

Reply to Review Comments

We are grateful for the constructive comments and helpful suggestions of the two Referees. Below are detailed responses to all the comments and corresponding explanations of the revisions made to the manuscript. Line numbers cited in the replies (highlighted) refer to the revised manuscript document version with tracked changes displayed

Referee 1

Dear authors, this was a pleasure to review your manuscript. It raises a very interesting topic of application of stable isotope studies for better understanding of soil N cycle. The manuscript presents a few of very original analytical approaches, like NO and NO₂⁻ isotopic analyses (as one of the very first for soil studies) and application of D17O to trace NO₃ and NO₂ soil transformations. The combination of all the approaches and the construction of the NO isotope model is very complex and challenging to present in an understandable form, but authors managed this very well. The manuscript is well organised, the results are well documented and supplement contains a lot of additional information precious for the readers who will further apply or develop the presented approach.

Reply: We thank Referee #1 for the positive feedback.

Comment 1: I could have one suggestion of expanding the analytics, maybe useful for your future studies. Since you used Chilian NO₃ with the D17O anomaly you could also monitor this anomaly in NO₂⁻ (this may be difficult due to low concentrations) or in NO or N₂O. This would allow you to determine the extend O-exchange and no further consideration of two scenarios: with and without O-exchange will be needed. This will bring more clarity to the whole study. An example of using D17O of N₂O to determine O-exchange can be found in Lewicka-Szczebak et al. (2016, BG).

Reply: We agree with the Referee that $\Delta^{17}\text{O}$ analysis of NO₂⁻ could provide valuable insights into the degree of oxygen isotope exchange between NO₂⁻ and H₂O during the anoxic incubation, thereby offering more constraints and confidence to the isotopic modeling. However, we had concerns about the feasibility of $\Delta^{17}\text{O}$ -NO₂⁻ analysis in this case because NO₂⁻ in water samples can undergo oxygen isotope exchange with H₂O during sample processing, preservation, and storage (e.g. even for samples frozen under -20°C) (Casciotti et al., 2007). Therefore, measuring soil NO₂⁻ for its $\Delta^{17}\text{O}$ values is not trivial, and will require comprehensive efforts to demonstrate its robustness throughout the sequence of soil extraction, extract processing, and sample storage. These efforts can be largely facilitated by development of $\Delta^{17}\text{O}$ -NO₂⁻ reference materials, which are currently lacking.

Analysis and interpretation of $\Delta^{17}\text{O}$ of soil NO are confounded by the ozone oxidation of NO to NO₂ during the NO collection and the fact that NO₂ is collected in the triethanolamine (TEA) solution as both NO₂⁻ and NO₃⁻. Therefore, $\Delta^{17}\text{O}$ or $\delta^{18}\text{O}$ of NO₂⁻/NO₃⁻ collected from soil emitted-NO does not contain direct information about soil NO turnover. These technical aspects have been extensively discussed in our original method paper (Yu and Elliott, 2017).

We have revised the manuscript to include Lewicka-Szczebak et al. (2016) and to note that our understanding of NO₂⁻ oxygen isotope exchange and reaction reversibility can benefit from robust soil $\Delta^{17}\text{O}$ -NO₂⁻ determination and calibration in the future (Line 569-575).

I have just a few very minor comments:

Comment 2: - Fig. 6 - do you assume that the abiotic NO cannot be further reduced to N₂O?

Reply: Due to lack of direct observational constraints, we did not assume any specific production or consumption pathways for NO yield from abiotic NO₂⁻ reactions in the isotopologue-specific model. As such, the model simulates net NO production, rather than gross rates. Specifically, based on the results from the abiotic incubation, we assumed that the net abiotic NO production from NO₂⁻ followed a pseudo-first order kinetics with respect to NO₂⁻ with an apparent stoichiometric coefficient for net NO production from NO₂⁻ of 0.52 (Line 510-513 of the original manuscript). This modeling parameterization implicitly accounts for parallel or competing abiotic NO production pathways in the soil, as well as potential NO consumption through abiotic reactions (e.g., chemo-denitrification of NO to N₂O; Line 365-380 of the original manuscript). In the revised manuscript, we have revised Fig 6 and its caption to clarify that the modeled abiotic NO production represents net NO yield, rather than gross NO production.

Comment 3: - L 609 - what do you mean here with "modified isotopologue-specific model" - this term was not used before in the manuscript and it is not clear if you just refer to the presented NO isotope model or sth else

Reply: It is mentioned in the original manuscript that the isotopologue-specific model we used to simulate co-occurring denitrification and NO₂⁻ re-oxidation was modified from a model of co-occurring nitrification and NO₃⁻ consumption we developed previously for well-aerated soils (Line 492-495 of the original manuscript). We have removed "modified" here to prevent any confusion.

Comment 4: - L 624 - what is "more normal" isotope effect?

Reply: In this study, we follow the convention to define kinetic isotope effect (Line 78-82 of the original manuscript). Under this definition, a normal kinetic isotope effect occurs when reaction rate constant of light isotopologues is higher than that of heavy isotopologues. Thus, normal kinetic isotope effects are expressed by positive η (η) values in this study, in opposition to inverse kinetic isotope effects, which have negative η values. Here, our estimated isotope effect for nitric oxide reduction ($^{15}\eta_{\text{NOR}}$) is between -8‰ and 2‰, higher than the previously reported $^{15}\eta_{\text{NOR}}$ for fungal nitric oxide reductase (i.e. -14‰). We have revised the manuscript to clarify that "more normal" is used here to describe our estimated $^{15}\eta_{\text{NOR}}$ being closer to zero (Line 631).

Comment 5: - Section 4.3 - I wonder why you do not consider NO₂⁻ oxidation to NO₃⁻ for oxic and suboxic conditions. If this process was so intensive under anoxic conditions, why it should not be active under oxic and suboxic conditions?

Reply: We did not explicitly consider aerobic NO₂⁻ oxidation to NO₃⁻ under oxic and hypoxic conditions because NO₂⁻ concentration was below the detection limit in both incubations (Line 315-317 of the original manuscript), suggesting that the two steps of nitrification (i.e. NH₄⁺ oxidation to NO₂⁻ and NO₂⁻ oxidation to NO₃⁻) were tightly coupled under these conditions (Line 651-653 of the original manuscript). Therefore, in the isotopologue-specific model of co-occurring nitrification and NO₃⁻ consumption, the two nitrification steps were lumped into a gross flux of NH₄⁺ oxidation to NO₃⁻ (Line 655-659 of the original manuscript; Text S5 in the Supplement) (Yu and Elliott, 2018). The excellent agreement between the modeled and

measured data (i.e., NH_4^+ and NO_3^- concentrations and $\Delta^{17}\text{O}-\text{NO}_3^-$; Figure 3) under both oxic and hypoxic conditions confirms that this model configuration is appropriate.

The NXR-catalyzed anaerobic NO_2^- re-oxidation and/or $\text{NO}_3^-/\text{NO}_2^-$ interconversion, which prevailed in the anoxic incubation, are considered not important in the oxic and hypoxic incubations. The results from the anoxic incubation, together with findings from previous studies (e.g. Wunderlich et al., 2013), suggest that NO_2^- accumulation coupled with O_2 deprivation is the key trigger of anaerobic NO_2^- re-oxidation by nitrite-oxidizing bacteria (NOB). This point has been emphasized in multiple places throughout the manuscript (Line 502-505, 598-604, and 839-846 of the original manuscript). The lack of NO_2^- accumulation in the oxic and hypoxic incubations suggests that NOB mainly performed aerobic NO_2^- oxidation to gain energy. No revision was made based on this comment.

Reply to Dr. Wolfgang Wanek

The paper reports on the isotopic fractionation of source and sink processes underlying soil NO emissions, NO emissions being important for atmospheric chemistry and as a soil N loss pathway. I am impressed by this study, proving in-depth analysis of isotopic constraints on formation and consumption pathways of soil NO, and partitioning the contribution of nitrifiers and denitrifiers as well as abiotic reactions. The approach taken with aerobic, suboxic and anoxic soil incubations combined with inorganic N additions in live and sterile soils, N and O isotope measurements in inorganic soil N and NO, amended by isotope fractionation and flux modeling provides a most complete assessment of NO source and sink processes. This study therefore highlights that stable isotope measurements in inorganic soil N with those in NO and N_2O can help in source attribution of these important atmospheric gases.

Reply: We are grateful for the encouraging remarks and positive feedback.

Comment 1: Minor corrections can be found in the annotated PDF.

Reply: We have incorporated all the corrections and edits into the revised manuscript. Thank you.

Comment 2: Lines 59-61: There are also complete ammonia oxidizing Nitrospira, that catalyze the whole nitrification reaction sequence from ammonia to nitrate in one organism (comammox bacteria).

Reply: We agree with Dr. Wanek that recent breakthrough in discovering completely nitrifying Nitrospira has broadened our understanding of microbial nitrification (Daims et al., 2015). However, to our best knowledge, studies on trace gas production (mainly as N_2O) by comammox bacteria are just starting (Kits et al., 2019), and whether and how free NO can be produced and released from complete nitrification remain unknown. There is also postulation that the revealed high affinity of comammox bacteria to ammonia may indicate a better adaptation of comammox bacteria to low-nitrogen environments (Kits et al., 2017; Kuyper, 2017). Therefore, for the sake of simplicity, we prefer not to include comammox bacteria in the discussion. Importantly, because NO_2^- concentration was below the detection limit during the oxic and hypoxic incubations (Line 315-317 of the original manuscript), the two nitrification steps were lumped into a gross flux of NH_4^+ oxidation to NO_3^- in our isotopologue-specific model (Line

655-659 of the original manuscript; Text S5 in the Supplement) (Yu and Elliott, 2018). Thus, our modeling scheme of nitrification is not in conceptual conflict with complete nitrification.

Comment 3: Line 80 and throughout the MS: it should always be kinetic isotope fractionation and equilibrium isotope fractionation.

Reply: Agreed. We have revised the manuscript to adopt a consistent use of isotope terminology.

Comment 4: Line 189: please provide xg (RCF) instead of rpm.

Reply: We have converted rpm (2000) to RCF (3400g) in the revised manuscript (Line 190).

Comment 5: Line 374: The reference Zhu-Baker et al. (2015) is missing in the reference list and should be Zhu-Barker.

Reply: Thank you. We have corrected this mistake and double-checked the entire reference list to ensure its accuracy.

References

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