New insights into the organic carbon export in the Mediterranean Sea from 3D modeling

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Abstract. The Mediterranean Sea is one of the most oligotrophic regions of the oceans, and nutrients have been shown to limit both phytoplankton and bacterial activities, resulting in a potential major role of dissolved organic carbon (DOC) export in the biological pump. Strong DOC accumulation in surface waters is already well-documented, though measurements of DOC stocks and export flux are still sparse and associated with major uncertainties. This study provides the first basin-scale overview and analysis of organic carbon stocks and export fluxes in the Mediterranean Sea through a modeling approach based on a coupled model combining a mechanistic biogeochemical model (Eco3M-MED) and a high-resolution (eddy-resolving) hydrodynamic simulation (NEMO-MED12). The model is shown to reproduce the main spatial and seasonal biogeochemical characteristics of the Mediterranean Sea. Model estimations of carbon export are also of the same order of magnitude as estimations from in situ observations, and their respective spatial patterns are mutually consistent. Strong differences between the western and eastern basins are evidenced by the model for organic carbon export. Though less oligotrophic than the eastern basin, the western basin only supports 39 % of organic carbon (particulate and dissolved) export. Another major result is that except for the Alboran Sea, the DOC contribution to organic carbon export is higher than that of particulate organic carbon (POC) throughout the Mediterranean Sea, especially in the eastern basin. This paper also investigates the seasonality of DOC and POC exports as well as the differences in the processes involved in DOC and POC exports in the light of intracellular quotas. Finally, according to the model, strong phosphate limitation of both bacteria and phytoplankton growth is one of the main drivers of DOC accumulation and therefore of export.

1 Introduction

The biological pump is recognized as a major component of carbon export in the ocean and plays a significant role in the carbon cycle as a whole (Siegenthaler and Sarmiento, 1993). The sinking of organic particles has long been identified as the main process involved in the biological pump, thereby sustaining the vertical carbon and nutrient gradients in the ocean (Eppley and Peterson, 1979; Sarmiento and Gruber, 2006). Considerable attention has therefore been paid to the export of organic carbon in its particulate form.

Advances in the characterization of dissolved organic pools have led to a better knowledge of the role of the dissolved organic carbon (DOC) compartment in the ocean carbon cycle. As a non-sinking tracer, the fate of DOC is strongly linked to physical processes and its export occurs via vertical mixing and/or downwelling when it lies in intermediate waters, and via oceanic overturning circulation in deep water (Hansell et al., 2002, 2009). If the early works of Copin-Montégut and Avril (1993) in the Mediterranean Sea and Carlson et al. (1994) in the Sargasso Sea were the first attempts to quantify DOC export flux below the euphotic zone, estimation of detrital particulate organic carbon (POC) export began years before with the deployment of sediment traps and isotopics measurements (Buesseler, 1991).

The seasonal variability of DOC in the euphotic zone has been widely recorded in the sub-tropical and temperate areas of the ocean (Carlson et al., 1994; Avril, 2002; Hansell and Carlson, 2001; Santinelli et al., 2013). The results of these studies indicate a time lag between DOC production and consumption, causing summer accumulation in the upper layers due to both biotic and abiotic processes, which either alter DOC bioavailability or reduce bacterial activity. The inefficiency of the microbial loop in organic carbon mineralization - the so-called malfunctioning microbial loop (Thingstad et al., 1997) - induces an accumulation of bioavailable DOC. This inefficiency is directly related to low phosphate availability in the upper waters of the Mediterranean Sea (Moutin et al., 2002; Van Wambeke et al., 2002; Thingstad et al., 2005; Santinelli et al., 2013).

In this paper, our aim is to investigate the pathways of organic carbon (OC) at the scale of the Mediterranean Sea, and more specifically to characterize OC export fluxes since this is crucial to determine the efficiency of the biological pump. High resolution 3D modeling using the biogeo-chemical mechanistic model Eco3M-MED (Alekseenko et al., 2014) forced by the physical model NEMO-MED12 (Beuvier et al., 2012b) was chosen to address this question. Major modeling work has already been done to estimate organic carbon export using box models (e.g. Toggweiler et al., 2003), ocean carbon-cycle models (e.g. Bopp et al., 2001; Sarmiento et al., 1998; Maier-Reimer et al., 1996; Sarmiento and Gruber, 2006) and ecosystem models coupled with hydrodynamic models (e.g. Le Quéré et al., 2010). Several coupled models have also been developed to study the whole of the Mediterranean Sea, starting with the early simulation by Crispi et al. (1998) and Crise et al. (1998). The number of models designed for this purpose is increasing (Lazzari et al., 2013; Mattia et al., 2013; Macías et al., 2014), but to our knowledge, no modeling work has yet focused on or-

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ganic carbon fluxes for the entire Mediterranean Sea. Here, our aim is to focus on OC export in the Mediterranean Sea by characterizing and quantifying the associated fluxes, studying their temporal and spatial variability, and providing the first estimations at this scale of the respective contributions of DOC and POC (which refers to the detrital particulate organic carbon only in the present paper) to carbon export. We also aim to analyze the processes involved in DOC and POC production export in the light of the intracellular quotas of planktonic organisms calculated by Eco3M-MED. The paper is organized as follows: After the introduction (Sec. 1), a succinct overview (Sect. 2) of the hydrodynamical model NEMO-MED12 (Sec. 2.1) and the biogeochemical model Eco3M-MED (Sec. 2.2) is provided, given that they are described in detail in the aforementioned papers. Simulation set-up and datasets used for model comparison are also presented. Sect. 3 first focuses on the results related to organic carbon inventory and export at the scale of the Mediterranean Basin, and for the purpose of discussion results on intracellular quotas in phytoplankton and bacteria as well as on exudation fluxes are also presented. In Sect. 4 results on export are discussed in the context of previous POC and DOC export evaluations in the Mediterranean Sea, and in the light of processes and intracellular quotas in phytoplankton and bacteria. Finally, ~~an appendix~~ a Supplementary Material is associated with this paper containing the assessment of the biogeochemical model outputs (nutrients, chlorophyll, primary production and DOC) through comparison with available data and analysis of the main discrepancies.

2 Material and methods

2.1 The hydrodynamic model

The physical run used in this work is described in Beuvier et al. (2012b). It has been simulated by the regional circulation model NEMO-MED12 Beuvier et al. (2012a) which is part of a suite of Mediterranean regional versions of OPA and NEMO (Madec and The-NEMO-Team, 2008) as OPA-MED16 (Béranger et al., 2005), OPAMED8 (Somot et al., 2006) and NEMO-MED8 (Beuvier et al., 2010).

Model resolution is 1/12 ° ($\approx$ 8 km) which means that most of the mesoscale features are explicitly resolved, and the domain includes the whole of the Mediterranean Sea as well as the Atlantic Ocean West of 11°W (Fig. 2). More details of the model and its parametrization are given in Beuvier et al. (2012a).

The simulation was initiated in October 1958 with temperature and salinity data representative of the 1955–1965 period using the MEDATLAS dataset (MEDAR/MEDATLAS-Group 2002, Rixen et al., 2005). Atmosphere forcings are applied daily and come from the ARPERA dataset (Herrmann and Somot, 2008), a 55-year simulation at 50 km and daily resolutions. SST-relaxation and water-flux correction terms, as well as fresh water input from rivers and the Black Sea and Atlantic exchanges are the same as described in Beuvier et al. (2010, 2012a).
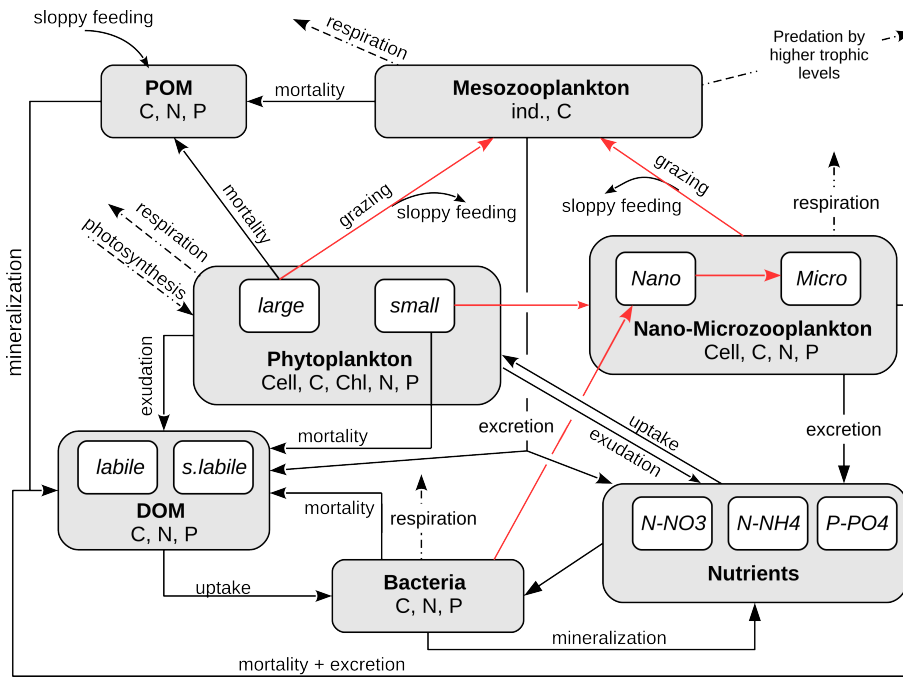
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Figure 1. Conceptual diagram of the biogeochemical model Eco3M-MED. Grey boxes represent major compartments and white boxes sub-compartments. State variables for each sub-compartment are listed at the bottom of compartment boxes. Red arrows indicate grazing processes from the prey to the predator.

2.2 The biogeochemical model

The biogeochemical model Eco3M-MED is embedded in the Eco3M modular numerical tool (Baklouti
95 et al., 2006b), and its structure is similar to the model presented in Alekseenko et al. (2014). Fig.
1 summarizes the interactions between the state variables through the biogeochemical processes.
We chose to represent three different element cycles C, N and P in order to reproduce the different
limitations and co-limitations observed in the Mediterranean Sea. Silicium, potentially limiting in
some regions (Leblanc et al., 2003) is not represented in the model, as P and N limitations are the
100 most common ones in the Mediterranean Sea. Six different planktonic functional types (P.F.T., see
Le Quéré et al. (2005) for a proper definition) are represented : 2 primary producers (phytoplankton),
1 decomposer (heterotrophic bacteria) and 3 consumers (nano-, micro- and meso-zooplanktons). The
structure of the trophic web thereby includes the main P.F.T.s of the Mediterranean Sea (Siokou-
Frangou et al., 2010).
105 Each P.F.T. of the model is represented through several state variables, namely C, N, P (and Chl
for producers) concentrations and a cell number (i.e. an abundance), except for meso-zooplankton
which is only represented through its C concentration and its abundance (in individuals per unit
volume). Intracellular ratios (i.e. the ratio between two elemental concentrations) as well as intra-
cellular quotas (i.e. the quantity of a given element per cell) can therefore be calculated dynamically

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110   by the model. Intracellular ratios are indicators of plankton stoichiometry, i.e. of its C:N:P elemental composition. Early biogeochemical models (NPZD models) have considered a constant C:N:P ratio in plankton given by the canonical Refield ratio of 106:16:1 (Redfield, 1958). Based on Droop's work (e.g. Droop, 1968, 1975), an increasing number of biogeochemical models (e.g. Baretta et al., 1995; Geider et al., 1998) have in recent decades assumed flexible plankton stoichiometry. Though

115   Droop's original quota function relating growth rate to the intracellular quota of the limiting element was based on cell quotas, these biogeochemical models have used intracellular ratios instead of quotas to regulate the rate of biomass synthesis (and other process rates) with quota functions similar to that of Droop. These flexible stoichiometry models have been widely used in the framework of theoretical batch or chemostat studies (e.g. Geider et al., 1998; Baklouti et al., 2006b) or for large-

120   scale studies with ERSEM (Baretta et al., 1995), BFM (Vichi et al., 2007) or others (e.g. Moore et al., 2002) models. In such models, substrate uptake and biomass synthesis are decoupled, but cell division is not explicitly represented.

     Intracellular quotas (or cell quotas) as they are defined in the present paper are indicators of the C, N and P cellular content of plankton. They are an original feature of the Eco3M-MED model in

125   the category of 3D coupled physical-biogeochemical models. This model is based on the assumption that there are a minimum ($Q_X^{\min}$) and a maximum ($Q_X^{\max}$) intracellular content for each element X among (C, N, P). $Q_X^{\min}$ can be interpreted as the amount of element X used in cellular structure and machinery, and the accumulated surplus as storage for future growth (Klausmeier et al., 2008). The variability in cell quotas has indeed been widely evidenced through several experimental and in situ

130   studies (e.g. Brown and Harris, 1978; Fukuda et al., 1998; Lovdal et al., 2008; Heldal et al., 2003; Bertilsson et al., 2003; Wilhelm et al., 2013).

     The use of cell numbers as state variables and of the associated intracellular quotas offers several advantages: firstly, it makes it possible to distinguish between cell division, which is described by a specific equation, see Eq. 1), biomass synthesis, and uptake. Second, intracellular quotas are indica-

135   tive of the actual internal status of cells, i.e. they indicate whether cells are rich or depleted in a given element, while intracellular ratios only provide relative values. In other words, a given value of intracellular ratio Q_{XY} can correspond to several different cell statuses (for example, a given C:N ratio can be obtained with an infinity of pairs of C and N intracellular concentration values). Thus, intracellular ratios can only provide information on the internal relative quantity of X as compared to that

140   of Y, while intracellular quotas inform on intracellular absolute quantities. The latter information is very useful for the analysis of plankton dynamics since it is informative about the nutritional status of each P.F.T. of the trophic web (see the Discussion section). It is also a good proxy of the quality of the prey available for zooplankton (i.e. whether prey are rich or depleted in a given element). Thirdly, the parameters determined at cell level can be used without using conversion factors. For

145   example, uptake rate measured at cell level (Talarmin et al., 2011), or grazing parameters expressed

in number of prey per predator per unit time, such as the ones provided in Christaki et al. (2009) for HNF and ciliates can be used directly.

Intracellular quotas have already been used in previous modeling studies to study phytoplankton growth (Klausmeier et al., 2004) or the dynamics of the planktonic food web (Thingstad et al., 2005). In the latter study, however, cell quotas of carbon were assumed to be fixed in the protozoa, while fixed C:N-ratios were assumed for bacteria and phytoplankton. Moreover, this model was used without being coupled with a physical model (i.e. for the simulation of microcosm and lagrangian experiments).

In the model, the producers are split into two different P.F.T.s according to their theoretical size, i.e. large phytoplankton (> 10 μm) mainly encompassing diatoms, and small phytoplankton (< 10 μm) which includes picophytoplankton and the remaining nanophytoplankton. The two P.F.T.s have different parameters, distinct predators and fuel different detritic pools (Fig. 1). Decomposers are represented by heterotrophic bacteria and are responsible for the organic matter mineralization, including hydrolysis of particles. Zooplankton is divided into three different size groups, heterotrophic nanoflagellate (HNF) which feeds on bacteria and small phytoplankton, ciliate which feeds on small phytoplankton and HNF, and mesozooplankton (copepods) which feeds on ciliate, HNF and large phytoplankton. Copepods are the only metazoans in the model, and mechanisms such as individual growth, egg production or reproduction are implicitly represented (Alekseenko et al., 2014).

The processes used in the model are extensively described in the aforementioned reference. However, for the purposes of the present paper, we recall that POC is fueled by the natural mortality of largest organisms (mesozooplankton, diatoms and ciliates) and by the egestion of fecal pellets and sloppy feeding by mesozooplankton, and consumed by POC hydrolysis to DOC. The DOC pool has many inputs (phytoplankton exudation, zooplankton excretion, mortality of small organisms, POC hydrolysis) and a single output (uptake by bacteria). The formulations of most of the biogeochemical processes, for which details are extensively given in Baklouti et al. (2006a, 2011); Mauriac et al. (2011), and Alekseenko et al. (2014), follow cell level mechanistic considerations. Intracellular ratios (Q_{XY}) and intracellular quotas (Q_X) are used to regulate growth via Droop's quota function (Droop, 1968) and net uptake and grazing rates via Geider's limitation formulation (Geider et al., 1998). For example, the specific growth rate (i.e. the division rate) μ of all unicellulars in the model is given by the following equation:

$$\mu = \mu^{\text{max}} \min_{X \in \{\text{C, N, P}\}} \left(1 - \frac{Q_X^{min}}{Q_X} \right) \tag{1}$$

where μ^{max} is the maximum division rate and $Q_X^{\min}$ the minimum intracellular X quota.

Grazing, primary production and uptake rates are controlled firstly by the organism's environment (either prey or nutrient concentration, or light availibility). Secondly, the internal cell status represented by intracellular quotas and ratios drives a feedback regulation of the net incorporated biomass

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through quota functions. Hence, the uptaken surplus (which becomes more and more significant as the intracellular quota approaches $Q^{\max}$) is either released in its initial form or exuded in the form of DOM. In the same way, excretion and fecal pellet production fluxes are proportional to the grazing flux and to a quota function the value of which increases as the quota approaches $Q^{\max}$. Furthermore, 10 % of the material grazed by mesozooplankton directly fuels the particulate organic matter stock, to represent sloppy feeding. Respiration rates are estimated via energy costs for every plankton activity (Alekseenko et al., 2014). Nitrification is represented through first order kinetics while particulate hydrolysis function depends on bacteria intracellular quotas (POC hydrolysis increases with bacterial C-limitation). Grazing by higher trophic levels is implicitly taken into account via quadratic mortality affecting only mesozooplankton. Grazing function is a Holling II type (Holling, 1959; Kooijman, 2000) for multiple prey. The only difference with the configuration of Alekseenko et al. (2014) lies in the formulation used to represent predator preferences for multiple prey. We here used the "Kill The Winner" (KTW) formulation depicted in Vallina et al. (2014), which combines active-switching (i.e. the preference of a predator for a given prey depends on prey density) and an ingestion rate always increasing with the total biomass of prey. This active-switching formulation was used to preserve foodweb diversity (e.g Prowe et al., 2012) and to prevent unrealistic predator-prey oscillations.

Since the model relies on a mechanistic basis, parameters are mainly physiological (and measurable) and they were either taken from literature or derived from other parameters on the basis of physiological considerations and in the interests of greater consistency between parameters. For example, maximum intracellular quotas are inferred from minimum ones as done in Thingstad et al. (2005). Another example lies in the relationship between the maximum uptake rate of a given element, which is the product of the maximum specific growth rate and the maximum intracellular quota in that element. Other examples as well as the whole set of parameters are given in Alekseenko et al. (2014).

2.3 Model coupling

The models NEMO and Eco3M-MED have been associated for the first time. The coupling between the hydrodynamic and biogeochemical models is offline, i.e. biological retroaction on the physics is not taken into account. Daily-averaged water velocities were used for the advection of biogeochemical tracers, using a MUSCL scheme (horizontal and vertical diffusion fluxes are calculated according to a centered scheme). The time-step used for the numerical integration of the tracer conservation equations equals 1200 s. A sinking velocity of 2 m d^{-1} is applied only on the particulate organic pool (i.e. the detrital compartment). The aim of this compartment is to represent particles with different sizes and sinking velocities and the value of 2 m d^{-1} is within the usual range found in the litterature (Vichi et al., 2007; Fasham et al., 2006). Light attenuation formulation in the water column is based on Morel (1988) results.
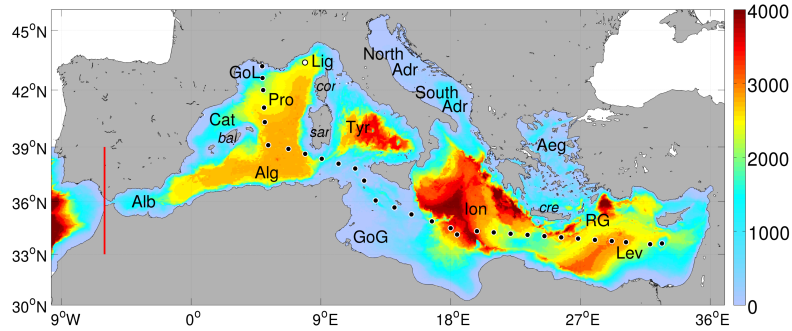
Figure 2. Bathymetry of the grid in meters, black dots represent the BOUM cruise stations (Moutin et al., 2012a) while white dot is DyFaMed position (Marty and Chiavérini, 2010). The area west of the red line constitutes the buffer-zone. Acronyms indicate different sub-basin names and islands (in italic). Terminology is taken from Millot and Taupier-Letage (2005). From west to east, **Alb** stands for Alboran Sea, **Cat** for Catalan Sea, **GoL** for Gulf of Lions, **Pro** for Provencal sub-basin, **Alg** for Algerian basin, **Lig** for Ligurian Sea, **Tyr** for Tyrrhenian Sea, **GoG** for Gulf of Gabes, **North Adr** and **South Adr** for north and south Adriatic Sea respectively, **Ion** for Ionian sub-basin, **Aeg** for Aegean Sea, **Lev** for Levantine sub-basin and **RG** for Rhodes Gyre. Major islands names are also plotted, *bal* stands for the Balearic islands, *sar* for Sardinia, *Cor* for corsica, *cre* for Crete.

2.4 Initial and boundary biogeochemical conditions

Initial nutrient and chlorophyll fields are derived from annual means of Mediterranean Sea climatological data (Schaap and Lowry, 2010). The remaining biogeochemical variables are derived from chlorophyll using conversion factors derived from published works (see Alekseenko et al. (2014) for details).

A "buffer-zone" has been defined between the domain western boundary and the Gibraltar Strait (from 11°W to 6°W), in which a damping procedure towards Atlantic conditions has been applied. The restoring time is 2 days west of 7.5°W, linearly increasing to 90 days from 7.5°W to 6°W (Fig. 2). Atlantic nutrient concentrations come from the World Ocean Atlas monthly climatology (Garcia et al., 2006), so that the nutrients damping in the "buffer-zone" takes into account the nutrients' monthly variability. Given the inaccuracies in phosphate measurements, we decided to compute phosphate profiles from that of nitrate by imposing a redfield ratio of 16 in order to be more consistent with observed $NO_3:PO_4$ ratios in this region (Gómez, 2003). Chlorophyll concentrations were not provided in this database. We therefore used in situ data from the SeaDataNet database to create a mean vertical chlorophyll profile for the Atlantic, and then used a climatology of surface chlorophyll from the GlobColour product in this region to represent an annual cycle of the chlorophyll vertical

profile. The remaining Atlantic biogeochemical variables were derived from chlorophyll using the same procedure as for initial conditions.

235   Nutrient (NO_3 and PO_4) inputs from riverine influx and coastal runoffs are derived from Ludwig et al. (2009), following the same procedure as for the riverine freshwater inputs in the circulation model (Beuvier et al., 2010, 2012b). The nutrient influx of the 29 rivers included in the RivDis database (Vörösmarty et al., 1996) are taken into account in the simulation, while the nutrients of the remaining rivers from the Ludwig et al. (2009) database are averaged for every sub-basin

240   and distributed along their respective sub-basin's coast as coastal runoffs. Dissolved organic carbon inputs in the Mediterranean Sea are distributed in every sub-basin according to the riverine DOC estimates of Ludwig (1996) (a total of $\sim$ 1.8 Tg C y^{-1} in the whole of the Mediterranean Sea). Sub-basin DOC inputs were then distributed among fluvial estuarine and coastal runoffs to match circulation model freshwater geographical distribution (Palmiéri, 2014; Palmiéri et al., in prep).

245   Mass exchanges with the Black Sea in the Dardanelles Strait are treated as river inputs, with nutrients and DOC input concentrations provided by the SESAME project (Tugrul and Besiktepe, 2007; Meador et al., 2010). But, since NO_3 budget indicates a negative net flux of NO_3 the Dardanelles Strait (i.e. exiting from the Mediterranean), NO_3 flux at Dardanelles is set to zero and the outcome is transferred on the Aegean sub-basin's runoffs. These runoffs are artificially reduced in order to keep

250   the riverine budget of NO_3 in the Aegean sub-basin realistic.

2.5   Simulation set-up

Using the biogeochemical initial conditions defined in Sect. 2.4, we have conducted a 5 years simulation using physical forcings from the years 1973-1977. This first simulation was considered as a 'spin-up', in order to reduce the impact of state variables adjustment in the simulations. It has delib-

255   erately been done long enough before the Eastern Mediterranean Transient period (starting around 1991) which is not stable enough to be chosen as a spin-up period. Moreover, due to high computational costs, it was not possible to run this first simulation until the year 1996. We therefore used the final biogeochemical state of this spin-up as initial conditions for a second simulation running from 1996 to 2012. In this second simulation, only the years following 1998 are considered, since the first

260   3 years were treated as an additional spin-up beyond which the stability of the run was ensured (i.e. no drift could be observed).

2.6   Data description

The aim of the present work is to study and to quantify organic carbon export fluxes using a 3D physical-biogeochemical model. For this purpose, our first aim was to assess the reliability of our

265   model by examining the agreement between different model outputs and corresponding available data : chlorophyll, nutrients, DOC concentrations and primary production rates.

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Three type of comparisons were undertaken : (i) at basin scale, using surface chlorophyll fields provided by satellite for comparisons (ii) at basin scale, using BOUM cruise transect as a "snapshot" to compare nutrients and DOC vertical profiles during the stratified period (iii) at a local scale using the time series data collected at DyFaMed station

2.6.1 Chlorophyll data derived from satellite

Among the specificities of the Mediterranean Sea, its strong oligotrophy and the major influence of colored dissolved organic matter, make the use of classical satellite chlorophyll products difficult (e.g. Claustre et al., 2002). Several algorithms have already been developed (Bosc et al., 2004; D'Ortenzio et al., 2002; Volpe et al., 2007), using different satellite reflectances and datasets. Here, we used a daily surface chlorophyll product delivered by the Myocean project (http://www.myocean.eu). In this product, chorophyll concentrations have been derived using the MEDOC4 algorithm developped by Volpe et al. (2007). This algorithm was built using a large dataset of chlorophyll concentrations collected in situ and reflectance measurements from 3 satellites (Seawifs, MERIS and MODIS), constituting a homogeneous series from September 1997 to March 2012.

2.6.2 The BOUM cruise data

The BOUM cruise took place during summer 2008 (from June 16 to July 20) and traversed both the western and eastern basins of the Mediterranean Sea (Moutin et al., 2012a). The data acquired during this cruise provide a unique picture of the biogeochemical status of the Mediterranean Sea since many biogeochemical variables were observed. Measurements of nutrients and DOC concentrations were used to perform a basin-scale comparison during the summer stratified period with the model outputs obtained at the same dates as the cruise, and averaged over this period.

2.6.3 The DyFaMed station data

The DyFaMed station is located in the Ligurian Sea at 7.9°E and 43.4°N (Fig. 2) and is isolated from coastal inputs by the Mediterranean Northern Current. A strong winter mixing is observed in this area, although it is less intensive than the deep convection occuring in the Provencal sub-basin (Marshall and Schott, 1999). Nutrients (Pasqueron de Fommervault et al., 2015), chlorophyll (Marty et al., 2008), dissolved organic carbon (Avril, 2002) and primary production rates (Marty et al., 2008) time series were used for comparison. The comparison of the model outputs with DyFaMed time series can be done through different methods. The simplest consists in using a single grid point which is the nearest to the DyFaMed station location. This implies that the model perfectly reproduces spatial patterns in this region, which is obviously never the case. On the other hand, the use of model outputs averaged on several grid points around the DyFaMed station amounts to dampening signal variability. We finally chose to use the nearest gridpoint to the DyFaMed station, while assessing spatial variability in the 8 neighbouring grid points (Table **??**).
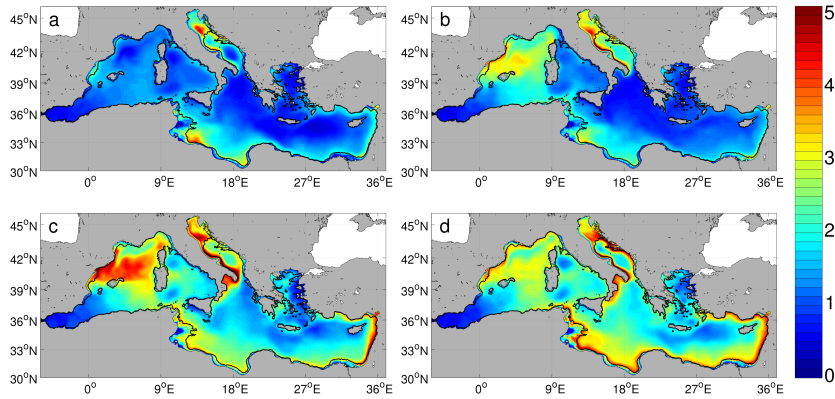
Figure 3. Modeled dissolved organic carbon inventory (mol m^{-2}) integrated over the first 100 m. Maps are averaged over the 2000-2012 period in (a) winter (Dec.-Feb.), (b) spring (Mar.-May), (c) summer (Jun.-Aug.), (d) autumn (Sept.-Nov.).

3 Results

3.1 Organic carbon inventory and export

3.1.1 Dissolved organic carbon inventory

In the following section, mDOC refers to the modeled dissolved organic carbon integrated over the
first 100 m of the water column. Seasonal variations of mDOC are given in Fig. 3. Low mDOC values
(< 1 mol m^{-2}) are observed throughout the year in the Alboran Sea (and up to the Balearic Islands),
the North Levantine basin, and in some well marked structures in the Tyrrhenian Sea. In contrast,
very high mDOC values (up to 5 mol m^{-2}) can be found throughout in the North Adriatic Sea and
along the Lybian Coast. Apart from these regions, mDOC is low everywhere (below 2 mol m^{-2})
in winter (Fig. 3 a), and this is also true in spring except in the region of the spring bloom in the
Provencal sub-basin. In the western basin, highest DOC concentrations are generally observed in
summer, with values reaching 4 mol m^{-2} in the bloom region of the Liguro-Provencal sub-basin. In
the eastern basin, they are reached in autumn and mostly concern the Adriatic Sea, and the regions
along the southern and eastern coasts.

3.1.2 Particulate organic carbon inventory

In what follows, mPOC refers to the modeled particulate organic carbon integrated over the first
100 m of the water column. Seasonal variations of mPOC are given in Fig. 4. Unlike mDOC, mPOC
highest values are observed in winter and spring. This is mostly true for the western basin since, in
the eastern basin, mPOC remains low (< 0.05 mol m^{-2}) all over the year, except for the Adriatic
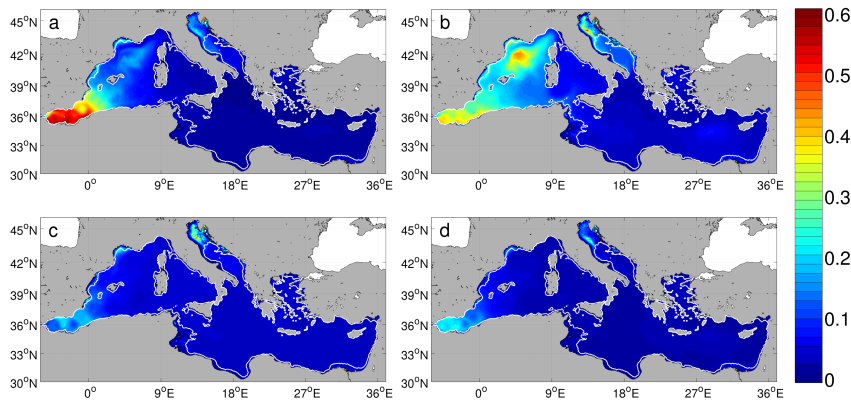
Figure 4. Modeled particulate organic carbon inventory (mol m^{-2}) integrated over the first 100 m. Maps are averaged over the 2000-2012 period in (a) winter (Dec.-Feb.), (b) spring (Mar.-May), (c) summer (Jun.-Aug.), (d) autumn (Sept.-Nov.). White lines are the 0 m and 100 m isolines.

320 Sea and a local maximum in the Rhodes Gyre distinguishable in spring. During winter (Fig. 4 a), the highest values of mPOC (> 0.5 mol m^{-2}) are found in the region of the Alboran Sea and the surrounding Balearic Islands and also in the Liguro-Provencal sub-basin though with much lower concentrations. In the Adriatic Sea, mPOC is in the range [0.1;0.2] mol m^{-2}. Elsewhere, mPOC is low (< 0.2 mol m^{-2}). During spring (Fig. 4 b), the maximum mPOC is observed in the region of the

325 bloom in the Provencal sub-basin (≈ 0.4 mol m^{-2}) and the North Adriatic Sea. During summer and autumn (Fig. 4 c and d), overall values are low (< 0.05 mol m^{-2}), except in the Alboran Sea (where values reach 0.3 mol m^{-2}) and in the North Adriatic Sea.

3.1.3 Dissolved and particulate organic carbon export

Organic carbon fluxes are computed by adding the contribution of advection (vertical velocity and

330 settling velocity for POC) and vertical diffusion (implicitly representing turbulent and convective mixing) fluxes across an horizontal section of the grid. Negative fluxes account for downward fluxes. For clarity, modeled fluxes will be referred to as F_{DOC} , F_{POC} and F_{OC} as the sum of the latter two. F_{DOC} and F_{POC} have been computed at 100 m and 200 m so as to include most of the productive layer and to allow the comparison in space and time between regions. These depths are also used in

335 several other modeling studies (Lévy et al., 1998; Bopp et al., 2001).

The yearly amount of mOC export at 100 m is equal to 48.4 MtC y^{-1}. The eastern basin is the main contributor to this export with a total export of 28.7 against 19.7 MtC y^{-1} for the western basin. mDOC export is equal to 38.8 MtC y^{-1}, and comparatively, river inputs of mDOC are equal to 1.8 MtC y^{-1}, thereby representing less than 5% of the exported mDOC. mDOC contribution to

340 the total organic carbon flux is dominant. In the western basin, the total amounts of exported mPOC

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and mDOC below 100 m are respectively 7.0 MtC y^{-1} and 12.7 MtC y^{-1}, meaning that 64 % of this export is due to DOC. In the eastern basin, DOC is responsible for 90 % of the organic carbon export below 100 m, with an annual flux of 26.1 (against 2.6 for POC) MtC y^{-1}.

3.1.4 Spatial variability of export fluxes

345   Mean F_{OC} over the whole basin equals -22.8 gC m^{-2} y^{-1}, but a wide spatial variability can be observed in Fig. 5. Hence, the main regions of mOC export are the Liguro-Provencal sub-basin, the Alboran Sea, the southern continental slopes and the Adriatic Sea.

In the western basin, high positive values (i.e. upward) of F_{DOC} are simulated along the French and Spanish coasts, the entrance to the Sicilian Strait and north-eastern Excluding these areas, the

350   highest downward fluxes of DOC are calculated in the Provencal sub-basin (especially in the region of deep convection), the north of the Balearic Islands and along the Algerian slope, where downward F_{DOC} can be higher than 60 gC m^{-2} y^{-1}.

In the eastern basin, the complexity of topography and hydrodynamic regimes in the Aegean Sea may explain the high heterogeneity of the fluxes calculated in this region that are difficult to

355   interpret. Highest downward F_{DOC} values are located along the continental slope from the Libyan to the Turkish coasts and in the Adriatic Sea. Elsewhere (i.e. in the open sea), F_{DOC} distribution is more homogeneous, with a median of -17 gC m^{-2} y^{-1}.

A strong difference exists between the western and eastern basins regarding F_{POC} at 100 m. The mean value of downward F_{POC} throughout the western basin is -9.8 $gC.m^{-2}.y^{-1}$ against -2.4

360   gC m^{-2} y^{-1} in the eastern basin (Fig. 5 bottom).

In the western basin, F_{POC} is the highest in the Alboran Sea, particularly in the south east of the easily identifiable anticyclonic eddies. Following the pathway of the Atlantic waters, downward F_{POC} values decrease to reach absolute values lower than 5 gC m^{-2} y^{-1} in the Tyrrhenian Sea. In the Provencal basin high POC fluxes linked to the deep convection, with values ranging from -15

365   to -30 gC m^{-2} y^{-1} have been modeled. Throughout the eastern basin, F_{POC} is low except in the Adriatic Sea.

Finally, as suggested in Fig. 5, the spatial correlation between POC and DOC fluxes is weak almost everywhere. Regions of high POC or DOC export generally do not match. The only areas associated with both high POC and DOC exports are the Algerian coast, the Adriatic coast, the regions of deep

370   convection and a band east of the Balearic Islands.

3.1.5 Seasonal variability

The seasonal variability and the spatial distribution of F_{DOC} and F_{POC} differ significantly (Fig. 6 and 7). In winter (Fig. 6a), F_{DOC} values are high in almost all of the Mediterranean Basin except the Alboran Sea, with maximum values that can be observed in the Provencal sub-basin and along

375   the continental slopes, especially along the southern and eastern coasts of the eastern basin. F_{DOC}
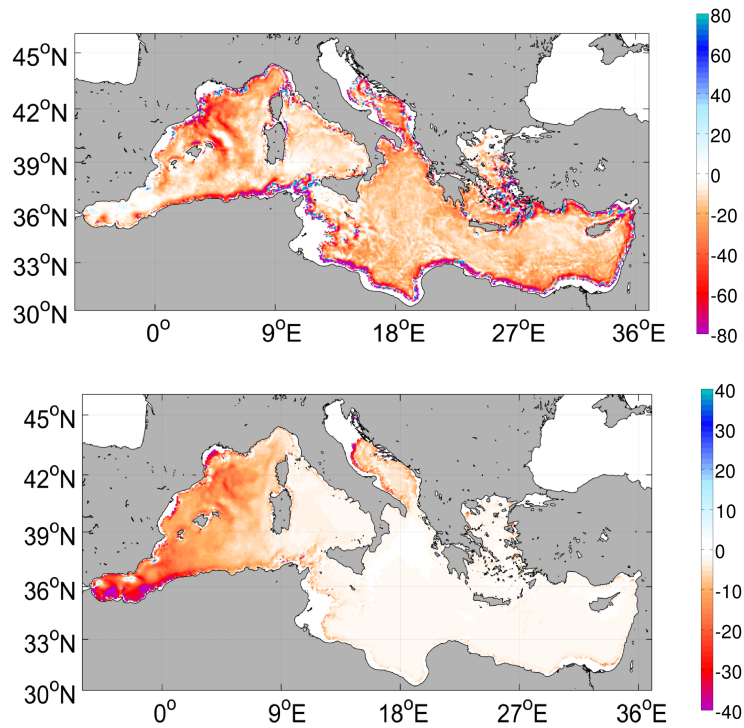
13

Figure 5. Maps of modeled annual DOC fluxes (top) and POC fluxes (bottom) below the 100 m layer in gC m^{-2} y^{-1}. Note the colorscale differences. Negative (red) means a downward flux.

distribution is quite similar in autumn, though with values that are significantly lower everywhere. During the rest of the year, F$_{DOC}$ values are very low in spring nearly everywhere, and almost null in summer. In several areas (Tyrrhenian and Adriatic Seas, Levantine and Ionian basins), high downward F$_{DOC}$ values are observed in winter while they are almost null during the rest of the year.

380   High downward POC fluxes at 100 m were calculated from winter to spring west of 7°E, namely in the Alboran Sea and the Provencal sub-basin (Fig. 7). In these regions, maximum values are reached in late winter (February-March) in the Alboran Sea, and in spring (March-April) in the Algerian Sea and the Provencal sub-basin. POC export in the eastern basin (excluding the Adriatic Sea) is very weak (even in the Rhodes Gyre) all year long. Maximum values can however be identified in

385   spring in the Tyrrhenian Sea, the Levantine basins (except for the Rhodes Gyre where the maximum is earlier in winter) and in the Adriatic Sea.
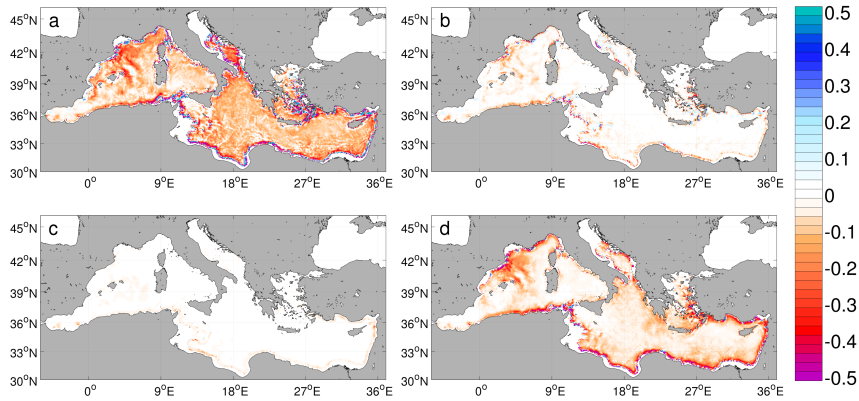
14

Figure 6. Maps of modeled DOC fluxes across the 100 m layer (F_{DOC}) in gC m^{-2} d^{-1} in (a) winter (Dec.-Feb.), (b) spring (Mar.-May), (c) summer (Jun.-Aug.), (d) autumn (Sept.-Nov.). Negative (red) means a downward flux.
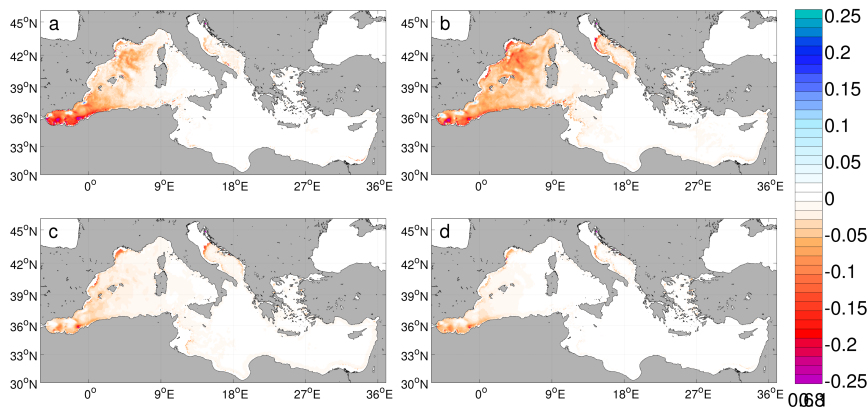

Figure 7. Maps of modeled POC fluxes across the 100 m layer F_{POC} in gC m^{-2} d^{-1} in (a) winter (Dec.-Feb.), (b) spring (Mar.-May), (c) summer (Jun.-Aug.), (d) autumn (Sept.-Nov.). Negative (red) means a downward flux.

3.1.6 Export below 200 m

Below 100 m, organic carbon is progressively consumed via the bacterial activity and respiration. At 200 m, the calculated mean export fluxes of total organic carbon are reduced by almost 87 % and 64 % compared to those at 100 m, respectively in the western and eastern basins. However, the ratio between export at these two depths is highly variable, depending on the region (see Fig. 8).

For POC (Fig. 8 a), if we consider first the regions where the annual F_{POC} values are significant, i.e. west of 7°E, (see Fig. 5 bottom), the 200 m to 100 m ratio is lower than 0.25 (i.e. only 25 % of
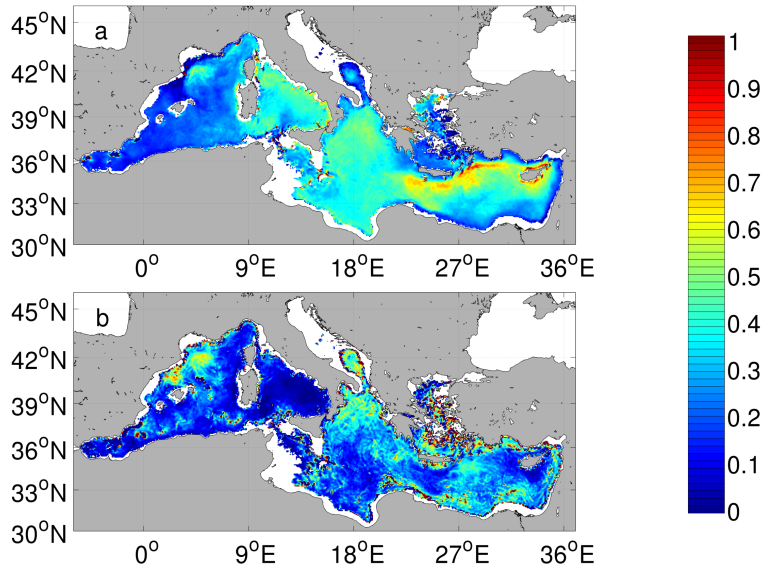
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Figure 8. Ratio between export fluxes at 200 m and at 100 m (a) for POC, (b) for DOC.

the carbon exported at 100 m goes below 200 m) in a region including the Alboran Sea, the western Algerian Sea and the Balearic Sea. This ratio is slighly higher but still below 0.3 for the central Algerian Sea and the Adriatic Sea. The Provencal sub-basin is the only region of high export below 200 m with a ratio about 0.4. In regions of low annual POC export (i.e. east of 7°E), the ratio ranges between 0.4 and 0.8 in the Tyrrhenian Sea, the Ionian and Levantine basins.

For DOC (Fig. 8 b), the ratio is more spatially variable, and in some regions the ratio is higher than 0.4, namely in the Provencal sub-basin, along the coasts of the Levantine basin, in the North Ionian basin, the Rhodes Gyre and the Adriatic Sea. Some patches of high ratios are also visible close to the Algerian Coast. Elsewhere the ratio ranges from almost zero (Tyrrhenian Sea, the Alboran Sea) to 0.2 in the eastern basin.

3.2 Intracellular quotas in bacteria and phytoplankton

Intracellular quotas in phytoplankton and bacteria are required for a further analysis of POC and DOC export fluxes and are presented in the following section. Carbon quota (Q_C) in small phytoplankton is maximum (> 0.7) in spring and summer in almost all of the Mediterranean Sea, though Q_C values are slightly lower in spring than in summer in the western basin, especially in the bloom region (Fig. 9). In autumn, though Q_C has decreased in nearly all of the Mediterranean Sea, Q_C values along the southern and eastern coasts of the eastern basin are significantly higher than in the rest of the open sea. In winter, Q_C values are even lower, with local maximum located in the Balearic Sea and in the south of the eastern basin.
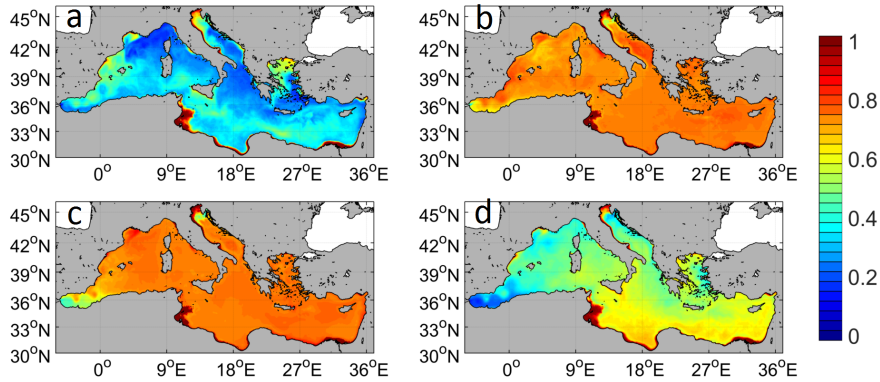
16

Figure 9. Seasonal variations of mean 0-50 m carbon relative quotas in small phytoplankton: (a) winter (Dec.-Feb.), (b) spring (Mar.-May), (c) summer (Jun.-Aug.), (d) autumn (Sept.-Nov.). Relative quotas are equal to 0 when the quota is mimimum (i.e. when $Q_C = Q_C^{min}$) and equal to 1 when the quota is maximum (i.e. when $Q_C = Q_C^{max}$)
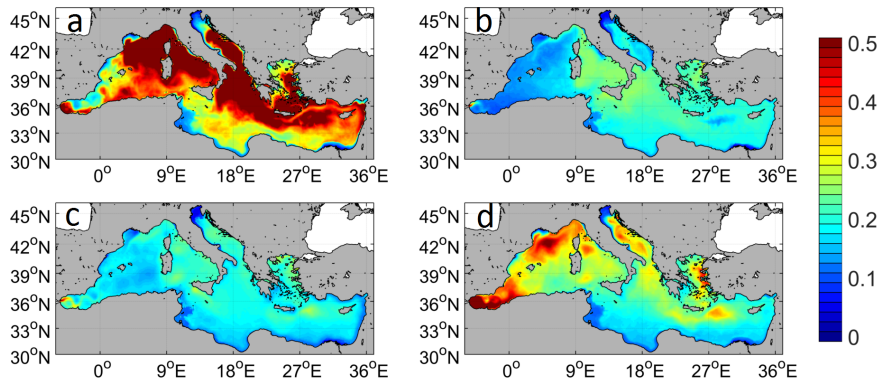

Figure 10. Seasonal variations of mean 0-50 m phosphorous relative quotas in small phytoplankton: (a) winter (Dec.-Feb.), (b) spring (Mar.-May), (c) summer (Jun.-Aug.), (d) autumn (Sept.-Nov.). Relative quotas are equal to 0 when the quota is mimimum (i.e. when $Q_P = Q_P^{min}$) and equal to 1 when the quota is maximum (i.e. when $Q_P = Q_P^{max}$)

The seasonal signal of the P quota (Q_P) in small phytoplankton is nearly the opposite of that of Q_C values in autumn and mostly in winter in nearly the whole of the Mediterranean Basin, and the lowest ones in spring and summer (Fig. 10). All year long, Q_P values are lower along the southern and eastern coasts than in the rest of the eastern basin.

Bacteria Q_C generally increases from winter to summer in most of the Mediterranean Basin (Fig. 11). In autumn, the decrease in Q_C is observed everywhere except throughout the same already identified region (namely along the southern and eastern coasts of the eastern basin). All year round,
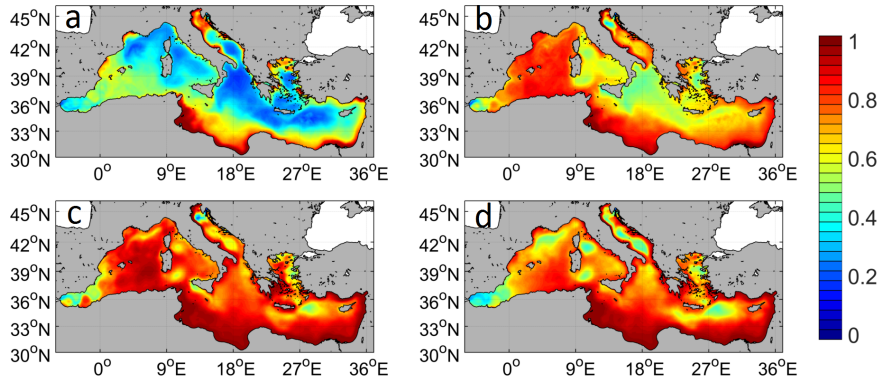
17

Figure 11. Seasonal variations of mean 0-50 m carbon relative quotas in bacteria: (a) winter (Dec.-Feb.), (b) spring (Mar.-May), (c) summer (Jun.-Aug.), (d) autumn (Sept.-Nov.). Relative quotas are equal to 0 when the quota is mimimum (i.e. when $Q_C = Q_C^{min}$) and equal to 1 when the quota is maximum (i.e. when $Q_C = Q_C^{max}$)

420   Q_C values are higher in this region than in the rest of the basin and even reach the Q_C^{max} value in summer and autumn thus indicating that carbon needs for bacteria growth are fully satisfied. In the deep convection regions (Liguro-Provencal sub-basin, Adriatic, Rhodes Gyre region), and in some eddies well identified in the Alboran and Tyrrhennian seas, the carbon quota is generally lower than in the surrounding waters, especially in autumn.

425   Bacteria Q_P values are very low everywhere in spring and summer except in the latter regions. The minimum Q_P values (i.e. the highest bacterial P-limitation) are observed in spring in the western basin, while they are reached in summer in the eastern basin. As for phytoplankton, Q_P values are lower all year round along the southern and eastern coasts than in the rest of the eastern basin.
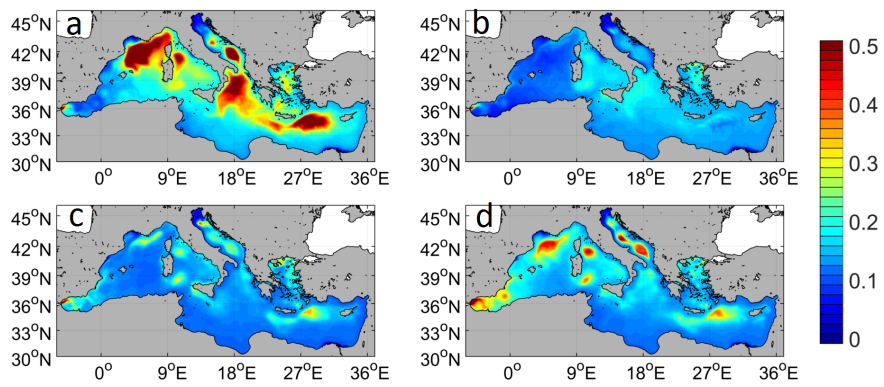
18

Figure 12. Seasonal variations of mean 0-50 m phosphorous relative quotas in bacteria: (a) winter (Dec.-Feb.), (b) spring (Mar.-May), (c) summer (Jun.-Aug.), (d) autumn (Sept.-Nov.). Relative quotas are equal to 0 when the quota is mimimum (i.e. when $Q_P = Q_P^{min}$) and equal to 1 when the quota is maximum (i.e. when $Q_P = Q_P^{max}$)
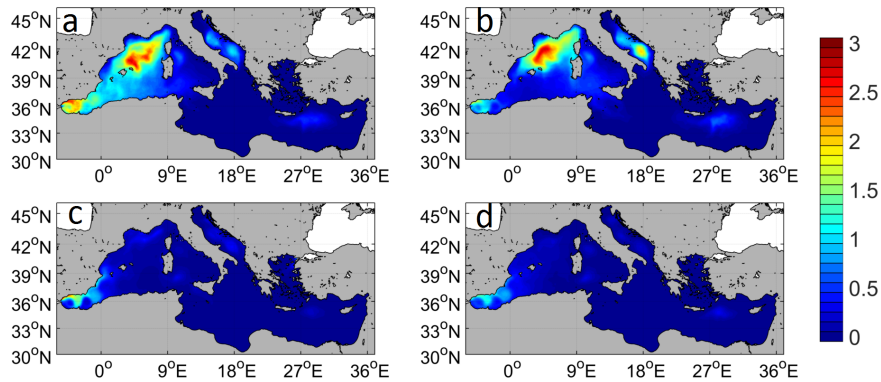
Figure 13. Seasonal variations of DOC mean 0-100 m exudation accumulated flux by large phytoplankton (in mol C.m^{-2}).

3.3 DOC exudation by phytoplankton

430 DOC exudation by large phytoplankton mainly occurs in the bloom region of the western basin (especially in the deep convection zone), in (late) winter and spring where accumulated fluxes are up to 2.8 mol C.m^{-2} (Fig. 13). Elsewhere, exudation fluxes are very low throughout the year, except in the Alboran Sea, two eddies of the Adriatic Sea and in the Rhodes gyre region.

The seasonality and the spatial patterns of DOC exudation flux by small phytoplankton are rather
435 different. The highest mDOC exudation fluxes are modeled in spring in the western basin, especially in the Gulf of Lions and the deep convection zone where accumulated fluxes up to 3 mol C.m^{-2} are calculated. In the eastern basin, the highest fluxes are observed in spring and summer. During these seasons, apart from the Adriatic Sea (especially in the north and along the eastern coast where accumulated fluxes also reach 3 mol C.m^{-2}) and some hot spots (Rhodes gyre, Nile plume),
440 mDOC exudation seems homogeneous though a north-south gradient is present. Hot spots of mDOC exudation are also present nearly all year long in the plumes of the main rivers.
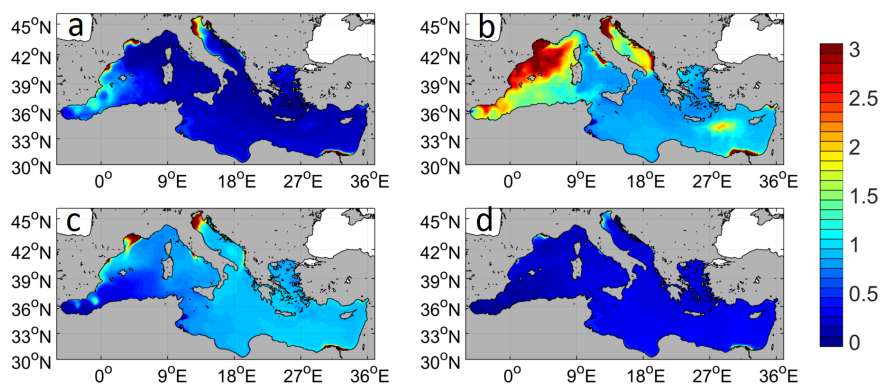
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Figure 14. Seasonal variations of mDOC mean 0-100 m exudation accumulated flux by small phytoplankton (in mol C.m^{-2}).

4 Discussion

4.1 The dissolved fraction in the organic carbon export is predominant at the scale of the Mediterranean Sea

445 One of the main results of this study is that mDOC export exceeds mPOC export in the whole of the Mediterranean Basin, with the exception of the Alboran Sea (west of 3°W). This is consistent with the comparisons between POC and DOC exports performed in the Tyrrhennian, North Ionian and Ligurian seas by Copin-Montégut and Avril (1993); Santinelli et al. (2013) or by Lefèvre et al. (1996) who estimated that DOC was the main source of remineralization processes in the aphotic

450 layer. In the western basin, the ratio of mDOC over mPOC export fluxes ranges between 2 and 5, and is approximately equal to 4 at the DyFaMed grid point. Observations at the DyFaMed station led to a oDOC export estimation of about 11.9 gC m^{-2}.y^{-1}, markedly higher than oPOC export estimations at 200 m (Avril, 2002, and references herein). Moreover, oPOC fluxes calculated by Miquel et al. (2011) during the 2001-2005 period ranged from 1.6 to 2.6 gC m^{-2}.y^{-1}. For comparison, mPOC

455 export flux was in the range [1.5;3.1] gC m^{-2}.y^{-1} during the same period. In the northwestern basin, the modeled ratio is about 2 at 100 m and 200 m, while in the same area a modeling study (Herrmann et al., 2014) led to a ratio at 200 m which ranged from 0.9 to 1.8, even though the corresponding export fluxes were higher than in the present study.

The ratio between modeled DOC and POC exports at 100 m ranges from 2 to 8 in the Adriatic
460 Sea. In the same region, a oDOC flux of 15.4 (against 23 for mDOC) gC m^{-2}.y^{-1} was estimated from observations by Santinelli et al. (2013). This is nearly 5 times higher than the measured oPOC export flux estimated by Boldrin et al. (2002) under the euphotic zone of 3.3 (against 4.5 for mPOC export at 100 m) gC m^{-2}.y^{-1}. These oDOC and oPOC fluxes were however estimated at different periods.

465 In the eastern basin, mDOC export is regularily more than 10 times that of mPOC, due to the very weak mPOC export and to the high mDOC export along the coasts and in the open sea. Few observations and estimations are available for this region. In the northern Ionian Sea, Boldrin et al. (2002) reported annual oPOC fluxes at 150 m of 2.4 gC m^{-2}.y^{-1}, which are in the same order of magnitude as the annual mPOC fluxes calculated in the same area but for a different period (1.2 gC
470 m^{-2}.y^{-1} and 0.6 gC m^{-2}.y^{-1} at 100 m and 200 m, respectively).

DOC predominance in the OC export flux is first due to the higher DOC gross production fluxes as compared to those of POC, and this still holds if the POC to DOC hydrolysis flux is ruled out (i.e. if the DOC inputs due to POC hydrolysis are not taken into account). At the scale of the Mediterranean Basin as a whole, mDOC and mPOC gross production fluxes are indeed respectively equal to 20
475 10^{12} and 2.7 10^{12} molC.y^{-1}. In the western basin, mDOC predominance in the export of OC still holds though to a lesser extent, with mDOC and mPOC gross production fluxes respectively equal

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to 8.7 10^{12} and 1.9 10^{12} molC.y^{-1}. In the following section, the reasons for these differences will be further analyzed in the light of the processes associated with DOC and POC production.

4.2   POC and DOC exports are characterized by different processes and timing

480   Strong disparities can be identified between the spatial patterns of the annual DOC and POC export fluxes (figure 5), with rather homogeneous DOC export fluxes across the Mediterranean Sea (though with well identified regions of maximum export that will be analyzed later), contrasting with the high east-west gradient in POC export. This is consistent with in situ measurements of daily POC export across the Mediterranean Sea at 200 m that showed much lower POC export in the eastern

485   basin than in the western basin (Moutin and Raimbault, 2002).

There are also considerable differences in the seasonality of DOC and POC export fluxes (Fig. 6 and 7). Over the whole of the Mediterranean Sea, 88 % of DOC export occurs between November and February, which is consistent with observations at the DyFaMed station where 90% of annual DOC export was linked to winter mixing (Avril, 2002). By contrast, POC export is more even

490   throughout the year, and during the same period only 23 % of POC is exported.

In the model, only the detrital compartment (POC) is allowed to sink. The sinking process is therefore the only source of explicit distinction between POC and DOC exports, but it is probably not sufficient to explain the strong aforementioned differences. The main source of difference lies in the biogeochemical processes that fuel or consume POC and DOC pools (see section 2.2). In the

495   model, POC is fueled by the natural mortality of the largest organisms (mesozooplankton, diatoms and ciliates) and by the egestion of fecal pellets and sloppy feeding by mesozooplankton. Thus, higher concentrations of large organisms in the western basin, primarily due to the spring bloom in the Liguro-Provencal sub-basin associated with high primary production rates is the main reason for the higher POC production and export in this basin. Hence, POC export is at a maximum in spring

500   (i.e. from March to May in figure 7) since it is the period including the maximum and the end of the bloom during which detrital concentrations of large organisms are highest. Moreover, according to the model, mortality is the main process that fuels the POC pool, far ahead of the egestion and sloppy feeding processes. More generally, a strong correlation between annual primary production and POC export has been evidenced at basin scale (Spearman's rank correlation coefficient is 0.84),

505   while this is not the case for DOC export (correlation below 0.01).

As shown in the Results section, the regions of high POC or DOC export are generally not the same, except for the regions characterized by high primary production rates during the spring bloom, namely the Alboran Sea, the bloom region in the NW Mediterranean Sea and the south of the Adriatic Sea (see also section 4.3). Apart from these regions, the annual DOC export at 100 m is relatively

510   high in almost all of the Mediterranean Basin, particularly in autumn and winter, and is the consequence of DOC accumulation in the 0-100 m layer during summer and autumn (Fig. 3). DOC export

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does indeed take place when DOC rich surface waters plunge or are mixed with poorer deeper waters.

This accumulation of DOC is primarily due to water stratification that results in nutrient depletion in the 0-100 m layer. As a result, the pool of DOC in phytoplankton is saturated with newly synthesized organic compounds since photosynthesis (i.e. carbon production), which is not controlled by P-availability, takes place more rapidly than is required to supply the needs of growth (cell division being limited by the intracellular quota of P). This results in high DOC exudation by phytoplankton, which is the main source of DOC in the model. The contribution of zooplankton excretion is at a maximum in spring in the bloom region of the NW Mediterranean, but remains always much lower than that of exudation (results not shown). Similarly, the annual contribution of POC hydrolysis to the DOC production flux is weak (around 10 %). Bacteria are the first consumers of DOC, and the second ingredient for DOC accumulation is therefore a strong nutrient limitation that will highly restrict the bacteria growth rate (see Eq. 1). In this situation, DOC availability may exceed bacteria needs and result in DOC accumulation when DOC production by phytoplankton exceeds DOC uptake by bacteria. This process is enhanced in hydrodynamic situations where the surface layers are isolated from the deep waters (i.e. stratification period). Such a mechanism of DOC accumulation due to a malfunctioning microbial loop has already been described in Thingstad et al. (1997) and is also the main driver of DOC accumulation in the model. Destratification in autumn leads to a net export as well as an increase of DOC consumption through bacterial activity, driven by nutrient supply from deep water.

4.3 DOC accumulation in the light of intracellular quotas

The regions of highest DOC export fluxes correspond to the regions where the highest DOC accumulation occurs. It is therefore informative to analyze the occurrence of DOC accumulation in the light of intracellular quotas. Geographical and hydrological considerations are indeed not sufficient for a full understanding of the DOC accumulation pattern at the scale of the Mediterranean Sea.

It has already been said that, according to the model, phytoplankton exudation is the primary source of DOC. High DOC exudation by phytoplankton occurs in nutrient-depleted waters. In such a situation N and/or P phytoplankton nutrient quotas are low and limit growth rate (i.e. the cell division rate). In the model, phytoplankton (and bacteria) cell division rate is indeed controlled by the most strongly limiting element among C, N and P (see Eq. 1). In other words, the intracellular quota which is the closest to its minimum value controls the division rate. When P (and/or N) are the most strongly limiting, growth will proceed at low rate and the carbon input due to photosynthesis will rapidly meet phytoplankton needs, thus resulting in an increase in the carbon quota Q_C. Since DOC exudation flux per cell increases with Q_C through a Geider et al. (1998) non-linear quota function, DOC exudation flux will highly increase as the quota approaches its maximum value $Q_C^{\max}$. Phytoplankton carbon quota is therefore a good indicator of DOC exudation.

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In the oligotrophic Mediterranean Sea, nutrient (and mostly P in the model) depletion is at a maximum at the end or just after the spring bloom, or under well established conditions of water stratification, thus leading to maximum exudation fluxes (see Fig. 13 and 14). In the rest of the Mediterranean, DOC exudation is at a maximum in (late) spring and summer, and mainly due to small phytoplankton. The latter is indeed characterized by low phosphorous quotas (see Fig. 10) and high carbon quotas (see Fig. 10).

The driving processes of DOC accumulation are not the same in the western and the eastern Mediterranean. In the western Mediterranean, and especially in the enlarged bloom region, large phytoplankton blooms first and is rapidly P-limited (as early as February) and the same occurs for small phytoplankton though later (i.e. only in spring, see Fig. 10). This is consistent with observations performed in the NW Mediterranean Sea (Gulf of Lions) (Diaz et al., 2001). In this situation, the high phytoplankton exudation fluxes are not only due to phytoplankton carbon quotas that are relatively high (around 50-60%, see the small phytoplankton carbon quota in Fig. 9), resulting in relatively high exudation flux per cell, but to the high phytoplankton abundance. Though exudation fluxes are high in (late) winter due to large phytoplankton (Fig. 13a), the high bacteria P-quotas (Fig. 12a) combined with winter mixing prevents DOC accumulation (Fig. 3a). In spring, and mostly in late spring, bacteria are strongly P-limited (Fig. 12b) since the bloom has rapidly consumed the available nutrients and vertical mixing has stopped. As a result, DOC accumulation starts in this region (Fig. 3b) and reaches its maximum in summer (Fig. 3c) during the stratification period since DOC exudation by phytoplankton still proceeds (though at a lower rate) and bacteria are still strongly P-limited (Fig. 12c). Finally, the end of the stratification in autumn will not only dilute the DOC-rich surface concentrations with DOC-poor deep waters, but allow the P-enrichment of surface waters (see the increase in bacteria Q_P in Fig. 12d).

In the eastern Mediterranean, DOC accumulation is mainly visible along the southern and eastern coasts. Moreover, it starts later than in the western Mediterranean (i.e. in summer against spring for the west), and is at a maximum in autumn. In the model, the Atlantic waters that flow along the coast are less dense (with densities slightly underestimated as compared to in situ measurements (Beuvier, 2011)) and therefore strongly isolated from the rest of the water column. As a result, their nutrient content will be progressively consumed and these waters become more and more oligotrophic as they flow along the southern coast of the basin, and always remain more oligotrophic than the rest of the eastern basin. In summer and autumn, they can even be considered as ultra-oligotrophic (see the phytoplankton Q_P in Fig. 10c and d). Moreover, they extend over a layer of around 100 m in thickness in which concentrations are roughly homogeneous. During summer and autumn, bacteria are also strongly P-limited but more and more carbon-rich (see Fig. 11) since phytoplankton exudation still proceeds (though at extremely low rates in autumn). Moreover, the vertical mixing that starts in autumn is not sufficiently deep to reach the nutrient-rich waters since the MLD is shallower than the bottom of these Atlantic waters. In addition, since DOC concentration is high over the

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585 whole layer, DOC surface concentrations are not diluted by the mixing. As a result, accumulation still proceeds until winter during which higher MLD will allow the P-enrichment in surface waters and dilute surface DOC concentrations as well.

Furthermore, DOC concentrations (as well as DOC annual export flux though this is more difficult to see in Fig.5) are negligeable throughout the year in some well-identified regions, namely the two
590 cyclonic structures in the Tyrrhenian Sea, the south of the Adriatic Sea (excluding the coastal zones), and the region of the Rhodes Gyre in the Levantine basin. All these structures are characterized by regular input of nutrients from deep waters, resulting in an absence of strong P-limitation in bacteria. Under such conditions, the bacteria carbon quota is rather low and DOC accumulation and export cannot occur.

595 Finally, the strong link between low phosphate availability in the upper surface water of the Mediterranean Sea and DOC accumulation due to nutrient limitation of bacterial production that is evidenced in this modeling study is consistent with previous in situ (Moutin et al., 2002; Van Wambeke et al., 2002) and modeling (Thingstad et al., 1997) studies and is shown to apply at the scale of the whole of the Mediterranean Sea, with the exception of the aforementioned specific regions.

600   **4.4 Robustness of results**

Though difficult to achieve in a rigorous way, the robustness of our main results will be discussed in the following section. As shown in section (2.2), the model includes many DOC and POC production and consumption processes. A sensitivity study on all the parameters they involve is obviously impossible to achieve, though some steps towards this goal have already been made in Baklouti
605 et al. (2006b). Moreover, accounting for the fact that most of the parameters used have a physiological significance (including cell size considerations), and constitute a coherent set that remains unchanged for the different studies undertaken with Eco3M-MED (even outside the Mediterranean), we consider that their values are reasonably reliable. However, the POC to DOC degradation (i.e. hydrolysis) rate and the sinking velocity are not physiological parameters and their impact on the
610 results will be discussed later.

The comparison of DOC stocks with the few available results (see ~~Appendix~~ the Supplementary Material) showed that, though the shape of the modeled DOC vertical profiles were quite different (but the values were in the same order of magnitude) from those measured, modeled and measured integrated DOC stocks over the 0-100 m layer showed much better agreement. Furthermore, when
615 compared to in situ estimations of DOC export from the DyFaMed station (Avril, 2002) and the Adriatic and Tyrrhenian seas (Santinelli et al., 2013), the model always provides higher DOC export values. These differences in DOC export may be partly attributable to the model failures discussed ~~in section (??)~~ the Supplementary Material but, as already mentioned, in situ estimations also involve considerable uncertainties. Hence, according to Santinelli et al. (2013), DOC export computations
620 from stock differences below the euphotic layer probably underestimate the real flux. This is also

the conclusion we came to by using model outputs to compute export fluxes with our method and with the in situ method. If we assume, however, that the different in situ estimations are consistent with each other, it appears that the highest DOC export occurs in the Adriatic Sea, followed by the DyFaMed station (Ligurian Sea) and then by the Tyrrhenian Sea, and the same order can be inferred from the model outputs.

Two parameters are essential in POC export, namely POC to DOC degradation rate and the sinking velocity.

Since our model includes a single detrital compartment, an intermediate value of 2 m d^{-1} has been used for the sinking velocity. This value is intended to be representative of the high sinking rates (> 100 m/day) of very large particles as well as the very low sinking rates of small particles. It may however reflect an underestimation of the actual mean value though this is difficult to verify. In several other models (e.g. Lévy et al., 1998; Lacroix and Gregoire, 2002; Herrmann and Somot, 2008), two detrital compartments are used, thus making it possible to differentiate between low and high sinking rates of detrital particules. However, in these models, the large detrital compartment (to which high sinking rates are affected) is only fueled by zooplankton fecal pellets (Lévy et al., 1998; Herrmann and Somot, 2008) and by mesozooplankton mortality in Lacroix and Gregoire (2002). These fluxes, except the latter, are probably weak compared to the other POC sources in our model (which is dominated by the mortality of the largest organisms). Finally, in these models, the remaining sources of POC fuel the small detrital compartment for which the sinking velocities are lower than that used in our model. To conclude on this point, adding complexity in a given model generally leads to the multiplication of the number of state variables and parameters. When these parameters and/or these new processes are not well known, this also adds uncertainty, and in this case complexity does not necessarily lead to a better agreement between model outputs and observations as this is suggested by several studies (e.g. Muller et al., 2009; Kriest et al., 2010; Paudel and Jawitz, 2012). With the addition of a class of large detrital particles, the sinking velocity associated with this detrital compartment as well as the definition of the processes that would have fueled this additional compartment (the latter are not the same in the different aforementioned models that use two detrital compartments) would have added a source of uncertainty in the model. As a consequence, the particulate carbon fluxes wouldn't necessarily have been more realistic than the ones provided by this study. The same conclusion may apply as regards the aggregation models which formulations are generally empirical and associated with parameters that are hardly measurable.

More importantly, it can be considered that the likely underestimated sinking velocity used in the present model is compensated by the very low POC degradation rate. In our model, its maximum value is set at 0.03 d^{-1} but it is modulated by the bacteria carbon quota. In substance, the higher the carbon quota, the more the degradation rate decreases and eventually becomes 0 when the bacteria carbon quota is maximum. As a result, the effective POC degradation rate is always less than 0.03 d^{-1} in the model, and it is lower in the surface layers since bacteria are more rich in carbon than

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in deep waters. It is also lower than all the values used in the aforementioned models. Concerning in situ data for the degradation rate, Sempéré et al. (2000) have determined values at 50 and 200 m for labile and less labile POC in three regions of the Mediterranean Sea, showing that, for the labile POC (which represent a significant part in the latter study), the degradation rate can be up to 100 times higher than that used in the present study.

Apart from these two parameters, it has been seen that the model underestimates Chl concentrations at the DCM (mainly due to a lack of large phytoplankton) and this may also lead to an underestimation of POC export. However, the 0-100 m mIPP values are consistent with oIPP thereby suggesting that this DCM underestimation has only a limited impact on carbon production. Overall, the annual POC export flux at 100 m provided by the model is around 8% of the annual primary production, a value that is consistent with in situ estimations (Miquel et al., 1994).

Between 100 m and 200 m, however, the mean bacteria carbon quota is lower since POC hydrolysis and bacteria and heterotrophic nanoflagellate mortalities are the only sources of DOC, resulting in higher hydrolysis rates and in lower POC export at 200 m. Looking at the vertical attenuation of POC fluxes, it is common to use a power law expressed as $F(z) = F(z = z_0) * (\frac{z}{z_0})^{-b}$, where F(z) is the depth-dependent POC flux and b a positive coefficient whose values may vary according to the location or the period. In regions of significant export, b values inferred from the model outputs fluctuate between 0.9 in the Provencal sub-basin and 2.3 for the Algerian basin. Values of b derived from observations tend to be lower, i.e. respectively equal to 0.92 and 1.0 for the western and eastern moorings (Gogou et al., 2014), or 0.75 in the Alboran Sea (Zúñiga et al., 2007). This again suggests that the attenuation of POC export flux between 100 m and 200 m is too great in the model. Furthermore, when compared to the few available data for POC export fluxes, the model always underestimates the export flux in the eastern basin. However, all the in situ estimations we could find in the literature were done at 150 m or 200 m depth, which means in the 100-200 m layer where the modeled POC export is more likely to be underestimated. In summary, all this suggests that the underestimation of POC export fluxes is more to be the case at 200 m than at 100 m depth though the comparison at the DyFaMed station shows that the mean mPOC export rate (5.6 gC.m^{-2}.y^{-1} and 2.2 gC.m^{-2}.y^{-1} at 100 m and 200 m respectively) is within the range of the measured rate at 200 m (i.e. [1.6;2.6] gC.m^{-2}.y^{-1} (Copin-Montégut and Avril, 1993; Miquel et al., 2011)). Finally, it is very unlikely that these uncertainties could shed doubt on the predominance of DOC in the OC export in the eastern basin. This conclusion also applies in the western basin (though with less certainty), all the more so in that in situ measurements allow the same conclusion to be drawn in the sampled stations of the NW Mediterranean (Copin-Montégut and Avril, 1993; Avril, 2002; Miquel et al., 2011).

5   Conclusions

A 14-year simulation combining a high resolution physical model (NEMO-MED12) and a mechanistic biogeochemical model (Eco3M-MED) has been developed to study carbon organic production
695   and fate at the scale of the Mediterranean Sea.

A preliminary work presented in the ~~Appendix~~ Supplementary Material focused on the Model Skill Assessment through an extensive comparison of different model outputs (i.e. chlorophyll, nutrients, primary production and DOC profiles) with available data at various time and space scales. This work allowed to verify the model's ability to represent the main features of the biogeochemical
700   functioning of the Mediterranean Sea. In the Results section, carbon export fluxes are investigated. Previous estimations of DOC export in the Mediterranean Sea were restricted to specific regions of the Mediterranean (e.g. the Ligurian, Adriatic, Tyrrhenian Seas). We here propose the first Mediterranean-scale view of annual DOC and POC export fluxes, as well as an analysis of their spatial and seasonal variations in the light of plankton intracellular quotas.

705   The two major results of this modeling study lie in (i) the predominance of the eastern basin in OC export (with nearly 60 % of the OC export occuring in the eastern basin), and (ii) in the crucial role of the dissolved fraction in the total organic carbon export. At Mediterranean scale, DOC export represents about four fifths of total organic carbon fluxes, thereby attesting to its major role in the carbon cycle and the biological pump in the Mediterranean Sea. The concept of a malfunctioning
710   microbial loop (Thingstad et al., 1997), due to high P-limitation of both phytoplankton and bacteria leading to high DOC exudation fluxes beyond bacterial needs, also applies in the present study though it is generalized to the whole of the Mediterranean Basin, except for some specific P-rich regions (see Results and Discussion). Export in the eastern basin is markedly high despite its lower productivity compared to the western basin. By contrast, POC export is closely associated with
715   regions characterized by high productivity. As a consequence, total carbon export in the eastern basin is considerably higher than expected as regards its low primary productivity. Results also show high spatial variability in organic carbon fluxes and a temporal uncoupling between POC and DOC exports. This is attributable to the differences in the processes involved in the production and export of POC and DOC.

720   Further comparisons with observations are clearly necessary to confirm these results, which emphasizes the need for in situ temporal monitoring to properly quantify organic carbon export. This study also highlights the need to examine the microbial food web in detail in order to further investigate the carbon cycle in the Mediterranean Sea. Furthermore, the implementation of an explicit inorganic carbon compartment in the biogeochemical model would close the carbon budget and help
725   in the full characterization of the biological pump.

In conclusion, the strong link between low phosphate availability in the upper surface water of the Mediterranean Sea and DOC accumulation due to nutrient limitation of bacterial production already identified by previous modeling (Thingstad et al., 1997) and in situ (Moutin et al., 2002;

Van Wambeke et al., 2002) studies, is confirmed by this modeling study, which may therefore be
730   of interest for other oceanic regions. The low phosphate availability of the upper waters has been
identified in other oceanic regions such as the Sargasso Sea (Wu et al., 2000), the North Pacific and
the South West Pacific (Van Den Broeck et al., 2004), and high DOC accumulation has also been
reported in some of these areas (Carlson et al., 1994). This work may therefore be of interest for
these oceanic regions. Finally, in the context of climate change, the enhanced stratification and the
735   probable geographical extension of low phosphate availability in upper waters (Karl et al., 1997;
Moutin et al., 2008) is expected to result in an increase in DOC production (Santinelli et al., 2013;
Lazzari et al., 2013), and thereby further increase the importance of DOC in the biological carbon
pump.

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