A quantitative model of nitrogen fixation in the presence of ammonium
Inomura, Keisuke; Bragg, Jason; Riemann, Lasse; Follows, Michael J.

*Published in:*  
P L o S One

*DOI:*  
10.1371/journal.pone.0208282

*Publication date:*  
2018

*Document Version*  
Publisher's PDF, also known as Version of record

*Citation for published version (APA):*  
A quantitative model of nitrogen fixation in the presence of ammonium

Keisuke Inomura1*, Jason Bragg2, Lasse Riemann3, Michael J. Follows1

1 Department of Earth, Atmospheric and Planetary Sciences, Massachusetts Institute of Technology, Cambridge, MA, United States of America, 2 National Herbarium of New South Wales, The Royal Botanic Gardens and Domain Trust, Sydney, NSW, Australia, 3 Marine Biological Section, Department of Biology, University of Copenhagen, Helsingør, Denmark

* Current address: School of Oceanography, University of Washington, Seattle, WA, United States of America

* ki24@uw.edu

Abstract

Nitrogen fixation provides bioavailable nitrogen, supporting global ecosystems and influencing global cycles of other elements. It provides an additional source of nitrogen to organisms at a cost of lower growth efficiency, largely due to respiratory control of intra-cellular oxygen. Nitrogen-fixing bacteria can, however, utilize both dinitrogen gas and fixed nitrogen, decreasing energetic costs. Here we present an idealized metabolic model of the heterotrophic nitrogen fixer Azotobacter vinelandii which, constrained by laboratory data, provides quantitative predictions for conditions under which the organism uses either ammonium or nitrogen fixation, or both, as a function of the relative supply rates of carbohydrate, fixed nitrogen as well as the ambient oxygen concentration. The model reveals that the organism respires carbohydrate in excess of energetic requirements even when nitrogen fixation is inhibited and respiratory protection is not essential. The use of multiple nitrogen source expands the potential niche and range for nitrogen fixation. The model provides a quantitative framework which can be employed in ecosystem and biogeochemistry models.

Introduction

Nitrogen fixation, the conversion of dinitrogen gas to bioavailable nitrogen, is carried out by diverse prokaryotes termed diazotrophs. It plays a fundamental role in maintaining the productivity of aquatic and terrestrial environments both locally and globally. The development of quantitative, mechanistic models with which to interpret and predict the biogeography and rate of nitrogen fixation and their relationship to environmental factors is a significant challenge for carbon cycle and climate modeling [1–4].

Diazotrophy in the presence of fixed nitrogen

The ability to fix nitrogen provides a clear ecological advantage in oligotrophic environments where nitrogen is limiting. This is at a cost of lower growth efficiency than an organism using fixed nitrogen [5] due to the direct cost of reducing N₂ as well as the indirect cost of managing
intra-cellular oxygen, which de-activates nitrogenase [6–10]. Strategies to manage oxygen include enhanced respiration [7,11–13], specialized heterocyst cells, and (in photo-autotrophic diazotrophs) temporal separation of nitrogen and carbon fixation [14–17]. The presence of reduced nitrogen also impacts nitrogen fixation. On the one hand, ammonium or nitrate offer a “cheaper” source of nitrogen for an organism [18] and the presence of fixed nitrogen can inhibit nitrogen fixation [19,20]. On the other hand, there is significant evidence showing that diazotrophs can and do often use both fixed nitrogen and dinitrogen [21].

Uptake of ammonium reduces the cell’s electron potential, which may inhibit the flow of reducing equivalents to nitrogenase [19]. High intracellular concentrations of ammonium have been observed to cause downregulation of nif genes and decrease synthesis of the nitrogenase complex [20], presumably because preferential consumption of fixed nitrogen leads to a more efficient growth. Despite this, active nitrogen fixation is observed in pelagic, mesopelagic and benthic marine environments where fixed nitrogen is present [21–23] and in marine waters, shelf-sediments and salt marshes where it is abundant [21,24–30]. In an extensive review [21], compiled evidence from diverse marine environments showing that, while sustained high concentrations of fixed nitrogen lead to the suppression of N$_2$ fixation, it does not always exclude it. In benthic environments, nitrogen fixation occurs at significant rates at fixed-nitrogen concentrations exceeding 100 $\mu$M [31]. Nitrogen fixing organisms do utilize fixed nitrogen sources. For example, Trichodesmium, a marine, phototrophic diazotroph has been shown to assimilate nitrate in laboratory studies [21,32,33] and heterotrophic diazotrophs are seen to switch between, or simultaneously use, both dinitrogen and ammonium [32,34,35].

**Significance for ecological and biogeochemical models**

Major current models of nitrogen fixation in the marine environment assumes that diazotrophs never utilize fixed nitrogen [36–39]. The ability to use fixed nitrogen extends the viable range for nitrogen fixation beyond highly nitrogen starved environments. It is observed to occur in nature, related to the availability of other substrates including phosphorus and organic carbon [21]. In order to appropriately capture the flexible nitrogen use of diazotrophs in ecological and biogeochemical models, enabling accurate evaluations of the biogeography and integrated rates of nitrogen fixation, we need a framework to predict when it is advantageous to a cell to fix nitrogen, use fixed nitrogen, or a combination of both.

As a step towards developing a quantitative, prognostic framework for modeling nitrogen fixation in diverse environments, here we bring together a quantitative model of the nitrogen fixing bacterium *Azotobacter vinelandii* [40] and published data from laboratory studies [34,41] in which it was grown in continuous culture with a sucrose and ammonium-based medium. We choose to develop the model around this system because *Azotobacter vinelandii* is a well-studied model organism (studied over a century [42]) and because the laboratory studies [34,41] are sufficient to quantitatively test and constrain a dynamic model of an organism using both N$_2$ and ammonium under a variety of environmental conditions.

**Key findings from laboratory studies**

In the laboratory, the expression and rate of nitrogen fixation by the heterotrophic diazotroph *Azotobacter vinelandii* were examined as a function of relative rates of supply of organic carbon and fixed nitrogen [34]. They quantified this by the molar ratio of sucrose to ammonium in the incoming medium, C/N, where C/N = 1 is equivalent to the elemental ratio C:N = 12 (Table 1). Below a threshold C/N, nitrogen fixation was absent. Above the threshold, nitrogen fixation occurred at a rate which increased with C/N and eventually saturated (Fig 1). The threshold C/N increased and the nitrogen fixation rate decreased as the oxygen concentration...
increased [34,41]. Below the threshold, the organism’s sole nitrogen source was ammonium. At higher C/N, carbon in excess of the requirements to assimilate the NH\(_4^+\) was used to deplete intra-cellular oxygen and fuel nitrogen fixation.

The aim of this study is to develop and apply a framework for interpreting and quantifying the cost and rates of nitrogen fixation in the presence of fixed nitrogen. In the Methods section we describe the modeling framework; further details are given in the supplementary material. In the Results section we simulate and interpret the nitrogen assimilation strategy of the heterotrophic diazotroph \textit{Azotobacter vinelandii} under a variety of environmental conditions. In the Discussion, we discuss the implications for the conditions under which heterotrophic nitrogen fixation may be viable in real world environments.

### Table 1. Nomenclature of the used symbols in this main text.

They are listed roughly in the order of appearance.

<table>
<thead>
<tr>
<th>Symbol</th>
<th>Definition</th>
<th>Unit</th>
</tr>
</thead>
<tbody>
<tr>
<td>C/N</td>
<td>Sucrose to ammonium ratio</td>
<td>mol sucrose mol N(^{-1})</td>
</tr>
<tr>
<td>C:N</td>
<td>Carbohydrate (in carbon) to ammonium ratio; C:N = 12C/N</td>
<td>mol C mol N(^{-1})</td>
</tr>
<tr>
<td>(R_{C/N}^f)</td>
<td>Lowest (R_{C/N}) at which nitrogen fixation is detected</td>
<td>mol sucrose mol N(^{-1})</td>
</tr>
<tr>
<td>(f_{N2})</td>
<td>The ratio of nitrogen fixation to the entire nitrogen source</td>
<td>dimensionless</td>
</tr>
<tr>
<td>(R_{C/N}')</td>
<td>(R_{C/N}') in mol C mol N(^{-1}): (R_{C/N}' = 12R_{C/N})</td>
<td>mol C mol N(^{-1})</td>
</tr>
<tr>
<td>(\varepsilon_m)</td>
<td>Diffusivity coefficient of the cell membrane layers</td>
<td>dimensionless</td>
</tr>
<tr>
<td>(\varepsilon)</td>
<td>Energy transfer efficiency</td>
<td>dimensionless</td>
</tr>
</tbody>
</table>

https://doi.org/10.1371/journal.pone.0208282.t001

---

**Fig 1.** Nitrogen fixation rate per unit protein for various C/N (sucrose/ammonium) supply ratios and two oxygen concentrations in a continuous culture of \textit{Azotobacter vinelandii}. Dilution rate is 0.15 (h\(^{-1}\)) with constant ammonium supply of 0.375 mol m\(^{-3}\) h\(^{-1}\). Here 100% O\(_2\) equals 225 μM thus approximately O\(_2\) saturation under normal air composition at 30˚C. Points are data redrawn from [34]; "+" based on acetylene reduction and "×" based on the rate of total nitrogen incorporation [34].

https://doi.org/10.1371/journal.pone.0208282.g001
Methods

Idealized metabolic model

The framework for this study is a previously published, simplified metabolic model of the heterotrophic soil bacterium *Azotobacter vinelandii* [40]. The model is depicted schematically in Fig 2 (and S1 Fig for detailed carbon fluxes), full details are provided in the Supplementary Material (S1 Text), and the code is available in Zenodo at https://zenodo.org/record/846300 (doi: 10.5281/zenodo.846300). Here we provide a brief description of the key elements of the model of [40] and the extensions developed for this study.

The framework consists of a simplified biochemical network to represent cellular metabolism constrained by mass, energy and electron balance at steady state [18]. The previously published model coupled diffusive uptake of organic carbon (sucrose), dinitrogen gas, and oxygen into a spherical cell. The uptake of the most limiting resource provided bounds on the growth rate. Growth efficiency was modeled using a balanced stoichiometric network which, following [18], combined half-reactions describing aerobic respiration, nitrogen fixation and synthesis of “biomass” with a prescribed elemental stoichiometry. Mass, electron and energy flow are conserved. The model is, in essence, a highly idealized “Flux Balance Analysis” rooted in the same principles as more detailed, genome-scale metabolic models. The published model [40] had two free parameters, energy conversion efficiency and the permeability of the cell wall, which were constrained by fitting the model to laboratory data when growing diazotrophically in the absence of fixed nitrogen [43]. Nitrogen fixation was assumed only possible when the intracellular oxygen was drawn down to very low concentrations, if necessary, by respiration beyond energetic demands (respiratory protection of nitrogenase, which is highly sensitive to...
The model successfully and quantitatively captured the relationship between nitrogen fixation, ambient oxygen concentration and the rate of supply of organic carbon substrate [40]. Here we extend the model to introduce ammonium assimilation as an additional nitrogen source. A complete description of the extended model is provided in the supplementary material. The previous model was simple enough to have a single metabolic configuration. The extended model has three possible metabolic configurations: exclusive diazotrophy, exclusive ammonium assimilation, or a combination of both. In the case of combined ammonium and dinitrogen use, the model organism can use the nitrogen sources in any combined ratio provided the uptake rates will allow it. We define the fraction of total nitrogen assimilation by nitrogen fixation as $f_{N2}$. The fraction of nitrogen assimilation from ammonium is thus $(1 - f_{N2})$. Thus an infinite variety of nitrogen sources is possible and we use an optimization approach to determine which combination will result. Since the laboratory data we use to constrain the model are continuous cultures, growth rate cannot be used to optimize “fitness” (as is often assumed). Instead we seek the combination of nitrogen sources which maximizes the standing stock of biomass (or protein) at the given growth (dilution) rate and environmental conditions. Since nitrogen fixation requires additional energy and electrons, relative to ammonium assimilation, it is intuitive that with low organic carbon and high ammonium supply rates, ammonium assimilation will be favored. Here, constrained also by the laboratory data from [34,41] we systematically examine and model the optimal nitrogen assimilation strategy under a variety of carbon and fixed nitrogen supply ratios, as well as a variety of ambient oxygen concentrations.

More specifically, we simulate *Azotobacter vinelandii* grown in a chemostat culture with a supply of ammonium based on laboratory experiments [34,41] for a range of C/N ratios (0–15 mol sucrose mol ammonium$^{-1}$, with constant ammonium at 2.5 mM) and oxygen concentrations (5%, 15%, 30% and 60%), at a dilution rate of 0.15 h$^{-1}$.

Model parameters are adapted from [40], which were either estimated from the literature, or fitted to the data by tuning (two parameters: diffusivity coefficient of the cell membrane layers, $\varepsilon_m$, and energy transfer efficiency, $\varepsilon$). We employ a higher transfer efficiency, $\varepsilon$, for ammonium-uptake based biomass production ($\varepsilon = 0.54$) by calibrating the model using laboratory data on microbes grown with glucose and ammonium compiled in [5].

Laboratory data of nitrogen fixation rates, respiration rates and protein concentrations were extracted from [34] (doi: https://doi.org/10.1007/BF00414840) and [41] (doi: https://doi.org/10.1007/BF00414819). The model developed in this paper has been written in Python 3 and uploaded in Zenodo at https://zenodo.org/record/846300 (doi: 10.5281/zenodo.846300).

**Results and discussion**

First we simulate the conditions of the laboratory studies and obtain steady-state solutions at a fixed dilution rate of 0.15 h$^{-1}$ across a range of input C/N and ambient oxygen concentrations. Specifically, we impose an ammonium concentration of 2.5 mM in the incoming medium and varied sucrose concentration from 0 to 37.5 mM, giving a C/N supply range of 0–15 (equivalent to an elemental C:N supply ratio of 0–180). Predicted nitrogen fixation and respiration rates are in good agreement with the laboratory data across a range of ambient oxygen concentrations, as shown in Fig 3A and 3B. The model also predicts observed protein concentration (S2 Fig) supporting the model assumption of maximizing culture biomass.

The model captures salient properties of the observed responses of nitrogen fixation to C/N ratio and oxygen concentration (Fig 3A). Nitrogen fixation occurs only above a threshold C/N
ratio ($R_{C/N}$) and the protein-specific nitrogen fixation rate increases in concert with C/N because carbohydrate in excess of that needed to assimilate the ammonium fuels nitrogen fixation and oxygen scavenging. At a given C/N, the nitrogen fixation rate is negatively impacted by increasing oxygen concentration (Fig 3A) as the cell must devote more carbohydrate towards protective respiration (Fig 3B, Fig 4). At higher C/N ratios, the proportion of the cellular nitrogen requirement accounted for by nitrogen fixation, here defined as $f_{N2}$, approaches 1 (= 100% in Fig 3A), and the nitrogen fixation rate at different oxygen levels converges to the same maximum rate. Below an oxygen-dependent threshold C/N, nitrogen fixation does not
Nitrogen fixation rates based on three metabolic phases

At a given oxygen concentration, based on the maximization of the protein-specific biomass production rate, the model identifies three distinct metabolic regimes corresponding to carbohydrate limited growth (low C/N), ammonium limited growth (moderate C/N), and combined ammonium and dinitrogen-based growth (high C/N), respectively, illustrated schematically in Fig 5.

Below the oxygen-dependent threshold C/N (Fig 5A), carbohydrate is the growth limiting factor and the ammonium in the culture is not fully consumed (Fig 5A and 5D). Since ammonium assimilation requires far less energy and fewer electrons than nitrogen fixation, optimal biomass production occurs in the absence of nitrogen fixation. Hence at low C/N delivery, growth efficiency and specific biomass production rate favor ammonium-based growth. With increased carbohydrate input, the cell density increases leading to a higher protein and biomass concentration (Fig 3C and S2 Fig, lowest C/N).

At moderate C/N ratios (Fig 5B) the cells are ammonium limited. Though all the excess carbohydrate is expended on respiration in excess of energetic demands, it is still insufficient to scavenge oxygen sufficiently to keep the cell cytosol nearly anoxic and protect nitrogenase. Hence nitrogen fixation is still inhibited (see section "High respiration during non-nitrogen fixing phase" for further discussion). Since the incoming ammonium concentration is fixed occur either in the laboratory data or model. The model provides an interpretation of these results in terms of three metabolic regimes.
and ammonium is consumed to an undetectable level \([34,41]\) (and the incoming sucrose concentration varied), the protein and biomass concentration stays constant in both laboratory experiments and simulations (Fig 3C and S2 Fig, constant lines).

At high C/N ratio (Fig 5C), due to excess carbohydrate relative to ammonium, ammonium in the medium is drawn down to its subsistence concentration (Fig 5D) and there is sufficient

---

**Fig 5. Nitrogen metabolism at different C/N ratios.** As C/N ratio increases the cell metabolism shifts from carbohydrate limitation at low C/N supply (A), to ammonium limitation at moderate C/N (B) then to combined ammonium assimilation and nitrogen fixation at high C/N supply which is also carbon limited provided the N\(_2\) concentration is sufficiently high (C). Black arrows, nutrient uptake; red arrows, respiration; blue arrows, biomass production; green arrow, nitrogen fixation. CH and BIO indicate the intermediate intracellular store of carbohydrate and cellular biomass, respectively. Respiration per unit protein, the ammonium concentration in the medium and nitrogen fixation rate per protein transitions in three phases are shown in (D).

https://doi.org/10.1371/journal.pone.0208282.g005
additional carbon with which to completely scavenge oxygen (Flat part in Figs 3B and 4) and fuel nitrogen fixation (Figs 3A, 4 and 5D). In this regime, growth is based on both ammonium assimilation and nitrogen fixation and the modeled nitrogen fixation rate increases non-linearly with the C/N ratio (Fig 3A), consistent with the laboratory data [34]. The experimental data are consistent with a model in which the cells allocate carbohydrate to ammonium assimilation and nitrogen fixation (with its attendant respiratory protection) in a manner that maximizes the standing stock of biomass and biomass production. Increasing carbohydrate resource leads to higher nitrogen fixation increasing protein and biomass concentration in the culture (higher end of C/N in Fig 3C and S2 Fig).

The diffusion of oxygen into the cells negatively influences the nitrogen fixation rate. As the ambient oxygen concentration increases, so does respiration rate (Fig 3B) as carbon is channeled into respiratory protection (Fig 4), at the expense of nitrogen fixation (Fig 3A). At an oxygen concentration at 60% of saturation (here 100% saturation equals 225 μM; about 21% atmospheric composition at a temperature of 30˚C), more than half of nitrogen for growth is supplied by ammonium if the C/N (sucrose/ammonium) supply ratio is lower than 14 (Fig 3A). On the other hand, at a low oxygen saturation of 5%, nitrogen fixation accounts for about 80% of nitrogen in biomass at C/N ratios higher than 8 (Fig 3A). The threshold C/N above which nitrogen fixation occurs is oxygen dependent (Fig 3A, Fig 4). The higher the ambient oxygen concentration the greater the required expenditure on respiratory protection, and the greater the C/N threshold for nitrogen fixation. This is qualitatively and quantitatively captured by the idealized metabolic model.

Nitrogen fixation as a function of carbohydrate, oxygen, and ammonium

The laboratory data and simulations discussed above considered a constant ammonium input (ammonium concentration of 2.5 mM and dilution rate of 0.15 hr^-1). Having demonstrated the consistency of the model and its parameters with these data, we now extrapolate beyond the range of laboratory studies to consider the variation of nitrogen fixation in *Azotobacter vinelandii* over a range of both ammonium and sucrose input concentrations, and at two levels of oxygen saturation: 5% and 30% (Fig 6).

As above, the nitrogen fixation rate per unit protein is controlled by the C/N ratio and by oxygen concentration (Fig 6i and 6ii). Note that contours of constant C/N have constant per protein nitrogen fixation rates. The critical C/N ratio for the onset of nitrogen fixation (white line, transition from regime B to regime C) increases with the oxygen concentration but even in a high oxygen environment (Fig 6ii), as long as the C/N ratio is sufficiently high, the model predicts high nitrogen fixation rate per protein.

Nitrogen fixation rate per unit volume of culture, however, varies along lines of constant C/N (compare regime C in Fig 6i and 6iii) since higher carbohydrate supply supports higher densities of cells at the same dilution rate (S3 Fig). Since ammonium does not suppress cell density but rather increases it in some regimes (S3 Fig), the negative influence of ammonium on nitrogen fixation per volume is relatively limited. The nitrogen fixation rate per volume is more severely suppressed by oxygen (Fig 6iii vs. 6iv); at 30% oxygen, the nitrogen fixation rate per volume is significantly decreased due to expenditure on respiratory protection which lowers the cell concentration (S3 Fig).

High respiration during non-nitrogen fixing phase

Nitrogen fixation requires low intra-cellular oxygen concentrations since nitrogenase is sensitive to oxygen. *Azotobacter* manages intracellular oxygen using respiratory protection [7,11] which has also been indicated in other nitrogen fixers including *Crocosphaera, Trichodesmium*...
and *Anabaena* [13,15,45,46]. Interestingly, comparison of models and data (Fig 7) suggests *Azotobacter* constitutively respires any carbohydrate in excess of the energetic demand for synthesizing biomass, even if it is insufficient to fully deplete intracellular oxygen.

To examine the level of respiration during ammonium limited growth, we compare two simulations. One simulation (Sim1) assumes that all 'excess' carbohydrate will be respired under all growth conditions: that is, the metabolic configuration is identified that maximizes biomass concentration. In the other simulation (Sim2), 'excess' respiration only occurs when the cells are fixing nitrogen, reflecting the possibility that respiratory protection is regulated and repressed when the cell is not actively fixing nitrogen. We find that model Sim1 predicts
an increase in respiration with increasing C/N, consistent with observations (Compare Sim1 and data in Fig 7i and 7ii), whereas Sim2 substantially underestimates rates of respiration during growth on ammonium (Compare Sim2 and data in Fig 7i and 7ii). The concordance of Sim1 with laboratory measurements suggests that oxygen-scavenging respiration is performed even when the cells are not able to fix nitrogen.

There are several possible explanations for these observations. One is that nitrogen fixation is so central to the biology and ecology of these organisms, that they perform excess respiration (respiratory protection) constitutively, at least to the point where it does not have a growth cost due to limitation by carbon. This might reflect adaptation of other enzymes or intracellular processes to low oxygen conditions, or might reduce the need for regulation of respiratory protection pathways. The latter might be advantageous particularly in dynamic or patchy environments, and might allow cells to preserve nitrogenase when they are not fixing nitrogen. In this case, cells might be able to commence nitrogen fixation rapidly when conditions allow [12].

**Wider implications**

**Simultaneous N$_2$ and fixed-N assimilation and relative fitness.** The experimental results of [34,41] demonstrate that nitrogen fixing organisms can, and will, simultaneously use both fixed and molecular nitrogen as nitrogen source. In their experiments, *Azotobacter vinelandii* drew down ammonium concentrations to undetectable levels even when fixing nitrogen, provided the supply of organic carbon was sufficient relative to the supply of fixed nitrogen. Our model captures this threshold and suggests that this is consistent with a metabolic organization which maximizes the per protein biomass production rate [47]. This is also consistent with numerous lines of evidence showing that, while the availability of fixed nitrogen can suppress nitrogen fixation in diazotrophs, it does not always do so [21]. Many of these examples are phototrophic nitrogen fixers in marine environments [21]. In the oligotrophic surface ocean,
light and energy are in excess supply relative to nitrogen (analogous to the excess organic car-
bon supply to Azotobacter). Under these circumstances, where the direct and indirect (oxygen
management) costs of nitrogen fixation by phototrophs can be easily met, it is still a fitness
advantage to utilize both N₂ and fixed nitrogen: by doing so, the diazotrophs increase its over-
all growth efficiency as assimilating fixed nitrogen is energetically cheaper. It also denies some
of the limiting nitrogen resource for the non-diazotrophs, effectively increasing the size of
the niche, or potential range, for the nitrogen fixer at the expense of the non-diazotrophs. A
simple theoretical perspective to demonstrate this is outlined in the Supplementary Material
(S2 Text).

Inferences on the viability of marine heterotrophic nitrogen fixation. We chose to
model a heterotrophic, diazotrophic soil bacterium, Azotobacter vinelandii, because of the
comprehensive laboratory data sets available. Analogous organisms are present in the oceans.
Nitrogenase genes associated with heterotrophic bacteria are almost ubiquitous in marine
waters and commonly dominate the nifH gene pool throughout the water column when eval-
uated from PCR amplicon libraries [48,49]. While the rates of nitrogen fixation associated with
these organisms is generally considered to be lower than that of photoautotrophic nitrogen fix-
ers, there are observations of heterotrophs actively fixing nitrogen both in surface and deeper
waters [50–54]. Whether or not their contribution to global nitrogen fixation is large, the exis-
tence of these organisms suggests an ecological niche and an evolutionary advantage. How-
ever, it is still not clear how exactly these organisms carry out nitrogen fixation in oxygenated
marine environments. Recent studies show that a model organism BAL361 responds to oxygen
and ammonium similarly to Azotobacter vinelandii [35,55] and here, we apply our model to
consider the potential limitations on their niche.

Consider the critical atomic ratio of carbon to nitrogen in the substrate: \( R_{C:N}^f = 12 \times R_{C:N}^f \),
(mol C mol N⁻¹) which for sucrose-fueled Azotobacter lies between ~15 at an oxygen concen-
tration of 5% of saturation, and ~100 at 60% of saturation. This C:N is higher than that of most
observed particulate organic matter in marine systems [56,57], suggesting that heterotrophic
nitrogen fixation will be unlikely to occur.

However, using the model we predict \( R_{C:N}^f \) to be as low as 6.5 at 1% oxygen saturation;
lower than many observed samples of particulate organic matter [56,57] as well as the observed
elemental ratio of dissolved organic matter, DOM [56,58,59]. The model also suggests that at
this oxygen concentration respiratory protection is not necessary since energetically balanced
respiration for nitrogen acquisition and biosynthesis fully deplete the intracellular oxygen.
This implies that heterotrophic nitrogen fixation may occur even in the presence of ammo-
nium at the low oxygen concentrations which can exist within particles due to active heterotro-
phic respiration [60]. The composition and nutritional value of natural DOM and POM is
poorly characterized and variable, and the analogy between Azotobacter and marine heterotro-
phic diazotrophs like BAL361 is not proven. However, the relationships described by the
model above provide clear hypotheses which could be tested with laboratory cultures of
marine organisms, employing organic matter of known stoichiometry and controlled oxygen
conditions.

Summary
Using a simplified metabolic model, we simulated continuous cultures of the heterotrophic
nitrogen fixer Azotobacter vinelandii, identifying and quantifying relationships between the
availability of fixed nitrogen and nitrogen-fixation. We find that using both nitrogen sources is
optimal for the diazotroph when energy and other resources allow and identify the transitions
between nitrogen and carbon limited regimes. We show that Azotobacter uses organic carbon
sources in excess of the requirements for respiratory protection even when there is insufficient excess carbon to deplete intra-cellular oxygen and enable nitrogen fixation. The model suggests that using both fixed nitrogen and dinitrogen gas, marine diazotrophs increases their fitness and the size of their fundamental niche. Finally, the model provides a testable hypothesis that heterotrophic nitrogen fixation may occur in marine particles despite the presence of ammonium.

Supporting information

S1 Text. Model details.  
(PDF)

S2 Text. Fitness advantage of using fixed nitrogen and dinitrogen.  
(PDF)

S1 Fig. Schematic of detailed carbon fluxes in a modeled cell.  
(PDF)

S2 Fig. Simulated concentrations of biomass in continuous cultures of *Azotobacter vinelandii*.  
(PDF)

S3 Fig. Cell density proxied by protein concentration for various sucrose and ammonium resources with different oxygen saturations.  
(PDF)

Acknowledgments

We thank Eric Stansifer for his help on Python programming, Lauren Woodruff for useful discussions on nitrogen fixers, and members of Writing and Communication Center at MIT for their advice on writing.

Author Contributions

**Conceptualization:** Keisuke Inomura, Jason Bragg, Lasse Riemann.

**Data curation:** Keisuke Inomura.

**Formal analysis:** Keisuke Inomura.

**Funding acquisition:** Keisuke Inomura, Lasse Riemann, Michael J. Follows.

**Investigation:** Keisuke Inomura.

**Methodology:** Keisuke Inomura, Michael J. Follows.

**Project administration:** Keisuke Inomura, Michael J. Follows.

**Resources:** Keisuke Inomura, Michael J. Follows.

**Software:** Keisuke Inomura.

**Supervision:** Michael J. Follows.

**Validation:** Keisuke Inomura.

**Visualization:** Keisuke Inomura.

**Writing – original draft:** Keisuke Inomura.
Writing – review & editing: Keisuke Inomura, Jason Bragg, Lasse Riemann, Michael J. Follows.

References


42. Noar JD, Bruno-Bárcena JM. *Azotobacter vinelandii*: the source of 100 years of discoveries and many more to come. Microbiology. 2018; 164: 421–436. https://doi.org/10.1099/mic.0.000643 PMID: 29533747


46. Murry MA, Wolk CP. Evidence that the barrier to the penetration of oxygen into heterocysts depends upon two layers of the cell envelope. Arch Microbiol. 1989; 151: 469–474.


