way coupled free run with the physical data assimilative run, the chlorophyll assimilative run and the run assimilating both physical data and chlorophyll. To consistently compare the model simulations with the VIIRS/in situ SST, the model outputs were masked wherever there were missing satellite data. The missing satellite data are due to the movements of clouds and atmospheric disturbances and the missing values are responsible for the small time-scale fluctuations in the different curves shown in the three panels. We do not show the one-way coupled runs assimilating chlorophyll, as those have by definition no impact on the simulated temperature.

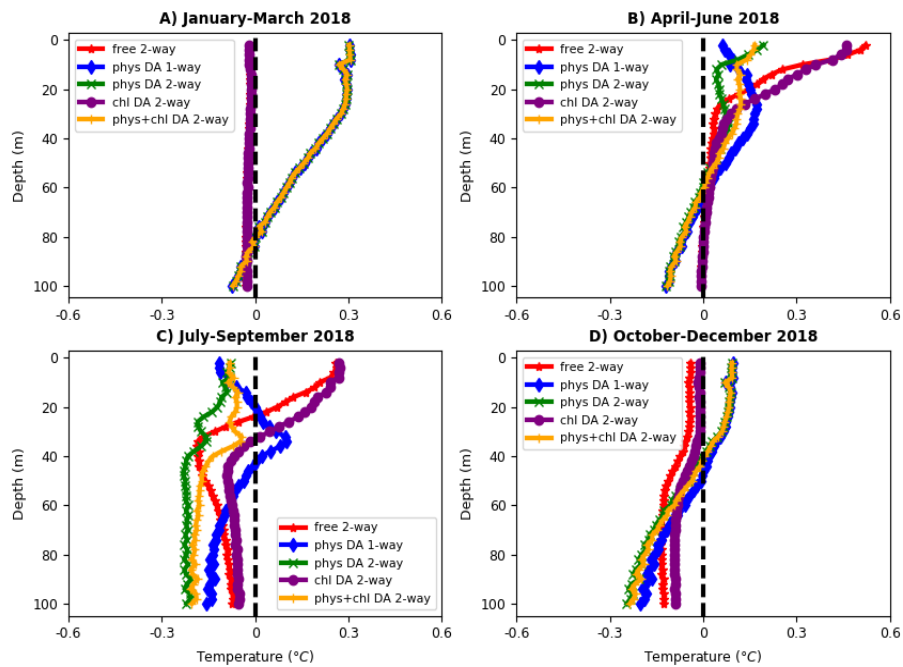


Figure S4. The seasonal differences in temperature (x-axis, °C) between the two-way coupled, or assimilative runs and the reference, one-way coupled free run. The differences are shown as a function of depth (y-axis, m), and averaged throughout the seasonal period and the NWE Shelf.

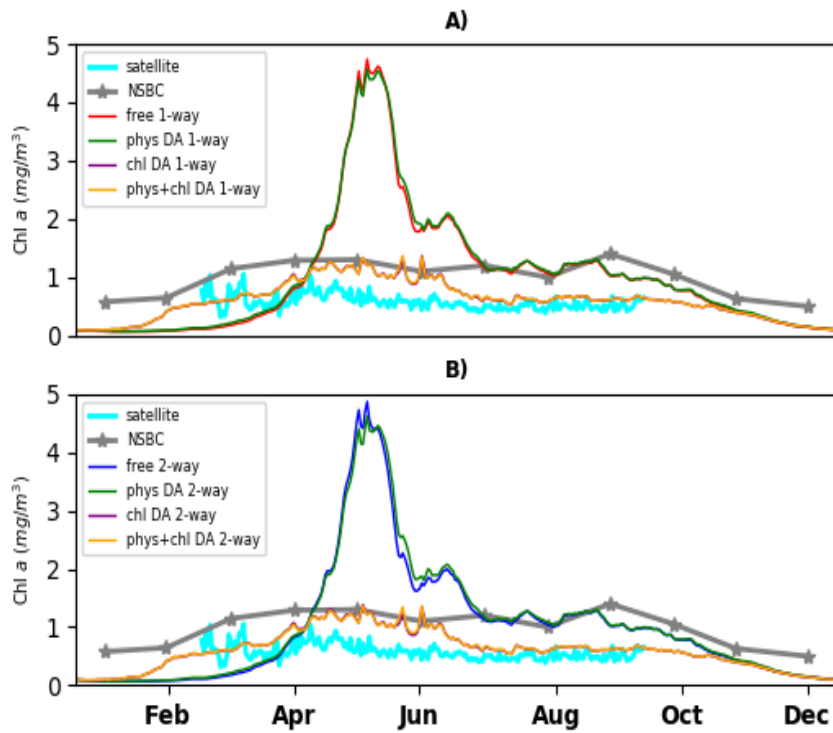


Figure S5. The 2018 time-series of surface chlorophyll *a* concentrations (mg/m^3) averaged throughout the NWE Shelf compared between the different one-way, two-way coupled, free, or assimilative simulations and the satellite data, as well as with the NSBC climatological dataset. Panel A compares the different one-way coupled runs, i.e. the one-way coupled free run with the physical data assimilative run, the chlorophyll assimilative run and the joint physical data-chlorophyll assimilative run, panel B compares the different two-way coupled runs, i.e. the two-way coupled free run with the physical data assimilative run, the chlorophyll assimilative run and the joint physical data-chlorophyll assimilative run. The chlorophyll assimilative run from both panels A and B is hard to see, as the line is nearly identical with the joint physical-chlorophyll assimilative run. The satellite data were considered only in the March-September period as the data outside this period are scarce and limited only to the southern part of the NWE domain. The small time-scale fluctuations in the satellite data are due to the missing values caused by the movement of clouds and atmospheric disturbances.

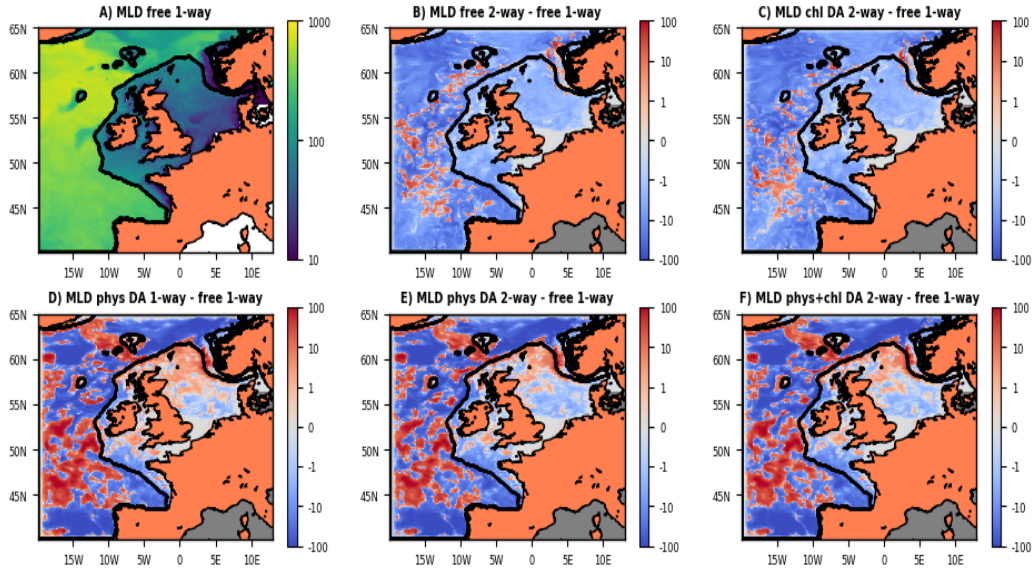


Figure S6. Panel A shows the mixed layer depth (MLD, in m) of the one-way coupled free run (the reference run). The MLD values are averaged for the spring bloom period between March-May 2018. The panels B-F show the relative changes (relative to the one-way coupled free reference run, in m) in MLD carried by the two-way coupled free run (panel B), chlorophyll assimilation into the two-way coupled model (panel C), physical data assimilation into the one-way coupled (panel D) and into the two-way coupled model (panel E) and the joint physical data-chlorophyll assimilation into the two-way coupled model (panel F). All panels B-F show the difference between the MLD of the specific two-way coupled, or assimilative simulation and the one-way coupled free run (panel A). The black line shows the boundary of the continental shelf (bathymetry $< 200m$).

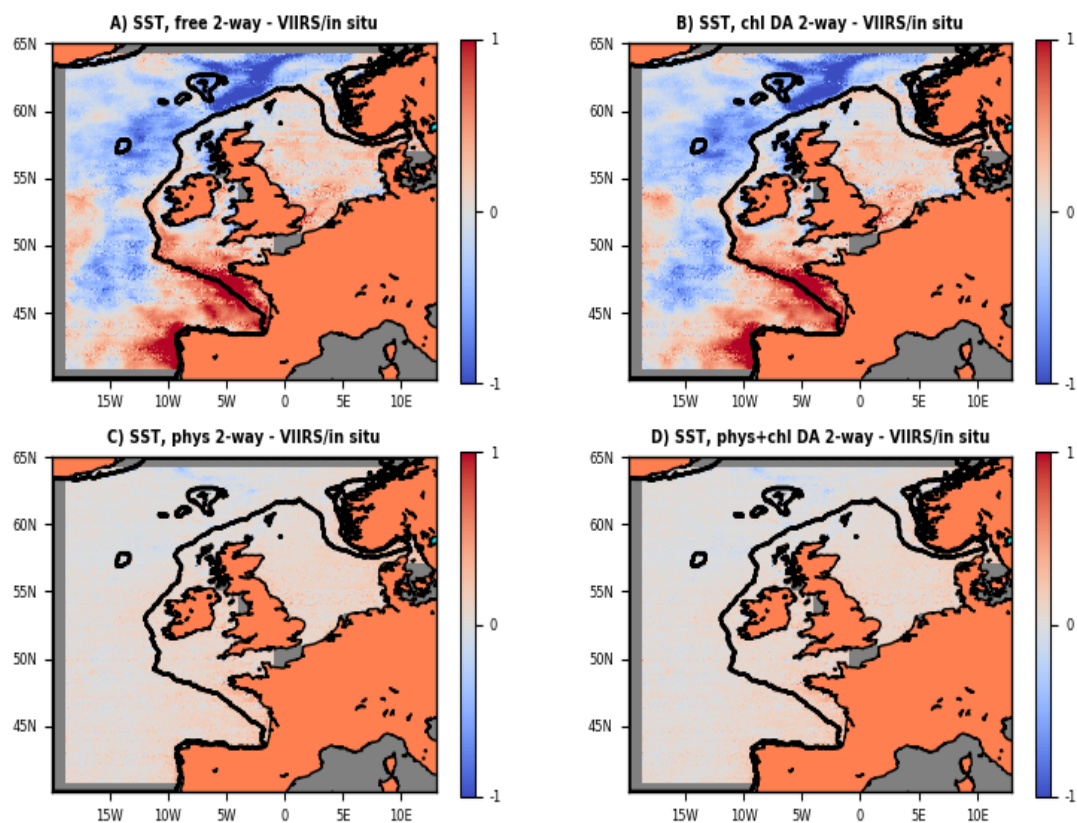


Figure S7. The model to VIIRS/in situ SST differences in $^{\circ}\text{C}$. The differences are shown for the: free two-way coupled model (panel A), physical data assimilation into the two-way coupled model (panel B), chlorophyll assimilation into the two-way coupled model (panel C), and joint physical data-chlorophyll assimilation into the two-way coupled model (panel D).

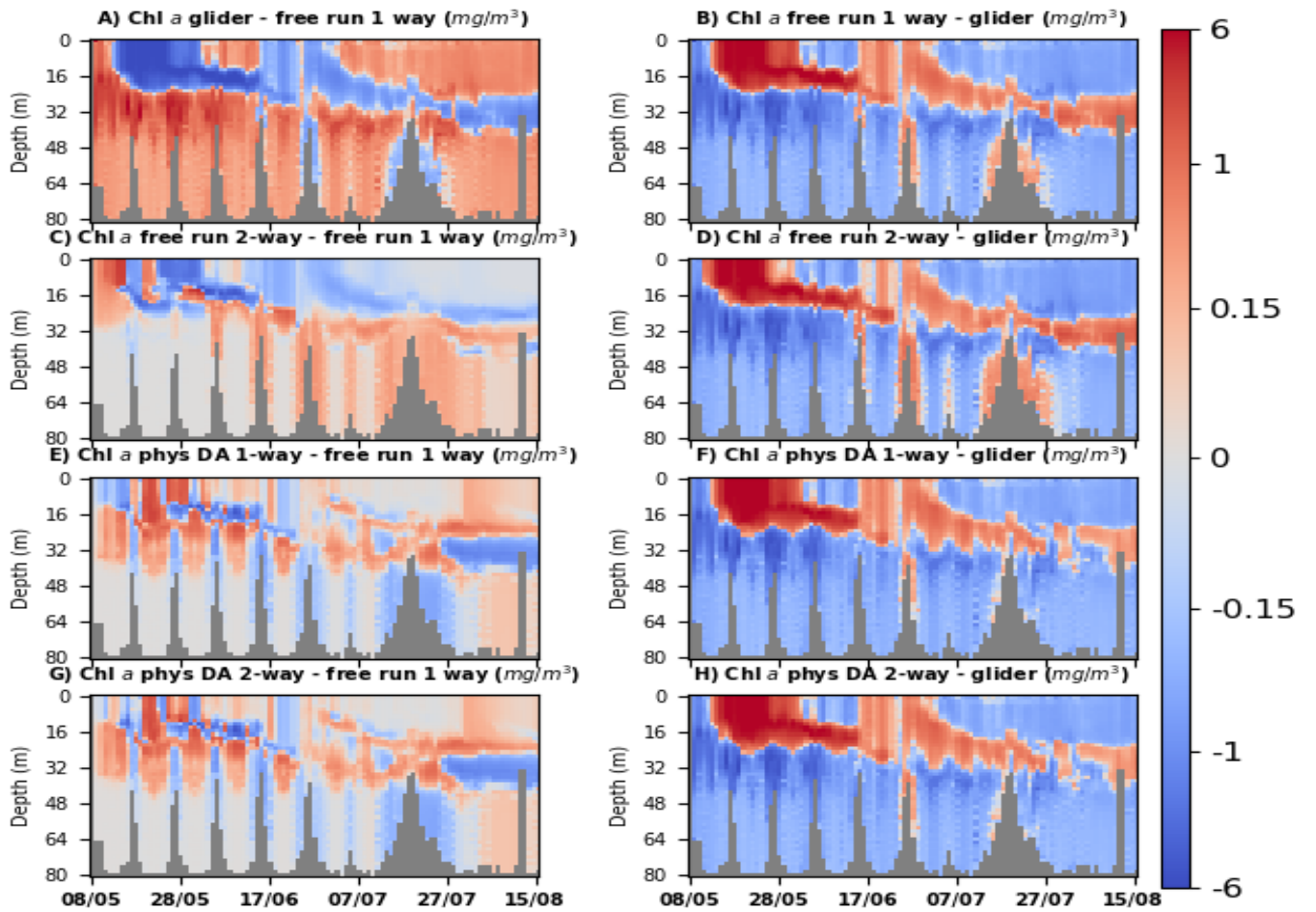


Figure S8. Hovmöller diagram for chlorophyll concentrations (mg/m^3) measured by the Cabot glider in the central North Sea during an early May to mid-August 2018 mission. The right-hand panels (B,D,F,H) show the chlorophyll differences between the free one-way coupled model run (panel B), free two-way coupled model run (panel D), the physical data assimilation into the one-way coupled model (panel F), the physical data assimilation into the two-way coupled model (panel H), and the Cabot glider observations (model minus glider). The left hand panels show the differences between the observations, or model simulations and the reference, free one-way coupled model run. The purpose of the left-hand panels is to show the desired changes to the one-way coupled model (panel A) and how these changes are realized by the biogeochemical feedback in the free run (panel C) and in the physical data-assimilative runs (panels E and G).

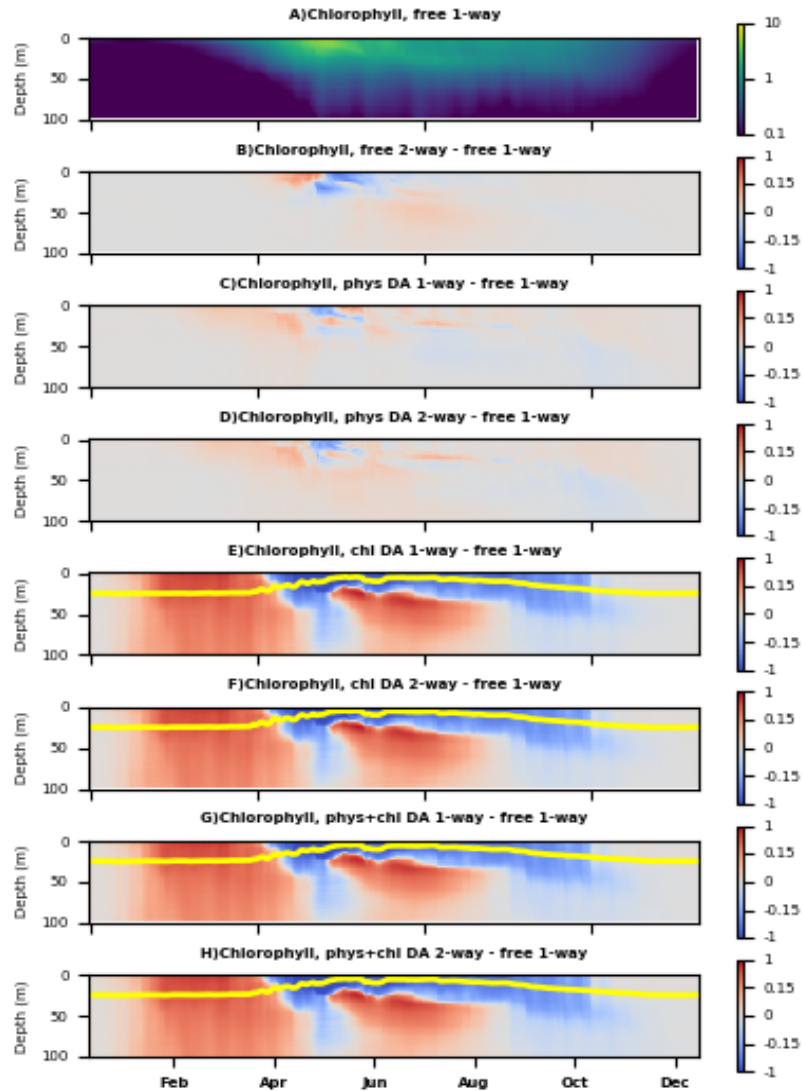


Figure S9. Impact of two-way coupling and assimilation on the simulated chlorophyll concentrations (mg/m^3). The panel A shows Hovmöller diagram (time on the x-axis vs depth on the y-axis) for the one-way coupled model free run, where the values for each day and depth represent the horizontal spatial averages throughout the NWE Shelf (bathymetry $< 200m$). Panels B-H show the same Hovmöller diagrams, but for the differences between the two-way coupled, or assimilative runs and the reference, free one-way coupled run. The yellow lines in the panels E-H show the mixed layer depth, providing the boundary of the region in which the ocean color assimilation directly updates the simulated chlorophyll.

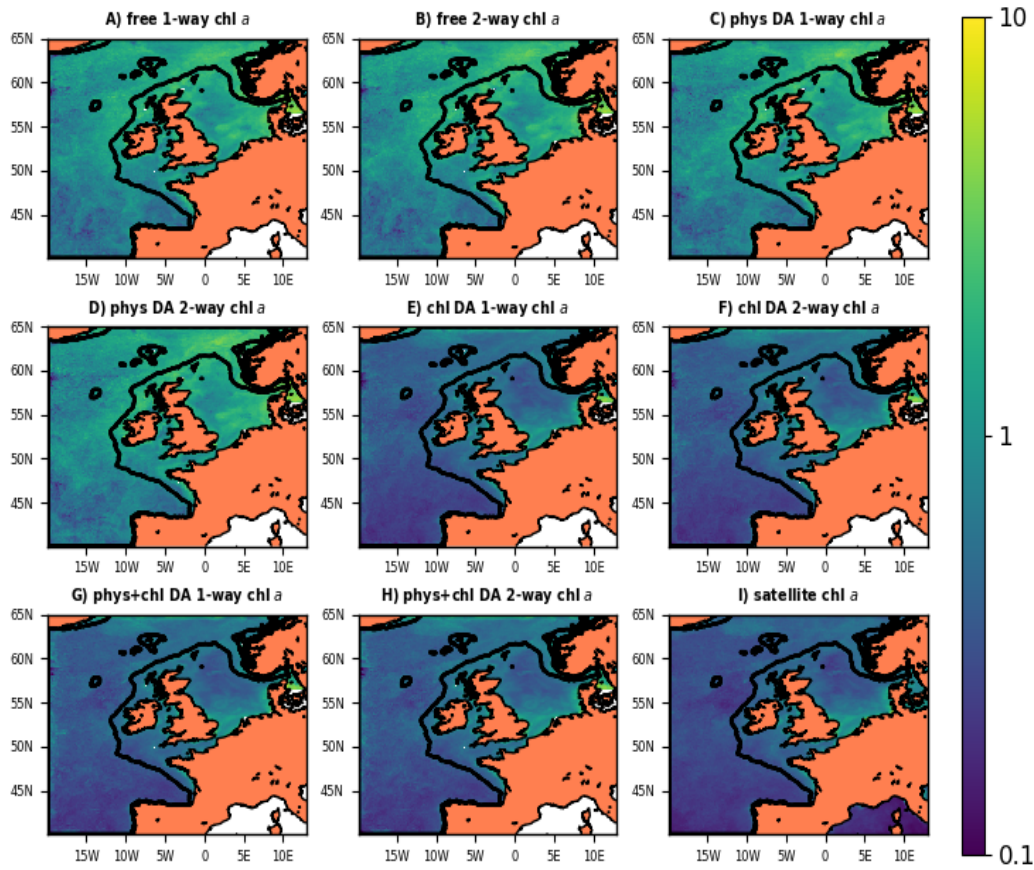


Figure S10. The 2018 mean surface chlorophyll concentrations (in mg/m^3). The different panels compare: the one-way coupled model free run (panel A), the two-way coupled model free run (panel B), the physical data assimilation into the one-way coupled model (panel C), the physical data assimilation into the two-way coupled model (panel D), the chlorophyll assimilation into the one-way coupled model (panel E), the chlorophyll assimilation into the two-way coupled model (panel F), the joint physical data-chlorophyll assimilation into the one-way coupled model (panel G), the joint physical data-chlorophyll assimilation into the two-way coupled model (panel H), and the assimilated satellite ocean color observations (panel I). The black line shows the continental shelf boundary (bathymetry < 200m).