Dear Dr Paul Halloran and Reviewers,

Thank you for your comments and suggestions about this work. Please note that in this document (pdf file), it is included a point-by-point response to the reviews of Reviewer 1, and Reviewer 2. Finally, it is also added a marked-up manuscript with all the changes done through this revision.

Kind regards,

Bahamondes Dominguez et al.

<u>Referee 1</u> General comments

Referee comment: The manuscript describes improvements made to a 1D nutrients-phytoplankton model in order to better describe the dynamics of the pelagic ecosystem in terms of phytoplankton growth and zooplankton biomass. Model simulations that included flexible stoichiometry and photo acclimation mechanisms for phytoplankton and variable grazing by zooplankton were compared with an ample set of observations collected in a temperate shelf sea. The integration of modelled and observed estimates of phytoplankton biomass and physiology, and zooplankton biomass, constitutes the main strength of the work.

However, from my perspective, it is not clear under the current state of the manuscript what are the main findings of the work and why are they relevant. Model improvements are not innovative per se, and it is not clear how these improvements have contributed to provide insight into the dynamics of the ecosystem in such shelf sea.

Also, the model description is not very detailed and this makes difficult to understand which are exactly the model improvements that the authors are testing in this work. The description of sampling procedures is not complete, since relevant information about samples' collection and analysis techniques is missing.

Response: Thank you for your overall comment in this work, which provides a general overview of what was done and comments on the novelty of this paper. GMD is a scientific journal dedicated to the public discussion of the description, development, and evaluation of numerical models as described in the website. This paper was uploaded as a "Development and technical paper", for which the aim is to describe technical developments and model improvements. We think that this paper meets the requirements to be published under the standards of GMD and the manuscript type selected as the aim is to show the improvement in a published model (S2P3 v7.0) based on comparison to observations, which are demonstrated to have a better agreement with the dataset used for the CCS location. We demonstrate an improvement in assessing the dynamics of the selected region, increasing confidence in the results with the model (S2P3 v8.0). Furthermore, this paper also highlights the relative importance of certain processes (e.g. zooplankton grazing and photo-acclimation), which under the general perspective of shelf sea modeling are processes that can be better understood with the sensitive analysis done in this work and which should be taken into account for accuracy in the representation of the shelf sea ecosystem. However, we understand that the initial manuscript might have had some confusing sections based on the comments of this reviewer and we will revise the manuscript to allow clearer understanding of the model development and specific relevance to the biogeochemistry of shelf seas.

<u>**Changes in manuscript:**</u> Please see our responses to following specific comments and the associated changes.

Specific comments

Introduction

Referee comment: My main comment about the introduction is that is not clear from the text what is the point of the work, what does it add to the existing knowledge?

Response: The point of the work is substantial development of the biological component of the model, as previously reported in Marsh et al. (2015). This is already clearly articulated in the Introduction (see L 26-42).

<u>Changes in manuscript:</u> None needed.

Referee comment: L50-54 This paragraph is difficult to understand. The authors say that they implemented flexible stoichiometry in the model and photoacclimation in the chlorophyll description, but they cite Droop which suggest that they have also included some functional relationship between growth rate and internal nutrient status or quotas. I think the authors should clarify which are exactly the modifications done in the model, only flexible stoichiometry or a full Droop's model.

Response: As stated in the indicated section, we implemented the Geider et al. 1998 model, which does indeed link growth to internal cellular nutrient status. We have clarified this in the text

Changes in manuscript: Moreover, changes in nutrient availability can further alter cellular chlorophyll and nitrogen quotas (Droop, 1983; Geider et al., 1998). The combined representation of these two processes in the physiological model of Geider et al. (1998) has been widely implemented in biogeochemical models (Moore et al. 2002). We term this new version of S2P3 v7.0 as S2P3-Photoacclim, which relates phytoplankton growth rates to cell quota (Droop, 1983), through linking the light-, nutrient-, and temperature-dependencies of phytoplankton growth rate to varying ratios of N : C : Chl (Geider et al., 1998).

<u>Referee comment:</u> L65 The authors mention here several models, but it is easy to get lost at this point. Perhaps a list of the models compared would be useful.

<u>Response</u>: We agree that some context is helpful here.

<u>Changes in manuscript:</u> Each characteristic is now related to a model in this sentence to about confusion. Reference to Figure 1 also allows clarification.

<u>Referee comment:</u> L69 "the effects of photo-acclimation and flexible stoichiometry" on what?

<u>Response</u>: We refer to the effects of these processes in the CCS location and to the response of phytoplankton to changes in their environment.

<u>Changes in manuscript</u>: Further comments and details have been added (see L69-72).

Methods

Referee comment: L107 Is that really internal nitrogen (m3 as phyto biovolume) or it is just phytoplankton nitrogen concentration in sea water? I believe is the latter, as in zooplankton (L108), so the sentence should state nitrogen and not internal nitrogen.

<u>Response</u>: Yes, it corresponds to phytoplankton nitrogen concentration.

<u>Changes in manuscript</u>: We have amended descriptions as 'internal nitrogen', rather than 'nitrogen'.

Referee comment: L101-118 This model description is not very clear. Does S2P3-NPZ includes flexible stoichiometry? I guess it does since phytoplankton is described with Chla and N. If the ratio is fixed there is no need to compute both. The authors mentioned in the introduction that the approach used was a Droop model, where the growth rate has to depend on the internal quotas of nutrients. The way the authors describe the model it seems that S2P3-Photoacclim allows flexible stoichiometry but it is not clear how growth depends on quotas. It is also the growth dependent on quotas in S2P3-NPZ? I would suggest the authors to include the equations of the phytoplankton growth model they are using, to ensure clarity and reproducibility.

Response: S2P3-NPZ includes a variable Chl:N as in S2P3 v7.0. For S2P3-Photoacclim the Chl:N:C varies, according to Geider et al. 1998 and in the source code of the model provided in Code and Data availability section. The choice of not including equations here and providing details of the equations with Figure 1

was decided based on the already published equations from different manuscripts (Geider et al., 1998, Sharples et al. 2006, Marsh et al., 2015). But we agree that further information about the S2P3 v8.0 should be provided with the equations.

<u>Changes in manuscript:</u> We will provide additional model details in Supplementary material (see Appendix A).

Referee comment: Figure 1. I imagine the idea of Figure 1 is to illustrate those equations without having to display them. Figure 1 is useful but not all symbols are explained. For instance, u, Z (zooplankton?) and QP are not described. Also, uptake of N in a) is u x PhytoChla but in b) to d) is u x PhytoC, I would say that a Q is necessary somewhere to convert units. And also in Figure 1, it is not clear how growth or uptake depend on quotas, if they do. Since one of the main goals of the manuscript is to account for flexible stoichiometry and hence simulate elemental quotas, maybe for improving clarity the authors could give clear names or symbols to each quota and refer to them throughout the manuscript.

Response: It is difficult to describe all those variables and parameters with one figure.

<u>**Changes in manuscript:</u>** As noted above, we will provide additional model details in Supplementary Material.</u>

Referee comment: L126. The description of the sampling performed during the SSB programme needs improvement: L129. Samples collected for what? Chlorophyll? L131. Please, indicate where this mooring is located. L132. I don't understand what info gives the term "stainless and titanium". L133. How these discrete samples were collected? with bottles? L134. The details about the CDT deployment in the other locations (not CCS) are missing.

<u>Response</u>: We agree that details of observations are appropriate.

Changes in manuscript: For this set of questions about section 4.1, more details are provided for the SSB programme in terms of surface chlorophyll, indicating where the mooring was located (lat, lon), how and where the discrete samples were collected (Niskin bottles), and further details of other CTD deployments in the Celtic Sea are provided in Table A1. The term of 'stainless and titanium' has been removed as an unnecessary detail.

Referee comment: L136. The description of the sampling of zooplankton biomass needs improvement: L138. How this zooplankton samples are collected? With nets? If so, please, specify the type of net and the type of trawl. L140. How

zooplankton biomass was measured?

Response: Again, we agree that further measurement details here are informative, although we note that a fuller description is provided in the published paper describing this data.

<u>**Changes in manuscript:</u>** A more detailed description of the zooplankton biomass sampling in section 4.2 has been added. This further detailed description includes information about how the zooplankton was collected (using nets of different sizes). Also, it is explained how zooplankton biomass was measured (using a FlowCam and ZooScan).</u>

Referee comment: L146. The description of the sampling for phytoplankton is also very vague: L148. Phytoplankton samples were collected with bottles? At "a number of stations" seems very vague.

<u>Response</u>: Again, we agree that further measurement details here are informative.

<u>Changes in manuscript</u>: Further description of the physiological observations is detailed in section 4.3 now. Phytoplankton were collected at different stations in the Celtic Sea and at different depths using Niskin bottles. Please see these added details in first paragraph of section 4.3.

Results

<u>Referee comment:</u> L177 The behaviour of S2P3-Photoacclim (blue line) seems very different to S2P3- NPZ, similar maybe in magnitude but not in timing.

Response: Given further consideration about the physiological observations used, it was decided that the stations CS2 and N1 should not be used for the validation of the S2P3-Photoacclim and S2P3 v8.0 models because one of the important assumptions of these models is that they do not consider advective fluxes and these stations are very close to the shelf edge where horizontal advection is relevant. This allowed better constraint of S2P3-Photoacclim and, therefore, it presents an improved representation of the ecosystem (timing and magnitude of the spring bloom are more similar to the buoy observations, Figures 3a, 4).

<u>Changes in manuscript:</u> For new values constrained in this model see Table 1. Table 3 values of S2P3-Photoacclim are also amended.

<u>Referee comment:</u> L185 Zooplankton biomass increased 1 month later than phytoplankton in 2014 but definitely not in 2015.

<u>Response</u>: We agree that this is worthy of note.

<u>**Changes in manuscript:**</u> Section 4.4, L207, sentence rephrased specifying that the 1 month difference occurs during 2014, while in 2015 there is not.

Referee comment: L190 The expression "have not reached the SPB" is ambiguous. Maybe they haven't reached the peak of the bloom, but, at least for S2P3-NPZ, the accumulation of biomass has already started accordingly to Figure 3. It would be necessary to clarify to which event of the SPB the authors refer.

<u>Response</u>: The timing of the spring phytoplankton bloom corresponds to a specific threshold.

<u>**Changes in manuscript:**</u> We now clarify and detail further as the term 'reaching the spring phytoplankton bloom' according to different definitions in the literature. Based on this definition, dates for the spring phytoplankton bloom being reached by each model are now also specified. The corresponding sentence is rephrased (L215).

<u>Referee comment:</u> Figure 4. I miss the same figure for 2014, are CTD data not available for this year?

Response: CTD observations were available for the CCS location in 2014 only during November, otherwise the year 2014 would have been included to compare CTD casts during spring and summer as in the year 2015.

Changes in manuscript: None needed.

Referee comment: L203. The description of results regarding phytoplankton physiology seems a bit short and does not give much information about the agreement of the model and observations: L205 If it is a 1D model, are not the vertical gradients the only ones available? Figure 5. The vertical gradients are difficult to see for observations, especially for Pmax.

<u>Response</u>: We agree that further details should be provided in the description of the results that compare phytoplankton physiology between the model and observations.

<u>**Changes in manuscript</u>**: Further description of the extent to which the modelled values and vertical gradients in Figure 5 match the data will be added.</u>

Referee comment: L218. To which location correspond Figures 6 and 7?

<u>Response</u>: All this paper is based at the CCS location, with all models calibrated

for that specific region.

<u>**Changes in manuscript:**</u> This is specified again in this sentence, which is also rephrased (L244).

Referee comment: L236. In Figure 7 I don't feel it is possible to see which event follows which. Maybe a mean seasonal cycle with some metric for interannual variability would be more easy to interpret.

<u>Response</u>: We agree to include details of the seasonal cycle and associated variability.

<u>Changes in manuscript</u>: Figure 7 has been modified to show the seasonal cycle of each variable in the S2P3 v8.0 model (black lines) and with the red lines representing the 95% quantiles (i.e. 95% of the data lie between these lines) of each variable over the 51 years of simulation to represent the inter-annual variability present in the dynamics of the ecosystem.

Referee comment: Figure 7 legend. Does this "forced with all meteorological components" refer to the whole model result? I imagine it does, but here it seems it refers to NPP only.

<u>Response</u>: We agree that clarification is helpful.

<u>Changes in manuscript:</u> Legend of Figure 7 rephrased and more details added to avoid this confusion.

Referee comment: L244. Parameters listed in Table 1. The reader does not know where these parameters fit into the model. They are not in Figure 1 and no equation that includes them is shown in the manuscript.

Response: See response to earlier comments.

<u>**Changes in manuscript:</u>** As noted above, we will provide additional model details in Supplementary Material (see Appendix A).</u>

<u>Referee comment:</u> Not all A Figures are described, are all necessary?

Response: We agree that further details were needed in the description of those figures. They were all described but not explicitly referred to which figure it corresponded.

<u>Changes in manuscript:</u> Reference to each figure has been added and further details to show the relevance of including each of them in the Appendix (see

Section 5).

Referee comment: L276. The meaning of this sentence is not clear to me.

<u>Response</u>: Noted. This sentence has been re-written in a clearer way

<u>Changes in manuscript:</u> See L322 for changes in this sentence.

Referee comment: Table 2. It is not clear what this "timing of the SPB" means. Is it day of the onset, day of the peak? DW is dry-weight? If I am not wrong, it is not explained anywhere in the manuscript.

<u>Response</u>: Timing of the spring phytoplankton bloom is referred as the quantitative definition for the start of the spring phytoplankton bloom.

<u>Changes in manuscript</u>: This is now described in section 4.4, L217. In the description of Table 2, it has been specified that corresponds to such definition. For the DW units of zooplankton, they are now explained in section 4.2, for zooplankton biomass dataset, L165.

<u>Referee comment:</u> To which location/area correspond the metrics in Table 2 and Table 3?

Response: For the CCS location.

<u>**Changes in manuscript:**</u> Specified now in the legend of Tables 2, 3, and in the description of each table (L298, LL314).

Conclusions

Referee comment: L300. I am afraid I wouldn't call this work innovative. NPZ models that include grazing are commonly used. Flexible stoichiometry I would say is almost the norm when describing phytoplankton pools. And the description of photoacclimation through the parameterization proposed by (Geider et al. 1998) is virtually standard in biogeochemical models. L305. This last sentence is difficult to understand. And also, I miss one sentence or paragraph that clearly states why this work is an advance in our knowledge.

<u>Response</u>: As we explained in response to general comments, the advance of this work is substantial development of a model originally described in GMD.

Changes in manuscript: None needed.

Technical corrections

Referee comment: L18 SSB acronym not defined.

<u>Change to manuscript:</u> Amended. Defined now in L18.

<u>Referee comment:</u> I would say "data" is plural (datum is the singular) and has been used as singular in several places (L20, L75, L82).

<u>**Change to manuscript:**</u> Noted. Lines where 'data' is used, it is changed as plural in their description. These include L20, L75, L82.

Referee comment: Sections 2 and 3 use mainly present tense, but there are several pasts in between that maybe can be reviewed (L76, L82, L103).

<u>Change to manuscript:</u> Noted and amended in sections 2 and 3.

<u>Referee comment:</u> L203 "Moreover" and "also" in the same sentence sound redundant.

<u>Change to manuscript:</u> Sentence amended. Moreover only included (L234).

<u>Referee comment:</u> L250. I am not sure, but I feel it is more clear to refer to ingestion rate of zooplankton and not phytoplankton.

<u>Response</u>: We understand why it is confusing thanks to your comment. To avoid further confusion, we will refer to Rm as: zooplankton maximal grazing rate.

<u>Change to manuscript:</u> Amended in L250, L253, Table 1.

Referee 2

Dear Sirs (English version):

Referee comment: The model presented by Angela Bahamondes Dominguez is interesting and it is related to the scope of the journal (Geoscientific Model Development). Overall, I find the model showing a considerable fit with the field data, and this suggest that could be useful to improve our understanding about how phytoplankton blooms take place in situ. For all these reasons, I think this article is within the standards of excellence of the journal.

<u>Response</u>: Thank you for your comment regarding the standards of this paper. We think that describing developments in a model at a technical level complies with the aims of Geoscientific Model Development.

Referee comment: The model presented (S2P3 v8.0) is a modification of a previous NPZ model (S2P3 v7.0) to which the photo-acclimatization of phytoplankton is introduced. This new model improves the previous model, as the results of the new model are better adjusted to in situ observations. However, the temperature is taken into account only in the respiration process, while it is not considered in photoacclimation and grazing, and both processes depend on it (Sarmento et al . 2010, Vázquez-Domínguez et al. 2013 and references therein). Furthermore, the model is slightly decoupled to the field data, as it happens in the second period (Fig. 3a) or the zooplankton biomass (Fib 3b), and besides it presents a mismatch with nitrogen (Figure 4c). This should be discussed.

<u>Response</u>: We appreciate the points raised by the reviewer. Firstly, we are aware of the effects of temperature on phytoplankton physiology and grazing. Within the developed model iteration we explicitly include the representation of temperature on phytoplankton physiology using the model of Geider et al. (1998). We acknowledge that representation of temperature effects on zooplankton grazing could also have been included. However, we note that there are always further processes which could be included and decided that the development step of the model described here was substantive enough to be worth a formal description within GMD. We note that there remains discrepancy between the model and the data, indicating further potential avenues for development.

<u>Changes to manuscript</u>: We will add the full equations of S2P3 v8.0 in Supplementary Material (see Appendix A). Furthermore, regarding the notemperature dependencies of grazing and photo-acclimation in this model, a sentence is added regarding this assumption in L216. Finally, in L225-226 is explained about the differences in DIN for Figure 4c.

Referee comment: Two additional questions: 1) it would be good to unify units in figure 6 since they are mixed (Chla, N, C), and this does not allow to estimate a transfer efficiency between trophic levels. If a scale with mmol C m-3 is added, the efficiency in carbon transfer can be estimated between phytoplankton-zooplankton, and if the same is done with N we would know the transfer

between N-P-Z. This is important at the biogeochemical level; and, 2) similarly, perhaps two columns could be added in Table2, indicating the mg C m-2 of phytoplankton and zooplankton.

Response: We agree that showing the transfer efficiency between trophic levels should be added.

<u>Changes to manuscript:</u> Figure 7 is changed to more easily represent this transfer efficiency, by adding a subpanel of phytoplankton N. Therefore, the transfer efficiency between trophic levels can be seen in terms of mmol N m-3 between phytoplankton-zooplankton-nutrients.

Referee comment: Finally, figure 7 (a) shows an inter-annual change in the intra-annual temperature variability, which may be due to temperature changes at the decadal level. All these changes seem to affect the biomass of phytoplankton, but they are not so apparent in the remaining variables. Perhaps, you have an explanation of these differences.

<u>Response</u>: We agree that this variability can be more clearly presented.

<u>Changes in manuscript</u>: Figure 7 has been modified to give the reader better insights about the dynamics of the model. The new figure shows the annual seasonal cycle of each variable (black line) and with the red lines representing the 95% quantiles (i.e. 95% of the data lie between these lines) of each variable over the 51 years of simulation to show the inter-annual variability of each variable. It is more apparent in this figure that not only biomass of phytoplankton shows changes through each year, but also all the remaining variables.

Constraining the response of phytoplankton to zooplankton grazing and photo-acclimation in a temperate shelf sea with a 1-D model towards S2P3 v8.0

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Abstract. An established 1-dimensional model of Shelf Sea Physics and Primary Production (S2P3) has been developed into three different new models: S2P3-NPZ which includes a Nutrient-Phytoplankton-Zooplankton (NPZ) framework, where the grazing rate is no longer fixed, but instead varies over time depending on different functions chosen to represent the predator-prey relationship between zooplankton and phytoplankton; S2P3-Photoacclim which includes a representation of the process

- 5 of photo-acclimation and flexible stoichiometry in phytoplankton; and S2P3 v8.0 which combines the NPZ framework and the variable stoichiometry of phytoplankton at the same time. These model formulations are compared to buoy and CTD observations, as well as zooplankton biomass and *in situ* phytoplankton physiological parameters obtained in the Central Celtic Sea (CCS). Models were calibrated by comparison to observations of the timing and magnitude of the spring phytoplankton bloom, magnitude of the spring zooplankton bloom, and phytoplankton physiological parameters obtained throughout the
- 10 water column. A sensitivity study was also performed for each model to understand the effects of individual parameters on model dynamics. Results demonstrate that better agreement with biological observations can be obtained through the addition of representations of photo-acclimation, flexible stoichiometry, and grazing provided these can be adequately constrained.

1 Introduction

Shelf seas are ocean regions where water depth is less than a few hundred metres (~ 200 m) and represent only ~ 10% by area
of the global ocean. However, these systems have a disproportionate importance because of their exceptionally high biological productivity (Holt and Proctor, 2008), being responsible for 15 to 30% of the total oceanic primary production (PP) (Wollast, 1998; Muller-Karger et al., 2005; Davis et al., 2014). Research vessels and remote, autonomous vehicles have been used to study shelf sea regions such as in the SSB-Shelf Sea Biogeochemistry (SSB) research programme (https://www.uk-ssb.org/), whose aim was to increase the understanding of how physical, chemical and biological processes interact on UK and European

20 shelf seas, collecting observations throughout 2014 and 2015 in different regions of the UK shelf sea, although this data is these data are not synoptic (i.e. it is they are not sampled at different locations simultaneously). To complement the available data from research vessels, ocean models have been used to study and understand marine biogeochemistry, including a variety of high spatial resolution models to represent the biogeochemistry of shelf seas with high complexity and horizontal spatial resolution (Sharples, 1999, 2008; Edwards et al., 2012; Marsh et al., 2015).

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Different models have been developed to study plankton communities, ranging from very simple ones, e.g. the Lotka-Volterra competition model (Volterra, 1926; Lotka, 1932) to more sophisticated ones, adding more degrees of complexity by including representation of the physical processes of advection and diffusion, or more complexity in ecosystem functions through representation of different groups of organisms and/or size structure. For example, coupled models such as the Nucleus for European

- 30 Modelling of the Ocean (NEMO) and European Regional Seas Ecosystem Model (ERSEM) (Edwards et al., 2012), Regional Oceanic Modelling System (ROMS) (Shchepetkin and McWilliams, 2005), and Finite Volume Community Ocean Model (FV-COM) (Chen et al., 2003). Although complexity can be useful for describing the interacting behaviour of multiple system components, incomplete understanding of the ecology and key processes of the organisms, and the lack of data for validation (Anderson, 2005) can reduce the reliability of predictions. Moreover, simulations of models like NEMO-ERSEM, ROMS, and
- 35 FVCOM rely on high-performance computing resources and running multiple sensitivity analyses and experiments is difficult. In contrast, simpler models like the Shelf Sea Physics and Primary Production (S2P3) model (Sharples, 1999; Simpson and Sharples, 2012) have been used to study the dynamics of shelf seas and to simulate seasonal stratification with greater computational efficiency by using a 1-D Nutrient-Phytoplankton (NP) model to represent physical and biogeochemical processes in the water column. In temperate shelf seas, away from advective sources such as the shelf break or plumes from rivers, horizontal
- 40 processes can be neglected in comparison to vertical processes, thus, although S2P3 does not consider advective fluxes it can make a good representation of the dynamics in the water column for temperate shelf seas (Sharples et al., 2006; Sharples, 2008; Marsh et al., 2015).

The S2P3 v7.0 model was introduced and developed in the work of Marsh et al. (2015), where it was outlined that further 45 development of that model would include resolving phytoplankton physiology. In this study, the S2P3 v7.0 model is developed by allowing variations in light intensity to produce phenotypic adjustments in the phytoplankton cells by changing the chlorophyll content of the phytoplankton and, therefore, the cellular absorption cross-section (Macintyre et al., 2002). This phenotypic change in response to variations in the photon flux density is called photo-acclimation (Falkowski and Laroche, 1991; Moore et al., 2006). The main property of photo-acclimation is the reduction of photosynthetic pigment content in response

- 50 to increased irradiance (Falkowski and Laroche, 1991). Moreover, changes in nutrient availability can further alter cellular chlorophyll and nitrogen quotas (Droop, 1983; Geider et al., 1998), incorporating a combined representation of these two processes (Geider et al., 1998), this new version of S2P3 v7.0 we term S2P3-Photoacclim and relates phytoplankton growth rates to cell quota (Droop, 1983), describing the light-, nutrient-. and temperature-dependencies of phytoplankton growth rate to varying ratios of N : C : Chl (Geider et al., 1998). On the other hand, simpler models using an NPZ or NPZD framework, with
- 55 the use of nutrients, phytoplankton, zooplankton, and detritus as the main model components (e.g., Steele, 1974; Wroblewski et al., 1988; Anderson, 2005) have shown good agreement with observations in terms of chlorophyll and PP, by simulating the timing and magnitude of the spring phytoplankton bloom in different regions of the ocean. Despite their relative simplicity, NPZ models can be a better option to approach an understanding of the physics and biology of an ecosystem, which lead to

a further development of the S2P3 v7.0 model where the simplest assumption of a fixed proportion of phytoplankton being

- grazed and remineralised into the DIN pool (grazing rate) is developed into an NPZ framework (S2P3-NPZ model), using a 60 Holling Type 2 or Ivlev grazing functional response of zooplankton grazing on phytoplankton (Franks, 2002), which shows a saturating response to increasing food.
- A combination of photo-acclimation, flexible stoichiometry (S2P3-Photoacclim model), and NPZ framework (S2P3-NPZ 65 model) is then performed to produce a newly developed model called S2P3 v8.0 (Figure 1). This paper presents a thorough analysis in terms of sensitivity to biological parameter values in each new developed model, resulting in differences of the model structure. A comparison between each model demonstrates how structural differences influence the representation of the spring phytoplankton bloom and annual PP. The aim of this paper is to provide a better understanding about the predatorprey relationship between zooplankton and phytoplankton, the effects of photo-acclimationand flexible stoichiometry and
- 70 thus the importance of these processes for accurate representation of phytoplankton dynamics in shelf sea modelling, flexible stoichiometry in a simple 1-D model. Model outputs are compared with observations of phytoplankton responses to physical forcing to illustrate importance of different processes representation.

2 Study region and model setup

- This study is focused on the Central Celtic Sea (CCS), a region located in the North-Western (NW) European Shelf, which 75 is characterised by its tidally dynamic environment and summer stratification (Pingree et al., 1978; Sharples and Holligan, 2006; Hickman et al., 2012). Daily meteorological data, provided by available from the National Centers for Environmental Predictions (NCEP) Reanalysis data (http://www.esrl.noaa.gov/psd) was are used to force the model at the CCS site, located at $49.4^{\circ}N$, $8.6^{\circ}W$. Wind speed (m s⁻¹), cloud coverage (%), air temperature (°C), and relative humidity (%) variables from this dataset were are all used to force each model version. 80

The following description of model setup is applied to each model structure developed in this work. Tidal components consist of the u-component (semi-major axis) and the v-component (semi-minor axis) for the M_2 , S_2 , and N_2 tidal constituents. This data was Tidal data are obtained from a fine mesh (12km resolution) covering the UK shelf. Tidal currents are predicted

- 85 using the Proudman Oceanographic Laboratory Coastal Ocean Modelling Systems (POLCOMS) 3-D shelf model (Holt et al., 2009; Wakelin et al., 2009) with an output extracted for the CCS location. Moreover, each model is initialised on 1st January of the first year of simulation with a temperature of 10.10 °C at all depths, and water column presumed mixed throughout, including a vertical resolution of 1m (i.e. 140 vertical levels). Initial values of physical variables are consistent with former studies (Sharples, 1999, 2008; Marsh et al., 2015), whereas initial values of biological variables are based on observations
- of zooplankton biomass at the CCS location over winter months of 0.02 mmol N m^{-3} (Giering et al., 2018); phytoplankton 90 chlorophyll correspond to a typical winter value for the CCS location of 0.2 mg Chl m^{-3} , and the DIN initial value is 7 mmol

 $m N~m^{-3}$. The initialised variables are only set up at the start of each simulation and do not reset in between years.

3 Model development

- 95 The S2P3 v7.0 model can be divided into two different components: a physical part and a biological part. For this research, the model is an improved version of the original described in Sharples et al. (2006) to be compiled and executed in a Unix environment (Marsh et al., 2015), allowing a 1-D representation of physical and biological processes in shelf seas by simulating the seasonal cycle of phytoplankton, water column stratification, and PP at a selected location defined by water depth and tidal current amplitude. The physical part of the model has been greatly described in many other studies (Sharples, 1999; Sharples et al., 2006; Sharples, 2008; Simpson and Sharples, 2012; Marsh et al., 2015) and is not described in this section again. This
- model uses the turbulence closure scheme based on Canuto et al. (2001). Likewise, the biological part of the S2P3 v7.0 model is described in Marsh et al. (2015) (Figure 1a).
- In order to explicitly account for the influence of zooplankton grazing and, hence, predator-prey dynamics, the S2P3 v7.0 model was-(Sharples et al., 2006; Marsh et al., 2015) is developed into an NPZ framework. This new version of the model (S2P3-NPZ) includes zooplankton as a state variable, contrary to the S2P3 v7.0 model where grazing was (*G*) is calculated as a fixed seasonal cycle which was represented as a sink term in the phytoplankton tendency equation. Addition of an explicit grazer also allows comparison to zooplankton biomass observations. As within S2P3 v7.0, the biological part of the S2P3-NPZ model calculates phytoplankton biomass in chlorophyll currency (*Phyto_{chl}*; mg Chl m⁻³). Similar to the S2P3 v7.0
- 110 model, the growth of phytoplankton biomass (μ) can be either nutrient-limited or temperature-limited, with the maximum growth rate of phytoplankton being related to temperature through an Eppley function and being modified by a nutrient quota (Q), which corresponds to a varying ratio between phytoplankton nitrogen and phytoplankton chlorophyll biomass ($Q = Phyto_N/Phyto_{chl}$). Additionally, phytoplankton are modelled in terms of internal nitrogen (mmol N m⁻³) represented by $Phyto_N$. Zooplankton biomass and external DIN are likewise modelled in terms of nitrogen (mmol N m⁻³; Figure 1b).
- 115

The S2P3-Photoacclim model is a new version of the S2P3 v7.0 model, incorporating a more complete representation of phytoplankton physiology (Geider et al., 1998). This model allows phytoplankton to acclimate to changes in light and nutrients, therefore, the ratios of N : C : Chl and characteristics of phytoplankton physiology can vary, allowing direct comparison to physiological data. The biological part of the S2P3-Photoacclim model uses three currencies of phytoplankton biomass: car-

120 bon $(Phyto_C; \text{mg C m}^{-3})$, nitrogen $(Phyto_N; \text{mg N m}^{-3})$, and chlorophyll $(Phyto_{chl}; \text{mg Chl m}^{-3})$. The S2P3-Photoacclim model calculates phytoplankton growth as a function of both nitrogen assimilation and carbon fixation (i.e. variable Chl : N and Chl : C ratios). It is assumed that respiration (R) is equal for all cellular components as a function of temperature: $R_C = R_n = R_{chl} = R_{ref}T_{function}$, where R_{ref} (d⁻¹) is a degradation rate constant at a reference temperature (Figure 1c). 125 The S2P3 v8.0 model describes a combination of zooplankton and physiological acclimation components in order to provide a more realistic representation of the ecosystem dynamics (Figure 1d). Full details, model equations, and parameters are listed in Appendix A.



Figure 1. Structure of the phytoplankton growth formulations: (a) S2P3 v7.0 model, with constant $Phyto_N : Phyto_{chl}$ ratio (Q); (b) S2P3-NPZ model, including explicit zooplankton with an associated ingestion rate of phytoplankton (I) based on the Ivlev grazing type; (c) S2P3-Photoacclim model, with varying ratios of N : C : Chl, with phytoplankton chlorophyll content regulated by a coefficient of chlorophyll synthesis (ρ_{chl}), which reflects the ratio of energy assimilated to energy absorbed (Geider et al., 1996), and with phytoplankton carbon regulated through a cost associated with biosynthesis ($u\zeta$) (Vries et al., 1974; Geider, 1992; Geider et al., 1998), respiration, and grazing (G); (d) S2P3 v8.0 model, including two different varying quotas for $Phyto_N : Phyto_c$ and $Phyto_N : Phyto_{chl}$.

4 Validation of the models: observations

130 To calibrate or tune each model, they were adjusted on a trial-and-error basis until disagreement with the *in situ* observations was minimised, allowing investigation of the sensitivity of each model to changes in the parameters listed in Table 1.

4.1 UK SSB programme

Observations were collected throughout the NW European shelf during the spring of Time-series of surface chlorophyll-a
concentrations (mg Chl m⁻³) from long-term mooring deployments including the Carbon and Nutrient Dynamics and Fluxes over Shelf Systems (CaNDyFloSS) Smartbuoy (Mills et al., 2003) were collected at the CCS location (49.4°N, 8.6°W, depth 145.8 m), gathering data for 5 minutes every 30 minutes during the years 2014 and 2015 as part of the research cruise expeditions DY026, DY018, DY029 and DY033. Samples were collected at the CCS location (Figure 2) during the pre-bloom, peak bloom, and post-bloom conditions of the area. Time-series of surface The phytoplankton community fluorescence from the water samples were calculated as a proxy for chlorophyll-a concentrations (mg Chl m⁻³) from long-term mooring deployments

including Smartbuoy (Mills et al., 2003) were collected. Moreover, CTD (stainless and titanium) and calibrated taking into account daytime fluorescence quenching which results in a reduction of fluorescence per unit chlorophyll; the chlorophyll fluorescence has an initial rise but then it declines (within less than a second) and finally increases again over a period of several minutes. For this study, day time data was removed.

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<u>CTD</u> casts were performed in different locations of the NW European shelf from the CCS location to the shelf break, with discrete samples of temperature, DIN, and chlorophyll-*a* collected using Niskin bottles as part of the research cruise expeditions DY029 and DY033. At the CCS location, the CTD samples were collected from pre-dawn to midday with a 1m vertical resolution over the whole water column (140m depth) during the year 2015. CTD casts for the CCS location were chosen to validate the model during spring and summer. Relevant information about dates and positions from other CTD casts taken during spring

150 the model during spring and summer. Relevant information about dates and positions from other CTD casts taken during spring and summer of the year 2015 are listed in Table A1. CTD casts chosen to compare the each model are marked in bold in Table B1.

4.2 Zooplankton biomass

- 155 Zooplankton biomass samples were collected at the CCS location (49°25 N, 8°35 W, ~ 150 m water depth) during four periods: 5th 12th August 2014, 10th 29th November 2014, 3rd 28th April 2015, and 13th 31st July 2015 for the cruises DY026, DY018, DY029, and DY033, respectively (Giering et al., 2018). Zooplankton were fractionated into microplankton, small mesozooplankton, and large mesozooplankton by using different mesh sizes. For the zooplankton biomass samples, net rings of 57 cm diameter were used and fitted with two different mesh sizes of 63 μ m and 200 μ m. The nets had a
- 160 closing mechanism when deployed, sampling zooplankton biomass during daytime and night-time at different depth: above and below the thermocline, and, when present, across the deep chlorophyll maximum (DCM; determined based on fluorescence measurements). The thermocline and DCM were determined from CTD casts immediately prior to the net deployments. The 63 μ m and 200 μ m mesh nets were hauled at 0.2 m s⁻¹ and 0.5 m s⁻¹, respectively.

- 165 For the S2P3-NPZ and the S2P3 v8.0 models, only mesozooplankton biomass were considered, with a community composition that included: amphipods, appendicularian, chaetogratha, copepods, euphausiacea, polychaeta, and others (e.g. cladocerans, dinoflagellates, echinoderm, eggs, foraminifera, gymnosomata, unidentified larvae, nauplii, ostracods and radiolarian, all of which contributed < 3% in all samples). The A FlowCam (Fluid Imaging Technologies Inc.) and a ZooScan were used to scan zooplankton individuals with images processed using ZooProcess 7.19 and Plankton Identifier 1.3.4 softwares (Gorsky et al.,</p>
- 170 2010). From these images, biovolume spectra were calculated and converted into image-derived dry weight (DW). The total 246 net hauls collected for biomass samples provided 44 vertical depth profiles, integrating zooplankton biomass typically between 0 and 120 m at the CCS location. The complete data set can be obtained from the British Oceanographic Data Centre (BODC), (http://www.bodc.ac.uk/data).

175 4.3 Physiological observations

Samples were collected at a number of stations Water samples were collected in the Celtic Sea from the cruises JR98 and CD173 using 10-liter Niskin bottles for four of the sites: CS1, CS2, CS3, and IS1 (Figure 2) -during 24-h periods on 31st July, 29th July, 05th August, and 2nd August, respectively. Multiple samples were collected in the surface and in deeper layers (in the surface mixed layer (SML) and the DCM) to obtain different phytoplankton populations throughout the photoperiod. The

- 180 JR98 cruise was undertaken from 24th July to 14th 24th July to 14th August of 2003. Stations, from the Irish Sea to the Celtic Sea shelf break, ranged in characteristics from very strong, narrow thermoclines in the southern Celtic Sea (CS1), to the weak, deep surface layer associated with internal wave mixing at the shelf edge (CS2). On the other hand, the CD173 cruise was undertaken from 15th July to 6th 15th July to 6th August of 2005, from the stratified region of the Celtic Sea shelf (stations D2, CS1, CS3, U2) and shelf break (stations CS2, N1). For this work, observations from both cruises were used, considering
- 185 only the stations from the seasonally stratified sites (B2, CS1, CS3, D2, JB1, OB, P1, U2, and ctd16) and excluding stations CS2 and N1 which are close to the shelf edge where advective fluxes are more relevant than in the stations nearer to the CCS location and none of the models used here consider advective fluxes.

Photosynthesis versus irradiance (P vs E) experiments were conducted in short-term incubations (2 - 4h) using a photosynthesis there is the conducted in the chlorophyll-*a* specific maximum light-saturated photosynthesis rate P^{Chl}_{max} (mg C (mg Chl – a)⁻¹ h⁻¹) and the maximum light utilisation coefficient, α^{chl} (mg C (mg Chl-a)⁻¹ h⁻¹ (μE m⁻² s⁻¹)⁻¹) (Jassby and Platt, 1976; Hickman et al., 2012). Multiple samples were collected in the surface and in deeper layers to obtain different phytoplankton populations throughout the photoperiod. Values of α^{chl} and the light saturation parameter, E_k (μE m⁻² s⁻¹) (given by E_k = P^{Chl}_{max}/α^{chl}) were spectrally corrected to the *in situ* irradiance at the sample depth according to the phytoplankton light absorption (Moore et al.,

2006). For this work, observations from both cruises were used considering only the stations from the seasonally stratified sites (B2, CS1, CS2, CS3, D2, JB1, N1, OB, P1, U2, and etd16). The maximum light utilisation coefficient (α^{chl}) was constrained for the S2P3-Photoacclim and the S2P3 v8.0 models by finding the mean from all observations: $\alpha^{chl} = 9.16 \times 10^{-6}$ mg C (mg

Chl-a)⁻¹ h⁻¹ ($\mu E m^{-2} s^{-1}$)⁻¹ (ranges of this parameter accounted from 3.58×10^{-6} to 3.59×10^{-5} mg C (mg Chl-a)⁻¹ h⁻¹ 200 ($\mu E m^{-2} s^{-1}$)⁻¹; std = 4.38×10^{-6}), and therefore, the values of P^{Chl}_{max} and E_k were used as variables for comparison with equivalent modelled values.



Figure 2. Map for study area and stations for the JR98 and CD173 cruises, including the CCS location (in red colour). Image created with Matlab using the repository data for gridded bathymetry provided by General Bathymetric Chart of the Oceans (GEBCO). Bathymetric data only considered for the shelf sea region (0 to 300m depth) with open ocean depth neglected (deeper than 300m). Continents considered in black colour (over 0m elevation).

Parameters (Units)	Definition	S2P3-NPZ	S2P3-Photoacclim	S2P3 v8.0
	Grazing inefficiency or 'messy feeding'			
γ_1 (dimensionless)	(0.0-1.0), returns a fraction of grazed mate-	0.2		0.1
	rial back into the DIN pool			
	Fraction of dead zooplankton (0.0-1.0) that			
γ_2 (dimensionless)	goes into the sediments or higher trophic	0.5		0.4
	levels			
$(mmol N m^{-3})^{-1}$	Rate at which saturation is achieved with in-	0.75.0.052		0 2 0 014
	creasing food levels	0.75 0.055		0.2.0.014
\mathbf{D} (1-1)	Maximum ingestion rate of phytoplankton	2.5		3.5
n _m (d)	Zooplankton maximal grazing rate	2.3		5.5
(1-1)	Loss rate of zooplankton due to predation	0.05		0.02
	and physiological death	0.05		0.02
$\mathbf{p}^{\mathbf{C}}$ (d^{-1})	Maximum value of the carbon-specific rate		2.0	2.5
r _{max} (u)	of photosynthesis		2.0	5.5
$Q_m \text{ (mmol N (mg C)}^{-1})$	Maximum value of the cellular nutrient		0.2.0.028	0.45 <u>0.032</u>
	quota		0.20.028	
$A^{\rm N}$ (mg Chl (mmol N) ⁻¹)	Maximum value of the chlorophyll : phyto-		0 3 4 2	0 15 2 1
v_{max} (ing Cin (initio N))	plankton nitrogen ratio		0.5 4. 2	0.13 2.1
$R_{C} = R_{n} = R_{chl} (d^{-1})$	Respiration rates		0-0.02	0.02

Table 1. List of parameter values, including units and definitions for the calibrated S2P3-NPZ, S2P3-Photoacclim, and S2P3 v8.0 models.

4.4 Model calibration

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For this study approximately two years of phytoplankton chlorophyll data were available for the CCS location, while in the case of the zooplankton biomass observations, these were collected only during certain days per year allowing only a discrete representation of the seasonal cycle of zooplankton. Finally, profiles of physiological data were only collected during summertime of the years 2003 and 2005. The calibrated version of the S2P3-NPZ model shows differences in the timing of the spring phytoplankton bloom for the year 2015 in comparison to observations of surface chlorophyll (Figure 3a), with a later bloom from the S2P3-NPZ model, reaching a peak bloom about a month later. Additionally, the magnitude of the spring phytoplankton bloom is

210 also higher in the model in comparison to observations. Phytoplankton are able to escape grazing control in April and early May, with the spring zooplankton bloom occurring about a month later (Figure 3b). Similar differences can be observed between the calibrated S2P3-Photoacclim model, demonstrating that constraining the timing of the spring phytoplankton bloom is a complex process. Tuning model parameters to generate earlier blooms modifies the magnitude of the spring phytoplankton bloom by increasing it to unrealistic levels. Figure 3 shows that between the three models the best agreement found in comparison to

- buoy observations correspond to the S2P3 v8.0 model, although the timing and magnitude of the spring phytoplankton bloom show some remaining small differences, with higher concentrations of surface chlorophyll-*a* during spring ($\sim 10 \text{ mg Chl m}^{-3}$). Moreover, the timing of the spring phytoplankton bloom during the year 2014 matches the observations but a delayed bloom is shown during the year 2015. On the other hand, zooplankton bloom happening approximately a month later than the spring phy-
- 220 toplankton bloom during the year 2014, but this difference is less during the year 2015, with the start of the zooplankton bloom happening about half a month after the start of the spring phytoplankton bloom. Remaining mismatches between the calibrated models and observations may be driven by water column processes including advection and diffusion that were not considered in these 1-D models, but affects the real water column where the observations were taken. Furthermore, photo-acclimation and grazing depend on changes in temperature (Geider, 1987; Vázquez-Domínguez et al., 2013) and other ecosystem processes
- 225 which are not explicitly represented, potentially explaining remaining mismatches between the observations and the calibrated <u>S2P3 v8.0 model.</u>

For the S2P3-NPZ and To provide a quantitative index of bloom timing we consider a threshold criteria (Siegel et al., 2002; Greve et al., 200). In this study, the spring phytoplankton bloom is defined as when surface chlorophyll reaches more than $1.5 \text{ mg Chl m}^{-3}$ (see

- 230 Table B2). For the S2P3-Photoacclim models, CTD observations show that during spring the models have model in comparison to CTD cast during spring (Figure 4b), the model has not yet reached the spring phytoplankton bloom whereas observations indicate this is more advanced (Figure 4b), therefore, in the models there are very the spring phytoplankton bloom has already started; the S2P3-Photoacclim model shows low chlorophyll-*a* concentrations at the surfaceand inorganic nutrients have not yet been, with DIN concentrations not being depleted at this stage (Figure 4c). Similar results can be seen for the S2P3-NPZ.
- 235 although Table B2 shows that the spring phytoplankton bloom has already started in this model, but it is still later than in the observations. The S2P3 v8.0 model, on the contrary, shows a good better agreement in terms of the timing of the spring phytoplankton bloom with the CTD observations. During summer months, the models are able to reproduce the sub-surface mixed layer observed in the CTD profile (Figure 4e), with a similar magnitude but shallower in the case of the S2P3-NPZ and the S2P3 v8.0 model approximately 20 m. On the other hand, the physical structure of the model shows a good agreement with
- 240 the observations during spring (Figure 4a), but there are differences during summer (Figure 4d). The lack of a marked mixed layer depth in all the models is likely related to short term meteorology forcing: during the 24/07/2015, the air temperature was high ($\sim 20 \,^{\circ}C$) and wind speed was low ($\sim 5 \,\mathrm{m} \, s^{-1}$). However, the thermocline shows a sharp development in the CTD observations that can not be constrained better in the models by parameterising values of the turbulent closure scheme, light attenuation in the water column, and mixing control parameters (data not shown).

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Moreover, the The S2P3-Photoacclim and the S2P3 v8.0 models were also evaluated further compared with phytoplankton physiological variability observations (Figure 5), both showing. Figure 5 shows that near the sea surface, where light levels are high, photo-acclimation of phytoplankton, values of P_{max}^{Chl} and E_k are higher than in deeper layers of the water column.

The observed P_{max}^{Chl} ranges approximately $0.5 \cdot 2.5 \times 10^{-3}$ (mg C (mg Chl – a)⁻¹ s⁻¹) in the surface waters (first 5m), while this range is smaller in deeper layers ($0 \cdot 1.0 \times 10^{-3}$ mg C (mg Chl – a)⁻¹ s⁻¹), e.g. at 40m depth. Similar variability can be observed with E_k, having lower values in deeper layers of the water column, but the largest variability occurs in the surface layer ($\sim 100 - 250 \,\mu\text{Em}^{-2} \,\text{s}^{-1}$). The S2P3-Photoacclim and S2P3 v8.0 models show a good agreement with the observations, with plausible values of P_{max}^{chl} and E_k found through the water column particularly in terms of the magnitude of the vertical gradients. However, the version of the S2P3 v8.0 model which has the best parameterisations when tuned to fit the timing of

- 255 the spring bloom has remaining discrepancies when compared to the physiological observations, showing an overestimation of P_{max}^{Chl} and E_k at depth. Calibration of these models against physiological data is are relatively novel as such comparisons remain rare. Greater complexity allowed the S2P3 v8.0 model to resolve a more diverse range of biogeochemical dynamics, explicitly accounting for zooplankton biomass and for the dynamics of internal quotas of phytoplanktonic cells, with phytoplankton biomass being in carbon, nitrogen, and chlorophyll currencies, allowing the decoupling of nutrient uptake from carbon fixation
- 260 (Klausmeier et al., 2004; Flynn, 2008; Bougaran et al., 2010; Bernard, 2011; Mairet et al., 2011; Ayata et al., 2013). Including additional parameters in models can add more unconstrained degrees of freedom (Ward et al., 2010), but also allows for more parameter combinations and, therefore, more flexibility to constrain S2P3 v8.0 in order to reproduce observations. The new added parameters and variables in the S2P3 v8.0 model, were carefully chosen to allow the model to be constrained against additional data, specifically zooplankton abundances and photosynthetic physiological measurements, therefore, a higher com-
- 265 plexity allowed better representations of the temporal dynamics. Additionally, despite having more sophisticated formulations of the ecosystem, the S2P3 v8.0 model continues to be a 1-D model, allowing multiple experiments to be run at the same time with relatively low computational cost.



Figure 3. (a) SSB observations of surface chlorophyll-*a* (black line), along with the modelled surface chlorophyll-*a* for the S2P3-NPZ (red line), S2P3-Photoacclim (cyan line), and S2P3 v8.0 (green line) calibrated models. (b) Observations of zooplankton biomass presented as discrete points for day nets (red dots) and night nets (blue dots) taken during the cruises DY026, DY018, DY029, and DY033; modelled zooplankton biomass from the S2P3-NPZ (red line) and S2P3 v8.0 (green line) calibrated models.



Figure 4. CTD observations from the SSB programme (black lines) including data for: springtime (20/04/2015) (a) temperature, (b) chlorophyll-*a*, and (c) DIN (black dots); for summertime (24/07/2015) (d) temperature, (e) chlorophyll-*a*, and (f) DIN (black dots) along the S2P3-NPZ (red line), S2P3-Photoacclim (cyan line), and S2P3 v8.0 (green line) calibrated models.



Figure 5. Observations from the cruises CD173 and JR98 for: (a) chlorophyll-a specific maximum light-saturated photosynthesis rate (P_{max}^{Chl}) in different locations of the Celtic Sea and for the calibrated S2P3-Photoacclim model (cyan lines) and the S2P3 v8.0 model (green lines); (b) observations of the light saturation parameter (E_k) for different stations across the Celtic Sea and for the calibrated S2P3-Photoacclim model (cyan lines) and the S2P3 v8.0 model (green lines). The data from both models were plotted for the same days that the observations were collected.

The ealibrated version behaviour of the S2P3 v8.0 model calibrated for the CCS location is displayed in Figures 6 and 7. Figure 6 shows contour plots from daily profiles of the S2P3 v8.0 model for temperature (Figure 6a), phytoplankton chlorophyll-a (Figure 6b), zooplankton biomass (Figure 6c), phytoplankton chlorophyll : phytoplankton carbon ratio (Figure 6d), and DIN (Figure 6e) for the years 2014 and 2015 which correspond to the observation period of the SSB programme. Figure 6 allows a more detailed overview of the model dynamics, with yellow colours representing high values of each variable and blue colours representing the lowest values. Water column temperature increases from April of each year, reaching a maximum value at the surface during summer months. At the same time, the spring phytoplankton bloom can be observed during April, reaching ~ 10 mg Chl m⁻³. Additionally, the spring zooplankton bloom can be observed approximately a month after the spring phytoplankton bloom is developed, with zooplankton being able to grow during summer months and decreasing until a minimum value during winter. These spring blooms also mark the start of DIN depletion at the surface, a state that lasts until the end of summer. Finally, the phytoplankton chlorophyll : phytoplankton carbon ratio shows the highest values during winter months

- 280 when irradiance levels are low, with the Chl : N ratio decreasing until the end of summer due to the lower concentrations of chlorophyll in the cell to avoid internal damage due to high irradiance during this period, highlighting the photo-acclimation of phytoplankton and flexible stoichiometry of the S2P3 v8.0 model.
- Figure 7 provides a general overview of the dynamics of the full-calibrated S2P3 v8.0 model, showing daily time-series from
 representing the inter-annual variability of each variable using the median (black lines), and lower and upper quantiles (red lines; 95% of the data distribution) during 1965 to 2015 for the calibrated version of this model. 2015. Figure 7a shows the start of thermal stratification during early spring with observable inter-annual variability in the extent of stratification. Once the water column is stratified, the spring phytoplankton bloom can be observed (Figure 7b), followed by the start of the spring zoo-plankton bloom (Figure 7c). As phytoplankton grows, DIN concentrations at the surface start to deplete reaching a minimum value during spring and summer months, and increasing when thermal stratification breaks down during winter months (Figure 7d). Finally, net primary production (NPP) time-series show seasonal and inter-annual phytoplankton dynamics (Figure 7e).
- All variables of the S2P3 v8.0 model shown in Figure 7 provide the spectrum of inter-annual variability of 95% of the data using the upper and lower quantile values (red lines), demonstrating that the inter-annual variability of thermal stratification provides variability for the timing and magnitude of the spring phytoplankton bloom as well as summer growth.

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Figure 6. Contoured daily vertical profiles for the start of 2014 to the end of 2015 for the calibrated version of the S2P3 v8.0 model including: (a) temperature (°C), (b) phytoplankton chlorophyll-a (mg Chl m⁻³), (c) zooplankton biomass (mmol N m⁻³), (d) phytoplankton chlorophyll : phytoplankton carbon ratio (mg Chl (mg C)⁻¹), and (e) DIN (mmol N m⁻³).



Figure 7. Annual representation of the median (black lines) calculated from 1965-2015 for the S2P3 v8.0 model calibrated for the CCS location and forced with all meteorological components (i.e. wind speed, cloud coverage, air temperature, and relative humidity). Red lines represent the annual lower and upper quantiles for each variable of the model (95% distribution of the data) over 1965-2015. (a) Surface temperature minus bottom temperature (°C), (b) surface chlorophyll-a (mg Chl m⁻³), (c) surface zooplankton biomass (mmol N m⁻³), (d) surface DIN (mmol N m⁻³), and (e) net primary production (NPP) (mg C m⁻² day⁻¹).

5 Sensitivity studies

An analysis was performed to assess the sensitivity of the model to selected parameter values for the S2P3 v8.0 model. Each parameter listed in Table 1 was varied in turn in order to understand how sensitive the model is to those changes and the effect that they have on the modelled ecosystem dynamics at the CCS location. In each case parameters were varied from the best calibrated value by +50% and -50%. Sensitivity studies are important tools to improve the accuracy of shelf sea models (Chen et al., 2013), but developing these analyses has to be done carefully in order to identify which processes are responsible for the observed model behaviour (Ward et al., 2013). A direct comparison was calculated in terms of the S2P3 v8.0 model attributes considering: the timing and magnitude of the spring phytoplankton bloom, and the total annual zooplankton biomass for the calibrated version of the model and each experiment, providing better insights on the effects that each parameter produces in the behaviour of each model. Table 2 only shows the experiments involving the maximum ingestion rate of phytoplankton

- 305 in the behaviour of each model. Table 2 only shows the experiments involving the maximum ingestion rate of phytoplankton zooplankton maximal grazing rate (R_m), mortality of zooplankton (m), and the maximum value of the Chl : N ratio (θ_{max}^N), as they were demonstrated to be the most significant parameters in terms of the sensitivity of the model according to the attributes calculated, with the rest of the experiments omitted for this discussion. The S2P3 v8.0 model is strongly influenced by NPZ parameters, with the maximum ingestion rate of phytoplankton zooplankton maximal grazing rate (R_m) (Stegert et al., 2007)
- 310 and zooplankton mortality rate (m) having the largest effect in the magnitude of the spring phytoplankton bloom (Figures A1a, A4aC1a, C4a) and in the total annual zooplankton biomass (Figures A1b, A4bC1b, C4b). The effects of R_m implies that lower values in the maximum ingestion rate of phytoplankton can produce earlier and larger spring phytoplankton blooms compared to the calibrated S2P3 v8.0 model. On the other hand, a higher value of R_m shows a delayed spring phytoplankton bloom (Figure A2C2) compared to the CTD observations. Additionally, zooplankton mortality produced differences in the timing of
- 315 the spring phytoplankton bloom, with delays of 30 days (year 2014) and 35 days (year 2015) when there is less zooplankton mortality, affecting the timing and magnitude of the spring zooplankton bloom, therefore, generating low values of surface chlorophyll-*a* during spring (Figure A5C5). It is well known that zooplankton are key players in the biogeochemical cycling of carbon and nutrients in marine ecosystems (Beaugrand and Kirby, 2010; Beaugrand et al., 2010), influencing the export of organic matter to the deep ocean (González et al., 2009; Juul-Pedersen et al., 2010). Additionally, grazing responses comprise
- the dominant losses for phytoplankton in the ocean (Banse, 1994), influencing plankton stocks and primary production (Franks et al., 1986).

Table 2 presents differences for the S2P3 v8.0 model calibrated for the CCS location and for selected sensitivity experiments in terms of: the timing of the spring phytoplankton bloom (days), defined as in Table B2, using a threshold for phytoplankton

325 biomass (>1.5 mg Chl m⁻³); the magnitude of the spring phytoplankton bloom (mg Chl m⁻³); and the total annual zooplankton biomass (g DW m⁻²). Representation of phytoplankton physiology had an important influence on the timing of the spring phytoplankton bloom (Table 2), with $\theta_{\text{max}}^{\text{N}}$ affecting the S2P3 v8.0 model the most in terms of this attribute of the model structure, showing less productive and delayed spring blooms when $\theta_{\text{max}}^{\text{N}}$ is lower . The (Figures C7a, C8b). Changes in the timing and magnitude of the spring zooplankton bloom coincides with the timing changes of the timing and magnitude of the phyto-

- plankton blooms Ayata et al. (2013) (Figure C7b). These changes in the plankton communities over the year due to different values of θ^N_{max}, also have an effect on the values of DIN (Figure C8c,f), with the largest differences shown during springtime at the surface (Figure C8c). These differences agree with the results found by Ayata et al. (2013), where it was demonstrated that taking into account photoacclimation and variable stoichiometry of phytoplankton growth in marine ecosystem models, produce qualitative and quantitative differences in phytoplankton dynamics. Moreover, these quota formulations in S2P3 v8.0
 were compared to the available dataset of physiological observations (Figure A9C9). It is interesting to note that the sensitivity analysis of NPZ parameters produced differences in the physiological variables P^{chl}_{max} and E_k, specially at the surface (Figures A3, A6C3, C6), suggesting that the predator-prey interactions are indirectly influencing phytoplankton physiology, presumably through feedbacks whereby zooplankton mediated nutrient cyclingand between zooplankton and the nutrient cycling, which subsequently have an effect on phytoplankton physiology due to the dependency of nutrient quotas to the availability of
- 340 <u>inorganic nutrients</u>. A key point from this study showed that the greater variety of data used in a model, the more constrained the parameter choices seem to get.

Experiments	Years	Timing spring phytoplankton bloom (date)	Magnitudespringphytoplanktonbloom (mg Chl m ⁻³)	Total annual zooplankton biomass $(g DW m^{-2})$	
Calibrated S2D2 vg 0	2014	3 rd April	137.7	6110	
Calibrated S2F5 V8.0	2015	9 th April	172.6	6094	
aN	2014	19 th May	103.5	3055	
$\theta_{\max}^{-}\downarrow$	2015	21 st May	103.1	3514	
$ heta_{ m max}^{ m N}$ \uparrow	2014	25 th March	110.2	6733	
	2015	$27^{ m th}$ March	211.5	6797	
	2014	29 th March	292.5	6840	
K _m ↓	2015	3 rd April	359.2	6812	
	2014	14 th April	61.4	5283	
$R_{m}\uparrow$	2015	25^{th} April	159.5	4992	
m↓	2014	3 rd May	238	8362	
	2015	14^{th} May	203.5	8369	
$m\uparrow$	2014	31 st March	104.4	4311	
	2015	6 th April	134.5	4368	
Table 2. List of the most sensitive experiments run for the S2P3 v8.0 model calibrated for the CCS location, including the year of observa-					

tions, timing and magnitude of the spring phytoplankton bloom, and total annual zooplankton biomass values.

6 Comparison of overall model performance

The four calibrated models were further tested \$2P3 v7.0 (Sharples et al., 2006; Marsh et al., 2015), \$2P3-NPZ, \$2P3-Photoacclim,
and \$2P3 v8.0 models were calibrated for the CCS location and further analysis was undertaken by running each of the extended
period-model for a extended period (1965 - 2015), to evaluate the statistics of productivity, partitioned between spring and summer. Table 3 shows that for the \$2P3 v7.0 model, on average, 69.2% of the annual phytoplankton production occurs during the spring phytoplankton bloom. On the other hand, for the \$2P3-NPZ model, on average, only 37.8% of the annual production occurs during spring months, showing that the predator-prey relationship has a strong influence on the magnitude of the spring

- 350 phytoplankton bloom every year. Additionally, the S2P3-Photoacclim model shows a very strong spring phytoplankton bloom, corresponding to 97.490% of the total annual NPP. Finally, for the S2P3 v8.0 model, only 67.4% of the annual production corresponds to the spring bloom period. It is clear that, on average, the least productive model overall was S2P3 v8.0, followed by the S2P3-NPZ, S2P3 v7.0, and S2P3-Photoacclim. This shows the impact and complexity that the predator-prey relationship has on the model dynamics, with the addition of explicit zooplankton and their grazing activity as one of the main losses of
- 355 phytoplankton (Franks et al., 1986). On the other hand, the S2P3 v7.0 model has, on average, more total annual NPP than the S2P3-NPZ model, suggesting that the influence of a constant grazing rate is not as strong in comparison to the one provided by the zooplankton grazing (NPZ framework), because the predator-prey relationship can not be entirely represented in the S2P3 v7.0 model. Additionally, S2P3-Photoacclim had, on average, the highest total annual NPP, demonstrating that the phenotypic responses of phytoplankton to changes in the environment and the addition of variable stoichiometric ratios of N : C : Chl,
- 360 allows for more efficient phytoplankton, which use the available resources to grow and fix carbon more efficiently.

Model	Characteristic (Units)	Mean	Maximum	Minimum	STD
S2D27 0	Total spring NPP ($g C m^{-2} yr^{-1}$)	39.6	54.7	33.5	5.8
52P3 V7.0	Total summer NPP ($g C m^{-2} yr^{-1}$)	17.2	22.9	56.9	5.1
	Total annual NPP ($g \ C \ m^{-2} \ yr^{-1}$)	57.1	61.1	53.1	1.8
CADA NDZ	Total spring NPP (g $C m^{-2} yr^{-1}$)	21.1	49.3	7.2	9.8
52P3-NPZ	Total summer NPP ($g C m^{-2} yr^{-1}$)	28.2	39.7	6.8	7.7
	Total annual NPP ($g \ C \ m^{-2} \ yr^{-1}$)	55.7	60.0	52.7	1.5
S2P3-Photoacclim	Total spring NPP ($g \ C \ m^{-2} \ yr^{-1}$)	66.9 <u>35.</u> 6	76.3 40.7	59.9 25.7	3.2.4.8
	Total summer NPP ($g C m^{-2} yr^{-1}$)	1.1-3.8	7.5-<u>11.4</u>	0.8 - <u>0.6</u>	1.1.4.2
	Total annual NPP ($g \ C \ m^{-2} \ yr^{-1}$)	68.6 - <u>39.4</u>	77.6 42.4	61.1-37.0	3.11<u>.7</u>
C2D2 0.0	Total spring NPP ($g \ C \ m^{-2} \ yr^{-1}$)	25.5	39.5	14.5	3.7
S2P3 v8.0	Total summer NPP ($g C m^{-2} yr^{-1}$)	10.3	12.9	6.8	1.1
	Total annual NPP (g $C m^{-2} ur^{-1}$)	37.8	47.8	33.4	2.2

of the total spring NPP, total summer NPP, and total annual NPP calculated from 1965 to 2015, including the mean, maximum, minimum, and STD values.

7 Conclusions

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This study demonstrates that the combination of an NPZ framework, photo-acclimation, and flexible stoichiometry of phytoplankton in one model produces a better representation of the ecosystem based on the comparison to observations. This combined framework offers a novel and innovative improvement to the S2P3 v7.0 model, for application to the CCS location and more broadly within shelf sea systems. The model validation, using both zooplankton biomass and physiological rates of phytoplankton observations are rarely found in the literature, providing a novel contribution to the marine biogeochemistry modelling field of shelf seas. The development of the S2P3 v8.0 model provides a better fit to observations in comparison to the S2P3 v7.0, S2P3-NPZ, and S2P3-Photoacclim models. Improved confidence in the S2P3 v8.0 model thus suggest improved insights in studies about the effects of physical forcing through tides (Sharples, 2008), intra and inter-annual variations in meterology (Sharples et al., 2006) and other drivers on PP, and phytoplankton dynamics would be possible.

Appropriate parameterisations to represent shelf seas is a subject that should be further supported by fieldwork campaigns and future work should aim to include additional datasets with longer time-series (Friedrichs et al., 2007; Ward et al., 2010). For this study, constraining the seasonal cycle of the phytoplankton physiology is not possible due to the lack of physiological

observations during other periods of the year, furthermore, phytoplankton and zooplankton biomass datasets only include the years 2014 and 2015 but longer time-series would help to improve the model calibration. Many model parameters quantities are poorly constrained observationally mainly due to the fact that model state variables are highly integrated pools, which are affected by biotic and abiotic factors in the environment, making them difficult to be determined by in situ measurements (Fennel et al., 2001).

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The tuning and sensitivity analysis performed in this work allows a better understanding of the ecosystem dynamics represented in the model and how it is influenced by each parameter. By considering the timing and magnitude of the spring phytoplankton bloom, the annual zooplankton biomass, and summer phytoplankton photosynthetic physiology as features to be calculated, a quantitative and clearer comparison between each model could be developed. Finally, the model calibration will never be 385 in perfect agreement with all the observations, particularly in this case, where only one type of phytoplankton and zooplankton were considered. Thus responses must represent some typical or average dynamics and cannot represent any effects of competition between types. Additionally no stages of zooplankton growth were taken into account, which might affect the predator-prey interactions (Wroblewski, 1982; Fennel, 2001). Despite such potential limitations, the S2P3 v8.0 model is able to reasonably represent the integrated behaviour of the mixture of species that inhabit the NW European Shelf Sea. 390

Code and data availability. S2P3 v8.0

The S2P3 v8.0 framework, comprising source code along with example scripts and output, is available online from :current version of model is available from https://doi.org/10.5281/zenodo.3600467 under the Creative Commons Attribution 4.0 International license. The exact version of the model used to produce the results used in this paper is archived on Zenodo

395 (https://zenodo.org), as are input data and scripts to run the model and produce the plots for all the simulations presented in this paper.

Unzipped and uncompressed, the directory /s2p3v8.0 contains several sub-directories:

- 400 - /main contains the source code, s2p3 v8.f90, which is compiled "stand-alone", and executed using accompanying scripts.
 - /domain contains, location (latitude and longitude), tidal components, and bathymetry (total depth) for the Central Celtic Sea in the western English Channel (s12_m2_s2_n2_h_tim.dat).
 - /met contains 2000 2015 meteorological forcing (Falmouth_met_i143_j17_2000 2015 ASCII file).
- 405 - /output contains example output data from experiments that show the calibrated version of each model (S2P3v8, S2P3-Photoacclim, and S2P3-NPZ).

 - /plotting contains MATLAB scripts for plotting time-series data from experiments S2P3v8, S2P3-Photoacclim, and S2P3-NPZ. This folder also include all the dataset from observations used to validate each model.

The ancillary files needed for simulations are available on request from the author (e-mail aab1g15@soton.ac.uk). 410

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

Appendix A: Model equations

The S2P3 v8.0 model can be divided into two different components: a physical part and a biological part. The physical model has been previously described in other studies (Sharples et al., 2006; Marsh et al., 2015) and it will not be described again here.

420 The biological part of the S2P3 v8.0 model calculates changes in phytoplankton carbon biomass (*Phyto_C*) over time as:

$$\frac{\partial Phyto_C}{\partial t} = \frac{\partial}{\partial z} \left(K_Z \frac{\partial Phyto_C}{\partial z} \right) + Phyto_C(\mu - R_C T_{func} - u\zeta) - I \frac{Z}{Q_P}.$$
(A1)

Phytoplankton biomass is also modelled in terms of internal nitrogen, Phyto_N. S2P3 v8.0 calculates Phyto_N as:

$$\frac{\partial Phyto_N}{\partial t} = \frac{\partial}{\partial z} \left(K_Z \frac{\partial Phyto_N}{\partial z} \right) + uPhyto_C - Phyto_N (R_n T_{func}) - IZ.$$
(A2)

The rate of change of phytoplankton biomass in terms of chlorophyll (*Phyto_{cbl}*) is described by:

$$425 \quad \frac{\partial Phyto_{chl}}{\partial t} = \frac{\partial}{\partial z} \left(K_Z \frac{\partial Phyto_{chl}}{\partial z} \right) + u\rho_{chl}Phyto_C - Phyto_{chl}(R_{chl}T_{func}) - I\frac{Z}{Q_z}$$
(A3)

The change in time of external DIN is calculated as:

$$\frac{\partial N}{\partial t} = \frac{\partial}{\partial z} \left(K_Z \frac{\partial N}{\partial z} \right) + \gamma_1 IZ + \gamma_2 mZ + Phyto_N (R_n T_{func}) - uPhyto_C.$$
(A4)

Zooplankton grazing on phytoplankton response depends on a Holling Type II or Ivlev grazing (Franks, 2002), with the ingestion rate of zooplankton (I) described as:

430 $I = R_m (1 - e^{(-\lambda Phyto_N)}),$

where R_m is the zooplankton maximal grazing rate (d⁻¹) and λ is the rate at which saturation is achieved with increasing food levels (mmol N m⁻³)⁻¹.

Zooplankton biomass is, therefore, modelled as:

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$$\frac{\partial Z}{\partial t} = \frac{\partial}{\partial z} \left(K_Z \frac{\partial Z}{\partial z} \right) + (1 - \gamma_1) I Z - m Z.$$
(A5)

The carbon-specific, light-saturated photosynthetic rate depends on the internal nitrogen of phytoplankton described as an f-ratio (f) based on the work of Moore et al. (2001):

$$P_m = P_{max}^C f, \tag{A6}$$

where

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$$f = \frac{Q - Q_{min}}{Q_m - Q_{min}}.$$
(A7)

Nitrogen assimilation is calculated as a Michaelis-Menten function based on the development of the f-ratio:

$$u = u_m \left(\frac{1-f}{1.015-f}\right) \left(\frac{N}{k_n + N}\right),\tag{A8}$$

The carbon-specific photosynthesis is a saturating function of irradiance and it is calculated as:

$$\mu = P_m (1 - e^{-\left(\frac{\alpha^{chl}I_{PAR}\theta}{P_m}\right)}).$$
(A9)

445 Finally, chlorophyll-*a* synthesis depends on the rates of photosynthesis and light absorption

$$\rho_{chl} = \theta_{max}^{N} \left(\frac{\mu}{\alpha^{chl} I_{PAR} \theta} \right). \tag{A10}$$

Note that in this model, the following condition is met: $R_C = R_N = R_{Chl} = R_{ref}T_{function}$, where R_{ref} (d⁻¹) is a degradation rate constant at a reference temperature.

Variables	Definition	Units
IPAR	Photosynthetically available radiation	Wm ⁻²
K_{z}	Depth-dependant vertical eddy diffusivity	$m^2 s^{-1}$
\widetilde{N}	External DIN concentration	$\underbrace{\mathrm{mmol}\mathrm{N}\mathrm{m}^{-3}}_{\overset{\overset{}{}}{}}$
$\underline{Phyto}_{C_{\sim}}$	Phytoplankton biomass in carbon currency	$mg C m^{-3}$
Phytochl~	Phytoplankton biomass in chlorophyll currency	$\underline{\text{mg}} \underline{\text{Chl}} \underline{\text{m}}^{-3}$
$\underline{Phyto}_{N_{\sim}}$	Phytoplankton biomass in nitrogen currency	$\underline{\text{mmol N m}^{-3}}$
Z_{\sim}	Zooplankton biomass in nitrogen currency	$\underline{\text{mmol N m}^{-3}}$
I_{\sim}	Ingestion rate of phytoplankton	d_{\sim}^{-1}
Pure	Carbon-specific rate of photosynthesis	d_{\sim}^{-1}
\underbrace{u}_{\sim}	Phytoplankton carbon-specific nitrate uptake rate	$\frac{\text{mmol N}(\text{mg C})^{-1} \text{d}^{-1}}{\text{mmol N}(\text{mg C})^{-1} \text{d}^{-1}}$
$Q_{P_{c}}$	Cellular nutrient quota (N:C)	$\underline{\text{mmol } N (\text{mg } C)^{-1}}$
Q	Internal nitrogen : phytoplankton chlorophyll ratio	$\underline{\text{mmol N}(\text{mg Chl})^{-1}}$
z	Vertical coordinate (positive upwards)	m
θ	Chlorophyll : phytoplankton carbon ratio	$\underline{\operatorname{mg}} \operatorname{Chl} (\underline{\operatorname{mg}} \operatorname{C})^{-1}$
L.	Carbon-specific rate of photosynthesis	d_{\sim}^{-1}
Repl	Chlorophyll synthesis regulation term	$\frac{\text{mg Chl}(\text{mmol N})^{-1}}{\text{mg Chl}(\text{mmol N})^{-1}}$

 Table A1. List of all variables for the biological part of the S2P3 v8.0 model.

Parameters	Definition	Units	Value
k_n	Half-saturation constant for nitrate uptake	$\underbrace{\text{mmol N m}^{-3}}_{\sim\sim}$	0.014
$P^C_{wax_\sim}$	Maximum value of the carbon-specific rate of photosynthesis	d_{\sim}^{-1}	3.5
Qm.	Maximum value of the cellular nutrient quota	$\underline{\text{mmol } N (\text{mg } \text{C})^{-1}}$	0.032
Q _{min}	Minimum value of the cellular nutrient quota	$\underline{\text{mmol } N (\text{mg } \text{C})^{-1}}$	0.0028
RC	Respiration rate constant	d^{-1}	0.02
Richl	Chlorophyll degradation rate constant	d_{\sim}^{-1}	0.02
$R_{n_{\sim}}$	Nitrate remineralisation rate constant	d_{\sim}^{-1}	0.02
$u_{m_{\sim}}$	Maximum value of the phytoplankton carbon-specific nitrate uptake rate	$\underline{\text{mmol } N (\text{mg } \text{C})^{-1} \text{d}^{-1}}$	0.004
α^{chl}	Chlorophyll-specific initial slope of the photosynthesis-light curve	$\underbrace{\underset{m}{\operatorname{mg}} C (\operatorname{mg} Chl)^{-1} d^{-1} (W)}_{{\operatorname{m}}^{-2})^{-1}}$	$\underbrace{1.99 \times 10^{-6}}_{}$
٤	Cost of biosynthesis	$\underline{\operatorname{mg}} C (\underline{\operatorname{mmol}} N)^{-1}$	$\widetilde{\mathbb{Q}}$
θ_{max}^N	Maximum value of the chlorophyll : phytoplankton nitrogen ratio	$\underline{mg} Chl (mmol N)^{-1}$	2.1
Tfunc	Temperature-response function	dimensionless	1.0
<u>m</u>	Loss rate of zooplankton due to predation and physiological death	d_{\sim}^{-1}	0.02
R_{m}	Zooplankton maximal grazing rate	d_{\sim}^{-1}	3.5
r	Grazing inefficiency or 'messy feeding' (0.0-1.0), returns a fraction of grazed material back into the DIN pool	dimensionless	0.1
Xz	Fraction of dead zooplankton (0.0-1.0) that goes into the sediments	dimensionless	0.4
λ_{\sim}	Rate at which saturation is achieved with increasing food levels	$(\underline{\text{mmol } N \text{ m}^{-3}})^{-1}$	0.014

 Increasing food levels

 Table A2. List of parameters, with their respective definitions, units, and initialised values for the biological part of the S2P3 v8.0 model.

450 Appendix B: Model calibration

<u>Cruise name</u>	Date	Latitude (°N)	Longitude (°W)	Depth (m)
DY029	03/04/2015	49.38	8.59	147
<u>DY029</u>	04/04/2015	49.38	8.59	146
<u>DY029</u>	05/04/2015	49.38	8.59	146
DY029	06/04/2015	49.4	8.58	147
DY029	11/04/2015	49.39	8.58	145
<u>DY029</u>	15/04/2015	49.4	8.59	147~
DY029	20/04/2015	49.4	8.6	147
DY029	21/04/2015	49.4	8.62	148~
<u>DY029</u>	25/04/2015	49.4	8.59	148~
DY029	26/04/2015	49.4	8.58	146
DY029	28/04/2015	49.4	8.58	146
DY033	13/07/2015	49.43	8.59	144~
DY033	14/07/2015	49.42	8.54	144
DY033	15/07/2015	49.37	8.61	145
DY033	24/07/2015	49.36	8.62	145
<u>DY033</u>	25/07/2015	49.41	8.59	148~
DY033	29/07/2015	49.42	8.57	147
<u>DY033</u>	30/07/2015	49.4	8.57	148~
DY033	01/08/2015	49.38	8.58	146

 Table B1. List of relevant CTD casts for the CCS location from DY029 and DY033 cruises considering the date, location (latitude and longitude), and depth. CTD casts in red are the ones chosen in this work to validate each model during spring and summer.

Years	Buoy	S2P3-NPZ	S2P3-Photoacclim	S2P3 v8.0
2014	5 th April	14 th April	25 th April	<u>3rd April</u>
2015	16 th March	24 th April	6 th May	9 th April

 Table B2. Quantitative comparison of the timing of the spring phytoplankton bloom between the buoy observations, S2P3-NPZ,

 S2P3-Photoacclim, and S2P3 v8.0 models.

Appendix C: Sensitivity studies



Figure C1. (a) SSB observations of surface chlorophyll-*a* (red line), along with the modelled surface chlorophyll-*a* for the calibrated S2P3 v8.0 model (green line), and experiments $R_m = 1.75$ (black line), and $R_m = 5.25$ (cyan line). (b) Observations of zooplankton biomass presented as discrete points for day nets (red dots) and night nets (blue dots) taken during the cruises DY026, DY018, DY029, and DY033; modelled zooplankton biomass from the calibrated S2P3 v8.0 model (green line) and experiments $R_m = 1.75$ (black line), and $R_m = 5.25$ (cyan line).



Figure C2. CTD observations from the SSB programme (red line) including data for: springtime (20/04/2015) (a) temperature, (b) chlorophyll-*a*, and (c) DIN (red dots); for summertime (24/07/2015) for (d) temperature, (e) chlorophyll-*a*, and (f) DIN (red dots) along the calibrated S2P3 v8.0 model (green line), experiments: $R_m = 1.75$ (black line), and $R_m = 5.25$ (cyan line).



Figure C3. Observations from the cruises CD173 and JR98 in different locations of the Celtic Sea, including the calibrated S2P3 v8.0 model (green lines), experiments $R_m = 1.75$ (black lines), and $R_m = 5.25$ (cyan lines) for: (a) chlorophyll-*a* specific maximum light-saturated photosynthesis rate (P_{max}^{Chl}) and (b) light saturation parameter (E_k).



Figure C4. (a) SSB observations of surface chlorophyll-*a* (red line), along with the modelled surface chlorophyll-*a* for the calibrated S2P3 v8.0 model (green line), and experiments m = 0.01 (black line), and m = 0.03 (cyan line). (b) Observations of zooplankton biomass presented as discrete points for day nets (red dots) and night nets (blue dots) taken during the cruises DY026, DY018, DY029, and DY033; modelled zooplankton biomass from the calibrated S2P3 v8.0 model (green line) and experiments m = 0.01 (black line), and m = 0.03 (cyan line).

Figure C5. CTD observations from the SSB programme (red line) including data for: springtime (20/04/2015) (a) temperature, (b) chlorophyll-*a*, and (c) DIN (red dots); for summertime (24/07/2015) for (d) temperature, (e) chlorophyll-*a*, and (f) DIN (red dots) along the calibrated S2P3 v8.0 model (green line), experiments: m = 0.01 (black line), and m = 0.03 (cyan line).

Figure C6. Observations from the cruises CD173 and JR98 in different locations of the Celtic Sea, including the calibrated S2P3 v8.0 model (green lines), experiments m = 0.01 (black lines), and m = 0.03 (cyan lines) for: (a) chlorophyll-*a* specific maximum light-saturated photosynthesis rate (P_{max}^{Chl}) and (b) light saturation parameter (E_k).

Figure C7. (a) SSB observations of surface chlorophyll-*a* (red line), along with the modelled surface chlorophyll-*a* for the calibrated S2P3 v8.0 model (green line), and experiments $\theta_{\text{max}}^{\text{N}} = 0.075$ (black line), and $\theta_{\text{max}}^{\text{N}} = 0.225$ (cyan line). (b) Observations of zooplankton biomass presented as discrete points for day nets (red dots) and night nets (blue dots) taken during the cruises DY026, DY018, DY029, and DY033; modelled zooplankton biomass from the calibrated S2P3 v8.0 model (green line) and experiments $\theta_{\text{max}}^{\text{N}} = 0.075$ (black line), and $\theta_{\text{max}}^{\text{N}} = 0.225$ (cyan line).

Figure C8. CTD observations from the SSB programme (red line) including data for: springtime (20/04/2015) (a) temperature, (b) chlorophyll-*a*, and (c) DIN (red dots); for summertime (24/07/2015) for (d) temperature, (e) chlorophyll-*a*, and (f) DIN (red dots) along the calibrated S2P3 v8.0 model (green line), experiments $\theta_{max}^{N} = 0.075$ (black line), and $\theta_{max}^{N} = 0.225$ (cyan line).

Figure C9. Observations from the cruises CD173 and JR98 in different locations of the Celtic Sea, including the calibrated S2P3 v8.0 model (green lines), experiments $\theta_{\max}^{N} = 0.075$ (black lines), and $\theta_{\max}^{N} = 0.225$ (cyan lines) for: (a) chlorophyll-*a* specific maximum light-saturated photosynthesis rate (P_{\max}^{Chl}) and (b) light saturation parameter (E_k).

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