

We thank Dr Silvio Pantoja for his constructive criticism and valuable comments. In the following we address the points brought up, with editor comments in boldface and author responses in normal typeface.

**1) Reviewer 1. “The authors report the N:P but not with P relative to one. I think this is confusing.. For example, page 10002, lines 14- 16. N:P 6.35/1.10, 12.00:1.25. It would be better if this was written as: 5.77 and 9.6 respectively and details of the concentration be inserted into a table, for example.”**

**Response. “We prefer to report the N:P ratios as they are. That way, mesocosms with the same initial concentrations of N or P (center points) are easier to distinguish for the reader.”**

**I agree with reviewer. We are not interested in a mesocosm result by itself but inferences about ocean’s response, and for that it is easier for readers to have ratios relative to one since this article is based on deviations of canonical N:P. Please change accordingly.**

We have changed N:P ratios in the text as suggested (e.g. lines 249-253). However, we cannot change the mesocosm treatment ID’s as suggested. Due to our experimental design, there is more than one mesocosm in *varied N* and *varied P* with N:P ratios of 16:1 (see Table 1), which would be indistinguishable from each other if we only reported the N:P ratio relative to one.

**2) “9. The authors switch between N and P and NO<sub>3</sub> and PO<sub>4</sub>-. This needs to be fixed and made consistent throughout.”**

**Response: We made this consistent throughout the manuscript.”**

**I need to ask you to use proper nomenclature throughout the text. For general statements such as the ones in lines 47-52 of v2, it could be fine to use N or P, but regarding your results, you must refer to what you are experimenting with (nitrate or NO<sub>3</sub>-), but not N since nitrogen (N) involves a variety of chemical species you have not studied. One example is shown in conclusions in lines 328-329 of version 2 (“...ability of the plankton community to rapidly and intensively react to N availability. These results indicate that the ultimate limiting nutrient for phytoplankton production in our experiment was N.”).**

We have made the appropriate changes to the manuscript and figures.

**3) Line 402 v2. Replace “measured” with “detected”. It makes a more clear argument.**

We changed this as suggested.

**4) Reviewer 2. “Given the extensive measurements of the P pools (see e.g. Figure 10), it would have been useful to see an attempt at mass balance.**

**We addressed this issue by data presented in Fig 10. Mass balances were not subject of this manuscript but will be addressed in a follow up study. More extensive presentation of mass balances would be beyond the scope of this study.”**

**Reviewer is referring to mass balance as part of the method. It cannot be “beyond the scope of this study” rather, how balance or lack of balance affects your interpretation.**

We included a mass balance in version 2 of the submitted manuscript (see lines 287-291).

**5) Why nitrate plus nitrite increase during the first day?**

There is an increase because of our artificial nutrient addition between day 1 and 2.

**6) Please write nitrate or  $\text{NO}_3^-$  (using subscripts and superscripts) throughout text and figures. Same format for nitrite, silicic acid.**

We changed this accordingly.

We thank anonymous Referee #1 for his/her constructive criticism and valuable comments. In the following we address the points brought up, with referee comments in boldface and author responses in normal typeface.

**1. The motivation for this study, ocean deoxygenation due to climate change, and thus reduction of the N:P ratio, is not the only process that will alter then N:P ratio in the ocean. Addition of anthropogenic nitrogen (e.g. see Kim et al 2014), as the potential to perturb the system by adding N in excess of P, thus intensifying or expanding phosphate limited regions and thus force the system the other way. They authors should really alter their motivation to cover both sides of the story here.**

We agree that ocean deoxygenation is not the only process that will likely alter N:P ratios in the ocean. We clarified this in the introduction. However, in the eastern tropical North Atlantic (ETNA)  $N_2$  fixation is supposed to be the dominant process for the input of new N over the next decades to centuries (Duce et al. 2008) compared to the North Pacific Ocean (Kim et al. 2014). Moreover, it is debatable how much of the atmospheric anthropogenic nitrogen input is affecting the open ocean of the ETNA. If the input is mostly restricted to the coastal upwelling region, biological production would be fueled, export enhanced and degradation of organic matter at depth would increase. The oxygen inventory of the ETNA OMZ would shrink further, thereby enhancing N loss processes, leading to a decrease of N:P ratios in the water column. The fertilization with anthropogenic N would thus be compensated by a negative feedback cycle. In any case, the significance of atmospheric anthropogenic N inputs into the ETNA is unclear. The expansion of the oxygen minimum zone in the ETNA, on the other hand, has been ongoing for the past decades and is expected to continue in the future (Stramma et al. 2009). As the original goal and motivation for this experiment was to study changes in the N:P ratio due to ocean deoxygenation in the ETNA, we focused on the description of our experiment in this context.

**2. The ability of nitrogen fixation to modify primary production is likely to be small. If you multiple nitrogen fixation rates by a C:N ratio of ~ 6, then compare the carbon fixed by diazotrophs to total carbon fixed, it is quite small. Instead, the switch to DDAs will impact carbon export, which is the potentially important here.**

We agree and thank for this valuable input. The export of carbon might indeed be influenced by a shift of the diazotrophic community towards DDAs and included this point into the discussion. We interpret from our data that primary production might be affected by a shift within the diazotrophic community, especially in the oligotrophic open ocean of the ETNA, where diazotrophs are the dominant primary producers. We clarified this point in the manuscript.

**3. The author needs to be clear when they refer to P limitation. It is likely phosphate limitation and not phosphorus limitation considering the ability of organisms to access DOP. Please be more explicit about this in the manuscript.**

We agree with the referee and made the appropriate changes in the manuscript.

**4. You argue that there is low O<sub>2</sub> and N:P ratios (not levels as you have stated) below the MLD in mode water eddies. How typical do you think this is? Is this a wide spread occurrence and if so, how many mode water eddies exist in the Atlantic. It is not clear if this is just a local feature/unique phenomenon or widespread.**

We changed "(...) with an accompanied decrease in N:P levels (...)" to "(...) with an accompanied decrease in N:P ratios (...)" (p. 9994, line 5-6).

Mode water eddies with very low oxygen concentrations were only recently discovered in the ETNA. These eddies can last several month, transporting shelf water signals offshore (Karstensen et al. 2015). In the ETNA, 5 such events were observed between 2007 and 2012 (Karstensen et al. 2015). We believe that the effect of mode water eddies is not basin-wide but rather of local importance. We clarified this in the manuscript.

**5. What measures were taken to prevent contamination of the sample with either nutrients or trace metals? Was the nutrient or iron concentration at the oceanic sample collection site monitored between collection and use in the mesocosms?**

Containers for water transport were first rinsed with diluted HCL and several times with deionized water. Nutrients in all mesocosms were measured before nutrient manipulation.  $\text{NO}_3^-$  and  $\text{NO}_2^-$ ,  $\text{PO}_4^{3-}$  and  $\text{Si}(\text{OH})_4$  were all below the detection limit and far below the manipulation levels (see Fig. 2). We therefore conclude that no contamination with these nutrients occurred during water sampling, transport and mesocosm filling. This information was added to the manuscript.

**6. Nutrients: considering you are reporting the change in nutrient concentrations over time and between treatments, you should really report the precision of analysis as well as limits of detection in section 2.2. Also, the instrument and methods are described for DOP (section 2.4) but not nutrients. This needs to be fixed to be consistent. Again, what was the precision and limits of detection for DOP? This should be reported.**

The precision of analysis and detection limits for nutrient and DOP analysis was added. The paragraph describing the nutrient measurements was changed as follows: "Samples (10 mL) for dissolved inorganic nutrients ( $\text{NO}_3^-$ ,  $\text{NO}_2^-$ ,  $\text{PO}_4^{3-}$ ,  $\text{Si}(\text{OH})_4$ ) were taken daily from each mesocosm and measured directly using a QuAatro Autoanalyzer according to Grasshoff et al. (1999). The detection limit of nutrient analyses were  $0.01 \mu\text{mol L}^{-1}$  for  $\text{NO}_2^-$  and  $\text{PO}_4^{3-}$  and  $0.03 \mu\text{mol L}^{-1}$  for  $\text{NO}_3^-$ ."

**7. In section 2.7, what was the atom percent enrichment of the  $^{15}\text{N}_2$  water added to the incubations and the final atom percent enrichment at the start of the incubations? This should be reported here.**

The preparation of the  $^{15}\text{N}_2$ -enriched seawater was performed as described in Mohr et al. (2010). Degassed seawater was filled into evacuated gas-tight 3L Tedlar<sup>®</sup> bags without a headspace. Addition of  $^{15}\text{N}_2$  gas was (depending on the exact water volume in the Tedlar bag) around 10 ml  $^{15}\text{N}_2$  per 1 L seawater. Dissolution of the  $^{15}\text{N}_2$  gas was achieved by 'slapping' the bubble with a ruler. After complete dissolution of the added  $^{15}\text{N}_2$  gas ( $^{15}\text{N}_2$ -enriched seawater), an aliquot of the  $^{15}\text{N}_2$  enriched water was collected for each preparation of enriched seawater and stored in an Exetainer, the isotopic composition was measured by membrane-inlet mass spectrometry. The  $^{15}\text{N}_2$  concentration in the prepared batches of enriched water was determined to be  $250 \mu\text{mol L}^{-1}$ , which translates in an  $^{15}\text{N}$ -enrichment of about 2 % in the 4.5 L bottle incubations, when adding 100 mL enriched seawater (depending on temperature and salinity). These details were added to the methods section.

**8. Section 2.8: Model selection. Why did you use a model here? What was the goal? This needs to be stated. Was it necessary? Also, the model description is quite confusing. What is gam and gamm? What is Akaike Information Criterion? This section needs to be edited to clarify the goal of the model and perhaps reduce the detail here and refer to other manuscripts where this sort of analysis has been done already. As a follow up on this, on getting to the end of this manuscript, I do not believe the modelling component adds any real value to this manuscript. The study is data rich and there are plenty of interesting and important points to make without including the model.**

As both referees state that they don't see the added value of the model we introduce in the manuscript, we decided to remove it from our manuscript. Instead, we show the original transcript data obtained from the study (new Fig. 8) in order to avoid confusion and make the manuscript more intelligible to the reader (see also response to referee 2).

**9. The authors switch between N and P and  $\text{NO}_3^-$  and  $\text{PO}_4^{3-}$ . This needs to be fixed and made consistent throughout.**

We made this consistent throughout the manuscript.

**10. The authors report the N:P but not with P relative to one. I think this is confusing. For example, page 10002, lines 14- 16. N:P 6.35/1.10, 12.00:1.25. It would be better if this was written as: 5.77 and 9.6 respectively and details of the concentration be inserted into a table, for example.**

We prefer to report the N:P ratios as they are. That way, mesocosms with the same initial concentrations of N or P (center points) are easier to distinguish for the reader.

**11. I suggest changing the use of the word 'build up' to 'accumulation'. Also avoid using words like rise and drop, should be increase and decrease respectively.**

We agree and made the appropriate changes in the manuscript.

**12. Note that similar observations of the C:N:P ratio for POM were observed by Davis et al 2014 in GRL.**

Thank you for this valuable input! We will add the suggested reference and the observations of this study to our discussion.

**13. I suggest reducing the precision on ratios reported, e.g. change 38.8 to 39 and 21.9 to 22. The decimal places don't add value here.**

We removed the decimal places as suggested.

We thank anonymous Referee #2 for his/her constructive criticism and valuable comments. In the following we address the points brought up, with referee comments in boldface and author responses in normal typeface.

**In various places throughout the manuscript (e.g. Page 9993, line 26, Page 994, line 11 etc.) the authors refer to limitation when making inferences on the basis of ratios of available or supplied inorganic nutrients. Actually their own experiments suggest that this link is far from straight forward and I would encourage them to clarify where possible, maybe stating that the dissolved ratios indicate the ‘potential for one nutrient to becoming limiting before the other’ or sticking to the use of terms like ‘deficiency’, ‘deficit’, ‘excess’ etc. (see e.g. Page 9995, line 1).**

We agree and revised the manuscript as recommended. For example, the sentence on Page 9993, line 18 was changed as follows: „Nevertheless, the nitracline tends to be deeper than the phosphocline in the ETNA (Hauss et al., 2013; Sandel et al., 2015), which also points towards a deficiency of N over P in the euphotic zone.

**Experimental methods and statistical analysis need to be further described in places. In particular, although clear through consulting Table S1, the number of replicate mesocosms for individual treatments should be more clearly indicated to the reader, e.g. through stating in the text on Page 9997. Additionally, on Page 10001 the authors introduce a complex statistical model for the interpretation of the data without providing any justification for why this was required or chosen. Overall I was not sure why the statistical model was required as it appeared to largely be used just for the analysis of the nifH gene/transcript data and it wasn’t clear that it added much to the interpretation of this data. Additionally it wasn’t clear to me whether the analysis presented in Figure S1 was based on the GLM modelling performed or simple correlation analysis? Additionally, why is Figure S1 in supplementary rather than within main body of manuscript?**

We added information about treatment replicates to the Material and Method section as follows: „In the first experiment, the P supply was changed at constant N supply (*varied P*) in thirteen of the sixteen units, while in the second experiment the N supply was changed at constant P supply (*varied N*) in twelve of the sixteen units. Each of these nutrient treatments was replicated 3 times. In addition to this, “cornerpoints” were chosen, where both the N and P supply was changed. The „cornerpoints“ were not replicated.“

Due to the fact that both referees don’t see the added benefit in introducing a model to interpret our data, we decided to remove the model from our study and instead show the original transcript data (please also see the comment to referee 1).

**Although an entirely feasible explanation, I think any potential causal link between the accumulation/availability of DOP and enhanced N<sub>2</sub> fixation needs to be treated with caution on the basis of the data presented and experiment(s) performed. e.g. Page 10005, lines 11-20, an alternative interpretation might be that both the accumulation of DOP and the enhancement of N<sub>2</sub> fixation are occurring within the ‘varied P’ experiments independently simply as a result of the addition of inorganic P. The authors may argue that the time series of DOP, P, POP, N<sub>2</sub> fixation might argue against this (e.g. Figure 10), but given only 2 sampling time points for N<sub>2</sub> fixation I would argue this remains equivocal. I would suggest the authors may simply wish to acknowledge this potential caveat.**

We agree that the existence of only two sampling points for N<sub>2</sub> fixation has to be emphasized more when interpreting and discussing our data set. The text now reads:

“In our experiments a significant increase in N<sub>2</sub> fixation rates was only measured in *varied P*. In mesocosms with highest N<sub>2</sub> fixation rates, DIP was depleted after day 5 or 6 while POP increased until the end of the experiment. After DIP depletion, DOP concentrations declined, which indicates that DOP served as P source until the end of the

experiment. It has to be noted that N<sub>2</sub> fixation rates were only measured at the beginning and the end of our experiment and possible fluctuations over time cannot be accounted for. However, increasing diazotrophic transcript abundances of *Richelia intracellularis* in symbiosis with the diatom *Rhizosolenia* (Het I) were also detected over the course of the *variable P* experiment. While the diatom abundance was probably favored by replete amounts of silicate added at the beginning of the experiment, no increase in diatom-diazotroph associations (DDAs) was detected in the *varied N* experiment. Measured N<sub>2</sub> fixation rates and transcript abundances leads us to speculate that DDAs were favored in the *varied P* experiment, where diazotrophs in the mesocosms utilized DOP resources in order to supply P to themselves and/or their symbiont.”

Concerning the alternative explanation suggested by the referee, we do not believe that N<sub>2</sub> fixation was solely enhanced by the addition of inorganic P in our experiment, since N<sub>2</sub> fixation was not measured in all treatments in *varied P*, but only in those treatments where inorganic P was depleted after a couple of days and DOP served as an alternative P source.

**Given the extensive measurements of the P pools (see e.g. Figure 10), it would have been useful to see an attempt at mass balance.**

We addressed this issue by data presented in Fig 10. Mass balances were not subject of this manuscript but will be addressed in a follow up study. More extensive presentation of mass balances would be beyond the scope of this study.

**Page 9995, line 26: ‘. . .are regarded as key factors. . .’**

This was changed.

**Page 10004, line 11-15: This text does not appear to be fully consistent with the content of Figure 8? i.e. nifH Fil do not appear to be dominant for either experiment in this figure?**

Due to the removal of the model from the manuscript (see comment above), Fig. 8 and 9 were dismissed. A new Fig. 8 was added to show the original transcript data. We ensured that the text describes the figure appropriately.

**Page 10006, line 12 (and elsewhere): it is worth noting that the POC, PON, POP data reported will not just reflect that of ‘primary producers’ but actually will represent average values for the whole microbial community.**

We agree and made the appropriate changes. For example, the sentences on Page 10006 now reads: “There is a large difference between the supply ratio of inorganic nutrients and the PON:POP ratio of the plankton community in our study.”

**Page 10008, line 24: The authors could be more specific here. They are specifically discussing excess inorganic P. Related, the authors should use the more specific term DIP to refer to dissolved inorganic P when appropriate throughout (compare Page 9999 line 13 with Page 10008, line 24).**

As already stated in the response to referee 1, we made the use of DIP, PO<sub>4</sub><sup>3-</sup> and P consistent throughout the manuscript in order to avoid confusion.

**Page 10009, line 9: do the authors mean P\* here? i.e. DIP – DIN/16 or some similar definition c.f. Deutsch et al. 2007? If so I don’t think the term has been defined to this point in the manuscript.**

Yes, we refer to P\* as described by Deutsch et al, 2007. The term P\* has been introduced in the Introduction, Page 9993, line 16.

**Page 10009, line 29: ‘. . .locally prior to offshore transport.’**

This has been corrected.

**A number of the figure captions (and associated statistics) require work and/or better description and figures could be clarified in places:**

**Figure 1: please explain error bars (standard deviations? Standard errors?)**

The figure caption now reads: "Experimental design and initial nutrient supply conditions during *varied P* (blue circles) and *varied N* (red diamonds). "Cornerpoints" during *varied P* and *varied N* are depicted as grey circles and white diamonds, respectively. Error bars denote the standard deviation of replicated (n=3) treatments.

**Figure 2: shaded areas were a bit difficult to make out**

We decreased the transparency of the shaded areas to make them better visible.

**Figures 3, 4 & 6: error bars for data points need explanation, regression lines also need to be described in caption. Also were the fits model I or model II type regressions?**

The figure captions now read:

"Figure 3: Maximum POC, PON and POP build-up as a function of the initial supply of N, P and N/P. Maximum  $\delta$ POM is defined as peak POM concentration subtracted by the initial (day 1) POM concentration. Treatments in *varied P* are depicted as blue circles; treatments in *varied N* are depicted as red diamonds. Error bars denote the standard deviation of replicated (n=3) treatments. Regression lines (continuous lines) indicate linear correlations between the initial nutrient supply and POM accumulation."

"Figure 4: PON/POP stoichiometry during (A) the exponential growth phase and (B) the stationary growth phase of the experiment. The grey line visualizes the Redfield Ratio. The color code, symbols and lines are the same as in Fig. 3."

"Figure 6. Positive linear correlation between maximum DOP build-up (defined as peak DOP concentration subtracted by the initial DOP concentration) and initial P supply during *varied P* (blue circles) and *varied N* (red diamonds)."

The information about the type of regression was added to the Material and Methods section.

**Figure 7: error bars again need description. Additionally what statistical test was being used here?**

The figure caption now reads: "Figure 7: Mean  $N_2$  fixation rates measured on day 2 and day 8 of both experiments. Because of the high variance between replicates we omitted  $N_2$  fixation rates from un-replicated treatments. Asterisks indicate a significant difference between day 2 and day 8 (paired t-test). Error bars denote the standard deviation."

**Figure 10: error bars.**

The figure caption now reads: "Dynamics of  $PO_4^{3-}$ , POP and DOP in all mesocosms. Because of the high variance between replicates we omitted  $N_2$  fixation rates from un-replicated treatments. Error bars denote the standard deviation."

**Figure S1, caption and figure do not appear to match. Caption refers to 'a' and 'b' parts when there only appears to be one part in figure?**

Please see comment above.

**Changing nutrient stoichiometry affects phytoplankton production, DOP ~~accumulation~~<sup>build-up</sup> and dinitrogen fixation – a mesocosm experiment in the eastern tropical North Atlantic**

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1 **Abstract:**

2 Ocean deoxygenation due to climate change may alter redox-sensitive nutrient cycles in the marine  
3 environment. The productive eastern tropical North Atlantic (ETNA) upwelling region may be  
4 particularly affected when the relatively moderate oxygen minimum zone (OMZ) deoxygenates further  
5 and microbially-driven nitrogen (N) loss processes are promoted. Consequently, water masses with a  
6 low nitrogen to phosphorus (N:P) ratio could reach the euphotic layer, possibly influencing primary  
7 production in those waters. Previous mesocosm studies in the oligotrophic Atlantic Ocean identified  
8 nitrate availability as a controlling of primary production, while a possible co-limitation of nitrate and  
9 phosphate (P) could not be ruled out. To better understand the impact of changing N:P ratios on  
10 primary production and ~~on~~ N<sub>2</sub> fixation in the ETNA surface ocean, we conducted land-based  
11 mesocosm experiments with natural plankton communities and applied a broad range of N:P ratios  
12 (2.67 – 48). Silicic acidate was supplied at 15 μmol Lkg<sup>-1</sup> in all mesocosms. We monitored nutrient  
13 drawdown, ~~bloom formation~~, biomass accumulation ~~build-up~~ and diazotrophic feedback ~~nitrogen fixation~~  
14 in response to variable nutrient stoichiometry. Our results confirmed nitrate to be limiting to the key  
15 factor determining primary production. We found that excess phosphate ~~P~~ was channeled through  
16 particulate organic matter (POP) into the dissolved organic matter (DOP) pool. In mesocosms with low  
17 inorganic phosphate ~~P~~ availability, DOP was utilized while N<sub>2</sub> fixation increased, suggesting a link  
18 between those two processes. Interestingly this observation was most pronounced in mesocosms  
19 where inorganic N ~~nitrate~~ was still available, indicating that bioavailable N does not necessarily ~~has to~~  
20 ~~have a negative impact on~~ suppress N<sub>2</sub> fixation. We observed a shift from a mixed  
21 cyanobacteria/~~proteobacteria~~ dominated active diazotrophic community towards a diatom-  
22 diazotrophic ~~-diatom associations~~ symbionts of the *Richelia-Rhizosolenia* symbiosis. We hypothesize  
23 that a potential change in nutrient stoichiometry in the ETNA might lead to a general shift within the  
24 diazotrophic community, potentially ~~modifying primary productivity~~ influencing primary productivity and  
25 carbon export.

## 26 1 Introduction

27 Eastern boundary upwelling systems are characterized by cold, nutrient-rich water masses that are  
28 transported from intermediate water layers towards the surface. ~~The r~~Resulting extensive primary  
29 production forms the basis for high biomass development and a productive food ~~webchain~~  
30 (Pennington et al., 2006). At the same time, biological degradation at depth and weak interior  
31 ventilation cause permanently low oxygen concentrations in intermediate water masses (100 – 900 m,  
32 Karstensen et al., 2008). These low oxygen conditions support denitrification and anammox that  
33 remove bioavailable nitrogen (N) from the water column (e.g. Codispoti et al., 2001; Lam et al., 2009;  
34 Kalvelage et al., 2011). Oxygen minimum zones (OMZs) also influence the availability of inorganic  
35 phosph~~ateorus~~ (P), silic~~onate~~ (Si) and trace elements such as iron (Fe), which are released at the  
36 sediment-water interface under oxygen deficient conditions (Ingall and Jahnke, 1994; Hensen et al.,  
37 2006). Subsequently, the elemental stoichiometry of inorganic ~~nitrogen-to-phosphorus~~nutrients (N:P)  
38 in upwelled water masses is below the Redfield ratio of 16:1 (Redfield, 1958), which manifests itself as  
39 an excess of P (P\*) relative to N ( $P^* = PO_4^{3-} - NO_3^-/16$ , after Deutsch et al. (2007)).

40 In the Eastern Tropical North Atlantic (ETNA) nutrient concentrations ~~and stoichiometry~~ within the  
41 euphotic layer cover a wide range. Water masses in coastal regions feature low N:P ratios mainly as a  
42 result of benthic N-loss along with P leaching from the sediment (Trimmer and Nicholls, 2009;  
43 Jaeschke et al., 2010; Schafstall et al., 2010) suggesting an N limitation of primary production in OMZ-  
44 influenced surface waters (Deutsch et al. 2007). In the transition zone between coastal upwelling and  
45 open ocean, N:P ratios approach Redfield proportions (Moore et al., 2008). Nevertheless, the  
46 nitracline tends to be deeper than the phosphocline in the ETNA (Hausse et al., 2013; ~~Sandel et al.,~~  
47 ~~2015~~), which also points towards ~~a deficiency of N over P in the euphotic zone-N being the limiting~~  
48 ~~nutrient for primary production in the euphotic zone~~. In the Central and West Atlantic, N:P ratios  
49 beyond 30:1 can be reached (Fanning, 1992; Moore et al., 2008), suggesting a severe P limitation of  
50 primary producers (Ammerman et al., 2003; Mills et al., 2004). ~~Additional input of atmospheric~~  
51 ~~anthropogenic nitrogen into the open ocean could further increase this P deficit in the future (Duce et~~  
52 ~~al., 2008)~~. Oxygen concentrations within the oxygen minimum in the ETNA are usually above  
53  $40 \mu\text{mol kg}^{-1}$  and thus considered too high to support N loss processes in the water column  
54 (Karstensen et al., 2008; Löscher et al., 2012; Ryabenko et al., 2012). However, recent observations  
55 of very low oxygen levels just below the mixed layer associated to anticyclonic modewater eddies  
56 suggest a potential for ~~localized~~ denitrification – with an accompanied decrease in N:P ~~ratios~~levels – in  
57 the open ocean of the ETNA (Karstensen et al., 2015).

58 Discrepancies from the canonical N:P ratio are known to influence productivity and composition of  
59 primary producers (Grover, 1997). Since the average elemental composition of ~~C,~~N and P in  
60 seawater as well as in phytoplankton is ~~106:~~16:1, a deviation of dissolved inorganic nutrients from this  
61 ratio could indicate which nutrient ~~is ultimately limiting for phytoplankton growth~~can potentially become  
62 ~~limiting before the other~~ (Lagus, 2004; Moore et al., 2013). Transferring this concept to upwelling  
63 regions with inorganic N:P ratios below Redfield, one would expect that the limiting nutrient for  
64 phytoplankton growth in those areas is N. It has been shown, however, that certain functional  
65 ecotypes of phytoplankton differ in their required nutrient ratio, as specific cellular entities (e.g.  
66 chlorophyll, proteins or rRNA) of primary producers have a unique stoichiometric composition

67 | deviating from the classical Redfield stoichiometry (Geider and La Roche, 2002; Quigg et al., 2003;  
68 | Arrigo, 2005). Thus, surface waters adjacent to OMZs potentially provide a niche for certain types of  
69 | primary producers, whose growth strategy and metabolic requirements are favored by low ratios of  
70 | N:P. Arrigo (2005) refers to them as „bloomers“ and characterizes them as organisms adapted to  
71 | exponential growth, which contain high amounts of ribosomes and P-rich rRNA. Those organisms  
72 | build their biomass in non-Redfield-proportions and exhibit low cellular N:P ratios. The deficit in  
73 | inorganic N of water masses adjacent to OMZs would thus be reduced by this non-Redfield production  
74 | and N:P ratios further offshore would approach Redfield conditions.

75 | Another concept of phytoplankton growth in N deficient waters is that inorganic nutrients are taken up  
76 | in Redfield proportion by primary producers, which leaves the surface water masses enriched in P.  
77 | Excess phosphate presence has been hypothesized to favor N<sub>2</sub>-fixation (Deutsch et al., 2007). The  
78 | conversion of readily available dissolved N<sub>2</sub> into bioavailable forms of fixed N by diazotrophs could  
79 | replenish the N-deficit in surface waters adjacent to OMZs.

80 | Previous bioassay studies that were conducted to identify controlling factors for primary production in  
81 | the eastern Atlantic using inorganic N, P and dissolved Fe addition, determined N as the key limiting  
82 | nutrient (e.g. (Graziano et al., 1996; Mills et al., 2004; Moore et al., 2008). These findings are in  
83 | accordance with an on-board mesocosm study from the same area, where phytoplankton growth  
84 | depended on the initial supply of N rather than on the N:P ratio and where a combined addition of N  
85 | and P did not further increase biomass production compared to the addition of N sources alone (Franz  
86 | et al., 2012). Additionally, the authors deduced that at low N:P ratios excess P was assimilated by  
87 | non-diazotrophic phytoplankton and was channeled into dissolved organic phosphorus (DOP). As  
88 | DOP might serve as an additional source of P for bacteria and phytoplankton (Mahaffey et al., 2014  
89 | and references therein) and is preferentially taken up by the filamentous diazotrophic cyanobacterium  
90 | *Trichodesmium* (Dyhrman et al., 2006; Sohm and Capone, 2006), it has been proposed that N<sub>2</sub>  
91 | fixation might be stimulated by an enhanced DOP supply under low N:P ratios (Franz et al., 2012).

92 | Until recently, oceanic N<sub>2</sub> fixation was mainly attributed to phototrophic cyanobacteria, such as  
93 | *Trichodesmium* or *Crocospaera*, which are restricted to nutrient depleted surface to subsurface  
94 | waters due to their light demand (Capone et al., 1997; Zehr and Turner, 2001). However, several  
95 | groups of non-cyanobacterial diazotrophs and cyanobacterial symbionts have been detected in  
96 | various oceanic regions, thus demonstrating the ubiquity and high diversity of diazotrophs (Foster et  
97 | al., 2009; Farnelid et al., 2011; Loescher et al., 2014). Despite the growing awareness of diazotrophic  
98 | diversity and distribution, the environmental conditions controlling diazotrophy are still not well  
99 | understood. However temperature, Fe and P availability and dissolved oxygen concentrations are  
100 | regarded as key factors for diazotrophic distribution and partly for active N<sub>2</sub> fixation (e.g. Sohm et al.,  
101 | 2011). The presence of high amounts of fixed N is thought to inhibit N<sub>2</sub> fixation (Weber and Deutsch,  
102 | 2014), since diazotrophs are either outcompeted by fast growing phytoplankton species such as  
103 | diatoms (Bonnet et al., 2009; Monteiro et al., 2011), or they themselves take up bioavailable forms of  
104 | N rather than use the energy consuming process of N<sub>2</sub> fixation (Mulholland and Capone, 2001;  
105 | Mulholland et al., 2001; Dekaezemacker and Bonnet, 2011).

106 | In the ETNA, upwelling of N depleted waters along with high Fe input via Saharan dust deposition  
107 | (Gao et al., 2001) sets a classical niche for N<sub>2</sub> fixation, while high N:P ratios beyond the upwelling

108 region of the ETNA point towards P limitation of diazotrophs (Ammerman et al., 2003; Mills et al.,  
109 2004). Nevertheless, a diverse community of cyanobacterial diazotrophs such as *Trichodesmium*  
110 (Capone et al., 1997; Tyrrell et al., 2003), a variety of unicellular cyanobacterial diazotrophs (Groups A,  
111 B, C, diatom-symbionts) (Falcon et al., 2002; Langlois et al., 2005) as well as non-cyanobacterial  
112 diazotrophs such as different clades of proteobacteria are abundant and widely distributed (e.g.  
113 (Langlois et al., 2005; 2008). Those diazotrophs have previously been demonstrated to actively fix N<sub>2</sub>  
114 in the ETNA (Langlois et al., 2005; 2008; Foster et al., 2009), showing highest rates in nutrient  
115 depleted surface to subsurface waters (Großkopf et al., 2012).

116 | We investigated the effect of variable ~~nitrate~~<sup>N</sup> and ~~phosphate~~<sup>P</sup> supply on phytoplankton growth and  
117 addressed the diazotrophic response to changes in N:P stoichiometry over time in two consecutive  
118 mesocosm experiments. In order to extend the design of previous mesocosm experiments (Franz et  
119 al., 2012), N and P supply ratios were varied while keeping either nitrate or phosphate at constant  
120 | concentrations. High N:P ratios were applied to investigate potential inhibition of ~~marine~~ N<sub>2</sub> fixation,  
121 while low N:P supply ratios were applied to unravel the role of excess P and consecutively formed  
122 DOP on primary production and diazotrophy. Direct N<sub>2</sub> fixation rate measurements as well as  
123 determination of *nifH* gene and transcript abundances were carried out to characterize the  
124 diazotrophic community and their response to the chosen treatment levels. The experimental design  
125 and response variables were chosen in order to assess responses of the phytoplankton community to  
126 possible changes in oceanic nutrient stoichiometry as a consequence of ocean deoxygenation.

127

## 128 2 Methods

### 129 2.1 Experimental Setup

130 In October 2012 we conducted two 8-day mesocosm experiments at the Instituto Nacional de  
131 Desenvolvimento das Pescas (INDP), Mindelo, Cape Verde. The night before the start of each  
132 experiment, surface water was collected with RV *Islândia* south of São Vicente (16°44.4'N, 25°09.4'W)  
133 and transported to shore using four 600 L food safe intermediate bulk containers. Containers for water  
134 transport were first rinsed with diluted HCl and several times with deionized water. The experimental  
135 setup comprised sixteen plastic mesocosm bags, which were distributed in four flow-through water  
136 baths. Blue, transparent lids were added to reduce the light intensity to approximately 20 % of surface  
137 irradiation. The collected water was evenly distributed among mesocosm bags by gravity, using a  
138 submerged hose to minimize bubbles. The volume inside each mesocosm was calculated after adding  
139 1.5 mmol silicic acidate and measuring the resulting silicic acidate concentration. The volume ranged  
140 from 105.5–145 liters. Nutrients in all mesocosms were measured before nutrient manipulation. Nitrate  
141 (NO<sub>3</sub><sup>-</sup>), nitrite (NO<sub>2</sub><sup>-</sup>), phosphate (PO<sub>4</sub><sup>3-</sup>) and silicic acid (Si(OH)<sub>4</sub>) were all below the detection limit and  
142 far below the manipulation levels (see Fig. 2). We therefore conclude that no contamination with these  
143 nutrients occurred during water sampling, transport and mesocosm filling. Experimental manipulation  
144 was achieved by adding different amounts of ~~inorganic N~~nitrate and phosphateP. In the first  
145 experiment, the phosphateP supply was changed at constant nitrateN supply (*varied P*) in thirteen of  
146 the sixteen units, while in the second experiment the nitrateN supply was changed at constant  
147 phosphateP supply (*varied N*) in twelve of the sixteen units. Each of these nutrient treatments was  
148 replicated 3 times. In addition to this, “cornerpoints” were chosen, where both the nitrateN and  
149 phosphateP supply was changed. The „cornerpoints“ were not replicated. These treatments were  
150 repeated during both experiments (see Fig. 1 for experimental design). Four cornerpoints should have  
151 been repeated, but due to erroneous nutrient levels in mesocosm 10 during *varied N*, this mesocosm  
152 also was adjusted to the center point conditions. Experimental treatments were randomly distributed  
153 between the four water baths. Initial sampling was carried out immediately after filling of the  
154 mesocosms on day 1. After nutrient manipulation, sampling was conducted on a daily basis between  
155 09:00 and 10:30 for days 2 to 8. Nutrient levels were set between 2 and 20 μmol L<sup>-1</sup> for nitrate, 0.25  
156 and 1.75 μmol L<sup>-1</sup> for phosphate and 15 μmol L<sup>-1</sup> for silicic acidate. Table S1 gives the target nutrient  
157 concentrations and corresponding measured concentrations in the mesocosms.

158 It has to be noted, that no algal bloom developed in mesocosm 5 during *varied N* (target  
159 concentrations: 17.65 μmol L<sup>-1</sup> NO<sub>3</sub><sup>-</sup>, 0.40 μmol L<sup>-1</sup> PO<sub>4</sub><sup>3+</sup>). Thus, it was not included in the analysis and  
160 data are not presented.

161 Although we refer to our experimental approach as mesocosm experiment, this label might be  
162 disputable depending on the definition of the term mesocosm. Sometimes, experimental enclosures  
163 are only defined by size, where our approach would fall into the range of a microcosm experiment  
164 (<1 m<sup>3</sup>; Riebesell et al., 2010). Independent of its size, a mesocosm can also be defined as a confined  
165 body of water, where environmental factors are manipulated at the community or ecosystem level  
166 (Stewart et al., 2013). In contrast, microcosm experiments are often used to manipulate factors at the  
167 population level and often lack the realism to extrapolate results to natural systems (Stewart et al.,  
168 2013). Although our experimental enclosures are limited in size, we consider itfeel justified ~~in~~

169 | ~~ususinging~~ the term mesocosm, as we conducted our experiments ~~s~~ with ~~a~~ natural communit~~iesy~~  
170 | consisting of at least 3 ~~different~~ trophic levels (bacteria, phytoplankton, microzooplankton).

171

### 172 | **2.3 Nutrients**

173 | Samples (10 mL) for dissolved inorganic nutrients ( $\text{NO}_3^-$ ,  $\text{NO}_2^-$ ,  $\text{PO}_4^{3-}$ ,  $\text{Si}(\text{OH})_4$ ) were taken daily from  
174 | each mesocosm and measured directly using a QuAatro Autoanalyzer (Seal Analytic) according to  
175 | Grasshoff et al. (1999). The detection limits of nutrient analyses were  $0.01 \mu\text{mol L}^{-1}$  for  $\text{NO}_2^-$  and  $\text{PO}_4^{3-}$ ,  
176 |  $0.03 \mu\text{mol L}^{-1}$  for  $\text{NO}_3^-$  and  $0.04 \mu\text{mol L}^{-1}$  for  $\text{Si}(\text{OH})_4$ .

177

### 178 | **2.4 Chlorophyll a**

179 | For chlorophyll a (Chl a) analyses, water samples (0.5 – 1 L) were vacuum-filtered (200 mbar) onto  
180 | Whatman GF/F filters (25 mm,  $0.7 \mu\text{m}$ ) before adding 1 ml of ultrapure water. Filters were immediately  
181 | stored frozen for at least 24 hours. 9 ml acetone (100 %) was then added to each sample and the  
182 | fluorescence was measured with a Turner Trilogy fluorometer, which was calibrated with a Chl a  
183 | standard dilution series (*Anacystis nidulans*, Walter CMP, Kiel, Germany). Chl a concentrations were  
184 | determined according to Parsons et al. (1984).

185

### 186 | **2.5 Dissolved organic phosphorus**

187 | Water samples for analyses were filtered through pre-combusted ( $450^\circ\text{C}$ , 5 hours) Whatman GF/F  
188 | filters (25 mm,  $0.7 \mu\text{m}$ ). The filtrate was stored in acid-clean 60 ml HDPE bottles (5 % HCl) for at least  
189 | 12 hours) and frozen at  $-20^\circ\text{C}$  until further analysis.

190 | Prior to analysis of total dissolved phosphorus (TDP) one metering spoon of the oxidizing reagent  
191 | Oxisolv (Merck) was added to 40 ml of sample, which was hereupon autoclaved for 30 minutes.  
192 | Samples were then analysed spectrophotometrically (Autoanalyzer QuAatro Seal Analytic), following  
193 | Bran and Luebbe AutoAnalyzer Method No. G-175-96 Rev. 13 ( $\text{PO}_4^{3-}$ ). The detection limit was  
194 |  $0.2 \mu\text{mol L}^{-1}$  and analytical precision was  $\pm 8.3\%$ .

195 | DOP concentrations were calculated as:

196

197 |  $\text{DOP} = \text{total dissolved phosphorus (TDP)} - \text{dissolved inorganic phosphorus-phosphate (DIP)} (\text{DIP} \text{PO}_4^{3-})$   
198 | (1)

199

### 200 | **2.6 Particulate organic matter**

201 | Particulate organic matter concentrations were determined by filtering 0.5 – 1 L seawater through pre-  
202 | combusted ( $450^\circ\text{C}$  for 5 hours) Whatman GF/F filters (25 mm,  $0.7 \mu\text{m}$ ) under low pressure (200 mbar).  
203 | Filters were immediately frozen and stored until analysis.

204 | Prior to analysis, particulate organic carbon (POC) and nitrogen (PON) filters were fumed with HCl  
205 | (37 %, for 24 hours) in order to remove inorganic carbon. After drying, filters were wrapped in tin cups  
206 | ( $8 \times 8 \times 15 \text{ mm}$ ) and measured according to Sharp (1974) using an elemental analyzer (Euro EA,  
207 | EuroVector, Milan, Italy).

208 | For particulate organic phosphorus (POP) measurements, filters were autoclaved with the oxidation  
209 | reagent Oxisolv (Merck) and 40 ml of ultrapure water for 30 min in a pressure cooker. Then,

210 orthophosphate was analyzed photometrically according to Hansen and Koroleff (1999).

211  
212 Relationships of dissolved and particulate organic matter accumulation to the inorganic nutrient supply  
213 ratios were determined using Model I regression analyses (SigmaPlot, Systat).

## 215 **2.7 Molecular methods**

216 Samples for the extraction of DNA/RNA were taken by filtering a volume of 1–2 L (exact volumes and  
217 filtration times were determined and recorded continuously) of seawater through 0.2 µm  
218 polyethersulfon membrane filters (Millipore, Billerica, MA, USA). The filters were frozen and stored at -  
219 80 °C until analysis. Nucleic acid extraction was performed using the Qiagen DNA/RNA All prep Kit  
220 (Qiagen, Hilden, Germany) according to the manufacturer's protocol. The extracted RNA was reverse  
221 transcribed to cDNA using the Superscript III First Strand synthesis Kit (Invitrogen) following the  
222 manufacturer's protocol with primers nifH2 and nifH3 (Langlois et al., 2005; Zani et al., 2000). *NifH*  
223 clusters were quantified from DNA and cDNA by quantitative Real Time PCRs as previously described  
224 by Church et al. (2005) and Langlois et al. (2008). TaqMan® qPCRs were set up in 12.5 µl reactions  
225 and were performed in technical duplicates in an ABI ViiA7 qPCR system (Life technologies, Carlsbad,  
226 CA, USA). For each primer and probe set, standard curves were obtained from dilution series ranging  
227 from 10<sup>7</sup> to 10 gene copies per reaction; standards were constructed using plasmids containing the  
228 target *nifH* gene. Sequences of primers and probes are given in Table 1. To confirm purity of RNA,  
229 non-template qPCRs were performed using the corresponding RNA.

## 231 **2.8 <sup>15</sup>N<sub>2</sub> seawater incubations**

232 Seawater incubations were performed in triplicates from each mesocosm on day 1 and day 8 of both  
233 experiments as previously described by Mohr et al. (2010) and Großkopf et al. (2012). Degassed  
234 seawater was filled into evacuated gas-tight 3L Tedlar® bags without a headspace. Addition of <sup>15</sup>N<sub>2</sub>  
235 gas was (depending on the exact water volume in the Tedlar® bag) around 10 ml <sup>15</sup>N<sub>2</sub> per 1 L  
236 seawater. Dissolution of the <sup>15</sup>N<sub>2</sub> gas was achieved by 'slapping' the bubble with a ruler. After  
237 complete dissolution of the added <sup>15</sup>N<sub>2</sub> gas (<sup>15</sup>N<sub>2</sub>-enriched seawater), an aliquot of the <sup>15</sup>N<sub>2</sub> enriched  
238 water was collected for each preparation of enriched seawater and stored in an Exetainer. Seawater  
239 samples were filled headspace-free; 100 ml of seawater were exchanged with previously degassed  
240 seawater containing a defined concentration <sup>15</sup>N<sub>2</sub> and <sup>13</sup>C-NaCO<sub>3</sub>. Incubations were performed in 4.5 L  
241 polycarbonate bottles closed with Teflon®-coated butyl rubber septum caps. The <sup>15</sup>N<sub>2</sub> concentration in  
242 the prepared batches of enriched water was determined to be 250 µmol L<sup>-1</sup>, which translates in an  
243 <sup>15</sup>N-enrichment of about 2 % in the 4.5 L bottle incubations, when adding 100 mL enriched seawater  
244 (depending on temperature and salinity). Water samples were incubated for 24 hours in the  
245 mesocosm water baths, thus at the same temperature and light regime, followed by a filtration on  
246 Whatman GF/F filters, which were analyzed using mass spectrometry as previously described in  
247 Loescher et al. (2014).

## 249 **2.9 Statistical evaluation**

250 Relationships of dissolved and particulate organic matter build up to the inorganic nutrient supply  
251 ratios were determined using regression analyses (SigmaPlot, Systat).

252 **Data selection.** Statistical modelling was performed with a subset of the dataset generated by  
253 removal of (i) variables with missing measures, (ii) incomplete observations, (iii) variables left with all  
254 zero data. In order to minimize collinearity of explanatory variables in the statistical models, a set of  
255 environmental variables was chosen according to their variance inflation factor (VIF): Starting with a  
256 linear model that included all variables of interest, the variable with the highest VIF was iteratively  
257 determined and removed from the model until all remaining explanatory variables had a VIF <2.5. The  
258 set of explanatory variables with minimal collinearity was used in model selection (see supplemental  
259 material).

260 **Model selection.** Concentrations of genes and transcripts, respectively, as determined by qPCR,  
261 were fitted to the selected explanatory variables in generalized linear additive mixed models  
262 employing functions *gam* and *gamm* of the R package *mge* v1.8.4 (Wood, 2004; 2011). Multivariate  
263 analysis of counts was realized by a factor of gene/transcript names in the model. Temporal variation  
264 of gene/transcript counts was modeled by cubic spline smoothers. Each combination of  
265 gene/transcript and Run\_ID (i.e. *varied N* or *varied P*) was given its own smoothing function. Other  
266 explanatory variables were standardized to zero mean and unit variance and added as covariates, or  
267 —if a non-linear response was to be expected— as cubic spline smoothers.

268 Candidate models were compared by means of the Akaike Information Criterion (AIC) and validated  
269 following the protocol of Zuur et al. (2009). Briefly, this comprised: (i) choice of an appropriate variance  
270 structure for the full model containing all selected explanatory variables and relevant interaction terms  
271 fitted with restricted maximum likelihood (REML); (ii) choice of the optimal fixed structure by  
272 subsequent removal of insignificant model terms fitted with maximum likelihood; (iii) refitting the model  
273 with REML and validation of model prerequisites. Regression models were visualized with the R  
274 packages *lattice* v0.20-30 and *latticeExtra* v0.6-26 (Sarkar, 2008).

275

### 276 3. Results

#### 277 3.1 Bloom development and nutrient dynamics in the mesocosms

278 In both consecutive experiments (*varied P* and *N*) a bloom formation was observed following nutrient  
279 manipulation. Nitrate $\text{O}_3^-$  and phosphate $\text{PO}_4^{3-}$  were readily taken up by the plankton community and  
280 nutrient concentrations thus declined until the end of the experiment (Fig. 2).  $\text{NO}_3^-$  was fully depleted  
281 in all mesocosms at days 6–8 in both runs, except in the mesocosms with highest N:P ratios of 48:1  
282 (treatment 12.00N/0.25P in *varied P*-) and 44:1 (treatment 17.65N/0.40P in *varied N*). Residual  $\text{PO}_4^{3-}$   
283 was still detectable at the end of the experiments (day 8) in all mesocosms with initial N:P values <10  
284 (treatments in *varied P*: 6.35N/1.10P, 12.00N/1.25P, 12.00N/1.75P; treatments in *varied N*:  
285 2.00N/0.75P, 4.00N/0.75P, 6.00N/1.03P) indicating a limitation of primary productivity dependent on  
286 the N:P stoichiometryratio.

287 Although initial Chl *a* concentrations were slightly higher in *varied P* than in *varied N* (~0.38  $\mu\text{g L}^{-1}$  and  
288 0.2  $\mu\text{g L}^{-1}$ , respectively), the increase in Chl *a* concentration was 5–10-fold until days 5/6 in *varied P*  
289 compared to 10–50-fold in *varied N*. After the bloom at days 5 and 6 Chl *a* declined again to 0.05–0.7  
290  $\mu\text{g L}^{-1}$  and 0.6–1.7  $\mu\text{g L}^{-1}$  in *varied P* and *varied N*, respectively (Fig. 2). ~~In both runs, bloom formation~~

291 | ~~was initially independent from nutrient supply, however, the applied statistical model showed a slight~~  
292 | ~~positive correlation between Chl a and initial N:P (see supplemental material, figure S1).~~

### 294 | **3.2 Particulate organic matter (POM) ~~buil~~accumulation~~ed-up~~ and stoichiometry**

295 | Temporal dynamics of POM were similar during both experiments. Initial concentrations of POC, PON  
296 | and POP were 10–17  $\mu\text{mol L}^{-1}$ , 1.5–2  $\mu\text{mol L}^{-1}$  and 0.05–0.12  $\mu\text{mol L}^{-1}$ , respectively (Fig. 2). In *varied*  
297 | *P*, POC and PON reached a maximum on day 6, while POP increased until the end of the experiment.  
298 | In *varied N* POM ~~accumulation~~~~build-up~~ also peaked on day 6 or 7 in most mesocosms, but differences  
299 | between N:P treatments were more pronounced in *varied N* compared to *varied P*. Our results indicate  
300 | that POM ~~accumulation~~~~build-up~~ was independent of the initial nutrient supply ratio in both experiments  
301 | (Fig. 3). We observed a significantly positive regression coefficient between maximum POC and PON  
302 | concentrations (defined as peak POC and PON concentration subtracted by the initial (day 1) POC  
303 | and PON concentration) to the initial  $\text{NO}_3^-$  supply (POC:  $r^2 = 0.64$ ,  $p = 0.0006$ ; PON:  $r^2 = 0.80$ ,  $p <$   
304 |  $0.0001$ ) while POP ~~accumulation~~~~build-up~~ showed a significantly positive regression coefficient to initial  
305 |  $\text{PO}_4^{3-}$  supply ( $r^2 = 0.31$ ,  $p = 0.048$ ).

306 | Mean PON:POP ratios during the exponential growth phase appeared to be independent of the initial  
307 | N:P supply ratio in both experimental runs (Fig. 4). With ratios between ~~176.6~~ and ~~232.8~~, the  
308 | PON:POP ratios were above, but close to Redfield proportion in all treatments during the first 5 days  
309 | of the experiments, consistent with an observed initial uptake of N:~~and~~P in Redfield proportions in all  
310 | mesocosms. During the post bloom phase, mean PON:POP ratios were positively correlated with the  
311 | initial nutrient supply ratio ( $r^2 = 0.73$ ,  $p < 0.0001$ ). Nevertheless, stoichiometry of POM (N:P between  
312 | ~~165.9~~ and ~~324.9~~) ~~mostly~~ exceeded Redfield proportions, even in treatments with lowest N:P ratios.

### 314 | **3.3 Dissolved organic phosphorus dynamics**

315 | Initial DOP concentrations during *varied P* were 0.14 ( $\pm 0.009$ )  $\mu\text{mol L}^{-1}$ . ~~In most mesocosms, except~~  
316 | ~~for the one with lowest initial  $\text{PO}_4^{3-}$  supply (12.00N/0.25P), DOP concentrations increased~~  
317 | ~~progressively until the end of the experiment (Fig. 5). and increased in all mesocosms, except in the~~  
318 | ~~one with lowest initial P supply (12.00N/0.25P). Maximum~~~~Highest~~ DOP concentrations of around  
319 | 0.4  $\mu\text{mol L}^{-1}$  were determined in mesocosm 12.00N/0.75P on day 5 and decreased again afterwards.  
320 | ~~In all other mesocosms DOP concentrations increased progressively until the end of the experiment~~  
321 | ~~(Fig. 5).~~ Maximum DOP ~~accumulation~~~~build-up~~ (defined as described for maximum POM  
322 | ~~accumulation~~~~build-up~~, section 3.2) was significantly correlated to the initial  $\text{PO}_4^{3-}$  supply (Fig. 5;  $r^2 =$   
323 |  $0.63$ ,  $p = 0.0007$ ), ~~which was also in accordance with the applied statistical model (Fig. S1).~~

324 | In *varied N* initial DOP concentrations in the mesocosms were 0.2 ( $\pm 0.038$ )  $\mu\text{mol L}^{-1}$  and increased  
325 | slightly until day 3. Afterwards DOP concentrations remained rather constant, although with  
326 | considerable variability in the data (Fig. 5).

327 | ~~A simple mass balance (Table S2) showed that part of the phosphorus pool, i.e. the sum of  $\text{PO}_4^{3-}$ ,~~  
328 | ~~DOP and POP, remained unaccounted for (P pool<sub>x</sub>) at the end of the experiment (P pool<sub>x</sub> in *varied P*~~  
329 | ~~~25% of the initial P pool, P pool<sub>x</sub> in *varied N* ~14%). This undetermined P pool is most likely due to~~  
330 | ~~wall growth, which became visible towards the end of the experiment. However, only in two~~  
331 | ~~mesocosms the difference between P pools sizes on day 2 and day 8 was significant.~~

332

### 333 **3.4 Importance of the *Richelia-Rhizosolenia* symbiosis for diazotrophy**

334 Directly measured rates of N<sub>2</sub> fixation showed an increase with time in *varied P*, while no statistically  
335 significant increase could be observed in *varied N* (Fig. 6).

336 A molecular screening of the diazotrophic community in the initial water batch used for *varied P* using  
337 the *nifH* gene as functional marker gene showed a dominance of filamentous cyanobacterial  
338 diazotrophs related to *Trichodesmium* accounting for ~54% of the diazotrophic community (results  
339 from qPCR), followed by proteobacterial diazotrophs (~36%) in *varied P* (data not shown). The high  
340 abundance of filamentous cyanobacterial diazotrophs indicated the presence of a bloom in the initial  
341 water batch in *varied P*. In *varied N*, the initial community consisted mainly of proteobacterial  
342 diazotrophs (~88%), followed by UCYN-B (9%) and filamentous cyanobacteria (3%).

343 ~~Predictability of gene abundances by time and Run ID (i.e. *varied N* or *varied P*) was assessed at the  
344 response scale by plotting the values fitted by the model with their 95% confidence bands (Fig. 7).  
345 Except for *nifH*\_Fil, which displayed no temporal changes at all, gene count levels were generally  
346 higher in *varied P* compared to *varied N*. This was especially true for all time points in case of the  
347 gamma-proteobacterial *nifH*\_AO (significance bands overlapped by less than one half, Cumming et  
348 al., 2007). In unicellular cyanobacterial clusters *nifH*\_UA (UCYN-A) and *nifH*\_CR (UCYN-B), gene  
349 abundances differed only intermittently between the two experimental runs: While *nifH*\_UA  
350 abundances were higher in *varied P* at days 3-4 and 6-8, *nifH*\_CR abundances were higher in *varied*  
351 *P* at days 2 and 4-6.~~

352 Changes in transcript abundance over time (Fig. 8) showed mainly different response patterns  
353 ~~between~~ were most intense for *Richelia-Rhizosolenia* (Het I) transcripts. At day 2, *nifH*\_Het I transcript  
354 abundances were higher in *varied N* conditions compared to *varied P*. This relation changed over the  
355 course of the experiments, with a pronounced increase of *nifH*\_Het I transcript abundances between  
356 day 6 and 8 in *varied P* (Fig. 8).

357 Thus, all classical *nifH* clusters (filamentous cyanobacteria, UCYN-A, -B, -C and proteobacteria  
358 diazotrophs) decreased in abundance of genes and gene transcripts down to the detection limit in both  
359 experiments (Figs. 7, 8, S3, S4), whereas diazotrophs of the *Richelia-Rhizosolenia* symbiosis (Het I)  
360 were the only diazotrophs that showed an increase in *nifH* transcripts over the course of the  
361 experiment, exclusively in *varied P* (Fig. 8, S2).

362 ~~In *varied P*, we observed an accumulation of DOP. In contrast, d~~uring *varied N*, *nifH* gene and  
363 transcript abundance of the *Richelia-Rhizosolenia* cluster was close to the detection limit and DOP  
364 ~~accumulation~~ build-up was rather negligible, thus a potential impact of DOP on diazotrophy was  
365 ~~hypothesized. In contrast, we~~ ~~in *varied P*, we observed an accumulation of DOP; i~~ ~~In contrast, n~~ *varied*  
366 ~~To unravel a potential impact of DOP on N<sub>2</sub> fixation, we investigated temporal DOP patterns, which~~  
367 ~~appeared strongly non-linear. At standard scores below -1 (~0.19 μM) DOP tended to contribute~~  
368 ~~negatively to overall *nifH* transcript abundance levels, whereas at standard scores around -0.5~~  
369 ~~(~0.24 μM) the effect of DOP on *nifH* transcript abundance was positive. At higher concentrations,~~  
370 ~~DOP tended again to a neutral or negative impact on transcript abundance indicating an optimum of~~  
371 ~~DOP concentration on *nifH* transcript abundance.~~

372 ~~In varied P.~~ Here, mesocosms with a significant increase in N<sub>2</sub> fixation (12.00N/0.25P and  
373 12.00/0.75P) were also the ones where DOP was used as ~~phosphorus~~P-source for biomass build up  
374 after PO<sub>4</sub><sup>3-</sup> was depleted (Fig. 9). In mesocosm 12.00N/0.75P, PO<sub>4</sub><sup>3-</sup> concentrations were below the  
375 detection limit after day 5. This coincided with a decrease of DOP after day 5, while POP  
376 concentrations increased until the end of the experiment. In mesocosm 12.00N/0.25P, POP also  
377 increased beyond the point of PO<sub>4</sub><sup>3-</sup> depletion and highest POP ~~accumulation~~build-up exceeded  
378 values that could be explained by PO<sub>4</sub><sup>3-</sup> incorporation alone. Thus a potential impact of DOP on  
379 diazotrophy is hypothesized. In mesocosms without a significant increase in N<sub>2</sub> fixation, POP and DOP  
380 concentrations increased until the end of the experiment and no apparent uptake of DOP could be  
381 observed.

382

## 383 4 Discussion

### 384 4.1 Controls on ~~plankton~~Primary pProduction

385 In order to understand potential consequences of changes in nutrient regimes, it is necessary to  
386 determine the factors that control and limit ~~primary-microbial~~ production. In our experiments,  
387 amendments of NO<sub>3</sub><sup>-</sup> significantly increased chlorophyll concentrations and enhanced the  
388 ~~accumulation~~build-up of POM, indicating the ability of the plankton community to rapidly and intensively  
389 react to ~~nitrate~~N availability. These results ~~indicate-suggest~~ that the ultimate limiting nutrient for-  
390 the phytoplankton ~~community-production~~ in our experiment was NO<sub>3</sub><sup>-</sup>. N<sub>2</sub> fixation was measurable in all  
391 initial samples, which indicates the presence of a niche for diazotrophs in the Cape Verde region. For  
392 the upwelling region as well as for the oligotrophic open ocean of the ETNA, ~~nitrate~~N limitation of the  
393 phytoplankton community has previously been reported (Davey et al., 2008; Moore et al., 2008; Franz  
394 et al., 2012). Additionally, Moore et al. (2008) observed a co-limitation of ~~nitrate~~N and ~~phosphate~~P  
395 during nutrient addition bioassay experiments in the ETNA. In our experiment, however, only POP  
396 ~~accumulation~~build-up was positively affected by PO<sub>4</sub><sup>3-</sup> supply. This argues against a secondary  
397 limitation by ~~phosphate~~P, but rather points towards a mechanism of accumulating and storing  
398 phosphate as polyphosphate within the cell (Schelske and Sicko-Goad, 1990; Geider and La Roche,  
399 2002; Martin et al., 2014).

400 There is a large difference between the supply ratio of inorganic nutrients and the PON:POP ratio of  
401 ~~primary-producersthe plankton community~~ in our study. Although initial N:P ratios in our mesocosms  
402 covered a wide range, PON:POP ratios reached maximum values of ~21 in both experiments during  
403 the exponential growth phase. During stationary growth, maximum PON:POP values of ~~398.8~~ in *varied*  
404 *N* and ~~224.9~~ in *varied P* were measured. However, during growth phases in both experiments  
405 PON:POP ratios did never ~~fall~~drop below ~~165.9~~. Very similar results were obtained by Franz et al  
406 (2012) off the Peruvian coast. However, two experiments conducted by Franz et al. (2012) in the  
407 ETNA and ~~off West Africa~~A showed a different response of the phytoplankton community. In these two  
408 cases, N:P supply ratio and PON:POP were highly correlated and PON:POP ratios as low as 6.0 (~~±±~~  
409 1.4) were observed in the stagnant phase. This shows that the stoichiometry of phytoplankton  
410 communities is flexible to a certain extent, but ~~sometimes~~ does not necessarily reach dimensions  
411 observed in laboratory experiments (Hecky et al., 1993) and implied by theoretical approaches (e.g.  
412 Geider and La Roche, 2002; Klausmeier et al., 2004). This may result from differences in the initial

413 community composition ~~tested that sometimes might if it~~ lacks organisms able to assemble a P-rich  
414 growth machinery (Klausmeier et al., 2004; Arrigo, 2005) ~~or in other as yet unresolved factors~~. It has  
415 been reported that cellular N contents ~~s~~ seems relatively inflexible in ~~parts of the some~~ phytoplankton  
416 ~~community groups~~, thus restricting the maintenance of metabolic processes at low dissolved inorganic  
417 nitrogen ~~(DIN)~~ concentrations (Moore et al., 2013). In contrast, ~~phosphate~~P requirements seem to be  
418 comparably flexible, as certain cellular components containing P (e.g. phospholipids) can be replaced  
419 by non-phosphorus containing compounds (Moore et al., 2013). This can also be deduced from our  
420 experiments, where higher N:P ratios lead to increasing PON:POP ratios, possibly due to the flexibility  
421 to substitute P compounds within the biomass. In contrast, lower N:P ratios lead to lower biomass  
422 accumulation, as the plasticity of PON:POP seems to be constrained by the availability of ~~nitrate~~N in  
423 our experiments.

424

#### 425 **4.2 The impact of bioavailable N on N<sub>2</sub> fixation**

426 The ability of diazotrophs to grow independent of a fixed N source ~~in principle~~ gives them ~~in principle~~  
427 an advantage to thrive under conditions where their competitors are limited by N availability. At the  
428 same time, diazotrophs are considered disadvantaged when competing with faster growing non-  
429 diazotrophs for nutrients under N replete conditions (Tyrrell, 1999; Ward et al., 2013). Contrary to this  
430 classical view, we could not detect a direct influence of reactive N compounds on N<sub>2</sub> fixation in our  
431 experiments. Despite a wide spectrum of applied ~~nitrate~~N concentrations in *varied N*, no significant  
432 ~~change-difference~~ in N<sub>2</sub> fixation rates could be detected. Evidence from culture experiments also  
433 suggests that inorganic N compounds do not ~~universally-always~~ repress N<sub>2</sub> fixation. While NO<sub>3</sub><sup>-</sup>  
434 addition in *Trichodesmium* spp. (Mulholland et al., 2001; Holl and Montoya, 2005) and NH<sub>4</sub><sup>+</sup> addition in  
435 *Crocospaera watsonii* (Dekaezemacker and Bonnet, 2011) reduced N<sub>2</sub> fixation rates, NO<sub>3</sub><sup>-</sup> addition  
436 did not reduce N<sub>2</sub> fixation rates in *C. watsonii* and *Nodularia* spp. cultures (Sanz-Alférez and del  
437 Campo, 1994; Dekaezemacker and Bonnet, 2011). Moreover, recent field surveys demonstrated the  
438 occurrence of N<sub>2</sub> fixation in nutrient rich water masses of the eastern tropical South Pacific (ETSP)  
439 and equatorial Atlantic upwelling regions (Fernandez et al., 2011; Subramaniam et al., 2013; Loescher  
440 et al., 2014) and also modelling ~~efforts-studies~~ predict high N<sub>2</sub> fixation rates in waters containing  
441 measurable amounts of ~~fixed-reactive~~ N (Deutsch et al., 2012; Weber and Deutsch, 2014). Clearly, the  
442 degree of feedback concerning the inhibition of N<sub>2</sub> fixation by reactive N compounds is not universal  
443 and there is evidence that the absence of P and Fe in seawater is a stronger indicator for limitation of  
444 N<sub>2</sub> fixation than the presence of inorganic N compounds (Weber and Deutsch, 2014).

445

#### 446 **4.3 The role of excess P and DOP as controls on N<sub>2</sub> fixation**

447 Deutsch et al. (2007) suggested that N<sub>2</sub> fixation is favored in upwelling regions, where N loss in  
448 adjacent OMZ waters and P leaching from the sediment lead to upwelling of waters enriched in P. This  
449 excess P is thought to be consumed by diazotrophs, thus replenishing the N-deficit in the vicinity of  
450 upwelling regions.

451 As nutrients were taken up in Redfield or above Redfield proportions in our experiments we would  
452 have expected excess ~~phosphate~~P in mesocosms with N:P supply ratios below Redfield. Instead,  
453 excess ~~phosphate~~P was absent and our data point towards a channeling of PO<sub>4</sub><sup>3-</sup> through the

454 | particulate pool into DOP, as an increase in  $\text{PO}_4^{3-}$  supply significantly increased the concentration of  
455 | DOP. Why phytoplankton synthesize and excrete higher levels of DOP under excess phosphateP  
456 | conditions remains unclear, but enhanced  $\text{PO}_4^{3-}$  uptake (followed by DOP accumulation) is thought  
457 | to hamper P limitation when sudden boosts in N are encountered (Mackey, 2012). In accordance with  
458 | our study, mesocosm experiments from the ETNA and eastern tropical south Pacific (ETSP) open  
459 | ocean (Franz et al., 2012) and measurements from shelf regions of the ETNA (Reynolds et al., 2014)  
460 | and Celtic Sea (Davis et al., 2014) showed the accumulation of DOP under excess phosphateP supply.  
461 | Although the composition and bioavailability of the DOP pool needs to be further evaluated, DOP may  
462 | act as a source of P for prokaryotic primary producers, either exclusively or in addition to  $\text{PO}_4^{3-}$  DIP  
463 | (Björkman and Karl, 2003; Dyhrman et al., 2006; Mahaffey et al., 2014; Reynolds et al., 2014). This  
464 | indicates that the ability to utilize DOP may give diazotrophs a competitive advantage when  
465 | bioavailable forms of N are depleted and either  $\text{PO}_4^{3-}$  or DOP concentrations are sufficient.  
466 | In our experiments a significant increase in  $\text{N}_2$  fixation rates was only measured-detected in *varied P*.  
467 | In mesocosms with highest  $\text{N}_2$  fixation rates,  $\text{DIP-PO}_4^{3-}$  was depleted after day 5 or 6 while POP  
468 | increased until the end of the experiment. After  $\text{DIP-PO}_4^{3-}$  depletion, DOP concentrations declined,  
469 | which indicates that DOP served as phosphorusP source until the end of the experiment. It has to be  
470 | noted that  $\text{N}_2$  fixation rates were only measured at the beginning and the end of our experiment and  
471 | possible fluctuations over time cannot be accounted for. However, #increasing diazotrophic transcript  
472 | abundances of *Richelia intracellularis* in symbiosis with the diatom *Rhizosolenia* (Het I) were also  
473 | detected over the course of the *same-variable P* experiment. While the diatom abundance was  
474 | probably favored by replete amounts of silicic acidate added at the beginning of the experiment, no  
475 | increase in diatom-diazotroph associations (DDAs) was detected in the *varied N* experiment.  
476 | Measured  $\text{N}_2$  fixation rates and transcript abundances lead~~This leads~~ us to speculate that DDAs were  
477 | favored in the *varied P* experiment, where diazotrophs in the mesocosms utilized DOP resources in  
478 | order to supply P to themselves and/or their symbiont. The ability to utilize DOP has previously been  
479 | shown for *R. intracellularis* (Girault et al., 2013) and our observations suggest that they may not only  
480 | provide their symbionts with N via  $\text{N}_2$  fixation but also with P via DOP utilization.  
481 | DDAs in our experiment were favored by replete amounts of silicic acidate and DOP and were – in  
482 | contrast to the classical view – not restrained by reactive N compounds. These findings suggest that  
483 | DDAs have the potential to actively fix nitrogen in shelf waters of upwelling regions. Therefore, the N-  
484 | deficit of upwelled water-masses could already be replenished locally prior to offshore transport.  
485 | A shift within the diazotrophic community towards DDAs could also exert controls on carbon export.  
486 | Grazing, particle aggregation and export likely increase when filamentous and proteobacterial  
487 | cyanobacteria are replaced by DDAs (e.g. Karl et al., 2008; 2012). The enhanced strength and  
488 | efficiency of the biological pump would therefore increase the potential for carbon sequestration in the  
489 | ETNA.

490

## 491 | **5 Conclusions and future implications for ETNA**

492 | Our findings add to the growing evidence that diminished N:P ratios in upwelling waters in the ETNA  
493 | will either decrease the biomass of non-diazotrophic primary producers, specifically due to the decline  
494 | of bioavailable N, or lead to a community shift towards primary producers that are able to adapt to

495 | changing N:P conditions. As a considerable amount of DOP was produced under excess **phosphateP**  
496 | conditions, changes in the N:P ratio of waters could exert profound control over DOP production rates  
497 | in the ETNA. Our results indicate that enhanced DOP production in upwelling regions will likely fuel N<sub>2</sub>  
498 | fixation, with an advantage for those diazotrophs capable of DOP utilization. We propose that N<sub>2</sub>  
499 | fixation in the ETNA might not only be restricted to the oligotrophic open ocean but can occur in  
500 | nutrient-rich upwelling regions as previously demonstrated for the tropical Pacific (Löscher et al.,  
501 | 2014) and the Atlantic equatorial upwelling (Subramanian et al., 2013), as N<sub>2</sub> fixation in DDAs seems  
502 | to be favored by the presence of silicic **acidate** and DOP, and not by the absence of fixed N  
503 | compounds.

504

505

### 506 **Acknowledgements**

507 | The authors thank their colleagues from the INDP, Cape Verde for their assistance with setting up the  
508 | experiment. We further acknowledge the captain and crew of RV Islandia. We thank Ulrike Panknin,  
509 | João Gladek, Ivanice Monteiro, Nuno Viera, Elizandro Rodriguez, Miriam Philippi and Chris Hoffmann  
510 | for technical assistance; further, we thank Alexandra Marki, Jasmin Franz, Harald Schunck and  
511 | Markus Pahlow for helpful discussion of the results. This study is a contribution of the DFG funded  
512 | Collaborative Research Center 754 ([www.sfb754.de](http://www.sfb754.de)).

513

#### 514 Authors' contribution

515 | HH and RK designed the experiment with input from JM, CRL, AL, CB, UR, RAS; led the logistics and  
516 | the study on site and provided nutrient and hydro-chemical datasets. JM, RK, AFR, AL, CB and HH  
517 | conducted the sampling of particulate and dissolved matter. JM and AFR performed DOM and POM  
518 | measurements, CRL performed N<sub>2</sub> fixation and molecular experiments and measurements. **SCN**  
519 | ~~performed the statistical modelling of the datasets.~~ JM and CRL wrote the manuscript with input from  
520 | all co-authors.

521

522 | All data will be uploaded at [www.pangaea.de](http://www.pangaea.de) upon publication.

523

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776 Springer, New York. 2009.~~
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- 780 **Tables**
- 781 Table 1: Primers and Probes used in *nifH* TaqMan qPCR assays.

Target Group	Reverse Primer (5'-3')	Forward Primer (5'-3')	Probe (5'-3')
Filamentous (Fil)	GCAAATCCACCGCAAACAAC	TGGCCGTGGTATTATTACTGCT ATC	AAGGAGCTTATACAGATC A
UCYN-A	TCAGGACCACCGGACTCAAC	TAGCTGCAGAAAGAGGAACTGT AGAAG	TAATTCCTGGCTATAACA
UCYN-B	TCAGGACCACAGATTCTACACACT GGTATCCTTCAAGTAGTACTTCGTCT	TGCTGAAATGGGTTCTGTTGAA TCTACCCGTTTGATGCTACACA	CGAAGACGTAATGCTC AACTACCATTCTTCACT
UCYN-C	AGCT AACAAATGTAGATTTCTGAGCCTTATT	CTAA	GCAG
GamAO	C	TTATGATGTTCTAGGTGATGTG	TTGCAATGCCTATTCCG TCCGGTGGTCCCTGAGCC
Het I (Rich-Rizo)	AATACCACGACCCGCACAAC	CGGTTTCCGTGGTGTACGTT	GTGT TCTGGTGGTCCCTGAGCC
Het II (Rich-Hemi)	AATGCCGCGACCAGCACAAC	TGGTTACCGTGATGTACGTT	GTGT

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### 783 Figure captions

784 Figure 1: Experimental design and initial nutrient supply conditions during *varied P* (blue circles) and  
785 *varied N* (red diamonds). “Cornerpoints” during *varied P* and *varied N* are depicted as grey circles and  
786 white diamonds, respectively. Error bars denote the standard deviation of replicated (n=3) treatments.

787 Figure 2: Temporal development of (A)  $\text{NO}_3^-$  and  $\text{NO}_2^-$ , (B)  $\text{PO}_4^{3-}$ , (C) Chl a, (D) POC, (E) PON and  
788 (F) POP within all treatments of both experimental runs. Standard errors-deviations are depicted as  
789 shaded error bands.

790 Figure 3: Maximum POC, PON and POP accumulation build-up as a function of the initial supply of  
791  $\text{NO}_3^-$ ,  $\text{PO}_4^{3-}$  and N/P. Maximum  $\delta$  POM is defined as peak POM concentration subtracted by the initial  
792 (day 1) POM concentration. Treatments in *varied P* are depicted as blue circles; treatments in *varied N*  
793 are depicted as red diamonds. Error bars denote the standard deviation of replicated (n=3) treatments.  
794 Regression lines (continuous lines) indicate significant linear correlations between the initial nutrient  
795 supply and POM accumulation.

796 Figure 4: PON/POP stoichiometry during (A) the exponential growth phase and (B) the stationary  
797 growth phase of the experiment. The grey line visualizes the Redfield Ratio. The color code, symbols  
798 and lines are the same as in Fig. 3. The color code is the same as in Fig. 3.

799 Figure 5: Temporal development of DOP with standard errors-deviations depicted as shaded error  
800 bands.

801 Figure 6: Positive linear correlation between maximum DOP accumulation (defined as peak DOP  
802 concentration subtracted by the initial DOP concentration) and initial  $\text{PO}_4^{3-}$  supply during *varied P*  
803 (blue circles) and *varied N* (red diamonds). Maximum DOP build-up (defined as peak DOP  
804 concentration subtracted by the initial DOP concentration) as a function of initial P supply during  
805 *varied P* (blue circles) and *varied N* (red diamonds).

806 Figure 7: Mean  $\text{N}_2$  fixation rates measured on day 2 and day 8 of both experiments. Because of the  
807 high variance between replicates we omitted  $\text{N}_2$  fixation rates from un-replicated treatments. Asterisks  
808 indicate a significant difference between day 2 and day 8 (t-test). Error bars indicate the standard  
809 deviation.

810 Figure 8: Temporal development of transcript abundances for (A) *Richelia-Rhizosolenia* (Het I) and  
811 filamentous cyanobacteria related to *Trichodesmium* (Fil). Standard deviations are depicted as shaded  
812 error bands.

813 Figure 8: Selected gene count models over time. Predicted counts (solid lines) with 95% confidence  
814 intervals (CI; dashed lines) are plotted along with measured count data. The predictive model is based  
815 on the original (untransformed) gene counts. For better visualization, values were square root-  
816 transformed prior to plotting.

817 Figure 9: Selected transcript count models over time. Predicted counts (solid lines) with 95%  
818 confidence intervals (CI; dashed lines) are plotted along with measured count data. The predictive  
819 model is based on the original (untransformed) transcript counts. For better visualization, values were  
820 square root transformed prior to plotting.

821 Figure 940: Dynamics of  $\text{PO}_4^{3-}$ , POP and DOP and  $\text{N}_2$  fixation rates in all mesocosms during *varied P*.  
822 Because of the high variance between replicates we omitted  $\text{N}_2$  fixation rates from un-replicated  
823 treatments.