Constraining oxygen consumption by nitrite oxidation in oceanic oxygen minimum zones

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Abstract

Oceanic oxygen minimum zones (OMZs) occur where microorganisms deplete dissolved oxygen (DO) to exceptionally low levels, and are globally significant sites of biogeochemical cycling. Amid the intense competition for DO and other substrates occurring in these metabolically challenging environments, aerobic nitrite oxidation may consume significant amounts of DO, but this has not been examined comprehensively. Using parallel measurements of oxygen consumption rates and ^{15}N-nitrite oxidation rates applied to water column profiles and to oxygen manipulation experiments, we show that nitrite oxidation is a substantial sink for DO in the ocean's largest OMZ. The contribution of nitrite oxidation to overall DO consumption increased at low DO concentrations, tracking gradients and variations within and across multiple stations in the eastern tropical North Pacific Ocean. Oxygen manipulation experiments produced highly consistent effects, with nitrite oxidation responsible for progressively more DO consumption (up to 97%) as DO was experimentally decreased. Natural abundance stable isotope data indicated coupling of nitrite oxidation and nitrate reduction, while 16S rRNA and metagenome sequencing revealed that *Nitrospina* ecotypes possessing high-affinity cytochrome oxidase genes were prevalent and active within the OMZ. Collectively, our results demonstrate that nitrite oxidation consumes significant amounts of DO, and that this proportion increases as DO declines— indicating that nitrite oxidation is critically important to the formation and maintenance of OMZs.

Significance

Oceanic oxygen minimum zones (OMZs) are naturally-occuring regions of low oxygen found in select areas of the ocean. Lack of dissolved oxygen has important implications for both the distribution of marine organisms and global biogeochemical cycles, yet we have a limited understanding of how oxygen is depleted to such low levels. Here we comprehensively quantify the contribution of nitrite oxidation to oxygen depletion in the ocean's largest OMZ. We observed highly consistent patterns across depth profiles, and in multiple experiments where we manipulated oxygen concentrations, finding that nitrite oxidation consumes progressively more oxygen at lower oxygen concentrations. Our findings demonstrate that nitrite oxidation plays a pivotal role in exhausting oxygen to the low levels found in OMZs.

Introduction

Aerobic nitrite oxidation is pervasive throughout much of the oceanic water column, playing a central role in deep ocean chemoautotrophy and carbon cycling (1, 2), as well as in the oceanic nitrogen (N) cycle (3, 4). Nitrite oxidation rates are typically undetectable using $^{15}NO_2^-$ isotopic tracer in the sunlit euphotic zone, peak at the base of the euphotic zone, and subsequently decline with depth (5). However, nitrite oxidation rate profiles deviate from this pattern in oceanic oxygen minimum zones (OMZs) that are depleted in dissolved oxygen (DO): rapid rates have been reported despite low DO concentrations (6–9), and nitrite oxidation itself may consume DO to low levels. OMZs are ultimately generated by high sinking fluxes of organic matter combined with reduced ventilation at depth, and are typically defined by DO concentrations <20 µM (10). Regions of the tropical oceans containing no measurable DO—termed anoxic marine zones (AMZs; ref. 11)—are further distinguished by the accumulation of nitrite to comparatively high (>1 µM) concentrations (12). These secondary nitrite maxima (SNM) result from anaerobic nitrate reduction to nitrite under low DO (11–13). Accumulated nitrite may be subsequently reduced via denitrification and anaerobic ammonium oxidation (14)—or, alternatively, oxidized by NOB. Nitrite is therefore rapidly produced and consumed via multiple processes in OMZs, linking the oceans' N, carbon, and oxygen cycles.

Despite its centrality in OMZ biogeochemical cycles, nitrite oxidation is still poorly understood in these regions of the ocean. In particular, high rates of nitrite oxidation where DO levels are intrinsically low indicates that marine nitrite-oxidizing bacteria (NOB) may contribute substantially to DO depletion and the ultimate generation and maintenance of OMZs. However, this contribution to O_2 drawdown has not been directly quantified. While nitrite oxidation has been measured in several OMZs (6–9), oxygen consumption rate (OCR) measurements in OMZs are rare (15). Yet simultaneous measurements are necessary to directly quantify the contribution of nitrite oxidation to oxygen consumption.

An additional distinguishing feature of AMZs is the presence of a deep chlorophyll maximum (DCM) where distinct ecotypes of the globally important cyanobacterium, *Prochlorococcus*, produce DO within low DO waters (16–18). DCM-based DO production can overlap with nitrite supply via nitrate reduction, but the degree to which nitrite oxidation (versus other processes) consumes DO in the DCM remains poorly constrained (18).

Our understanding of how NOB oxidize nitrite under low O_2 is also evolving. NOB gain

energy by oxidizing nitrite to nitrate while using dissolved oxygen (DO) as a terminal electron

acceptor. Adaptations to low DO are evident in different NOB, and both *Nitrospina* and

Nitrococcus may contribute to varying degrees (19, 20). But although *in situ* nitrite oxidation

rates are typically highest under low DO, experimental oxygen manipulations show the opposite

pattern: declining rates with declining DO (21). This may be explained by variations in genomic

content among different NOB that allow them to occupy distinct, highly-resolved DO niches

throughout the water column (4, 7, 21)—including those that occur from the edge (20 µM) to the

core (<5 nM) of OMZs. These details are essential to understand in the context of ocean

deoxygenation, as seemingly small variations in DO may have significant biogeochemical

implications if different ecotypes within different functional groups—such as nitrite oxidizers,

nitrate reducers, and organisms respiring organic matter aerobically—have varying sensitivities

to DO.

Here we address these open questions through parallel measurements of nitrite oxidation

and overall OCR in the oceans' largest OMZ, the eastern tropical North Pacific Ocean (ETNP).

Nitrite oxidation rate measurements (using $^{15}NO_2^-$), OCR measurements (using optical sensor

spots), and ammonia oxidation rate measurements (using $^{15}NH_4^+$) were made along depth

profiles at six stations in the ETNP, including three OMZ stations and three AMZ stations

(Figure 1). At each station, we sampled in the upper 100 m to capture the primary nitrite

maximum and an expected peak in nitrite oxidation rates at the base of the euphotic zone (EZ).

We then sampled a range of DO values (from 200 µM to the anoxic SNM at AMZ stations) to

quantify rate variations in response to vertical gradients in DO. How nitrite oxidation varies

from the OMZ edge to its anoxic core is critical to our understanding of ocean biogeochemical

cycles; we therefore conducted oxygen manipulation experiments and examined the response of

nitrite oxidation and OCR. Combined with natural abundance stable isotope data, 16S rRNA

sequences, and metagenomes, we show that nitrite oxidation is a substantial, and occasionally

even dominant, sink for oxygen in the ETNP.

Results and Discussion

Nitrite oxidation rates in the ETNP

We sampled six stations in the ETNP OMZ with low (<20 µM) DO concentrations at depth (Figs

1 and 2). Three stations extending out from the coast of Mexico (Stations 1-3) were AMZ

stations with distinct DCMs and accumulations of nitrite in SNMs indicative of anaerobic N

cycling (Figure 2A and B). We expected that these would be hotspots of nitrite oxidation given

oxygen supply via photosynthesis in the DCM overlapping with nitrite supply via nitrate

reduction. Across all stations and depths, we observed the highest nitrite oxidation rates at the

AMZ stations (Fig. 2C). Station 1 is located nearest the coast and had a shallow OMZ at the

time of our sampling, with 20 µM DO at 28 m depth, and nitrite > 1 µM at 100 m. Chlorophyll

concentrations were also high in the upper water column (up to 5 mg m^{-3} at 20 m), with a DCM

spanning 75-125 m. Nitrite oxidation displayed a local maximum at the base of the EZ at Station

1 (20-30 m), and then increased to higher levels (>100 nmol L^{-1} d^{-1}). This increase at 100 m

corresponded with the overlap between the bottom of the DCM and the top of the SNM. Nitrite

oxidation rates then reached higher values at 125 and 150 m within the SNM at Station 1.

Stations 2 and 3 displayed similar nitrite oxidation rate profiles to each other, including elevated

rates in the DCM and SNM. Nitrite oxidation rates were similar in magnitude, and peak values

at the base of the EZ and in the OMZ were also similar (69-96 nmol L^{-1} d^{-1}). Depth patterns

tracked oceanographic differences across the three AMZ stations, as the depth of all features

increased moving offshore from Station 1 to 2 to 3 (Fig. 2). For example, the DCM extended

from 105-155 m at Station 2, while nitrite concentrations began to increase below 100 m; nitrite

oxidation rates were elevated at 140 m and declined slightly with increasing depth. At Station 3,

the DCM (120-180 m) and SNM (>140 m) depths were deeper, and nitrite oxidation rates

increased from 100 to 200 m.

In contrast to these three AMZ stations (Stations 1-3), rate profiles at Stations 4-6 showed

peaks at the base of the EZ and decreases with depth, with no subsurface increase within the

OMZ (*SI Appendix,* Figure S1). Parallel measurements of ammonia oxidation rates also showed

this type of pattern at all stations (*SI Appendix*, Fig. S1). Subsurface maxima in ammonia

oxidation tracked variations in the EZ across all six stations, but rates were not elevated in

OMZ/AMZ waters—again contrasting with nitrite oxidation rate profiles at the AMZ stations.

These data accord with earlier work in OMZs showing contrasting ammonia and nitrite oxidation

rate profiles, and particularly high rates of nitrite oxidation in OMZ waters (6–8, 22-24).

Elevated nitrite oxidation below the DCM (> 120 m at Station 1, >155 m at Station 2, and >180

m at Station 3), where little to no DO is available, could have a number of possible explanations

discussed below. Within the DCM, our data support the idea that nitrite oxidation contributes to

'cryptic' oxygen cycling (18)—i.e., that DO produced via oxygenic photosynthesis is rapidly

consumed.

Oxygen consumption via nitrite oxidation

We determined the contribution of nitrite oxidation to overall oxygen consumption via parallel

measurements of oxygen consumption rates (OCR) using *in situ* optical sensor spots—which are

non-invasive, compare favorably with other low-level measurement approaches, are the only

cost-effective means of achieving of substantial replication, and for which sensitivity increases

under low DO (25, 26). Decreases in DO were measured in both nitrite and ammonia oxidation

rate sample bottles, as well as in three additional replicates, to leverage statistical power for

increased sensitivity to low-level DO consumption (see Materials and Methods). Water column

OCR profiles at all stations showed exponential declines with depth and decreasing DO

concentrations (Fig. 2D and S1). OCR magnitudes were similar to the limited previous

measurements that have been conducted in OMZs (e.g., 161-2600 nmol L^{-1} d^{-1} in ref 15). OCR

also tracked variations in DO across stations, with progressively steeper declines in OCR with

depth from Station 6 through Station 1. This pattern of declining OCR with increasing depth and

decreasing DO contrasted with that of nitrite oxidation rates, which were notably elevated under

low DO (Fig. 2).

OCR was not significantly different from zero at >125 m at Station 1 and >140 m at

Stations 2 and 3, obviously owing to the initial lack of DO at these depths at the AMZ stations.

However, typically 1-2 out of 5 replicates did contain DO introduced during sampling and

incubation. Short of conducting *in situ* incubations, such DO contamination is essentially

unavoidable. However, all microorganisms and metabolic processes exposed to DO may

capitalize on its introduction and directly compete for DO. We directly compared nitrite

oxidation rates with OCR, assuming that each mole of nitrite is oxidized using ½ mole of O_2 (5).

(As the O added to nitrite to form nitrate comes from water, and ^{15}N is used to measure the rate,

it is possible that oxidants other than DO are used; see below.) We found that nitrite oxidation

increased as a proportion of overall OCR at lower DO levels (Figure 3A and B). Nitrite

oxidation was responsible for up to 69% of OCR at Station 1, although most values were closer

to 10-40% at Stations 2 and 3 (Fig. 3). In contrast, ammonia oxidation contributed no more than

5% in the OMZ (*SI Appendix*, Fig. S1).

Effects of oxygen manipulation experiments on nitrite oxidation and OCR

These data indicate that NOB are poised to rapidly consume any DO introduced to sample

bottles, although they may be partly affected by sample collection and manipulation. We

therefore verified these observations by experimentally manipulating DO concentrations to

measure the response of OCR, as well as $^{15}NO_2^-$ nitrite oxidation rates, to changing DO (Figure

4). We conducted experiments on the edge of the OMZ (depth of 20 µM DO) at Stations 1-5, in

the DCM at Stations 2 and 3, and in the SNM at Station 3 and an additional sampling Station 3.5

(Table 1). (We could not conduct additional experiments at Stations 1 and 2 owing to a

hurricane in the region at the time of sampling in 2018, and added Station 3.5 as a result.)

Oxygen manipulations were designed to quantify nitrite oxidation and OCR across the spectrum

of DO concentrations that occur from the OMZ edge to its core—rather than solely probing their

lower limits—in order to constrain rate responses to changing DO within OMZs.

Oxygen manipulation experiments were consistent with water column profiles and

provide additional evidence that nitrite oxidation can be a substantial oxygen sink. In all

experiments, we found that OCR declined with experimentally decreased DO (Fig. 4). All

curves displayed Michaelis-Menten type forms; however, we note that mixed assemblages of

microorganisms drive this overall pattern by using DO to oxidize a variety of substrates. These

substrates can include nitrite, as well as different forms of organic matter, reduced sulfur

compounds, and possibly methane (27). Consistent with earlier work (21), nitrite oxidation rates

also displayed Michaelis-Menten-type curves in most of our experiments (Fig. 4). However,

nitrite oxidation rates were less sensitive to declining DO than OCR. Nitrite oxidation was

therefore responsible for progressively higher proportion of overall OCR as DO was

experimentally decreased (Fig. 3C). In fact, we found that 97% of OCR in the DCM at Station 2

could be explained by measured nitrite oxidation rates, while most values were closer to 10-20%.

This pattern in experiments was notably similar to results from rate profiles (Fig. 3B and C).

Based on these data, we calculated the DO level at which nitrite oxidation may be

expected to consume all DO. As the data above indicate, this is obviously variable across

experiments: our experiment in the DCM at Station 2 showed that nitrite oxidation can consume

essentially all available DO (Fig. 4 C), while the experiment with 20 μM [DO] water at Station 2

suggests that, even at low DO levels, nitrite oxidation would not consume all DO (Fig. 4B). We

fit Michaelis-Menten-type curves to nitrite oxidation rates and OCR individually (Table 1;

although these are, again, mixed microbial assemblages). Calculated DO affinities (K_s values)

were substantially lower for nitrite oxidation compared with OCR, particularly in the DCM

(Stations 2 and 3) and SNM (Stations 3 and 3.5). Based on these relationships, OCR always

exceeds nitrite oxidation on the OMZ edge, but nitrite oxidation is estimated to consume all DO

below concentrations of 35-96 nM in the SNM and DCM.

These data tie together multiple aspects of OMZ biogeochemistry into a coherent picture.

Although nitrite oxidation rates decline as DO is experimentally decreased at individual depths—

and so with particular assemblages of NOB—different depths/assemblages display different

properties (Fig. 4). More importantly, this decline is always less severe for nitrite oxidation than

for overall OCR (Figs. 2-4). NOB are therefore highly effective at scavenging low levels of DO

(Figs. 3 and 4; (18), which is consistent with the idea of a nitrogen-oxygen feedback loop in

OMZs driven partly by oxygen depletion via nitrite oxidation (28).

Isotopic, 16S rRNA, and metagenomic constraints on nitrite oxidation

To further verify our findings, we applied natural abundance stable isotope measurements, 16S

rRNA sequencing, and metagenome sequencing to water samples and nucleic acid samples

collected in parallel with nitrite oxidation rate measurements (Figure 5). The dual N and O

isotopic composition of dissolved nitrate ($\delta^{15}N$ and $\delta^{18}O$, respectively; *SI Appendix* Fig. S2) is

thought to provide an effective constraint on nitrite oxidation under low DO conditions due to

isotopic 'overprinting' by nitrite oxidation (29, 30). While the respiratory reduction of nitrate

(denitrification) leads to equal 1:1 increases in $\delta^{15}N$ and $\delta^{18}O$ (31, 32), deviations in 1:1 N:O

isotope behavior arise from a decoupling of the N and O systems. These deviations are widely

interpreted as reflecting cryptic re-oxidation of nitrite under low DO, where the reduction of

nitrate to nitrite removes an O atom, and the subsequent re-oxidation of nitrite to nitrate appends

a new O atom derived from ambient water (33, 34). Deviations from the 1:1 relationship are

represented via $\Delta(15,18)$ values (35), where more negative values represent larger departures

from the 1:1 relationship.

At Stations 1-3, dual nitrate isotope values exhibited non-linear trends (Figure 5B and *SI*

Appendix Fig. S3) that are consistent with isotopic overprinting by nitrite oxidation (29, 30).

$\Delta(15,18)$ values were lowest (i.e., deviations were largest) at 50-150 m at Station 1, 100-140 m at

Station 2, and 110-160 m at Station 3. The lower portions of these depth ranges overlapped with

the upper portion of the DCM (75-125 m, 105-155 m, and 120-180 m, respectively) and tracked

depth variations between stations. Peak deviations at Stations 1 and 2 were just above the DCM

(75 and 100 m), while the peak at Station 3 corresponded closely with the DCM (140 m).

Throughout these ETNP OMZ sites, our nitrate isotope data are consistent with previous

observations and interpretations of rapid recycling between nitrate and nitrite (29, 35, 36),

ultimately evidenced by isotopic overprinting of the nitrate reduction signal by nitrite oxidation.

16S rRNA gene and transcript sequencing revealed a similar pattern to rate profiles,

experiments, and isotopic data, while also enabling identification of NOB that are abundant and

active in the ETNP OMZ. Based on DNA, *Nitrospina* 16S rRNA amplicon sequence variants

(ASVs; ref. 37) comprised up to 3.4-5.4% of all ASVs at Stations 1-3 (Fig. 5C), and relative

abundances were strikingly consistent with the isotopic anomalies—abundances and anomalies

were well-correlated (r^2=0.70-0.84, P<0.05) at Stations 1 and 3, for instance. These patterns

were accentuated for RNA samples, but with discrete peaks in 16S rRNA transcripts generally

occurring at deeper depths (Fig. 5D). *Nitrospina* 16S rRNA peaked at 125 m at Sta 1 (1.8% of

all 16S rRNA transcripts; with an additional upper water column peak at 50 m), 140 m at Station

2 (3.2%), and particularly 140-180 m at Station 3 (1.6-7.1%). All of these depths lie within the

DCM at each station. Although 20 *Nitrospina* ASVs were identified, three were dominant (each

>1% of all ASVs, together constituting 72-100% of all *Nitrospina* 16S rRNA gene sequences

and transcripts). One of these ASVs was found only in OMZ samples at Stations 1-3, and was

not detected above the OMZ or at Stations 4 and 5. This was also the lone ASV observed in the

OMZ at Station 3, where *Nitrospina* were most active based on their comparatively high

percentages of all 16S rRNA transcripts. In contrast, no *Nitrococcus* ASVs and only one low-

abundance *Nitrospira* ASV were identified out of >11,000 ASVs and >3.5 million 16S

sequences from 73 DNA and 73 RNA samples. In line with earlier work in the ETNP (7, 19),

other OMZs (6, 38), and the varaitions in DO affinity observed across oxygen manipualtion

experiments (Fig. 4), these results indicate that particular *Nitrospina* ecotypes—and one ASV in

particular—are significant for low-oxygen nitrite oxidation, while other ASVs may be more

important at different depths and DO concentrations.

To validate the genetic potential for nitrite oxidation by *Nitrospina* under low DO

concentrations, we sequenced metagenomes collected on the OMZ edge, at the DCM, and within

the OMZ core at Stations 1, 2, and 3. Nitrite oxidoreductase genes from *Nitrospina* were

prevalent at all stations and depths, establishing the genomic potential for nitrite oxidation

throughout the ETNP OMZ (Table 2). In addition, multiple high-affinity cytochrome C oxidase

genes from *Nitrospina* were present in all samples—indicating that *Nitrospina* are capable of

consuming DO at the low concentrations found within the OMZ. These genes were more

common in DCM and SNM metagenomes than at the OMZ edge, consistent with *Nitrospina*

ecotype distributions, and with the lower DO concentrations found at the DCM and SNM.

Cytochrome *bd*-type oxidase genes from *Nitrospina* were also detected in all metagenomes

except the OMZ edge sample at Station 1, although these lack quinol binding sites and may not

functional as canonical oxidases (19, 39). Sun et al. (19) also suggested that chlorite dismutase

genes may be relevant for anaerobic metabolism in *Nitrospina*, and these were present in all

metagenomes. Finally, *Prochlorococcus* genes were prevalent from the OMZ edge to core, and

especially within the DCM, supporting the idea that oxygenic photosynthesis and cryptic oxygen

cycling occur throughout the ETNP OMZ. Overall, metagenomic data are consistent with

experimental data and 16S data, and indicate *Nitrospina* are tightly tuned to DO concetrations in

the ETNP.

Oxygen consumption by nitrite oxidation in OMZs

Assembled together, our results provide a comprehensive view of nitrite oxidation and its

contribution to oxygen consumption in the ocean's largest OMZ. Rate measurements, oxygen

manipulation experiments, stable isotopic data, and molecular data all converge on the fact that

nitrite oxidation is active from the base of the euphotic zone into the OMZ. Nitrite oxidation was

particularly significant within the DCM at Stations 1-3: nitrite oxidation rates were elevated,

isotopic overprinting of nitrate by nitrite oxidation was evident, and *Nitrospina* were abundant

and active (Fig. 2C and 5). All of these data tracked consistent progression in the depth of the

DCM across stations, and indicate that nitrite oxidation is important is important for DO

utilization in the DCM. Based on rate profiles, nitrite oxidation ranged from 10-47% of OCR in

the DCM, while experiments indicate maximum percentages of 13-97% (Fig. 3B and 3C).

Stations 4-6—which lack a DCM and SNM—provide a notable contrast, as do ammonia

oxidation rate profiles (Fig. S1).

Peaks in the magnitude of departure from a 1:1 nitrate reduction signal (e.g., isotopic

overprinting by nitrite oxidation), and in *Nitrospina* ASV abundances, further indicate that nitrite

oxidation is active above and below the DCM (Fig. 5). At Stations 2 and 3, nitrite oxidation

rates above the DCM were similar to rates deeper in the water column. Station 1, in contrast,

showed a rate increase within the AMZ. While rate measurements may reflect consumption of

introduced DO, *Nitrospina* 16S rRNA transcripts increased in relative abundance within the

AMZ. RNA samples were collected from the CTD rosette and rapidly filtered, so if these data

capture a response to oxygen exposure, this response is exceptionally rapid and consistent.

Alternatively, these data indicate that *Nitrospina* are active within the AMZ. One possibility is

that *Nitrospina* survive anaerobically for periods of time until provided DO (40). Alternative

oxygen-evolving mechanisms from nitrite have also been proposed for anaerobic methane-

oxidizing bacteria (41) that are active in OMZs (42). Finally, the presence of *Prochlorococcus*

below the DCM may allow low-level DO production via oxygenic photosynthesis (Table 2).

Regardless of the mechanism, isotopic data and 16S rRNA data are consistent with transient

nitrite oxidation coupled to nitrate reduction below the DCM.

However, our primary focus was on oxygen consumption via nitrite oxidation and its

implications for the formation, maintenance, and expansion of OMZs. Two clear and consistent

patterns emerged from both water column profiles and oxygen manipulation experiments: (i)

overall OCR declined with decreasing DO in profiles and experiments, while (ii) nitrite oxidation

increased as a proportion of OCR as DO declined in both profiles and experiments (Fig. 3B and

3C and 4). Variation in the proportion of OCR explained by nitrite oxidation from depth to

depth and experiment to experiment was superimposed on these two patterns. Nitrite oxidation

was significant throughout the OMZ and experiments—typically ranging from 10-40% of

OCR—and sometimes dominant. For example, nitrite oxidation responded rapidly to introduced

DO in the SNM at Station 1, and evidently consumed all DO produced in the DCM at Station 2.

Variation in the proportion of overall OCR attributable to nitrite oxidation could reflect

fluctuations through time (28), as well as differences in substrate availabilities and affinities

relative to DO for different processes consuming DO. Other than nitrite oxidation, aerobic

respiration of organic matter is probably most significant for OCR given relative substrate

concentrations (organic C > CH_4 or reduced S compounds; refs. 42–44), as well as high affinities

for DO among heterotrophic bacteria (45–47). Notably, many aerobic heterotrophs are also

facultatively anaerobic, switching to nitrate reduction under low DO (13, 48). Along with a

general decline in aerobic respiration rates with decreasing DO, this switch to nitrate reduction

may explain the steep declines in OCR that we observed in our experimental data. Production of

nitrite from nitrate reduction may also jumpstart nitrite oxidation, and is consistent with isotopic

data. We suggest that such cryptic nitrite/nitrate cycling may explain the adaptations of

particular *Nitrospina* ASVs to AMZs: although these are low-DO environments, they are also

high-nitrite environments, representing a trade-off in the availability of electron donor and

acceptor. The costs of life at low DO may be offset by the advantages of consistently elevated

nitrite concentrations.

In line with this idea, we found that the proportion of DO consumed by nitrite oxidation

increased at progressively lower DO concentrations in both profiles and experiments (Fig. 3B

and C). This consistent pattern demonstrates that nitrite oxidation increases in relative

importance as DO declines, and suggests that nitrite oxidation may play a critical role in the shift

from low oxygen (an OMZ) to functional anoxia (an AMZ). Moreover, our results indicate that

nitrite oxidation is carried out by just a few *Nitrospina* ASVs—in contrast with the diversity of

other microbial groups that may be involved in overall oxygen consumption (4, 49). Put another

way, even when nitrite oxidation represents a relatively modest proportion of OCR, just one to

three ASVs may be responsible for this contribution to overall oxygen consumption. *Nitrospina*

are therefore instrumental in DO consumption, and disproportionately important as DO declines.

When coupled to nitrate reduction, this could result in an oxygen consumption feedback loop

fueled by cryptic nitrite/nitrate cycling. Rate profiles, oxygen manipulation experiments, natural

abundance measurements, 16S rRNA sequencing, and metagenomic data all favor this idea: as

DO declines, OCR decreases precipitously while nitrite oxidation does not; the low-oxygen

adaptations of particular *Nitrospina* support high abundances and activity; and isotopic

anomalies are consistent with strong coupling between nitrite oxidation and nitrate reduction.

Our data convincingly establish that nitrite oxidation is pivotal in oxygen and nitrogen cycling in

OMZs, and so central in the creation and maintenance of these biogeochemically-important

regions of the ocean.

Materials and Methods

Sample Collection. Samples were collected in April 2017 and June 2018 aboard the *R/V Oceanus*. At each station, conductivity/salinity, temperature, depth, pressure, chlorophyll fluorescence, and photosynthetically active radiation (PAR) were measured by a SeaBird SBE-9plus CTD, SBE-3F temperature sensor, SBE-43 DO sensor, WetLabs ECO-FLR Fluorometer, and Biospherical QCP2200 PAR sensor. Initial casts were used to measure DO and nutrient profiles to guide subsequent sampling. Nutrient samples were analyzed for NH_4^+ and NO_2^- aboard the ship, with additional shore-based analyses of combined $NO_3^- + NO_2^-$ and PO_4^{3-} at the University of California Santa Barbara Marine Science Institute Analytical Lab (*SI Appendix, Supplementary Materials and Methods*).

At all stations, water samples were collected in the upper 100 m to capture variation in the upper water column, and we then sampled across a range of DO levels and nitrite levels to capture variation in the OMZ. Samples were collected at 200, 100, 50, 20, 10, 5, and 1 μM [DO] at all stations. Because the OMZ is shallower and more intense moving south and nearshore in the ETNP (Fig. 1), these DO levels can occur within the upper 100 m, such that upper 100 m and DO-based sampling overlapped. This overlap was greater at Station 1, followed by 2 and 3. At these AMZ stations, samples were also collected at three depths between the 1 μM [DO] level and the SNM.

OCR Measurements and Oxygen Manipulations. Two types of sensor spots were used to measure OCR and manipulate DO, one with a wider range and detection limit of 100 nM (Fibox, Loligo Systems, Viborg, Denmark), as well as trace-level spots with a detection limit of 10 nM (FireSting, Pyroscience, Aachen, Germany) (26). Loligo sensor spots were used for water column profiles given the wider DO range, while FireSting trace-level spots were used to establish deoxygenation experiments and measure OCR within them. Optical sensors were calibrated using DO-saturated water and sodium sulfite-saturated and He-purged water. On the upper end, Fibox sensor spot DO measurements were highly correlated with CTD DO values (r^2=0.996, slope=0.997) for [DO] > 1 μM. On the lower end, we regularly checked the '0 nM' concentration with repeated measurements of sodium sulfite-saturated water and He-purged water, and our lowest values measured in experiments were consistently within 5 nM of 0. In

both profiles and experiments, OCR was measured as the linear decrease in DO over the course

of experiments (25, 50). Experiments were conducted for 24 hours in the dark in a cold van

adjusted to ambient temperature, with separate incubations (and therefore temperatures)

conducted for samples from the upper 100 m, for oxygen/nitrite-based sampling, and for

deoxygenation experiments at each station.

Water column profiles of OCR were measured using 5 replicates at each depth, including

one each with tracer level (5-10% *in situ* concentration measured at sea) addition of $^{15}NH_4^+$ or

$^{15}NO_2^-$ to measure ammonia/nitrite oxidation (see below). Loligo sensor spots were attached to

the inside of 300 mL Wheaton BOD glass bottles using silicone glue prior to the cruise. At sea,

bottles were filled to at least three times overflowing via slow laminar flow, were rapidly capped

and transferred to the cold van, and initial DO measurements were made. After incubation,

endpoint DO measurements were made and then 50 mL samples from relevant bottles were

frozen for nitrite and ammonia oxidation rate measurements.

Oxygen manipulation experiments were conducted in 500 mL serum bottles with

attached FireSting sensor spots. For each experiment, a total of 24 bottles were filled with water

collected at a specific depth, sealed, and then bubbled with ultrapure He gas while DO was

monitored. 8 bottles had tracer-level $^{15}NO_2^-$ additions, 8 bottles had tracer-level $^{15}NH_4^+$

additions, and 8 were unlabeled. In each set of 8 bottles, we established initial DO values

typically ranging from 10s to 1000s of nM (Figure 4). OCR was measured in all bottles based on

starting and ending DO values, and samples for nitrite oxidation were collected from $^{15}NO_2^-$

labeled bottles at the end of the experiments. For all experiments, dedicated bottles were used

for the different ^{15}N labels.

Nitrate isotopic measurements. Both natural abundance and ^{15}N-tracer-based nitrate isotopes

were quantified by use of the denitrifier method (51, 52), in which 20-30 nmoles of NO_3^- were

quantitatively converted to N_2O by a culture of the denitrifying bacterium, *Pseudomonas*

aureofaciens. Product N_2O was cryogenically purified and trapped under flow of ultra-high

purity He, before being introduced to an IsoPrime 100 isotope ratio mass spectrometer. Isotope

ratios were normalized to international reference materials (USGS 32, USGS 34, USGS 35) and

reported using standard delta notation. Any nitrite present in the sample was removed by

addition of sulfamic acid (31). In samples from the upper 100 m, where NO_3^- concentrations

were low, 'carrier' NO_3^- of known isotopic composition was added and sample compositions

were calculated by mass balance. Precision of natural abundance isotope ratio measurements

was ±0.3‰ and ±0.4‰ for $\delta^{15}N$ and $\delta^{18}O$, respectively. $\delta^{15}N$ and $\delta^{18}O$ values were used to

calculate the Δ(15,18) deviation from the 1:1 line following Sigman et al. (35).

Nitrite Oxidation Rates. Nitrite oxidation rates were measured by adding 98 atom percent

(atom%) $^{15}NO_2^-$ to a final concentration of 12-184 nmol L^{-1}—representing 5-10% of *in situ* NO_2^-

concentrations (with the exception of some samples in the upper 100m with no measurable

nitrite)—and measuring the accumulation of ^{15}N label in the NO_3^- pool following Beman et al.

(7). After incubation for ~24 hours, samples were frozen at sea. Upon thawing in the laboratory,

excess $^{15}NO_2^-$ was removed following Granger and Sigman (31). In brief, sulfamic acid (~8-10

$\mu L\ mL^{-1}$) was added, the samples were shaken and allowed to sit for >5 minutes, and then

samples were neutralized by adding NaOH (4 M, ~11 $\mu L\ mL^{-1}$) prior to analysis.

Rates of $^{15}NO_2^-$ oxidation ($^{15}R_{ox}$) were calculated using equation 1 from Beman et al. (7)

adapted from Ward et al. (53):

443
$$^{15}R_{ox} = \frac{\left(n_t - n_{oNO_3^-}\right) \times \left[NO_3^-\right]}{\left(n_{NO_2^-}\right) \times t} \tag{1}$$

where n_t is the atom% ^{15}N in NO_3^- measured at time t, $n_{oNO_3^-}$, is the measured atom% ^{15}N of

unlabeled NO_3^-, $[NO_3^-]$ is the concentration of the NO_3^- pool, and $n_{NO_2^-}$ is the exponential

average atom% of NO_2^- over time t. $n_{NO_2^-}$ was calculated by isotope mass balance based on

initial NO_2^- concentrations, measured isotopic composition, added ^{15}N-labeled NO_2^-, and

measured $^{15}NH_4^+$ oxidation rates (which produce unlabeled NO_2^-; ammonia oxidation rate

measurements followed Beman et al. (23)).

RNA/DNA Extractions, 16S rRNA Sequencing and Analysis, and Metagenome Sequencing

and Analysis are reported in the *SI Appendix, Supplementary Materials and Methods.*

Sequence data are available in the Sequence Read Archive under BioProjects PRJNA192803

(16S data), and PRJNA634212 (metagenomes).

Statistical analyses were carried out in the R statistical environment (RStudio Version 1.0.136).

Linear regressions were performed using the 'lm' function in R; Michaelis-Menten curve fits

were performed using the 'drm' function in the R package drc (54).

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Figures (for initial submission)

Figure 1: Station locations (numbered black circles) in the ETNP plotted on dissolved oxygen

concentrations (in μM) at 100 m depth from the World Ocean Atlas.

Figure 2: Depth profiles of (A) DO (solid lines) and nitrite (data points connected by dashed lines), (B) chlorophyll a, (C) nitrite oxidation rates, and (D) OCR (error bars denote standard deviation of five replicates) show consistent variation across Stations 1-3 (denoted by different colors). Maximum chlorophyll values at Station 1 plot off-axis.

Figure 3: DO consumption via nitrite oxidation as a percentage of OCR with (A) depth, and as a function of DO in (B) depth profiles and (C) oxygen manipulation experiments. Colors denote different stations. Only measurements for DO <18 μM are included in panel B. All experimental bottles with dual nitrite oxidation and OCR are shown in C; and different symbols denote different experiments.

Figure 4: Nitrite oxidation rates (filled symbols) and OCR (open symbols) as a function of DO in oxygen manipulation experiments. Colors denote different stations, with panels displaying data from different experiments: A) Station 1 OMZ edge (20 µM DO), B) Station 2 OMZ edge, C) Station 2 DCM, D) Station 3 DCM, E) Station 3 SNM, F) Station 3 OMZ edge, G) Station 3.5 SNM, and H) Station 4 OMZ edge. Note differences in vertical axes between experiments, and differences in horizontal axes in panels A and B compared with the remaining panels.

Figure 5: Depth profiles of (A) DO (solid lines) and nitrite (data points connected by dashed

lines), (B) Δ(15,18) dual-isotope deviations in nitrate, and *Nitrospina* ASVs as a percentage of

(C)16S rDNA sequence libraries and (D) 16S rRNA sequence libraries. Colors denote different

sampling stations.

Table 1: Oxygen manipulation experiments and calculated parameters for OCR and nitrite

oxidation rates

Sta.	Depth (m)	Type	OCR			Nitrite oxidation		
			v_{max} (nM d^{-1})	K_s (nM)	r^2	v_{max} (nM d^{-1})	K_s (nM)	r^2
1	46	OMZ edge	12005	13390	.892	94.5	611	.861
2	90	OMZ edge	4164	3262	.878	23.3	1876	.347
2	140	DCM	5426	5674	.811	43.9	185	.445
3	130	DCM	2765	18040	.569	149	63.8	.665
3	170	SNM	5973	6382	.757	71.2	38.4	.019
3	110	OMZ edge	4025	4247	.868	23.0	346	.910
3.5	80	SNM	1612	1621	.846	122	34.1	.269
4	100	OMZ edge	2075	1738	.776	47.5	1015	.825

636   **Table 2:** Relative abundance of *Nitrospina* reads, functional genes from *Nitrospina*, and

637   *Prochlorococcus* reads within metagenomes (expressed per million reads)

Sta.	Depth	Type	*Nitrospina*	Nitrite oxido- reductase	Cyto- chrome c	Cyto- chrome bd	Chlorite disumtase	*Prochloro- coccus*
1	25	OMZ edge	1084	2	0	0	1	865
1	87.5	DCM	31920	65	14	2	11	47992
1	100	SNM	21196	37	22	2	7	10765
2	89	OMZ edge	22442	38	12	10	14	57194
2	130	DCM	33290	57	20	4	11	33522
2	160	SNM	21725	37	18	2	4	4157
3	123	OMZ edge	18647	34	10	2	11	54709
3	140	DCM	34227	55	18	4	12	37202
3	180	SNM	17617	30	17	2	4	4881