

Review of Frey et al.

General comments:

Frey et al. used a combined biogeochemical and microbial ecology approach to investigate N₂O production in the ETSP off Peru. More specifically, they used ¹⁵N-labeled incubations to measure N₂O production during ammonium oxidation as well as denitrification and their regulation by O₂ and particulate organic matter. They also measured *archaeal amoA* and *nirS* genes abundances and activity. Overall the manuscript is well written and will make a great contribution to the field as the exact mechanisms regulating N₂O production in ODZs are still not well understood. Of particular importance, few studies have looked at the effects of organic matter addition on N₂O production. I recommend publication after addressing the generally minor considerations below.

Specific comments

Abstract:

Another important finding is that hybrid N₂O formation represented 70-86% of the N₂O production during ammonium oxidation, regardless of the ammonium oxidation rate or O₂ concentrations. One sentence about this should be added to the abstract.

Introduction

Lines 70-75: The distinction between hybrid N₂O production by ammonia oxidizing archaea and chemodenitrification (e.g. nitrite reduction coupled to iron II oxidation) should be better made. Hybrid N₂O formation (mediated by AOA) has been observed in the ODZ water-column, but not chemodenitrification (also referred to as abiotic N₂O production; Wankel et al., 2017), likely due to substrate limitation (Fe, Mn).

line 78: Correct nitrifier-denitrification for denitrification.

Lines 79-81: It should be noted that Frame and Casciotti (2010) only observed higher yields at decreasing O₂ concentrations for high starting cell densities. At lower cell densities (closer to values found in ODZs), the impact of decreasing O₂ on N₂O yield was much lower than observed in other studies.

Lines 102-104: Charpentier et al (2007) also suggested that nitrifier-denitrification is enhanced by high concentration of organic particles, which creates high NO₂- and low-O₂ microenvironments.

Lines 113-114: It would also be relevant to look at *nor* genes which are encoding nitric oxide reductase.

Materials and methods:

Line 136: It is not clear why a 3 mL He helium headspace is created before incubating, since it will impact in-situ O₂ concentrations.

Line 136-137: I assume purging is done to avoid O₂ contamination? What is the O₂ threshold defining anoxia here? One potential problem with purging is that it also removes other gases (e.g., H₂S) involved in autotrophic denitrification (for instance, see Callbeck *et al.*, 2018).

Lines 150-153: How did O₂ vary during the incubations? These data should perhaps be included as part of the supplementary materials.

Line 153: Explain the rationale for using particles >50 μm.

Lines 192-219: Plots showing increase in ¹⁵N labeled products over time should be included in the supplementary materials. Were the relationships always linear?

Lines 228-229: These nirS primers exclude epsilon-proteobacteria (Murdock, *et al.*, 2017). Epsilon proteobacteria are often the dominant portion of autotrophic sulfur oxidizers in sulfidic waters (e.g., Grote *et al.*, 2008), thus this aspect should be discussed.

Line 256: Add accession number.

Results:

Lines 282-283: Could a contour plot of chlorophyll concentration added to the supplementary material for reference?

Lines 334-335: This result is a bit puzzling as previous studies (e.g., Dalsgaard *et al.*, 2014), observed fifty percent inhibition of N₂O production by denitrification at about 300 nM O₂. These observations are also unlike results from their *in situ* O₂ gradient experiments.

Lines 349-350: It is also surprising to observe the highest yield for N₂O production at highest O₂ concentrations, for which N₂O production should be inhibited (Dalsgaard *et al.*, 2014).

Discussion:

Lines 421-426: Some of these are likely causal relationships.

Lines 425-426: This suggests that when NO₃⁻ is abundant, denitrifying bacteria are less likely to use NO₂⁻ (either from their internal pool or outside the cell) for N₂O production during denitrification.

Line 441: What is the detection limit for [N₂O]?

Lines 441-444: Bourbonnais *et al.* (2017) used biogeochemical tracers (N₂O concentrations and isotopes) that integrates over longer timescales compared to ¹⁵N-labeled incubations, which are more like taking a snapshot in time. Therefore, discrepancies between N₂O production rate is expected and should be discussed in this context.

Line 451: Cite Fassbender et al. (2018) that discusses impacts of eddies on biogeochemical processes at different scales.

Lines 443: The error on this higher rate estimate seems rather large (in Figure 3, p).

Lines 458-460: This part is confusing. The O₂ threshold for reductive N₂O production should be higher than for N₂O consumption, not the converse. In other words, nitric oxide reductase should be more O₂ tolerant than nitrous oxide reductase (Dalsgaard et al., 2014). Otherwise, N₂O would not accumulate.

Lines 445-446: I do not understand this statement.

Lines 479-481: This hypothesis is also supported by a rather long turnover time for NO₂⁻ as inferred from the δ₁₈O of NO₂⁻, which is generally fully equilibrated with water in offshore waters (Bourbonnais et al., 2015). This is not the case in coastal waters, where NO₂⁻ seems to be more dynamic (see and cite Hu et al., 2016).

Lines 495-496: How can these contrasting results be reconciled?

Lines 522-524: If hybrid N₂O formation during AOA is purely (or even partly) abiotic, then measured rates would be overestimated as HgCl₂ would not stop N₂O production at the end of the incubations. For how long were these samples stored before being measured? This point should be better discussed.

Lines 565-566: What was the chlorophyll concentration in the center of the eddy?

Lines 641-643: N₂O emission to the atmosphere are possible only if the water is upwelled.

Lines 649-652: Temporal variability is particularly not well captured in observational studies.

Figure legends:

Rename Figure 7: N₂O production after additions of...

Figures 2 and 3 are too small. Legend (station #) is almost impossible to read.

Figure 5: Samples impacted by denitrification should be more clearly indicated (by a circle or rectangle and in the Figure legend) in Figure 5b.

Supplements:

Figures S1: I recommend expanding the scale at lower O₂ concentrations since this is the focus of the paper.

Figure S5: Add linear regression and r-square for natural samples in the zoom up plot.

Figure S6: Since there are only a few data points for $[\text{NO}_2^-]/[\text{NO}_3^-]$ higher than 0.10, I don't think the outlier (light gray dot) can be removed. There is much more scatter in Figure 5 in Ji et al. (2018) for the same relationship.

Other references:

Callbeck, C. M., Lavik, G., Ferdelman, T. G., Fuchs, B., Gruber-Vodicka, H. R., Hach, P. F., ... & Schunck, H. (2018). Oxygen minimum zone cryptic sulfur cycling sustained by offshore transport of key sulfur oxidizing bacteria. *Nature communications*, 9(1), 1729.

Charpentier, J., Farias, L., Yoshida, N., Boontanon, N., & Raimbault, P. (2007). Nitrous oxide distribution and its origin in the central and eastern South Pacific Subtropical Gyre. *Biogeosciences Discussions*, 4(3), 1673-1702.'

Fassbender, A. J., Bourbonnais, A., Clayton, S., Gaube, P., Omand, M., & Franks, P. J. S. (2018). Interpreting mosaics of ocean biogeochemistry. *Eos*, 99(10.1029).

Hu, H., Bourbonnais, A., Larkum, J., Bange, H. W., & Altabet, M. A. (2016). Nitrogen cycling in shallow low-oxygen coastal waters off Peru from nitrite and nitrate nitrogen and oxygen isotopes. *Biogeosciences*, 13(5), 1453-1468.

Murdock, S. A., & Juniper, S. K. (2017). Capturing compositional variation in denitrifying communities: a multiple-primer approach that includes Epsilonproteobacteria. *Appl. Environ. Microbiol.*, 83(6), e02753-16.