1	A year in the life of the Eastern Mediterranean:
2	Monthly dynamics of phytoplankton and bacterioplankton in an ultra-oligotrophic sea
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23	eukaryotes.

24 Highlights

- Bacterioplankton dynamics were assessed monthly in the Eastern Mediterranean Sea
- Small-sized picophytoplankton numerically dominated the phytoplankton community
- Seasonal phytoplankton dynamics are similar to BATS and Red Sea, but not to HOT
- Annual primary productivity is among the lowest in the world's oceans
- Bacterial to primary production ratio is higher than most oligotrophic seas

30 Abstract

31 The Eastern Mediterranean Sea (EMS) is a poorly studied ultra-oligotrophic marine 32 environment, dominated by small-size phyto- and bacterioplankton. Here, we describe the 33 dynamics of a single annual cycle (2018-19) of phyto- and bacterioplankton (abundances, 34 pigments and productivity) in relation to the physical and chemical conditions in the photic 35 water column at an offshore EMS site (Station THEMO-2, ~1,500m depth, 50km offshore). We show that phytoplankton biomass (as chlorophyll a), primary and bacterial productivity 36 differed between the mixed winter (January-April) and the thermally stratified (May-37 38 December) periods. Prochlorococcus and Synechococcus numerically dominated the 39 picophytoplankton populations, with each clade revealing different temporal and depth changes indicative to them, while pico-eukarvotes (primarily haptophytes) were less 40 abundant, yet likely contributed significant biomass. Estimated primary productivity (~32 gC 41 m⁻² y⁻¹) was lower compared with other well-studied oligotrophic locations, including the 42 43 north Atlantic and Pacific (BATS and HOT observatories), the western Mediterranean 44 (DYFAMED observatory) and the Red Sea, and was on-par with the ultra-oligotrophic South Pacific Gyre. In contrast, integrated bacterial production (~11 gC m⁻² y⁻¹) was similar to other 45 46 oligotrophic locations. Phytoplankton seasonal dynamics were similar to those at BATS and 47 the Red Sea, suggesting an observable effect of winter mixing in this ultra-oligotrophic 48 location. These results highlight the ultra-oligotrophic conditions in the EMS and provide, for 49 the first time in this region, a full-year baseline and context to ocean observatories in the 50 region.

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1 Introduction

Convective mixing of the water column is one of the main mechanisms responsible 53 54 for transport of nutrients to the photic zone of oligotrophic oceans and seas, often resulting in increased phytoplankton biomass and activity (Behrenfeld, 2010). This process usually 55 56 occurs during wintertime upon the progressive cooling of the sea surface, although other 57 mechanisms may also deliver nutrients to the mixed layer such as physical upwelling along shorelines, frontal systems and gyres (Anabalón et al., 2016), nutrient runoff from rivers 58 (Jickells, 1998), and atmospheric deposition (Guieu et al., 2014). Stratification is established 59 60 during springtime, nutrients gradually become depleted, which, together with other processes such as increased predation, lead to a decline in algal biomass and productivity (Behrenfeld 61 62 and Boss, 2014). This cycle has been extensively studied over decades in the major 63 oligotrophic gyres (e.g. HOT and BATS) (Steinberg et al., 2001), as well in other locations including the Western Mediterranean and Red Seas (Genin et al., 2018; Marty et al., 2002; 64 65 Marty and Chiavérini, 2010), and is considered a fundamental driving force of marine 66 ecosystem structure and function.

67 The offshore water of the Eastern Mediterranean Sea are considered ultra-oligotrophic 68 (Berman-Frank and Rahav, 2012; Siokou-Frangou et al., 2010). The oligotrophic nature of 69 this system, and especially of the Eastern Mediterranean Sea (EMS), is mainly driven by its 70 general anti-estuarine circulation (Pinardi and Masetti, 2000). Additionally, the relatively 71 stable density stratification throughout most of the year (winter mixing rarely exceeds the 200 72 m depth, (D'Ortenzio et al., 2005) has been suggested to result in a very low supply of deep 73 water nutrients to the euphotic zone (Hazan et al., 2018). Finally, modern riverine inputs into 74 the EMS are extremely low, especially since the Nile was dammed in the 1960s (Krom et al., 2014). The abovementioned conditions makes the EMS, and specifically the easternmost 75

76 Levantine basin, among the warmest, saltiest and least productive waters in the world (Ozer77 et al., 2017).

78 Despite the importance of the EMS coastline in providing ecosystem services to over 70 79 million people e.g. (Peled et al., 2018), it lacks the continued high resolution records 80 available from other oligotrophic regions such as the North Atlantic (the Bermuda Atlantic 81 Time Series, BATS), North Pacific (The Hawaii Ocean Time-Series station ALOHA, HOT), 82 the Western Mediterranean e.g. station DYFAMED (Marty et al., 2002; Marty and 83 Chiavérini, 2010) and, to a lesser extent, the Red Sea (Shaked and Genin, 2017). Specifically, 84 detailed phytoplankton and bacterioplankton time-series are missing, and are important in order to both understand the current system and the ways it may be impacted by local, 85 86 regional and global change (e.g. (Marty and Chiavérini, 2010). Thus, studying the dynamics of the microorganisms at the base of the marine food-web over time are of great ecological 87 88 importance. 89 Studies from the early 1980's showed a clear seasonal cycle of chlorophyll a (a proxy of algal 90 biomass) and primary production in surface waters of both neritic and pelagic locations, with 91 the changes attributed primarily to picophytoplankton (Azov, 1986; Berman et al., 1984; 92 Kimor and Wood, 1975; Yacobi et al., 1995). More recent studies of the EMS show similar 93 seasonal trends using remote-sensing of surface chlorophyll a (Rosenberg et al., 2020) and in 94 measurements of cell numbers, production and dinitrogen fixation in coastal waters (Rahav et 95 al., 2018; Raveh et al., 2015). However, to the best of our knowledge, no detailed temporal 96 (monthly) measurements have been presented of microbial processes and phytoplankton 97 community structure at the offshore EMS waters. In this study, we followed the composition 98 and activity of phytoplankton and bacterioplankton from a pelagic location (station THEMO-

2) in the offshore EMS over a full year at monthly temporal resolution, and at high depth

100 resolution (~20 samples, ~12 of them across the photic zone). These measurements were

101 performed as part of the SoMMoS (Southeastern Mediterranean Monthly cruise Series) 102 campaign, which compared an open-ocean station with one at the edge of the continental 103 shelf (Figure 1A). Additional studies from this cruise series will focus on the carbonate 104 system (Juntao et al. *in prep*), nutrient dynamics (Ben Ezra et al., 2021), coccolithophore 105 dynamics (Keuter et al. in prep) and a detailed comparison of the offshore and coastal 106 stations (Krom et al. in prep). Together, these studies provide a baseline allowing improved 107 interpretation for future research from the two long-term ocean observatories recently 108 established in the EMS: the DeepLev (Katz et al., 2020) and THEMO (Diamant et al., 2020), 109 as well as comparison to long-term monitoring activities (Ozer et al., 2017; Rahav et al.,

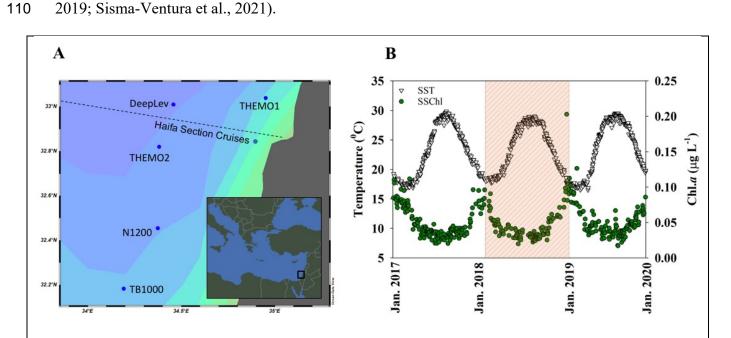


Figure 1: Overview of the SoMMoS sampling locations context. A) A map of the Levantine Basin (EMS) showing the location of the THEMO-2 sampling site (the focus of this study), as well as of additional observatories and locations of long-term studies discussed here: THEMO-1, DeepLev (Katz et al., 2020), N-1200 (Rosenberg et al., 2020), TB1000 (Yogev et al., 2011) and the Haifa section transect cruises (Ozer et al., 2017). At the bottom right corner an inset of the EMS with a black frame indicating the study site. B) Changes in satellite-derived sea surface temperature (orange) and sea

surface chlorophyll a (green) from the 1 km² region surrounding THEMO-2 between 2017-2020. Data were extracted from the Copernicus Marine Environment Monitoring Service (CMEMS, https://marine.copernicus.eu/). The period of this study is shaded.

111

112 2 Materials and Methods

113 **2.1** Cruises and sample collection

114 Water samples were collected as part of the SoMMoS project during twelve cruises, from February 2018 until January 2019. Each cruise took samples at two stations: THEMO2, an 115 116 open ocean station at a depth of ~1500 m (~50 Km from the coast, 32.820 N, 34.380 E), and THEMO1 which is positioned at the edge of the continental shelf at ~125m depth (~10 Km 117 118 from the coast, 33.040 N, 34.950 E). THEMO2 was sampled from 12:00-22:00 (local time), while THEMO1 was sampled during the night, typically 00:00-02:00 (local time). Here, we 119 120 will present information from data collected at the offshore THEMO2 station. Samples were 121 collected using a 12-bottle rosette with 8 L Niskin bottles. Samples were collected at 20-24 122 depths across the entire water column (11-12 bottles between the surface and 200 m, which 123 we define here as the photic zone). Sampling depths were selected based on real-time data of 124 Conductivity, Temperature, Depth (CTD) profiler (Seabird 19 Plus) from the down-cast before each sample collection in the up-cast, an oxygen optode (on some cruises) and a 125 fluorescence meter (Turner designs, Cyclops-7). The continuous data was processed using the 126 127 Sea-Bird data conversion software, and minimized using bin averaging. One bin of data lines 128 was defined as a change of 1 decibar (db) between each bin. The first two meters of 129 measurements were compiled together to account for sensors and CTD pump adjustment time 130 and rosette depth while at sea surface. Each cast allocated water for analysis in the following 131 order to give priority to the more 'sensitive' parameters: dissolved methane, pH, alkalinity, 132 dissolved inorganic carbon, inorganic nutrients, total and dissolved organic carbon, cell

count, primary and bacterial production, DNA and algal pigment markers. Pre-filtered 133 134 inorganic nutrient samples were analyzed fresh (unfrozen) the day after the cruise, using a 135 SEAL AA-3 autoanalyzer system, and are described in detail elsewhere (Ben-Ezra et al., 2021). A summary of all of the currently-available measurements can be found in 136 137 Supplementary Table S1, and the BCO-DMO (acronym: SoMMoS) and ISRAMAR 138 (https://isramar.ocean.org.il/isramar2009/) databases. Mixed layer depth (MLD) was calculated using a temperature difference of $\Delta 0.3$ °C (Mena et al., 2019). Calculations based 139 140 on a density difference of $\Delta 0.15 \text{ kg/m}^3$ yielded similar results. During several months 141 (February-April 2018 and January 2019), the density plots revealed a progressive increase in 142 density without a clear pycnocline but with multiple "bumps", indicative of water column 143 instability down to below 200 m (Supplementary Figure S1). At these times, the MLD calculations based on a defined difference in temperature or salinity from the surface 144 145 preclude a robust estimate of the mixed layer depth, as they may underestimate the actual 146 values. Based on these calculations, we divide the study period into 'a generally mixed 147 period' during January-April (winter/spring), and a 'stratified period' during May-December (summer/autumn, Table 1). Estimates based on the vertical distribution of inorganic nutrient 148 149 concentrations suggest the mixing period may have begun as early as November 2018 (Ben-150 Ezra et al., 2021). We note that the monthly sampling resolution likely to 'miss' short-lived 151 deep mixing events, as can be observed from mooring operations (e.g. (Gunn et al., 2020). 152

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153 **2.2 Bacterial and primary productivity**

154 Heterotrophic prokaryotic productivity (hereafter referred to as bacterial productivity, BP)

155 was estimated using the ³H-leucine incorporation method (Simon and Azam, 1989).

156 Triplicate 1.7 ml of ocean water were taken from each sampled depth and incubated with a

157 7:1 mixture of 'cold' leucine and 'hot' ³H-leucine (final concentration 100 nmol leucine L⁻¹)

for 4 h at room temperature in the dark immediately after sampling. Preliminary experiments 158 159 show that this was a saturating level of leucine in the offshore water of the SE Mediterranean 160 Sea. After incubation, incorporation was terminated by adding 100 µl trichloroacetic acid (TCA). As a negative control for non-specific binding, another set of triplicates were sampled 161 162 from a surface layer and treated with TCA immediately after the addition of the radioactive 163 tracer. At the end of each cruise, the samples were processes using the micro-centrifugation protocol and 1ml scintillation cocktail (ULTIMA-GOLD) was added to all samples before 164 165 counted using TRI-CARB 2100 TR (PACKARD) scintillation counter. A conversion factor 166 of 3 kg C per mole of leucine incorporated and an isotopic dilution of 2.0 were used to 167 calculate the C incorporated (Simon and Azam, 1989). Net daily photosynthetic carbon fixation rates were estimated using the ¹⁴C incorporation 168 169 method (Nielsen, 1952), with several modifications (Hazan et al., 2018). Triplicate 50 ml 170 samples were taken from each depth within the photic zone and from one aphotic depth using 171 sterile vials and kept at surface light and temperature conditions. The 'dark' sample served as 172 blank and was kept under the same temperature as the 'light' samples. Radioactive spiking was done at ~08:00 AM the day following of the cruise in order to start a 24h incubation for 173 174 all samples (including those collected at THEMO-1 station, not shown) at the same time. 175 Early work by Letelier and colleagues (1996) at station HOT showed that prolonged on-deck 176 incubations, similarly to the protocol used in this study, may result in underestimated PP rates 177 as it cannot precisely mimic the temperature and illumination levels in-situ. Our preliminary 178 tests concur with this conclusion and found that ashore incubations underestimate PP rates by 179 up to $\sim 20\%$ compared to incubations onto a mooring rope tied to the ship (Figure S2). Samples were spiked with 50µl (5 µCi) of NaH¹⁴CO₃ tracer and were incubated for 24 h 180 181 under 3 light regimes: surface illumination (samples from the upper mixing depths), 50% 182 illumination (samples from below the mixing depth to the DCM) and $\sim 1\%$ illumination

183	(samples from the DCM and below). Shading was performed using neutral density nets, thus
184	changing light intensity but not spectral properties. Water samples were then filtered through
185	GF/F filters (0.7 μ m nominal pore size, 25 mm diameter) using low vacuum pressure (< 50
186	mmHg) and rinsed 3 times with filtered sea water. Filters from each sample were then put in
187	scintillation vials where 50 μl of 32%HCl solution was immediately added in order to remove
188	excess ¹⁴ C-bicarbonate and kept overnight for incubation. After incubation 5 mL scintillation
189	cocktail (ULTIMA-GOLD) was added to the samples and counted using TRI-CARB 2100
190	TR (PACKARD) scintillation counter. Three random aliquots were counted immediately
191	after the addition of the radiotracer (without incubation) with ethanolamine to serve as added
192	activity measurements.
193	
194	2.3 Picophytoplankton abundance using flow-cytometry
195	Triplicates water samples (1.5 ml) were collected from each sampling depth, put in cryo-vials
196	(Nunc), and supplemented with 7.5 μ l 25% glutaraldehyde (Sigma). Vials were incubated in
197	the dark for 10 min, flash-frozen in liquid nitrogen, and stored in -80 °C freezer. Before
198	analysis, samples were thawed in the dark at room temperature. Each sample was run twice
199	on a BD Canto II flow-cytometer with 2 μm diameter fluorescent beads (Polysciences,
200	Warminster, PA, USA) as a size and fluorescence standard. In the first run three types of
201	phytoplankton cells were identified based on their natural auto-fluorescence:
202	Prochlorococcus, Synechococcus and picoeukaryotes. Cells were differentiated based on cell
203	chlorophyll (Ex482nm/Em676nm, PerCP channel) and phycoerythrin fluorescence
204	(Ex564nm/Em574nm), and by the size of cell (forward scatter). Before the second FCM, run
205	samples were stained with SYBR Green I (Molecular Probes/ ThermoFisher) to enable
206	counting followed by detection at Ex494nm/Em520nm (FITC channel). This provided counts
207	of the total bacterial population (phytoplankton + heterotrophic bacteria and archaea) as well

as a distinction between cells with High or Low DNA content (not shown). Data were
processed using FlowJo software. Flow rates were determined several times during each
running session by weighing tubes with double-distilled water, and counts of the standard
beads were used to verify a consistent flow rate.

212

213 2.4 Algal pigment markers

Eight litters of seawater were collected from all photic sample depths and one from a dark 214 depth (depth varies between cruises). Water was filtered onto GF/F filters (0.7 µm nominal 215 216 pore size, 47mm diameter, Waters) using a peristaltic pump until either all 8 L were filtered 217 or the filter became blocked, in which case the volume filtered was recorded. Filters were placed in cryo-vials and flash frozen in liquid nitrogen until they could be stored in a -80 °C 218 219 freezer. Pigments were extracted in 1ml 100% methanol for 3 h at room temperature and 220 clarified using syringe filters (Acrodisc CR, 13 mm, 0.2 µm PTFE membranes, Pall Life 221 Sciences). Total chlorophyll was measured spectrophotometrically using a NanoDrop 2000c 222 (Thermo Sciences) at 632, 652, 665 and 695 nm, and the concentration of chlorophyll a was calculated (Ritchie 2008). Ultra high-pressure Liquid Chromatography (UPLC) was 223 224 performed on an ACOUITY UPLC system (Waters) equipped with a photodiode array 225 detector. A C8 column (1.7 µm particle size, 2.1 mm internal diameter, 50 mm column 226 length, ACQUITY UPLC BEH, 186002877) was used. The chromatography method was adapted for UPLC from the LOV method (Hooker et al., 2005). Samples were preheated to 227 228 30 °C and column to 50 °C before each run. Running buffers were a 70:30 mixture of 229 methanol and 0.5M ammonium acetate (buffer A) and 100% methanol (buffer B). The 230 program consisted of an isocratic run using a 80:20 mixture of buffers A:B for 2min, 231 followed by a linear gradient to 50:50 for 7 minutes and an increase to 100% solvent B. The 232 flow rate was 0.5ml/min. Pigment standards from DHI (Denmark) were used to identify the

UPLC peaks (chlorophyll a, divinyl-chlorophyll a, chlorophyll b, chlorophyll c2, zeaxanthin,
beta-carotene, diatoxanthin, fucoxanthin, peridinin, 19'- butanoyloxyfucoxanthin and 19'hexanoyloxyfucoxanthin). Due to potential degradation of the pigment standards, we present
the total chlorophyll measured spectrophotometrically and the pigment ratios within each
UPLC run.

238

239 **3 Results**

240 **3.1** Physical and chemical properties of the water column

241 Between January 2016 and December 2019, a clear pattern was observed in satellite-derived 242 sea surface temperature at the THEMO-2 location (Figure 1B), which was mirrored during the monthly cruise measurements (February 2018-January 2019, Figure 2A). The measured 243 244 *in-situ* sea surface temperatures increased by $> 11^{\circ}$ C from the winter minimum of $\sim 17.9 \circ$ C to 245 the summer maximum of 29.1 °C (Figure 2A, Supplementary Figure S3, Table 1). Sea 246 surface salinity also increased from a winter minimum of ~39.3 psu to a summer maximum 247 of 39.8 psu (Figure 2B, Supplementary Figure S3). Both temperature and salinity minima were higher than the climatological minima (~15.2 °C and 38.9 psu, measured between 2002-248 249 2020 (Herut et al., 2020), suggesting that the sampling period represents a relatively warm 250 and salty year. The temporal changes in sea surface temperature and salinity led to 251 differences in the water density profiles (Figure S1), indicative of stratification of the upper water layer between May and December 2018 (Mixed Layer Depth = 15-49 m) and mixed 252 253 between February-April 2018 and January 2019 (MLD not determined, see materials and 254 methods) (Table 1). Inorganic nutrient (NO₃+NO₂) concentrations began to increase in the 255 mixed layer already during November, suggesting that the stratification had started to erode 256 earlier than observed based on the density profiles, possibly due to short-term mixing events 257 (e.g. (Gunn et al., 2020).

- 258 Dissolved oxygen concentrations were at or above 100% saturation throughout the whole
- 259 photic layer (0-200 m), ranging from ~180 μ mol L⁻¹ to ~240 μ mol L⁻¹ (Figure 2C). Where,
- soluble reactive phosphorus (SRP) concentrations at the surface water were at or close to the
- 261 limit of detection ~0.006 μ M, (~6 nM) throughout the year, whereas nitrate+nitrite (NOx)
- were close to the detection limits from August to November (limit of detection 0.013 µM) but
- reached 0.3-0.5 µM during the mixed period (Table 1, and see Ben Ezra et al., 2021 for more
- 264 information).

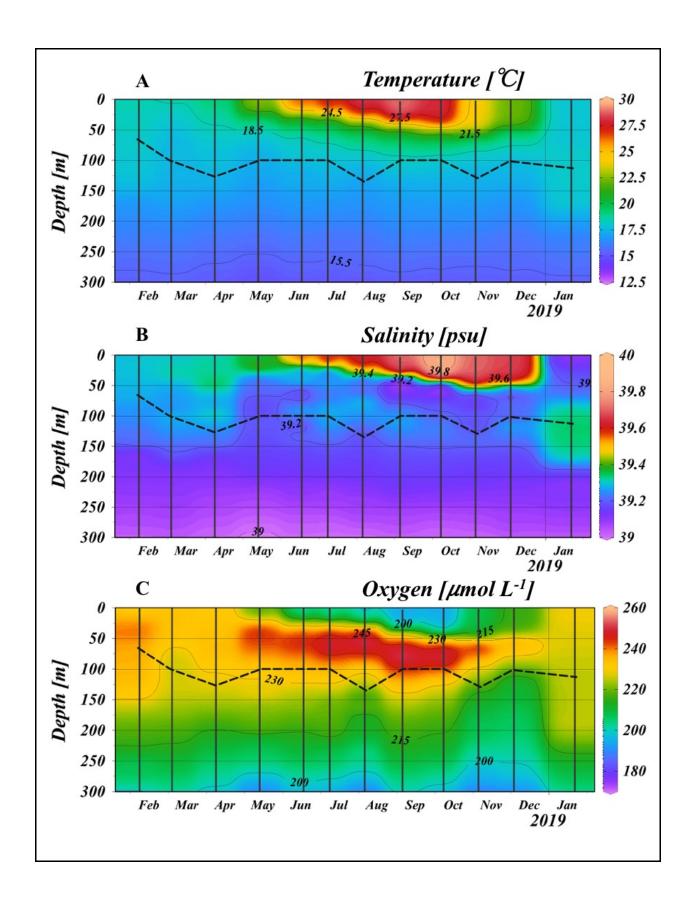


Figure 2: Temperature (A), Salinity (B) and oxygen concentrations (C) at THEMO2 station measured monthly between February 2018 to January 2019. The dashed line represents the depth of the Deep Chlorophyll Maximum (DCM). See Supplementary Figure S3 for the full depth profiles to ~1,400m.

265

Table 1: Median and range of major measured oceanographic parameters during the mixed

and stratified periods. BDL – Below detection limit. ND – not determined (see materials and

268 methods and Supplementary Figure S1 for details). Depth-integrated values are from 0-200m.

Parameter	Mixed period (Jan-Apr)	Thermally Stratified period (May-Dec)
SST (°C)	18.6 (17.9-19.6)	24.2 (22.1-29.1)
Mixed layer depth (m)	ND	26.5 (15-49)
DCM depth (m)	105 (67-125)	100 (100-135)
Surface (10 m) PO ₄ [µM]	BDL (≤0.006)	BDL (≤0.006)
Surface (10 m) NOx [µM]	0.3 (0.3-0.6)	0.1 (0.03-0.5)
Surface Si [µM]	0.8 (0.8-1)	0.7 (0.6-1)
Chl $a \text{ [mg m}^{-2}\text{]}$	14.6 (8.6-17.8)	8.8 (5.4-24.6)
<i>Synechococcus</i> [cells m ⁻²]	$1 \times 10^{12} (5.8 \times 10^{11} - 1.6 \times 10^{12})$	5x10 ¹¹ (5x10 ¹⁰ -8.5x10 ¹¹)
Prochlorococcus [cells m ⁻²]	$7.1 \times 10^{11} (2.5 \times 10^{11} - 2.2 \times 10^{12})$	$2.3x10^{12} (1x10^{12} - 3.2x10^{12})$
Pico-eukaryotes [cells m ⁻²]	1.7x10 ¹¹ (BDL-7.67x10 ¹¹)	8.6x10 ⁹ (7.2x10 ⁷ -2.3x10 ¹¹)
Heterotrophic prokaryotes [cells m ⁻²]	$6.6 \times 10^{13} (6 \times 10^{13} - 1.03 \times 10^{14})$	$5.4 \text{ x} 10^{13} (3.1 \text{ x} 10^{13} 1.1 \text{ x} 10^{14})$
PP [mgC m ⁻² d ⁻¹]	113 (69-142)	60 (48-111)
BP [mgC m ⁻² d ⁻¹]	54 (32-71)	18 (7-27)

270 **3.2** Chlorophyll a, Primary Productivity and Bacterial Productivity

- 271 While satellite-derived sea surface chlorophyll *a* showed a clear seasonal cycle over three
- 272 years, with lower levels during the stratified period (Figure 1B), the entire photic zone
- 273 displayed more complex evolution in depth and chlorophyll concentrations (Figure 3A, B). A
- 274 prominent Deep Chlorophyll Maximum (DCM) was observed year-round, ranging from
- depths of 60 m (February 2019) to 120 m (April 2019). The concentration of total
- 276 chlorophyll a at the DCM was always higher than that measured at the surface, and ranged
- 277 between ~0.15 μ g L⁻¹ during February-July, decreasing to ~0.07 μ g L⁻¹ during August-
- 278 November. These values are within the range or somewhat lower than those measured in
- other studies in the EMS (~0.09-0.42 μ g L⁻¹) (Christaki et al., 2001; Yacobi et al., 1995), and
- are ~50-80% lower than at BATS (Steinberg et al., 2001). Due primarily to the changes in
- chlorophyll a at the DCM, the integrated chlorophyll did not follow the same temporal
- variability as observed with the sea-surface satellites measurements (Figure 1B).

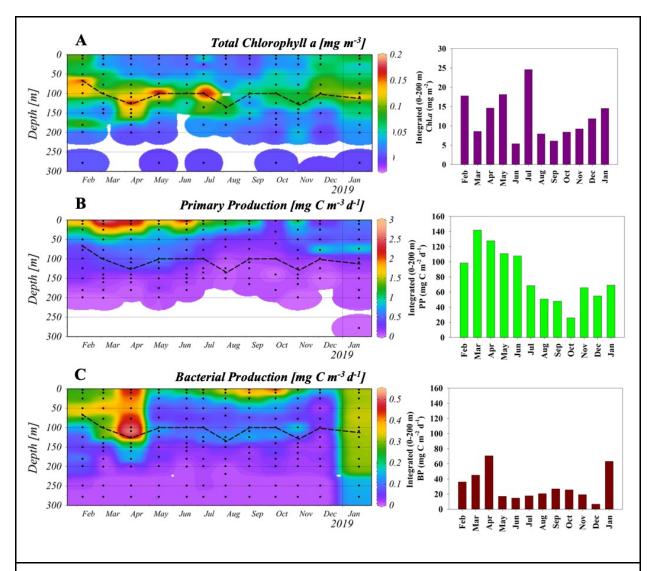


Figure 3: Seasonal changes in depth-resolved (left panels) and depth-integrated (right panels) chlorophyll *a* (A), primary productivity (B) and bacterial productivity (C). Black dots in panels A-C represent sampling points from each cruise. See Supplementary Figure S4 for the full depth profile of BP to ~1,400m.

284

Primary productivity (PP) was highest at the surface most of the year (1-3 mg C m⁻³ d⁻¹),

286 declining with depth, with no observable maximum during most of the year at the DCM

287 $(0.08-0.47 \text{ mg C m}^{-3} \text{ d}^{-1}, \text{ Figure 3B})$. This is consistent with the chlorophyll maximum in

- 288 ultra-oligotrophic oceans being decoupled from the PP maximum (Lazzari et al., 2012). An
- exception to this decoupling was observed during December 2019 and January 2020, where a

peak of PP was observed at around 75 m (0.93-0.96 mg C m⁻³ d⁻¹), slightly above the DCM 290 291 (Figure 3B). The observed values at the surface are within the range or somewhat lower than 292 previously measured values in the EMS (typically around $\sim 2 \text{ mg C m}^{-3} \text{ day}^{-1}$, but with values as high as ~18 mg C m⁻³ day⁻¹ recorded), (Hazan et al., 2018; Rahav et al., 2013). The 293 integrated PP values ranged between 69-142 mg C m⁻² d⁻¹ during the mixing period, and from 294 295 48 to 113 mg C m⁻² d⁻¹ during the stratified months (Figure 3B). These values are within the 296 ranges observed in other studies of this region (reviewed in (Berman-Frank and Rahav, 2012; 297 Siokou-Frangou et al., 2010) and discussed below). Our measurements result in an annual PP of \sim 32 gC m⁻², which is lower by \sim 50% than most estimates from the EMS (Boldrin et al., 298 299 2002; Psarra et al., 2000). These annual estimates highlight the ultra-oligotrophic 300 characteristics of the easternmost Levantine Basin. During most of the year BP was highest at the surface (~0.2-0.5 mg C m⁻³ d⁻¹, Figure 3C, 301 302 Supplementary Figure S4), within the range of previous observations at this area (Rahav et 303 al., 2019; Tanaka et al., 2007; Van Wambeke et al., 2000). However, BP differed from PP in 304 several aspects. First, a significant increase in BP was observed during April, with the highest 305 rate observed at ~110 m where the DCM was detected. Indeed, a smaller secondary peak in 306 BP was observed at a depth corresponding to the DCM throughout the year (Figure 3C). 307 Second, surface BP increased during summer from a minimum in May to a maximum in 308 September-October, before decreasing again. Third, during January 2019 a large increase was observed in BP, which was more-or-less homogenously distributed throughout the water 309 column down to ~200 m. Depth-integrated BP ranged from a maximum of ~70 mg C m⁻² d⁻¹ 310 during April 2018 and January 2019 to a minimum of ~20 mg C m⁻² d⁻¹ during May-311 December 2018, within the range of previous measurements $\sim 10-45$ mg C m⁻² day⁻¹ (e.g. 312 (Christaki et al., 2011; Robarts et al., 1996; Van Wambeke et al., 2000). The integrated PP to 313

BP ratio also differed between the mixed and stratified periods. Where, during the mixed
season the PP:BP ratio was ~3:1, while during the stratified period it reached ~5:1.

316

317 **3.3** Phytoplankton abundance and specific functional groups

318 Picocyanobacteria (Prochlorococcus and Synechococcus) were the most abundant

319 picophytoplankton cells throughout the year (Figure 4, Supplementary Figure S5).

320 *Prochlorococcus* was more abundant during the stratified period, and predominated below

321 the mixed layer (Figure 4A). Based on the ratio of divinyll chlorophyll a to total chlorophyll a

322 (Figure 5A), *Prochlorococcus* contributed up to ~45% of the total phytoplankton biomass at

323 the DCM during late fall (Figure 5A). *Synechococcus* was more abundant during the mixed

324 period and in the surface waters (Figure 4B). Pico-eukaryotes were present at up to 10⁴ cells

325 ml⁻¹ during February-June (Figure 4C), but there was no clear group-specific signal in the

326 tested photosynthetic pigments during this period (Figure 5B-D), and thus the specific pico-

327 eukaryotic groups present at this time could not be identified. In contrast, the presence of 19-

328 hexanoyloxyfucoxanthin and 19-butanoyloxyfucoxanthin (19-Hex+19-But), primarily during

329 August to November (Figure 5B), suggests the presence of haptophytes, and this was

330 corroborated with microscopic identification of coccolithophores (see a more detailed

discussion in Keuter et al, *in prep*). Finally, peridinin and fucoxanthin peaked during January,

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332 suggesting the presence of dinoflagellates and diatoms, respectively (Figure 5C,D).
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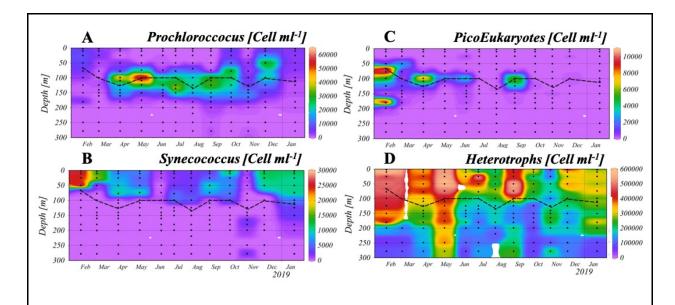


Figure 4: Annual dynamics of monthly measured picophytoplankton and heterotrophic bacterial abundance: *Prochlorococcus* (A), *Synechococcus* (B), PicoEukaryotes (C) and heterotrophic bacteria (D) at the THEMO2 station. Black dots represent sample point from each cruise. No data were available from the March cruise for total microbial counts. Note the differences in the color scale for each plot. Dashed line represents the DCM. See Supplementary Figure S5 for the full depth profiles to ~1,400m.

333

Throughout the year, total prokaryotic microbial counts (Sybr-stained cells, thus including cyanobacteria, heterotrophic bacteria and archaea) ranged between 3×10^5 to 6×10^5 cells ml⁻¹, with higher values observed during February and May (Figure 3SB). Heterotrophic bacteria were much more abundant than the combined phototrophs, the latter forming 2.4-8.5% of the total Sybr-stained cell counts (Figure 4D). Below the photic layer, cell counts were typically lower, ranging from 1.9×10^4 - 4.5×10^5 cells ml⁻¹ (Supplementary Figure S5).

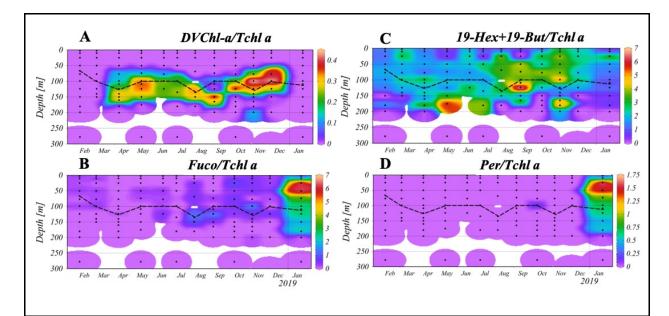


Figure 5: Monthly changes in the ratios ofmajor accessory pigments to Total chlorophyll *a*. A) divinyl chlorophyll *a* (DVChl-a). B) Combined 19-hexanoyloxyfucoxanthin and 19butanoyloxyfucoxanthin (19-Hex+19-But). C) Fucoxanthin (Fuco). D) Peridinin. Black dots represent sampling depths. Dashed line represents the DCM.

340

341 4 Discussion

342 4.1 How oligotrophic is the EMS compared to other oligotrophic oceans?

343 The EMS has previously been suggested to be one of the most oligotrophic marine systems 344 on Earth, including a claim for the "deepest Secchi-depth world record" - 53 m during 345 summertime (Berman et al., 1984). A comparison of published values of PP, including those 346 measured in this study, supports this notion, with the median integrated PP being ~66% lower 347 than Bermuda, the Red Sea and the Western Mediterranean Sea (WMS), and ~80% lower 348 than station ALOHA (HOT) (Figure 6A). Our estimates of PP during the SoMMoS cruises are among the lowest in the EMS (Figure 6A), yielding an annual PP of \sim 32-39 gC m⁻² y⁻¹, 349 350 although similar values have previously been reported (Dugdale and Wilkerson, 1988). The 351 annual PP during the SoMMoS cruises is approximately half of that reported above the continental slope of the Cretan Sea, ~59 gC m⁻² y⁻¹, (Psarra et al., 2000), as well as in other 352

more western locations in the EMS, ~62 gC m⁻² y⁻¹ (Boldrin et al., 2002). It should be noted, 353 354 though, that our incubation approach do not precisely mimic the *in-situ* illumination and 355 temperature at the time of sampling, which, based on our previous work, may underestimate 356 the actual PP rates by up to $\sim 20\%$ (Figure S2). Under these circumstances, the annual PP we report may reach \sim 39 gC m⁻² y⁻¹, approximately \sim 33% lower than previously reported from 357 358 the EMS (Psarra et al., 2000; Boldrin et al., 2002). Moreover, comparison between different studies that used similar, yet not identical, methodology for PP estimates, should be done 359 360 with care. For example, in the comparison presented in Figure 6A many of the studies 361 measured PP from dawn to dusk while in this study we used longer incubations which account for the whole day (net PP). These differences may also, partly, account for the 362 363 changes between the different oceanic locations, yet even with these caveats taken into 364 account the PP in the EMS is still lower than other oceanic regions (Figure 6A). 365 The values of integrated chlorophyll (Table 1, Figure 2A) and phytoplankton cell counts are 366 also somewhat lower than previously-published measurements from the same region (Figure 367 4, Figure 7) (Berman et al., 1984; Robarts et al., 1996). Both the temperature and salinity minima during the SoMMoS cruise series were higher than the 2002-2020 climatological 368 369 average (Herut et al., 2020), and thus we currently cannot determine whether the lower PP 370 values measured during this annual cruise series are due to inter-annual variability, or 371 whether these are due to the possible underestimation of the PP values discussed above 372 (section 2.2). Nevertheless, it is clear that the EMS is one of the most oligotrophic locations 373 on Earth, as defined using primary production, and that key seasonal dynamics are similar 374 between studies performed a decade apart (compare Figures 7D, E). 375 In addition to low PP, the EMS exhibits low sinking fluxes of particulate organic carbon 376 (POC). The POC flux measured from the bottom of the photic layer (180 m) at the DeepLev mooring station, which is in the same region as the THEMO-2 site (Figure 1A, ~10Km north 377

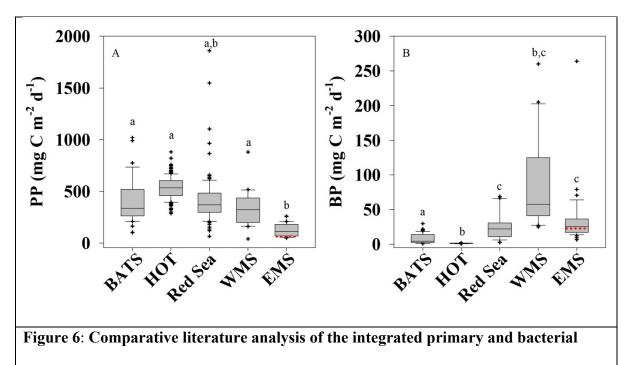
378	of our study site, Lat: 32.820 N; 34.380 E), is ~0.2 gC m ⁻² during the summer months (May-
379	September 2018) and ~0.8 gC m ⁻² per year (Alkalay et al., 2020). This POC flux to the
380	aphotic layer is less than 2% of the PP measured during summertime (~12 gC m ⁻² in 5
381	months) or 2.5% from the annual PP rates (~32 gC m ⁻² y ⁻¹). The fraction of PP exported as
382	particulate matter in the EMS is therefore lower than that reported in other oligotrophic
383	systems; i.e., HOT and BATS stations ~5% (Karl and Church, 2014; Steinberg et al., 2001).
384	This low contribution provides an additional biogeochemical implication of the extreme
385	oligotrophic conditions of the EMS. It also points to that the dissolved organic carbon
386	fraction derived from PP is rapidly recycled in the photic layer by heterotrophic/mixotrophic
387	bacteria as often observed in LNLC environments, as reviewed in (Santinelli, 2015). We
388	note, however, that estimates of export production based on nitrate+nitrate loss from the
389	photic zone, calculated from the same data, are significantly higher (172 mmol N m ⁻² year ⁻¹ ,
390	corresponding to ~13.6g C m ⁻² year ⁻¹ , presented in a companion paper, Ben Ezra et al.,
391	2021). The reason for this discrepancy is unclear, but may be due to over-estimation of the
392	export production rates (e.g. if part of the nitrate+nitrite was taken up by heterotrophic
393	organisms) and/or underestimation of the sinking carbon flux from sediment traps, and/or
394	non-Redfield C:N conversion factors that may prevail in the EMS. Moreover, such an
395	underestimation may be due to sinking flux from organisms too large to be caught in the
396	sediment traps such as fish or jellyfish (Edelist et al., 2020).
397	In contrast to PP, integrated BP in the EMS was higher than estimates at BATS and HOT,
398	and similar to the Red Sea (Figure 6B, which compares only measurements obtained using
399	the ³ H-leucine incorporation method). Yet, values obtained in the South Pacific Gyre, 43-61
400	mg C m ⁻² d ⁻¹ , (Berman-Frank et al., 2016) do come close to our measurements in the EMS.
401	Thus, differences in PP between oligotrophic regions are not necessarily coupled to BP
402	measurements. Previous studies, including from HOT and BATS, have shown that BP and PP

403 are decoupled across multiple time-scales (e.g hours-months) (Viviani and Church, 2017). 404 Similarly, PP and BP were uncoupled in the cyclonic Rhodes Gyre and the anti-cyclonic 405 Cyprus Eddy of the EMS (Rahav et al., 2013) and also in our study when BP increased in 406 surface waters between June and November, while PP rates declined (Figure 3). One 407 potential reason for this discrepancy is that PP is often measured on filtered samples 408 (particulate PP), and thus does not include the fraction of dissolved organic carbon (DOC) 409 fixed through PP that is released from the cells due to exudation or lysis (Viviani et al., 410 2015).

411 When phytoplankton have sufficient carbon and light for photosynthesis yet nutrients such as 412 P and N are limiting, photosynthesis is uncoupled from growth and high concentrations of 413 DOC are often released to the environment (Berman-Frank and Dubinsky, 1999). Previous 414 studies from natural samples and from cultures of the numerically-dominant phytoplankton in 415 the EMS, *Prochlorococcus* and *Synechococcus*, suggest a potentially high but variable 416 fraction of released DOC (typically 20-70% of PP, with one study suggesting >90% in lab 417 cultures, (Roth-Rosenberg et al., 2020; Viviani et al., 2015). It is currently unclear whether 418 autochthonous production (that is, local PP) is sufficient to support the growth requirements 419 of heterotrophic bacteria, given what is known about the growth efficiency of bacteria. A 420 study from the Cretan Sea suggesting that this is possible if the release of DOC from 421 phytoplankton exceeds 40% of PP (Anderson and Turley, 2003). 422 The EMS is affected by coastal intrusions of relatively chlorophyll-rich waters (Efrati et al., 423 2013), which could contribute also additional DOC, yet the importance of these processes in 424 determining BP and PP requires additional studies, as they occur on temporal and spatial 425 scales not captured by the SoMMoS cruise series. Such studies are also needed to understand 426 the relative contribution of dissolved compared to particulate carbon to export processes in 427 oligotrophic regions.

An additional explanation for the uncoupling between PP and BP was previously suggested 428 429 following an *in-situ* phosphorus addition experiment at an anti-cyclonic eddy in the EMS, 430 where orthophosphate addition led to chlorophyll a decrease while heterotrophic biomass and 431 activity increased (Thingstad et al., 2005). The authors postulated two possible scenarios to 432 explain their observation; 1) fast growing bacteria out-competed phytoplankton for the added 433 phosphorus. Then, the accumulated heterotrophic biomass was quickly channeled toward larger consumers (the 'bypass hypothesis'). 2) luxury uptake of P, mainly by heterotrophic 434 435 bacteria (and to a lesser extent by small-size picophytoplankton), formed a phosphorus-rich 436 diet for grazers, resulting in increased egg production (the 'tunneling hypothesis'). 437 Additionally, a scenario where an increase in temperature drives increased growth by 438 heterotrophic bacteria (Luna et al., 2012), resulting in a draw-down of inorganic nutrients, 439 could explain the increased BP in surface waters during summer, when no increase in PP was 440 observed (Figure 3).

441



productivity (PP and BP, respectively) in the EMS and other well-studied oligotrophic

locations (Bianchi et al., 1999; Bonnet et al., 2011; Casotti et al., 2003; Christaki et al., 2011; Decembrini et al., 2009; Fernández et al., 1994; Gasol et al., 1998; Gaudy et al., 2003; Hazan et al., 2018; Ignatiades et al., 2002; Lemée et al., 2002; Lohrenz et al., 1988; Morán and Estrada, 2001; Moutin and Raimbault, 2002; Pedrós-Alió et al., 1999; Rahav et al., 2019; Eyal Rahav et al., 2013; Robarts et al., 1996; Siokou-Frangou et al., 2002; Van Wambeke et al., 2004; Vidussi et al., 2000; Wambeke et al., 2002; Zervoudaki et al., 2007). Box-Whisker plots show the interquartile range (25th–75th percentile) of the data set. The horizontal line within the box represents the median value. The red dashed line in the EMS boxes shows the median values for the SoMMoS cruises described here. The letters above the box-plots represent significant differences (ANOVA, p < 0.05) for mean values between sampling sites. The BP compilation includes only measurements obtained using the ³H-leucine incorporation method. Note the different Y axis.

442

443 4.2 Who are the main phytoplankton in the EMS?

444 During the thermally mixed period, Synechococcus were more abundant than 445 Prochlorococcus at the upper ~100 m (Figures 4). This period was characterized by 446 measurable N at the surface while P concentrations were still extremely low – possibly 447 causing P limitation (December-July, Table 1, Ben Ezra et al., 2021). In contrast, the 448 extreme surface nutrient scarcity during the stratified period, when both N and P were below the limit of detection and were potentially co-limiting (Ben Ezra et al., 2021), likely 449 450 contributed to the dominance of the small-size cyanobacterium Prochlorococcus over the 451 somewhat larger-cell Synechococcus (Figures 4). It is generally accepted that, as the nutrient 452 concentrations decrease (i.e., the water becomes more oligotrophic), so does the average size 453 of phytoplankton (Margalef and Kinne, 1997). Additionally, while most Synechococcus 454 genomes encode the genes required for nitrate uptake and utilization, this trait is found only

in a subset of the *Prochlroococcus* genomes (Berube et al., 2018), and single-cell analysis 455 456 suggests that more Synechococcus cells in nature utilize nitrate compared to Prochlorococcus 457 (Berthelot et al., 2019). Thus, the type of nutrient limitation, P or N+P, (Krom et al., 1991; 458 Thingstad et al., 2005; Zohary et al., 2005) may also affect the temporal dynamics of these 459 two clades in the water column. 460 The abundances of these two cyanobacterial groups also differed with depth. Synechococcus was more abundant in the surface waters than Prochlorococcus (Fig 4). This was especially 461 462 evident from May to September when Prochlorococcus cells could not be identified in the 463 surface layer using either flow cytometry or pigment analysis (Figures 4 and 5). While the low Prochlorococcus abundance at the surface could be partly due to the limit of detection of 464 465 the flow cytometer used, it is supported by the lack of observed DVChl-a (Figure 5a) and by previous genetic analyses (Rosenberg et al., 2020). It has been suggested that 466 467 Prochlorococcus' unique photosynthetic pigments (DVChl-a and DVChl-b) provide a 468 competitive advantage for these organisms at deeper layers of the water column, leading to a 469 niche separation between these two pico-cyanobacteria (Moore et al., 1995). The observed 470 niche (depth) separation between Prochlorococcus and Synechococcus has been previously 471 observed including in the first description of Prochlorococcus from the North Atlantic and 472 North Pacific (Chisholm et al., 1988), and the Eastern South Pacific (BIOSOPE cruise, 473 (Huang et al., 2015)). However, this is not a universal observation. At the HOT station in the 474 North Pacific, Prochlorococcus and Synechococcus often show similar depth distributions, 475 both being more abundant at the surface, with Prochlorococcus more abundant numerically 476 and in terms of biomass (van den Engh et al., 2017). Similarly, in the Indian Ocean during 477 May-June 2003 Prochlorococcus dominated the surface waters whereas Synechococcus were 478 much less abundant (Huang et al., 2015). Indeed, a global analysis of >35,000 quantitative 479 measurements of Prochlorococcus and Synechococcus found that Prochlorococcus were

often more abundant than Synechococcus at high PAR and temperature values representative 480 481 of surface waters in tropical waters (Flombaum et al., 2013). Thus, the reasons why 482 Prochlorococcus and Synechococcus partitioned by both depth and season in the EMS, and 483 why this is not necessarily observed elsewhere, remain unclear. Other factors probably also 484 impact distribution, as was suggested for surface populations of Prochlorococcus that 485 appeared to be negatively affected by atmospheric deposition through biological (e.g., airborne viruses or other biological agents that infect the cells) and chemical (e.g., trace-486 487 metals toxicity) processes (Rahav et al., 2020). 488 While the pico-cyanobacteria Prochlorococcus and Synechococcus were the most numerically abundant phytoplankton at THEMO-2, when biomass is examined the 489 490 contribution of picoeukaryotes can be as high as that of the two pico-cyanobacteria, or even 491 higher (Supplementary Figure S6, and see below). A previous study of the diversity of pico-492 phytoplankton in the EMS showed that pico-eukaryotes were not very common in terms of 493 their DNA sequences or flow cytometry signals, but were often dominant in terms of RNA 494 sequences, one potential proxy for photosynthetic activity (Man-Aharonovich et al., 2010). In 495 that study, pico-eukaryotes were suggested to comprise up to 60% of the photosynthetic 496 picoplankton biomass in surface waters, and were mainly composed of Haptophytes 497 (primarily Prymnesiophytes) and Stramenopiles (Man-Aharonovich et al., 2010). In 498 agreement with these observations, the photosynthetic pigment 19'-hexanoyloxyfucoxanthin (19'-hex), which is considered (together with 19'-butanoyloxyfucoxanthin, or 19'-but) to be a 499 500 diagnostic pigment for prymnesiophytes, was found year-round in the water column, but was 501 more abundant relative to the total phytoplankton biomass (i.e. in relation to total 502 chlorophyll) during August-October (stratified period). Thus, both molecular evidence (Man-503 Aharonovich et al., 2010) and biochemical evidence (pigment analysis) point to the 504 importance of prymnesiophytes in the EMS. A companion study from the SoMMoS cruises

505 provides an overview of seasonal dynamic and impact of calcified haptophytes

506 (coccolithophores) during this yearly survey (Keuter et al, *in prep*).

507 Interestingly, we often observed 19'-hex around or below the DCM, including at depths

508 where light intensities are very low and photosynthesis may not provide enough carbon or

509 energy to support growth (Fig 5). Recent studies have unveiled that haptophytes, including

510 coccolithophores, are mixotrophic - capable of acquiring prey by phagotrophy or organic

511 compounds by osmotrophy in addition to photosynthesis (e.g. (Avrahami and Frada, 2020;

512 Godrijan et al., 2020; Tillmann, 1998). Indeed, haptophytes can contribute to large fractions

513 of total bacteriovory in marine settings (Frias-Lopez et al., 2009; Unrein et al., 2014). Further

analysis of the DNA samples collected on the SoMMoS cruises may help identify

515 mixotrophy within the phytoplankton clades found at the base of the photic layer (e.g.

516 (Yelton et al., 2016) and help illuminate the metabolic capacities required to support

517 phytoplankton growth in this environment.

518 Pigments associated with diatoms and dinoflagellates, which are often dominant

519 phytoplankton groups (Andersen et al., 1996), were observed at relatively high concentrations

only during January 2019 (Figure 5C, D). At this time DVChl-*a* measurement were below the

521 detection limit in the water column, suggesting a succession from pico-cyanobacteria to

522 larger eukaryotic organisms, which occurred relatively rapidly - over several weeks. This

523 occurred concomitantly with a drop in surface temperature (Figure 2A) and thus a break in

524 stratification, an increase in nitrate+nitrite (Ben Ezra et al., 2021), and a shift in

525 coccolithophore population towards r-selected species such as *Emiliania huxleyi* which are

526 indicative for a higher nutrient regime (Keuter et al, *in prep*). While chlorophyll-a

527 concentrations increased both at the surface (including by satellite sensing, Figure 1B) and at

528 the DCM (Figure 3A), no concomitant increase was measured in PP (Figure 3B). Yet, a

529 major increase was observed in BP (Figure 3C). As described in detail in the materials and

- 530 methods, our measurements of PP likely underestimate productivity at depth, and the
- 531 pigments were observed primarily below the mixed layer. It is possible that the
- 532 phytoplankton bloom actually occurred several days before our cruise, and that the pigments
- 533 observed are from stressed or dead phytoplankton. In this case, the major increase in BP
- observed at the time may represent heterotrophic bacteria utilizing dissolved and particulate
- 535 organic carbon produced during the bloom.

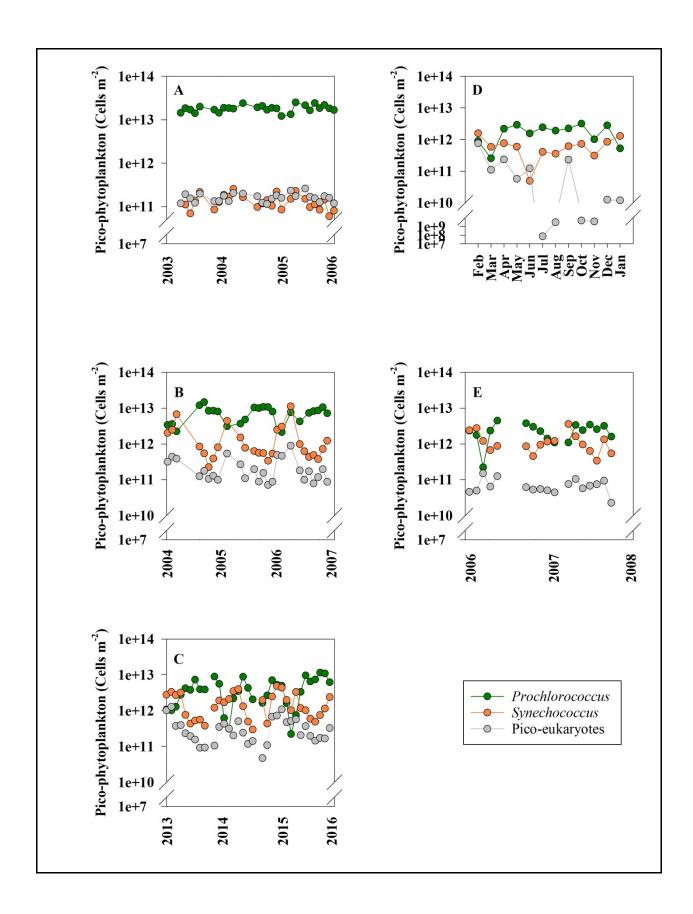


Figure 7: Integrated cell count time series of *Prochlorococcus* (green), *Synechococcus* (orange) and picoeukaryotes (gray) at stations HOT (A), BATS (B), the Northern Red Sea (C), and the EMS (D, E – this study and (Yogev et al., 2011) respectively). Abundance measurements were carried out using flow cytometry and are presented in a log scale. Data for stations HOT and BATS were compiled from (Malmstrom et al., 2010) and from the Northern Red Sea from the Eilat National Monitoring Program (Shaked and Genin, 2017).

536

537 4.3 Phytoplankton composition and seasonality in the EMS in comparison to HOT,

538 BATS and the Red Sea

539 The north Atlantic and Pacific gyres, the northern Red Sea and the EMS are all considered 540 Low-Nutrient Low-Chlorophyll marine systems, with phytoplankton numerically dominated by pico-cyanobacteria throughout most of the year. Nevertheless, there are differences in the 541 compositions and the seasonal dynamics of the different phytoplankton clades among these 542 543 LNLC location (Fig. 7, Sup Fig. 5). Specifically, BATS, the Red Sea and the EMS all show 544 seasonal succession patterns in the abundance of the main phytoplankton groups, although 545 there are subtle differences between these sites: i) The period when integrated Synechococcus 546 numbers are higher than Prochlorococcus is consistently longer in the Red Sea and EMS (3-4 547 months) compared to BATS (1-2 months); ii) Prochlorococcus reached higher absolute cell abundances during the stratified period at BATS ($\sim 10^{13}$ cells m⁻²) compared to the Northern 548 Red Sea or the EMS (~ 2.5×10^{11} cells m⁻²); iii) In terms of calculated biomass, the Red Sea is 549 550 dominated by pico-eukaryotes for much of the year, whereas in BATS and the EMS the 551 contribution of Prochlorococcus, Synechococcus and pico-eukaryotes to the total biomass is 552 similar, and the dominant groups change over the year (Supporting Fig S6). The absolute nutrient concentrations and the identity of the limiting nutrient(s) are different 553 554 between these locations. For example, BATS is typically considered P limited, whereas the

EMS is potentially P limited during winter and co-limited by N and P during summer 555 556 (Zohary et al., 2005). There may also be differences in the length of the mixed period, its 557 intensity, or the composition of the mixed water (e.g. deep water in the EMS have an N:P ratio of ~28:1, compared to the canonical 16:1 Redfield ratio (Redfield, 1934). Future work 558 559 may use the observed differences in phytoplankton dynamics as a sensitive "readout" of the 560 system, enabling a better understanding of the oceanographic processes underlying the differences between BATS, the Red Sea and the EMS. 561 562 In contrast to BATS, the Red Sea and the EMS, seasonality at HOT in is much less 563 pronounced (Figure 7). *Prochlorococcus* are about 2-orders of magnitude more abundant than Synechococcus and pico-eukaryotes, and are also the most dominant in terms of biomass (Sup 564 565 Fig. 5). The drivers of phytoplankton temporal dynamics at HOT are thus likely fundamentally different compared to BATS, the Red Sea and the EMS. It has been suggested 566 that a-periodic events such as mesoscale eddies, Fe deposition from winds, and biological 567 568 processes such as dinitrogen fixation, may be major drivers of phytoplankton dynamics at 569 HOT (Karl and Church, 2014) and less so in the EMS (Berman-Frank and Rahav, 2012; Yogev et al., 2011). 570

571

572 5 Closing remarks

573 The year of data presented here show that the EMS is one of the most oligotrophic regions in

the global oceans, as estimated based on PP (32 mg C m⁻² year ⁻¹), chlorophyll a (median ~ 10

575 mg m⁻²), and dominance of small-size picophytoplankton with overall low abundances

576 compared to other LNLC regions (in some cases ~2 orders of magnitude lower). In contrast,

577 BP in the EMS is higher than at HOT and BATS, and similar to the Red and Western

578 Mediterranean seas. This suggests that the system is overall heterotrophic, and that different

579 factors might limit phytoplankton and heterotrophic bacteria (e.g. (Thingstad et al., 2005).

Despite these observations, there is a clear seasonal cycle in the phytoplankton populations 580 581 and their activities, reminiscent of BATS and the Northern Red Sea (but not of HOT), 582 suggesting that mixing (and potentially other seasonal processes such as dust deposition) 583 drive similar processes across many, but not all, LNLC regions. 584 Our results also highlight important knowledge gaps regarding the EMS. For example, we do 585 not know to what extent the bacterial populations or their physiological traits (e.g. bacterial carbon demand or growth efficiency) differ seasonally or vertically in the EMS, as well as 586 587 compared to other LNLC locations. Detailed studies are also lacking of higher trophic levels 588 in this region (dominant micro-grazer species, diversity, biomass, grazing rates, etc.). These 589 are critical for understanding ecosystem dynamics including top-down vs. bottom-up 590 regulations etc. Biogeochemical characterization of the EMS will also benefit from more 591 explicit characterization of carbon-per-cell estimates, used to derive the biomass 592 contributions from flow-cytometry counts (i.e. Supplementary Figure S6) can vary depending 593 on taxonomy and physiological state (Kirchman, 2013). Future studies of these aspects, 594 together with other data emerging from the SoMMoS cruise series and the ocean observatories in the EMS, should provide the needed background for governments and other 595 596 stakeholders to employ science-based environmental policies in this rapidly changing region.

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- 607

608 Author Contributions

- MDK, YL, ER and DS initiated and designed study; TR, TBE, NB, DRR, OB, MDK, ER and
- 610 DS collected samples; TR, TBE, NB and AT analyzed samples with help from MDK, DA,
- 611 DRR and SG; TR, ER and DS wrote the manuscript with input from all co-authors.

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953