

1 **Impact of dust enrichment on Mediterranean plankton**

2 **communities under present and future conditions of pH and**

3 **temperature: an experimental overview**

4 Frédéric Gazeau¹, Céline Ridame², France Van Wambeke³, Samir Alliouane¹, Christian Stolpe¹,
5 Jean-Olivier Irisson¹, Sophie Marro¹, Jean-Michel Grisoni⁴, Guillaume De Liège⁴, Sandra Nunige³,
6 Kahina Djaoudi³, Elvira Pulido-Villena³, Julie Dinasquet^{5,6}, Ingrid Obernosterer⁶, Philippe Catala⁶,
7 Cécile Guieu¹

8 ¹ Sorbonne Université, CNRS, Laboratoire d'Océanographie de Villefranche, LOV, 06230

9 Villefranche-sur-Mer, France

10 ² CNRS-INSU/IRD/MNHN/UPMC, LOCEAN: Laboratoire d'Océanographie et du Climat:

11 Expérimentation et Approches Numériques, UMR 7159, 75252 Paris Cedex 05, France

12 ³ Aix-Marseille Université, Université de Toulon, CNRS/INSU, IRD, MIO, UM 110, 13288,

13 Marseille, France

14 ⁴ Sorbonne Université, CNRS, Institut de la Mer de Villefranche, IMEV, 06230

15 Villefranche-sur-Mer, France

16 ⁵ Scripps Institution of Oceanography, University of California San Diego, USA

17 ⁶ CNRS, Sorbonne Université, Laboratoire d'Océanographie Microbienne, LOMIC, F-66650

18 Banyuls-sur-Mer, France

19 Correspondence to: Frédéric Gazeau (f.gazeau@obs-vlfr.fr)

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22 Abstract

23 In Low Nutrient Low Chlorophyll areas, such as the Mediterranean Sea, atmospheric fluxes
24 represent a considerable external source of nutrients likely supporting primary production especially
25 during stratification periods. These areas are expected to expand in the future due to lower nutrient
26 supply from sub-surface waters caused by enhanced stratification, likely further increasing the role
27 of atmospheric deposition as a source of new nutrients to surface waters. Yet, whether plankton
28 communities will react differently to dust deposition in a warmer and acidified environment
29 remains an open question. The impact of dust deposition both in present and future climate
30 conditions was assessed through three perturbation experiments in the open Mediterranean Sea.
31 Climate reactors (300 L) were filled with surface water collected in the Tyrrhenian Sea, Ionian Sea
32 and in the Algerian basin during a cruise conducted in May/June 2017 in the frame of the
33 PEACETIME project. The experimental protocol comprised two unmodified control tanks, two
34 tanks enriched with a Saharan dust analog and two tanks enriched with the dust analog and
35 maintained under warmer (+3 °C) and acidified (-0.3 pH unit) conditions. Samples for the analysis
36 of an extensive number of biogeochemical parameters and processes were taken over the duration
37 of the experiments (3-4 d). Here, we present the general setup of the experiments and the impacts of
38 dust seeding and/or future climate change scenario on nutrients and biological stocks. Dust addition
39 led to a rapid and maximum input of nitrate whereas phosphate release from the dust analog was
40 much smaller. Our results showed that the impacts of Saharan dust deposition in three different
41 basins of the open Northwestern Mediterranean Sea are at least as strong as those observed
42 previously in coastal waters. However, interestingly, the effects of dust deposition on biological
43 stocks were highly different between the three investigated stations and could not be attributed to
44 differences in their degree of oligotrophy but rather to the initial metabolic state of the community.

45 Finally, ocean acidification and warming did not drastically modify the composition of the
46 autotrophic assemblage with all groups positively impacted by warming and acidification,
47 suggesting an exacerbation of effects from atmospheric dust deposition in the future.

48 1. Introduction

49 Atmospheric deposition is well recognized as a significant source of micro- and
50 macro-nutrients for surface waters of the global ocean (Duce et al., 1991; Jickells et al., 2005;
51 Moore et al., 2013). The potential modulation of the biological carbon pump efficiency and the
52 associated export of carbon by atmospheric deposition events are still poorly understood and
53 quantified (Law et al., 2013). This is especially true for Low Nutrient Low Chlorophyll (LNLC)
54 areas where atmospheric fluxes can play a considerable role in nutrient cycling and that represent
55 60% of the global ocean surface area (Longhurst et al., 1995) as well as 50% of global carbon
56 export (Emerson et al., 1997). These regions are characterized by a low availability of
57 macronutrients (N, P) and/or metal micronutrients (e.g. Fe) that can severely limit or co-limit
58 phytoplankton growth during large periods of year. The Mediterranean Sea is a perfect example of
59 these LNLC regions and exhibits chlorophyll *a* concentrations of less than $0.2 \mu\text{g L}^{-1}$ all year round
60 over most of its area, except in the Ligurian Sea where relatively large blooms can be observed in
61 late winter-early spring (e.g. Mayot et al., 2016). Recent assessments showed that the atmospheric
62 input of nutrients in the Mediterranean Sea is of the same order of magnitude as riverine inputs
63 (Powley et al., 2017), making the atmosphere a considerable external source of nutrients (Richon et
64 al., 2018). Atmospheric depositions are mostly in the form of pulsed inputs of aerosols from both
65 natural (Saharan dust) and anthropogenic origins (e.g. Bergametti et al., 1989; Desboeufs et al.,
66 2018). Dust deposition is mainly associated with wet deposition and occurs in the form of extreme
67 events (Loÿe-Pilot and Martin, 1996). Ternon et al. (2010) reported on an average annual dust flux
68 over four years of $11.4 \text{ g m}^{-2} \text{ yr}^{-1}$ (average during the period 2003–2007) at the DYFAMED station
69 in the Northwestern Mediterranean Sea. In this region, the most important events reported in the
70 2010 decade amounted to $\sim 22 \text{ g m}^{-2}$ (Bonnet and Guieu, 2006; Guieu et al., 2010b). Atmospheric

71 deposition provides new nutrients to the surface waters (Guieu et al., 2010b; Kouvarakis et al.,
72 2001; Markaki et al., 2003; Ridame and Guieu, 2002), Fe (Bonnet and Guieu, 2006) and other trace
73 metals (Desboeufs et al., 2018; Guieu et al., 2010b; Theodosi et al., 2010), that represent significant
74 inputs likely supporting the primary production especially during the stratification period (Ridame
75 and Guieu 2002, Bonnet et al. 2005), although no clear correlation between dust and ocean color
76 could be evidenced from long series of satellite observation (Guieu and Ridame, 2020).

77 Experimental approaches have shown that wet dust deposition events in the Northwestern
78 Mediterranean Sea (the dominant deposition mode in that basin) present a highest positive impact,
79 by supplying bioavailable new nutrients, compared to dry deposition on all tested parameters and
80 processes (Guieu et al., 2014a), except for N₂ fixation (Ridame et al., 2014). This so-called
81 fertilizing effect has been experimentally shown using microcosms or mesocosms where the wet
82 deposition of Saharan dust analog strongly stimulated primary production and phytoplankton
83 biomass (Guieu et al., 2014a; Ridame et al., 2014) but also modified the phytoplankton diversity
84 (Giovagnetti et al., 2013; Lekunberri et al., 2010; Romero et al., 2011). However, besides
85 phytoplankton, dust deposition modified also the bacterial community assemblage and led to even
86 stronger enhancements of production and/or respiration rates (Pulido-Villena et al., 2014). The
87 budgets established from four artificial seeding experiments during the DUNE project (Guieu et al.,
88 2014a) showed that by stimulating predominantly heterotrophic bacteria, atmospheric dust
89 deposition can enhance the heterotrophic behavior of these oligotrophic waters. This has the
90 potential to reduce the fraction of organic carbon that can be exported to deep waters during the
91 winter mixing period (Pulido-Villena et al., 2008) and ultimately limit net atmospheric CO₂
92 drawdown.

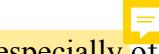
93 Another effect induced by Saharan dust deposition is the export of particulate organic
94 carbon (POC) as lithogenic particles can aggregate and ballast dissolved organic matter (Bressac et

95 al., 2014; Desboeufs et al., 2014; Louis et al., 2017a; Ternon et al., 2010). This so-called lithogenic
96 carbon pump can represent a major part of the carbon export following a dust deposition event (up
97 to 50% during the DUNE experiment; Bressac et al., 2014). Recently, Louis et al. (2017a) showed
98 that Saharan dust deposition triggers the abiotic formation of transparent exopolymeric particles
99 (TEP), leading to the formation of organic-mineral aggregates, a formation process that is highly
100 dependent on the quality and quantity of TEP-precursors initially present in seawater.

101 In response to ocean warming and increased stratification, open ocean nutrient cycles are
102 being and will be perturbed in the next decades with a high confidence of having regionally variable
103 impacts on primary producers (IPCC, 2019). Overall, LNLC areas are expected to expand in the
104 future (Irwin and Oliver, 2009; Polovina et al., 2008) due to lower nutrient supply from sub-surface
105 waters (Behrenfeld et al., 2006), likely further increasing the role of atmospheric deposition as a
106 significant source of new nutrients to surface waters. The ongoing warming and acidification of the
107 global ocean (IPCC, 2019), both also evidenced in the Mediterranean Sea (e.g. Kapsenberg et al.,
108 2017; The Mermex group, 2011) raise the question on whether plankton communities will react
109 differently to dust deposition in a warmer and acidified environment. Although dependent on
110 resource availability, it is well known that remineralisation by bacteria is subject to positive
111 temperature control (López-Urrutia and Morán, 2007). Under severe nutrient limitation, there is no
112 evidence that warming will lead to an enhancement of primary productivity (Marañón et al., 2018),
113 further pushing the balance towards net heterotrophy in oligotrophic areas.

114 With respect to ocean acidification, an *in situ* mesocosm experiment conducted during the
115 summer stratified period in the Northwestern Mediterranean Sea showed that the plankton
116 community was rather insensitive to this perturbation under strong nutrient limitation (Maugendre
117 et al., 2017, and references therein). This is coherent with results from Maugendre et al. (2015),
118 based on a batch experiment, showing that, under nutrient-depleted conditions in late winter, ocean

119 acidification has a very limited impact on the plankton community and that small species (e.g.
120 Cyanobacteria) might benefit from warming with a potential decrease of the export and energy
121 transfer to higher trophic levels. In contrast, in more eutrophic coastal conditions, Sala et al. (2016)
122 showed that ocean acidification exerted a positive effect on phytoplankton, especially on pico- and
123 nano-phytoplankton. Similarly, Neale et al. (2014) showed in a coastal ecosystem of the Alboran
124 Sea that ocean acidification could lead, although moderately, to high chlorophyll levels under low
125 light conditions with an opposite effect under high irradiance.

126 To date and to the best of our knowledge, there  has been no attempts to evaluate the
127 behavior of plankton communities after the deposition of atmospheric particles in the context of
128 future levels of temperature and pH. Yet, following the recommendation from Maugendre et al.
129 (2017), any perturbation experiment for future climate conditions in the Mediterranean Sea should
130 consider atmospheric deposition as a source of new nutrients and consider both temperature and pH
131 as external forcings. Such experiments were conducted in the frame of the PEACETIME project
132 (ProcEss studies at the Air-sEa Interface after dust deposition in the MEditerranean sea;
133 <http://peacetime-project.org/>) during the cruise on board the R/V “Pourquoi Pas?” in May/June
134 2017. The project aimed at extensively studying and parameterizing the chain of processes
135 occurring in the Mediterranean Sea after atmospheric deposition,  especially of Saharan dust, and to
136 put them in perspective of on-going environmental changes (Guieu et al., 2020). During that cruise,
137 three perturbation experiments were conducted in climate reactors (300 L tanks) filled with surface
138 water collected in the Tyrrhenian Sea (TYR), Ionian Sea (ION) and in the Algerian basin (FAST;
139 Fig. 1). Six tanks were used to follow simultaneously and with a high temporal resolution, the
140 evolution of biological activity and stocks, nutrients stocks, dissolved organic matter as well as
141 particles dynamics and export, following a dust deposition event simulated at their surface, both
142 under present environmental conditions and following a realistic climate change scenario for 2100



143 (ca. +3 °C and -0.3 pH units; IPCC, 2013). In this manuscript, we will present the general setup of
144 the experiments, the impacts of dust seeding and/or future climate change scenario on nutrients and
145 biological stocks. Among several other manuscripts related to these experiments that are introduced
146 here, a companion paper will be focusing on plankton metabolism (primary production,
147 heterotrophic prokaryote production) as well as on carbon budget.

148 **2. Material and Methods**

149 **2.1. General setup**

150 Six experimental tanks (300 L; Fig. 2) in which the irradiance spectrum and intensity can be
151 finely controlled and in which future ocean acidification and warming conditions can be fully
152 reproduced were installed in a temperature-controlled container. The tanks are made of high-density
153 polyethylene (HDPE) and were trace-metal free in order to avoid contaminations, with a height of
154 1.09 m, a diameter of 0.68 m, a surface area of 0.36 m² and a volume of 0.28 m³. All tanks were
155 cleaned before the experimental work following the protocol described by Bressac and Guieu
156 (2013). A weak turbulence was generated by a rotating PVC blade (9 rpm) in order to mimic natural
157 conditions. Each tank was equipped with a lid containing six rows of LEDs (Alpheus©). Each of
158 these rows were composed of blue, green, cyan and white units in order to mimic the natural sun
159 spectrum. At the conical base of each tank, a polyethylene (PE) bottle collecting the exported
160 material from above was screwed onto a polyvinyl chloride (PVC) valve that remained open during
161 the duration of the whole experiment. Photosynthetically active radiation (PAR; 400-700 nm) and
162 temperature were continuously monitored in each tank using respectively QSL-2100 Scalar PAR
163 Irradiance Sensors (Biospherical Instruments©) and pt1000 temperature sensors (Metrohm©)
164 connected to a D230 datalogger (Consort©).

165 The experimental protocol comprised two unmodified control tanks (C1 and C2), two tanks
166 enriched with Saharan dust (D1 and D2) and two tanks enriched with Saharan dust and maintained
167 under warmer (+3 °C) and acidified (-0.3 pH unit) conditions (G1 and G2). The atmosphere above
168 tanks C1, C2, D1 and D2 was flushed with ambient air (ca. 400 ppm, 6 L min⁻¹) and tanks G1 and
169 G2 were flushed with air enriched with CO₂ (ca. 1000 ppm, 6 L min⁻¹) in order to prevent CO₂

170 degassing from the acidified tanks. CO₂ partial pressure (*p*CO₂) in both ambient air and
171 CO₂-enriched air was monitored using two gas analysers (LI-820, LICOR©). The CO₂
172 concentration in the CO₂-enriched air was manually controlled through small injections of pure CO₂
173 (Air Liquide©) using a mass flow controller.

174 Three experiments were performed at the long duration stations TYR, ION and FAST. The
175 tanks were filled by means of a large peristaltic pump (Verder© VF40 with EPDM hose, flow of
176 1200 L h⁻¹) collecting seawater below the base of the boat (depth of ~ 5 m), used to supply
177 continuously surface seawater to a series of instruments during the entire campaign. In order to
178 homogeneously fill the tanks, the flow was divided into six HDPE pipes distributing the water
179 simultaneously into the different tanks. Overall, the filling of the six tanks took ~2 h (including
180 rinsing and initial sampling, see thereafter). At the three stations, tanks were always filled at the end
181 of the day before the start of the experiments: TYR (17/05/2017), ION (25/05/2017) and FAST
182 (02/06/2017). While filling the tanks, this surface seawater was sampled for the measurements of
183 selected parameters (sampling time = t-12h, see Table 1). After filling the tanks, seawater was
184 slowly warmed using 500 W heaters, controlled by temperature-regulation units (COREMA©), in
185 G1 and G2 overnight to reach an offset of +3 °C. ¹³C-bicarbonate was added to all tanks at 4:00 am
186 (local time; Gazeau et al., in preparation, this issue) and G1 and G2 were acidified by addition of
187 CO₂-saturated filtered (0.2 µm) seawater (~1.5 L in 300 L; collected when filling the tanks at each
188 station) at 4:30 am to reach a pH offset of -0.3. Sampling for many parameters took place prior to
189 dust seeding (sampling time = t0, see Table 1). Dust seeding was performed between 7:00 and 9:00
190 (local time) in tanks D1, D2, G1 and G2. The same dust analog was used and the same dust flux
191 was simulated as for the DUNE 2009 experiments described in Desboeufs et al. (2014). Briefly, the
192 fine fraction (< 20 µm) of Saharan soils collected in southern Tunisia, which is a major source of
193 dust deposition over the northwestern Mediterranean basin, was used in the seeding experiments.

194 The particle size distribution showed that 99% of particles had a size smaller than 0.1 μm , and that
195 particles were mostly made of quartz (40%), calcite (30%) and clay (25%; Desboeufs et al., 2014).
196 This collected dust underwent an artificial chemical aging process by addition of nitric and sulfuric
197 acid (HNO_3 and H_2SO_4 , respectively) to mimic cloud processes during atmospheric transport of
198 aerosol with anthropogenic acid gases (Guieu et al., 2010a, and references therein). To mimic a
199 realistic wet flux event of 10 g m^{-2} , 3.6 g of this analog dust were quickly diluted into 2 L of
200 ultrahigh-purity water (UHP water; 18.2 $\text{M}\Omega \text{ cm}^{-1}$ resistivity), and sprayed at the surface of the
201 tanks using an all-plastic garden sprayer (duration = 30 min). The N and P total contents in the dust
202 were $1.36 \pm 0.09\%$ of N and $0.055 \pm 0.003\%$ of P (see Desboeufs et al., 2014, for a full description
203 of dust chemical composition). The experimental protocol included the analysis of an extensive
204 number of biogeochemical parameters and processes, not all shown and discussed in this paper, that
205 are listed in Table 1. The experiment at stations TYR and ION lasted 72 h (3 days) whereas the last
206 experiment at station FAST was extended to four days. Seawater sampling was conducted 1 h (t1h),
207 6 h (t6h), 12 h (t12h), 24 h (t24h), 48 h (t48h) and 72 h (t72h) (+ 96 h = t96h for station FAST) after
208 dust addition. Acid-washed silicone tubes were used for transferring the water collected from the
209 tanks to the different vials or containers. For some parameters (e.g. nutrients, dissolved organic
210 carbon), sampled seawater was filtered online at the exit of the tanks on sterile membrane filter
211 capsules (gravity filtration with Sartobran \circledcirc 300; 0.2 μm).

212 **2.2. Analytical methods**

213 **2.2.1. Carbonate chemistry**

214 Seawater samples for pH measurements were stored in 300 mL glass bottles with a glass
215 stopper, pending analysis on board (within 2 h). Samples were transferred to 30 mL quartz cells and

216 absorbances at 434, 578 and 730 nm were measured at 25 °C on an Cary60 UV-Spectrophotometer
217 (Agilent©) before and after addition of 50 µL of purified meta-cresol purple provided by Robert H.
218 Byrne (University of South Florida, USA) following the method described by Dickson et al. (2007).
219 pH on the total scale (pH_T) was computed using the formula and constants of Liu et al. (2011). The
220 accuracy of pH measurements was estimated to 0.007 pH units, using a TRIS buffer solution
221 (salinity 35, provided by Andrew Dickson, Scripps university, USA).

222 Seawater samples for total alkalinity (A_T ; 500 mL) measurements were filtered on GF/F
223 membranes and analyzed onboard within one day. A_T was determined potentiometrically using a
224 Metrohm© titrator (Titrando 888) and a glass electrode (Metrohm©, ecotrode plus) calibrated using
225 first NBS buffers (pH 4.0 and pH 7.0, to check that the slope was Nernstian) and then using a TRIS
226 buffer solution (salinity 35, provided by Andrew Dickson, Scripps university, USA). Triplicate
227 titrations were performed on 50 mL sub-samples at 25 °C and A_T was calculated as described by
228 Dickson et al. (2007). Titrations of standard seawater provided by Andrew Dickson (Scripps
229 university, USA; batch 151) yielded A_T values within 5 µmol kg⁻¹ of the nominal value (standard
230 deviation = 1.5 µmol kg⁻¹, n = 40).

231 All parameters of the carbonate chemistry were determined from pH_T , A_T , temperature,
232 salinity, as well as phosphate and silicate concentrations using the R package seacarb¹. Propagation
233 of errors on computed parameters was performed using the new function “error” of this package,
234 considering errors associated with the estimation of A_T , pH_T as well as errors on dissociation
235 constants (Orr et al., 2018).

236 ¹ Seacarb: seawater carbonate chemistry with R. Gattuso, J.-P., J. M. Epitalon, H. Lavigne, J. C. Orr, B. Gentili, M.
237 Hagens, A. Hofmann, A. Proye, K. Soetaert and J. Rae, 2018. <https://cran.r-project.org/package=seacarb>

238 **2.2.2. Nutrients**

239 Seawater samples for dissolved nutrients were filtered online ($<0.2\text{ }\mu\text{m}$), collected in
240 polyethylene bottles and immediately analyzed on board. Nitrate + nitrite (NO_x) and silicate
241 (Si(OH)_4) measurements were conducted using a segmented flow analyzer (AAIII HR Seal
242 Analytical \circledR) according to Aminot and Kérouel (2007) with a limit of quantification of $0.05\text{ }\mu\text{mol}$
243 L^{-1} for NO_x and $0.08\text{ }\mu\text{mol L}^{-1}$ for Si(OH)_4 . In addition, for t-12h samples, the analysis of NO_x was
244 also performed by a spectrometric method in the visible at 540 nm, with a 1 m Liquid Waveguide
245 Capillary Cell (LWCC). The limit of detection was $\sim 10\text{ nmol L}^{-1}$ and the reproducibility was $\sim 6\%$.
246 Also from samples taken at t-12h, the measurement of ammonium concentrations was performed on
247 board using a Fluorimeter TD-700 (Turner Designs \circledR) according to Holmes et al. (1999). This
248 fluorimetric method is based on the reaction of ammonia with orthophtaldialdehyde and sulfite and
249 has a limit of quantification of $0.01\text{ }\mu\text{mol L}^{-1}$. Dissolved inorganic phosphorus (DIP) concentrations
250 were quantified using the Liquid Waveguide Capillary Cell (LWCC) method according to
251 Pulido-Villena et al. (2010). The LWCC was 2.5 m long and the limit of detection was 1 nmol L^{-1} .

252 **2.2.3. Pigments**

253 A volume of 2.5 L of sampled seawater was filtered onto GF/F filters, immediately frozen in
254 liquid nitrogen and stored at $-80\text{ }^\circ\text{C}$ pending analysis at the SAPIGH analytical platform at the
255 Institut de la Mer de Villefranche (IMEV, France). Filters were extracted at $-20\text{ }^\circ\text{C}$ in 3 mL
256 methanol (100%) containing an internal standard (vitamin E acetate, Sigma \circledR), disrupted by
257 sonication and clarified one hour later by vacuum filtration through GF/F filters. The extracts were
258 rapidly analyzed (within 24 h) on a complete Agilent \circledR Technologies 1200 series HPLC system.
259 The pigments were separated and quantified as described in Ras et al. (2008).

260 **2.2.4. Flow cytometry**

261 For the enumeration of autotrophic prokaryotic and eukaryotic cells, heterotrophic
262 prokaryotes and heterotrophic nanoflagellates (HNF) by flow cytometry, subsamples (4.5 mL) were
263 fixed with glutaraldehyde grade I 25% (1% final concentration), and incubated for 30 min at 4 °C,
264 then quick-frozen in liquid nitrogen and stored at -80 °C until analysis. Samples were thawed at
265 room temperature. Counts were performed on a FACSCanto II flow cytometer (Becton
266 Dickinson©) equipped with 3 air-cooled lasers: blue (argon 488 nm), red (633 nm) and violet (407
267 nm). The separation of different autotrophic populations was based on their scattering and
268 fluorescence signals according to Marie et al. (2010). *Synechococcus* spp. was discriminated by its
269 strong orange fluorescence (585 ± 21 nm), and pico- and nano-eukaryotes were discriminated by
270 their scatter signals of red fluorescence (> 670 nm). For the enumeration of heterotrophic
271 prokaryotes, cells were stained with SYBR Green I (Invitrogen – Molecular Probes) at 0.025% (vol
272 / vol) final concentration for 15 min at room temperature in the dark. Stained prokaryotic cells were
273 discriminated and enumerated according to their right-angle light scatter (SSC) and green
274 fluorescence at 530/30 nm. In a plot of green versus red fluorescence, heterotrophic prokaryotes
275 were distinguished from autotrophic prokaryotes. For the enumeration of HNF, staining was
276 performed with SYBR Green I (Invitrogen—Molecular Probes) at 0.05% (v/v) final concentration
277 for 15-30 min at room temperature in the dark (Christaki et al., 2011). Cells were discriminated and
278 enumerated according to their SSC and green fluorescence at 530/30 nm. Fluorescent beads (1.002
279 µm; Polysciences Europe©) were systematically added to each analyzed sample as internal
280 standard. The cell abundance was determined from the flow rate, which was calculated with
281 TruCount beads (BD biosciences©). Biomasses of each group were estimated based on conversion
282 equations and/or factors found in the literature (see section 2.3).

283 **2.2.5. Micro-phytoplankton and -heterotrophs**

284 At t-12h (i.e. seawater sampled during the filling of the tanks), a volume of 500 mL was
285 sampled in glass vials and immediately preserved in a 5% acidic Lugol's solution pending analysis.
286 At the Laboratoire d'Océanographie de Villefranche (LOV, France), 100 mL aliquots were
287 transferred to sedimentation chambers (Utermöhl) and counted under an inverted microscope at 200
288 to 400 magnifications.

289 **2.2.6. Mesozooplankton**

290 At the end of each experiment (t+72h for TYR and ION and t+96 h for FAST, after artificial
291 dust seeding), the sediment traps were removed, closed and stored with formaldehyde 4% (see
292 Gazeau et al., in preparation, this issue). The valve at the base of the tanks was then reopened to let
293 the remaining water inside the tanks (TYR 165-180 L; ION = 172.5 L and FAST = 150 L) pass
294 through a large PVC sieve (100 µm). The organisms retained on that mesh were gently removed
295 from the sieve using a washing bottle filled with filtered seawater (0.2 µm), and transferred directly
296 inside a 250 mL bottle. The bottle was filled with the sample (1/3 of the volume), and was
297 completed with formaldehyde 4%. The zooplankton digital images were obtained using a
298 ZooSCAN (Hydroptic©; Gorsky et al., 2010) at the PIQv-platform of EMBRC-France. The
299 identification of species was performed by automatic comparison with the library data set EcoTaxa
300 (<https://ecotaxa.obs-vlfr.fr/>, last access: 17/04/2020) and then all validated and corrected by a
301 human operator. 

302 2.3 Computations

303 The maximum percentage of dissolution from dust observed with respect to N and P was
304 calculated considering that these evapo-condensated dust contain $1.36 \pm 0.09\%$ of N and $0.055 \pm$
305 0.003% of P (Desboeufs et al., 2014). Based on maximal concentrations observed in the D and G
306 tanks after seeding (two discrete sampling within 6 h), one can estimate the maximal % of
307 dissolution of dust in seawater during the three experiments:

$$308 \quad \%_{dissolution} = \frac{CONC_{max} - CONC_{init}}{CONC_{dust}} \cdot 100 \quad (1)$$

309 where $CONC_{init}$ is the concentration of the corresponding nutrient in each tank before seeding (t0),
310 $CONC_{max}$ corresponds to the concentration of the corresponding nutrient in each tank when nutrient
311 concentration was at a maximum over the first 6 h after seeding as observed based on our discrete
312 sampling procedure, and $CONC_{dust}$ corresponds to the maximum input of each nutrient, if 100% of
313 its total concentration in the dust analog dissolves (as estimated based on dust chemical
314 composition; Desboeufs et al., 2014; see above).

315 As micro-phytoplankton counting was not performed throughout the experiment, as a first
316 approximation, autotrophic biomass was calculated as the sum of carbon contained in
317 *Synechococcus*, pico-eukaryotes and nano-eukaryotes (abundances based on flow cytometry) and is
318 therefore restricted to the fraction $< 20 \mu\text{m}$. For *Synechococcus*, conversion to carbon units were
319 done considering $250 \text{ fg C cell}^{-1}$ (Kana and Glibert, 1987), while the equation proposed by Verity et
320 al. (1992; $0.433 \text{ BV}^{0.863}$ where BV refers to the biovolume) was used for pico- and nano-eukaryotes
321 assuming a spherical shape and a diameter of 2 and 6 μm for the two groups, respectively.

322 Percentages of these different groups were calculated in order to estimate the composition of the
323 communities at the start and its evolution during the experiments. Furthermore, heterotrophic

324 biomass was computed as the sum of heterotrophic prokaryotes biomass and heterotrophic
325 nanoflagellates biomass. For heterotrophic prokaryotes, conversion to carbon units were done
326 considering 20 fg C cell⁻¹ (Lee and Fuhrman, 1987) and for heterotrophic nanoflagellates assuming
327 220 fg C μm⁻³ (Børshem and Bratbak, 1987), a spherical shape and a diameter of 3 μm. The ratio of
328 autotrophic and heterotrophic biomass during the experiments was used to evaluate the trophic
329 status of the investigated communities and its evolution. Finally, a proxy for micro-phytoplankton
330 biomass (B_{micro}) was estimated following Vidussi et al. (2001), as the sum of Fucoxanthin and
331 Peridinin.

332 **3. Results**

333 **3.1. Initial conditions**

334 Initial conditions of various measured parameters at the three sampling stations while filling
335 the tanks are shown in Table 2. pH_T and total alkalinity concentrations observed when pumping
336 seawater for the experiments (before ¹³C-bicarbonate addition and dust seeding: t=12h) followed a
337 west to east increasing gradient (8.03, 8.04 and 8.07; 2443, 2529 and 2627 µmol kg⁻¹ at FAST, TYR
338 and ION, respectively). NO_x concentrations were maximal at station FAST with a NO_x:DIP molar
339 ratio of ~ 4.6. Very low NO_x concentrations were observed at stations TYR and ION (14 and 18
340 nmol L⁻¹, respectively). DIP concentrations were the highest at station TYR (17 nmol L⁻¹) and the
341 lowest at the most eastern station (ION, 7 nmol L⁻¹). Consequently, the lowest NO_x:DIP ratio was
342 measured at TYR (0.8), compared to ION and FAST (2.8 and 4.6, respectively). Ammonium
343 concentrations were maximal at TYR (0.045 µmol L⁻¹), intermediate at ION (0.022 µmol L⁻¹), and
344 minimal at FAST (below detection limit). Silicate concentrations were similar at stations TYR and
345 ION (~ 1 µmol L⁻¹) and higher than at station FAST (0.64 µmol L⁻¹).

346 Very low and similar concentrations of chlorophyll *a* were measured at the three stations
347 (0.063 - 0.072 µg L⁻¹). The proportion of the different major pigments (Fig. 3) showed that
348 phytoplankton communities at stations TYR and ION were very similar with a dominance of
349 Prymnesiophytes (i.e. 19'-hexanoyloxyfucoxanthin; Ras et al., 2008) followed by Cyanobacteria
350 (i.e. Zeaxanthin; Ras et al., 2008). In contrast, at station FAST, the planktonic community was
351 clearly dominated by photosynthetic prokaryotes (i.e. Zeaxanthin and Divinyl-chlorophyll *a*;
352 Cyanobacteria and Prochlorophytes, respectively; Ras et al., 2008). At all three stations, the

353 proportion of pigments representative of larger species (i.e. Fucoxanthin and Peridinin; diatoms and
354 dinoflagellates respectively; Ras et al., 2008) were very small (< 5%).

355 Cellular abundances of all studied microorganisms (phytoplankton, micro-grazers,
356 heterotrophic bacteria) were the highest at FAST (Table 2). Picoeukaryotes, *Synechococcus* and
357 heterotrophic prokaryotes abundances followed an east to west increasing trend (ION < TYR <
358 FAST). In contrast, nano-eukaryotes abundance was similar at FAST and ION, and minimal at
359 TYR. The abundance of heterotrophic nanoflagellates (HNF) were similar at TYR and FAST
360 (~110-125 cells mL⁻¹), twice as high as the one observed at station ION. This east to west increasing
361 trend was also observed for micro-phytoplankton and micro-heterotrophs abundances (microscopic
362 analyses; Table 2). The ratio between autotrophic biomass and heterotrophic biomass was clearly in
363 favor of the heterotrophic compartment at stations TYR and FAST (~0.6 at the two stations) but the
364 opposite was found at station ION (ca. 1.3).

365 **3.2. Experimental conditions**

366 Irradiance levels during the experiments in controls were maximal at station ION and
367 minimal at station FAST (daily average maximum levels in controls: ~ 1050, ~ 1130 and ~ 1020
368 µmol photons m⁻² s⁻¹ at TYR, ION and FAST, respectively; Fig. 4). Decreases of water transparency
369 after dust addition was observed at all three stations with a maximum dust impact at station ION
370 and the lowest impact at station FAST where irradiance levels decreased by only 60 µmol photons
371 m⁻² s⁻¹ after dust addition (average between tanks D and G). At station TYR, a more pronounced
372 decrease was observed in acidified and warmed tanks (G1 and G2) with a decrease of daily average
373 maximum irradiance of ~ 60 and ~ 160 µmol photons m⁻² s⁻¹ as compared to dust-amended tanks D
374 and controls, respectively. Temperature control (Fig. 4) was not optimal showing deviations
375 between replicates of treatment G of up to 1.5 °C (station ION). Temperature in controls and D

376 tanks displayed a daily cycle with an increase during the day and a decrease at night. Overall, the
377 differences between the warmed treatment (G) and the other tanks were +3, +3.2 and +3.6 °C at
378 TYR, ION and FAST, respectively.

379 Addition of CO₂-saturated filtered seawater led to a decrease of pH_T from 8.05 ± 0.004
380 (average ± SD between C1, C2, D1 and D2 at t0) to 7.74 (average between G1 and G2) at station
381 TYR, from 8.07 ± 0.002 to 7.78 at station ION and 8.05 ± 0.001 to 7.72 at station FAST (Fig. 5).
382 pH_T levels remained more or less constant in ambient pH levels tanks during all three experiments
383 with no clear impact of dust addition in tanks D1 and D2. In lowered pH tanks, pH levels gradually
384 increased during the experiments with a systematic larger increase in one of the duplicates (G1). Yet
385 pH_T increases remained moderate thanks to the flushing of CO₂-enriched air above the tanks (*p*CO₂
386 of 1017 ± 11, 983 ± 96, 1023 ± 25 ppm at TYR, ION and FAST, respectively; data not shown).
387 Partial pressure of CO₂ in ambient air was similar at the three stations, i.e. 410 ppm (data not
388 shown). At all three stations, ¹³C-addition led to an increase of total alkalinity between 6 and 11
389 µmol kg⁻¹ and dust addition led to a decrease in tanks D and G between 8 and 16 µmol kg⁻¹ with no
390 apparent effects of warming and acidification. Overall, no large changes in this parameter were
391 observed during the experiments (Fig. 5).

392 **3.3. Changes in nutrient concentrations**

393 Dust addition in tanks D and G led to a rapid and maximum input of NO_x (as observed
394 during the first 6 h; Fig. 6; Table 3) of ~ 11 µmol L⁻¹ at all three stations with no differences
395 between both treatments. The corresponding dissolution percentage of N contained in the dust
396 analog was between 94 and 99%. In contrast, maximum DIP release (within 6 h after dust addition)
397 from the dust analog was much smaller and comprised between 20 and 37 nmol L⁻¹, with slightly
398 higher release at FAST (31-37 nmol L⁻¹) as compared to the other stations. Dissolution percentages

399 for DIP were estimated between 9.2 and 17.3% of total phosphorus contained in dust. As a
400 consequence of these contrasted dissolution of N and P, NO_x :DIP ratios increased from initial
401 values below 5 to above 300, within 6 h after dust seeding, in the dust amended (D and G) tanks
402 (Fig. 6).

403 After these rapid increases due to N and P releases in dust amended tanks, both variables
404 decreased with time. While nutrient variability was small in control tanks over the duration of the
405 experiments (NO_x and DIP variations below 20 and 3 nmol L⁻¹, respectively), large decrease of both
406 elements was measured in dust amended tanks (D and G; Table 4). For NO_x , similar linear
407 decreases were observed throughout the experiments at stations TYR and ION with no visible
408 differences between tanks D and G. In contrast, at station FAST, a more pronounced decrease in
409 NO_x was observed in dust-amended (D and G) tanks as compared to the other stations, with
410 detectable larger decreases in warmed and acidified tanks relative to the D treatment. Nevertheless,
411 at all stations, NO_x concentrations in D and G treatments remained far above ambient levels
412 throughout the experiments (> 9 $\mu\text{mol L}^{-1}$). Abrupt decreases in DIP were observed during the three
413 experiments after the initial increase. At station TYR, after 24 h, all DIP released from dust
414 decreased to initial levels in tanks G while it took two more days to reach initial levels in tanks D.
415 In contrast, at station ION, no clear difference in DIP dynamics was observed between treatments D
416 and G, with concentrations that decreased rapidly during the first 24 h but that remained above
417 initial levels until the end of the experiment. At station FAST, similarly to station TYR, DIP
418 decreased rapidly from t12h in treatment G, reaching levels close to initial conditions at the end of
419 the experiment. DIP decrease was much lower in treatment D (Table 4) with concentrations
420 maintained far above ambient levels throughout the experiment. As a consequence of these
421 differences between NO_x and DIP dynamics as well as differences among stations, NO_x :DIP ratio

422 increased during the experiments with clear differences between stations (Fig. 6) and remained
423 much higher than that in the controls over the duration of the three experiments.

424 Silicate dynamics showed at all stations higher concentrations in dust amended (D and G)
425 tanks relative to the controls. At TYR, while in control tanks, concentrations remained stable, they
426 increased linearly with time in the other tanks (D and G) with no apparent effect of the imposed
427 increase in temperature and decrease in pH (i.e. tanks G). Difference of Si(OH)₄ concentration
428 between dust amended treatments (D and G) and controls was ~0.1 µmol L⁻¹ at the end of the
429 experiment. At station ION, after an initial decrease of concentrations between t-12h and t0,
430 concentrations increased in all tanks until the end of the experiment with higher concentration in
431 dust amended tanks (D and G) than in controls (no difference between D and G treatments). In
432 contrast, at FAST, concentrations increased between t-12h and t0, and continued to increase in all
433 tanks (with higher values in dust amended tanks) until t48h and then decreased until the end of the
434 experiment. At the end of the experiment (t96h), Si(OH)₄ concentration was higher in the G
435 treatment than in the D treatment which was similar to the controls.

436 3.4. Changes in biological stocks

437 Regarding biological stocks, temporal dynamics showed very different patterns with respect
438 to the sampling station. At TYR, total chlorophyll *a* concentrations did not change in dust amended
439 tanks maintained under ambient levels of temperature and pH (Fig. 7) and even led to slightly
440 decreased values 24 h after dust addition (e.g. -35 to -38% in D1 and D2, respectively as compared
441 to controls; Table 5). No clear effect of dust addition (tanks D vs. C) were detectable for all groups
442 based on pigment analyses (Fig. 7). Results obtained based on flow cytometry counting (Fig. 8)
443 were coherent with these observations and showed stronger decreases in cell abundances for < 20
444 µm autotrophic groups in tanks D1 and D2 (-77 to -80%). In contrast, at this station, the abundance

445 of heterotrophic prokaryotes (HP) increased rapidly after dust addition both under ambient
446 (+53-68%) and future (+68%) environmental conditions, with no clear difference among those
447 treatments. In warmed and acidified tanks, strong discrepancies between the duplicates were
448 observed for pigments and autotrophic cell abundances. Indeed, tank G1 showed moderately higher
449 levels for all variables as compared to tanks C at the exception of pico-eukaryotes, while in G2 all
450 variables responded strongly to dust addition with maximum relative changes of > 300% (at the
451 exception of nano-eukaryotes: +119%). While HNF abundances responded positively to the
452 treatments in D1, D2 and G2 (+100-352%), abundances increased sharply in tank G1 towards the
453 end of the experiment (+1095%). At ION, a clear distinction between treatments could be observed
454 for almost all pigments and cell abundances (Fig. 7, Fig. 8). At the exception of nano-eukaryotes
455 and HNF, all variables (pigments and cell abundances) increased as a response to both dust addition
456 and warmed/acidified conditions (i.e. C < D < G). As an example (Table 5), the maximum relative
457 changes as compared to controls observed for total chlorophyll *a* were 109-183% and 399-426% in
458 tanks D and G, respectively. The highest stimulation to dust addition was observed for
459 *Synechococcus* with a +317-390% increase and +805-1425% increase in D and G tanks respectively
460 (Table 5). Abundances of nano-eukaryotes and HNF suggested no impact of dust addition under
461 ambient conditions but a positive impact in treatment G. In contrast to what was observed at TYR
462 for HP abundances, an effect of temperature and pH was observed at station ION with a higher
463 impact of dust addition under future environmental conditions. At station FAST, all above
464 mentioned variables related to biological stocks increased strongly after dust addition (Fig. 7, Fig. 8
465 and Table 5). For instance, total chlorophyll *a* increased following an exponential trend until the end
466 of the experiment reaching maximal values at t96h with slightly lower values observed under
467 ambient environmental conditions (+237-318% in D tanks vs. ~ +400% in G tanks).
468 Prymnesiophytes (i.e. 19'-hexanoyloxyfucoxanthin) and diatoms (i.e. Fucoxanthin) appeared as the

469 groups benefiting the most from dust addition with no large impacts of warming/acidification. In
470 contrast, Pelagophytes (i.e. 19'-butanoyloxyfucoxanthin) and green algae (i.e. Total Chlorophyll b)
471 responded much more in treatment G than in treatment D. Finally, although Cyanobacteria (i.e.
472 Zeaxanthin) responded faster to dust addition under future environmental conditions (tanks G), this
473 effect tended to attenuate towards the end of the experiment. In contrast to estimates based on
474 HPLC data, increases in cell abundances did not generally take place until the end of the
475 experiment. While abundances in pico-eukaryotes increased until t96h in treatment D, abundances
476 sharply declined between t72h and t96h for this group in treatment G. The same trend was observed
477 for *Synechococcus* during this experiment, although discrepancies between duplicates in treatment
478 D at sampling time t96h did not allow drawing conclusions on the behavior of this group at the end
479 of the experiment. Both under ambient and future conditions, abundances of nano-eukaryotes
480 declined sharply between t72h and t96h. The decline in HP abundances appeared even earlier
481 during the experiment with moderate maximum relative differences as compared to controls
482 observed at t48h. HP abundances declined very sharply between t48h and t96h in treatment G,
483 reaching control levels, while this decline was less sharp under ambient environmental levels.
484 Finally, HNF dynamics during this experiment was hard to evaluate with no clear effects of dust
485 addition or pH/temperature conditions and with a large increase in abundances in only one duplicate
486 of treatment G (t24h) followed by a gradual decrease.

487 Abundances of meso-zooplankton at the end of the experiments showed relatively similar
488 values at stations TYR and ION while much higher levels were observed at station FAST (Fig. 9).
489 As a consequence of large variability between duplicates at stations TYR and ION, no clear effects
490 of treatments were detected. At station FAST, although the sample size was too low to statistically
491 test for differences, higher total abundances of meso-zooplankton species were observed in the
492 dust-amended tanks with no differences between ambient and future conditions of temperature and

493 pH. However, differences in abundance were visible between these two treatments for specific
494 groups, with respectively higher abundance of Harosa and lower abundance of Crustacea (other
495 than copepods) and Mollusca in warmed and acidified tanks.

496 4. Discussion

497 4.1. Initial conditions

498 Overall, the three experiments were conducted with surface seawater collected during
499 typical stratified oligotrophic conditions typical of the open Mediterranean Sea at this period of the
500 year. However, at all three stations, initial concentrations of NO_x (14, 18 and 59 nmol L⁻¹) at TYR,
501 ION and FAST, respectively; Table 2) were lower than the ones reported by Manca et al. (2004) in
502 surface waters (5 m) in these areas in spring (0.036 ± 0.10 , 0.275 ± 0.358 and 0.183 ± 0.282 μmol
503 L⁻¹ for the areas corresponding to TYR, ION and FAST, respectively; <http://doga.ogs.trieste.it/medar/climatologies/>, last access: 28/04/2020). Similarly, surface DIP
504 concentrations as measured at the three stations were lower than values extracted from the
505 compilation of Manca et al. (2004) for the same period (0.072 ± 0.072 , 0.054 ± 0.035 and $0.115 \pm$
506 0.078 $\mu\text{mol L}^{-1}$ in the areas corresponding to TYR, ION and FAST, respectively). However, direct
507 measurements of NO_x and DIP concentrations using nanomolar techniques (as performed in our
508 study) are scarce in the Mediterranean Sea, limiting our ability to compare our results with these
509 published values which, in any case, show large interannual variability. Djaoudi et al. (2018)
510 reported low DIP values in the three studied basins. Furthermore, low observed concentrations of
511 NO_x and DIP at all three stations during our study were also in agreement with reported
512 concentrations in the coastal waters of Corsica during experiments using *in situ* mesocosms in June,
513 whether during the DUNE project (DIP ~ 5 nmol L⁻¹; Pulido-Villena et al., 2014; $\text{NO}_x < 30$ nmol L⁻¹;
514 Ridame et al., 2014) or during the MedSeA project ($\text{NO}_x \sim 50$ nmol L⁻¹ and DIP ~ 35 nmol L⁻¹;
515 Louis et al., 2017b). Furthermore, at all three stations, $\text{NO}_x:\text{DIP}$ molar ratios were well below the
516 Redfield ratio (16:1) and are consistent with ratios found in these previously cited studies. Both low
517 $\text{NO}_x:\text{DIP}$ ratio and low nutrient concentrations suggest that communities found at the three stations
518

519 experienced N and P co-limitation at the start of the experiments, as previously shown by Tanaka et
520 al. (2011). Some enrichment experiments in DIP, $\text{NO}_3 + \text{NH}_4$, glucose, alone or in combinations were
521 conducted using seawater sampled while filling the tanks. Bacterial production was mainly
522 stimulated by N+P addition at the three sites, although a slight stimulation was also detected after P
523 addition alone at TYR and ION (France Van Wambeke, pers. comm.). Initial concentrations of
524 dissolved Fe in the sampled seawater ranged from 1.5 nmol L⁻¹ (TYR) to 2.5 nmol L⁻¹ (ION;
525 Roy-Barman et al., in preparation, this issue). Such concentrations were unlikely limiting for
526 biological activity as previously shown in the Mediterranean Sea (Bonnet et al., 2005; Ridame et
527 al., 2014).

528 Total chlorophyll *a* concentrations of ~ 0.06 - 0.07 $\mu\text{g L}^{-1}$ (Table 2) were typical of
529 chlorophyll *a* levels found in these areas of the surface Mediterranean Sea at this period of the year,
530 as seen by satellite (Bosc et al., 2004), or from a database of *in situ* measurements (Manca et al.,
531 2004). During the DUNE and MedSeA projects cited above, chlorophyll *a* concentrations around
532 0.07 $\mu\text{g L}^{-1}$ were also encountered at the start of these experiments conducted in coastal waters
533 (Gazeau et al., 2017; Ridame et al., 2014). Although total chlorophyll *a* concentrations were rather
534 similar between the three tested stations, the composition of the phytoplankton communities, based
535 on HPLC pigment analyses, differed substantially. Indeed, while the communities were dominated
536 by nano-eukaryotic species at stations TYR and ION, both HPLC and flow cytometry data suggest
537 a larger contribution of pico-eukaryotes and Cyanobacteria at station FAST. Micro-autotrophs (e.g.
538 large diatoms and dinoflagellates) were slightly higher at station FAST. Due to their low
539 competitiveness during periods of nutrient limitation, the small contribution of large phytoplankton
540 cells at the start of the experiment is a fingerprint of LNLC areas and surface Mediterranean waters
541 at this period of the year (Siokou-Frangou et al., 2010). Autotrophic biomasses, as estimated based
542 on flow cytometry data (see Material and Methods) were similar at station TYR and ION (5.6 and

543 6.0 µg C L⁻¹) and maximal at FAST (7.7 µg C L⁻¹; Table 2). Although these estimates do not take
544 into account the contribution of micro-autotrophs, they appear to be in fair agreement with
545 estimates based on total chlorophyll *a* data, assuming a carbon to chlorophyll ratio of 70 (Bellacicco
546 et al., 2016), i.e. 4.4, 4.6 and 5.1 µg C L⁻¹ at TYR, ION and FAST, respectively. Furthermore, as
547 already mentioned, based on pigment analyses (HPLC), the sum of Fucoxanthin and Peridinin
548 (representative of diatoms and dinoflagellates, respectively) represented only ~10% of the total
549 chlorophyll *a* biomass at all stations. As biomass of both heterotrophic nanoflagellates and
550 prokaryotes followed a west to east gradient (FAST > TYR > ION), ratio of autotrophic vs
551 heterotrophic biomass appeared clearly in favor of the heterotrophic compartment at stations TYR
552 and FAST (ratio of 0.6) while a value above the metabolic balance was estimated at ION (ratio of
553 1.3). This is coherent with the highest net community production (NCP) rates being reported at this
554 station by Gazeau et al. (in preparation, this issue) showing that the initial community at the start of
555 this experiment was very close to metabolic balance (mean ± SE: -0.06 ± 0.09 µmol O₂ L⁻¹ d⁻¹). The
556 highest community respiration rates and consequently lowest NCP rates were measured at station
557 TYR (-1.9 µmol O₂ L⁻¹ d⁻¹) further suggesting that the autotrophic plankton community was not
558 very active (Ridame et al., in preparation, this issue) also confirmed by the lowest rate of CO₂
559 fixation; Ridame et al., in preparation, this issue), and relying on regenerated nutrients, as shown by
560 the highest level of NH₄⁺ measured at the start of this experiment. In contrast, the community at
561 station FAST although slightly heterotrophic (Gazeau et al., in preparation, this issue) and limited
562 by the low amount of nutrients (Table 2) was the most active as shown by the highest levels of ¹⁴C
563 production and heterotrophic prokaryote production (Gazeau et al., in preparation, this issue) as
564 well as N₂ fixation (Ridame et al., in preparation, this issue). Altogether, the heterotrophic signature
565 of the three investigated stations, although closer to metabolic balance for ION, reflected typical

566 natural biogeochemical conditions in the Mediterranean Sea during late spring to early summer
567 (Regaudie-de-Gioux et al., 2009).

568 **4.2. Experimental assessment**

569 The experimental tanks used in this study have already been validated in several studies
570 designed to investigate the inputs of macro- and micro-nutrients (e.g. NO_x, DIP, DFe) and the
571 export of organic matter, under close-to-abiotic conditions (seawater filtration onto 0.2 µm)
572 following simulated wet dust events using the same analog as used in our study (Bressac and Guieu,
573 2013; Louis et al., 2017a, 2018). Louis et al. (2017a, 2018) further investigated these impacts under
574 lowered pH conditions. During these experiments, no control of atmospheric *p*CO₂ was performed
575 and pH levels in the acidified filtered seawater rapidly increased due to CO₂ degassing (from ~7.4
576 to ~7.7 in six days). Prior to the cruise, we improved our experimental system to allow mimicking
577 future conditions by controlling atmospheric *p*CO₂ in addition to light and temperature (i.e. climate
578 reactors). During our experiments, thanks to the control of atmospheric *p*CO₂ (~ 1000 ppm), we
579 significantly reduced CO₂ degassing and maintained pH levels close to experimental targets.
580 However, as can be seen in Fig. 5, the regulation was consistently more efficient in tank G2 as
581 compared to G1. We attribute this small discrepancy (highest difference of 0.04 pH units between
582 the two G tanks at FAST) to a potential leak or a longer flushing time above tank G1. Nevertheless,
583 we do not anticipate this as an issue.

584 The lids above tanks, equipped with LEDs in order to reproduce sunlight intensity and
585 spectrum, were used for the first time during these experiments. The maximal intensity reached
586 under control conditions (C1, C2) was between 900 and 1000 µmol photons m⁻² d⁻¹. Although
587 slightly lower than estimates for the Northwestern Mediterranean Sea at 5 m depth in June (~1100
588 µmol photons m⁻² d⁻¹; Bernard Gentili, personal communication, 2017), simulated intensities were

589 fairly consistent between duplicates under control conditions (C1, C2) and under dust-amended
590 conditions (D1, D2). In contrast, larger differences were observed between warmed and acidified
591 tanks (G1 and G2; maximal differences of 100-200 $\mu\text{mol photons m}^{-2} \text{ d}^{-1}$ depending on the
592 experiment) that generally increased during each experiment. The reasons of these discrepancies are
593 not clear and could result from differences of light intensity generated by the lids, of PAR sensors
594 sensitivity and/or of the amount of particles remaining in the tanks. Unfortunately, although
595 replication appeared satisfactory for this treatment (except at station TYR; see below), we can not
596 fully exclude a potential impact of these technical issues on our results for this warmer and acidified
597 treatment. A similar conclusion can be drawn regarding temperature regulation in the container
598 where temperature was not spatially homogeneous, leading to significant differences among
599 replicates. After this study, experimental tanks were installed in a new container in order to solve
600 these problems.

601 The experimental strategy chosen during this study implied considering three different
602 treatments: control, simulation of dust deposition and simulation of dust deposition under future
603 projected environmental conditions. This unbalanced design, i.e. without the consideration of a
604 treatment without dust addition under future temperature and pH levels, was chosen for practical
605 reasons as only six tanks could be used for this study. Furthermore, as already mentioned, previous
606 experiments clearly showed very limited effects of these drivers when communities are strongly
607 limited by nutrient availability (Maugendre et al., 2017), therefore the objective of our study was to
608 test the impact of an external forcing (atmospheric deposition) under future conditions, without
609 discriminating warming from acidification effects. More importantly, the relatively low number of
610 experimental units that could be installed in an ambarcable clean container, implied considering
611 duplicated tanks for each treatment. This forced choice implied the impossibility to perform
612 statistical analyses on our results, as at least triplicates are necessary for most statistical tests.

613 Differences between duplicates were, for the vast majority of studied variables and processes, lower
614 than differences between treatments and appear acceptable considering the difficulty to incubate
615 plankton communities for which slight differences in their initial composition can translate into
616 very important differences in dynamics (Eggers et al., 2014). Unfortunately, while no large
617 replication issues were observed during the ION and FAST experiments, very important
618 discrepancies were detected for tanks of the warmed and acidified treatment at station TYR. The
619 reasons behind the different behavior of the autotrophic community in tanks G1 and G2 are not
620 fully understood but we strongly suspect that heterotrophic nano-flagellates, feeding mainly on
621 prokaryotic picoplankton (Sherr and Sherr, 1994), exerted a strong top-down control on this group
622 in tank G1 in which HNF abundance sharply increased during the experiment (+1100% in G1 vs. +
623 300% in G2). Interestingly, while autotrophic prokaryotes were clearly impacted, no differences
624 between the two tanks G1 and G2 were observed for heterotrophic prokaryotes although
625 nanoflagellates are known to feed to this group as well (Sherr and Sherr, 1994). Heterotrophic
626 nano-flagellates were likely not the only group of grazers which abundance increased during this
627 experiment in tank G1 as the biomass of diatoms (i.e. Fucoxanthin) did not increase in this tank.
628 Nevertheless, as no analyses of micro-grazer abundances were performed during the experiments,
629 this hypothesis can not be verified. All in all, these discrepancies for this treatment at station TYR
630 remain an issue and prevent us from drawing any conclusion on the combined effect of temperature
631 and pH on the dynamics of the community for that station.

632 **4.3. Impact of dust addition**

633 During the three experiments, the observed increases in NO_x and DIP few hours after dust
634 addition were rather similar to the enrichment levels obtained during the DUNE experiments at the
635 surface of the mesocosms ($\sim 50 \text{ m}^3$) after the simulation of a wet dust deposition using the same

636 dust analog and the same simulated flux (Pulido-Villena et al., 2014; Ridame et al., 2014). NO_x
637 levels moderately decreased over the course of our experiments due to biological uptake (50–1420
638 nmol L⁻¹, depending on the experiment). The opposite feature was observed for the DIP released by
639 dust that rapidly decreased during our experiments except at station FAST in the D treatment where
640 final concentrations did not reach initial levels. These enrichment levels, especially for NO_x, were
641 much higher than those observed by Pitta et al. (2017, and references therein) during land-based
642 mesocosm experiments in the Eastern Mediterranean Sea, in which a dry Saharan deposition was
643 simulated. In contrast to this experiment, the objective of our study was to assess the impact of wet
644 dust deposition, the main dust deposition pathway in the Western Mediterranean Sea (Loë-Pilot
645 and Martin, 1996). Furthermore, following observations of mixing between dust and polluted air
646 masses during their transport (e.g. Falkovich et al., 2001; Putaud et al., 2004), we chose to use an
647 evapo-condensed dust analog that mimics the processes taking place in the atmosphere prior to
648 deposition, essentially the adsorption of inorganic and organic soluble species (e.g. sulfate and
649 nitrate; see Guieu et al., 2010a, for further details). The imposed evapo-condensation processes are
650 responsible for the large nitrate releasing capacity of the dust particles used in our study. Regarding
651 the intensity of simulated wet deposition event, the 10 g m⁻² deposition event considered here
652 represents a high but realistic scenario, as several studies reported even higher short deposition
653 events in this area of the Mediterranean Sea (Bonnet and Guieu, 2006; Loë-Pilot and Martin, 1996;
654 Ternon et al., 2010).

655 Although NO_x and DIP increases after dust addition were rather similar during our three
656 experiments, interestingly the impacts on plankton community composition and functioning were
657 drastically different. Most experiments reporting on the effect of dust addition in the Mediterranean
658 Sea showed significant increases in chlorophyll *a* concentrations (mean 89% increase; Guieu and
659 Ridame, 2020). Such fertilization of primary producers was indeed observed at stations ION and

660 FAST under present conditions (maximum change in total chlorophyll *a* relative to the controls was
661 ~280% at FAST and ~150% at ION). The largest increase in chlorophyll *a* concentrations at station
662 FAST is coherent with the largest observed NO_x decrease following dust addition at this station.
663 Interestingly, following dust addition at this station, autotrophic production did not lead to DIP
664 exhaustion throughout the experiment as DIP concentrations were still above ambient conditions at
665 the end of the experiment. Maximal primary production rates (¹⁴C-incorporation) at this station at
666 the end of the experiment suggest a strong DIP recycling and the dominance of regenerated
667 production towards the end of the experiment (Gazeau et al., in preparation, this issue). Guieu et al.
668 (2014b) showed, based on the analysis of eight aerosols addition studies, that *Synechococcus* had in
669 most of the cases a weak responses to aerosol addition in contrast to nano- and
670 micro-phytoplankton, suggesting that aerosol deposition may lead to an increase in larger size class
671 phytoplankton. Yet, *Synechococcus* were well stimulated in some dust addition experiments (Herut
672 et al., 2005; Lagaria et al., 2017; Paytan et al., 2009), similar to what was observed at both stations
673 ION and FAST, where *Synechococcus* abundance was clearly enhanced by dust deposition. The
674 increase in *Synechococcus* abundance to dust-amended tanks was the highest relative to those of
675 pico- and nano-eukaryotes at these stations. This was especially true at station ION where no clear
676 response to nutrient enrichment was observed for nano-eukaryotes throughout the experiment.

677 However, it must be stressed that our experiments were performed over a relatively short period (3
678 to 4 days), and the sharp increase in Fucoxanthin paralleled by a decrease in silicates, at the end of
679 the experiment at station FAST where DIP limitation was not yet apparent, suggests a delayed
680 response of diatoms as compared to smaller groups (i.e. autotrophic prokaryotes, pico- and
681 nano-eukaryotes). Although this was not observed based on pigment analyses, the sharp decline in
682 nano-eukaryote abundances at the end of the FAST experiment following dust addition, further

683 suggests that this group, reacting quickly to nutrient enrichment was progressively grazed and/or
684 outcompeted by larger phytoplankton species.

685 In contrast to what was observed at stations ION and FAST, no stimulation of autotrophic
686 biomass and primary production rates (Gazeau et al., in preparation, this issue) was observed in the
687 dust treatments under present conditions at station TYR. To the best of our knowledge, this is the
688 first experimental evidence of a complete absence of response from an autotrophic community
689 following dust wet deposition. There is clear evidence that not only phytoplankton but also
690 heterotrophic bacteria are limited by inorganic nutrients, mainly DIP, in oligotrophic systems
691 (Obernosterer et al., 2003; Wambeke et al., 2002), thus suggesting that the supply of these resources
692 could explain variability in bacterial activity. Many recent studies have shown significant increase
693 in heterotrophic bacterial abundance, respiration and/or production following dust deposition in
694 oligotrophic ecosystems (Lekunberri et al., 2010; Pitta et al., 2017; Pulido-Villena et al., 2008,
695 2014; Romero et al., 2011). Most of the time, heterotrophic processes appear to be more stimulated
696 by dust pulses compared to autotrophic processes with increasing degree of oligotrophy, the
697 dominant response being modulated by the competition for nutrients between phytoplankton and
698 bacteria (Marañón et al., 2010). This is clearly what was observed at this station, with heterotrophic
699 prokaryotes reacting quickly and strongly to nutrient addition both in terms of abundances (max: +
700 53-68%) and production rates (max: + 787-946%; Gazeau et al., in preparation, this issue). The
701 absence of response from autotrophic stocks could be due to a tight top-down control from grazers
702 hiding potential responses from the autotrophic community (Lekunberri et al., 2010; Marañón et al.,
703 2010). Feliu et al. (2020, this issue) have shown that the mesozooplankton assemblage at TYR was
704 clearly impacted by a dust event that took place nine days before sampling at that station (François
705 Dulac, Pers. Com. 2019) and evidenced by  dust export in *in situ* deployed sediment traps (Bressac
706 et al., in preparation, this issue). This dust deposition likely stimulated phytoplankton growth and

707 consequently increased the abundance of herbivorous grazers (copepods) and attracted carnivorous
708 species. After the rapid increase observed a few hours after dust addition, DIP levels decreased to
709 reach similar levels than in control tanks at the end of this experiment (Fig. 6). Yet, heterotrophic
710 prokaryote abundances increased until the end of the experiment (Fig. 8) although production rates
711 reached a plateau after 24 h (Gazeau et al., in preparation, this issue). This is coherent with
712 measurements of the alkaline phosphatase activity that slightly increased at the end of the
713 experiment in dust-amended tanks suggesting the use of dissolved organic phosphorus by bacteria
714 to compensate for the increasing lack of DIP (Gazeau et al., in preparation, this issue). Altogether,
715 the strong stimulation of heterotrophic prokaryotes and the absence of detectable effects on the
716 autotrophic compartment drove the community towards a stronger net heterotrophic state as shown
717 by increases in community respiration and decreases in net community production rates in
718 dust-amended as compared to control tanks (Gazeau et al., in preparation, this issue).

719 At station FAST, the competition for nutrients between autotrophs and heterotrophs was
720 clearly in favor of autotrophs. While, as discussed above, all groups of primary producers benefited
721 from nutrient enrichment at this station, the increases in heterotrophic prokaryote abundances were
722 rather limited following dust deposition, leading to an increase of net community production rates
723 throughout this experiment to reach positive levels while control tanks remained below metabolic
724 balance (Gazeau et al., in preparation, this issue). At station ION, the situation was somewhat
725 intermediate with a parallel enhancement of both autotrophic and heterotrophic stocks and
726 processes, although the system was slightly in favor of net autotrophy at the end of the experiment
727 (Gazeau et al., in preparation, this issue).

728 Transfer of newly produced organic matter to higher trophic levels in the different
729 treatments was evaluated through the quantification of meso-zooplankton abundance at the end of
730 each experiment. Although we are fully aware that such an approach is certainly criticizable

731 considering the low incubation times (3 to 4 days), it may still be representative of lowered
732 mortality or faster growth. Altogether it does not appear as a surprise that an increase in
733 meso-zooplankton abundances was only detected at station FAST where the strongest enhancement
734 of primary production was observed. Such an increase in meso-zooplankton abundance in the
735 dust-amended as compared to control treatment was observed during land-based mesocosm
736 experiments in the Eastern Mediterranean Sea (Pitta et al., 2017).

737 Finally, although no clear effects of dust deposition under present conditions were detectable
738 on autotrophic prokaryotes at station TYR, the strongest increase in N₂ fixation rates was recorded
739 at this station (+434-503%, as compared to +173-256% and +41-49% at ION and FAST,
740 respectively; see Ridame et al., in preparation, this issue, for more details). However, the potential
741 impact of this process on NO_x concentration is highly negligible compared to the very large stock of
742 NO_x present in the dust-amended tanks, as less than 1 nmol L⁻¹ d⁻¹ of NO_x can be produced by this
743 process (Ridame et al., in preparation, this issue).

744 **4.4. Impact of warming and acidification**

745 Very few past studies have investigated the release and fate of nutrients from atmospheric
746 particles under climate conditions as expected for the end of the century, and, to the best of our
747 knowledge, our study represents the first attempt to test for the combined effect of ocean warming
748 and acidification on these processes. Louis et al. (2018) have already shown from an abiotic dust
749 experiment that even an extreme ocean acidification scenario (~ -0.6 pH units) does not impact the
750 bioavailability of macro- and micro-nutrients (NO_x, DIP and DFe) for surface phytoplankton
751 communities in the oligotrophic Northwestern Mediterranean Sea, using the same dust analog and
752 simulated flux as used during our experiments. Similar results were presented by Mélançon et al.
753 (2016) regarding the release of DFe from dust in high-nutrient low-chlorophyll (HNLC) waters of

754 the Northeastern Pacific, following a mild ocean acidification scenario of -0.2 pH units. Our results
755 agree with these previous findings and further highlighted the absence of direct effect of ocean
756 warming (+3 °C) on the release of nutrients from atmospheric particles.

757 The differences in nutrient consumption dynamics between ambient and warmed/acidified
758 tanks were substantially dependent on the considered nutrient and investigated station. Regarding
759 NO_x, while no impacts of warming and acidification could be observed at stations TYR and ION
760 due to large variability between the duplicates (Table 4), larger NO_x consumption rates were shown
761 under future climate conditions at the most productive station FAST as a consequence of strongly
762 enhanced biological stocks (see thereafter) and metabolic rates (Gazeau et al., in preparation, this
763 issue). The differences in DIP dynamics between the two dust-amended treatments were more
764 complex to interpret depending on the investigated station. A clear feature of our experiments is
765 that, in contrast to present day pH and temperature conditions, all the stock of DIP released from
766 dust was consumed at the end of the three experiments under future conditions. That being said, the
767 decreasing rates of DIP concentrations for that future conditions treatment differed depending on
768 the station (Table 4). While DIP dynamics were relatively similar between treatments at ION, a
769 clear effect of warming and acidification was shown at station TYR and FAST where the vast
770 majority of released DIP was consumed within 24 h ($\Delta\text{DIP} = -1.3$ and -1.1 to $-1.5 \text{ nmol L}^{-1} \text{ h}^{-1}$ at
771 TYR and FAST, respectively). An interesting outcome at station TYR was that, despite the
772 important discrepancies observed for autotrophic stocks and metabolic rates between the duplicates
773 G1 and G2 (see section 4.2), a very similar dynamics was observed for DIP concentrations in these
774 tanks. As heterotrophic prokaryote biomass and production rates (Gazeau et al., in preparation, this
775 issue) did not differ between these duplicate tanks, this further highlights the clear dominance of
776 heterotrophic processes at this station, a dominance which was exacerbated by dust addition and
777 future climate conditions.

778 At station ION, DIP consumption rates were similar following dust addition under present
779 and future conditions. This result appears surprising as large impacts of warming and acidification
780 have been observed, especially for primary producers, as shown by almost doubled chlorophyll *a*
781 concentrations as compared to dust amended tanks (D). At this station, all autotrophic groups
782 benefited from ocean acidification and warming. *Synechococcus* and to a lesser extent
783 pico-eukaryotes appeared as the most impacted ones. Yet these differences of sensitivity among
784 autotrophs did not lead to detectable changes in the composition of the autotrophic assemblage as
785 compared to ambient conditions, with still a large dominance of nano-eukaryote carbon biomass at
786 the end of this experiment (62% in treatment G vs. 64% in treatment D). Very contrasted results
787 have been shown on the effect of ocean acidification on small autotrophic species (e.g. Dutkiewicz
788 et al., 2015) while there are increasing evidences that small phytoplankton species will be favored
789 in a warmer ocean (e.g. Chen et al., 2014; Daufresne et al., 2009; Morán et al., 2010). As mentioned
790 earlier, our experimental protocol was not conceived to discriminate temperature from pH effects,
791 however results concur with those of Maugendre et al. (2015) which further suggested temperature
792 over elevated CO₂ as the main driver of increased picophytoplankton abundance. As heterotrophic
793 prokaryotes were also positively impacted by future environmental conditions, the similarity of DIP
794 dynamics between ambient and future conditions suggests a tight coupling between the autotrophic
795 and heterotrophic compartments at this station. This is further evidenced by the absence of
796 differences detected over the relatively short time duration of our experiment on meso-zooplankton
797 abundance and carbon export efficiency (Gazeau et al., in preparation, this issue).

798 At FAST, similar to what was observed at station ION, all phytoplanktonic groups were
799 positively impacted by warming and acidification with the strongest changes detected for
800 *Synechococcus* as compared to ambient conditions. However, in contrast to station ION, all groups
801 reached maximal abundances (and carbon biomass) after 3 days of incubations, thereafter

802 drastically decreasing most likely as a consequence of DIP limitation (see above). It must be
803 stressed that this pattern could not be observed through pigment dynamics as no sampling was
804 performed for these analyses after 3 days of incubation. Also, in contrast to station ION, the
805 abundance of heterotrophic prokaryotes in the warmer and acidified treatment reached a maximum
806 after 2 days of incubations and then strongly decreased to reach levels observed in the control
807 treatment. This suggests that the heterotrophic compartment was the first to suffer from DIP
808 limitation and further highlights the dominance of the autotrophic compartment in terms of nutrient
809 consumption at this station. The differential dynamics of these two compartments under warmer and
810 acidified conditions most likely led to an excess production of organic matter that translated, for
811 instance, in higher dissolved organic carbon concentrations in this treatment (Gazeau et al., in
812 preparation, this issue). This excess production as compared to ambient conditions did not seem to
813 reach higher trophic levels as no clear differences in meso-zooplankton abundances were observed.
814 We fully acknowledge that the duration of our experiments was certainly too short to carefully
815 assess the proportion of newly formed organic matter consumed by meso-zooplanktonic species and
816 its effect on their abundances, yet group-specific variations were observed. Finally, it appeared that
817 at least part of this excess organic matter was exported to the bottom of the tanks as a higher carbon
818 export efficiency was observed at this station under warmer and acidified conditions (Gazeau et al.,
819 in preparation, this issue).

820 5. Conclusion

821 These experiments conducted during the PEACETIME cruise represent the first attempt to
822 investigate the impacts of atmospheric deposition on surface plankton communities both under
823 present and future environmental conditions. Despite few experimental issues that are discussed, the
824 three experiments provided new insights on these potential impacts in the open Mediterranean Sea.
825 Interestingly, the effect of dust deposition was highly different between the three investigated
826 stations in the Tyrrhenian Sea, Ionian Sea and in the Algerian basin. As the initial conditions in the
827 sampled surface seawater at the three stations were very similar in terms of nutrient availability and
828 chlorophyll content, these differences rather seem to be a consequence of the initial metabolic states
829 of the community (autotrophy vs. heterotrophy). In all three cases, nutrient addition from dust
830 deposition did not strongly modify but rather exacerbated this initial state. Relative changes in main
831 parameters presented in this manuscript and processes presented in Gazeau et al. (in preparation,
832 this issue) as a consequence of dust addition under present and future environmental conditions are
833 shown in Fig. 10, and compared to the compilation of published data for the Mediterranean Sea
834 from Guieu and Ridame (2020). At station TYR, under conditions of a clear dominance of
835 heterotrophs on the use of resources, dust addition drove the community into an even more
836 heterotrophic state with no detectable effect on primary producers. At station ION, where the
837 community was initially closer to metabolic balance, both heterotrophic and autotrophic
838 compartments benefited from dust derived nutrients. At FAST, the most active station in terms of
839 autotrophic production, addition of nutrients boosted both compartments but heterotrophic
840 prokaryotes became quickly P-limited and overall larger effects were observed for phytoplankton.
841 Ocean acidification and warming did not have any detectable impact on the release of nutrients
842 from atmospheric particles. Furthermore, these external drivers did not drastically modify the

843 composition of the autotrophic assemblage with all groups benefiting from warmer and acidified
844 conditions. However, very large increases were observed both for autotrophic and heterotrophic
845 stocks and processes suggesting an exacerbation of effects from atmospheric dust deposition in the
846 future, rather than a change in the role of Mediterranean surface plankton community as a source or
847 a sink of CO₂ to or from the atmosphere.

848 **Data availability**

849 All data and metadata will be made available at the French INSU/CNRS LEFE CYBER database

850 (scientific coordinator: Hervé Claustre; data manager, webmaster: Catherine Schmechtig).

851 INSU/CNRS LEFE CYBER (2020)

852 **Author contributions**

853 FG and CG designed and supervised the study. FG, CG, CR and KD sampled seawater from the

854 experimental tanks during the experiments. JMG and GDL participated in the technical preparation

855 of the experimental system and all authors participated in sample analyses. FG, CR and CG wrote

856 the paper with contributions from all authors.

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1189 **Tables**

1190 Table 1. List of investigated parameters and processes during the three experiments at stations TYR,
1191 ION and FAST. Related manuscripts are indicated. pH_T: pH on the total scale, A_T: total alkalinity,
1192 ¹³C-C_T: ¹³C signature of dissolved inorganic carbon, NO_x: nitrate + nitrite, DIP: dissolved inorganic
1193 phosphorus, Si(OH)₄: silicate, DFe: dissolved iron, DAL: dissolved aluminium, Th-REE-Pa:
1194 Thorium (230Th and 232Th), Rare Earth elements and Protactinium (231Pa), POC: particulate
1195 organic carbon, DOC: dissolved organic carbon, ¹³C-DOC: ¹³C signature of dissolved organic
1196 carbon, TEP: transparent exopolymeric particles, NCP/CR: net community production and
1197 community respiration (oxygen based), ¹⁴C-PP: primary production based on ¹⁴C incorporation.

Sampling time	T-1	T0	T1	T2	T3	T4	T5	T6 / T7	Related manuscript
	Filling tanks	Before seeding, after warming / acidification)	+1 h	+6 h	+12 h	+24 h	+48 h	+72 h/+96 h	
Temperature									
		Continuous							This manuscript
Irradiance									
		Continuous							This manuscript
Carbonate chemistry									
pH _T									This manuscript
<i>A</i> _T									This manuscript
$\delta^{13}\text{C}$ -C _T									Gazeau et al. (in preparation)
Macro-nutrients									
NO _x									This manuscript
DIP									This manuscript
Si(OH) ₄									This manuscript
Micro-nutrients									
DFe									Roy-Barman et al. (in preparation)



	Gazeau et al. (in preparation)
	Gazeau et al. (in preparation)
Processes	
Amino acids	
Carbohydrates	
NCP/CR	
^{14}C -PP	
Heterotrophic production	
Ectoenzymatic activity	
N_2 fixation	
$^{13}\text{CO}_2$ -fixation	
Virus production, lysogeny	Dinasquet et al. (in preparation)

1200 Table 2. Initial conditions as measured while filling the tanks (initial conditions in pumped surface
1201 water; sampling time: t-12h). pH_T: pH on the total scale, NO_x: nitrate + nitrite, NH₄: ammonium,
1202 DIP: dissolved inorganic phosphorus, Si(OH)₄: silicate, TChla: total chlorophyll *a*, HNF:
1203 heterotrophic nanoflagellates. The three most important pigments in terms of concentration are also
1204 presented (19'-hexanoyloxyfucoxanthin, Zeaxanthin and Divinyl Chlorophyll *a*). Biomasses of the
1205 different groups analyzed through flow cytometry were estimated based on conversion equations
1206 and/or factors found in the literature (see section 2.3). Autotrophic biomass was, as a first
1207 approximation, estimated only based on flow cytometry data and therefore corresponds to the
1208 fraction < 20 µm. Heterotrophic biomass was estimated as the sum of heterotrophic prokaryote and
1209 HNF biomasses (see section 2.3). Values below detection limits are indicated as < dl.

	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 sampling (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
Carbonate chemistry	pH _T	8.04	8.07	8.03
Nutrients	Total alkalinity ($\mu\text{mol kg}^{-1}$)	2529	2627	2443
	NO _x (nmol L ⁻¹)	14.0	18.0	59.0
	NH ₄ ⁺ ($\mu\text{mol L}^{-1}$)	0.045	0.022	< dl
	DIP (nmol L ⁻¹)	17.1	6.5	12.9
Pigments	Si(OH) ₄ ($\mu\text{mol L}^{-1}$)	1.0	0.96	0.64
	NO _x /DIP (molar ratio)	0.8	2.5	4.6
	TChla ($\mu\text{g L}^{-1}$)	0.063	0.066	0.072
	19'-hexanoyloxyfucoxanthin ($\mu\text{g L}^{-1}$)	0.017	0.021	0.016
	Zeaxanthin ($\mu\text{g L}^{-1}$)	0.009	0.006	0.036
	Divinyl Chlorophyll <i>a</i> ($\mu\text{g L}^{-1}$)	~0	0	0.014

Flow cytometry	Pico-eukaryotes (abundance in cell mL ⁻¹ ; biomass in µg C L ⁻¹)	347.8; 0.5	239.9; 0.4	701.0; 1.0
	Nano-eukaryotes (abundance in cell mL ⁻¹ ; biomass in µg C L ⁻¹)	150.5; 3.9	188.8; 4.8	196.6; 5.0
	<i>Synechococcus</i> (abundance in cell mL ⁻¹ ; biomass in µg C L ⁻¹)	4972; 1.2	3037; 0.8	6406; 1.6
	Autotrophic biomass (µg C L ⁻¹)	5.6	6.0	7.7
	Heterotrophic prokaryotes abundance (x 10 ⁵ cell mL ⁻¹)	4.79	2.14	6.15
	HNF (abundance in cell mL ⁻¹)	110.1	53.6	126.2
	Heterotrophic biomass (µg C L ⁻¹)	9.9	4.5	12.7
	Pennate diatoms (abundance in cell L ⁻¹)	140	520	880
Microscopy	Centric diatoms (abundance in cell L ⁻¹)	200	380	580
	Dinoflagellates (abundance in cell L ⁻¹)	2770	3000	3410
	Autotrophic flagellates (abundance in cell L ⁻¹)	0	60	650
	Ciliates (abundance in cell L ⁻¹)	270	380	770

1212 Table 3. Maximum input of nitrate + nitrite (NO_x) and dissolved inorganic phosphorus (DIP)

1213 released from Saharan dust in tanks D and G as observed from the two discrete samplings

1214 performed over the first 6 h after seeding. The estimated maximal percentage of dissolution is also

1215 presented (see section 2.3 for details on the calculations).

	NO _x				DIP			
	D1	D2	G1	G2	D1	D2	G1	G2
Maximum input	$\mu\text{mol L}^{-1}$				nmol L^{-1}			
TYR	11.0	11.1	11.1	11.0	24.6	20.4	24.6	23.9
ION	11.2	11.6	11.2	11.3	23.3	22.0	19.6	22.9
FAST	11.3	11.1	11.1	11.2	30.8	31.3	36.9	29.8
Percentage of dissolution (%)								
TYR	95	96	95	94	12	10	12	11
ION	96	99	96	97	11	10	9	11
FAST	97	97	95	97	15	15	17	14

1216

1217 Table 4. Removal rate of nitrate + nitrite (NO_x) and dissolved inorganic phosphorus (DIP) in tanks
 1218 D and G during the three experiments (TYR, ION and FAST). For NO_x , decreasing rates were
 1219 estimated based on linear regressions between maximal concentrations (i.e. after dust enrichment, at
 1220 t1h or t6h) and final concentrations (t72 h for TYR and ION and t96h for FAST). For DIP,
 1221 decreasing rates were estimated based on linear regressions between maximal concentrations (i.e.
 1222 after dust enrichment at t1h or t6h) and concentrations measured at sampling times after which a
 1223 stabilization was observed. This sampling time is shown in parentheses. All rates are expressed in
 1224 $\text{nmol L}^{-1} \text{ h}^{-1}$.

	NO _x			DIP		
	TYR	ION	FAST	TYR	ION	FAST
D1	-6.5	-8.6	-14.3	-0.4 (t72h)	-0.5 (t48h)	-0.2 (t96h)
D2	-1.0	-8.6	-13.5	-0.3 (t72h)	-0.8 (t24h)	-0.2 (t96h)
G1	-6.7	-13.1	-21.6	-1.3 (t24h)	-0.8 (t24h)	-1.5 (t24h)
G2	-0.8	-1.6	-25.2	-1.3 (t24h)	-1.6 (t24h)	-1.1 (t24h)

1225

1226 Table 5. Maximum relative changes in tanks D and G as compared to controls (average between C1
1227 and C2), expressed as a %, for the three experiments (TYR, ION and FAST). The sampling time at
1228 which these maximum relative changes were observed is shown in brackets. Tchla refers to the
1229 concentration of total chlorophyll *a* and B_{micro} to the biomass proxy of micro-phytoplankton (sum of
1230 Fucoxanthin and Peridinin, see Material and Methods) based on high performance liquid
1231 chromatography (HPLC). HP and HNF refer to heterotrophic prokaryote and heterotrophic
1232 nanoflagellate abundances, respectively, as measured by flow cytometry.

Experiment	Tank	HPLC				Flow cytometry			
		TChla	B _{micro}	Pico-eukaryotes	Nano-eukaryotes	Synechococcus	HP	HNF	
TYR	D1	-35 (t24h)	-33 (t12h)	-75 (t72h)	-80 (t1h)	-71 (t48h)	68 (t72h)	352 (t72h)	
TYR	D2	-38 (t12h)	-39 (t24h)	-75 (t72h)	-80 (t1h)	-72 (t48h)	53 (t72h)	100 (t72h)	
TYR	G1	60 (t72h)	52 (t72h)	-75 (t1h)	89 (t72h)	76 (t72h)	67 (t72h)	1095 (t72h)	
TYR	G2	359 (t72h)	392 (t72h)	323 (t72h)	119 (t72h)	700 (t72h)	68 (t48h)	298 (t72h)	
ION	D1	183 (t72h)	157 (t72h)	126 (t72h)	89 (t72h)	317 (t72h)	128 (t72h)	44 (t72h)	
ION	D2	109 (t72h)	156 (t72h)	117 (t72h)	-59 (t1h)	390 (t72h)	133 (t72h)	27 (t72h)	
ION	G1	399 (t72h)	454 (t72h)	458 (t72h)	256 (t72h)	805 (t72h)	176 (t72h)	175 (t72h)	
ION	G2	426 (t72h)	612 (t72h)	510 (t72h)	292 (t72h)	1425 (t72h)	161 (t72h)	129 (t72h)	
FAST	D1	318 (t96h)	356 (t96h)	113 (t96h)	208 (t72h)	348 (t96h)	27 (t96h)	-38 (t96h)	
FAST	D2	237 (t96h)	322 (t96h)	91 (t96h)	219 (t72h)	197 (t96h)	40 (t48h)	-49 (t96h)	
FAST	G1	399 (t96h)	415 (t96h)	198 (t72h)	274 (t72h)	357 (t48h)	61 (t48h)	243 (t24h)	
FAST	G2	395 (t96h)	421 (t96h)	129 (t72h)	202 (t96h)	344 (t48h)	67 (t48h)	74 (t24h)	

1234 **Figure captions**

1235 Fig. 1. Map showing the sampling stations in the Mediterranean Sea along the transect performed
1236 onboard the R/V “Pourquoi Pas ?” during the PEACETIME cruise.

1237 Fig. 2. Scheme of an experimental tank (climate reactor).

1238 Fig. 3. Proportion of the different pigments, as measured by high performance liquid
1239 chromatography (HPLC) in pumped surface seawater for the three experiments (t-12h).

1240 Fig. 4. Continuous measurements of temperature and irradiance level (PAR) in the six tanks during
1241 the three experiments. The dashed vertical line indicates the time of dust seeding (after t0).

1242 Fig. 5. pH on the total scale (pH_T) and total alkalinity (A_T) measured in the six tanks during the
1243 three experiments. The dashed vertical line indicates the time of dust seeding (after t0). Error bars
1244 correspond to the standard deviation based on analytical triplicates.

1245 Fig. 6. Nutrients (nitrate + nitrite: NO_x , dissolved inorganic phosphorus: DIP, silicate: $Si(OH)_4$ as
1246 well as the molar ratio between NO_x and DIP, measured in the six tanks during the three
1247 experiments. The dashed vertical line indicates the time of seeding (after t0).

1248 Fig. 7. Concentrations of total chlorophyll *a* and major pigments, measured by high performance
1249 liquid chromatography (HPLC), in the six tanks during the three experiments. The dashed vertical
1250 line indicates the time of seeding (after t0).

1251 Fig. 8. Abundance of pico-eukaryotes, nano-eukaryotes, *Synechococcus*, heterotrophic prokaryotes
1252 (HP), and heterotrophic nano-flagellates (HNF), measured by flow cytometry, in the six tanks
1253 during the three experiments. The evolution of autotrophic biomass (see Material and Methods for

1254 details on the calculation) is also shown. The dashed vertical line indicates the time of seeding (after
1255 t0).

1256 Fig. 9. Abundances of meso-zooplankton species as measured at the end of each experiment.

1257 Fig. 10. Maximal relative change (%) of main biological stocks (TChla: total chlorophyll *a*, HP:

1258 heterotrophic prokaryotes) and processes (BP: bacterial production; PP: ^{14}C -based primary

1259 production; see Gazeau et al., in preparation, this issue; BR: bacterial respiration (no data from this

1260 study); and N_2 fixation, see Ridame et al., in preparation, this issue) as obtained during the present

1261 study at the 3 stations (TYR, ION and FAST) under ambient conditions of pH and temperature

1262 (open red squares) and future conditions (full green squares). Squares are delimited by the range of

1263 responses observed among the duplicates for each treatment. The dotted green squares for station

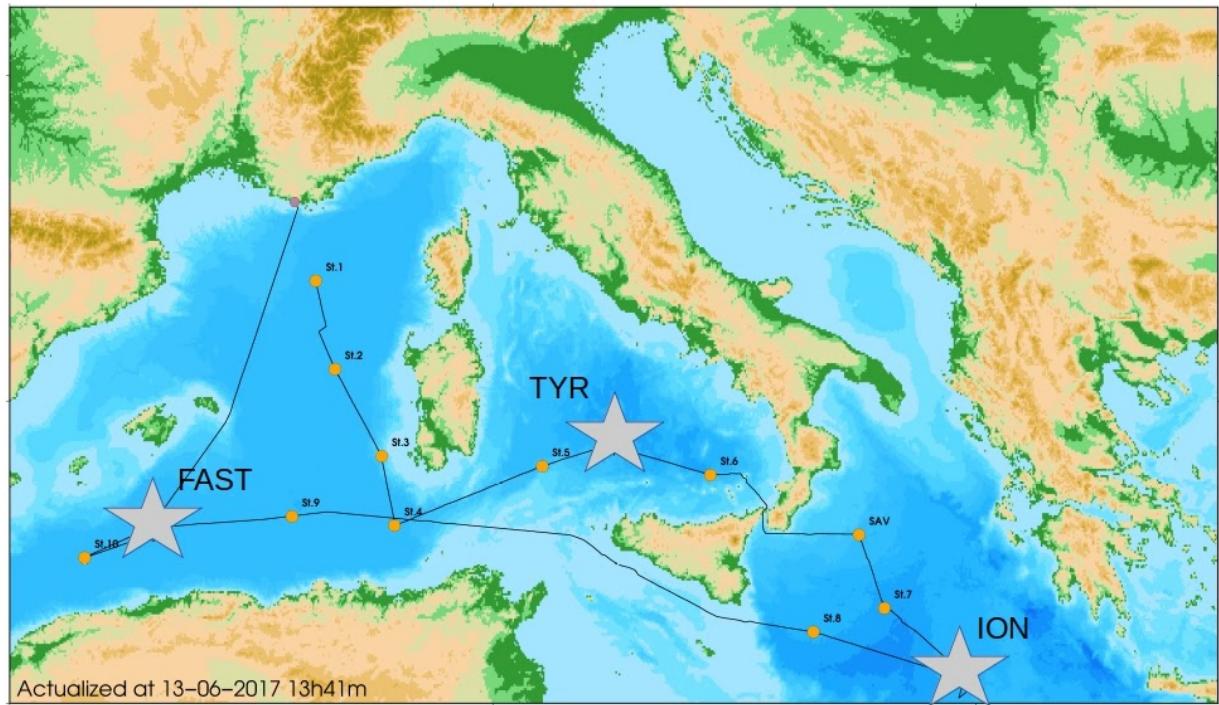
1264 TYR denote the large variability observed between duplicates for some parameters and processes

1265 that prevented drawing solid conclusions. Box-plots represent the distribution of responses

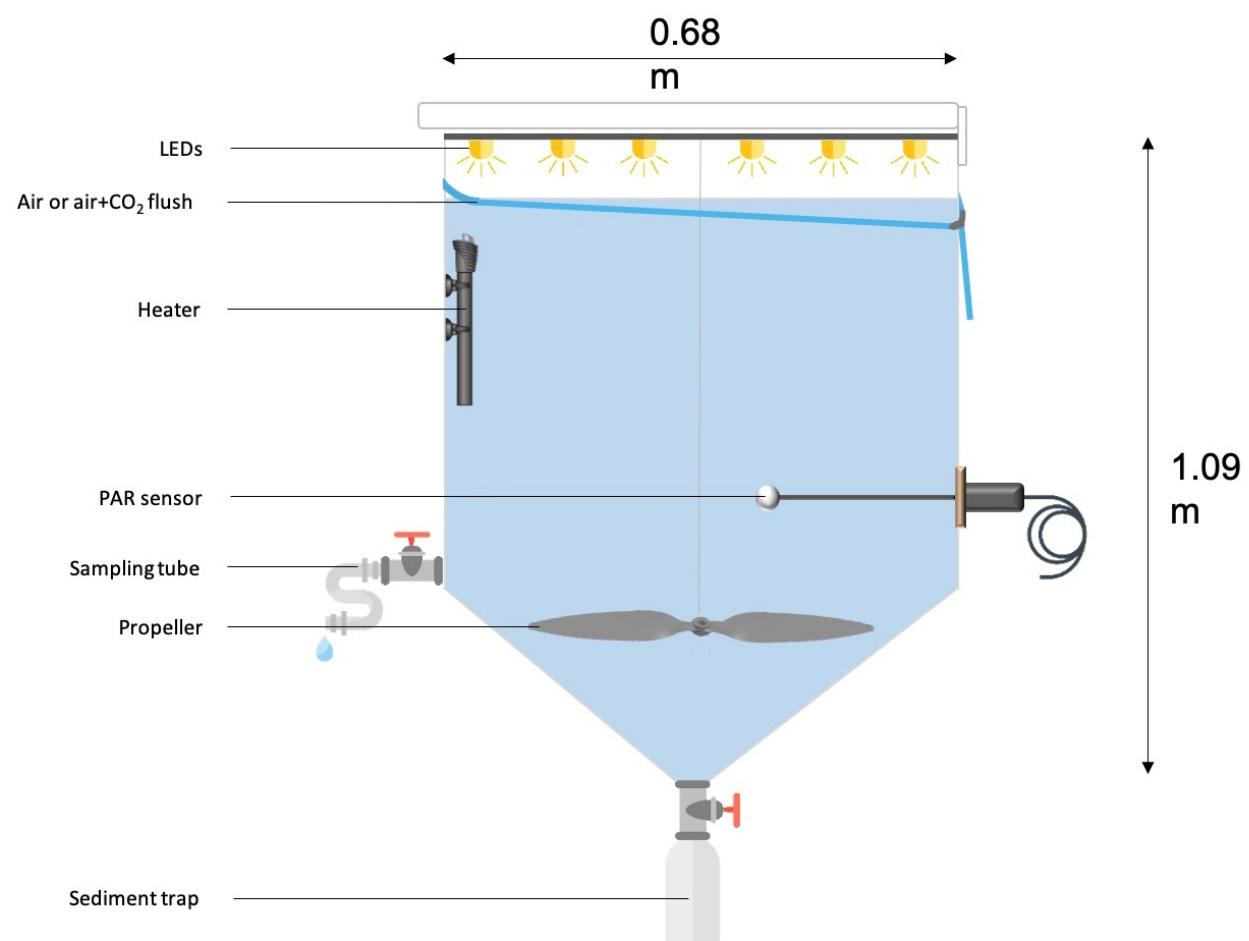
1266 observed from studies conducted in the Mediterranean Sea, as compiled by Guieu and Ridame

1267 (2020).

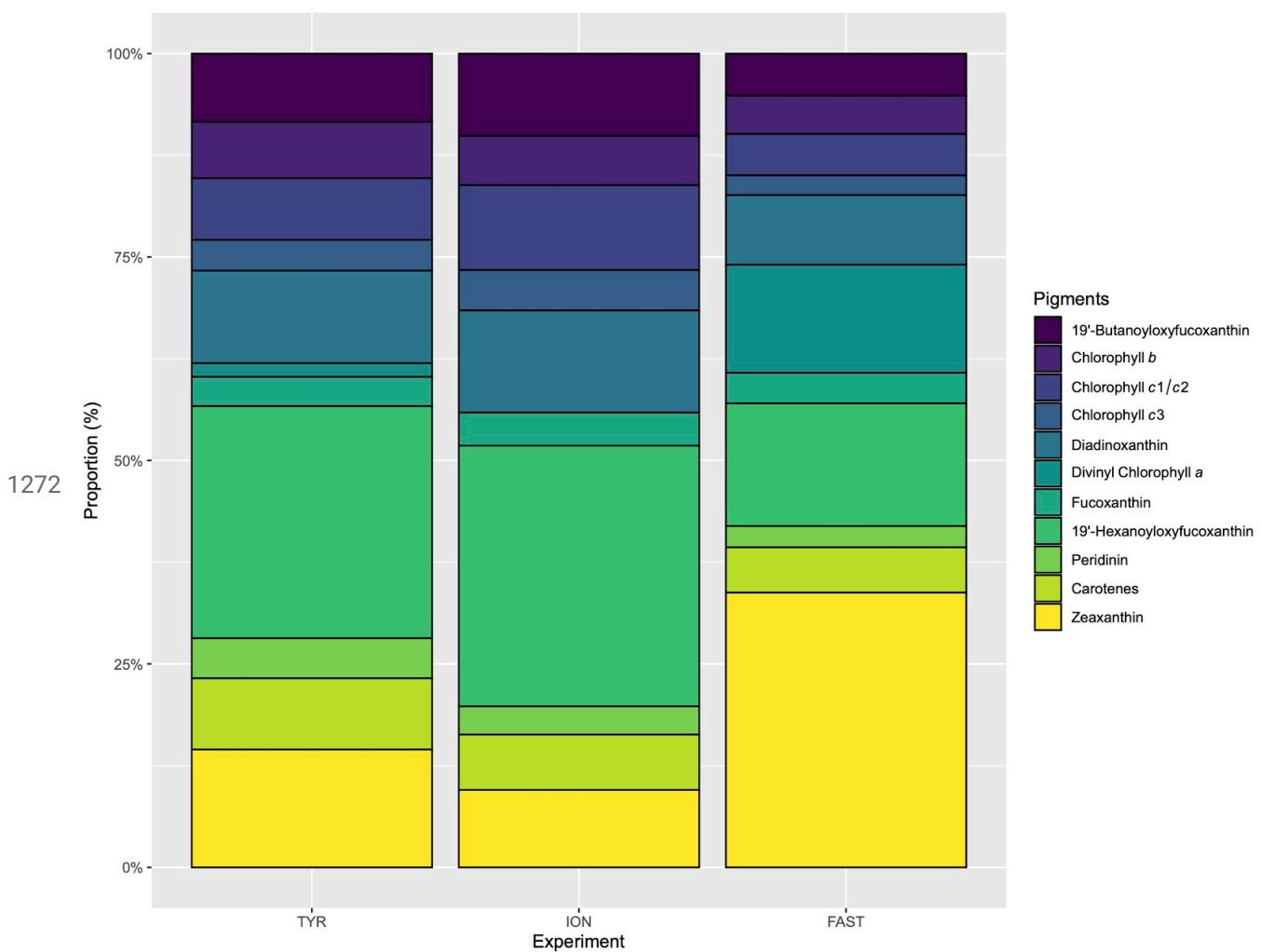
1268

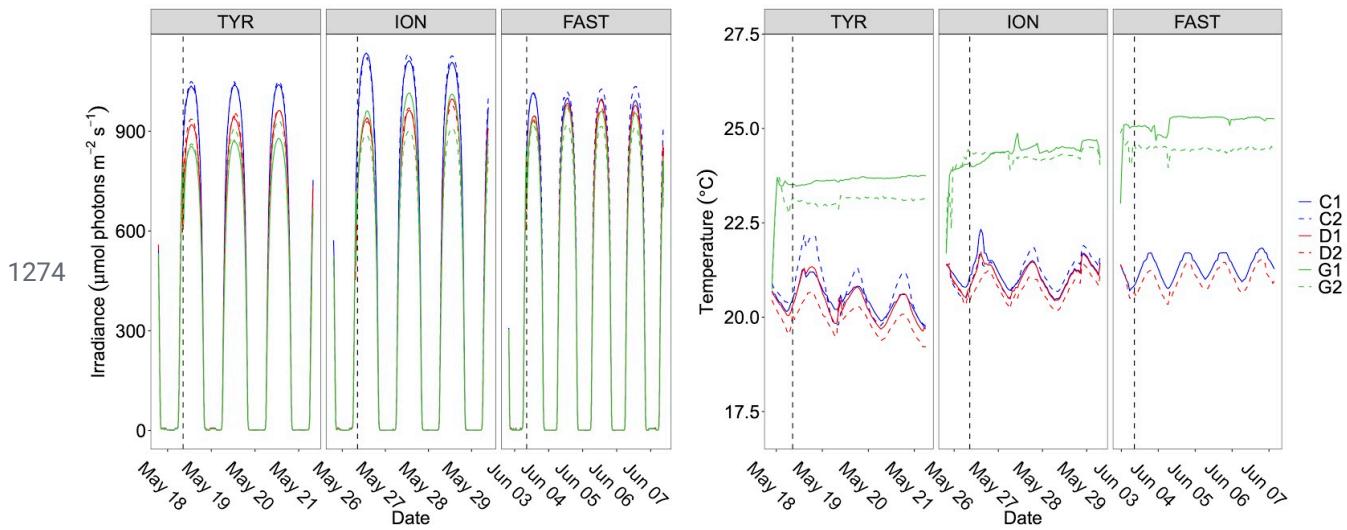


1269 Fig. 1.

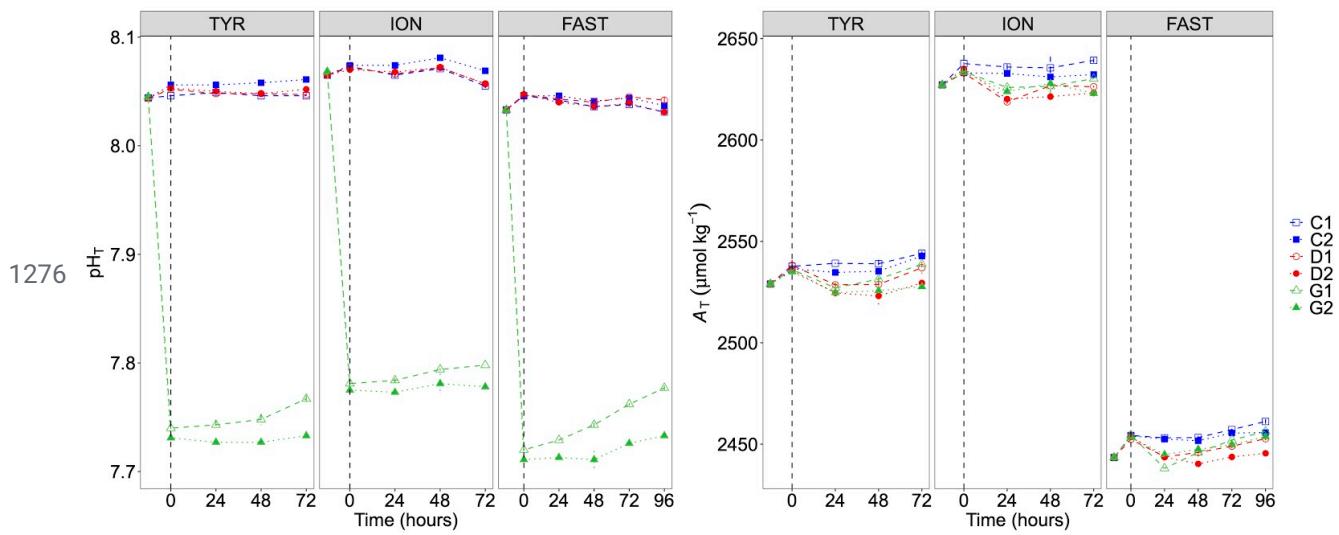


1271 Fig. 2.

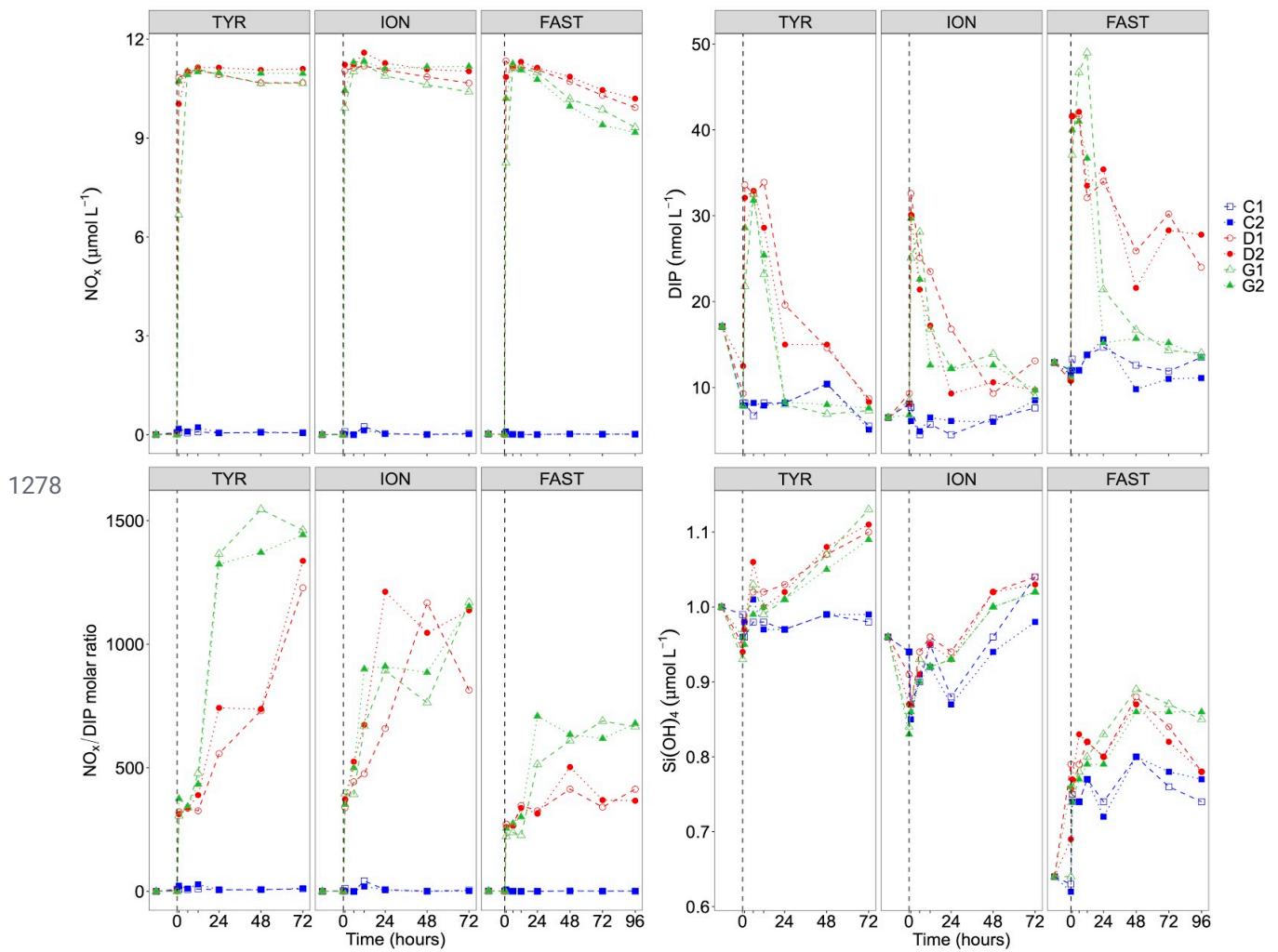




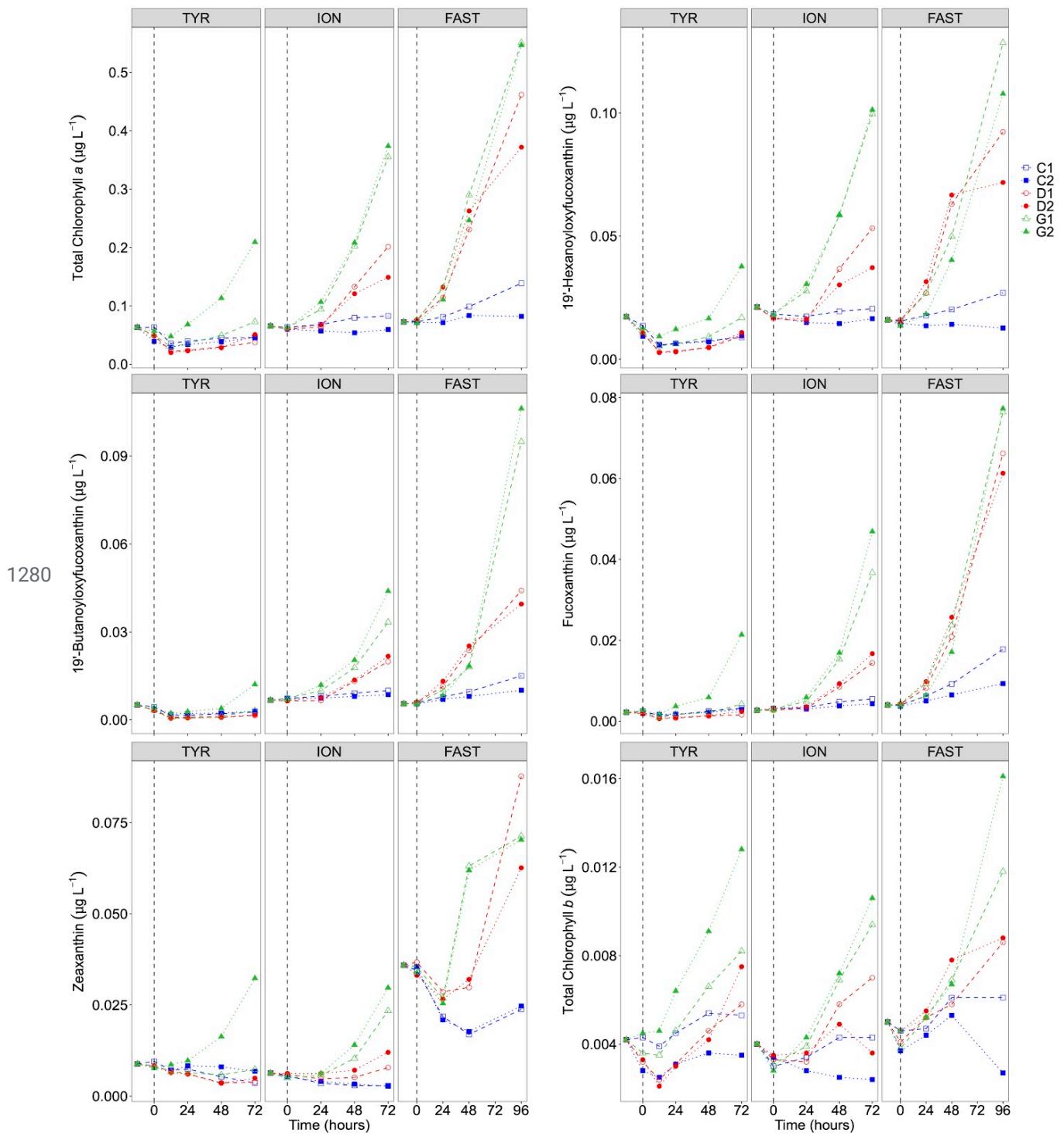
1275 Fig. 4.



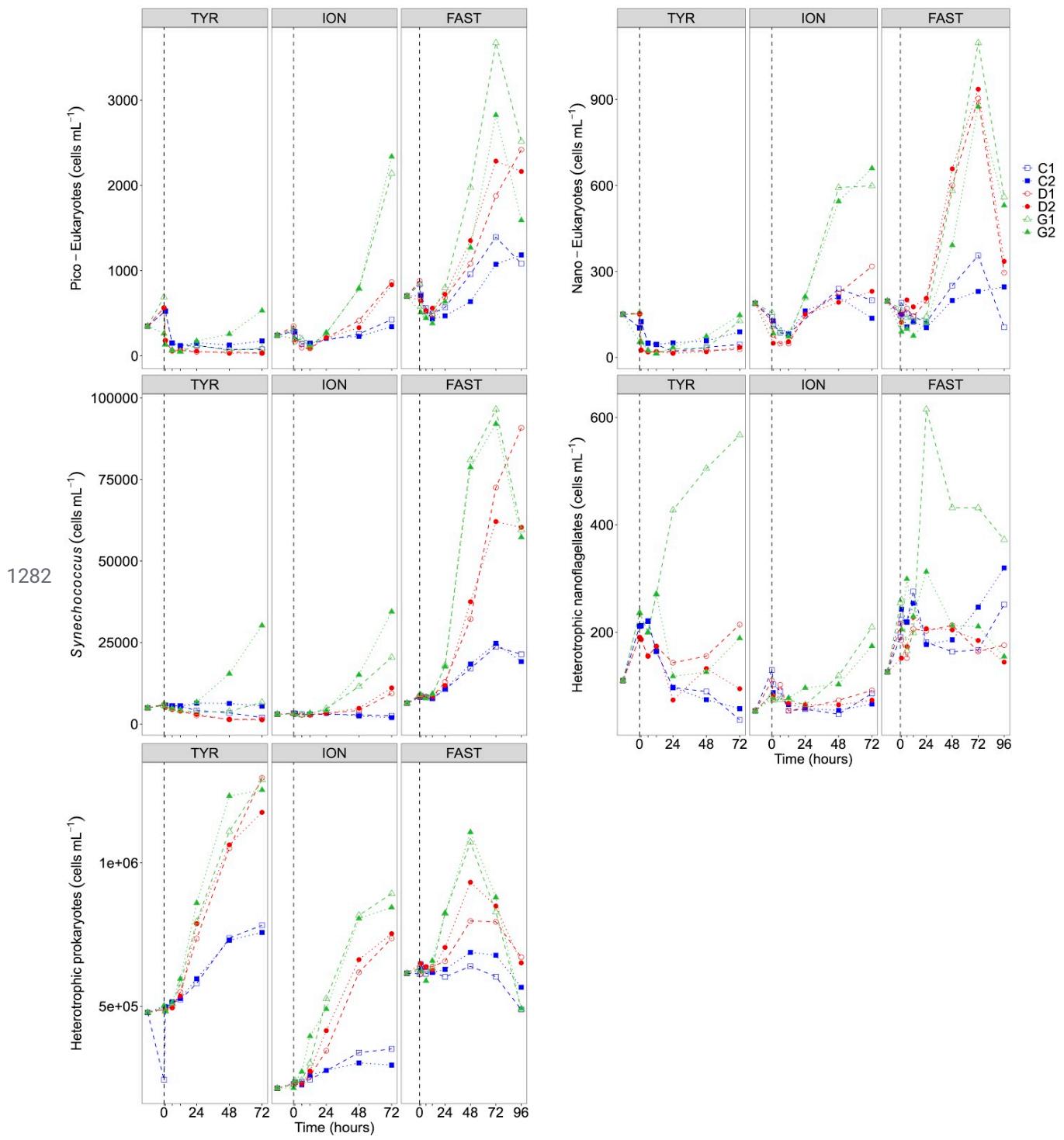
1277 Fig. 5.



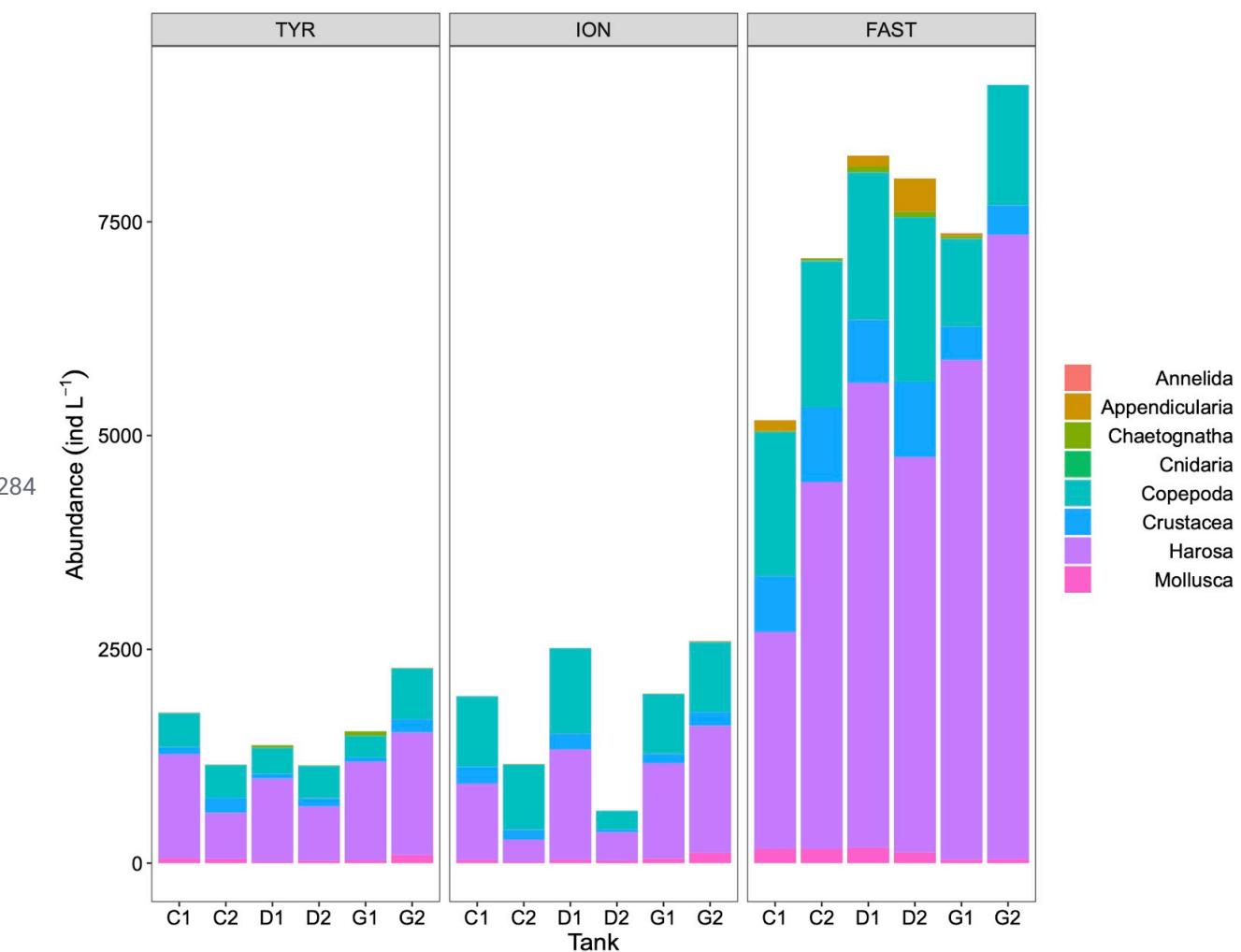
1279 Fig. 6.



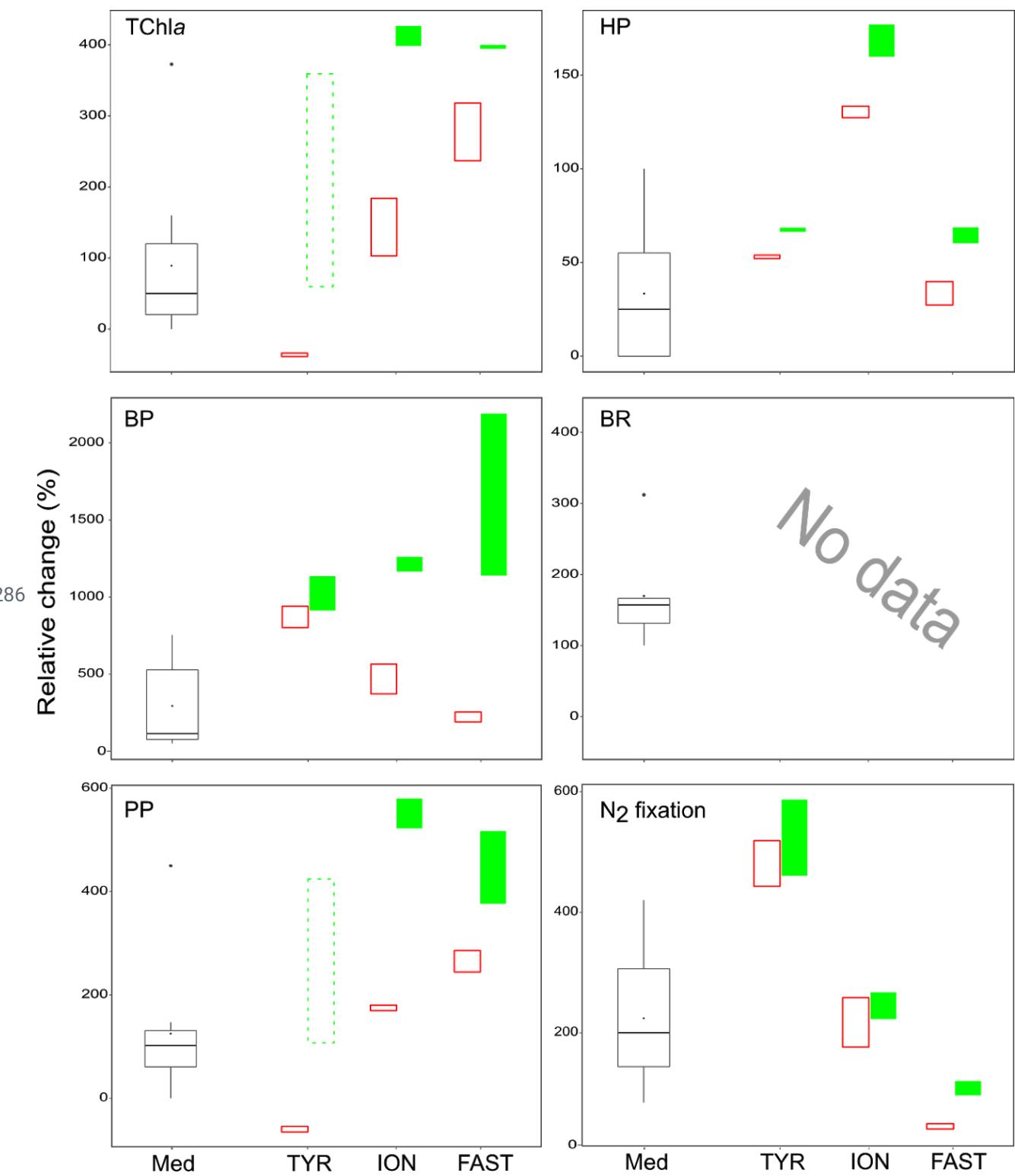
1281 Fig. 7.



1283 Fig. 8.



1285 Fig. 9.



1287 Fig. 10.