Impact of increasing carbon dioxide on dinitrogen and carbon fixation rates under oligotrophic conditions and simulated upwelling

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Dinitrogen (N2) fixation is a major source of bioavailable nitrogen to oligotrophic ocean communities. Yet, we have limited understanding how ongoing climate change could alter N2 fixation. Most of our understanding is based on short-term laboratory experiments conducted on individual N2-fixing species whereas community-level approaches are rare. In this longer-term in situ mesocosm study, we aimed to improve our understanding on the role of rising atmospheric carbon dioxide (CO2) and simulated deep water upwelling on N2 and carbon (C) fixation rates in a natural oligotrophic plankton community. We deployed nine mesocosms in the subtropical North Atlantic Ocean and enriched seven of these with CO2 to yield a range of treatments (partial pressure of CO2, pCO2 = 352–1025 μatm). We measured rates of N2 and C fixation in both light and dark incubations over the 55-day study period. High pCO2 negatively impacted light and dark N2 fixation rates in the oligotrophic phase before simulated upwelling, while the effect reversed in the light N2 fixation rates in the bloom decay phase after added nutrients were consumed. Dust deposition and simulated upwelling of nutrient-rich deep water increased N2 fixation rates and nifH gene abundances of selected clades including the unicellular diazotrophic cyanobacterium clade UCYN-B. Elevated pCO2 increased C fixation rates in the decay phase. We conclude that elevated pCO2 and pulses of upwelling have pronounced effects on diazotrophy and primary producers, and upwelling and dust deposition modify the pCO2 effect in natural assemblages.

The capacity of the marine microbes to fix atmospheric carbon dioxide (CO2) is largely governed by the availability of bioavailable nitrogen, such as nitrate (NO3–) and ammonium (NH4+), in the sunlit ocean (Falkowski 1997). Biological dinitrogen (N2) fixation, although an energetically expensive process, is a major source of bioavailable nitrogen to marine biota in oligotrophic waters in equatorial, tropical, and subtropical regions (Zhang et al. 2020). Despite recognizing the role of anthropogenic activities on the nitrogen cycle, the influence of ongoing climatic changes on N2 fixation is insufficiently understood (Jickells et al. 2017; Zehr and Capone 2020).

Partial pressure of CO2 (pCO2) in the atmosphere has increased from nearly 280 μmol mol−1 in 1750 to 416 μmol mol−1 in 2021 (Keeling and Keeling 2017). Under the high CO2 emission scenario RCP8.5, pCO2 may rise above 1000 μmol mol−1 by 2100 (Moss et al. 2010). The ocean is one of the largest sinks of CO2 and removes around 30% of the anthropogenic carbon (C) added to the atmosphere (Gruber et al. 2019). Consequential rise in oceanic pCO2 is considered a threat to marine ecosystems (Doney et al. 2020). While marine scientists have studied its impact on many ocean phytoplankton species (e.g., Riebesell 2004), its role on community level N2 fixation remains less well understood (Hutchins and Fu 2017). Previous studies on open ocean N2 fixers were carried out mainly on Trichodesmium and Crocosphaera (UCYN-B), two common diazotrophic species in the ocean, in monoclonal culture experiments (e.g., Hutchins et al., 2007; Fu et al., 2008; Wannicke et al., 2018). Those studies underscored the mixed physiological response on nitrogenase activity in individual species to elevated pCO2. However, laboratory experiments do not consider ecological interactions, e.g., competition for nutrients among diazotrophs and other phytoplankton, which result from both direct and indirect effects of high pCO2. Moreover, the most widely distributed cyanobacterial diazotroph, UCYN-A, has not been yet brought into culture (Zehr et al. 2008; Tang et al. 2019). As such, culture studies are not fully representative as they
overlook an important part of the diazotrophic community and their interactions in the environment. To address these limitations and gain insights into a natural community’s response to high pCO₂, we have sub-sampled an in situ mesocosm experiment with eight different pCO₂ levels ranging from the present day atmospheric value to levels projected for the end of this century under selected CO₂ emission scenarios (Bindoff et al. 2019).

Most autotrophic organisms, including diazotrophs possess CO₂ concentrating mechanisms, which promote C fixation by accumulating CO₂ around the CO₂-fixing enzyme RuBisCO (Badger et al. 2005). Thus, in a high pCO₂ ocean, these marine diazotrophs may be able to redirect their energy/resources to fix N₂. This has stimulated the hypothesis that overall N₂ fixation rates will increase at higher pCO₂ levels (Barcelos e Ramos et al. 2007; Hutchins et al. 2007). However, some unicellular diazotrophs such as UCYN-A lack the oxygen-evolving photosystem II and the metabolic ability to fix inorganic C (Zehr et al. 2008). Hence, these species may not directly benefit from high CO₂. Besides, these unicellular diazotrophs are often found in symbiosis with unicellular eukaryotic algae (Thompson and Zehr 2013). unicellular diazotrophic cyanobacterium clade UCYN-A and its symbiotic algae transfer nitrogen and C to each other (Thompson et al. 2012), thus high CO₂ may indirectly benefit them. unicellular diazotrophic cyanobacterium clade UCYN-B fix C themselves, so they could potentially benefit directly from elevated CO₂ (Mohr et al. 2010b; Rabouille et al. 2017). This study was aimed to test the hypothesis that N₂ fixation will increase with increasing pCO₂ levels in the natural environment.

In addition to altered pCO₂ levels in the future ocean, upwelling is predicted to intensify in many oceanic regimes (Narayan et al. 2010; Wang et al. 2015). The Canary Current System is regularly perturbed by natural upwelling via eddies or filaments (Santana-Falcón et al. 2020). Upwelling intensification may impact nutrient cycling in such regimes with a potential enrichment of surface waters with nutrients and thereby increasing primary productivity (Aure et al. 2007). For example, filament structures such as Cape Ghir in the Canary Current Upwelling System, which export 2–3 times more C compared to other filaments (Sangrà et al. 2015), might fix even more C with enhanced upwelling (Xiu et al. 2018). It is unclear what impact upwelling of nutrient-rich water has on surface N₂ fixation rates.

Growth of diazotrophy is limited by iron (Fe) and phosphorus nutrients (Mills et al. 2004). Upwelling and dust deposition are two important sources of such nutrients in the sunlit ocean. Rates of N₂ fixation associated with unicellular diazotrophs can be doubled due to dust deposition events in the Canary Islands (Benavides et al. 2013). Non-diazotrophic primary producers (e.g., diatoms) outcompete diazotrophs with regard to direct nutrient uptake during upwelling (Mills and Arrigo 2010). In addition, the nitrate: phosphate (NO₃⁻ : PO₄³⁻) ratio in surface waters has been proposed to play a major role for nutrient cycling and primary production (Tyrrell 1999). Sub-surface waters, when upwelled, have potential to change surface water nutrient stoichiometry (Franz et al. 2012), and, in turn, can influence N₂ fixation rates. From the biogeochemical modeling point of view, the mathematical expression for excess phosphorus, i.e., P* = ([PO₄³⁻] – [NO₃⁻])/16, which is based on the Redfield ratio, is useful for estimating the distribution of N₂ fixation rates where atmospheric dust deposition is sufficient (Deutsch et al. 2007; Knapp et al. 2016). Sub-surface water in the Atlantic is generally depleted in PO₄³⁻ (negative P*) compared to NO₃⁻ (Lomas et al. 2010; Singh et al. 2015). Introduction of high NO₃⁻ and low PO₄³⁻ water to a surface plankton community might inhibit N₂ fixation rates (Knapp et al. 2012). Hence, we tested a second hypothesis, which was that the addition of deep water (simulating future upwelling events) will decrease N₂ fixation rates due to higher bioavailable inorganic nitrogen concentrations.

**Methods**

**Mesocosm set-up**

A detailed methodology on the mesocosm study set-up and carbonate chemistry manipulation is provided in Taucher et al. (2017) but we give a brief account here on the mesocosm setup as well as a detailed explanation of incubations carried out for analysis of C and N₂ fixation rates. Nine ‘Kiel Off-Shore Mesocosms for Ocean Simulations’ (KOSMOS; Riebesell et al. 2013) were deployed in the northern part of Gando Bay (27°56' N, 15°22' W, eastern North Atlantic) in Gran Canaria at a water depth of ~20–25 m on 23 September 2014. Each mesocosm contained around 35,000 L (31.6–37.8 m³) seawater. Two mesocosms were set up as controls (369 and 352 μatm pCO₂, respectively) and therefore not treated with CO₂ additions, while others were enriched with CO₂ saturated water with a gradient of seven individual treatments (448, 563, 668, 716, 887, 976, and 1025 μatm pCO₂ averaged over the experimental duration, distributed randomly in seven mesocosms). The first CO₂ manipulation was done stepwise at the beginning of the study during 1–6 October 2014 (day 0 is counted from 1 October) to establish the treatments and allow some time for acclimation, followed by manipulations on days 21 and 38 to maintain targeted pCO₂ levels. Deep water was collected from 650 m depth at a location (27°57' N, 15°18' W) about 4 nautical miles north-east from the study site where the water depth was about 1000 m. We simulated upwelling by replacing ~20% of water in each mesocosm with the collected deep water during the night from day 24 to 25. Mesocosms were sampled over 0–13 m depth using an integrated water sampler (HydroBIOS, Kiel) before noon on each sampling day.

**Carbonate chemistry analyses, nutrient measurements and aerosol optical depth estimates**

Samples for dissolved inorganic carbon and total alkalinity were gently filtered (0.2 μm pore size) using a peristaltic pump and measured on the same day. Dissolved inorganic carbon
concentrations were determined by infrared absorption using a LI-COR LI-7000 on an AIRICA system (MARIANDA, Kiel) with an overall precision better than 5 μmol kg⁻¹. Total alkalinity was measured by potentiometric titration using a Metrohm 862 Compact Titrosampler and a 907 Titrandos unit following Dickson et al. (2003). The estimates of pCO₂ values were based on measured total alkalinity, dissolved inorganic carbon, silicate, and PO₄³⁻ concentrations as well as measured temperature and salinity using CO2SYS (Pierrot et al. 2011) and carbonate chemistry constants by Lueker et al. (2000).

Nutrient samples were collected in acid-cleaned (10% HCl) bottles (PETG), filtered (0.45 μm pore size filter, cellulose acetate filters, Whatman) immediately upon arrival of samples in the laboratory, and analyzed on the same day. Concentrations of NO₃⁻, Si(OH)₄, and PO₄³⁻ were measured with a SEAL Analytical QuAAtro AutoAnalyzer following Hansen and Koroleff (2007). Precision (detection limit) of nutrients (based on standard deviation of triplicates) was ±0.007 μM (0.03 μM) for NO₃⁻, ±0.003 μM (0.008 μM) for PO₄³⁻ and ±0.011 μM (0.05 μM) for Si(OH)₄.

Daily total and dust aerosol optical depth (AOD) at 550 nm over the study region (27°–28° N, 15°–16° W) has been derived from the Copernicus Atmosphere Monitoring Service (CAMS) model reanalysis (Inness et al. 2019). The CAMS reanalysis are global fields of atmospheric composition developed by the European Centre for Medium-Range Weather Forecasts (ECMWF). The CAMS model system assimilates a suite of satellite-based observations related to trace gases and aerosols with the Integrated Forecasting System. The CAMS reanalysis has improved spatial and temporal resolutions (80 km, 3-hourly), and this model has been shown to have smaller biases when compared with independent observational datasets (Inness et al. 2019).

Carbon and nitrogen fixation experiments

We did 14 dark incubations and 11 light incubations for C and N₂ fixation rate measurements without replication for each pCO₂ level on different days from day – 3 (28 September 2014, 3 days before the first CO₂ enrichment was done in the mesocosms) to day 55 (25 November 2014). We started light incubations from day 11. During each morning of the experimental day, water samples from the nine mesocosms and surrounding Atlantic water (for reference) were collected over 0–13 m depth using the integrated water sampler and transferred into 2.8 L polycarbonate bottles (Nalgene, Thermo Fisher Scientific, Waltham, MA) on board the sampling boats. Polycarbonate bottles were stored in a cool box to control temperature and to block sunlight until return to the on-shore laboratory.

Due to the low nutrient concentrations, surface Atlantic water was used to prepare dissolved ¹⁵N₂ solution (98% atom% ¹⁵N, Lot No. 1-18404, Cambridge Isotope Laboratories, MA) on each sampling day following a method modified from Mohr et al. (2010a). Dissolution of the ¹⁵N₂ gas was achieved by ~ 30 min of gentle slapping of the enriched gas bubble in a gas sampling bag (Tedlar bag with single polypropylene fitting) with a ruler. Each sample bottle was mixed gently by rotating around 10 times before 50 mL of water was removed to make space for the dissolved ¹⁵N₂ enriched seawater. Subsequently, 50 mL of this dissolved labeled ¹⁵N₂ tracer water was added to each polycarbonate bottle to achieve a final ¹⁵N enrichment of 2.35 atom %. Additionally, 1 mL (0.2 M concentration) of labeled NaH¹³CO₃ (99%, Cambridge Isotope Laboratories, MA) dissolved in deionized water was added to the same 2.8 L bottles for primary production rate measurements (C fixation rates). By mistake, we added 2 mL of the same concentration of NaH¹³CO₃ during the days 25 to 37 for C fixation rates. Carbon isotopic (¹³C) enrichment was estimated for each mesocosm based on the ambient dissolved inorganic carbon concentrations as reported by Taucher et al. (2017). The ¹³C enrichment varied from 4.06 to 7.08 atom % (higher enrichment were during the days 25 to 37). After addition of both the tracers, bottles were closed and gently rotated to mix the water.

For light incubations, sample bottles were placed in an incubator covered with 25% transmittance filter sheet (Blue Lagoon, LEE filters) for 24 h at the Spanish Bank of Algae, next to the land-based facilities of the Plataforma Oceánica de Canarias (PLOCAN). For the dark incubations, sample bottles were placed in a fully closed and covered cemented tank inside a laboratory at PLOCAN for 24 h in running seawater. Running seawater was used to maintain in situ temperature in both the light and dark incubations, which ranged from 26°C initially to 21.5°C towards the end of the study in November. After 24 h of incubation, a subsample of seawater from each incubation bottle (volume: 0.5–2.8 L, depending on sample biomass) was filtered onto pre-combusted (450°C, 5 h) 25 mm diameter glass fiber filters using vacuum filtration (<200 mbar pressure). We used 0.7 μm pore size filters (GF/F, Whatman, UK) for light incubations and 0.3 μm pore size filters (GF75, Advantec, Toyo Roshi Kaisha Ltd., Japan) for dark incubations, in order to capture smaller heterotrophic diazotrophs in dark incubations that can pass through the 0.7 μm GF/F (Bombar et al. 2018). All filtrations were completed within 2 h. Filters were dried at 50°C for 24 h and stored in cryogenic vials until mass spectrometric analysis. Samples for background isotope abundances in particular organic matter were collected from the mesocosms and filtered onto GF filters as described for the incubations.

For isotope analysis, the filters were stored over concentrated (36%) HCl for 4 h in a desiccator so that the acidic fumes removed carbonates present. Filters were then dried overnight in an oven at 50°C and later pelletized in tin cans. Samples were analyzed for particulate organic carbon (POC) and nitrogen (PON) and their isotopic composition (δ¹⁵N and δ¹³C) using an elemental analyzer coupled to a Conflo II and a Finnigan DeltaPlus isotope ratio mass spectrometer (IRMS) at GEOMAR, Kiel. International Atomic Energy Agency (IAEA)
standard materials (acetanilide, caffeine) were used for mass and isotope calibration of C and N (standard deviation < 0.2‰ δ15N and δ13C). Nitrogen fixation rates were calculated following Montoya et al. (1996):

\[ N_2 \text{ fixation} = \left( \frac{1}{t} \right) (A_{PN_i} - A_{PN_f}) / [PON_f] \]

where \( t \) = incubation time, \( A_{PN_i} = 15N \text{ atom}\% in PON at the end of incubation, \( A_{PN_f} = 15N \text{ atom}\% in PON at the start of the incubation.} \ A_N = 15N \text{ enrichment in the dissolved form after the tracer addition at the start of the incubation, and PON}_f refers to particulate organic nitrogen concentration at the end of the incubation. In the above equation, all the terms are measured except \( A_N \), which is defined as:

\[ A_N = \frac{(15N_{\text{tracer}} \times \text{tracer concentration} + 15N_{\text{natural}} \times \text{natural concentration})}{\text{tracer concentration} + \text{natural concentration}} \]

The natural concentration of dissolved N2 for a given temperature and salinity was calculated using the table given in Weiss (1970). Carbon fixation rates were calculated similar to N2 fixation rates following Slawyk et al. (1977). Taking into account all the sources of uncertainty (detection limits for POC and POC are 15 and 5 μg, respectively; precisions of δ13C and δ15N to be 0.2), our detection limits for C and N2 fixation rates were 0.12 and 0.05 nM N h⁻¹, respectively. Most of the samples for day 35 (belonging to dark incubations for nitrogen analysis only) were lost due to the technical issues with IRMS.

Molecular methods

Seawater samples were taken directly from the mesocosms on days 1, 3, 19, 25, 28, 35, 41, and 53 of the experiment runtime, in volumes between 1 and 2 L. The water was filtered on 0.2 μm pore size membrane filters (Millipore) and kept at ~80°C until further analysis. Nucleic acids were extracted using the Qiagen AllPrep DNA/RNA kit with modifications as described in Loescher et al. (2014). After spectrophotometric quality checks (NanoDrop), a PCR for the key functional marker gene for N2 fixation, nifH was performed following the protocol of Zani et al. (2000). The nifH PCR products were pooled, Topo TA cloned (Thermo Fisher Scientific) and Sanger-sequenced as a commercially available service, in order to identify major present diazotrophic clades. We performed qPCR analyses for filamentous cyanobacteria, UCYN-A, -B, -C, gammaproteobacteria of the AO and PO types, and the Het I and II clades (Langlois et al. 2005; Foster and Zehr 2006). The qPCRs were performed as described in Loescher et al. (2014) against plasmids containing an environmental copy of the target gene (standards obtained from Langlois et al. [2008]), in technical duplicates on a Biorad qPCR cycler. In addition, we enumerated *Trichodesmium* colonies using a microscope.

Statistical analysis

We analyzed the nifH gene abundances, C and N2 fixation rates for three different experimental phases separately: (1) the oligotrophic phase after CO2 addition until day 23, (2) the phytoplankton bloom (growth) phase after simulated upwelling between day 25 and day 35, and (3) the post-bloom (decay) phase from day 37 until the end of the study. Phases are delineated based on the concentrations of nutrients and chlorophyll—oligotrophic (low nutrients), growth (increase in chlorophyll due to deep water addition) and decay phase (decline of chlorophyll; Taucher et al. 2017). As done previously in such mesocosm experiments (Paul et al. 2015), a pCO2 gradient was chosen for the reasons described in Havenhand et al. (2010). We did linear regression analyses (significant at p < 0.05) to determine the effect of pCO2 on C and N2 fixation rates for each phase. Variables for each phase were averaged and plotted against the average pCO2 in each mesocosms in the corresponding phases. All the data points after day 0 are included in the analysis. Units for normalized rates changed to h⁻¹ (e.g., nM N h⁻¹/[PON] = h⁻¹ for N₂ fixation). Since rates in h⁻¹ were low, we report the rates in day⁻¹.

Results

**Variation in aerosol optical depth and inorganic nutrients**

Variations in the total and dust AOD suggested three dust events over the study region during the experimental duration (Fig. 1). Initiation of these dust events were on days 1, 18 and 31. The second dust event, which started before the growth phase, lasted longer (11 days) as compared to other events. The second dust event was recognized as a Calima event. Calima events are those events in the Canaries which are associated with hot, dry and thick dust layers in the lower atmosphere.

Environmental conditions changed from oligotrophic to eutrophic phase in the mesocosms after deep water addition (Fig. 2). Concentrations of NO₃⁻ were below 0.2 μM until day 23, thereafter peaked around 3.1 μM throughout the mesocosms following the deep water addition. Concentrations of PO₄³⁻ were close to detection limit (0.05 μM) before the deep water addition but increased up to 0.2 μM after addition. Unlike NO₃⁻ and PO₄³⁻, SiO₄ concentrations were relatively higher (0.5 μM) at the start of the experiment and decreased until day 23, followed by an increase of up to 2 μM after deep water addition. All nutrients reached their previous concentration levels in <4 days after deep water addition, suggesting rapid and almost complete assimilation of nutrients added through deep water. Concentrations of NO₂⁻ and PO₄³⁻ were 16.71 and 1.05 μM, respectively, in the deep water, i.e., having a slightly positive P* value (0.09 μM). Estimates of P* in the mesocosms suggested that PO₄³⁻ was in excess until the deep water addition, and decreased thereafter due to possible preferential consumption of PO₄³⁻ in the first part of growth phase.
Fig 1. Variation in the daily means of dust aerosol optical depth (AOD) and total AOD over the study area. Error bars represent standard deviation (1σ) of 3-hourly values. Day 0 (1 October 2014) indicates the day of the first CO2 manipulation, corresponding to the start of elevated pCO2 treatments. The dashed green line represents the time of deep water addition, which was done on the night of day 24. White, green, and gray-shaded areas represent oligotrophic, growth, and decay phases, respectively.

Fig 2. (a) NOx−, (b) PO4^{3−}, (c) Si(OH)4 concentrations, and (d) P* values (P* = [PO4^{3−}] - [NO3− + NO2−]/16). Day 0 (1 October 2014) indicates the day of the first CO2 manipulation, corresponding to the start of elevated pCO2 treatments. Legend numbers indicate pCO2 (μatm) averaged over the study period. The dashed green line represents the time of deep water addition, which was done on the night of day 24. White, green, and gray-shaded areas represent oligotrophic, growth, and decay phases, respectively.
Values of $P^*$ turned positive again as soon as the nutrients from the deep water addition were consumed (day 29).

**Influence of rising carbon dioxide on carbon and nitrogen fixation rates**

Phototrophic N$_2$ fixation rates from the light incubations varied between 0.05 and 1.35 nM N h$^{-1}$ (mean ± standard deviation, i.e., $\mu ± \sigma = 0.40 ± 0.28$ nM N h$^{-1}$, number of data points for all the mesocosms for all the sampling days, $n = 87$). Rates were below average in the oligotrophic phase, while they increased gradually during the growth phase and attained their peak during the decay phase in most mesocosms (Fig. 3a). Higher rates were associated with high p$CO_2$ mesocosms during the decay phase. Interestingly, rates in the surrounding (Atlantic) water also increased during the growth phase but decreased to the oligotrophic phase level in the decay phase. Regression analysis suggested a negative effect of p$CO_2$ levels on N$_2$ fixation rates in the oligotrophic phase, whereas it had a positive effect in the decay phase (Table 1). The CO$_2$ effect was negative in the growth phase as well but it was statistically insignificant.

Dark N$_2$ fixation rates were lower compared to phototrophic N$_2$ fixation rates (Fig. 3a, b), and varied between 0.07 and 0.41 nM N h$^{-1}$ ($\mu ± \sigma = 0.17 ± 0.07$ nM N h$^{-1}$). Like phototrophic N$_2$ fixation rates, regression analysis suggested that dark N$_2$ fixation rates were negatively influenced by p$CO_2$ levels during the oligotrophic phase (Table 1). The levels of p$CO_2$ did not change dark N$_2$ fixation rates during growth and decay phases.

Phototrophic C fixation rates (from the light incubation) varied between 7 and 678 nM C h$^{-1}$ ($\mu ± \sigma = 96 ± 129$ nM C h$^{-1}$). Like N$_2$ fixation rates, there was a positive effect of p$CO_2$ levels on C fixation rates during the decay phase (Table 1). Carbon fixation rates from dark incubations (dark C fixation) remained close to their detection limits (Fig. 3d) and varied between 0 and 14 nM C h$^{-1}$ ($\mu ± \sigma = 2 ± 2$ nM C h$^{-1}$).

We extended similar regression analysis on C and N$_2$ fixation rates normalized by corresponding POC and PON concentrations, respectively. Phase-wise patterns of normalized rates were somewhat similar to volumetric rates (Fig. 4) but the effect of p$CO_2$ on C (phototrophic) and N$_2$ fixation rates was not statistically significant (Table 1).

**Carbon and nitrogen fixation rates following deep water additions**

The simulated upwelling event had a profound impact on volumetric C and N$_2$ fixation rates from the light incubations (Fig. 3). Phototrophic N$_2$ fixation rates increased gradually following the addition of nutrient-rich deep water and dust deposition, and reached up to five times the initial rates after a week in all the mesocosms (Fig. 3a). Rates continued to increase in the two mesocosms with the highest CO$_2$ levels until day 40 (16 d after deep water addition). Likewise, dark
N2 fixation rates increased following deep water addition but these were barely twice the initial rates (Fig. 3b). The magnitude of the increase in dark N2 fixation rates were similar in different CO2 levels. Light C fixation rates increased immediately after simulated upwelling, peaking four days later and declining thereafter (Fig. 3c). There was no such observable change in the dark C fixation rates (Fig. 3d).

However, when the rates were normalized with their respective elemental concentrations, the impact of the simulated upwelling is less discernible on N2 fixation (both light and dark) rates (Fig. 4a,b). Normalized phototrophic C fixation rates were slightly higher, whereas normalized dark C fixation rates remain unchanged and close to detection limit after the deep water addition (Fig. 4c,d).

### Table 1. Summary of linear regression of pCO2 on different variables for each phase. The variables and pCO2 were averaged for each phase. Significant effects (p < 0.05) are listed in bold. Phase I = oligotrophic, phase II = bloom and phase III = decay. Total degrees of freedom, df = 8 for the first phase, and df = 7 for second and third phase as we lost one mesocom having average pCO2 = 976 μatm after the first phase.

| Phase | Variable* | p | r   | F   | Variable (normalized) | p  | R  | F   |
|-------|-----------|---|-----|-----|----------------------|----|----|-----|
| I     | N2 fixation (light) | 0.01 | -0.77 | 10.27 | N2 fixation (light) | 0.07 | -0.62 | 4.46 |
| II    | N2 fixation (light) | 0.15 | -0.56 | 2.69 | N2 fixation (light) | 0.19 | -0.52 | 2.33 |
| III   | N2 fixation (dark) | 0.02 | 0.78 | 9.05 | N2 fixation (dark) | 0.16 | 0.54 | 2.53 |
| I     | C fixation (light) | 0.001 | -0.91 | 34.47 | C fixation (light) | 0.11 | -0.56 | 3.28 |
| II    | C fixation (light) | 0.85 | -0.35 | 0.39 | C fixation (light) | 0.49 | -0.28 | 0.53 |
| III   | C fixation (light) | 0.45 | -0.31 | 0.65 | C fixation (light) | 0.20 | -0.51 | 2.51 |

*Statistics for C fixation (dark) rates is not presented as these rates were close to their analytical uncertainties.

Fig 4. Daily variation in concentration normalized (a) phototrophic (incubation conducted in light) N2 fixation rates, (b) N2 fixation rates (incubation conducted in dark), (c) C fixation rates (incubation conducted in light), and (d) C fixation rates (incubation conducted in dark). Legend numbers and colors are as described for Fig. 2.
Composition of the diazotrophic community
We observed a diazotrophic community typical for the eastern subtropical North Atlantic, including filamentous, unicellular cyanobacteria, and gammaproteobacteria matching the Atlantic Ocean type (Fig. 5). None of the diazotroph clades (i.e., UCYN-A, -B, -C, gammaproteobacteria) were significantly (at \( p < 0.05 \)) influenced by high \( pCO_2 \) in the different phases, so we have not listed the corresponding regression analysis. However, we observed a temporal increase in unicellular cyanobacteria of the UCYN-B clade, and to a lesser extent, of the gammaproteobacteria after deep water addition and dust pulse. *Trichodesmium* colonies (from microscopic observations) were never counted to be above 1 colony L\(^{-1} \) in any of the mesocosms and only detected in genetic analyses before deep water addition (data not shown).

Discussion
Variable carbon dioxide response dependent on nutrient concentrations and plankton bloom status
Our results suggest a variable effect of \( CO_2 \) on \( N_2 \) fixation rates. Under nutrient deplete conditions, both light and dark rates were negatively correlated with \( pCO_2 \), yet under decay conditions, the correlation was positive for light \( N_2 \) fixation rates. Overall the diazotrophic community was dominated by unicellular cyanobacteria, consistent with other observations in the eastern subtropical North Atlantic Ocean (Benavides et al. 2013). Nitrogen fixation by UCYN-A has been shown to be unaffected by high \( pCO_2 \) levels in a natural community of low dissolved Fe in oligotrophic conditions during short (5 d) acclimation experiments (Law et al. 2012). However, under double the modern \( pCO_2 \) levels and Fe replete conditions, rates of \( N_2 \) fixation by UCYN-B were reported to be enhanced by 80% in a culture experiment (Fu et al. 2008). A compilation of studies suggests that high \( pCO_2 \) affects \( N_2 \) fixation rates positively during Fe-replete conditions and negatively during Fe-depleted conditions (Table S1). We did not measure Fe in our samples, but our phototrophic \( N_2 \) fixation rates started increasing in all the treatments gradually after the deep water addition and dust deposition, both being important source of Fe. Rates in the waters from the Atlantic also started increasing around the same time. This might be due to the second Calima event before the deep water addition that may have introduced large amounts of dust (iron), triggering \( N_2 \) fixation in the surrounding waters (Fig. 1). However, a rate increase in the mesocosms may not be only due to likely Fe inputs during the Calima event, as the increase in most mesocosms was somewhat higher (1.35 nM N h\(^{-1} \)) compared to the surrounding Atlantic waters (0.74 nM N h\(^{-1} \)). Rates peaked at 1.35 nM N h\(^{-1} \), up to five times higher than in the oligotrophic phase in the highest \( pCO_2 \) treatments 17 days after the nutrient enrichment. This positive effect of high \( pCO_2 \) levels...
on N₂ fixation in the decay phase became insignificant in the biomass normalized rates (Fig. 4a, Table 1), suggesting that N₂ fixation rates are strongly dependent on the absolute planktonic community biomass present and to a lesser extent on the pCO₂ levels. Since diazotrophic biomass is a negligible fraction of planktonic community biomass in a natural community (Montoya et al. 2002), it could still be argued that the rates are community composition dependent.

In summary, N₂ fixation rates seem to be controlled by a complex interplay of different environmental conditions, including nutrient availability and stoichiometry. If our analysis is interpreted with respect to Fe-limited (during the oligotrophic phase) and Fe-rich (after deep water addition), then the observed CO₂ effects (negative during Fe-limitation, positive during Fe-replete conditions) are consistent with most of the laboratory/single-strain culture studies (Table S1). Hence, nutrient competition and other interactions in a natural plankton assemblage did not appear to modify the direct effects of elevated pCO₂ on N₂ fixation rates reported in culture studies.

Different response timing of nitrogen and carbon fixation to upwelling event

As per the Trichodesmium centric classical view, the first phase of the experiment would open up a niche for N₂ fixers, because of the oligotrophic (low bioavailable nitrogen) conditions (Capone et al. 1997). In the growth phase, increased NO₃⁻ from simulated upwelling would suppress phototrophic N₂ fixation (Holl and Montoya 2005). However, our observations (Fig. 3) are contrary to this classical view of high nitrogen (nutrient) concentrations inhibiting diazotrophic growth (Karl and Letelier 2008) as N₂ fixation rates and UCYN-B nifH copies gradually started increasing after nutrient-rich deep water addition and dust deposition. In fact, N₂ fixation is limited by Fe and PO₄³⁻ (Mills et al. 2004), and N₂ fixers are not strongly inhibited by inorganic nitrogen species such as NO₃⁻ (Fernandez et al. 2011; Loescher et al. 2014). In our study, we anticipate that biological nutrient uptake processes would have quickly increased because deep water contains ample amounts of nutrients for primary producers in this otherwise oligotrophic region. Hence slow growing N₂ fixers may have been outcompeted by rapidly growing phytoplankton such as diatoms as observed in the microscopy data (Taucher et al. 2018a) and in measured C fixation rates (light incubations). Indeed, there was a quick build-up of biomass leading to a phytoplankton bloom, which declined long before diazotrophy peaked on day 41 (figs. 7 and 8 in Taucher et al., 2017). The phytoplankton bloom rapidly resulted into nitrogen limitation around day 29 (Taucher et al. 2018b), creating an amplified diazotrophic niche. We found high C fixation rates associated with a phytoplankton bloom on day 29, followed by an increase in N₂ fixation rates associated with concurrent high UCYN-B copies. Low NO₃⁻, leftover PO₄³⁻ and Fe from the deep water and concurrent dust deposition might have contributed to N₂ fixation rates peaking on day 41 in high pCO₂ mesocosm. A recent meta-analysis consisting of published data on nine species, 18 different strains, and 18 experiments using natural plankton communities suggest an increase in N₂ fixation rates by 29 ± 4% under high pCO₂ scenarios (Wannicke et al., 2018). It has been noted that nutrient limitation (particularly by Fe) negates the positive effect of high pCO₂ on N₂ fixation (Wannicke et al. 2018). Integrating the results from all three phases of the present study, our data indeed indicate that N₂ fixation rates (in the light) are enhanced under elevated pCO₂ conditions if PO₄³⁻ is in excess and Fe is presumably available in higher concentrations. Overall, we suggest that the effects of CO₂ on N₂ fixation are regulated through changes in the plankton community structure and nutrient stoichiometry.

In addition to being the first in situ elevated pCO₂ mesocosm study in subtropical oligotrophic waters, this study is also the first large scale upwelling simulation primarily investigating its impact on C and N₂ fixation rates. Carbon fixation rates showed an increasing trend over 4 days upon upwelling simulation, followed by a decline, while N₂ fixation rates were still increasing. Concentrations of PO₄³⁻ were more than doubled from 0.07 μM before, to ~0.2 μM after the deep water addition and P* became negative (~0.06 μM). The increase in P* to above 0 within a week of the deep water addition suggests that deep water NO₃⁻ was preferentially and rapidly consumed by non-diazotrophs (Mills and Arrigo 2010; Singh et al. 2017), which created an environment of excess PO₄³⁻ where diazotrophs could thrive as found in a PO₄³⁻ enriched mesocosm experiment (Bonnet et al. 2016). Over the course of the study, we indeed encountered a non-diazotrophic cyanobacterial bloom in the growth phase as corroborated in HPLC data (Taucher et al. 2018a). Non-diazotrophic cyanobacteria consume less PO₄³⁻ relative to NO₃⁻ compared to fast growing phytoplankton and N:P in their cells is higher than the Redfield ratio (Bertilsson et al. 2003; Singh et al. 2015). Estimates of P* turned positive as soon as the deep water NO₃⁻ was consumed on day 28. This also coincided on day 29 with high N₂ fixation rates (average of all the light incubations 0.42 ± 0.12 nM N h⁻¹). A delayed increase in N₂ fixation rates after deep water addition is likely because the diazotrophs are slow growing (Berman-Frank et al. 2001) and the niche had to first be established by non-diazotrophic organisms (Mills and Arrigo 2010; Singh et al. 2017). In summary, our results suggest that the diazotrophic activity was stimulated potentially due to niche construction of excess PO₄³⁻ a few days after the addition of nutrient-rich deep water and dust deposition.

Rates of C fixation increased rapidly after deep water addition, indicating fast growing primary producers could take advantage of this nutrient input. In contrast, N₂ fixation rates peaked up to a week later. This lag likely reflects the slower growth rates of diazotrophic species in addition to construction of the niche by non-diazotrophic cyanobacteria. Additionally, this response lag could be due to the effect of
elevated $pCO_2$ on the symbiotic relationship of unicellular diazotrophs with diatoms (Thompson and Zehr 2013). Higher N$_2$ fixation rates occurred in the high $pCO_2$ (> 800 µatm) mesocosms, in which ecological and biogeochemical processes were dominated by a bloom of the toxic microalga *Vicicittus globosus* (Riebesell et al. 2018). It remains unclear if the toxicity has any significant influence on N$_2$ fixation rates, however it appears as though the fundamentally different plankton community in these two mesocosms at high $pCO_2$, along with deep water addition and dust deposition, created an environment that supported high N$_2$ fixation rates.

**Unicellular diazotrophic cyanobacteria driven dark nitrogen fixation rates**

As seen in our *nifH* gene copies, diazotrophs may have particularly benefited from the deep water addition, presumably via the associated PO$_4^{3-}$ and micronutrient supply. There is a chance that seed populations (such as gammaproteobacteria) were introduced into the mesocosms from the deep water as these were undetectable before the deep water addition. Diazotroph abundances were highest in samples from the decay phase. The latter may speak for a standing stock of these diazotrophs, which are able to persist for a certain time even after nutrient drawdown. One of the identified clades, UCYN-B, has been shown to primarily fix N$_2$ in the dark and C in the light (e.g., Großkopf and LaRoche, 2012; Wilson et al., 2010). However, if this photosynthetic clade is limited by light availability, the highly energy-demanding process of N$_2$ fixation would have to be sustained by respiration only, thus explaining the lower activity in dark incubations. Dark N$_2$ fixation rates, however, may be a result of the unicellular cyanobacteria of the UCYN-B clade fixing mainly at night (Mohr et al. 2010b).

**Effect of rising carbon dioxide on carbon fixation rates**

Previous studies suggest either a positive effect or no significant effect of high $pCO_2$ on plankton community C fixation rates (e.g., Schulz et al. 2017). Our results from the oligotrophic and eutrophic phases suggest no effect of elevated $pCO_2$ on phototrophic C fixation. Our C fixation analysis is consistent with the conclusions derived from a $^{14}C$ based primary production study conducted in the same mesocosm study for the decay phase, when $pCO_2$ seems to have a positive effect on C fixation rates (Hernández-Hernández et al. 2018). The magnitude of $^{14}C$ based C fixation rates during all the phases were comparable to those estimated based on $^{13}C$ tracer. Assuming photoautotrophs require 1 mol N to fix 6.6 mol C (Redfield 1958), the contribution of N$_2$ fixation to new nitrogen supply for phototrophic C fixation was moderate (mean value of 4% with highest contribution up to 13%). Dark C fixation rates varied within the analytical uncertainties suggesting absence of lithoautotrophs or chemoautotrophs in our experiments (Figs. 3d, 4d).

**Conclusion**

Under the applied $pCO_2$ range (352–1025 µatm), we observed variable effects of elevated $pCO_2$ on C and N$_2$ fixation that appeared to depend on nutrient concentrations and bloom status during our 55 day-long experiment. Interestingly, our results suggest positive effect of high $pCO_2$ on C fixation rates only in the decay phase. The strongest negative effect of CO$_2$ on N$_2$ fixation was detected during the oligotrophic phase, prior to the simulated upwelling event. The nutrient-rich deep water and dust deposition had a marked impact on both N$_2$ fixation and C fixation rates, magnitudes higher than the difference between CO$_2$ treatments. Our analysis indicated that higher NO$_3^-$ concentrations from addition of deep water in the middle of the study did not inhibit N$_2$ fixation rates, thus providing more support for challenging the classical paradigm in our understanding of N$_2$ fixation. Nitrogen fixation in our experiment seems to have responded rather to elevated PO$_4^{3-}$ from deep water (or another nutrient such as Fe from the deep water and dust deposition) despite excess NO$_3^-$. Preferential consumption of NO$_3^-$ during the growth phase may have amplified the diazotrophic niche in the decay phase. Unicellular diazotrophic cyanobacterium clade UCYN-B seems to have benefited the most from the deep water addition, while gammaproteobacterial abundances might have been introduced into the mesocosms with the deep water addition. We thus conclude that diazotrophic species and N$_2$ fixation rates respond to elevated $pCO_2$ in a complex way depending upon nutrient conditions. Our results suggest that N$_2$ fixation rates can respond positively to enhanced upwelling but after a time-lag of a few days, as diazotrophs may benefit from increased nutrients (PO$_4^{3-}$ and Fe) availability. Coupling of elevated $pCO_2$ and nutrient exhaustion, as observed in the decay phase and common in the oligotrophic ocean, will also positively impact N$_2$ fixation rates.

**Author contributions**

Arvind Singh and Allanah J. Paul designed incubation experiments, while Lennart T. Bach and Ulf Riebesell designed the mesocosm experiment along with the KOSMOS team. Caroline R. Löschner did the molecular analysis. Narendra Ojha analyzed model AOD data. Arvind Singh wrote the manuscript with inputs from all the coauthors.

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