Distribution of diverse nitrogen fixers in the global ocean

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[1] We employ a global three-dimensional model to simulate diverse phytoplanktonic diazotrophs (nitrogen fixers) in the oceans. In the model, the structure of the marine phytoplankton community self-assembles from a large number of potentially viable physiologies. Amongst them, analogs of Trichodesmium, unicellular diazotrophs and diatom-diazotroph associations (DDA) are successful and abundant. The simulated biogeography and nitrogen fixation rates of the modeled diazotrophs compare favorably with a compilation of published observations, which includes both traditional and molecular measurements of abundance and activity of marine diazotrophs. In the model, the diazotroph analogs occupy warm subtropical and tropical waters, with higher concentrations and nitrogen fixation rates in the tropical Atlantic Ocean and the Arabian Sea/Northern Indian Ocean, and lower values in the tropical and subtropical South Pacific Ocean. The three main diazotroph types typically co-exist in the model, although Trichodesmium analogs dominate the diazotroph population in much of the North and tropical Atlantic Ocean and the Arabian Sea, while unicellular-diazotroph analogs dominate in the South Atlantic, Pacific and Indian oceans. This pattern reflects the relative degree of nutrient limitation by iron or phosphorus. The model suggests in addition that unicellular diazotrophs could add as much new nitrogen to the global ocean as Trichodesmium.

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1. Introduction

[2] Nitrogen fixers or diazotrophs, organisms that fix nitrogen gas (N₂) into organic form, play an important role in the climate system as the availability of inorganic nitrogen can represent a significant limiting factor for marine primary production [Falkowski et al., 1998]. On the global scale, diazotrophy is the largest source of fixed nitrogen in the modern ocean [Galloway et al., 2004; Gruber, 2004], counterbalancing the sinks due to denitrification [Gruber and Sarmiento, 1997; Codispoti et al., 2001] and anammox [Devol, 2003]. It can also have important local consequences: In the oligotrophic North Pacific Ocean, nitrogen fixation is estimated to fuel up to 50% of the new production [Karl et al., 1997].

[3] Diazotrophs in the ocean are represented by a variety of autotrophic and heterotrophic organisms. The diazotrophic heterotrophic community is still poorly understood [Zehr et al., 2000; Langlois et al., 2005], so we focus here on the autotrophs. Three broad categories of marine diazotrophs are associated with photo-autotrophy, which are the cyanobacteria Trichodesmium [Carpenter and Romans, 1991; Capone et al., 1997; Karl et al., 2002; LaRoche and Breitharh, 2005], unicellular cyanobacteria [Zehr et al., 2001], and symbiotic diazotrophs associated with diatoms [Carpenter et al., 1999; Foster and Zehr, 2006].

[4] Much of the current understanding of marine nitrogen fixation comes from the study of Trichodesmium; large, filamentous and non-heterocystous cyanobacteria that often aggregate at the surface of the ocean to form colonies. Trichodesmium has been observed to grow in most warm oligotrophic subtropical and tropical regions of the ocean [Carpenter and Romans, 1991; Capone et al., 1997], apart from the western equatorial and Southeastern subtropical Pacific Ocean [Magie et al., 1974; Bonnet et al., 2008, 2009].

[5] The two other types of marine autotrophic diazotrophs are, as yet, not as well understood. However, it has been suggested that they might add as much new nitrogen to the oceans as Trichodesmium [Carpenter et al., 1999; Zehr et al., 2001]. Unicellular diazotrophs were recently discovered by the association of the nifH gene, encoding for the nitrogen-fixing enzyme nitrogenase, with small cells [Zehr et al., 1998]. Unicellular diazotrophs are between 10 to 1000 times smaller than Trichodesmium (less than 10 μm; Goebel et al. [2007]) but have been observed to have nitrogen fixation rates of similar magnitude to that of Trichodesmium in the tropical North Atlantic Ocean [Falcon et al., 2004]. Observations indicate the presence of unicellular diazotrophs...
in subtropical and tropical waters of the Atlantic Ocean [Zehr et al., 1998; Falcon et al., 2002, 2004; Montoya et al., 2007; Langlois et al., 2008; Foster et al., 2009] and Pacific Ocean [Zehr et al., 2001; Montoya et al., 2004; Church et al., 2005; Goebel et al., 2007; Bonnet et al., 2008; Church et al., 2008], as well as the Arabian Sea [Mazard et al., 2004].

[6] Both Trichodesmium and unicellular diazotrophs have been observed to have a relatively low maximum growth rate, high N:P and Fe:P ratios, and to be adapted to high light conditions and warm temperatures [Falcon et al., 2005; LaRoche and Breitbarth, 2005; Goebel et al., 2008]. These characteristics are probably related to the high energetic cost of breaking down the N₂ triple bond using the enzyme nitrogenase.

[7] Diatom-diazotroph associations (DDA) have been observed in the ocean for some decades though they are still not well understood due to the complexity of the symbiotic relationship [Zehr and Ward, 2002]. Observed DDAs are mostly between Richelia intracellularis autotroph diazotrophs and Hemiaulus hauckii or Rhizosolenia clevei diatoms [Venrick, 1974; Villareal, 1991; Carpenter et al., 1999; Gomez et al., 2005; Foster et al., 2007]. Observations indicate the presence of DDAs in the tropical and subtropical North Pacific Ocean [Venrick, 1974; Gomez et al., 2005; Church et al., 2008; Kitajima et al., 2009], the equatorial North Atlantic Ocean (in particular in the vicinity of the Amazon River plume) [Villareal, 1991; Carpenter et al., 1999; Foster and Zehr, 2006], in the Mediterranean Sea [Zeev et al., 2008] and the China Sea [Gomez et al., 2005].

[8] Ultimately, continued observations are necessary to define global habitats and nitrogen fixing contributions of each type of diazotroph. Models can help to synthesize and extrapolate the existing observations, as well as to explore hypothesized mechanisms of the regulators of community structure and habitat and to evaluate the integrated rate of nitrogen fixation in the global ocean. However, representing the diverse populations of marine microorganisms in models remains a challenge. Several published models include specific representations of a diazotrophic functional group [e.g., Fennel et al., 2002; Hood et al., 2004; Moore and Doney, 2007], typically based on physiological understanding of Trichodesmium. For the first time, Goebel et al. [2007] resolved multiple diazotrophic types in a model, evaluating nitrogen fixation rates based on nifH observations of three diazotroph types (Trichodesmium and unicellular Groups A and B), in a one-dimensional representation of station ALOHA in the subtropical North Pacific Ocean.

[9] Here we introduce diazotrophy as a possible physiological function, associated with a variety of other characteristics, into a three-dimensional ocean circulation, biogeochemistry and ecosystem model in which the photo-autotroph communities “self-assemble” from a broad range of initialized phytoplankton types according to their relative fitness in the modeled environment [Follows et al., 2007]. In previous studies, this approach successfully reproduced the distribution of the diverse community of the cyanobacterium Prochlorococcus in the Atlantic Ocean [Follows et al., 2007], a plausible biogeography of the dominant functional groups of phytoplankton in the global ocean [Dutkiewicz et al., 2009], as well as the latitudinal gradient in marine phytoplankton diversity [Barton et al., 2010], though diazotrophy was not previously represented.

[10] We first describe the physical, biogeochemical, and ecosystem model (section 2). Then, in order to evaluate the veracity of the modeled plankton communities and nitrogen fixation rates, we compile and compare published direct observations of biomass and nitrogen fixation rates for Trichodesmium, unicellular diazotrophs and diatom-diazotroph associations (section 3). We subsequently show that the model system exhibits plausible patterns of abundance and nitrogen fixation rates for the main phytoplanktonic diazotrophs (section 4). This work provides a first cut at a global synthesis of the likely distributions and contributions in fixing nitrogen of three of the main types of marine diazotrophs.

[11] This study focuses on describing the model diazotroph biogeography in the light of a compilation of published observations of diazotroph biomass and nitrogen fixation rates. A deeper investigation into the ecological controls on the emergent model ecosystem is too extensive to include in this manuscript but is the subject of a companion paper (F. M. Monteiro et al., Biogeographical controls on the marine nitrogen fixers, submitted to Global Biogeochemical Cycles, 2010).

2. Model Description

[12] Our modeling framework is based on that of Follows et al. [2007] and Dutkiewicz et al. [2009], where a three-dimensional global ocean model underpins biogeochemical and ecosystem model components with “self-assembling” phytoplankton community structure.

2.1. Ocean Circulation and Biogeochemistry Models

[13] The ocean model (MITgcm; Marshall et al. [1997]) is configured in a relatively coarse resolution (1° × 1° horizontally; 23 vertical levels), with sub-mesoscale eddy fluxes parameterized following the work of Gent and McWilliams [1990] and boundary layer turbulent mixing following that of Large et al. [1994]. We use the ECCO-GODAE state estimates [Wunsch and Heimbach, 2007] in which the model circulation has been constrained to closely match in situ and remote observations of the physical environment.

[14] The model ocean transports inorganic and organic forms of phosphorus (P), nitrogen (N), iron (Fe) and silica (Si). None of the external sources of these elements (atmospheric, river or sediments) are represented in the model, except for nitrogen fixation (see later) and the atmospheric deposition of iron, using the model estimates made by Mahowald et al. [2005]. Here we assume that 0.8% of deposited iron is bioavailable [Mahowald et al., 2009]. The dynamic representation of the marine iron cycle, following Parekh et al. [2005], includes complexation with an organic ligand and particulate scavenging. The biogeochemical and biological tracers interact through the formation, transportation and remineralization of organic matter. Photo-autotrophs are assumed to have fixed elemental ratios and may be consumed by two simple grazers while excretion and mortality, from both phytoplankton and zooplankton, transfer organic matter into sinking particulate and dissolved organic detritus pools. Heterotrophic bacteria are not explicitly
represented. Organic detritus is simply respired back to inorganic forms on a fixed timescale.

[15] Here we also include a more detailed representation of the nitrogen cycling, including explicit representation of nitrogen fixation by diazotrophic phytoplankton (see next paragraph). The combined effect of deep denitrification, anammox and (possibly) heterotrophic diazotrophy is represented in the model very simply, by restoring excess nitrate towards the observed climatology from the World Ocean Atlas 2005 [Garcia et al., 2006] below 200 m depth (details in Appendix A).

2.2. Ecosystem Model

[16] As in the study of Follows et al. [2007], we initialize many tens (here 78) of physiologically unique phytoplankton types with characteristics randomly generated from plausible ranges of possible physiologicals. The randomization places the organism in one of two nominal size classes (“large” or “small”) and assigns parameters regulating the sensitivity of growth to light, nutrients and temperature (Appendix A). Trade-offs are imposed for several characteristics as a function of size: small phytoplankton are presumed to be more efficient at taking up nutrients in oligotrophic conditions, to be able to cope with lower light (due to a lack of package effect), but to be more likely to become photoinhibited and to have slower maximum growth rates. Though, in a general sense growth rate might be expected to decrease with cell size, taxonomic differences can dominate, as seen when comparing laboratory cultures of small, pico-cyanobacteria Prochlorococcus and fast-growing, populations of larger diatom cells [Sarthou et al., 2005]. The high maximum growth rates observed for diatoms may in part reflect an enhanced internal store, which can be larger for bigger cell types [Droop, 1983] though not accounted for explicitly in our model. The model also randomly selects for the nutrient strategy of the phytoplankton types. For example, a “coin-flip” determines whether a cell type requires silicon; if so it is assumed to be a diatom analog. The rate of change of the biomass of each phytoplankton is determined as a combination of growth (which is a function of light, temperature and nutrient resources), sinking, grazing, mortality and transport by the fluid environment (see Dutkiewicz et al. [2009] for full prognostic equations).

[17] Here, we introduce to this framework the possibility for phytoplankton to fix nitrogen gas (N₂). Diazotrophy is also assigned at random and may be associated with any other set of physiological characteristics. The diazotrophs are then assumed to be nitrogen non-limited, to have a smaller specific maximum growth rate (high energetic cost of nitrogen fixing) and higher elemental N:P and Fe:P ratios (high iron requirements of nitrogenase enzyme) than non-diazotrophic phytoplankton (Appendix A). We nominally assume generated diazotrophic cells in the large size class which are not diatoms, to be analogs of Trichodesmium; model diatoms which can fix N₂ to be analogs of diatom-diazotroph associations (DDA); and small size-class cells which can fix nitrogen to be analogs of unicellular, cyanobacterial nitrogen fixers. In all simulations presented here, each of these three main diazotroph types is composed of several (typically 3 to 4) phytoplankton types, which differ slightly in physiology. Here we focus on the total population of each main diazotroph type (e.g. Trichodesmium), that is the sum of any generated phytoplankton that fall into that classification.

[18] As in our previous studies [Follows et al., 2007; Dutkiewicz et al., 2009], two class sizes of grazers are represented in the model, where small (or big) size class preferentially eats small (or big) phytoplankton. For simplicity, here we choose not to have different palatability for the diazotrophs. Though it is probable that Trichodesmium has lower grazing pressure than other phytoplankton [Capone et al., 1997], there is evidence that at least one copepod group feeds on Trichodesmium [O’Neil and Roman, 1994]. More research on grazing pressure would certainly help us to better shape the trade-offs of diazotroph grazers.

[19] Most simulations were integrated for ten model years, after which time a robust seasonal cycle of productivity and phytoplankton community structure has developed and persists for several decades. Over longer timescales the back-ground nutrient distributions drift slightly, driving minor adjustments in biogeography. One simulation was integrated for 100 years and the biogeography, in particular the diazotroph communities, remained relatively constant for that period. Here, all shown results are annual average of the tenth year of simulation. An ensemble of ten simulations was performed, each identical except for the randomization of physiological traits in the initialized organisms. The collection of the ten simulations is the “ensemble” and the individual simulations are the “ensemble members”. The biogeographical patterns that are established in the ensemble members were robust and broadly consistent and here we present the “ensemble mean”; the average of the results from the ten simulations.

[20] The resulting biogeography of the main phytoplankton functional groups is very similar to those found in the studies of Follows et al. [2007] and Dutkiewicz et al. [2009]. Here the model additionally resolves multiple diazotroph types. Though typically at relatively low abundance relative to the main contributors to biomass, the model diazotrophs have a robust distribution which we compare to a compilation of published observations in the following sections.

3. Observations of Diazotroph Biogeography and Nitrogen Fixation

[21] Diazotroph abundance and nitrogen fixation rates have been observed in the ocean for more than 35 years, with early studies related to Trichodesmium [Dugdale et al., 1964] and DDAs [Mague et al., 1974]. Published data on the distribution of Trichodesmium have been compiled for the global ocean [Carpenter, 1983] and the North Atlantic Ocean [Hood et al., 2004]. Here we extend these two compilations to include most currently available observations of occurrence, biomass and nitrogen fixation rates for the three main phytoplanktonic diazotroph types and for the global ocean.

[22] For diazotroph abundance, we compile traditional observations of Trichodesmium and DDA (net tow combined with microscopy), video plankton recorder observations of Trichodesmium and evaluations of the abundance of
the \textit{nifH} gene for \textit{Trichodesmium}, unicellular diazotrophs and DDAs. Measurement of nitrogen fixation rates comes from acetylene reduction assay or isotopic \textit{^{15}N}_{2} methods which have been used to evaluate areal rates (molN m\textsuperscript{-2} d\textsuperscript{-1}) for \textit{Trichodesmium}, unicellular diazotrophs and DDAs. (Some additional published measurements of nitrogen fixation rates are not included here as they integrate the contribution of diazotrophic heterotrophic bacteria which are not represented in the model.)

[23] To compare model results with these observations, we convert the published values into common units (\textmu{}molP m\textsuperscript{-1} for diazotroph biomass and \textmu{}molN m\textsuperscript{-2} d\textsuperscript{-1} for nitrogen fixation rates) and include both the original measurement and estimated conversion in tabular form in the auxiliary material (Tables S1–S8).\textsuperscript{1} The tabulated values include both the uncertainty of the conversion factor and that of the original measurement (e.g. for nitrogen fixation rates, the depth distribution of the diazotroph and the time period of fixation used in the calculation can lead to uncertainties [Lipschultz and Owens, 1996]). Here we outline the method of conversion used for the abundance of each diazotroph type.

3.1. \textit{Trichodesmium}

[24] By far the majority of observations on marine nitrogen fixation have been related to \textit{Trichodesmium}. We convert observations of \textit{Trichodesmium} abundances from filaments, cells, colonies or \textit{nifH} copies into units of phosphorus biomass (\textmu{}molP m\textsuperscript{-1}). To do so, we use average values of field and laboratory measurements of the amount of phosphorus present in the observed entity (see Text S1 and Tables S1 and S2). The resulting compilation shows that \textit{Trichodesmium} observations (Tables 1 and 2) have relatively good coverage in the North Atlantic Ocean, North East subtropical Pacific Ocean and China Sea, but are more sparse in central South Atlantic Ocean, tropical East Pacific Ocean, middle West subtropical Pacific Ocean and East Indian Ocean.

3.2. Unicellular Cyanobacteria Diazotrophs

[25] Measurements of the abundance of unicellular cyanobacteria diazotrophs in the ocean are made using \textit{nifH} gene sequencing methods. Here we convert the number of \textit{nifH} copies into biomass units (\textmu{}molP m\textsuperscript{-1}) for unicellular Groups A and B by assuming 1 \textit{nifH} copy per cell [Goebel et al., 2007]. Following Goebel et al. [2008], since there are no direct measurements of the phosphorus quota in unicellular diazotrophs, we scale \textit{Trichodesmium} phosphorus conversions to the unicellular diazotroph volume, using the allometric relationship for marine phytoplankton developed by Verity et al. [1992] (see auxiliary material, Text S1). This scaling introduces some additional uncertainty.

[26] Goebel et al. [2007] estimated cell volumes of \textit{Trichodesmium}, unicellular Group B and Group A to be between 200–1000 \textmu{}m\textsuperscript{3}, 4–270 \textmu{}m\textsuperscript{3}, and 0.2–4.0 \textmu{}m\textsuperscript{3} respectively. unicellular Group B cell volume is therefore on average two orders of magnitude larger than for unicellular Group A. We sum Group A and Group B biomass to estimate the total populations of unicellulars (Table 3). In most observations of unicellular diazotrophs, \textit{nifH} concentrations are higher for unicellular Group A than for unicellular Group B (see Tables S4 and S5). However the biomass of Group B is generally greater than for unicellular Group A due to the difference in cell volume, thus suggesting a greater contribution to global nitrogen fixation from Group B. Current observations of unicellular diazotrophs in the ocean are mostly restricted to the North Atlantic region, the East subtropical Pacific Ocean and the Arabian Sea (Tables 3 and 4).

3.3. Diatom-Diazotroph Associations

[27] Due to the complexity of symbiotic organisms, the understanding of diatom-diazotroph associations (DDA) remains limited. Detection of DDAs in the ocean relies on specific microscopy (transmitted light and epi-fluorescent) [Carpenter et al., 1999] and \textit{nifH} gene sequencing [Foster and Zehr, 2006]. Since no calibration of DDA \textit{nifH} copies to biomass is available, we do not attempt in this study to compare quantities of abundance between the model and the data, but simply make the distinction between presence or absence of DDAs (Table 5, completed by the sparse observations of DDA nitrogen fixation rate in Table 6). We note though that the model itself provides biomass values for the DDA analogs, presented in Figure 1d.

[28] In the following section we present the model results of the abundance and nitrogen fixing activity of the three major marine photo-autotrophic diazotroph types and compare with the compiled observations for occurrence, biomass and nitrogen fixation rates.

4. Model Results: Diazotroph Biogeography and Nitrogen Fixation

[29] Figure 1 presents the ensemble-average, annual mean biomass (\textmu{}molP m\textsuperscript{-1}) of the total diazotrophs, \textit{Trichodesmium}, unicellulars, and DDAs in the surface waters (0–50 m) for the tenth year of the ensemble simulation. Here we concentrate on the 0–50 m layer as most observations of diazotroph activity are taken in the upper water column. Figure 2 presents the corresponding model result for nitrogen fixation rates (\textmu{}molN m\textsuperscript{-2} d\textsuperscript{-1}). Concentrations and rates are presented on a log scale, as values span large ranges. The emergent diazotroph biogeography is relatively robust between the ten–ensemble-member simulations, illustrated in Figure 3 which displays the standard deviation, relative to the mean, of the simulation ensemble. (The standard deviation of nitrogen fixation rates is shown in Figure S1). Over the ten simulations, biomass and rates are consistent for total, \textit{Trichodesmium} and unicellular-diazotroph analogs which have a low standard deviation relative to the mean in most regions. However the standard deviation, relative to the mean is high for the DDAs indicating a greater role of the randomization of physiological characteristics in regulating the habitat of this diazotroph type. This reflects, in part, our inability to define clear trade-offs and the likelihood that we have over-simplified this complex symbiotic relationship in the model.

\textsuperscript{1}Auxiliary materials are available with the HTML. doi:10.1029/2009GB003731.
Table 1. Comparison Between Model and Observations of Abundance of *Trichodesmium* (µmolP m⁻³) in the Global Ocean: Biomass of *Trichodesmium*¹

| Regions                          | Model (µmolP m⁻³) | Observations (µmolP m⁻³) | References                                                   |
|---------------------------------|------------------|--------------------------|--------------------------------------------------------------|
| A West subtropical North Atlantic | 10⁻⁴-10⁻³        | 10⁻⁴-10⁻³                | Carpenter [1983], Carpenter and Romans [1991],              |
|                                 |                  |                          | Orecutt et al. [2001], LaRoche and Breitbarth [2005],       |
|                                 |                  |                          | Davis and McGillicuddy [2006], and Langlois et al. [2008]  |
| B East subtropical North Atlantic | ≤10⁻³            | 10⁻³-10⁻³                | Tyrrell et al. [2003], Davis and McGillicuddy [2006],       |
|                                 |                  |                          | and Langlois et al. [2008]                                  |
| C Gulf of Mexico                 | 10⁻⁶-10⁻⁴        | 10⁻³                     | Carpenter [1983] and Carpenter and Romans [1991]           |
|                                 |                  |                          | Carpenter [1983], Carpenter and Romans [1991],              |
|                                 |                  |                          | Carpenter et al. [2004], Foster et al. [2007],              |
|                                 |                  |                          | and Langlois et al. [2008]                                  |
| D Caribbean                      | 10⁻⁴-10⁻³        | 10⁻³                     | Carpenter and Price [1977], Carpenter [1983],               |
|                                 |                  |                          | and Carpenter and Romans [1991]                             |
| E West tropical North Atlantic   | 10⁻⁴-10⁻³        | 10⁻²-10⁻²                | Carpenter [1983], Carpenter and Romans [1991], Capone et al. |
|                                 |                  |                          | Carpenter et al. [2004], Foster et al. [2007],              |
| F East tropical North Atlantic   | 10⁻⁴-10⁻³        | 10⁻²-10⁻²                | Carpenter [1983], Vidal et al. [1999], Tyrrell et al. [2003], |
| G West tropical South Atlantic   | 10⁻⁵-10⁻³        | ≤10⁻⁴                    | Carpenter [1983] and Tyrrell et al. [2003]                  |
| H East tropical South Atlantic   | 10⁻⁶-10⁻³        | 10⁻³-10⁻³                | Carpenter [1983] and Tyrrell et al. [2003]                  |
| I West subtropical South Atlantic| 10⁻⁶-10⁻⁴        | 10⁻⁶-10⁻³                | Carpenter [1983] and Tyrrell et al. [2003]                  |
| J East subtropical South Atlantic| ≤10⁻⁶            | 10⁻⁶                     | Carpenter [1983]                                            |
| K East subarctic North Pacific   | <10⁻⁷            | below detection           | Church et al. [2008]                                        |
| L West subtropical North Pacific | 10⁻³-10⁻⁴        | 10⁻²-10⁻⁴                | Carpenter [1983], Chen et al. [2003], Kitaizima et al. [2009], |
| M East subtropical North Pacific | 10⁻⁶-10⁻⁴        | 10⁻²-10⁻⁴                | Shiozaki et al. [2009], Carpenter [1983], Letelier and Karl [1996], |
| N West tropical North Pacific    | 10⁻⁶-10⁻⁵        | 10⁻⁵                     | Carpenter [1983] and Shiozaki et al. [2009]                 |
| O Central tropical Pacific       | ≤10⁻⁶            | ≤10⁻⁴                    | Mague et al. [1974] and Church et al. [2008], 20S-4N, first 10 m |
| P West Equatorial Pacific (coastal)| ≤10⁻⁵            | 10⁻³                     | Mague et al. [1974] and Bonnet et al. [2009]                |
| Q West Equatorial Pacific (open ocean)| ≤10⁻⁷            | below detection           | Dupouy et al. [2000]                                        |
| R West tropical South Pacific (Great Barrier Reef)| <10⁻⁷| 10⁻²-10⁻³                | Church et al. [1991], Chang et al. [2000],                     |
|                                 |                  |                          | Mague et al. [1974] and Bonnet et al. [2008]                |
| S West tropical South Pacific    | <10⁻⁷            | below detection           | Carpenter [1983], Chang et al. [2000], Chen et al. [2003], |
| T East subtropical South Pacific | <10⁻⁷            | below detection           | Wu et al. [2003], Dong et al. [2008]                       |
| U China Sea                      | 10⁻²-10⁻³        | 10⁻²-10⁻³                | Somasundar et al. [1990] and Capone et al. [1998]           |
| V Arabian Sea                    | 10⁻⁴-10⁻³        | 10⁻⁴-10⁻³                | Carpenter [1983]                                            |
| W West tropical North Indian Ocean| 10⁻³             | 10⁻³                     | Carpenter [1983]                                            |
| X West tropical South Indian Ocean| 10⁻⁴-10⁻³        | 10⁻⁴                     | Carpenter [1983]                                            |
| Y West tropical South Indian Ocean| 10⁻⁴-10⁻³        | 10⁻²-10⁻¹                | Carpenter [1983], Carpenter and Romans [1991], Kronkamp et al. [1997], |
| Z Mediterranean Sea              | 10⁻⁴             | 10⁻⁴                     | Carpenter [1983]                                            |

¹The areas are labeled with letters and are reported in Figure 1b. The conversions used for *Trichodesmium* abundances are: P:col = 3.87 × 10⁻⁷ µmolP col⁻¹ [Mague et al., 1977], P:fil = 1.65 × 10⁻⁵ µmolP fil⁻¹ [Letelier and Karl, 1996], cell:fil = 76.6 cell fil⁻¹ [Letelier and Karl, 1996; Sanudo-Wilhelmy et al., 2001; Carpenter et al., 2004; LaRoche and Breitbarth, 2005; Dong et al., 2008], and nifH copies:cell = 1 [Goebel et al., 2007]. Data are detailed in the auxiliary material (Text S1 and Tables S1 and S2) and values have been rounded.

[30] We discuss now the modeled biogeography of diazotrophs both collectively and as separate types, in comparison with their observed distributions.

### 4.1. Sum of Diazotrophs

[31] Although there is no explicit demand that modeled diazotroph analogs grow only in warm waters, the total autotrophic diazotroph population occupies warm subtropical and tropical regions of the ocean (Figure 1a), as is generally observed. While some previous models parameterize their diazotrophic functional types to only grow in warm waters [i.e., Moore et al., 2004, 2006; Breitbarth et al., 2007; Moore and Doney, 2007] or stratification conditions [Hood et al., 2004], here this feature is an emergent result of the “self-assembling” model communities. Cold-adapted diazotrophs were initialized in the model but were selected against due to other factors. Collectively, the modeled diazotrophs occupy warm subtropical and tropical waters (Figure 1a for biomass and Figure 2a for nitrogen fixation rate), with higher concentrations and nitrogen fixation rates...
in the tropical Atlantic Ocean, the Arabian Sea, and the Northern Indian Ocean (≥10^{-2} \mu molP \text{ m}^{-3} \text{ d}^{-1} for biomass, and \geq 10^{2} \mu molN \text{ m}^{-2} \text{ d}^{-1} for nitrogen fixation rates). Much of the central (sub)tropical South Pacific Ocean has very low diazotroph activity. The low iron concentrations there (Figure 5d) and the high requirement for iron imposed on the model diazotrophs limit diazotroph growth in this region of the Pacific Ocean.

### 4.2. Trichodesmium Analogs

[32] The modeled habitat and nitrogen fixation rate of *Trichodesmium* are compared with observations on a regional basis (Tables 1 and 2). Each region is designated by a letter (see Tables 1–6) indicated on the map of *Trichodesmium* surface biomass (Figure 1b) and nitrogen fixation rate (Figure 2b). Both observed and modeled *Trichodesmium* biomass concentration and nitrogen fixation rates

| Regions | Model (\mu molP \text{ m}^{-3}) | Observations (\mu molP \text{ m}^{-3}) |
|---------|---------------------------------|----------------------------------|
| A West subtropical North Atlantic | 10^{-2} \text{ to } 10^{-4} | 10^{-1} |
| B East subtropical North Atlantic | \leq 10^{-4} | 10^{-1} |
| C West tropical North Atlantic | 10^{-5} \text{ to } 10^{-3} | 10^{-5} \text{ to } 10^{-3} |
| D East tropical North Atlantic | 10^{-5} \text{ to } 10^{-3} | 10^{-3} \text{ to } 10^{-5} |
| E Arabian Sea | 10^{-2} | 10^{-2} |
| F East tropical North Atlantic | 10^{-4} \text{ to } 10^{-2} | 10^{-2} |
| G West tropical South Atlantic | 10^{-6} \text{ to } 10^{-4} | 10^{-6} |
| H East subtropical/subtropical North Pacific | 10^{-5} \text{ to } 10^{-3} | 10^{-5} \text{ to } 10^{-3} |
| I Central tropical South Pacific | 10^{-6} | 10^{-6} |
| J West Equatorial Pacific | 10^{-7} | 10^{-7} |
| K East tropical South Atlantic | 10^{-7} | 10^{-7} |
| L 26° S 246° | 10^{-7} \text{ to } 10^{-6} | 10^{-7} |
| M Arabian Sea | 10^{-2} | 10^{-2} |

The areas are labeled with letters and are reported in Figure 1c. We assume a ratio for nifH copies:cell = 1 [Goebel et al., 2007] and use the allometric relationship by Verity et al. [1992] to convert phosphorus cell content from *Trichodesmium* estimate to unicellular Groups A and B. Data are detailed in the auxiliary material (Text S1 and Tables S4, S5, and S6) and values have been rounded.
vary by an order of magnitude or more even within regions (Tables 1 and 2). The large observed variability within regions may, in part, be due to seasonality, eddies, and sparse sampling. 

[33] The large scale patterns in modeled *Trichodesmium* biomass and nitrogen fixation rates is generally consistent with those observed: High *Trichodesmium* concentrations (on the order of $10^{-5}$ $\mu$molP l$^{-1}$, equivalent to $10^5$ fil m$^{-3}$) and nitrogen fixation rates (from $10^{-1}$ to $10^{-2}$ $\mu$molN m$^{-2}$ d$^{-1}$) occur in the tropical Atlantic Ocean, the Northern Indian Ocean and Arabian Sea. Somewhat lower values are found in the subtropical North Atlantic Ocean (concentrations around $10^{-5}$ to $10^{-4}$ $\mu$molP l$^{-1}$ and rates of $10^{-1}$ to $10^{-2}$ $\mu$molN m$^{-2}$ d$^{-1}$). In the subtropical South Atlantic Ocean, model and observations show a sharp gradient in concentrations from about $10^{-4}$ to $10^{-2}$ $\mu$molP l$^{-1}$ in the West, to almost nothing in the East of the basin. Low values characterize most of the North Pacific Ocean in both the model and the observations with about $10^{-5}$ $\mu$molP l$^{-1}$ of biomass (equivalent to $10^2$ fil m$^{-3}$) and 1 $\mu$molN m$^{-2}$ d$^{-1}$ of nitrogen fixation rate. Finally the model does not produce any *Trichodesmium* in the East South Pacific: Cruises in this region (Q and T) also did not find any *Trichodesmium* [Mague et al., 1974; Bonnet et al., 2008, 2009].

[34] A few regions in the model have lower biomass of *Trichodesmium* than what is observed. These include the Gulf of Mexico (C), some areas in the Western Pacific Ocean (L, P, R, S) and the Western part of the Indian Ocean (Y). Some of these (C, R, Y) are coastal regimes where tidal mixing, riverine inputs, and other influences unresolved in the model may be important. The modeled Western tropical and subtropical Pacific (L, P, R, S) have very low iron concentrations, potentially due to the lack of parameterization of sediment supply, which might also lead to unrealistically low diazotrophic activity. The Western tropical Indian Ocean (X) and the Mediterranean Sea (Z) have biomass of *Trichodesmium* which are too high in the model. Finally, the modeled nitrogen fixation rates are much lower than observed in the East subtropical North Pacific (M), though the modeled biomass concentrations in the region is similar to the observations. Similarly in the Central tropical Pacific Ocean (region Ø), the model reproduces observations of nitrogen fixation rates but underestimates *Trichodesmium* biomass. These discrepancies between biomass and rate of nitrogen fixers are not understood but might reveal an interesting dynamics present in the real ocean, where *Trichodesmium* fix more (or less) nitrogen than they actually need in the East subtropical North (or Central tropical) Pacific Ocean.

[35] Estimates of the occurrence of *Trichodesmium* blooms based on remote sensing [Westberry and Siegel, 2006] found high bloom concentrations in the tropical Atlantic Ocean, the Arabian Sea and the Western Indian Ocean; supporting the modeled distribution presented here. *Westberry and Siegel* [2006] also inferred high *Trichodesmium* concentrations in the Eastern tropical Pacific Ocean consistent with inferences

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**Table 4.** Comparison Between Model and Observations of Nitrogen Fixation Rates of Unicellular Diazotrophs ($\mu$molN m$^{-2}$ d$^{-1}$) in the Global Ocean: Nitrogen Fixation Rate of Unicellular Diazotrophs

| Regions | Model (\(\mu\text{molN m}^{-2} \text{ d}^{-1}\)) | Observations (\(\mu\text{molN m}^{-2} \text{ d}^{-1}\)) | References |
|---------|---------------------------------------------|---------------------------------------------|------------|
| a       | West subtropical North Atlantic             | $10^{-1} - 10^{1}$                         | $1 - 10^1$ | Falcon et al. [2004] and Foster et al. [2007] |
| b       | East subtropical/subtropical North Pacific  | $10^{2} - 10^{4}$                         | $10^1$     | Montoya et al. [2004] |
| c       | East tropical North Pacific                 | $1 - 10^1$                                 | $1 - 10^2$ | Zehr et al. [2001], Falcon et al. [2004], and Montoya et al. [2004] |
| d       | East tropical Indian Ocean (North Australia)| $10^4$                                     | $10^2$     | Montoya et al. [2004] |

*The areas are labeled with letters and are reported in Figure 2c. Data are detailed in the auxiliary material (Text S1 and Table S7) and values have been rounded.

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**Table 5.** Comparison Between Model and Observations of Occurrence of Diatom-Diazotroph Associations (DDA) in the Global Ocean: Occurrence of Diatom-Diazotroph Associations

| Regions | Model | Observations | References |
|---------|-------|--------------|------------|
| α       | Caribbean | present | present | Villareal [1991] |
| β       | West tropical North Atlantic | present | present | Carpenter et al. [1999] and Foster et al. [2007] |
| γ       | East tropical Atlantic | present | present | Foster et al. [2009] |
| δ       | East subtropical North Pacific | none | below detection | Verriek [1974] and Church et al. [2008] |
| ε       | West subtropical North Pacific | present | present | Gomez et al. [2005] and Kitajima et al. [2009] |
| ζ       | East subtropical North Pacific | present | present | Mague et al. [1974], Verriek [1974], and Church et al. [2005, 2008, 2009] |
| η       | West tropical North Pacific | present | present | Kitajima et al. [2009] |
| θ       | Central tropical North Pacific | none-presents | none-presents | Mague et al. [1974] and Church et al. [2008] |
| i       | West Equatorial Pacific | none | below detection | Bonnet et al. [2009] |
| κ       | East South Pacific | none | below detection | Bonnet et al. [2008] |
| λ       | 26°S 246° | none | present | Bonnet et al. [2008] |
| μ       | South China Sea | present | present | Gomez et al. [2005] |
| ν       | East Mediterranean Sea | none-presents | present | Zeev et al. [2008] |

*The areas are labeled with letters and are reported in Figure 1d.
based on geochemical diagnostics [Deutsch et al., 2007]. Here, however, our model solutions show almost no presence of *Trichodesmium*. It is difficult to reconcile the results of both studies by Deutsch et al. [2007] and Westberry and Siegel [2006] to the direct in situ observations, as so far very little autotrophic diazotrophy has been observed in these regions [Mague et al., 1974; Bonnet et al., 2008, 2009], and seems to support our model solutions.

[36] The model captures the seasonal cycle of *Trichodesmium* at the Bermuda Atlantic Time-series Station, with a notable late summer bloom (Figure 4, shown here for a single ensemble member). *Trichodesmium* biomass varies between about $2 \times 10^{-5} \, \mu \text{molP l}^{-1}$ in the late Spring to about $13 \times 10^{-5} \, \mu \text{molP l}^{-1}$ in the late summer, consistent with time-series observations [Orcutt et al., 2001]. The model predicts a similar, late summer bloom of *Trichodesmium* in the subtropical North Pacific Ocean, as observed at the Hawaii Ocean Time-Series [Goebel et al., 2007; Church et al., 2009]. Here though the seasonality is captured, the modeled abundances are gradually drifting due to a long term trend in the modeled iron concentrations in this region.

### 4.3. Unicellular Diazotroph Analogs

[37] Both model and observations suggest high concentrations of unicellulars (about $10^{-3} \, \mu \text{molP l}^{-1}$), with corresponding high nitrogen fixation rates, in most of the East North Atlantic Ocean and Arabian Sea/Indian Ocean (Figure 1c and Table 3 for concentrations, and Figure 2c and Table 4 for rates). Both model and observations suggest lower concentrations and fixation rates in the East subtropical North Pacific (about $10^{-4} \, \mu \text{molP l}^{-1}$ and $10^3 \, \mu \text{molN m}^{-2} \text{d}^{-1}$ respectively) as well as no unicellular diazotroph activity in the East South Pacific, except for a small area in the middle of the subtropical gyre.

[38] The model and observations only differ significantly in the Western subtropical North Atlantic Ocean (A). Here however, the observations provide a mixed perspective with a measured rate of nitrogen fixation of the same order of magnitude as in the North Pacific [Falcon et al., 2004; Foster et al., 2007], consistent with the model solution. In contrast, another cruise found no evidence of unicellular diazotroph abundance from *nifH* analysis [Langlois et al., 2005], whereas the model supports a population of unicellular diazotrophs in agreement with the modeled fixation rate.

[39] Here we treat Group A and Group B both as strict autotrophs. However recent discoveries about the physiology of unicellular Group A indicate that they must rely on organic carbon as they lack the PSII oxygenic photosynthetic apparatus [Zehr et al., 2008] and may participate in a symbiotic relationship [Tripp et al., 2010]. In the current model, we do not clearly represent Group A and their rather special physiology and ecology. As understanding increases, future models may need to treat these organisms differently, with potentially significant implications for overall nitrogen fixation rates.

### 4.4. Diatom-Diazotroph Association Analogs

[40] Since we are unable to convert the biomass observations of DDAs to model units, we compare patterns of presence or absence of the organisms (Figure 1d and Table 5). DDAs have been observed in the Western side of the (sub)tropical North Atlantic Ocean, the Eastern subtropical and Western tropical North Pacific, the Kuroshio current and the Eastern Mediterranean Sea. The model solutions are consistent except in the Eastern subtropical North Pacific and the Mediterranean Sea where there are no model DDAs. In the model, DDAs are most abundant in the Arabian Sea and the Indian Ocean, as well as the western tropical Atlantic Ocean. The nitrogen fixation rates of modeled DDAs are typically too low at any locations for which there are measurements (Figure 2d and Table 6), perhaps indicating that the parameterization of the symbiotic relationship of a DDA is too simplistic.

[41] Overall, the model captures the main features of the observed habitats for the three diazotrophy types and is broadly consistent with observed abundances and rates for *Trichodesmium* and unicellular diazotrophs. The model might underestimate nitrogen fixation by DDAs globally and by *Trichodesmium* in areas of the Western Pacific Ocean.

### 5. Discussion

#### 5.1. Biogeography of Marine Diazotrophs

[42] The three main diazotrophic types co-exist over much of the modeled subtropical ocean (Figure 5a). Unicellular-diazotroph analogs tend to dominate in the model in most of the regions inhabited by diazotrophs (Figure 5b), especially where iron is limiting (Figures 5c and 5d). The lower nutrient half-saturation constant, a consequence of their smaller size, confers in these regions an advantage on the unicellular diazotrophs. Unicellular-diazotroph analogs also dominate the diazotroph population in a small region of the equatorial Atlantic Ocean, which is characterized by low phosphate concentration rather than low iron concentration (Figures 5c and 5e). When the nutrient limitation is lifted,
Trichodesmium analogs are no longer disadvantaged. Trichodesmium dominate particularly in regions where iron is more plentiful (subtropical Atlantic Ocean and Arabian Sea). DDAs are found in the model to dominate only in small regions, though we note that the simulated distribution of DDA analogs is quite variable between ensemble members and we are less confident of these predictions.

[43] Trichodesmium analogs in our model are responsible for about 43% of the global total nitrogen fixation rate (25 TgN yr⁻¹; section 5.2), while unicellular-diazotroph analogs are responsible for about 49% (30 TgN yr⁻¹) of the global rate. Consistent with previous suggestions [Carpenter et al., 1999; Zehr et al., 2001], our global ocean model explicitly indicates that unicellular diazotrophs could be as important in the global nitrogen budget as Trichodesmium. DDA analogs contribute to about 8% of the global nitrogen fixation rate (5 TgN yr⁻¹), though we are not sufficiently comfortable with the parameterization of this symbiotic

![Figure 1.](image-url)

Annual average concentration of diazotroph analogs generated in the model (μmolP L⁻¹, log scale, first 50 m depth) for an ensemble of ten model simulations, each initialized with a different, randomized set of phytoplankton physiologies. Regions of published observations are indicated by letters and ellipses. A solid ellipse denotes observed presence of the organisms of interest. A dashed ellipse denotes that observations were made but the organism of interest was not seen. (a) Total diazotroph abundance. (b) Trichodesmium analogs. Letters represent regions with observations of Trichodesmium abundance, compared with model results in Table 1. (c) Unicellular-diazotroph analogs. Letters represent regions with observations of unicellular abundance, compared with model results in Table 3. (d) Diatom-diazotroph-association (DDA) analogs. Letters represent regions with observations of DDA occurrence, compared the model results in Table 5.
relationship and the robustness of their distribution in the ensemble to be confident of this result.

5.2. Nitrogen Fixation Rates

[44] The global nitrogen fixation rate of our ensemble average is 60 ± 15 TgN yr⁻¹ (where the uncertainty is the standard deviation of the ten ensemble members), within the range of published direct and indirect estimates, which vary from 5 to 135 TgN yr⁻¹.

[45] Direct in situ biological estimates increased from early estimates of 5–10 TgN yr⁻¹ [Capone and Carpenter, 1982; Carpenter et al., 1992] to more recent estimates of

Figure 2. Annual average nitrogen fixation rate of diazotroph analogs generated in the model (μmolN m⁻² d⁻¹, log scale, first 50 m depth) for an ensemble of ten model simulations, each initialized with a different, randomized set of phytoplankton physiologies. Regions of published observations are indicated by letters and ellipses. A solid ellipse denotes observed rates of the organisms of interest. A dashed ellipse denotes that observations were made but rates of the organism of interest were undetectable. (a) Total diazotrophs, representing a global nitrogen fixation average of 60 TgN yr⁻¹. (b) Trichodesmium analogs, representing 43% of global nitrogen fixation rate. Letters correspond to regions with observations of Trichodesmium rates, compared with the model results in Table 2. (c) Unicellular-diazotroph analogs, fixing 49% of the global nitrogen fixation rate. Letters represent regions with observations of unicellular diazotroph rates, compared with the model results in Table 4. (d) Diatom-diazotroph association (DDA) analogs, fixing 8% of the global nitrogen fixation rate. Letters represent regions with observations of DDA rates, compared with the model results in Table 6.
However these have accounted only for the contribution of *Trichodesmium*, extrapolating sparse measurements to a basin scale or global value for warm oligotrophic regions [Capone et al., 1997; Galloway et al., 2004]. Our model results (and we believe the data compilation) suggest that there are large regional variations in nitrogen fixation even in these warm oligotrophic waters. Thus we suggest that direct estimations of *Trichodesmium* activity [Capone et al., 1997; Galloway et al., 2004] extrapolated from a few measurements have large uncertainties. Our model suggests that *Trichodesmium* only contributes 25 TgN yr\(^{-1}\), though we underestimate *Trichodesmium* nitrogen fixation rates in the Western Pacific Ocean (where iron limitation is too strong) and in coastal zones.

[46] Geochemically based estimates of nitrogen fixation, relying on interpretations of N* and DIN\(_{waq}\), reflect contributions from the entire diazotrophic community (including heterotrophic nitrogen fixers). Geochemical estimates range between 110 and 135 TgN yr\(^{-1}\) globally, subject to significant uncertainties [Gruber and Sarmiento, 1997; Gruber, 2004; Hansell et al., 2004; Landolfi et al., 2008; Deutsch et al., 2007]. The difference with our model result might be in part due to the activity of diazotrophic heterotrophic bacteria which is not resolved here.

[47] Published biogeochemical models have estimated global nitrogen fixation rates ranging from 55 to 100 TgN yr\(^{-1}\) [Moore et al., 2002, 2004, 2006] with a clear relationship between rate of iron supply to surface ocean and global nitrogen fixation. Increasing dust-borne iron solu-
ability to 4% in our model induces a global nitrogen fixation rate of 80 ± 30 TgN yr⁻¹ but at the expense of fidelity in comparison to the observed data of diazotroph activity. The distribution of the combined diazotroph types in this model is similar to that in previous model studies [Moore et al., 2004; Moore and Doney, 2007], which represented a single, generic marine diazotroph based on Trichodesmium. Therefore it is possible that the first order global biogeochemical impact of autotrophic nitrogen fixers in the ocean can be captured without the level of diversity modeled in this study. The degree of nitrogen fixation may be rather dictated by the relative shortfall of nutrients (e.g. the available amount of iron and phosphate to the diazotrophs). However the spatial extent of the diazotroph habitat, the stability of the biogeochemical nitrogen system and response to variability and climatic changes are probably all affected by the level of the captured ecological resolution.

[48] The ensemble-mean nitrogen fixation rate over the North Atlantic basin (0°–60°N; 285°W–360°) is 12 ± 4 TgN yr⁻¹. Previous studies have suggested nitrogen fixation rate in the North Atlantic Ocean to vary between 1 and 35 TgN yr⁻¹ for direct biological estimates [Capone and Carpenter, 1982; Lipschultz and Owens, 1996; Capone et al., 2005], between 4 and 85 TgN yr⁻¹ for geochemical estimates [Michaels et al., 1996; Gruber and Sarmiento, 1997; Hansell et al., 2004; Deutsch et al., 2007; Hansell et al., 2007; Landolfi et al., 2008] and between 8 and 30 TgN yr⁻¹ for model estimates [Coles et al., 2004]. Though the North Atlantic Ocean is the most studied region for nitrogen fixation, the large range of estimates indicates the uncertainties associated with all these techniques. The latest and most accurate geochemical study by Landolfi et al. [2008] estimated nitrogen fixation rate for the North Atlantic region of 15 TgN yr⁻¹, which is within the standard deviation range of our model ensemble values.

6. Concluding Remarks

[48] We coupled a three-dimensional ocean model with a biogeochemical and ecosystem parameterization to simulate the diverse population of diazotrophic phytoplankton in the global ocean. In this “self-assembling” ecosystem model, analogs of Trichodesmium, unicellular diazotrophs and diatom-diazotroph associations (DDA) are enabled and are successful and abundant in each of the ten members of the ensemble of simulations. In addition, we compiled currently available observations of biomass and nitrogen fixation rates for Trichodesmium, unicellular diazotrophs and DDAs in order to evaluate the veracity of the model solutions.

[50] The simulations capture the broad character of the observed distribution of Trichodesmium, unicellular diazotrophs and DDAs, as well as the abundances and nitrogen fixation rates of Trichodesmium and unicellular diazotrophs. They suggest that the three main types of photo-autotrophic diazotrophs co-exist over most of the warm subtropical and tropical waters of the ocean with higher abundances and rates in the tropical North Atlantic Ocean and the Northern Indian Ocean, consistent with the data compilation, diagnostic studies and other models. In this model, the confinement of diazotrophs to warm oligotrophic waters is not imposed but is a result of selection from a broad variety of initialized diazotroph analogs including some adapted to cold temperatures, suggesting that other factors (resource availability) are more significant in defining the habitat. In the Eastern South Pacific Ocean, the model suggests an absence of diazotrophs attributable to iron limitation. This finding is at odds with some other models and diagnostic studies [Westberry and Siegel, 2006; Deutsch et al., 2007] but appears to be supported by observations [Mague et al., 1974; Bonnet et al., 2008, 2009]. Diazotroph activity is underestimated in our model in the Western Pacific Ocean because iron limitation is too strong (perhaps due to a lack of sedimentary input) and in the unresolved coastal zones. Our globally integrated nitrogen fixation rate (60 ± 15 TgN yr⁻¹) is about half that estimated by geochemical tracers. This may reflect, in part, the fact that we do not resolve heterotrophic diazotrophs and that some regions are too strongly iron limited. An inclusion of sedimentary iron sources and variable iron solubility may help for the latter issue.

[51] In the model, Trichodesmium analogs dominate the total diazotroph population in most of the North and tropical Atlantic Ocean and in the Arabian Sea, whereas unicellular-diazotroph analogs dominate in the South Atlantic, Pacific and Indian oceans where iron or phosphorus are less plentiful. The model predicts in particular a high level of nitrogen fixation by unicellular diazotrophs in the Indian Ocean. Observations are still too sparse to support or refute this result. Finally, the model suggests that unicellular diazotrophs globally contribute to add as much new nitrogen to the ocean as Trichodesmium, supporting Zehr et al. [2001] hypothesis on the importance of small diazotrophs in the
Figure 5. Diversity, dominance and nutrient limitations of diazotroph analogs in the model. (a) Diversity of diazotroph analogs (between *Trichodesmium*, unicellular diazotrophs and diatom–diazotroph associations (DDA)). (b) Dominant diazotroph analogs, calculated from the 0–50 m abundance shown in Figure 1. (c) Nutrient limitations between phosphate and iron for the total population of diazotroph analogs. (d) Modeled total iron concentration (nmol Fe L\(^{-1}\), log scale) between 0−50 m. (e) Modeled phosphate concentration (μmol P L\(^{-1}\), log scale) between 0−50 m.
marine nitrogen cycle. Thus, while *Trichodesmium* may still be the most observed diazotrophic phytoplankton, any attempt to estimate total global nitrogen fixation rates should include the contribution from other diverse diazotrophic communities. Much remains to be explored as new observations improve our understanding of these different nitrogen fixers and suggest better parameterizations in our model.

**Appendix A: Model Ecosystem Parametrization**

[52] The ecosystem model equations are similar to that used in the studies of Follows et al. [2007] and Dutkiewicz et al. [2009]. We use the exact same equations and parameter values as the multiple resource case experiment of Dutkiewicz et al. [2009], which are provided in their Appendix. Here we added changes for the nitrogen cycle to include the source of nitrogen fixation and the sink of denitrification (detailed below). We also used a smaller Fe solubility constant (α_{Fe} = 0.008), ideal to represent the distribution of *Trichodesmium* in the North Atlantic Ocean model, and in the range of the observations that vary between 0.01% and 80%, with most values in the lower range [Mahowald et al., 2009].

### A1. Nitrogen Fixation

[53] Nitrogen fixation in our model is represented by several autotrophic diazotroph types. When a phytoplankton is generated to be a diazotroph (Diaiz), we assume the growth to be nitrogen non-limited and to have a smaller maximum growth rate (\( \mu_{\text{max}}^{\text{Diaiz}} = 1/2 \mu_{\text{max}} \)) as well as higher N:P and Fe:P elemental ratios (N:P_{Diaiz} = 40 mol: mol; Fe:P_{Diaiz} = 3 \times 10^{-2} \text{ mol: mol}; see Text S1 in auxiliary material). Sensitivity studies were carried out in the model testing different values of \( \alpha_{Fe} \) and Fe:P_{Diaiz}. We used those values that provided the best match to the observations of diazotrophs. Diazotroph abundances are observed on average to be ten times smaller that other phytoplankton [LaRoche and Breitbarth, 2005]. In our model we initialize diazotroph concentrations to be ten times less than for other phytoplankton; this avoids a large, unrealistic spike of new nitrogen at the beginning of the simulation and therefore reduces the computational time to reach steady state. In addition, we assume a half-saturation constant for unicellular-diazotroph analogs to vary between 0.012 and 0.017 \( \mu\text{mol} \text{P} \text{ L}^{-1} \), which is between the range of Prochlorococcus analogs and other small organisms (such as *Synechococcus*). This is because we assume the half-saturation constant to be size-dependent and that unicellular diazotrophs are smaller than most small phytoplankton, and slightly bigger than Prochlorococcus.

### A2. Nitrate Restoring Scheme

[54] We restore in the model the concentrations of nitrate to the observations (World Ocean Atlas 2005, Garcia et al. [2006]) everywhere below 200 m, but only if the concentration exceeds the observation. This provides a crude representation of the combined effect of denitrification, anammox and (possibly) heterotrophic nitrogen fixation in the deep ocean. The model expression for the nitrate restoring scheme is equivalent to:

\[
\left( \frac{d\text{NO}_3^-}{dt} \right)_{\text{rest}} = \frac{1}{\tau_{\text{rest}}} (\text{NO}_3^- - \text{NO}_3^-_{\text{rest}}),
\]

\( \forall \text{ NO}_3^- > \text{NO}_3^-_{\text{rest}} \) & \( z > 200 \text{m} \)

[55] \( \tau_{\text{rest}} \) is the restoring time scale, set to 1 year for our presented results. Sensitivity studies reveal that varying \( \tau_{\text{rest}} \) from one year to a decade shows very similar distributions of diazotrophs in the global ocean model.

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