



Isotopologue effects during N₂O reduction in soils and in pure cultures of denitrifiers

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[1] Site preference (SP), the difference in $\delta^{15}\text{N}$ between the central and outer nitrogen atoms in N₂O, is a powerful approach for apportioning fluxes of N₂O from soils to nitrification and denitrification (Sutka et al., 2006). A critical aspect of the use of SP data to apportion sources of N₂O to nitrification and denitrification is the need to evaluate data for isotope shifts that may have occurred during N₂O reduction in soils prior to its escape to the atmosphere. We present data on the isotopologue effects during reduction of N₂O during anaerobic incubation of soils and pure cultures of denitrifying bacteria. Isotopic enrichment factors for N₂O reduction in soil mesocosms experiments varied between -9.2 and -1.8‰ for nitrogen and between -25.1 and -5.1‰ for oxygen. In pure cultures of *Psuedomonas stutzeri* and *Psuedomonas denitrificans* we observed isotopic enrichment factors for SP of -5.0 and -6.8‰ , respectively. We further find that N₂O consumption produces consistent relationships between $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ and $\delta^{18}\text{O}$ and the $\delta^{15}\text{N}$ of the central N atom in N₂O of 2.5 and 1.6, respectively, which are clearly diagnostic of this process. Our results indicate that SP may be altered during reduction of N₂O and thus bias evaluations of its origins. To understand the impacts of N₂O reduction in soil flux studies on source isotope signals we modeled the isotope effects of N₂O production occurring simultaneous with reduction and find increasingly curvilinear relationships between $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ and $\delta^{18}\text{O}$ and $\delta^{15}\text{N}^\alpha$ with increased reduction. Consequently, a deviation from the linear mixing relationship between soil-derived and atmospheric N₂O is an indication of extensive reduction. On the basis of our characterization of isotopic fractionation during N₂O reduction, we show that the rate of reduction would have to be substantially greater than 10% of that of production to impact SP estimates of N₂O from denitrification by more than a few percent. Nonetheless, reduction results in a small, but potentially important, increase in SP away from values proposed for bacterial denitrification (0‰) toward those associated with production from nitrification (33‰) (Sutka et al., 2006). On this basis, estimates of the proportion of N₂O derived from denitrification obtained from SP values are underestimates and therefore conservative.

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1. Introduction

[2] Nitrous oxide is an important greenhouse gas that is strongly influenced by microbial production and consumption. Agricultural soils annually add about 2.5 Tg of N₂O-N to the atmosphere [*Intergovernmental Panel on Climate*

Change, 2001] and N₂O abatement in agriculture has been identified as one of several high-impact strategies for stabilizing atmospheric greenhouse gas concentrations [*Caldeira et al.*, 2004]. Effective mitigation, however, requires a process-level understanding of N₂O sources, and even though agricultural soils represent >50% of anthropogenic N₂O emissions worldwide [*Mosier et al.*, 1998a, 1998b; *Robertson*, 2004], the microbial source of N₂O in most soils remains ambiguous. Both nitrification and denitrification produce N₂O, and because these processes are under very different environmental controls [*Robertson and Groffman*, 2006] they are responsive to very different management strategies [*Matson et al.*, 1989]. Understanding the microbial source of N₂O in high-flux soils is thus an important research objective.

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[3] Efforts to determine microbial pathways of N₂O production have primarily relied upon the application of nitrification inhibitors such as acetylene [Berg *et al.*, 1982; Robertson and Tiedje, 1987; Davidson, 1992], isotopically enriched substrates [Well *et al.*, 2006; Menyailo and Huntgate, 2006b], or observation of characteristic changes in the natural isotopic abundances of ¹⁵N and ¹⁸O in N₂O [Webster and Hopkins, 1996; Ostrom *et al.*, 2000; Yamulki *et al.*, 2000, 2001; Perez *et al.*, 2000, 2001; Wrage *et al.*, 2004a]. These approaches have significant limitations. Acetylene-inhibition is limited to short-term experiments and presents methodological challenges related to microsite heterogeneity, unequal diffusion of inhibitor into microsites, and unintended effects of inhibitors on microbial activity that collectively can bias results [Tiedje *et al.*, 1989; Tilsner *et al.*, 2003; Wrage *et al.*, 2004b, 2004c; Bateman and Baggs, 2005]. Use of isotopically enriched substrates can alter activity by changing substrate concentrations and can provide erroneous signals if substrates are being produced from natural abundance sources over the time course of measurement [Madsen, 1998]. Consequently, a number of studies have been directed at distinguishing N₂O derived from nitrification and denitrification based on the magnitude of fractionation during N₂O production in the absence of inhibitors [Webster and Hopkins, 1996; Mandernack *et al.*, 2000; Yamulki *et al.*, 2000, 2001; Perez *et al.*, 2000, 2001; Bol *et al.*, 2003, 2004; Tilsner *et al.*, 2003]. These efforts are based on determination of the difference in isotopic composition (fractionation) of the substrate (ammonium or nitrate) and N₂O during production by the soil microbial community. Quantitative estimates of the relative importance of nitrification and denitrification based on this approach are confounded because the $\delta^{15}\text{N}$ of the substrates can fluctuate spatially and temporally [e.g., Ostrom *et al.*, 1998] and the magnitude of isotopic fractionation for microbial N cycling pathways tend to be variable [e.g., Ostrom *et al.*, 2002].

[4] Recently, the intramolecular distribution of isotopes within N₂O has been used in addition to bulk isotope data to constrain origins and global budgets [Toyoda and Yoshida, 1999; Breninkenmeijer and Röckmann, 2000; Yoshida and Toyoda, 2000; Toyoda *et al.*, 2001]. Nitrous oxide is an asymmetric molecule consisting of two N atoms with unique covalent bonds. Equilibrium and kinetic isotope effects in natural reactions result in distinct abundances of ¹⁵N within the central (α) or outer (β) N atoms [Miller and Yung, 2000; Yoshida and Toyoda, 2000]. The difference in $\delta^{15}\text{N}$ between the α and β atoms in N₂O is termed site preference (SP). In pure microbial incubation studies, SP has been found to have unique values reflecting the microbial production pathway [Sutka *et al.*, 2003, 2006; Toyoda *et al.*, 2005]. Production of N₂O via nitrite or nitrate reduction, whether by nitrifying or denitrifying organisms, has been shown to result in particularly low SP values in most studies (-5‰ [Toyoda *et al.*, 2005]; average of 0‰ [Sutka *et al.*, 2006]) with one notable exception (-23‰) that may have been affected by inorganic reduction [Toyoda *et al.*, 2005]. Production of N₂O via hydroxylamine oxidation by nitrifying or methane oxidizing microorganisms, has been shown to have SP values that average 33‰ that is clearly distinct from the SP associated with production by nitrate or nitrite reduction [Sutka *et al.*, 2006]. Furthermore,

SP is constant over the course of reactions and independent of substrate isotopic composition whereas $\delta^{15}\text{N}$ and $\delta^{18}\text{O}$ values change markedly during reactions and are dependent upon the isotope values of the substrates [Sutka *et al.*, 2003, 2006; Toyoda *et al.*, 2005]. Thus, while pure culture studies of N₂O production using microorganisms using the full suite of reduction enzymes is still yet to be done [Schmidt *et al.*, 2004], SP is emerging as a noninvasive tool for apportioning N₂O flux to specific microbial pathways [Well *et al.*, 2006; Sutka *et al.*, 2006].

[5] While the SP values of N₂O have the potential to reveal microbial production pathways, consumption (or reduction) of N₂O during denitrification can impart an isotope effect that could bias or obscure source isotope values [Tilsner *et al.*, 2003; Wrage *et al.*, 2004a; Yamagishi *et al.*, 2007]. Rates of N₂O reduction independent of gross denitrification are rarely determined in soils; hence it is difficult to know the importance of this process a priori [e.g., Bandibas *et al.*, 1994; Cavigelli and Robertson, 2001]. Denitrification results in cleavage of the N and O covalent bond in N₂O, which based on kinetic isotope theory, is expected to result in an increase in the ¹⁵N content of the α position and, therefore, an increase in SP [Popp *et al.*, 2002; Yamagishi *et al.*, 2007]. The increase in SP in response to reduction would result in a shift away from values associated with denitrification (0‰) toward those associated with nitrification (33‰) [Sutka *et al.*, 2006] and result in an overestimate of the importance of nitrification.

[6] Characterization of fractionation during N₂O reduction may resolve the importance of this process and provide an ability to correct for resulting isotope shifts. Fractionation during N₂O reduction is, at minimum, a two step process involving diffusion into the cell and enzymatic reduction. Consequently, the degree of fractionation will reflect the relative importance of the isotope effect during diffusion, which is likely small, and that associated with enzymatic reduction which is expected to be large [e.g., Ostrom *et al.*, 2002]. The relative importance of diffusion and enzymatic reduction of N₂O is expected to vary with the degree of soil water saturation and other soil properties such that fractionation is small when soils are saturated (supply limited by diffusion) and large when soils are dry (diffusion nonlimiting). Thus an important goal of our research is to constrain fractionation during reduction of N₂O as a means to correct for the influence of isotope effects on source apportionment. In this study we use the term “isotopologue” to refer to both bulk nitrogen (average of $\delta^{15}\text{N}^{\alpha}$ and $\delta^{15}\text{N}^{\beta}$) and oxygen isotope values and the $\delta^{15}\text{N}$ of the individual nitrogen atoms within the N₂O molecule. Given that molecules of ¹⁵N¹⁴N¹⁶O and ¹⁴N¹⁵N¹⁶O differ in the position of ¹⁵N enrichment but not mass, these molecules are appropriately referred to as “isotopomers.” We present results on isotopologue effects during N₂O reduction in both pure culture and soil mesocosm experiments and an assessment of the importance of N₂O reduction on source apportionment studies.

2. Methods

2.1. Site Description

[7] Our field experiments were conducted at the Kellogg Biological Station Long Term Ecosystem Research

(KBS LTER) site located in southwest Michigan, USA (42°24' latitude, 85°24' longitude). Native vegetation in the area is beech-maple and oak-hickory forests interspersed with open oak savannas [Burbank *et al.*, 1992]. Most of the area was cleared for agriculture in the mid-1800s and modern agronomic yields are typical of those in the North Central region as a whole. Soils at the site are Kalamazoo (fine-loamy) and Oshtemo (coarse-loamy) mixed, mesic typic Hapludalfs developed from glacial outwash deposited at the end of the Wisconsin glacialation (J. R. Crum and H. P. Collins, KBS soils, 1995, available at www.lter.kbs.msu.edu/soil/characterization). The main experimental design of the KBS LTER site consists of a series of replicated ecosystems arranged along a management intensity gradient and this design facilitates an understanding of management history on N₂O fluxes and production pathways (<http://lter.kbs.msu.edu>).

2.2. N₂O Reduction Experiments Within Soil Mesocosms

[8] Soils for N₂O reduction experiments were collected from the KBS LTER in June of 2002 from 3 replicates of 3 different treatments: (1) a conventionally managed corn-soybean-wheat row-crop ecosystem; (2) a historically cultivated early successional field abandoned after spring planting in 1989; and (3) a late successional deciduous forest. Approximately 4 kg of soil were collected from each of the nine experimental plots by taking 20 cores from the upper 25 cm of soil from each of three different treatment plots from the replicated series of cropped and unmanaged ecosystems. A 4 mm sieve was used to homogenize soil. Aliquots were covered and allowed to air dry for ca. three weeks. Dry soil (100 g) was added to mesocosms (1 L glass Mason jars bearing lids fitted with butyl rubber septa) and packed to a volume of ~80 mL for a target bulk density of 1.2 g dry soil cm⁻³. Water was added to achieve ~85% water filled pore space [Bergsma *et al.*, 2002]. Heterogeneities in the soil, however, resulted in differences in water filled pore space and subsequent measurements revealed values between 91 and 100%. To prevent N₂O production over the course of the N₂O reduction experiments soils were incubated for 2 weeks in the absence of oxygen to exhaust alternative substrates (primarily nitrate) for denitrification after which production was not evident. To initiate N₂O reduction and maintain anoxic conditions, sealed jars were flushed with pure N₂ and then amended with 500 μL of pure N₂O that was delivered using a gas tight syringe (Hamilton). This resulted in a headspace concentration appreciably greater than that observed in typical soil environments but assured sufficient gas for isotopologue analysis. These conditions also assure that concentration is not a factor in contributing to the diffusional limitation of enzymatic reduction within cells (see section 3.1). Headspace samples of 500 μL were taken periodically and stored in vialtainers (BD Diagnostics) previously purged with pure N₂ and brought to atmospheric pressure. Samples were stored for no longer than 1 month prior to isotopic analysis.

2.3. N₂O Reduction Within Pure Culture Experiments

[9] *Pseudomonas stutzeri* (provided by J. M. Tiedje) and *Pseudomonas denitrificans* (ATCC 13867 purchased from ATCC, Manassas, Virginia) were cultured from a frozen stock and maintained in Citrate Minimal Medium (CMM)

[Anderson *et al.*, 1993]. These suspensions were derived from 100 mL cultures in CMM incubated at 30°C in the case of *P. denitrificans* and 25°C for *P. stutzeri*. A single culture of each species was grown to late log phase of growth and concentrated (centrifugation: 10,000 g for 10 min at 5°C). The cell pellet was washed twice (20 mL, sterile CMM) and resuspended in 20 mL of sterile CMM. For each culture, a 12 mL exetainer vial (Labco, UK) was prepared with 2 mL of the concentrated cell suspension and 1 mL of a 0.01 M KNO₃ stock solution. The tubes were sealed with a N₂ headspace and incubated at 30°C and 22°C for *P. denitrificans* and *P. stutzeri*, respectively. Isotopically characterized N₂O (20 μL) was injected into the headspace and an initial sample was taken after the headspace gases were allowed to mix for 2 min. The headspace was sampled for N₂O approximately every 20 min using a 100 or 500 μL gas tight syringe. The gas sample was immediately analyzed for isotopologue abundances.

2.4. Analysis of Isotopologues

[10] Gas samples were analyzed on a multicollector GV Instruments IsoPrime Mass Spectrometer interfaced with a continuous flow Trace Gas Inlet System for purification and concentration of N₂O. Isotopic analysis of N₂O involves removal of CO₂ and water using chemical scrubbers (Carbosorb and magnesium perchlorate) and cryogenic trapping followed by chromatographic separation on a Poraplot Q gas chromatographic column with He as the carrier gas within the Trace Gas system. The effluent from the Trace Gas system is subsequently allowed to enter the mass spectrometer for isotopic characterization. Determination of N₂O concentrations were based on the intensity of the ion current generated by the mass 44 detector in comparison to standards consisting of air spiked with known quantities of pure N₂O. Calibration was performed daily and precision of N₂O concentrations for replicate standards and samples were less than 5%. The multicollector mass spectrometer is able to simultaneously monitor 5 masses of interest for N₂O isotopologues; 30, 31, 44, 45 and 46. We follow the convention of Toyoda and Yoshida [1999] in defining the central and outer nitrogen atoms as α and β, respectively, elsewhere identified as atoms 2 and 1, respectively [Breninkenmeijer and Röckmann, 2000].

[11] Values for δ¹⁵N, δ¹⁸O and δ¹⁵N^α are obtained from the ratio of the 45:44, 46:44 and 31:30 ion beam ratios, respectively. We applied corrections for the contribution of ¹⁷O to masses 31 and 45 and for a small degree of rearrangement of ¹⁵N between the α and β positions within the ion source [Toyoda and Yoshida, 1999; Breninkenmeijer and Röckmann, 2000; Sutka *et al.*, 2003, 2004a]. The value of δ¹⁵N^β is calculated given that δ¹⁵N is the average of δ¹⁵N^α and δ¹⁵N^β [Toyoda and Yoshida, 1999; Breninkenmeijer and Röckmann, 2000]. Final δ¹⁵N, δ¹⁸O, δ¹⁵N^α and δ¹⁵N^β values were calculated using the approach outlined by Toyoda and Yoshida [1999] and reported with respect to Air as the international standard with the exception of δ¹⁸O values that are reported with respect to VSMOW.

[12] There is disagreement with respect to the δ¹⁵N^α and SP of tropospheric air [Toyoda and Yoshida, 1999; Kaiser *et al.*, 2004] that has recently been resolved [Westley *et al.*, 2007]. We report here our data correction method and internal laboratory standard values so that additional cor-

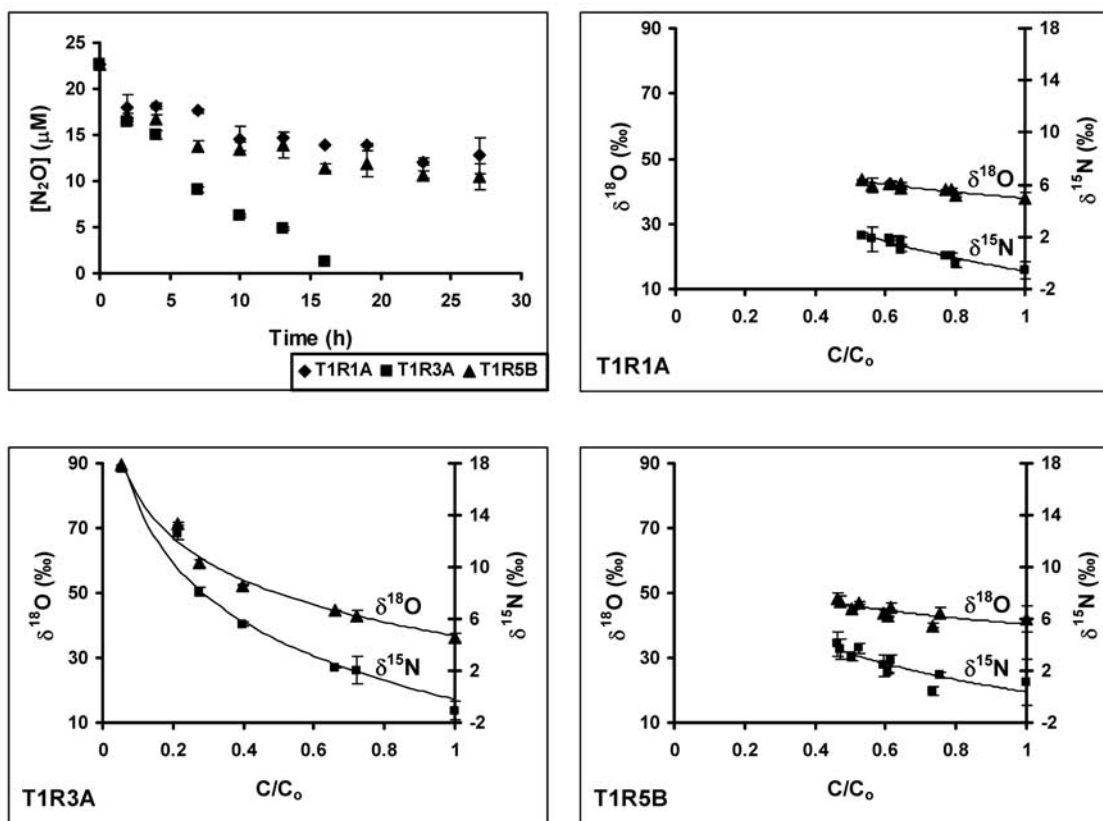


Figure 1. Concentration, $\delta^{15}\text{N}$, and $\delta^{18}\text{O}$ data for N₂O within soil mesocosm N₂O reduction experiments using soils under conventional agricultural treatment from three replicate plots within the KBS LTER main site. Isotope values are expressed in relation to the concentration (C) at any point in time divided by the initial concentration (C_0) following the Rayleigh model (equation (3)). The slope of a natural log fit to the data equates to the fractionation factor for each experiment and these values are listed in Table 1. Error bars represent 2 standard deviations around the mean of three replicates for each treatment. For most data points, the error bars are less than the size of the markers.

rections can be made, if needed. Our internal laboratory standard was calibrated in collaboration with S. Toyoda and N. Yoshida at the Tokyo Institute of Technology using the equations of *Toyoda and Yoshida* [1999], and we commonly obtain the values of $\delta^{15}\text{N}^\alpha$ and $\delta^{15}\text{N}^\beta$ of 16.4 and 2.4‰, respectively, reported by these authors for tropospheric air. The $\delta^{15}\text{N}$, $\delta^{15}\text{N}^\alpha$ and $\delta^{18}\text{O}$ of our laboratory pure N₂O standard is 1.6, 14.9 and 41.7‰, respectively.

[13] The multicollector mass spectrometer that we use is the first to determine all masses needed for determination of N₂O isotopologues simultaneously. On other instruments samples are analyzed twice with adjustments to ion source parameters between analyses [e.g., *Breninkenmeijer and Röckmann*, 2000; *Röckmann et al.*, 2003]. While the multicollector approach offers significant time savings it challenges the analyst to demonstrate appropriate accuracy in several isotope ratios over a wide range of sample concentrations (defined as “linearity”). We have found, in practice, that the optimal ion source parameters to assure precision and linearity for $\delta^{45/44}$ and $\delta^{46/44}$ are not optimal for $\delta^{31/30}$. Consequently, we have optimized analysis for $\delta^{31/30}$, since these are the masses of lowest intensity, which results in a small deterioration in the precision of $\delta^{45/44}$ and $\delta^{46/44}$ as

well as some loss in sensitivity. Similar challenges in isotopologue linearity were reported by *Röckmann et al.* [2003]. With this adjustment we are routinely analyzing samples as low as 3.5 nmol N₂O with precisions for $\delta^{15}\text{N}$, $\delta^{18}\text{O}$, $\delta^{15}\text{N}^\alpha$, $\delta^{15}\text{N}^\beta$, and SP better than 0.3, 0.7, 0.9, 0.9, and 1.3‰, respectively. Prior to the initiation of our revised ion source tuning protocol the instrument was linear for $\delta^{45/44}$ and $\delta^{46/44}$ but not linear for $\delta^{31/30}$ below 6 nmol N₂O. Samples from the soil mesocosm experiments did not reach this threshold. Consequently, we report $\delta^{15}\text{N}$ and $\delta^{18}\text{O}$ values for these samples but do not report $\delta^{15}\text{N}^\alpha$, $\delta^{15}\text{N}^\beta$ or SP.

3. Results and Discussion

3.1. N₂O Reduction in Soil Mesocosms

[14] The concentration of N₂O in all soil mesocosm experiments exhibited declines over the course of the experiment consistent with microbial reduction though rapid initial declines in some treatments may reflect initial equilibration between the headspace and soil (Figures 1 and 2). Correlations of isotopologue values with the natural log of the change in concentration is a good indication that reduction is the predominant process occurring in the mesocosms [*Ostrom et al.*, 2002]. A rapid decline in N₂O was evident in

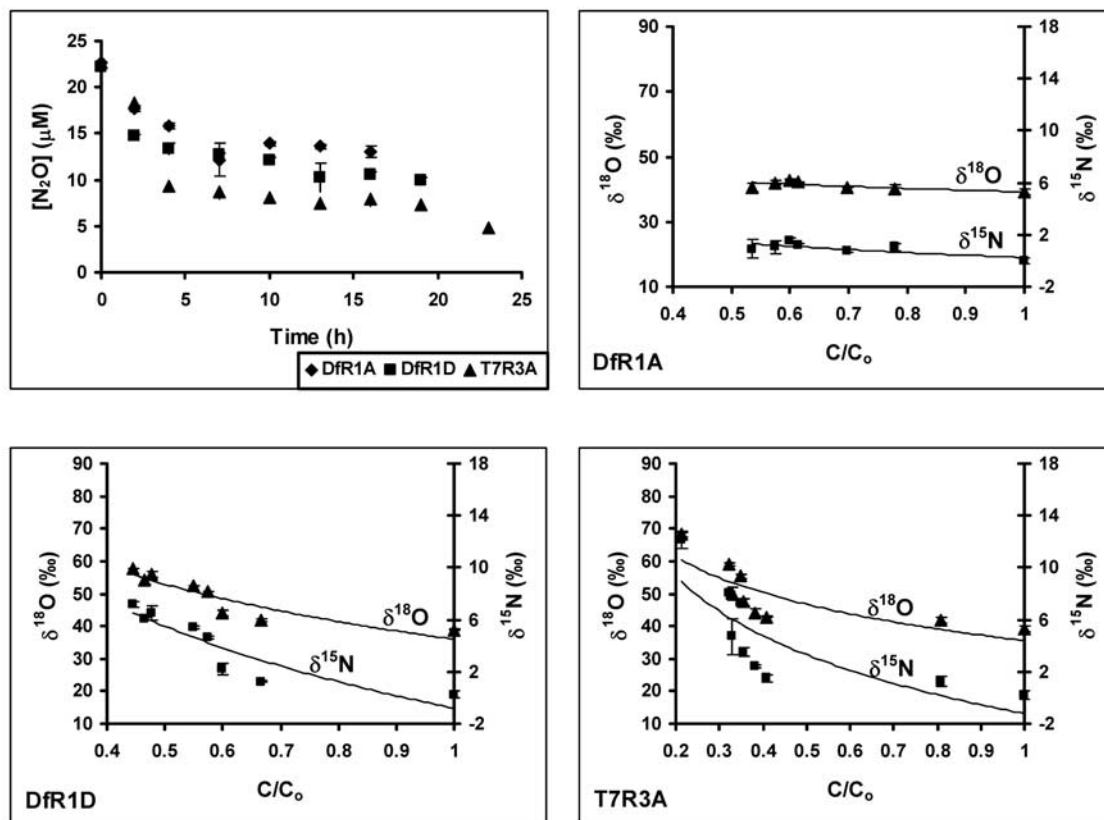


Figure 2. Concentration, $\delta^{15}\text{N}$, and $\delta^{18}\text{O}$ data for N₂O within soil mesocosm N₂O reduction experiments using soils from two replicate plots dominated by deciduous forest (DfR1A and DfR1D) and an early successional treatment (T7R3A) within the KBS LTER main site. Isotope values are expressed in relation to the concentration (C) at any point in time divided by the initial concentration (C₀) following the Rayleigh model (equation (3)). The slope of a natural log fit to the data equates to the fractionation factor for each experiment and are listed in Table 2. Error bars represent 2 standard deviations around the mean of three replicates for each treatment.

the soil mesocosm from the conventional agricultural field, T1R3A, which illustrates heterogeneity in N₂O reduction rates across plots with similar management histories. Rates of N₂O reduction ranged from 64 to 241 $\mu\text{mol}\cdot\text{h}^{-1}\cdot\text{m}^{-1}$ (Table 1), which are between 1 and 5 times the rates of denitrification previously measured by *Cavigelli and Robertson* [2000] in conventional agricultural fields at the KBS LTER. N₂O was the only oxidized N substrate available for denitrification in these mesocosms, so the rates of N₂O reduction observed also represent gross denitrification. Furthermore, the artificial conditions imposed by homogenization of soils, saturation, and maintenance of anaerobic conditions indicate that these rates should be considered potential rather than actual rates of N₂O reduction and, consequently, are likely in excess of what can be expected in the field. Despite their departure from field conditions, our experiments are appropriate for constraining the isotope effects associated with N₂O reduction by soil microbial communities.

[15] In this paper, we follow the convention of *Mariotti et al.* [1981] by defining the magnitude of isotopic fractionation during a specific reaction as the fractionation factor α ,

$$\alpha = k_2/k_1, \quad (1)$$

where k_1 and k_2 are the reaction rates for the light and heavy isotopically substituted compounds, respectively (although some authors use the inverse of this ratio). We further define an isotopic enrichment factor, ϵ , which approximates for N₂O reduction the difference in isotopic composition between N₂ and N₂O,

$$\epsilon = (\alpha - 1) \cdot 1000. \quad (2)$$

Isotopic fractionation during many microbial reactions, including denitrification, has been described using a Rayleigh distillation equation in which the isotopic composition of the residual substrate of a reaction (δ_s) is related to that of the initial substrate (δ_{s0}) an isotopic enrichment factor (ϵ), and the ratio of the observed to initial substrate concentration (C/C_0) [*Mariotti et al.*, 1981],

$$\delta_s = \delta_{s0} + \epsilon \ln(C/C_0). \quad (3)$$

Equation (3) describes a relationship by which the isotopic composition of the substrate, N₂O, is linearly related to the natural log of its concentration. Within each of the mesocosm experiments both $\delta^{15}\text{N}$ and $\delta^{18}\text{O}$ were correlated with the natural log of C/C_0 ($r^2 > 0.65$, $P < 0.05$) (Figures 1

Table 1. Rates of N₂O Reduction (Denitrification) and Isotopic Enrichment Factors for Mesocosm Experiments Using Soils From a Deciduous Forest (DfR1A and DfR1D), Early Successional (T7R3A), and Conventional Agricultural (T1R1A, T1R3A, and T1R5B) Treatment Histories^a

Soil Mesocosm	Rate, $\mu\text{mol m}^{-2} \text{h}^{-1}$	$\epsilon^{15}\text{N}$	$\epsilon^{18}\text{O}$
DfR1A	87.1 ± 22.8	-1.8 ± 0.4	-5.1 ± 1.5
DfR1D	114.3 ± 2.9	-9.2 ± 0.3	-25.1 ± 0.5
T7R3A	137.0 ± 3.5	-6.8 ± 0.8	-15.3 ± 3.4
T1R1A	63.6 ± 9.1	-4.5 ± 0.4	-8.6 ± 0.7
T1R3A	240.8 ± 4.4	-8.3 ± 1.8	-19.8 ± 1.3
T1R5B	91.5 ± 11.9	-4.7 ± 0.6	-11.1 ± 2.4

^aAll treatments were conducted in triplicate and the error shown represents 1 standard deviation.

and 2). Thus even while rates of reduction and isotope values differed markedly between mesocosm experiments, the significant correlations observed indicate that the fundamental constraints of the Rayleigh model, namely a unidirectional reaction and maintenance of a closed system, were followed.

[16] Under ideal conditions, an isotopic enrichment factor is a constant that should not vary with experimental conditions. Isotope effects controlled by biological processes, however, fundamentally violate the assumptions of the Rayleigh model in that transformations rarely consist of a single reaction. Denitrification, for example, consists of a series of steps involving diffusion of nitrate into the cell, reduction of nitrate to nitrite, nitrite to nitric oxide, nitric oxide to N₂O, and N₂O to N₂ [Firestone and Davidson, 1989]. Fractionation factors will vary depending on environmental conditions (e.g., degree of water saturation and availability of electron donors) and with which step in the overall denitrification process is rate limiting. The observed fractionation factor is most often that associated with the rate limiting step but may reflect variation in the relative importance of individual reaction steps [Bryan *et al.*, 1983; Ostrom *et al.*, 2002]. During N₂O reduction, the most important rate limiting steps are diffusion into the cell and enzymatic reduction. While fractionation during diffusion is generally small [e.g., Brandes and Devol, 1997] fractionation during enzymatic reduction is large [Yoshida *et al.*, 1984; Yamazaki *et al.*, 1987]. The low observed fractionation factors for nitrogen in our soil mesocosm experiments (-9.1 to -1.7‰) (Table 1) relative to those reported previously (-39 to -27‰) [Yoshida *et al.*, 1984; Yamazaki *et al.*, 1987] indicate that diffusion is important in limiting the expression of the enzymatic fractionation in our experiments. The variability in observed fractionation factors suggests that the balance between diffusion and enzymatic reduction in controlling fractionation is variable between mesocosm experiments. Such observed variability in the expression of fractionation is a confounding factor that often limits application of natural abundance stable isotope studies in source apportionment.

3.2. N₂O Reduction in Pure Culture Experiments

[17] Within pure cultures of *Pseudomonas stutzeri* and *Pseudomonas denitrificans* correlations between isotope values and the natural log of C/C₀ during N₂O reduction substantiate the use of the Rayleigh model requirements of a closed system (Figure 3). The observed isotopic enrichment

factors for $\delta^{15}\text{N}$ and $\delta^{18}\text{O}$ data (Table 2) were within the range of those obtained in soil mesocosms. Experimental conditions enabled the determination of isotopic enrichment factors for $\delta^{15}\text{N}^{\alpha}$, $\delta^{15}\text{N}^{\beta}$ and SP in addition to those for bulk isotopes. Isotopic enrichment for the β position were quite small (~2‰), whereas those associated with the α position were greater than those associated with the bulk $\delta^{15}\text{N}$. These results indicate that the majority of N isotope fractionation during reduction is associated with the α position in N₂O. This is consistent with cleavage of the N-O covalent bond during denitrification. The fractionation factor for SP during N₂O reduction was small (~6‰) but of sufficient magnitude that if reduction of N₂O is a prevalent process changes in SP can be expected. Based on the Rayleigh model using a fractionation factor for SP of 6‰, for example, 50% consumption of an existing N₂O pool would result in a 4‰ increase in SP. While small, this change is of sufficient magnitude to affect estimates of the relative importance of nitrification and denitrification to N₂O fluxes determined on the basis of SP.

3.3. Relationships Among Isotopic Enrichment Factors for N₂O Reduction

[18] A critical question for apportionment studies is whether changes resulting from N₂O reduction can be recognized and/or potentially corrected for. Our results indicate that reduction of N₂O prior to its evolution to the atmosphere has the potential to result in changes in $\delta^{15}\text{N}$, $\delta^{18}\text{O}$ and SP of sufficient magnitude to compromise calculation of sources based on these values. Correction of isotope values for fractionation during reduction is theoretically possible if the extent of the reaction and isotopic enrichment factors are known [Ostrom *et al.*, 2002]; however, this approach, based on bulk isotope data, is not possible in this case owing to the wide variation of ϵ values we observe for N₂O reduction. Furthermore, reduction results in ¹⁵N enrichment, which serves to decrease the difference in $\delta^{15}\text{N}$ between soil nitrate or ammonium and N₂O, which is the basis of the apportionment approach commonly used in bulk natural abundance isotopologue studies [Perez *et al.*, 2000, 2001; Bol *et al.*, 2003]. Reduction also results in a small but appreciable increase in SP, which shifts values away from the SP value for denitrification (0‰) and toward that of nitrification (33‰) [Sutka *et al.*, 2006]. Consequently, we conclude that the isotope effects resulting from reduction cannot be ignored in isotope based source apportionment studies.

[19] We note, however, that despite a wide range of fractionation factors evident for N₂O reduction within the soil mesocosm and pure culture data there is a remarkable and significant relationship between $\epsilon^{18}\text{O}$ and $\epsilon^{15}\text{N}$ that is characterized by a slope of 2.5 (Figure 4). Similarly, we find a relationship of 2.6 between $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ for all of the data in our reduction experiments (Figure 4) which is expected given that fractionation factors are derived from individual isotope values (equation 3). The slight difference between the slopes of $\delta^{18}\text{O}$ versus $\delta^{15}\text{N}$ and $\delta^{18}\text{O}$ versus $\epsilon^{15}\text{N}$ in Figure 4 is a consequence of unequal numbers of samples in experiments used to calculate fractionation factors and thus we believe the ratios of fractionation factors are a more accurate descriptor of the isotopologue relationships during N₂O reduction. The value of 2.5 we report is in

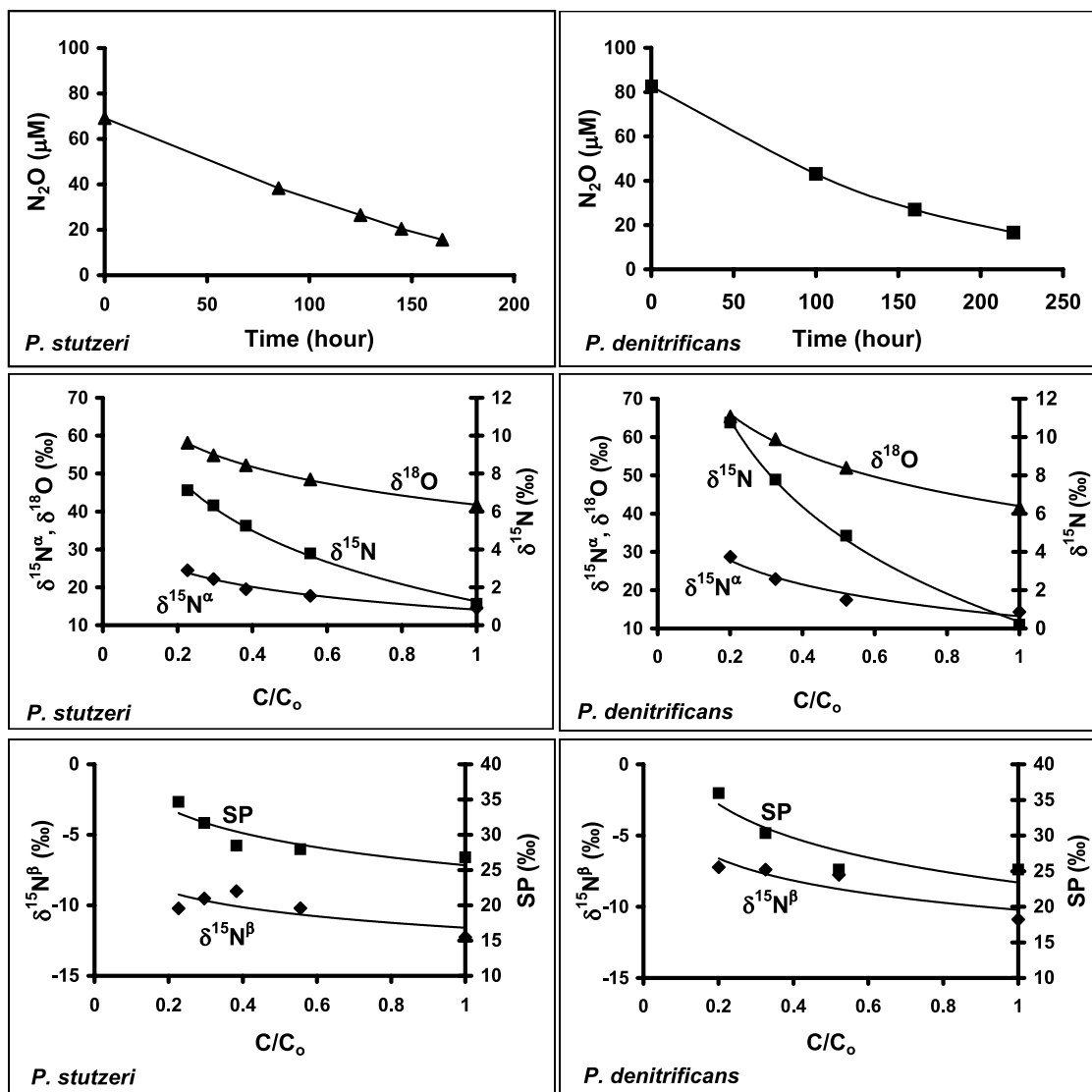


Figure 3. Concentration, $\delta^{15}\text{N}$, $\delta^{18}\text{O}$, $\delta^{15}\text{N}^{\alpha}$, $\delta^{15}\text{N}^{\beta}$, and SP values for N₂O reduction within pure culture experiments of *P. stutzeri* and *P. denitrificans*. Isotope values are expressed in relation to the concentration (C) at any point in time divided by the initial concentration (C₀) following the Rayleigh model (equation (3)). The slope of a natural log fit to the data equates to the fractionation factor for each experiment and isotopomer and are listed in Table 2.

remarkable agreement with the values of 2.5 reported for reduction in Siberian soils [Menyailo and Huntgate, 2006a], 2.0 reported for reduction in soils [Webster and Hopkins, 1996; Mandernack et al., 2000] and pure culture [Webster and Hopkins, 1996], and 2.6 that is obtained by taking the ratio of $\epsilon^{18}\text{O}$ and $\epsilon^{15}\text{N}$ obtained during reduction in anoxic ocean waters [Yamagishi et al., 2007]. Furthermore, ratios of $\epsilon^{18}\text{O}$ to $\epsilon^{15}\text{N}^{\alpha}$ and $\epsilon^{18}\text{O}$ to SP (which are analogous to the slope) within the pure culture experiments, while only based on two cultures, were in remarkable agreement with values of 1.7 and 2.2, respectively (Table 2), that is similar to a value of 1.5 in ocean waters [Yamagishi et al., 2007]. A similar constant relationship between oxygen and nitrogen isotope fractionation in nitrate for denitrification has been reported [Böttcher et al., 1990; Durka et al., 1994]. While fractionation factors for N₂O reduction experiments or in the field are susceptible to considerable variation, the

relationships between $\epsilon^{15}\text{N}$, $\epsilon^{15}\text{N}^{\alpha}$ and $\epsilon^{18}\text{O}$ covary in a predictable and diagnostic manner. Consequently, reduction of N₂O can be clearly recognized when relationships between $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$, $\delta^{18}\text{O}$ and $\delta^{15}\text{N}^{\alpha}$, and $\delta^{18}\text{O}$ and SP approach values of 2.5, 1.7 and 2.2, respectively.

3.4. N₂O Reduction and Isotopologue-Based Source Apportionment

[20] Mass balance mixing models are commonly used in flux chamber studies to determine the isotopologue compo-

Table 2. Isotopic Enrichment Factors for N₂O Reduction (Denitrification) in Pure Culture Experiments

Denitrifier Species	$\epsilon^{15}\text{N}$	$\epsilon^{18}\text{O}$	$\epsilon^{15}\text{N}^{\alpha}$	$\epsilon^{15}\text{N}^{\beta}$	ϵSP
<i>P. stutzeri</i>	-4.1	-10.9	-6.6	-1.6	-5.0
<i>P. denitrificans</i>	-6.6	-15.0	-9.1	-2.2	-6.8

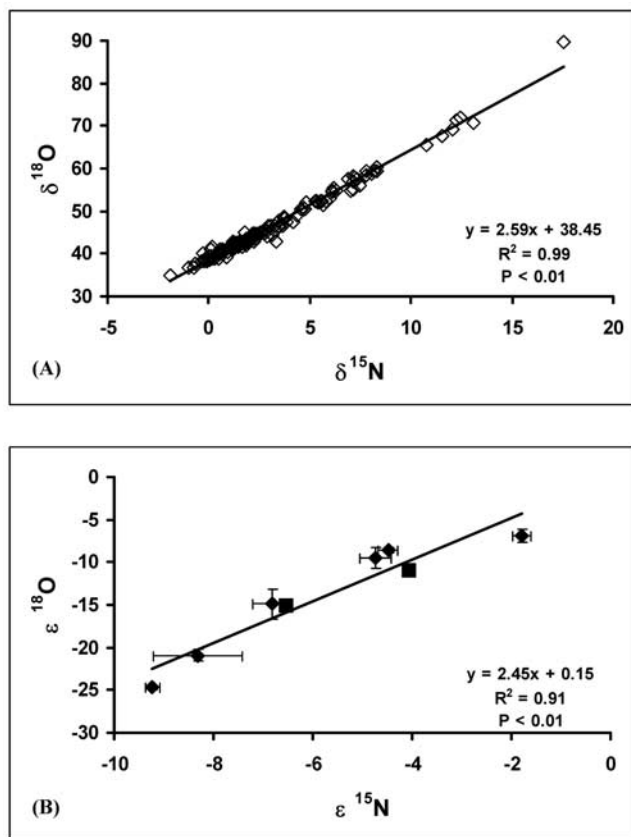


Figure 4. Relationships between (a) $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ values for all soil mesocosm (diamonds) and pure culture (triangles) N_2O reduction experiments and (b) between the fractionation factors of oxygen and nitrogen isotopes during N_2O reduction in the soil (diamonds) and pure culture (squares) experiments (ϵ values from Tables 1 and 2). The error bars in Figure 4b indicate 1 standard deviation based on triplicate experiments for each soil type in the mesocosms experiments. The pure culture experiments were not replicated.

sition of soil-derived N_2O [e.g., *Well et al.*, 2006]. If, however, reduction of N_2O is significant, the isotope effects associated with this process will result in enrichment in ^{15}N and ^{18}O and an inaccurate assessment of the isotopologue values for soil-derived N_2O . Consequently, it is desirable (1) to know when the isotope effects associated with reduction are important and (2) to correct for such shifts if possible. We can establish the affect of N_2O reduction in flux chamber studies on isotope values in a hypothetical manner based on the relationships between $\delta^{18}\text{O}$, $\delta^{15}\text{N}$ and $\delta^{15}\text{N}^\alpha$ established within our mesocosm and pure culture studies in comparison to flux chamber data collected previously at the KBS LTER [*Ostrom et al.*, 2004] (Figure 5). During a typical flux chamber deployment, the concentration of N_2O will rise steadily representing the addition of soil-derived N_2O to that initially present in the atmosphere (320 ppbv). With continued input of soil-derived N_2O the isotopologue composition of N_2O within the chamber will shift away from values characteristic of the atmosphere (labeled “troposphere” in Figure 5) toward values reflecting the isotopologue composition of soil-derived N_2O and data will lie along a line connecting the isotopologue values for

the troposphere and soil-derived N_2O (labeled “mixing line”). If N_2O reduction is the only process affecting the isotopic composition of N_2O then the concentration of N_2O would decline steadily from tropospheric values (320 ppbv)

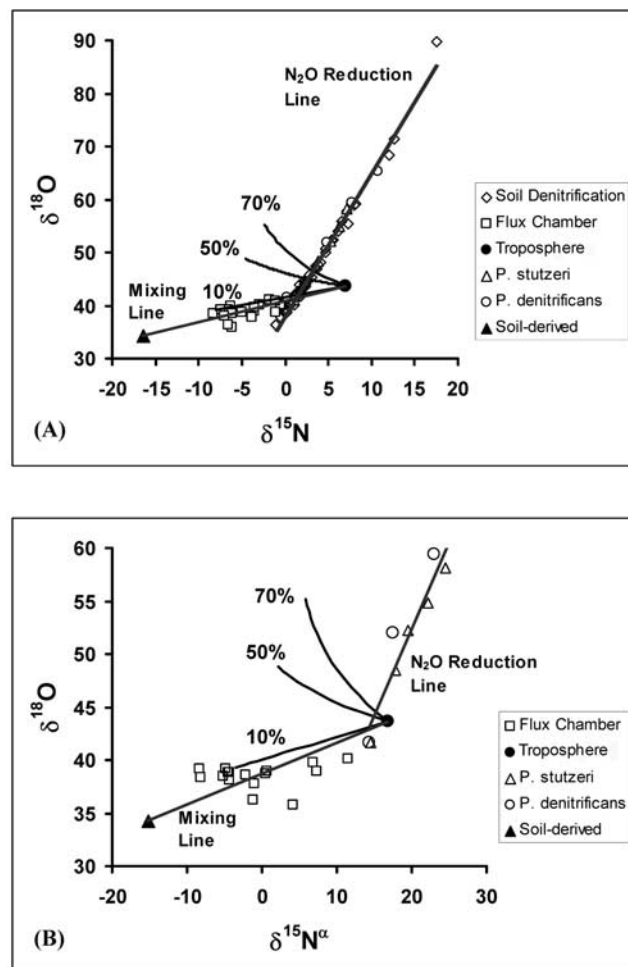


Figure 5. Relationships between the (a) $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ and (b) $\delta^{18}\text{O}$ and $\delta^{15}\text{N}^\alpha$ of N_2O for N_2O production and reduction in soils. Data from cultivated successional soil flux chambers at the KBS LTER [*Ostrom et al.*, 2004] are indicated in open squares and lie along a “mixing line” between the values for tropospheric N_2O (solid circle) [*Yoshida and Toyoda*, 2000] and calculated values for soil-derived N_2O (solid triangle). Isotopomer values from soil mesocosm (open diamond) and pure culture studies of *P. stutzeri* (open triangles) and *P. denitrificans* (open circle) combine to form the “ N_2O reduction line.” The slope of the N_2O reduction lines, 2.5 for $\delta^{18}\text{O}$ – $\delta^{15}\text{N}$ and 1.7 for $\delta^{18}\text{O}$ – $\delta^{15}\text{N}^\alpha$, would result when N_2O reduction occurs in the absence of production. The N_2O reduction line does not intersect the isotopomer values for the troposphere but originates with values of the pure N_2O standard that was used in the headspace of all reduction experiments. Solid lines represent the trends expected within flux chamber studies at the production rates and soil-derived isotopologue values from *Ostrom et al.* 2004 when simultaneous production and consumption occur, at 10, 50, and 70% the rate of production, based on a bulk nitrogen isotopic enrichment factor of 10‰.

and isotopologue values would rise steadily. These data would like on a line with a slope of 2.5 for the plot of $\delta^{18}\text{O}$ versus $\delta^{15}\text{N}$ and 1.7 for the plot of $\delta^{18}\text{O}$ versus $\delta^{15}\text{N}^\alpha$ (identified as “N₂O reduction line” in Figure 5). In that soil-derived N₂O is commonly depleted in ¹⁵N and ¹⁸O relative to tropospheric N₂O [Mandernack *et al.*, 2000; Perez *et al.*, 2000, 2001; Tilsner *et al.*, 2003; Wrage *et al.*, 2004a], samples representing a mixture of soil-derived and atmospheric sources lie on a mixing line with a slope less than 1 (0.41 in the flux chamber study shown in Figure 5). Linear relationships between $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ of less than 1 have been commonly associated with N₂O production within flux chamber studies [Webster and Hopkins, 1996; Mandernack *et al.*, 2000]. Similarly, the low $\delta^{15}\text{N}^\alpha$ values for soil-derived N₂O from the Ostrom *et al.* [2004] study also produce a linear $\delta^{18}\text{O}$ versus $\delta^{15}\text{N}^\alpha$ mixing relationship with atmospheric N₂O with a slope of 0.30 (Figure 5b). These values differ markedly from the slopes obtained in our reduction experiments that produce relationships between $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ and $\delta^{18}\text{O}$ and $\delta^{15}\text{N}^\alpha$ of 2.5 and 1.7, respectively (shown as “N₂O reduction lines” in Figure 5). Consequently, a qualitative indicator of N₂O reduction are slopes for the relationships between $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ and $\delta^{18}\text{O}$ and $\delta^{15}\text{N}^\alpha$ that approach values of 1 or greater.

[21] We can predict, in a theoretical manner, how the relationships between $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ and $\delta^{18}\text{O}$ and $\delta^{15}\text{N}^\alpha$ will respond if reduction of N₂O occurs simultaneously with production. In the absence of production, isotope data in flux chambers will follow the “mixing line” shown in Figure 5 owing to the addition of soil-derived N₂O to tropospheric N₂O. The precise changes in $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ are determined from a basic isotope mixing model [e.g., Well *et al.*, 2006],

$$\delta_{\text{meas}} = (\delta_{\text{ATM}}Q_{\text{ATM}} + \delta_{\text{SD}}Q_{\text{SD}})/Q_{\text{meas}}, \quad (4)$$

where, meas, ATM and SD refer to the isotope values (δ) of and quantity (Q) of N₂O measured, that is present initially in the atmospheric (ATM), and that added from soil-derived (SD) production, respectively. We use actual flux rate and soil-derived isotope values that were obtained from a series of flux chamber deployments at the KBS LTER [Ostrom *et al.*, 2004] with the assumption that this data is unaffected by N₂O reduction. On the basis of the average rate of production and isotopologue composition of soil-derived N₂O, we predict, from equation (4), how the isotopic composition of N₂O measured (δ_{meas}) changes as a function of time over a typical 1 hour closing of a flux chamber. The solution to δ_{meas} for each point in time becomes the initial isotopic composition of N₂O (δ_{so}) in equation (3) that evaluates the isotopologue shifts resulting from N₂O reduction. For the C/C_o term in equation (3) we use 0.1, 0.3 or 0.7 to represent a rate of reduction equivalent to 10, 30 or 70% of the rate of production. We use a bulk nitrogen fractionation factor ($\epsilon^{15}\text{N}$) for N₂O reduction of -10‰ to predict the resulting $\delta^{15}\text{N}$ values of the chamber N₂O. Fractionation factors for $\epsilon^{18}\text{O}$ and $\epsilon^{15}\text{N}^\alpha$ are based on the constant relationships between $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ and $\delta^{18}\text{O}$ and $\delta^{15}\text{N}^\alpha$ of 2.5 and 1.7, respectively, which were found in the soil and pure culture N₂O reduction experiments. A $\epsilon^{15}\text{N}$ of -10‰ was chosen as a reasonable worst-case-scenario; being slightly greater than the largest degree of fractionation

observed in the soil N₂O reduction experiments (Table 1). The net result of simultaneous production and consumption of N₂O is a deviation from a linear mixing relationship between soil-derived and atmospheric N₂O to progressively more curvilinear relationships between $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ and $\delta^{15}\text{N}^\alpha$ with increasing importance of reduction. Consequently, an important line of evidence for an alteration of isotope values due to reduction is the deterioration of a linear relationship between $\delta^{18}\text{O}$ and $\delta^{15}\text{N}$ and $\delta^{18}\text{O}$ and $\delta^{15}\text{N}^\alpha$. The magnitude of deviation from linearity may thus serve as a tool for potentially correcting isotopologue data for isotope shifts due to reduction.

[22] A small degree of reduction of 10% results in only a slight deviation from the mixing lines while maintaining a linear relationship (Figure 5). Over the course of 1 hour at the production rates observed the net affect of 10% reduction and a $\epsilon^{15}\text{N}$ of -10‰ is a shift in $\delta^{15}\text{N}$ of only 0.7‰. Given the proportional relationships between isotopologue enrichment factors we have established for reduction, this small shift in $\delta^{15}\text{N}$ corresponds to increases in $\delta^{18}\text{O}$ and SP of 1.9 and 0.9‰, respectively. Sutka *et al.* [2006] indicate that SP values of 0 and 33‰ can be used as conservative tracers of N₂O production from denitrification and nitrification, respectively. On this basis, the small shift in SP that would result from a reduction rate equivalent to 10% of the production rate would result in an underestimate of the proportion of N₂O from denitrification of only 2.7%. In this manner, the proportional relationships between isotopologues provides a basis to correct for shifts resulting from N₂O reduction, however, a precise correction requires knowledge of the soil-derived isotopologue values a priori which is unlikely. Nonetheless, our approach defines a means for establishing if the affect of N₂O reduction on isotopologue values in soil flux chamber studies is appreciable and provides a semi-qualitative approach for understanding the magnitude of such shifts.

[23] While our experiments are quite useful in quantifying isotope effects in response to N₂O reduction in natural soil microbial communities it is important to understand that our experimental design specifically addresses reduction of N₂O that enters microbial cells from the surrounding soil or headspace. Our design does not address N₂O reduction that occurs within the cells of nitrifying or denitrifying microbes following production of N₂O but prior to its escape from the cell. In other words, N₂O that is produced by soil bacteria can undergo 3 possible routes: (1) escape from the cell to the atmosphere, (2) reduction within the cell prior to its escape, or (3) reduction upon reentering the cell prior to escape to the atmosphere. Our study specifically simulates step 3 but does not address reduction within the cell prior to its escape (step 2). This distinction is subtle but important as studies commonly report the ratio of N₂O production to the sum of N₂O and N₂ production (N₂O/(N₂O + N₂)) [Dendooven and Anderson, 1995; Dendooven *et al.*, 1996; Bergsma *et al.*, 2002; Well *et al.*, 2006] which measures total N₂O reduction and, therefore, differs from our experimental design by including reduction via step 2 as well as step 3. Ratios of N₂O/(N₂O + N₂) across ecosystems have been shown to be quite variable [Dendooven and Anderson, 1995; Dendooven *et al.*, 1996; Well *et al.*, 2006] and our previous studies using soils from the KBS LTER found values between 0.3 to 0.4 [Bergsma *et al.*, 2002]. Our study does not

specifically address the isotopologue effects associated with step 2 but rather targets the isotope effects during step 3. Given that steps 2 and 3 are both measures of N₂O reduction within microbial cells we expect that the isotopologue effects associated with steps 2 and 3 will be quite similar and that our results apply to N₂O reduction whether or not the N₂O reduced originates from within or external to the cell.

[24] While it is clear from measurements of the N₂O/(N₂O + N₂) ratio that substantial rates of N₂O reduction relative to production can be expected in soils, we believe most isotopologue studies of N₂O soil flux are often biased by methodological limitations to periods of high flux when rates of N₂O reduction are minor and not of sufficient magnitude to alter isotopologue data. The low abundance of N₂O in the atmosphere creates an analytical challenge to mass spectrometrists such that isotopologue studies of soil flux occur most often when fluxes are high [Tilsner *et al.*, 2003]. Furthermore, error propagation in the calculation of soil-derived isotopologue values from mixing models (equation (3)) increases when the proportion of N₂O from soils versus the atmosphere is low. Reduction of N₂O clearly results in enrichment in ¹⁵N and ¹⁸O, however, our theoretical exercise indicates that minor rates of reduction relative to production have only a small affect on isotopologue data; most notably on SP. Thus we believe the tendency for soil-derived N₂O to be characterized by depletions in ¹⁵N and ¹⁸O relative to atmospheric N₂O [Mandernack *et al.*, 2000; Perez *et al.*, 2000, 2001; Yamulki *et al.*, 2001; Bol *et al.*, 2003; Tilsner *et al.*, 2003; Wrage *et al.*, 2004a] and slopes for the relationships between δ¹⁸O and δ¹⁵N and δ¹⁸O and δ¹⁵N^α of less than one are key lines of evidence for the lack of substantial isotopologue shifts resulting from reduction. Such depletions contrast markedly with studies in anoxic marine waters that show substantial enrichments in ¹⁵N and ¹⁸O that are clearly the result of N₂O reduction (e.g., δ¹⁸O > 100‰ [Yoshinari *et al.*, 1997; Naqvi *et al.*, 1998]). Furthermore, our theoretical exercise demonstrates that if reduction occurs at rates of production of 30% or more there is a substantial deviation from the linear relationships between δ¹⁸O and δ¹⁵N and δ¹⁸O and δ¹⁵N^α. Thus the linear relationships between δ¹⁸O and δ¹⁵N [Webster and Hopkins, 1996; Mandernack *et al.*, 2000; Menyailo and Huntgate, 2006a] and δ¹⁸O and δ¹⁵N^α [Sutka *et al.*, 2006] is strong evidence that reduction was a minor process in these studies. This conclusion is in contrast to studies of the N₂O/(N₂O + N₂) ratio that indicate rates of reduction in excess of 60% that were reported previously at our study site [Bergsma *et al.*, 2002], however, N₂O reduction may be enhanced in incubation studies (by the addition of glucose, for example) and there is a paucity of data on variation in the N₂O/(N₂O + N₂) ratio across broad temporal and spatial scales. Consequently, our expectation that N₂O reduction is a minor process during periods of high soil N₂O flux needs to be confirmed by rate measurements of both N₂O production and consumption across seasonal and spatial scales and, in particular, by isotopologue studies during periods of low N₂O flux.

[25] Resolving production rates and sources of N₂O production continues to be an ongoing effort of many researchers and we have established that SP is an important means for understanding microbial N₂O production and

consumption. While a natural abundance level approach, such as the use of SP, is advantageous to tracer level studies in that the artificial conditions imposed by incubation are not needed, the microbial N cycle adds considerable complexity. For example, dissimilatory nitrate reduction to ammonium and fungal denitrification have been implicated as sources of N₂O [Smith, 1982; Robertson and Tiedje, 1987; Laughlin and Stevens, 2002]. While we expect that the conditions that favor these processes (such as high soil organic content and lack of tilling) are not likely operative in our study clearly more research is warranted. Preliminary research in our laboratory indicates that the SP associated with N₂O production during fungal denitrification is uniquely enriched in ¹⁵N in the α position (>35‰) [Sutka *et al.*, 2004b]. Nonetheless, the low SP value for N₂O production from denitrification of approximately 0‰ is unique relative to all other production pathways measured thus far (nitrification via hydroxylamine production and fungal denitrification) [Sutka *et al.*, 2006]. Furthermore, reduction of N₂O results in an increase in the ¹⁵N content of the α position and SP. Consequently, low values for SP approaching 0‰ can only be the result of production from denitrification during times when an influence from N₂O reduction is minor. Consequently, calculations of the proportion of N₂O from denitrification based on SP values remain conservative as all other production pathways and N₂O reduction cause increases in SP.

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