Impact of dust addition on the metabolism of Mediterranean plankton communities and carbon export under present and future conditions of pH and temperature

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22

23 Abstract

24 Although atmospheric dust fluxes from arid as well as human-impacted areas represent a 25 significant source of nutrients to surface waters of the Mediterranean Sea, studies focusing on the 26 evolution of the metabolic balance of the plankton community following a dust deposition event 27 are scarce and none were conducted in the context of projected future levels of temperature and 28 pH. Moreover, most of the experiments took place in coastal areas. In the framework of the 29 PEACETIME project, three dust-addition perturbation experiments were conducted in 300-L 30 tanks filled with surface seawater collected in the Tyrrhenian Sea (TYR), Ionian Sea (ION) and 31 in the Algerian basin (FAST) onboard the R/V "Pourquoi Pas?" in late spring 2017. For each 32 experiment, six tanks were used to follow the evolution of chemical and biological stocks, 33 biological activity and particle export. The impacts of a dust deposition event simulated at their 34 surface were followed under present environmental conditions and under a realistic climate 35 change scenario for 2100 (ca. + 3 °C and -0.3 pH units). The tested waters were all typical of 36 stratified oligotrophic conditions encountered in the open Mediterranean Sea at this period of the 37 year, with low rates of primary production and a metabolic balance towards net heterotrophy. 38 The release of nutrients after dust seeding had very contrasting impacts on the metabolism of the 39 communities, depending on the station investigated. At TYR, the release of new nutrients was 40 followed by a negative impact on both particulate and dissolved ¹⁴C-based production rates, 41 while heterotrophic bacterial production strongly increased, driving the community to an even 42 more heterotrophic state. At ION and FAST, the efficiency of organic matter export due to 43 mineral/organic aggregation processes was lower than at TYR and likely related to a lower 44 quantity/age of dissolved organic matter present at the time of the seeding and a smaller

45 production of DOM following dust addition. This was also reflected by lower initial 46 concentrations in transparent exopolymer particles (TEP) and a lower increase in TEP 47 concentrations following the dust addition, as compared to TYR. At ION and FAST, both the 48 autotrophic and heterotrophic community benefited from dust addition, with a stronger relative 49 increase in autotrophic processes observed at FAST. Our study showed that the potential positive 50 impact of dust deposition on primary production depends on the initial composition and 51 metabolic state of the investigated community. This impact is constrained by the quantity of 52 nutrients added in order to sustain both the fast response of heterotrophic prokaryotes and the 53 delayed one of primary producers. Finally, under future environmental conditions, heterotrophic 54 metabolism was overall more impacted than primary production, with the consequence that all 55 integrated net community production rates decreased with no detectable impact on carbon 56 export, therefore reducing the capacity of surface waters to sequester anthropogenic CO₂.

57

58 1. Introduction

59 Low Nutrient Low Chlorophyll (LNLC) areas represent 60% of the global ocean surface 60 area (Longhurst et al., 1995; McClain et al., 2004). Although phytoplankton production in these 61 areas is limited by the availability of nitrogen, phosphorus and iron, it accounts for 50% of global 62 carbon export (Emerson et al., 1997; Roshan and DeVries, 2017). Atmospheric dust fluxes 63 represent a significant source of these nutrients to surface waters in LNLC regions and as such 64 could play a significant role in stimulating primary production (e.g. Bishop et al., 2002; Guieu et 65 al., 2014b; Jickells and Moore, 2015), potentially increasing the efficiency of the biological pump in the sequestration of atmospheric CO₂. However, as heterotrophic prokaryotes have been 66 67 shown to outcompete phytoplankton during nutrient addition experiments (e.g. Guieu et al., 68 2014a; Mills et al., 2008; Thingstad et al., 2005), dust deposition could induce even stronger 69 enhancements of heterotrophic bacterial production and/or respiration rates thereby reducing net 70 atmospheric CO₂ drawdown and the potential for carbon export outside the euphotic zone (Guieu 71 et al., 2014b). Indeed, several experiments conducted in the Atlantic Ocean and in the 72 Mediterranean Sea have shown a fast and dominant effect of dust additions on heterotrophic 73 bacterioplankton metabolism (Herut et al., 2005, 2016; Lekunberri et al., 2010; Marañón et al., 74 2010; Pulido-Villena et al., 2008, 2014). However, to the best of our knowledge, no study 75 focused on the evolution of the metabolic balance of the plankton community after such a dust 76 event in the open sea. The metabolic balance (or net community production, NCP) is defined as 77 the difference between gross primary production (GPP) of autotrophic organisms and community

respiration (CR) of both autotrophic and heterotrophic organisms, revealing the capacity of
 surface waters to absorb atmospheric CO₂.

80 The Mediterranean Sea is a perfect example of LNLC regions and receives anthropogenic 81 aerosols originating from industrial and domestic activities from all around the basin and other 82 parts of Europe and pulses of natural inputs from the Sahara (Desboeufs, 2022). These 83 atmospheric depositions, mostly in the form of pulsed inputs (Loÿe-Pilot and Martin, 1996), 84 provide new nutrients (Guieu et al., 2010; Kouvarakis et al., 2001; Markaki et al., 2003; Ridame 85 and Guieu, 2002) to the surface waters with fluxes that are of the same order of magnitude as 86 riverine inputs (Powley et al., 2017). These significant nutrient enrichments likely support 87 primary production especially during the stratification period (Bonnet et al., 2005; Ridame and 88 Guieu, 2002). However, no clear correlation between dust and ocean color have been evidenced 89 from long series of satellite observations (Guieu and Ridame, 2020). This raises the question on 90 which compartment (autotrophic or heterotrophic) benefits the most from these transient relieves 91 in nutrient (N, P) limitation.

92 In response to ocean warming and increased stratification, LNLC areas are expected to 93 expand in the future (Irwin and Oliver, 2009; Polovina et al., 2008) due to lower nutrient supply 94 from sub-surface waters (Behrenfeld et al., 2006). Furthermore, dust deposition could increase in 95 the future due to desertification (Moulin and Chiapello, 2006), although so far the trend for 96 deposition remains uncertain because the drying of the Mediterranean basin might also induce 97 less wet deposition over the basin (Laurent et al., 2021). Nevertheless, whether the fluxes 98 increase or not in the coming decades and centuries, new nutrients from atmospheric sources will 99 play an important role in a surface mixed layer even more stratified and isolated from the deeper 100 nutrient-rich layer. The question remains on how plankton metabolism and carbon export would

101respond in a warmer and more acidified ocean. Indeed, with an average annual anthropogenic102 CO_2 uptake, during the period 2010 to 2019, of 2.5 ± 0.6 GtC (~22.9% of anthropogenic103emissions; Friedlingstein et al., 2020), the oceans substantially contribute towards slowing down104the increase in atmospheric CO_2 concentrations, and therefore towards limiting terrestrial and105ocean warming. However, this massive CO_2 input induces global changes in seawater chemistry106referred to as "ocean acidification" because increased CO_2 concentration lowers seawater pH107(i.e. increases its acidity).

108 Although the response of plankton metabolism to ocean warming has been shown to be 109 highly dependent on resource availability (Lewandowska et al., 2014), both for heterotrophic 110 bacteria (Lopez-Urrutia and Moran, 2007) and phytoplankton (Marañón et al., 2018), it has been 111 suggested that ocean warming will substantially weaken the ocean biological CO₂ sink in the 112 future as a consequence of stronger increase in remineralization than in photosynthesis 113 processes, following the metabolic theory of ecology (MTE; Brown et al., 2004; Gillooly et al., 114 2001). Ocean acidification alone has been shown to exert no or very limited influence on 115 plankton metabolism in the Mediterranean Sea (Maugendre et al., 2017a; Mercado et al., 2014). 116 To the best of our knowledge, only Maugendre et al. (2015) studied the combined impact of 117 ocean warming and acidification on plankton metabolism in the Mediterranean Sea. They found 118 a very limited impact of ocean acidification on the plankton community and a positive impact of 119 warming on small phytoplankton species (e.g. Cyanobacteria) with a potential decrease of the 120 export and energy transfer to higher trophic levels. Their study was conducted under nutrient 121 depleted conditions (Maugendre et al., 2017b). Hence, there is still a need to assess the combined 122 impact of warming and acidification on the metabolic balance of plankton communities in this 123 region, following a transient relief in nutrient availability.

124 So far there has been no attempt to evaluate the evolution of plankton metabolism and carbon 125 export following atmospheric deposition in the context of future levels of temperature and pH. 126 Such experiments were conducted in the frame of the PEACETIME project (ProcEss studies at the 127 Air-sEa Interface after dust deposition in the MEditerranean sea; http://peacetime-project.org/) 128 during the cruise on board the R/V "Pourquoi Pas?" in May/June 2017 (Guieu et al., 2020a, b). 129 The project aimed at extensively studying and parameterizing the chain of processes occurring in 130 the Mediterranean Sea after atmospheric deposition, especially of Saharan dust, and to put them in 131 perspective of on-going environmental changes. During this cruise, three perturbation experiments 132 were conducted in 300-L tanks filled with surface seawater collected in the Tyrrhenian Sea (TYR), 133 Ionian Sea (ION) and in the Algerian basin (FAST; Fig. 1). Six tanks were used to follow the 134 evolution of chemical and biological stocks, biological activity and export, following a wet dust 135 deposition event simulated at their surface, both under present environmental conditions and 136 following a realistic climate change scenario for 2100 (ca. + 3 °C and -0.3 pH units; IPCC, 2013). 137 A companion paper presents the general setup of the experiments and the impacts of dust under 138 present and future environmental conditions on nutrients and biological stocks (Gazeau et al., 139 2021). In this paper, we show that the effects of dust deposition on biological stocks were highly 140 different between the three investigated stations and could not be attributed to differences in their 141 degree of oligotrophy but rather to the initial metabolic state of the community. We further 142 demonstrated that ocean acidification and warming did not drastically modify the composition of 143 the autotrophic assemblage with all groups positively impacted by warming and acidification. 144 Here, we focus on the impacts of dust seeding on plankton metabolism (e.g. primary production, 145 heterotrophic prokaryote production) and carbon export.

147 2. Material and Methods

148 **2.1. General set-up**

149 The general set-up of the experiments is fully detailed in Gazeau et al. (2021). Briefly, 150 three experiments were performed at the long duration stations TYR, ION and FAST during the 151 Peacetime cruise onboard R/V "Le Pourquoi Pas?" (Fig. 1). During these experiments (3 to 4 days each), seawater was incubated in 300-L tanks (Fig. S1) installed in a temperature-controlled 152 153 container, in which the irradiance spectrum and intensity can be finely controlled and in which 154 future ocean acidification and warming conditions can be fully reproduced. The tanks were made 155 of high-density polyethylene (HDPE) and were trace-metal free in order to avoid contaminations, 156 with a height of 1.09 m, a diameter of 0.68 m, a surface area of 0.36 m² and a volume of 0.28 m³. 157 The conical base of the tanks was equipped with a sediment trap that was left open during the 158 duration of the experiments and removed at the end. The experimental protocol comprised two 159 unmodified control tanks (C1 and C2), two tanks enriched with Saharan dust (D1 and D2) and 160 two tanks enriched with Saharan dust and maintained simultaneously under warmer (+ 3 °C) and 161 acidified (-0.3 pH unit) conditions (G1 and G2). At the three stations, tanks were always filled at 162 the end of the day before the start of the experiments: TYR (17/05/2017), ION (25/05/2017) and 163 FAST (02/06/2017). The tanks were filled by means of a large peristaltic pump (Verder© VF40 164 with EPDM hose, flow of 1200 L h⁻¹) collecting seawater below the base of the boat (depth of ~ 165 5 m), used to supply continuously surface seawater to a series of instruments during the entire 166 campaign. While filling the tanks, seawater was sampled for the measurements of selected 167 parameters (sampling time = t-12h). After filling the tanks, seawater was slowly warmed

168	overnight using 500 W heaters, controlled by temperature-regulation units (COREMA©), in G1
169	and G2 to reach an offset of $+$ 3 °C. ¹³ C-bicarbonate was added to all tanks at 4:00 am (all times
170	in local time) and G1 and G2 were acidified by addition of CO ₂ -saturated filtered (0.2 $\mu m)$
171	seawater (~1.5 L in 300 L; collected when filling the tanks at each station) at 4:30 am to reach a
172	pH offset of -0.3. Sampling for many parameters took place prior to dust seeding (sampling time
173	= t0). Dust seeding was performed between 7:00 and 9:00 in tanks D1, D2, G1 and G2. The same
174	dust analog was used and the same dust flux was simulated as for the DUNE 2009 experiments
175	described in Desboeufs et al. (2014). Briefly, the fine fraction (< 20 μ m) of Saharan soils
176	collected in southern Tunisia, which is a major source of dust deposition over the northwestern
177	Mediterranean basin, was used in the seeding experiments. The particle size distribution showed
178	that 99% of particles had a size smaller than 0.1 μ m, and that particles were mostly made of
179	quartz (40%), calcite (30%) and clay (25%; Desboeufs et al., 2014). This collected dust
180	underwent an artificial chemical aging process by addition of nitric and sulfuric acid (HNO ₃ and
181	H ₂ SO ₄ , respectively) to mimic cloud processes during atmospheric transport of aerosol with
182	anthropogenic acid gases (Guieu et al., 2010, and references therein). To mimic a wet flux event
183	of 10 g m ⁻² , 3.6 g of this analog dust were quickly diluted into 2 L of ultrahigh-purity water
184	(UHP water; 18.2 M Ω cm ⁻¹ resistivity), and sprayed at the surface of the tanks using an all-
185	plastic garden sprayer (duration = 30 min). The intensity of this simulated wet deposition event
186	(i.e. 10 g m ⁻²) represents a high but realistic scenario, as several studies reported even higher
187	short wet deposition events in this area of the Mediterranean Sea (Bonnet and Guieu, 2006;
188	Loÿe-Pilot and Martin, 1996; Ternon et al., 2010).

Depending on the considered parameter or process, seawater sampling was conducted 1 h
(t1h), 6 h (t6h), 12 h (t12h), 24 h (t24h), 48 h (t48h) and 72 h (t72h) after dust additions in all

three experiments with an additional sample after 96 h (t96h) at FAST). Acid-washed silicone tubes were used for transferring the water collected from the tanks to the different vials or containers.

194 **2.2. Stocks**

195 **2.2.1. Dissolved and particulate organic carbon**

196 The concentration of dissolved organic carbon (DOC) was determined from duplicate 10 197 mL GF/F (pre-combusted, Whatman[©]) filtered subsamples that were transferred to pre-198 combusted glass ampoules, acidified with H_3PO_4 (final pH = 2) and sealed. The sealed glass 199 ampoules were stored in the dark at room temperature until analysis at the Laboratoire 200 d'Océanographie Microbienne (LOMIC). DOC measurements were performed on a Shimadzu© 201 TOC-V-CSH (Benner and Strom, 1993). Prior to injection, DOC samples were sparged with 202 CO_2 -free air for 6 min to remove inorganic carbon. Sample (100 μ L) were injected in triplicate 203 and the analytical precision was 2%. Standards were prepared with acetanilid. 204 Seawater samples for measurements of particulate organic carbon concentrations (POC; 2 205 L) were taken at t-12h, t0, t12h, t24h, t48h and t72h (or t96h for station FAST), filtered on pre-206 combusted GF/F membranes, dried at 60 °C and analyzed at the Laboratoire d'Océanographie de 207 Villefranche (LOV, France) following decarbonatation with a drop of HCl 2N, on an elemental 208 analyzer coupled with an isotope ratio mass spectrometer (EA-IRMS; Vario Pyrocube-Isoprime 209 100, Elementar[©]). A caffeine standard (IAEA-600) was used to calibrate the EA-IRMS.

210 **2.2.2. Total hydrolysable carbohydrates and amino acids**

211 For total hydrolysable carbohydrates and amino acids, samples were taken at t0, t6h, 212 t24h, t48h and t72h at all stations. For total hydrolysable carbohydrates (TCHO) > 1 kDa, 213 samples (20 mL) were filled into pre-combusted glass vials (8 h, 500 °C) and stored at -20 °C 214 pending analysis. Prior to analysis, samples were desalted with membrane dialysis (1 kDa 215 MWCO, Spectra Por) at 1 °C for 5 h. Samples were subsequently hydrolyzed for 20 h at 100 °C 216 with 0.8 M HCl final concentration followed by neutralization using acid evaporation (N₂, for 5 h at 50 °C). TCHO were analysed at GEOMAR using high performance anion exchange 217 218 chromatography with pulsed amperometric detection (HPAEC-PAD), on a Dionex© ICS 3000 219 ion chromatography system following the procedure of Engel and Händel (2011). Two replicates 220 per TCHO sample were analyzed. The variation coefficient between duplicate measurements was 221 7% on average.

222 For total hydrolysable amino acids (TAA), samples (5 mL) were filled into pre-223 combusted glass vials (8 h, 500 °C) and stored at -20 °C. Samples were hydrolyzed at 100 °C for 224 20 h with 1 mL 30% HCl (Suprapur®, Merck) added to 1 mL of sample, and neutralized by acid 225 evaporation under vacuum at 60 °C in a microwave. Samples were analyzed by high 226 performance liquid chromatography (HPLC) using an Agilent 1260 HPLC system following a 227 modified version of established methods (Dittmar et al., 2009; Lindroth and Mopper, 1979). 228 Separation of 13 amino acids with a C18 column (Phenomenex Kinetex, 2.6 µm, 150 x 4.6 mm) 229 was obtained after in-line derivatization with o-phthaldialdehyde and mercaptoethanol. A 230 gradient with solvent A containing 5 % acetonitrile (LiChrosolv, Merck, HPLC gradient grade) 231 in sodium dihydrogenphosphate (Suprapur®, Merck) buffer (pH 7.0) and solvent B being 232 acetonitrile was used for analysis. A gradient from 100% solvent A to 78% solvent A was 233 produced in 50 min. Two replicates per TAA sample were analyzed. The variation coefficient

between duplicate measurements was 8% on average. For TCHO and TAA, instrument blanks were performed with MilliQ water. The detection limit was calculated as 3x the blank value, which is \sim 1 nmol L⁻¹ for both parameters.

237 2.2.3. Transparent exopolymer particles

238 Samples for transparent exopolymer particles (TEP) were taken at t0, t24h and t72h at all 239 stations. The abundance and area of TEP were microscopically measured following the procedure given in Engel (2009). Samples of 10-50 mL were directly filtered under low vacuum 240 241 (< 200 mbar) onto a 0.4 µm Nucleopore membrane (Whatman[©]) filter, stained with 1 mL Alcian Blue solution (0.2 g l^{-1} w/v) for 3 s and rinsed with MilliQ water. Filters were mounted on 242 243 Cytoclear[©] slides and stored at -20 °C until analysis. Two filters per sample with 30 images each 244 were analyzed using a Zeiss Axio Scope.A1 (Zeiss[©]) and an AxioCam MRc (Zeiss[©]). The 245 pictures with a resolution of 1388 x 1040 pixels were saved using AxioVision LE64 Rel. 4.8 246 (Zeiss[©]). All particles larger than 0.2 μ m² were analyzed. ImageJ[©] and R were subsequently 247 used for image analysis (Schneider et al., 2012). The coefficients of variation between duplicate 248 filters averaged 28%.

Filters prepared with 10 mL MilliQ water instead of samples served as a blank. Blanks were always <1% of sample values. The carbon content of TEP (TEP-C) was estimated after Mari (1999) using the size-dependent relationship:

$$252 \quad TEP-C = a \,\Sigma_i \, n_i \, r_i^D \tag{1}$$

with n_i being the number of TEP in the size class i and r_i being the mean equivalent spherical radius of the size class. The constant $a = 0.25 * 10^{-6}$ (µg C) and the fractal dimension of

aggregates D = 2.55 were used as proposed by Mari (1999). To relate to organic carbon concentration in seawater, data for TEP-C are given as μ mol L⁻¹.

257 **2.3. Processes**

258 2.3.1. Dissolved and particulate ¹⁴C incorporation rates

259 The photosynthetic production of particulate ($< 0.2-2 \mu m$ and $> 2 \mu m$ size fractions) and 260 dissolved organic matter was determined from samples taken at t0, t24h, t48h and t72h (or t96h 261 at station FAST) with the ¹⁴C-uptake technique. From each tank, four polystyrene bottles (70 262 mL; three light and one dark bottles) were filled with sampled seawater and amended with 40 263 µCi of NaH¹⁴CO₃. Bottles were incubated for 8 h in two extra 300 L tanks maintained under 264 similar light and temperature regimes as in the experimental tanks (ambient temperature for C1, 265 C2, D1 and D2 and ambient temperature + 3 °C for G1 and G2). Incubations were terminated by 266 sequential filtration of the sample through polycarbonate filters (pore sizes 2 μ m and 0.2 μ m, 47 267 mm diameter) using low-pressure vacuum. Filters were exposed for 12 h to concentrated HCl fumes to remove non-fixed, inorganic ¹⁴C, and then transferred to 4 mL plastic scintillation vials 268 269 to which 3.5 mL of scintillation cocktail (Ultima Gold XR, Perkin Elmer[©]) were added. For the 270 measurement of dissolved primary production, a 5 mL aliquot of each sampling bottle was 271 filtered, at the end of incubation, through a 0.2 µm polycarbonate filter (25 mm diameter). This 272 filtration was conducted, under low-pressure vacuum, in a circular filtration manifold that allows 273 the recovery of the filtrate into 20 mL scintillation vials. The filtrates were acidified with 200 μ L 274 of 50% HCl and maintained in an orbital shaker for 12 h. Finally, 15 mL of liquid scintillation 275 cocktail was added to each sample. All filter and filtrate samples were measured onboard in a

liquid scintillation counter (Packard© 1600 TR). ¹⁴C-based production rates (PP; in µg C L⁻¹ h⁻¹)
were calculated as:

278
$$PP = C_{T} x \left(\frac{DPM_{sample} - DPM_{dark}}{DPM_{added} x t} \right)$$
(2)

where $C_{\rm T}$ is the concentration of total dissolved inorganic carbon (µg C L⁻¹), DPM_{sample} and DPM_{dark} are the radioactivity counts in the light and dark bottle, respectively, DPM_{added} is the radioactivity added to each sample, and t is the incubation time (h).

282 The percentage extracellular release (PER%) was calculated as:

$$283 \quad \text{PER}\% = \frac{\text{PPd}}{\text{PPd} + \text{PPp}} \times 100 \tag{3}$$

where PPd refers to ¹⁴C-based dissolved production and PPp refers to ¹⁴C-based particulate production (sum of < 2 and > 2 μ m size fractions).

286 2.3.2. Integrated ¹³C incorporation

Addition of ¹³C-bicarbonate (NaH¹³CO₃ 99%; Sigma-Aldrich[©]) was performed in each 287 288 tank before t0 in order to increase the isotopic level (δ^{13} C signature) of the dissolved inorganic 289 carbon pool to ca. 350‰. We followed the time evolution of the δ^{13} C signature in dissolved 290 inorganic carbon (δ^{13} C- C_T), dissolved organic carbon (δ^{13} C-DOC) and particulate organic carbon 291 pools (δ^{13} C-POC). For the analysis of the actual δ^{13} C-C_T, 60 mL of sampled seawater (at t-12h, 292 t0, t12h, t24h, t48h and t72h; + t96h at station FAST) was gently transferred to glass vials 293 avoiding bubbles. Vials were sealed after being poisoned with 12 µL saturated HgCl₂ and stored 294 upside-down at room temperature in the dark pending analysis. At the University of Leuven, a 295 helium headspace (5 mL) was created in the vials and samples were acidified with 2 mL of

296	phosphoric acid (H ₃ PO ₄ , 99%). Samples were left to equilibrate overnight to transfer all C_T to
297	gaseous CO ₂ . Samples were injected in the carrier gas stream of an EA-IRMS (Thermo©
298	EA1110 and Delta V Advantage), and data were calibrated with NBS-19 and LSVEC standards
299	(Gillikin and Bouillon, 2007).
300	At the same frequency as for δ^{13} C- C_T , samples for δ^{13} C-DOC were filtered online (see
301	above), transferred to 40 mL pre-cleaned borosilicate amber EPA vials with septa caps (PTFE-
302	lined silicone) and stored in the dark pending analysis at the Ján Veizer Stable Isotope
303	Laboratory (Ottawa, Canada).
304	At t-12h, t0, t12h, t24h, t48h and t72h (or t96h at station FAST), the δ^{13} C-POC was
305	obtained based on the same measurements as described above for POC, on an elemental analyzer
306	coupled with an isotope ratio mass spectrometer (EA-IRMS; Vario Pyrocube-Isoprime 100,
307	Elementar [©]).

308 Carbon isotope data are expressed in the delta notation (δ) relative to Vienna Pee Dee
309 Belemnite (VPDB) standard (REF?). The carbon isotope ratio was calculated as:

310
$$R_{\text{sample}} = \left(\frac{\delta^{13}C_{\text{sample}}}{1000} + 1\right) x R_{\text{VPDB}}$$
(4)

311 with $R_{VPDB} = 0.011237$.

2.3.2. Community metabolism (oxygen light-dark method)

313	At the same frequency as for ${}^{14}C$ incorporation, from each tank, a volume of 2 L was
314	sampled in plastic bottles and distributed in 15 biological oxygen demand (BOD; 60 mL)
315	borosilicate bottles. Five BOD bottles were immediately fixed with Winkler reagents (initial O_2
316	concentrations), five BOD bottles were incubated in the dark for the measurement of community
317	respiration (CR) in two incubators maintained respectively at ambient temperature for C1, C2,
318	D1 and D2 and at ambient temperature + 3 °C for G1 and G2. Additionally, five BOD bottles
319	were incubated for the measurement of net community production (NCP) in the same tanks as
320	described above for ¹⁴ C-incorporation. Upon completion of the incubations (24 h), samples were
321	fixed with Winkler reagents. Within one day, O2 concentrations were measured using an
322	automated Winkler titration technique with potentiometric endpoint detection. Analyses were
323	performed on board with a Metrohm© Titrando 888 and a redox electrode (Metrohm© Au
324	electrode). Reagents and standardizations were similar to those described by Knap et al. (1996).
325	NCP and CR were estimated by regressing O2 values against time, and CR was expressed as
326	negative values. Gross primary production (GPP) was calculated as the difference between NCP
327	and CR. The combined standard errors were calculated as:

$$328 \qquad SE_{xy} = \sqrt{SE_x^2 + SE_y^2} \tag{5}$$

329 2.3.4. Heterotrophic prokaryotic production and 330 ectoenzymatic activities

331 At all sampling times, heterotrophic bacterial production (BP, sensus stricto referring 332 to heterotrophic prokaryotic production) was determined onboard using the microcentrifuge 333 method with the ³H- leucine (³H-Leu) incorporation technique to measure protein production 334 (Smith and Azam, 1992). The detailed protocol is in Van Wambeke et al. (2021). Briefly, 335 triplicate 1.5 mL samples and one blank were incubated in the dark for 1-2 h in two 336 thermostated incubators maintained respectively at ambient temperature for C1, C2, D1 and 337 D2 and at ambient temperature +3 °C for G1 and G2. Incubations were ended by the addition 338 of TCA to a final concentration of 5%, followed by three runs of centrifugation at 16000 g 339 for 10 min. Pellets were rinsed with TCA 5% and ethanol 80%. A factor of 1.5 kg C mol leucine⁻¹ was used to convert the incorporation of leucine to carbon equivalents, assuming no 340 341 isotopic dilution (Kirchman et al., 1993). 342 Ectoenzymatic activities were measured fluorometrically, using fluorogenic model 343 substrates that were L-leucine-7-amido-4-methyl-coumarin (Leu-MCA) and 4 344 methylumbelliferyl - phosphate (MUF-P) to track aminopeptidase activity (LAP) and 345 alkaline phosphatase activity (AP), respectively (Hoppe, 1983). Stocks solutions (5mM) 346 were prepared in methycellosolve and stored at -20 °C. Release of the products of LAP and 347 AP activities, MCA and MUF, were followed by measuring increase of fluorescence (exc/em 348 380/440 nm for MCA and 365/450 nm for MUF, wavelength width 5 nm) in a 349 VARIOSCAN LUXmicroplate reader calibrated with standards of MCA and MUF solutions. 350 For measurements, 2 mL of unfiltered samples from the tanks were supplemented with 100

351 µL of a fluorogenic substrate solution diluted so that different concentrations were 352 dispatched in a black 24-well polystyrene plate in duplicate $(0.025, 0.05, 0.1, 0.25, 0.5, 1 \,\mu\text{M})$ 353 for MUF-P, 0.5, 1, 5, 10, 25 µM for MCA-leu). Incubations were carried out in the same 354 thermostatically controlled incubators than those used for BP and reproducing temperature 355 levels in the experimental tanks. Incubations lasted up to 12 h long with a reading of 356 fluorescence every 1 to 2 h, depending on the intended activities. The rate was calculated from the linear part of the fluorescence versus time relationship. Boiled-water blanks were 357 358 run to check for abiotic activity. From varying velocities obtained, we determined the 359 parameters Vm (maximum hydrolysis velocity) and Km (Michaelis-Menten constant which 360 reflects enzyme affinity for the substrate) by fitting the data using a non-linear regression on 361 the following equation:

$$362 \qquad \mathbf{V} = \mathbf{V}_{\mathrm{m}} \, \mathbf{x} \frac{\mathbf{S}}{\mathbf{K}_{\mathrm{m}} + \mathbf{S}} \tag{6}$$

363 where V is the hydrolysis rate and S the fluorogenic substrate concentration added.

2.3.5. Inorganic and organic material export

At the end of each experiment (t72h for TYR and ION and t96 h for FAST, after artificial dust seeding), the sediment traps were removed, closed and stored with formaldehyde 4%. Back in the laboratory, after the swimmers were removed, the samples were rinsed to remove the salts and then freeze-dried. The total amount of material collected was first weighted to measure the total exported flux. Several aliquots were then weighted to measure the following components: total carbon and organic carbon, lithogenic and biogenic silicates and calcium. Total carbon was measured on an elemental analyzer coupled with an isotope ratio mass spectrometer (EA-IRMS; 372 Vario Pyrocube-Isoprime 100, Elementar[©]). Particulate organic carbon (POC) was measured in 373 the same way after removing inorganic carbon by acidification with HCl 2N. Particulate 374 inorganic carbon (PIC) was obtained by subtracting particulate organic carbon from particulate 375 total carbon. Calcium concentrations were measured by ICP-OES (Inductively Coupled Plasma -376 Optic Emission Spectrometry; Perkin-Elmer[©] Optima 8000) on acid digested samples (the 377 organic matrix was removed by HNO₃ while the mineral aluminosilicate matrix was eliminated 378 with HF). Biogenic silica (BSi) and Lithogenic silica (LSi) were measured by colorimetry 379 (Analytikjena[©] Specor 250 plus spectrophotometer) after a NaOH/HF digestion, respectively 380 (Mosseri et al., 2005). The carbonate fraction of the exported material was determined from 381 particulate calcium concentrations (%CaCO₃ = $5/2 \times$ (%Ca). The organic matter fraction was 382 calculated as 2 x %POC (Klaas and Archer, 2002) The lithogenic fraction was calculated as 383 [total mass – (organic matter + $CaCO_3$ + opal) and was very comparable to the lithogenic 384 fraction calculated from LSi (taking Si concentration in dust analog used for seeding from 385 Desboeufs et al., 2014; ca. 11.9%). In the controls, the amount of material exported was low and 386 the entire content of the traps was filtered in order to measure total mass and organic matter mass 387 fluxes.

388

389 **2.4. Data processing**

All metabolic rates were integrated over the duration of the experiments using trapezoidal
 integrations and the relative changes (in %) in tanks D and G as compared to the controls
 (average between C1 and C2) were computed following:

393 Relative change =
$$\left(\frac{\text{Rate}_{\text{Treatment}} - \text{Rate}_{\text{Controls}}}{\text{Rate}_{\text{Controls}}}\right) \times 100$$
 (7)

394 Where Rate_{Treatment} is the integrated rate measured in treatments D and G (D1, D2, G1 or G2) and 395 Rate_{Controls} is the averaged integrated rates between the duplicate controls (treatment C). Daily 396 rates of ¹⁴C-based production were computed from hourly rates assuming a 14 h daylight period. 397 As incubations performed from samples taken at t0 (before dust addition) do not represent what 398 happened in the tanks between t0 and t24h, as a first assumption, we considered a linear 399 evolution between these rates and those measured from samples at t24h, and recomputed an average value for the time interval t0 - t24 h. At FAST, no incubations were performed for ¹⁴C 400 401 incorporation and oxygen metabolism between t72h and t96h, again an average rate between 402 rates measured from samples taken at t48h and t96h was used for this time interval. Since 403 bacterial respiration rates were not measured, bacterial growth efficiency (BGE, expressed as a 404 percentage) was estimated based on BP (carbon units) and community respiration (CR, oxygen 405 units). As BP was determined more often than CR during the first 48 h, hourly BP rates were 406 integrated using trapezoidal integrations during the time period when CR was measured. We 407 assumed that heterotrophic prokaryotes were responsible for 70% of CR (BR/CR ratio; Lemée et 408 al., 2002) and used a respiratory quotient (RQ) of 0.8 (del Giorgio and Williams, 2005), 409 following the equation:

410 BGE =
$$\left(\frac{BP}{CR \times \frac{BR}{CR} \operatorname{ratio} \times RQ + BP}\right) \times 100$$
 (8)

411 When BP varied following an exponential growth, we calculated growth rates (μ_{BP}) from linear 412 least square regression of ln BP rates versus time.

413 **3. Results**

414 **3.1. Initial conditions**

415 Initial conditions in terms of the chemical and biological standing stocks measured while 416 filling the tanks at the three stations are fully described in Gazeau et al. (2021). Briefly, the three 417 experiments were conducted with surface seawater collected during stratified oligotrophic 418 conditions typical of the open Mediterranean Sea at this period of the year (Table 1). 419 Nevertheless, a dust event took place nine days before sampling at station TYR as evidenced 420 from particulate inventory of lithogenic proxies (Al, Fe) in the water column (Bressac et al., 421 2021), likely stimulating phytoplankton growth before the start of the experiment. Nitrate + 422 nitrite (NO_x) concentrations were maximal at station FAST with a NO_x to dissolved inorganic 423 phosphate (DIP) molar ratio of ~ 4.6. Very low NO_x concentrations were observed at stations TYR and ION (14 and 18 nmol L⁻¹, respectively). DIP concentrations were the highest at station 424 425 TYR (17 nmol L^{-1}) and the lowest at the most eastern station (ION, 7 nmol L^{-1}). Consequently, 426 the lowest NO_x:DIP ratio was measured at TYR (0.8), compared to ION and FAST (2.8 and 4.6, 427 respectively). Silicate (Si(OH₄)) concentrations were similar at TYR and ION (~1 μ mol L⁻¹) and 428 the lowest at FAST (~0.6 µmol L⁻¹). Both POC and DOC concentrations were the highest at 429 station TYR (12.9 and 72.2 µmol L⁻¹, respectively) and the lowest at FAST (6.0 and 69.6 µmol 430 L^{-1} , respectively). Very low and similar concentrations of chlorophyll *a* were measured at the 431 three stations (0.063 - 0.072 µg L⁻¹). Phytoplankton communities at stations TYR and ION were 432 dominated by Prymnesiophytes followed by Cyanobacteria, while, at station FAST, the 433 phytoplanktonic community was clearly dominated by photosynthetic prokaryotes. At all three 434 stations, the proportion of pigments representative of larger species was very small (< 5%;

Gazeau et al., 2021). Heterotrophic prokaryotes were the most abundant at station FAST (6.15 x 10^5 cells mL⁻¹) and the least abundant at station ION (2.14 x 10^5 cells mL⁻¹).

437 Relatively similar ¹⁴C-based particulate production rates were measured at the start of the 438 experiments (t0) in the control tanks (C1 and C2) at station ION and FAST (ca. 0.014 - 0.015 µg 439 C L⁻¹ h⁻¹). At both stations, ca. 80% of the production was attributed to larger (> 2 μ m) cells and 440 the percentage of extracellular release (%PER) did not exceed 45%. Lower rates were estimated at station TYR (total particulate production of 0.08 µg C L⁻¹ h⁻¹) from which 87.5% was due to 441 large cells $> 2 \mu m$. A larger amount of ¹⁴C incorporation was released as dissolved organic 442 443 matter at station TYR compared to the two other stations (PER ca. 60%). Metabolic balance 444 derived from oxygen measurements showed that, at all three stations, the community was net 445 heterotrophic with a higher degree of heterotrophy at station TYR (NCP were -1.9, -0.2, -0.8 446 µmol O₂ L⁻¹ d⁻¹ at TYR, ION and FAST, respectively, as measured in the controls from seawater 447 sampled at t0). CR and GPP rates were respectively the highest and the lowest at station TYR 448 compared to the other two stations. Finally, BP rates were the highest at station FAST (35.8 ng C $L^{-1} h^{-1}$, intermediate at ION (26.1 ng C $L^{-1} h^{-1}$) and the lowest at TYR (21.3 ng C $L^{-1} h^{-1}$). 449

450 **3.2. Changes in biological stocks**

451 DOC concentrations showed a general positive trend during the three experiments and a 452 large variability between duplicates (Fig. 2). This variability appeared as soon as 1 h after dust 453 seeding (t1h) while the range of variation at t0 (before dust seeding) was rather moderate 454 (difference between minimal and maximal values in all tanks of 1.3, 6.2 and 4.3 μ mol C L⁻¹ at 455 station TYR, ION and FAST, respectively). As a consequence of this variability, no clear impact 456 of dust seeding (D) could be highlighted at station TYR and FAST. Indeed, DOC concentrations 457 in the two duplicates (D1 and D2) were higher than values in the controls (C1 and C2) in only 458 33% of the samples along the experiments (after dust seeding). In contrast, at station ION, DOC 459 concentrations appeared impacted by dust seeding as higher concentrations were almost 460 systematically (83% of the time after dust seeding) measured for this treatment as compared to 461 control tanks at the same time. At all stations, this impact was somewhat exacerbated under 462 conditions of temperature and pH projected for 2100 (G1 and G2) as DOC concentrations were 463 almost all the time higher in these tanks than in control tanks (83 - 100% of the samples after 464 dust seeding, depending on the station).

465 Total hydrolysable carbohydrates and amino acids concentrations along the three 466 experiments are shown in Fig. S2. TCHO concentrations were quite variable between tanks before dust seeding (t0; 649 - 954, 569 - 660 and 600 - 744 nmol L⁻¹ at station TYR, ION and 467 468 FAST, respectively) and no visible impact of the treatments were visible at station TYR (TCHO 469 tended to decrease everywhere). In contrast, at station ION and FAST, values in dust amended 470 tanks increased and appeared higher than in control tanks towards the end of the experiments 471 although the large variability between duplicates tended to mask this potential effect. An impact 472 of dust seeding was much clearer for TAA concentrations that showed larger increases 473 throughout the three experiments in tanks D1 and D2 as compared to control tanks, this effect 474 being exacerbated for warmer and acidified tanks (G1 and G2). The ratio between TAA and 475 DOC concentrations (Fig. 2) showed positive trends in tanks D and G during all three 476 experiments with a clear distinction between treatments at the end of the experiments (G > D >477 C). The strongest increase was observed at station FAST in tanks G where final TAA/DOC ratios 478 were above 3%.

479 Particulate organic carbon (POC) concentrations strongly decreased at all stations
480 between t-12h and t0, this decrease being the largest at station TYR where concentrations

481 dropped from 25.7 to 9.6 - 13.2 µmol C L⁻¹ (Fig. 3). After dust seeding, POC concentrations did 482 not show clear temporal trends for the three experiments although a slight general increase could 483 be observed at station FAST. Furthermore, no impact of dust seeding and warming/acidification 484 could be observed on POC dynamics. While concentrations of transparent exopolymer particles 485 (TEP-C) were rather constant through time in control tanks at the three stations, a large increase 486 was observed in dust-amended tanks (D and G) with TEP-C reaching values up to $\sim 2 \mu mol C L^{-1}$ 487 in tank G1 at station TYR after 24 h (i.e. ~17% of POC concentration, Fig. 3). In all cases except 488 for tank G2 at station ION, TEP-C further decreased towards the end of the experiments although 489 concentrations remained well above those observed in the controls. As the variability between 490 duplicated tanks G was rather high, no impact of warming/acidification on TEP dynamics could 491 be highlighted at the three stations.

492 **3.3. Changes in metabolic rates**

¹⁴C-based particulate production rates as measured during the different time intervals at the three stations were low in control tanks (maximal total particulate production of $0.34 \ \mu g \ L^{-1}$ h⁻¹ at station FAST) and did not show any particular temporal dynamics (Fig. 4). In these tanks, the vast majority of particulate production was attributed to cells above 2 μ m (65 - 89%). The percentage of extracellular release (%PER) was overall maximal at station TYR and minimal at station FAST with a tendency to decrease with time at the three stations although large variations were observed between duplicates.

500 Dust addition alone did not have any clear positive impact on all ¹⁴C-based rates at 501 station TYR, with even an observable decrease in production rates from larger cells (> 2 μ m) 502 compared to the controls. In contrast, at this station, dust seeding under warmer and acidified 503 conditions (tanks G) had a positive effect on particulate production rates, this effect being 504 particularly visible for cells $< 2 \mu m$ and to a lesser extent on dissolved production with a general 505 decrease of %PER. An important discrepancy between the duplicates of treatment G was 506 observable at the end of the experiment with much larger rates measured in tank G2. 507 In contrast to station TYR, an enhancement effect of dust addition was clearly visible at 508 station ION where all rates increased towards the end of this experiment reaching a maximal 509 total particulate production of 0.6 - 0.7 µg L⁻¹ h⁻¹ in tanks D1 and D2. Since this positive effect 510 was similar between small and larger cells, dust addition alone had no effect on the partitioning 511 of production at this station, with cells > 2 μ m representing ~80% of total production. Although 512 being also positively impacted and increasing with time, dissolved production appeared less 513 sensitive than particulate production leading to an overall decrease of %PER at station ION 514 following dust addition. These positive impacts of dust seeding on ¹⁴C-based particulate 515 production rates were even more visible at this station under warmer and acidified conditions 516 (tanks G) with maximal rates more than doubled compared to those measured under present 517 conditions of temperature and pH (1.5 - 1.6 µg L⁻¹ h⁻¹). Dust seeding under warmer and acidified 518 conditions had a slight impact on the partitioning of particulate production at station ION with 519 smaller cells benefiting the most from these conditions. %PER remained between 20 and 30%. 520 At station FAST, similarly to station ION, total particulate production rates were clearly 521 enhanced by dust addition (tanks D) reaching maximal values during the incubation time interval 522 t48 - 56h. No clear increase was observed for total particulate production on the next incubation 523 (t96 - 120h) while production rates of cells larger than 2 µm increased and rates of smaller cells 524 decreased. However, in contrast to station ION, there was much less impact of 525 warming/acidification on all measured rates at station FAST although rates measured on smaller 526 cells (< 2 µm) did not decrease at the end of the experiment as observed under present

527	environmental conditions. %PER under both present conditions of temperature and pH (tanks D)
528	decreased during this experiment reaching values lower than in the controls and in tanks G.
529	The initial enrichment of the tanks in ¹³ C-bicarbonate led to an increase in the ¹³ C
530	signature of dissolved inorganic carbon (δ^{13} C- C_T) of above 300‰, with generally lower values
531	measured in warmer and acidified tanks (G; Fig. S3). After this initial enrichment, δ^{13} C- C_T levels
532	decreased linearly in all tanks. At stations TYR and ION, the isotopic signature of dissolved
533	organic carbon (δ^{13} C-DOC; Fig. S3) increased with time, although these increases were rather
534	low and limited to $\sim 4\%$ over the course of the experiments. In contrast to station TYR, at ION,
535	an enhanced incorporation of ¹³ C into DOC was visible after 24 h in tanks D and G in
536	comparison to control tanks. A similar observation was done at station FAST, especially at the
537	end of the experiment, although much more variability was observed at this station.
538	The incorporation of ¹³ C into particulate organic carbon (δ^{13} C-POC) is shown in Fig. 5.
539	At all stations, δ^{13} C-POC increased with time but reached lower enrichment levels at station
540	TYR as compared to ION and FAST. At station TYR, incorporation rates appeared smaller in
541	dust-amended tanks under present environmental conditions (tanks D). As for ¹⁴ C-based
542	production rates, an important discrepancy was observed between duplicates under future
543	conditions of temperature and pH (tanks G) with much higher final δ^{13} C-POC at the end of the
544	experiment in tank G2. At station ION, enrichment levels obtained at the end of the experiment
545	were more important in dust-amended tanks reaching maximal levels of 73‰ in tank G2 at t72h.
546	This enhancement effect was even more visible at station FAST with maximal enrichment levels
547	of 146‰ (tank D2 at t96h). Since no sampling occurred at t72h, these enrichment levels cannot
548	be directly compared to what was measured at station TYR and ION. However, by interpolating

values at t72h assuming a linear increase between these time intervals, enrichment levels
appeared similar although slightly higher for tanks D between station ION and FAST.

551 NCP rates as measured using the O₂ light-dark method showed that, under control 552 conditions, the communities remained the vast majority of the time throughout the three 553 experiments in a net heterotrophic state (NCP ≤ 0 ; Fig. 6). This was especially true at station 554 TYR where the lowest NCP rates were measured. At this station, dust addition whether under 555 present or future conditions of temperature and pH did not switch the community towards net 556 autotrophy but even drove the community towards a stronger heterotrophy. This was related to 557 the fact that while gross primary production rates were not positively impacted, community 558 respiration increased in tanks D and G. At station ION, dust addition alone (tanks D) led to a 559 switch from net heterotrophy to net autotrophy after two days of incubation due to a stronger 560 positive effect of dust on GPP than on CR. Under future environmental conditions (tanks G), the 561 same observation was made with higher NCP and GPP rates than in tanks D. CR rates reacted 562 quickly to these forcing factors in tanks G and initially (first incubation) drove the community 563 towards a much stronger heterotrophy as compared to the other tanks. Finally, at station FAST, 564 similarly to what was observed at ION, the community became autotrophic after two days of 565 incubation in dust amended tanks as, although both GPP and CR were positively impacted by 566 dust addition, this impact was less important for CR. Warming and acidification had a limiting 567 impact on this enhancement, with a lower final NCP in tanks G compared to tanks D, a 568 difference that can be related to an absence of effects of these environmental stressors on GPP 569 while CR clearly increased at higher temperature and lower pH.

570 While BP remained constant or gradually increased in control tanks depending on the 571 station, a clear and quick fertilization effect was observable following dust addition (treatment D

572 and G) at all stations (Fig. 7). At station TYR, BP rates sharply increased to reach maximal 573 values at t24h, with an even stronger increase observed under warmer and acidified conditions 574 (tanks G). After this initial increase, rates slightly decreased towards the end of the experiment. 575 This fertilization effect appeared less important at station ION where lower maximal rates were 576 obtained after 24 h as compared to station TYR. Nevertheless, the same observations can be 577 made, namely, 1) higher rates were measured under future temperature and pH levels and 2) after 578 this initial sharp increase, rates gradually decreased towards the end of the experiment especially 579 in tanks G. At station FAST, a much stronger effect of warming/acidification was observed with 580 an important increase of BP in tanks G until 24 or 48 h post-seeding, depending on the duplicate. 581 A sharp decline was observed for this treatment until the end of the experiment although rates 582 remained higher than those measured in tanks C and D. The impact of dust addition under 583 present environmental conditions (tanks D) was somehow more limited than at the other stations 584 with a gradual increase until t72h with maximal rates $\sim 40 - 100\%$ higher than rates measured in 585 the controls. However, BP increased exponentially between t0 and t12h in all tanks including 586 controls, and in all experiments (Table 2). The growth rate of BP (μ_{BP}) in control tanks was the 587 highest at TYR, intermediate at ION and the lowest at FAST. µ_{BP} increased significantly in all 588 dust amended tanks compared to controls. Under future environmental scenarios, μ_{BP} tended to 589 increase compared to treatment D but with a variable relative change (Table 2). 590 BGE increased in dust amended tanks under present environmental conditions (treatment 591 D) at TYR and ION, while no changes were detectable at station FAST due to a strong 592 discrepancy between control duplicates and overall higher BGE at this station in the controls

593 (Table 3). In contrast, warming and acidification exerted the strongest effect at station FAST

594 with a doubling of BGE between treatment G and D. Although an increase in BGE was also

595	observed at the two other stations in treatment G as compared to present environmental
596	conditions (treatment D), this increase was more limited (ca. 1 to 1.4-fold increase).
597	The alkaline phosphatase Vm (AP Vm) increased in all experiments after dust seeding,
598	with amplified effects in G treatments (Fig. S4). Note that AP Vm increased also in the controls
599	at TYR and FAST. In contrast, leucine aminopeptidase Vm (LAP vm) showed succession of
600	peaks instead of a continuously increase (Fig. S4). It was higher in dust alone treatment (D) as
601	compared to the controls at TYR and FAST. A larger variability between duplicates at ION
602	prevents such an observation. At all stations, maximum velocities were measured under future
603	environmental conditions (G). Vm being possibly influenced by enzyme synthesis but also by the
604	number of cells inducing such enzymes, we computed also specific AP Vm per heterotrophic
605	bacterial cell (Fig. 7). Specific AP Vm slightly increased during all experiments in controls and
606	dust-amended tanks (D) with no visible differences between these treatments, a clear over-
607	expression of this enzyme was observed under warmer and more acidified conditions (treatment
608	G) especially at station FAST where velocities were enhanced by a ~8-fold at t96h.

609

3.4. Inorganic and organic material export

610 Both total mass and organic matter fluxes, as measured from analyses of the sediment 611 traps at the end of each experiment, were extremely low under control conditions (Fig. 8). Only 612 less than 30% of the dust introduced at the surface of the tanks were recovered at the end of the 613 experiment (3 or 4 days after) in the sediment traps with TYR>ION>FAST. The composition of 614 the exported material was quite similar for each experiment with no significant difference 615 between D and G treatments with 3-5% opal, 4% organic matter, 35-36% CaCO₃ and 48-54% 616 lithogenic (Fig. S5). Additions of dust in tanks D and G led to a strong increase in both fluxes 617 with a large variability between the duplicates of treatment D at ION. No clear changes between

- 618 tanks maintained under present and future conditions of temperature and pH could be
- 619 highlighted.

620 **4. Discussion**

621 4.1. Initial conditions of the tested waters and evolution in

622 controls

623 As discussed in details in the companion paper from Gazeau et al. (2021), the three 624 sampling stations were typical of stratified (mixed layer depth of 10-20 m) oligotrophic 625 conditions encountered in the open Mediterranean Sea in late spring / early summer. Briefly, the 626 low NO_x:DIP ratios nutrient concentrations suggest that communities found at the three stations 627 experienced N and P co-limitation at the start of the experiments. The composition of the smaller 628 size phytoplankton communities differed substantially, with autotrophic nano-eukaryotes 629 dominating at stations TYR and ION and a larger contribution from autotrophic pico-eukaryotes 630 and Cyanobacteria at station FAST. The observed low total chlorophyll *a* concentrations and the 631 small contribution of large phytoplankton cells at the start of the three experiments are 632 characteristic of LNLC areas in general, and of surface Mediterranean waters in late spring and 633 summer (Siokou-Frangou et al., 2010). DOC concentrations at the start of the experiments were in the same range (60 - 75 μ mol C L⁻¹) as those measured from samples collected in surface 634 635 waters using clean sampling procedures (Van Wambeke et al., 2021), revealing no contamination 636 issues from our sampling device. TAA concentrations as measured in the tanks at t0 were also 637 consistent with measurements from surface water samples (Van Wambeke et al., 2021) with an 638 average across stations and treatments of 254 ± 36 nmol L⁻¹ (Fig. S2). In contrast, TCHO 639 appeared higher at t0 (average across stations and treatments of 681 ± 98 nmol L⁻¹) than 640 concentrations based on clean *in situ* sampling (average of 595 ± 43 nmol L⁻¹; Van Wambeke et 641 al., 2021). The decrease in POC concentrations between pumping (t-12h) and t0 for the three

642 experiments, especially at station TYR (likely linked to higher initial concentrations), was likely 643 a consequence of sedimentation of senescent cells and/or fecal pellets in our experimental 644 systems, which are designed to evaluate the export of matter thanks to their conical shape. TEP 645 concentrations were not quantified at t-12h and therefore there is no possibility to evaluate if 646 sedimentation of these particles occurred before t0 in our tanks. At t0, larger and more abundant 647 TEP were measured at station TYR compared to the two other stations (data not shown). As a consequence of a very low availability in inorganic nutrients, TChla and ¹⁴C-based 648 649 production rates were very low, all typical of oligotrophic conditions. Nano- and micro-650 phytoplanktonic cells (> 2 μ m) contributed most of the ¹⁴C-based particulate production (~ 80%), 651 as found also on several on-deck incubations at the three stations (on average $73 \pm 6\%$; Marañón 652 et al., 2021). %PER values were also very similar to those measured during these on-deck 653 incubations (~40-45%; see Marañón et al., 2021). This suggests no significant impact of our experimental protocol on rates and partitioning of ¹⁴C-based production rates (i.e. sampling from 654 655 the continuous seawater supply, delay of 12 h before initial measurements, artificial light etc.). 656 The low values of chlorophyll stocks as well as of ¹⁴C-based production rates are consistent with 657 previous estimates based on direct measurements, satellite observations and modelling

approaches in the same areas in late spring / early summer (e.g. Bosc et al., 2004; Lazzari et al.,

659 2016; Moutin and Raimbault, 2002).

660 The metabolic balance was in favor of net heterotrophy at all stations at the start of the 661 experiments (NCP < 0). Net heterotrophy in the open Mediterranean sea at this period of the year 662 has been reported by Regaudie-de-Gioux et al. (2009) and Christaki et al. (2011) in agreement 663 with our measurements at t0 in control tanks (Table 1). The lowest NCP and the highest CR rates 664 were measured at station TYR, suggesting that the autotrophic plankton community was not very

665 active at this station. This was confirmed by the ¹⁴C-based particulate production rates, which 666 were about half the ones measured at the other two stations. The community at TYR was most 667 likely relying on regenerated nutrients, as shown by the highest levels of ammonium (NH4⁺) 668 measured at the start of this experiment (Gazeau et al., 2021). As discussed in Guieu et al. 669 (2020a), a dust deposition event took place several days before the arrival of the vessel in this 670 area, likely on May 10-12. This dust event was confirmed by inventory of particulate aluminium 671 in the water column at several stations of the Tyrrhenian Sea including TYR, 6 to 9 d after the 672 event (Bressac et al., 2021). This dust deposition likely stimulated phytoplankton growth and 673 POC accumulation shortly after the deposition and consequently increased the abundance of 674 herbivorous grazers (copepods) and attracted carnivorous species (Feliú et al., 2020), 675 subsequently driving the community towards a net heterotrophic state that characterized the 676 initial condition of the experiment at this station. The favorable conditions for BP growth at TYR 677 were also confirmed by the highest μ_{BP} growth rates obtained among the three experiments 678 (Table 2; $0.06 - 0.07 h^{-1}$) in controls tanks. 679 The two other stations, although both also showing a slight net heterotrophic state, were 680 clearly different from each other in terms of initial biological stocks and metabolic rates. Indeed,

681 whereas TChla and abundances of pico- and nano-autotrophic cells (flow cytometry counts;

682 Gazeau et al., 2021) were higher at FAST compared to ION, the autotrophic community was not

683 more efficient at fixing carbon at FAST, as shown by similar initial ¹⁴C-based production rates.

In contrast, both heterotrophic prokaryotic abundances and BP were much higher at station

685 FAST as compared to ION, leading to initial higher CR and lower NCP. At ION, the initial NCP

686 closer to metabolic balance further suggests a tight coupling between heterotrophic prokaryotes

and phytoplankton at this station, as discussed by Dinasquet et al. (2021).

688 For most of the chemical and biological stocks (e.g. nutrients, pigments etc.) presented in 689 Gazeau et al. (2021), no major changes took place during the three experiments under control 690 conditions. Here, we further show that DOC, POC as well as TEP concentrations did not exhibit 691 strong changes during the experiments. For DOC, large variability between the duplicates (C1 692 and C2) potentially masked an increase towards the end of the experiments. The same holds true 693 for autotrophic metabolic rates, as ¹⁴C-based particulate production rates showed no marked 694 variations during the three experiments, although a slight increase was visible at FAST until 695 t48h. The communities at the three stations remained heterotrophic under the nutrient-limited 696 conditions in the controls. However, heterotrophic prokaryotes probably benefited from initial 697 inputs of available organic matter issued from other stressed eukaryotic organisms and/or POC 698 decay between t-12h and t0, which could be due to both sedimentation and degradation. This was 699 reflected in the progressive increase of BP, their variable initial growth rates (μ_{BP} ranged from 0.02 to 0.06 h⁻¹ in control tanks according to the experiment) as well as increasing TAA/DOC 700 701 ratios at the three stations. Finally, an initial increase of BP during incubations is generally 702 described and classically attributed to a bottle effect, which favours large, fast-growing bacteria 703 and often induces mortality of some phytoplankton cells (Calvo-Díaz et al., 2011; Ferguson et 704 al., 1984; Zobell and Anderson, 1936)

705 4.2. Impact of dust addition under present environmental

706 conditions

The addition of nitrogen and phosphorus in the experimental tanks through dust seeding (+ 11 to + 11.6 μ mol L⁻¹ and + 22 to + 30.8 nmol L⁻¹ for NO_x and DIP, respectively, in dust enriched, i.e. D1 and D2, versus controls; Gazeau et al., 2021) had very contrasting impacts on

710	the metabolism of the communities, depending on the station. At TYR, surprisingly, the relieving
711	of nutrient (N, P) limitation had a negative impact on ¹³ C incorporation as well as on both
712	particulate and dissolved ¹⁴ C-based production rates (as seen by the relative changes compared to
713	the control presented in Fig. 9). These observations are fully corroborated by the observed
714	relative decrease in GPP in these tanks (D1 and D2) relative to controls and by the negative
715	impact of dust-addition on TChla concentrations as discussed by Gazeau et al. (2021). Integrated
716	¹⁴ C-incorporation rates converted to P (using a C:P molar ratio of 245:1 determined in the
717	particulate organic matter in surface waters of the Northwestern Mediterranean Sea during
718	stratification; Tanaka et al., 2011) showed that phytoplankton P requirements in treatment D (~2
719	nmol P L ⁻¹) were much lower than the release of DIP through dust addition at station TYR (+
720	20.4 to + 24.6 nmol P L ⁻¹ ; Gazeau et al., 2020). This suggests that the observed strong decrease
721	of DIP at TYR following dust addition was due to an utilization by the heterotrophic
722	compartment. Indeed, in contrast to the autotrophic compartment, both heterotrophic prokaryotic
723	abundances (Gazeau et al., 2021) and BP (this study, Fig. 9) showed that heterotrophic
724	prokaryotes reacted quickly and strongly to the increase in DIP availability. Integrated BP
725	increased by almost 400% in tanks D1 and D2 as compared to controls (Fig. 9). Such relative
726	increases of BP surpassing by far the observed relative increases of CR suggest a much more
727	efficient utilization of resources by heterotrophic prokaryotes in this treatment (i.e. BGE
728	increased by 200% as compared to the controls; Fig. 9). As such, at TYR, the addition of dust
729	drove the community to an even more heterotrophic state. Such absence of response of the
730	autotrophic community despite the input of new N and P from simulated wet deposition was
731	never observed in dust enrichment experiments performed in the Mediterranean Sea (Guieu and
732	Ridame, 2020). To the best of our knowledge, it is the first time that a negative effect of dust

733 addition is experimentally demonstrated on the metabolic balance. The apparent utilization of 734 nutrients, especially DIP (Gazeau et al., 2021), by heterotrophic prokaryotes was extremely fast, 735 starting right after dust addition and driving DIP concentrations back to control levels at the end 736 of the experiment (t72h). While heterotrophic prokaryotic abundances increased until the end of 737 the experiment, BP rates increased exponentially during the first 24h, and then BP reached a 738 plateau. Heterotrophic prokaryotes appeared limited by nutritive resources although DIP 739 concentrations were not yet back to their initial level and no relative increase of the AP Vm per 740 cell compared to the control was observed in these tanks. Independent nutrient experiments 741 showed a direct stimulation of BP in the dark after addition of DIP (Van Wambeke et al., 2021), 742 suggesting a great competition with phytoplankton for DIP utilization at TYR. After 24 h, 743 abundances of heterotrophic prokaryotes continued to increase while BP stabilized, suggesting a 744 less extent of lysis and viral control than in the other experiments (abundances of heterotrophic 745 nanoflagellates decreased; Dinasquet et al., 2021). This limitation of BP was potentially a 746 consequence of relatively less available access to labile DOC sources, as ¹⁴C-based production 747 rates decreased relative to the controls at t24h and t48h although BP increased by 200 - 800%. 748 The very tight coupling between phytoplankton and bacteria at all stations investigated was 749 further confirmed by the absence of an important ¹³C incorporation into DOC (Fig. S3). 750 At stations ION and FAST, in contrast to TYR, both the autotrophic and heterotrophic 751 community benefited from dust addition relative to the controls (Fig. 9). Interestingly, while the 752 relative increase in integrated autotrophic processes (GPP and all ¹⁴C-based production rates)

was more important at FAST than at ION, the opposite was observed for BP. Estimated BGE

values even suggest an absence of response to dust addition at station FAST compared to the

controls. The different (relative) responses of BP at the two stations could be partly explained by
the dynamics of BP in the control tanks as no clear pattern could be observed at ION while a continuous increase was observed at FAST. As shown by Gazeau et al. (2021), at FAST, abundances of heterotrophic prokaryotes were much higher at the start of the experiment, further increased until t48h and then declined until the end of the experiment.

760 We can rule out a potential limitation of BP from DIP availability at station FAST as DIP 761 levels remained much higher in tanks D than in the controls (Gazeau et al., 2021). Furthermore, 762 the amount of maximum DIP reached before its decline compared to TYR and ION showed a 763 less important direct DIP uptake, suggesting that communities were not as much P limited at 764 FAST compared to the other stations at the start of the experiment. Finally, no increase of 765 specific AP Vm was observed in these tanks as compared to the controls (Fig. 7), suggesting no 766 particular additional needs for AP synthesis per unit cell following dust addition. A potential 767 explanation resides in the competition between heterotrophic bacteria and phytoplankton for DIP 768 utilization. At station ION, P requirements of the autotrophic community were low compared to the initial input of DIP following dust seeding (~9 nmol P L^{-1} as compared to an input of + 22 to 769 770 + 23.3 nmol P L⁻¹; Gazeau et al., 2021). In contrast, at FAST, the autotrophic community 771 consumed a much larger proportion of the initial DIP input (~25 nmol P L⁻¹ as compared to an input of 30.8 - 31.3 nmol P L⁻¹) and phytoplankton appeared as a winner for the utilization of 772 773 DIP towards the end of the experiment at this station. It seems that heterotrophic bacteria and 774 phytoplankton were more in balance and less stressed at the start of the experiment at FAST, i.e. 775 phytoplankton abundances showed no decrease between t-12h and t0 and BP did not increase as 776 much as during the other two experiments, suggesting a strong predation pressure (μ_{BP} was the lowest of the three experiments: ca. 0.02 h⁻¹ in the controls). 777

778 The explanation for the observed differential responses of the autotrophic community at 779 the two stations (FAST > ION) is not evident and further complicated by the fact that the 780 sampling strategy differed between the two stations (i.e. no sampling at t72h, replaced by a 781 sampling at t96h). It is however unlikely that this different sampling strategy was responsible for 782 the different changes in computed integrated autotrophic rates at the two stations. As a maximal 783 increase in nano-eukaryote abundance was observed at t72h at FAST (followed by a drastic 784 reduction at t96h; Gazeau et al., 2021), excluding this sampling point in the calculation of 785 autotrophic metabolic rates would most likely have led to an underestimation of these rates rather 786 than an overestimation. Furthermore, a similar partitioning of ¹⁴C-based production rates 787 throughout the two experiments did not provide clear insights on which size-group benefited the 788 most at station FAST compared to ION. Two non-exclusive explanations could be proposed: (1) 789 as mentioned above, a less important immediate consumption of DIP by heterotrophic bacteria 790 leading to a higher availability of new DIP for phytoplankton growth at FAST (+ 31 vs + 22 to + 23 nmol L⁻¹ at FAST and ION, respectively; Gazeau et al., 2021) along with (2) the presence of a 791 792 potentially more active community at the start of the experiment at FAST with a much higher 793 contribution from smaller cells (i.e. pico-eukaryotes, Synechococcus; Gazeau et al., 2021) that 794 are well known to be better competitors for new nutrients (e.g. Moutin et al., 2002) and that were 795 less stressed at the start of the experiments.

During both experiments at ION and FAST, communities switched from net heterotrophy to net autotrophy between 48 and 72 h following dust addition (Fig. 6), leading to a positive integrated NCP at both stations (Fig. 9). This is an important observation since, to the best of our knowledge, the present study constitutes the first investigation of the community metabolism response to dust addition. However, it is important to discuss the timing of such a switch in

801 community metabolism. Since heterotrophic prokaryotes reacted faster than autotrophs to the 802 relief of nutrient (N, P) limitation (i.e. BP already increased by 150-500% at t24 h, while 14 C-803 based production rates increased only after 48-72 h), NCP was first lower (and negative) in the 804 dust-amended tanks as compared to the controls. Marañón et al. (2010) and Pulido-Villena 805 (2008, 2014) have already reported on a much faster response of the heterotrophic prokaryote 806 community to dust enrichment in the central Atlantic Ocean and Mediterranean Sea, 807 respectively. As DIP concentrations at the completion of their 48 h incubations did not differ 808 from that in the controls, it is unlikely that primary production rates and consequently NCP 809 would have further increased. In contrast, during our experiments, DIP concentrations in dust-810 amended tanks (D) reached initial levels only after 72 h at TYR and ION and remained far above 811 ambient levels at FAST until the end of the experiment (t96h). During the PEACETIME cruise, 812 high frequency sampling of CTD casts allowed following the evolution of biogeochemical 813 properties and fluxes before and after wet dust deposition that took place in the area around 814 FAST on June 3-5 (Van Wambeke et al., 2020). As in our experiment, a rapid increase in BP was 815 responsible for the observed *in situ* decline in DIP concentrations in the mixed layer following 816 the rain with no detectable changes in primary production (Van Wambeke et al., 2020). The 817 intensity of the wet deposition event that was simulated during our experiments was, by far, 818 higher, but still representative of a realistic scenario (Bonnet and Guieu, 2006; Loÿe-Pilot and 819 Martin, 1996; Ternon et al., 2010).

The most intriguing result concerning the export of inorganic and organic matter is that these fluxes were maximal at the end of the experiment at TYR in the dust-amended tanks despite the fact that ¹⁴C-based production was relatively low and not enhanced by dust addition. Based on previous studies (Bressac et al., 2014; Louis et al., 2017; Ternon et al., 2010), organic

824	matter export was most likely mainly due to the formation of organic-mineral aggregates
825	triggered by the introduced lithogenic particles (referred thereafter to as POC _{litho}). Indeed, Louis
826	et al. (2017) showed that such an aggregation process occurs within 1 h after dust deposition.
827	These authors further demonstrated the key role of TEP as the conversion of dissolved organic
828	matter (DOM) to POC was mediated by TEP formation/aggregation activated by the introduction
829	of dust. As TEP concentrations were only measured on two occasions after seeding with the first
830	measurement occuring at t24h,), it prevents studying in detail the dynamics of these particles.
831	Nevertheless, it is very likely that the sharp decrease of TEP-C (Fig. 3) between t24h and t72h
832	was related to POC_{litho} export. The coefficient linking POC_{litho} to $Litho_{flux}$ (i.e. the mass of
833	sedimented particles) measured here (0.02) is consistent with values reported for other
834	experiments conducted in the Mediterranean Sea (Louis et al., 2017).
835	Even though ¹⁴ C-based production rates were enhanced in the dust-amended tanks at
836	stations ION and FAST, the amount of POC exported at the end of these experiments remained
837	lower than at TYR, with fluxes ~ 10-20 mg C m ⁻² d ⁻¹ .
838	The recovery of the introduced dust (traced by the lithogenic mass recovered in the traps)

Ŋ 839 was low (27% at TYR, ~20% at ION and 13-19% at FAST) reflecting that a majority of the dust 840 particles (the smaller ones that are the most abundant according to the particle size distribution of 841 the dust) still remained in the tanks after 3 or 4 days following dust addition. This has been 842 already observed in pelagic mesocosms (Bressac et al., 2012) as those small particles can 843 aggregate to organic matter and eventually sink. The higher export efficiency observed 844 (TYR>ION>FAST) is likely linked to the higher initial abundance and higher production of 845 TEPs during the experiment (Fig. 3). At TYR, impacted by a strong dust event several days before the experiment started (see above), the likely stimulation of the autotrophs after this in 846

847 *situ* event should have been followed by the production of a fresh and abundant DOM,

comparable to the "post-bloom situation" in Bressac and Guieu (2013).

849 **4.3. Impact of dust addition under future environmental**

850 conditions

851 Warming and/or acidification had a clear impact on most evaluated stocks and metabolic 852 rates. Gazeau et al. (2021) have already discussed temperature/pH mediated changes in nutrient 853 uptake rates and autotrophic community composition in these experiments. Briefly, they showed 854 that warming and acidification did not have any detectable impact on the release of nutrients 855 from atmospheric particles. Furthermore, these external drivers did not drastically modify the 856 composition of the autotrophic assemblage with all groups benefiting from warmer and acidified 857 conditions. Here, we showed that the difference in the response of plankton community 858 metabolism to dust addition under present and future conditions of temperature and pH was highly dependent on the sampling station (Fig. 9). At all stations, ¹⁴C-based particulate 859 860 production rates were enhanced under future conditions as compared to those measured under present environmental conditions (treatment D) although this pattern was not observed for ¹³C 861 862 incorporation into POC at stations ION and FAST. At ION, no differences could be detected and 863 at FAST an even lower ¹³C-enrichment was measured at the end of the experiment. These 864 contrasting patterns between ¹⁴C-uptake rates and ¹³C-enrichment of POC are likely explained by 865 the fact that the latter covered the whole experimental period (including dark periods) and represents net community carbon production while ¹⁴C-based rates were measured over 8 h 866 incubations in the light, providing an estimate in between gross and net carbon production. 867

868 Similarly, the heterotrophic compartment was more stimulated, as BP rates increased 869 strongly at all stations under future conditions compared to treatment D. The relatively smaller 870 increase in CR rates, compared to BP, leading to higher BGE suggests a better utilization of 871 resources by heterotrophic prokaryotes under future environmental conditions. Overall, CR was 872 more impacted than GPP, with the consequence that all integrated NCP rates decreased under 873 future environmental conditions compared to present conditions (treatment D). At station TYR, 874 as discussed previously, dust addition under present conditions did not lead to a switch from net 875 heterotrophy to net autotrophy. This pattern was even more obvious under warmer/acidified 876 conditions, with a larger decrease in integrated NCP at this station. The decrease of integrated 877 NCP at station FAST relative to controls, as well as the smaller increase of all ¹⁴C-based 878 production rates relative to those observed at station ION must be taken with caution. As already discussed, the fact that for these processes (O₂ metabolism and ¹⁴C-incorporation), no samples 879 were taken at FAST at t72h when maximal cell abundances were recorded for all autotrophic 880 881 groups (pico- and nano-eukaryotes, autotrophic bacteria) must have artificially led to an 882 underestimation of these integrated metabolic rates. The question of the timing appeared even 883 more preponderant under warmer/acidified conditions, especially at station FAST, where the 884 very important increase in BP led to a full consumption of DIP before t48h (Gazeau et al., 2021) 885 and drove the community towards a strong heterotrophy. The metabolic balance further switched 886 to a slight autotrophy at t72h when heterotrophic bacterial activity appeared limited by nutrient 887 availability.

888 Both elevated partial pressure of CO_2 (pCO_2) and warming are major global change 889 stressors impacting marine communities. Elevated pCO_2 may directly facilitate oceanic primary 890 production through enhanced photosynthesis (Hein and Sand-Jensen, 1997; Riebesell et al.,

891	2007) although the effects appear to be species- and even strain-specific (e.g. Langer et al.,
892	2009). Warming affects organisms by enhancing their metabolic rates (Brown et al., 2004;
893	Gillooly et al., 2001). Although recent studies suggest large differences in temperature sensitivity
894	between phytoplankton taxa (Chen and Laws, 2017) and no significant overall difference
895	between algae and protozoa (Wang et al., 2019), mineralization rates are usually believed to be
896	more impacted by warming than primary production rates, potentially leading to a decline in net
897	oceanic carbon fixation (Boscolo-Galazzo et al., 2018; Garcia-Corral et al., 2017; Lopez-Urrutia
898	and Moran, 2007; Regaudie-de-Gioux and Duarte, 2012) and carbon export efficiency (Cael et
899	al., 2017; Cael and Follows, 2016). Overall, our experimental set-up did not allow discriminating
900	warming from acidification effects, precluding an evaluation of their potential individual
901	impacts. Nevertheless, we could speculate to which extent a 3 °C warming and a doubling of
902	CO ₂ can explain some of the observed differences between D and G (for instance, a 2-fold
903	increase in ¹⁴ C-based production rates at ION). For photosynthesis, meta-analysis studies
904	indicate minor effects of pCO_2 on most investigated species (Kroeker et al., 2013; Mackey et al.,
905	2015). Recent studies show a strong, although species-dependent, temperature sensitivity of
906	phytoplankton growth (Chen and Laws, 2017; Wang et al., 2019), suggesting that a 3 °C
907	warming could explain most of the increased carbon fixation in G compared to D. With respect
908	to NCP, our results are in line with the general view and suggest a weakening of the so-called
909	fertilization effect of atmospheric deposition in the coming decades.
910	In contrast, we did not observe an additional impact of future environmental conditions

912 order of magnitude for treatments D and G. This result is in agreement with the findings of a

on the export of organic matter after dust addition as, at each station, this export was of the same

911

913 similar experiment in coastal Mediterranean waters that considered only pH change (Louis et al.,

914 2017) but stands in contrast with the findings of Müren et al. (2005) who showed a clear 915 decrease in sedimentation following a 5 °C warming in the Baltic Sea. Only a few studies have 916 addressed the combined effect of both temperature and pH changes on aggregation processes and 917 export but none considered dust as the particulate phase. These studies, focused mainly on the 918 formation of TEP, were inconclusive on the impact of these combined factors (Passow and 919 Carlson, 2012, and references therein). The potential effect of warming and acidification on 920 biogenic carbon export was certainly, over the rather restricted duration of the experiments, 921 insignificant as compared to the large amount of carbon exported through the lithogenic pump. 922 Although a longer experimental period would likely be necessary to clearly support an impact of 923 future conditions on export, those changes occur on a long time scale that cannot be easily 924 mimicked by experimental approaches. Only *in situ* co-located observations (atmospheric flux 925 and export in sediment traps) over long temporal scales would be necessary to ascertain the 926 interactive effects of these stressors at the decadal time scale.

927 **5. Conclusion**

928 Although the three experiments were conducted under rather similar conditions in terms 929 of nutrient availability and chlorophyll stock of the tested seawater, contrasting responses were 930 observed following the simulation of a wet dust deposition event. Under present conditions of 931 temperature and pH, at the site where the community was the most heterotrophic (TYR), no 932 positive impact of new nutrients could be observed on autotrophs, while a fast and strong 933 response of heterotrophic bacteria drove the metabolic balance towards an even more 934 heterotrophic state. The situation was different at the two other stations where a more active 935 autotrophic community responded quickly to the relief in nutrient (N, P) limitation, driving the

936 community to an autotrophic state at the end of these experiments. In all tested waters, an overall 937 faster response of the heterotrophic prokaryote community, as compared to the autotrophic 938 community, was observed after new nutrients were released from dust. Phytoplankton could 939 benefit from nutrient inputs, only if the amount released from dust was enough to sustain both 940 the fast bacterial demand and the delayed one of phytoplankton. As our experimental protocol 941 consisted in simulating a strong, although realistic, wet dust deposition, further work should 942 explore at which flux a wet dust deposition triggers an enhancement of net community 943 production and therefore increases the capacity of the surface oligotrophic ocean to sequester 944 atmospheric CO₂. This question, of the utmost importance in particular for modelling purposes, 945 should be answered through future similar experiments as the ones considered in our study but 946 following a gradient approach of dust fluxes. As a consequence of a stronger sensitivity of 947 heterotrophic prokaryotes to temperature and/or pH, the ongoing warming and acidification of 948 the surface ocean will result in a decrease of the dust fertilization of phytoplankton in the coming 949 decades and a weakening of the atmospheric CO₂ sequestration capacity of the surface 950 oligotrophic ocean. The contrasting results obtained at the three stations during our study will 951 need to be translated into process parameterization. The important dataset presented in this 952 manuscript, covering a variety of tested waters, environmental stressors and responses, will 953 allow such parameterization to be used in biogeochemical models coupled to ocean dynamics in 954 order to depict the spatial and temporal dynamics of stocks and fluxes following dust deposition 955 in surface oligotrophic waters.

956 **Data availability**

- 957 Underlying research data are being used by researcher participants of the PEACETIME
- 958 campaign to prepare other manuscripts, and therefore data are not publicly acces- sible at the
- 959 time of publication. Data will be accessible (http://www.obs-
- 960 vlfr.fr/proof/php/PEACETIME/peacetime.phpTS4, https://doi.org/10.17882/75747, Guieu et al.,
- 961 2020b) once the special issue is completed (all papers should be published by fall 2021).

962 Author contributions

FG and CG designed and supervised the study. All authors participated in sample analyses. FGwrote the paper with contributions from all authors.

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1345 **Tables**

1346 Table 1. Initial chemical and biological stocks as measured while filling the tanks (initial conditions in pumped surface water;

1347 sampling time: t-12h). NO_x: nitrate + nitrite, DIP: dissolved inorganic phosphorus, Si(OH)₄: silicate, POC: particulate organic carbon,

- 1348 DOC: dissolved organic carbon, TEP: transparent exopolymer particles, TChla: total chlorophyll a. Values shown for ¹⁴C
- 1349 incorporation rates, percentages of extracellular release (%PER) as well as for net community production (NCP), community
- 1350 respiration (CR) and gross primary production (GPP) were estimated from samples taken at t0 in the control tanks. For heterotrophic
- 1351 bacterial production (BP), rates were estimated from seawater sampled at t-12h.

	Sampling station	TYR	ION	FAST
	Coordinates (decimal)	39.34 N, 12.60 E	35.49 N, 19.78 E	37.95 N, 2.90 N
	Bottom depth (m)	3395	3054	2775
	Day and time of pumping (local time)	17/05/2017 17:00	25/05/2017 17:00	02/06/2017 21:00
	Temperature (°C)	20.6	21.2	21.5
	Salinity	37.96	39.02	37.07
Stocks	NO _x (nmol L ⁻¹)	14.0	18.0	59.0
	DIP (nmol L ⁻¹)	17.1	6.5	12.9
	Si(OH)4 (µmol L ⁻¹)	1.0	0.96	0.64

	POC (µmol L ⁻¹)	12.9	8.5	6.0
	DOC (µmol L ⁻¹)	72.2	70.2	69.6
	TEP (x $10^6 L^{-1}$)	6.8	3.8	3.7
	TChla (μ g L ⁻¹)	0.063	0.066	0.072
	Heterotrophic prokaryotes abundance (x 10 ⁵ cell mL ⁻¹)	4.79	2.14	6.15
Processes	$^{14}\text{C}\text{-based}$ total particulate production (µg C L $^{-1}$ h $^{-1})$	0.08 ± 0.03	0.14 ± 0.04	0.15 ± 0.04
	¹⁴ C-based > 2 μ m particulate production (μ g C L ⁻¹ h ⁻¹)	0.07 ± 0.02	0.11 ± 0.02	0.11 ± 0.02
	$^{14}\text{C}\text{-based} < 2 \ \mu\text{m}$ particulate production ($\mu\text{g} \ \text{C} \ \text{L}^{\text{-1}} \ \text{h}^{\text{-1}}$)	0.01 ± 0.01	0.04 ± 0.02	0.05 ± 0.01
	%PER	60 ± 20	45 ± 3	32 ± 23
	NCP (μ mol O ₂ L ⁻¹ d ⁻¹)	-1.9 ± 0.3	$\textbf{-0.2}\pm0.2$	$\textbf{-0.8}\pm0.9$
	CR (µmol O ₂ L ⁻¹ d ⁻¹)	-2.6 ± 0.1	-1.2 ± 0.5	-1.9 ± 1.6
	GPP (μ mol O ₂ L ⁻¹ d ⁻¹)	0.7 ± 0.4	1.1 ± 0.3	1.1 ± 0.7
	BP (ng C L ⁻¹ h ⁻¹)	11.6	15.2	34.6

1353	Table 2. Heterotrophic bacterial production (BP) growth rates (μ_{BP} in h^{-1}) estimated from the
1354	exponential phase of BP growth, observable from at least four sampling points, between t0 and
1355	t12h, during the three experiments (TYR, ION and FAST) in the six tanks (controls: C1, C2; dust
1356	addition under present conditions of temperature and pH: D1, D2; dust addition under future
1357	conditions of temperature and pH: G1 and G2). Values \pm SE are shown.

	μ_{BP}				
	TYR	ION	FAST		
C1	0.076 ± 0.025	0.042 ± 0.007	0.020 ± 0.003		
C2	0.066 ± 0.018	0.041 ± 0.005	0.026 ± 0.004		
D1	0.117 ± 0.008	0.095 ± 0.020	0.089 ± 0.014		
D2	0.194 ± 0.020	0.145 ± 0.007	0.090 ± 0.007		
G1	0.164 ± 0.020	0.126 ± 0.011	0.124 ± 0.005		
G2	0.150 ± 0.003	0.137 ± 0.033	0.163 ± 0.014		

1359	Table 3. Estimated bacterial growth efficiency (BGE in %) during the course of the three
1360	experiments (TYR, ION and FAST) in the six tanks (controls: C1, C2; dust addition under
1361	present conditions of temperature and pH: D1, D2; dust addition under future conditions of
1362	temperature and pH: G1 and G2). BGE was calculated based on integrated heterotrophic
1363	bacterial production (BP) and community respiration (CR) rates by applying a bacterial
1364	respiration to CR ratio of 0.7 and a respiratory quotient of 0.8 (see Material and Methods).

Bacterial growth efficiency (BGE)				
	TYR	ION	FAST	
C1	11.1	9.8	15.4	
C2	11.7	14.5	22.0	
D1	31.8	21.0	17.3	
D2	32.3	30.6	19.9	
G1	39.3	35.2	37.6	
G2	32.5	34.8	38.1	

1367 Figure caption

1368 Fig. 1. Location of the sampling stations in the Mediterranean Sea on board the R/V Pourquoi

- 1369 Pas? during the PEACETIME cruise. Background shows satellite-derived surface chlorophyll a
- 1370 concentration averaged over the entire duration of the cruise (courtesy of Louise Rousselet).

1371 Fig. 2. Dissolved organic carbon (DOC) concentrations and ratio between total hydrolysable

1372 amino acids (TAA) and DOC concentrations measured in the six tanks (controls: C1, C2; dust

1373 addition under present conditions of temperature and pH: D1, D2; dust addition under future

1374 conditions of temperature and pH: G1 and G2) during the three experiments (TYR, ION and

1375 FAST). The dashed vertical line indicates the time of seeding (after t0).

1376 Fig. 3. Particulate organic carbon (POC) concentrations and transparent exopolymer particle

1377 carbon content (TEP-C) measured in the six tanks (controls: C1, C2; dust addition under present

1378 conditions of temperature and pH: D1, D2; dust addition under future conditions of temperature

1379 and pH: G1 and G2) during the three experiments (TYR, ION and FAST). The dashed vertical

1380 line indicates the time of seeding (after t0).

1381 Fig. 4. ¹⁴C-based production rates (A: $< 2 \mu m$ and B: $> 2 \mu m$ size fractions, C: total particulate)

estimated from 8 h incubations on samples taken in the six tanks (controls: C1, C2; dust addition

1383 under present conditions of temperature and pH: D1, D2; dust addition under future conditions of

- 1384 temperature and pH: G1 and G2) during the three experiments (TYR, ION and FAST). The
- 1385 percentage of extracellular release (D, %PER) is also shown.

1386 Fig. 5. Incorporation of ¹³C into particulate organic carbon (δ^{13} C-POC) in the six tanks (controls:

1387 C1, C2; dust addition under present conditions of temperature and pH: D1, D2; dust addition

1388 under future conditions of temperature and pH: G1 and G2) during the three experiments (TYR,

1389 ION and FAST). The dashed vertical line indicates the time of seeding (after t0).

1390 Fig. 6. A: Net community production (NCP), B: community respiration (CR) and C: gross

1391 primary production (GPP) rates estimated using the oxygen light-dark method (24 h incubations)

1392 on samples taken in the six tanks (C1, C2, D1, D2, G1 and G2) during the three experiments

1393 (TYR, ION and FAST).

1394 Fig. 7. Heterotrophic bacterial production rates (BP) and cell-specific maximum hydrolysis

1395 velocity (Vm) of the alkaline phosphatase (both over 1-2 h incubations) on samples taken in the

1396 six tanks (C1, C2, D1, D2, G1 and G2) during the three experiments (TYR, ION and FAST).

1397 Fig. 8. Total mass and organic matter fluxes measured in the sediment traps at the end of the

three experiments (TYR, ION and FAST) in the six tanks (C1, C2, D1, D2, G1 and G2).

1399 Fig. 9. Relative difference (%) between integrated rates measured in tanks D (D1, D2; dust

1400 addition under present conditions of temperature and pH) and G (G1, G2; dust addition under

1401 future conditions of temperature and pH) as compared to the controls (C1, C2) during the three

1402 experiments (TYR, ION and FAST). Vertical boxes represent the range observed between the

1403 two replicates per treatment.



1405 Fig. 1





1407 Fig. 2








1411 Fig. 4



1413 Fig. 5







1415 Fig. 6









1419 Fig. 8





- Dust addition under current conditions of temperature and pH
- Dust addition under conditions of temperature and pH projected for 2100 ΔT = + 3°C and ΔpH = -0.3

1421 Fig. 9

1420