



Nitrogen-Induced Changes in Soil Environmental Factors Are More Important Than Nitrification and Denitrification Gene Abundance in Regulating N₂O Emissions in Subtropical Forest Soils

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Subtropical regions are currently experiencing a dramatic increase in nitrogen (N) deposition; however, the contributions of nitrification and denitrification processes to soil N₂O emissions and the underlying mechanisms under increasing N deposition remain unclear. Therefore, a ¹⁵N-tracing laboratory experiment with four N application rates (0, 12.5, 25, and 50 μg ¹⁵N g⁻¹ soil) was conducted to investigate the response of nitrification- and denitrification-derived N₂O to N additions in an evergreen broad-leaved forest (BF) and a *Pinus* forest (PF) in the Wuyi Mountains in southeastern China. Moreover, the abundance of functional genes related to nitrification (*amoA*), denitrification (*nirK*, *nirS*, and *nosZ*), and soil properties were measured to clarify the underlying mechanisms. Results showed that nitrification-derived N₂O emissions were generally decreased with increasing N input. However, denitrification-derived N₂O emissions were a non-linear response to N additions, with maximum N₂O emissions at the middle N application rate. Denitrification-derived N₂O was the dominant pathway of N₂O production, accounting for 64 to 100% of the total N₂O fluxes. Soil NH₄⁺-N content and pH were the predominant factors in regulating nitrification-derived N₂O emissions in BF and PF, respectively. Soil pH and the *nirS* abundance contributed the most to the variations of denitrification-derived N₂O emissions in BF and PF, respectively. Our results suggest that N application has the potential to increase the contribution of denitrification to N₂O production in subtropical forest soils. Changes in soil chemical properties induced by N addition are more important than the abundance of nitrification and denitrification functional genes in regulating soil nitrification- and denitrification-derived N₂O emissions.

Keywords: nitrification, denitrification, microbial functional genes, N₂O emissions, N application

INTRODUCTION

Nitrogen (N) deposition is continuously increasing primarily due to increased anthropogenic fossil fuel combustion, industrialization, and N fertilizer application. Elevated N deposition is expected to not only alter N cycling in forest ecosystems but also enhance N gas loss from the soils (Lu et al., 2011; Fang et al., 2015; Niu et al., 2016; Tang et al., 2018). As one of the primary forms of N gas loss from the soils, nitrous oxide (N₂O) is more than 298 times as effective at trapping atmospheric heat as CO₂ and is also one of the largest stratospheric ozone-depleting substances (IPCC, 2013). Terrestrial ecosystems contribute ~60% of the global N₂O emissions (IPCC, 2013), of which about 15 to 55% are from tropical and subtropical forests that are believed to be the largest N₂O terrestrial source (Solomon, 2007; Zhang et al., 2019). Moreover, N deposition in tropical and subtropical areas has been steadily increasing in recent decades (ranging from 30 to 73 kg N ha⁻¹ year⁻¹), and this trend is projected to continue over the coming decades (Zhang et al., 2008; Liu et al., 2013; Zhu et al., 2015).

N₂O is produced primarily during the soil microbial processes of nitrification and denitrification, the rates of which are influenced by the inputs of N. Recent studies in subtropical forests demonstrated that the contribution of nitrification-derived N₂O to the total N₂O emissions was ≤50% and that of denitrification were 53 to 100% under aerobic conditions (Zhang et al., 2011; Zhu et al., 2013a; Han et al., 2018; Tang et al., 2018). In addition, the rates of denitrification have generally increased following N application (Zhu et al., 2013b; Morse et al., 2015; Niu et al., 2016; Han et al., 2018). However, few studies have focused on the response of nitrification-derived N₂O emissions to N application, especially in subtropical forests (Han et al., 2018). Therefore, whether the response of nitrification-derived N₂O emissions to N application is more pronounced than that of denitrification-derived N₂O emissions or whether the contribution of denitrification to N₂O production under elevated N input is still dominant over that of nitrification remains unclear. Thus, quantifying the underlying nitrification and denitrification contributions to the N₂O production will help us understand the changes in the N transformation process with increasing N input in forest ecosystems.

Soil nitrification and denitrification can be influenced by different soil microbial groups, soil properties, and climate conditions (Levy-Booth et al., 2014; Cheng et al., 2019). Ammonia oxidation is not only the first but also the rate-limiting step of nitrification, which is performed by both ammonia-oxidizing archaea (AOA) and bacteria (AOB) (Chen and Peng, 2020). On the other hand, denitrification is a four-step microbial facilitated reduction process, whereby NO₃⁻ is reduced to NO₂⁻, NO, N₂O, and eventually to N₂. The first step for the gas production during denitrification is the reduction of NO₂⁻ to NO, which is catalyzed by nitrite reductases encoded by *nirK* and *nirS* genes (Levy-Booth et al., 2014; Hu et al., 2015). N₂O can be further reduced to N₂ by nitrous oxide reductases, which are encoded by *nosZ* genes (Levy-Booth et al., 2014). While the importance of nitrification and denitrification microorganisms in regulating soil N₂O emission has been widely accepted (Chen and Peng, 2020), to what extent the abundance of nitrifier and

denitrifier functional gene can be a good predictor of N₂O emissions under elevated N inputs remain unclear. Moreover, several recent studies have reported that soil environmental factors (e.g., NH₄⁺, NO₃⁻, pH, TN/TP, and TC/TN) can explain better the changes in soil nitrification and denitrification activities than soil nitrification and denitrification functional genes in fertilized forest soils (Pett-Ridge et al., 2013; Shcherbak et al., 2014; Tang et al., 2018, 2019). Although tropical and subtropical forest soils are the largest N₂O terrestrial sources (Solomon, 2007; Soper et al., 2018; Zhang et al., 2019), only a few studies so far in these regions have attempted to directly link the nitrification- and denitrification-derived N₂O emissions in fertilized soils to N functional genes (Tang et al., 2018, 2019), and more importantly, to investigate the relative contribution of soil biotic and abiotic factors on the nitrification- and denitrification-derived N₂O emissions. Greater insight into the regulation of these processes can help us understand the underlying mechanisms and adopt management practices to lower N₂O emissions.

To evaluate the changes in soil N₂O emissions resulting from nitrification and denitrification processes under N application in subtropical forests and clarify the underlying mechanisms, we carried out a ¹⁵N-NO₃⁻ and ¹⁵N-NH₄⁺ labeling experiment under laboratory conditions. Mineral soils were collected from two subtropical forests, an evergreen broad-leaved forest (BF), and a *Pinus* forest (PF), which are located in the Wuyi Mountains in southeastern China. The evergreen BF was chosen because it is a product of the monsoon climate and it represents the climax vegetation type of subtropical regions (Zhou et al., 2013). *Pinus* forest was selected because of its strong contribution to afforestation purposes since the 1980s. It is worth highlighting that the selected pine species have a strong ability to adapt to poor soil conditions in the hilly red soil region of south China, which faced moderate to intense soil and water loss challenges in the 1980s. The forests have been widely planted since that time and account for about 30.5% of the subtropical forests in south China (Cao et al., 2015). Many previous studies have demonstrated that broadleaf/deciduous species produce more N₂O than conifers due to a large difference in the production and reduction rates of N₂O under the broadleaf/deciduous species (Zhang et al., 2008; Qin et al., 2019). However, some studies reported that coniferous stands emit more N₂O than the adjacent deciduous stands that were affected by N deposition. Therefore, the role of different forest types (or tree species) in regulating N₂O emissions remains unclear.

This study aimed to (1) investigate the responses of nitrification- and denitrification-derived N₂O emissions to elevated N inputs, (2) clarify the dominant factors that control nitrification- and denitrification-derived N₂O emissions, and (3) compare the differences in nitrification- and denitrification-derived N₂O emissions between the BF and PF. We hypothesized that (1) nitrification- and denitrification-derived N₂O emissions would be enhanced with the increased N application and that denitrification may contribute more to the increase in N₂O emissions than nitrification, (2) the abundances of functional genes could be more important than soil environmental factors in regulating nitrification- and denitrification-derived N₂O emissions, and (3) the nitrification- and denitrification-derived N₂O emissions would be greater in the BF than the PF.

MATERIALS AND METHODS

Study Site Descriptions and Soil Sampling Procedures

The experimental soil samples were collected at the Wuyishan National Nature Reserve (WNNR), located in Fujian Province (27°33′–27°54′N, 117°27′–117°51′E) in southern China. The climate of the WNNR (area: 56,527 ha) is humid monsoon, which is mainly characterized by the mean annual precipitation and temperature of 2,000 mm and 15°C, respectively, as well as 83.5% relative humidity and 100 fog days year⁻¹ (Xu et al., 2010; Wang et al., 2013). The WNNR has two typical vegetation types, that is, evergreen BF and pine forest (PF). The BF and PF are located at about 550 m a.s.l. and 1,100 m a.s.l., respectively. While *Castanopsis carlesii* Hay and *Castanopsis eyrie* Tutch are the dominant species in the BF, *Pinus taiwanensis* Hayata is the dominant species in the PF (Xu et al., 2010; Lyu et al., 2019). Background atmospheric N deposition in the WNNR is about 8 kg N ha⁻¹ year⁻¹ (Xu et al., 2016; Gurmessa et al., 2022).

In April 2018, three 20 m × 20 m plots without N addition (plots were about 10 m away from each other) were randomly established in each vegetation type, and soil samples were collected with a soil core sampler (8 cm in diameter) from the surface soil layer (0–20 cm) after litter removal. Ten soil cores were randomly collected from each plot and there were a total of 30 soil cores collected in each forest stand. Soil samples from the same forest type were thoroughly mixed to form a composite sample. All soil samples were sieved through a 2-mm mesh sieve and were then divided into two sub-samples. The first sub-sample was maintained at 4°C for 1 week before being used for the incubation experiment. The second sub-sample was air-dried for the measurements of soil chemical properties. The soil at both sites is classified as yellow-red soil based on the Chinese Soil Classification System, equivalent to an Ultisol according to the United States Department of Agriculture Soil Taxonomy classification system (Lyu et al., 2019). Soil chemical properties were shown in Table 1. The concentrations of soil SOC, TN, NH₄⁺-N, and NO₃⁻-N in the PF were significantly higher than those in the BF, whereas soil pH and δ¹⁵N were lower in the PF and no significant difference was found in the soil C/N ratio in these two forest soils (Table 1).

¹⁵N-Tracing Experiment and Gas Sampling Procedures

This study used a completely randomized factorial design experiment that consisted of four N application rate treatments (0, 12.5, 25, and 50 μg ¹⁵N g⁻¹ soil equivalent to 0, 16, 32, and 64 kg N ha⁻¹ year⁻¹). We applied N as ¹⁵NH₄NO₃ (99.10 atom

%) and NH₄¹⁵NO₃ (99.21 atom %). The rates of N application were in the range of those of N deposition (10 to 73 kg N ha⁻¹ year⁻¹) in subtropical forests (Zhang et al., 2011, 2019). For each forest soil type, 120 glass jars (500 mL) were prepared and divided into two groups: one group comprised of 60 glass jars containing ¹⁵NH₄NO₃ (4 N addition rates × 3 replications per N application rate × 5 destructive soil sampling) and the other group consisted of 60 glass jars containing NH₄¹⁵NO₃. A total of 30 g (oven-dry basis) of fresh soil was weighed into each glass jar and pre-incubated for 7 days at 25°C to allow the recovery of soil microbial activities. Then, ¹⁵NH₄NO₃ or NH₄¹⁵NO₃ was dissolved in 4 mL deionized water and added to the soil at a rate of 0, 12.5, 25, and 50 μg ¹⁵NH₄⁺-N g⁻¹ soil or ¹⁵NO₃⁻-N g⁻¹ soil. Soil without N addition was regarded as a control and was added with the same volume of deionized water to ensure its similarity in soil moisture content with the N-added soil. The soils were then incubated for 15 days at 25°C and their moisture content was maintained at 60% water-holding capacity throughout the entire experiment. Soils were corrected for water loss by weighing each glass jar plus soil every week and deionized water was added as required to maintain constant soil moisture. During gas sampling, three glass jars at each level of N treatment were randomly selected. The glass jars were closed with the gas-tight rubber lids, and gas sampling tubes were fixed into the middle of the rubber lids to enable the collection of the gas samples. Gas samples (20 mL) were collected from each glass jar with a 3-way stopcocks syringe on days 1, 2, 4, 9, and 15 after N application. Five millimeters were taken from each gas sample to determine the N₂O concentration (Shimadzu, GC-2014C, Japan) and the remaining 15 mL were used to determine the N₂O isotopic composition (IRMS, Isoprime 100, United Kingdom). Before each gas sampling, the glass jars were opened and flushed with ambient air for approximately 30 min and then resealed with stoppers for 30 min.

Quantification of the Contributions of Nitrification and Denitrification to N₂O Emissions

The fractions of N₂O derived from denitrification (FD) and nitrification (FN) processes were calculated using the following equations from Stevens et al. (1997):

$$FD = (a_m - a_n)/(a_d - a_n) \quad (1)$$

$$FN = 1 - FD \quad (2)$$

where a_m is the ¹⁵N atom % of N₂O, a_n is the ¹⁵N atom % of N₂O from the nitrification process (assumed to be equivalent to ¹⁵N

TABLE 1 | Soil chemical properties in the BF and PF.

Forest type	SOC (%)	TN (%)	C/N	δ ¹⁵ N (‰)	pH	NH ₄ ⁺ -N (mg kg ⁻¹)	NO ₃ ⁻ -N (mg kg ⁻¹)
BF	4.10 ± 0.07b	0.21 ± 0.01b	19.80 ± 0.47a	10.84 ± 1.24a	4.80 ± 0.03a	91.89 ± 1.13b	9.40 ± 0.06b
PF	5.67 ± 0.20a	0.27 ± 0.01a	20.63 ± 0.12a	9.97 ± 1.56b	4.72 ± 0.01b	126.66 ± 3.07a	15.37 ± 0.56a

BF and PF indicate the broad-leaved forest and Pinus forest, respectively. Values are means ± standard error, n = 3. Different lowercase letters in the same column represent the difference was significant at P < 0.05.

atom % of the soil NH₄⁺, and a_d is the ¹⁵N atom % of N₂O from denitrification (assumed to be equivalent to ¹⁵N atom % of the soil NO₃⁻).

The atom percent of ¹⁵N is defined as ¹⁵N atom%

$$= \frac{^{15}\text{N}}{(^{15}\text{N} + ^{14}\text{N})} \times 100\% \quad (3)$$

$$\text{Denitrification-derived N}_2\text{O flux} = FD \times \text{N}_2\text{O flux} \quad (4)$$

$$\text{Nitrification-derived N}_2\text{O flux} = FN \times \text{N}_2\text{O flux} \quad (5)$$

Soil Inorganic N, Net Nitrification Rate, Dissolved Organic Carbon, Dissolved Organic Nitrogen, and pH Measurements

Soil samples were also collected after gas sampling. Five times destructive soil sampling was used for the analysis of the soil NH₄⁺ and NO₃⁻ contents, as well as the δ¹⁵N values of soil NH₄⁺ and NO₃⁻. Dissolved organic carbon (DOC) and nitrogen (DON) contents and soil pH were also determined at the end of the experiment. Soil NH₄⁺ and NO₃⁻ contents were extracted using 2 mol L⁻¹ KCl for 1 h and then the extracted solutions were divided into two parts. One part was reacted with sodium salicylate (for NH₄⁺ solution) and hydrazine sulfate solution (for NO₃⁻ solution) and measured, respectively, at 660 and 550 nm using a discrete wet chemistry analyzer (SmartChem 200, AMS Alliance, Italy) to determine the concentrations of soil NH₄⁺-N and NO₃⁻-N. The other part was used for ¹⁵N measurements by distillation using MgO and Devarda's alloy (Zhang et al., 2011). The DOC in the soil was extracted with 2 mol L⁻¹ KCl at 1:5 w/v and measured using a TOC-TN analyzer (Shimadzu TOC, Japan). The soil DON concentration was determined by calculating the difference between soil total dissolved nitrogen (2 mol L⁻¹ KCl extracted at 1:5 w/v) and soil mineral nitrogen (Jones and Willett, 2006). Soil pH was measured at a 1:2.5 soil to solution ratio using a pH detector (PHS-3C, Sheng Ci, Shanghai, China).

Quantification of the *amoA*, *nirK*, *nirS*, and *nosZ* Gene Abundances

At the end of the incubation experiment, soil samples for molecular analyses were immediately stored at -80°C before the aforementioned analysis. DNA was extracted from 0.3 g of frozen soils using a Mo Bio PowerSoil DNA Isolation Kit (Mo Bio Laboratories, Carlsbad, CA, United States) according to the manufacturer's protocol. The concentration and purity of the extracted DNA were determined with a spectrophotometer at 260 nm (NanoDrop Technologies, United States). All extracted soil DNA samples were stored at -80°C before analysis. Quantitative polymerase chain reaction (qPCR) was used to estimate the abundance of nitrification and denitrification functional genes (*amoA*, *nirK*, *nirS*, and *nosZ*) using a real-time PCR detection system. The PCR primers used in this study are listed in **Supplementary Table 1**. The 10-fold serially diluted plasmids carrying each target gene were subjected to real-time PCR assays in triplicate to generate a standard curve. The qPCR efficiencies for AOA-*amoA*, AOB-*amoA*, *nirK*, *nirS*, and *nosZ* were 0.85, 0.94, 0.96, 0.91, and 0.87, respectively. The

corresponding determination coefficients of the standard curve for AOA-*amoA*, AOB-*amoA*, *nirK*, *nirS*, and *nosZ* were 0.998, 0.993, 0.990, 0.999, and 0.996, respectively.

Statistical Analyses

The gene copy numbers were log-transformed to meet the homogeneity of variance requirements. Three-way ANOVA was used to examine the effects of forest type, N application rate, sampling date, and their interactions on soil N₂O flux, nitrification-, and denitrification-derived N₂O emissions, as well as on soil NH₄⁺-N, NO₃⁻-N and their ratio (NH₄⁺-N/NO₃⁻-N). Two-way ANOVA with Duncan's test was performed to determine the effects of forest type, N application rate, and their interactions on the soil properties (soil pH, DOC, and DON) and the abundances of soil nitrification and denitrification functional genes (*amoA*, *nirK*, *nirS*, and *nosZ*). All of the above statistical analyses were performed using SPSS 16.0 (SPSS Inc., Champaign, IL, United States).

To estimate the direct and indirect effects of soil biotic and abiotic factors on soil nitrification- and denitrification-derived N₂O emissions in the BF and MF, structural equation models (SEM) were performed based on our knowledge of soil biotic and abiotic properties in relation to soil nitrification- and denitrification-derived N₂O. Before the SEM analyses, the units of the predictor and dependent parameters were adjusted to obtain comparable parameter variances according to the standardized method. The quality of the SEM model was assessed by using a low χ² value and $P > 0.05$, the root-mean-square error of approximation (RMSEA, the RMSEA ≤ 0.08 indicates a model fit), and the model fit index (GFI > 0.90). All SEM analyses were using Amos 17.0 (IBM, SPSS, NY, United States).

RESULTS

Effects of N Addition on Total and Nitrification- and Denitrification-Derived N₂O Emissions

Soil N₂O fluxes decreased with increasing incubation period (**Figure 1**) and were significantly affected by forest type, N application rate, sampling date, and their interactions (**Table 2**). The N₂O flux values in the BF ranged from 0.05 to 3.85 μg N kg⁻¹ day⁻¹ throughout the entire incubation period (**Figure 1**), with the average fluxes across all the treatments being about 2.4 times greater than those in the PF (**Figure 1**). Non-linear responses of the soil N₂O fluxes to N addition were observed in both the BF and the PF. Specifically, the highest amount of soil N₂O emissions was noted at low and moderate N applications in the BF and PF, respectively (**Figure 2A**).

The application rates of N fertilizer altered not only the total amount of soil N₂O fluxes but also the nitrification- and denitrification-derived N₂O fluxes (**Figures 1, 2**). The nitrification- and denitrification-derived N₂O emissions were significantly higher in the BF than in the PF (**Figure 2B**). The nitrification-derived N₂O fluxes were in the range of 0.03 to 2.66 μg N kg⁻¹ day⁻¹ and 0 to 0.45 μg N kg⁻¹ day⁻¹ in the BF

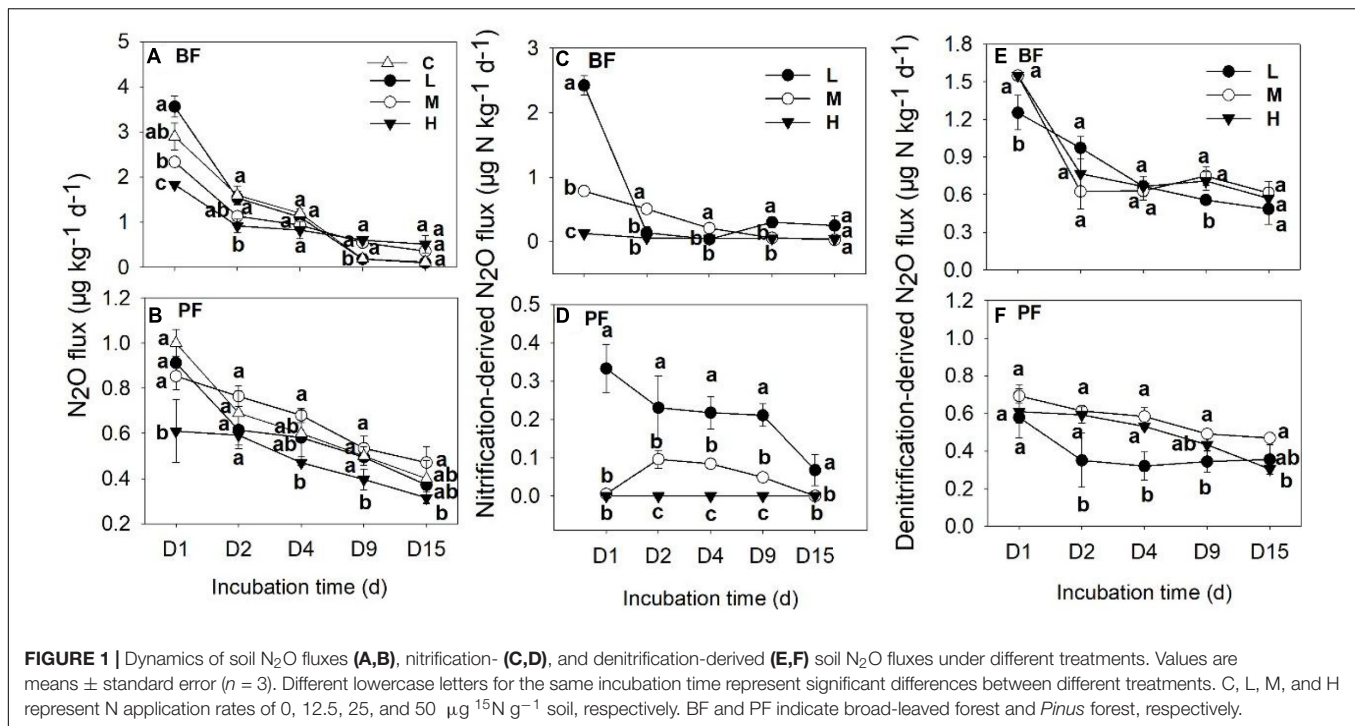


FIGURE 1 | Dynamics of soil N₂O fluxes (A,B), nitrification- (C,D), and denitrification-derived (E,F) soil N₂O fluxes under different treatments. Values are means ± standard error ($n = 3$). Different lowercase letters for the same incubation time represent significant differences between different treatments. C, L, M, and H represent N application rates of 0, 12.5, 25, and 50 $\mu\text{g }^{15}\text{N g}^{-1}$ soil, respectively. BF and PF indicate broad-leaved forest and *Pinus* forest, respectively.

TABLE 2 | *P*-values from Repeated-measure ANOVA of the effects of forest type (FT), N application rates (N), sampling date (D), and their interaction on the N₂O fluxes, nitrification-, and denitrification-derived N₂O emissions, soil NO₃⁻-N, and NH₄⁺-N, $n = 3$.

	FT	N	D	FT *N	FT *D	N *D	FT *N*D
N ₂ O fluxes	<0.001	<0.01	<0.001	0.157	<0.001	<0.001	<0.001
Nitrification-derived N ₂ O	<0.001	<0.001	<0.001	<0.001	<0.001	<0.001	<0.001
Denitrification-derived N ₂ O	<0.001	<0.05	<0.001	0.123	<0.001	0.274	<0.05
NO ₃ ⁻ -N	<0.001	<0.001	<0.001	0.605	<0.001	<0.001	<0.001
NH ₄ ⁺ -N	<0.001	<0.001	<0.001	0.554	<0.001	<0.001	<0.01
NH ₄ ⁺ -N/ NO ₃ ⁻ -N	0.222	<0.001	<0.001	<0.001	<0.001	<0.001	<0.01

and PF, respectively (Figures 1C,D), and these fluxes decreased significantly as N application rates increased in both forests. At high rates of N addition, the decrease in nitrification-derived N₂O fluxes was approximately 84.6% in the BF and 100% in the PF relative to those observed at low N addition rates (Figure 2B). The denitrification-derived N₂O fluxes were in the range of 0.4 to 1.7 $\mu\text{g N kg}^{-1} \text{ day}^{-1}$ and 0.2 to 0.8 $\mu\text{g N kg}^{-1} \text{ day}^{-1}$ in the BF and PF, respectively (Figures 1E,F) and maximum average of denitrification-derived N₂O emissions was observed in middle N application rate in both BF and PF (Figure 2B). Moreover, denitrification was the dominant pathway of soil N₂O production in the present study, accounting for 64 to 90% and 81 to 100% of the total N₂O fluxes in the BF and PF, respectively (Figure 2C).

Effects of N Addition on Soil Inorganic N, pH, Dissolved Organic Carbon, and Dissolved Organic Nitrogen

Soil NO₃⁻-N and NH₄⁺-N contents were significantly affected by forest type and N application rate (Table 2). The soil NO₃⁻-N and NH₄⁺-N contents increased from day 1 to day 9 and

then decreased significantly on the last day of the incubation experiment (Figure 3). These soil inorganic N contents increased with increasing N application rates, and they were significantly higher in the PF than in the BF (Figures 3A–D). Soil NH₄⁺-N contents in the control treatments were, respectively, ~9- and ~7-fold greater in the BF and PF compared with soil NO₃⁻-N contents (Figures 3E,F). Moreover, the ratio of soil NH₄⁺-N to soil NO₃⁻-N decreased with increasing N inputs. High N application significantly decreased soil pH in both forests. While high N addition significantly decreased soil DON content in PF, no significant effect of N addition on soil DOC was observed in this forest type (Figure 4).

Effects of N Addition on the Abundances of Nitrification and Denitrification Functional Genes

While the abundances of AOA- and AOB-*amoA* genes in the BF and PF were not significantly affected by N inputs (Figures 5A–D), those of *nirK* and *nirS* genes were increased substantially by N inputs (Figures 5E,F). The increase in abundance of *nirK* genes

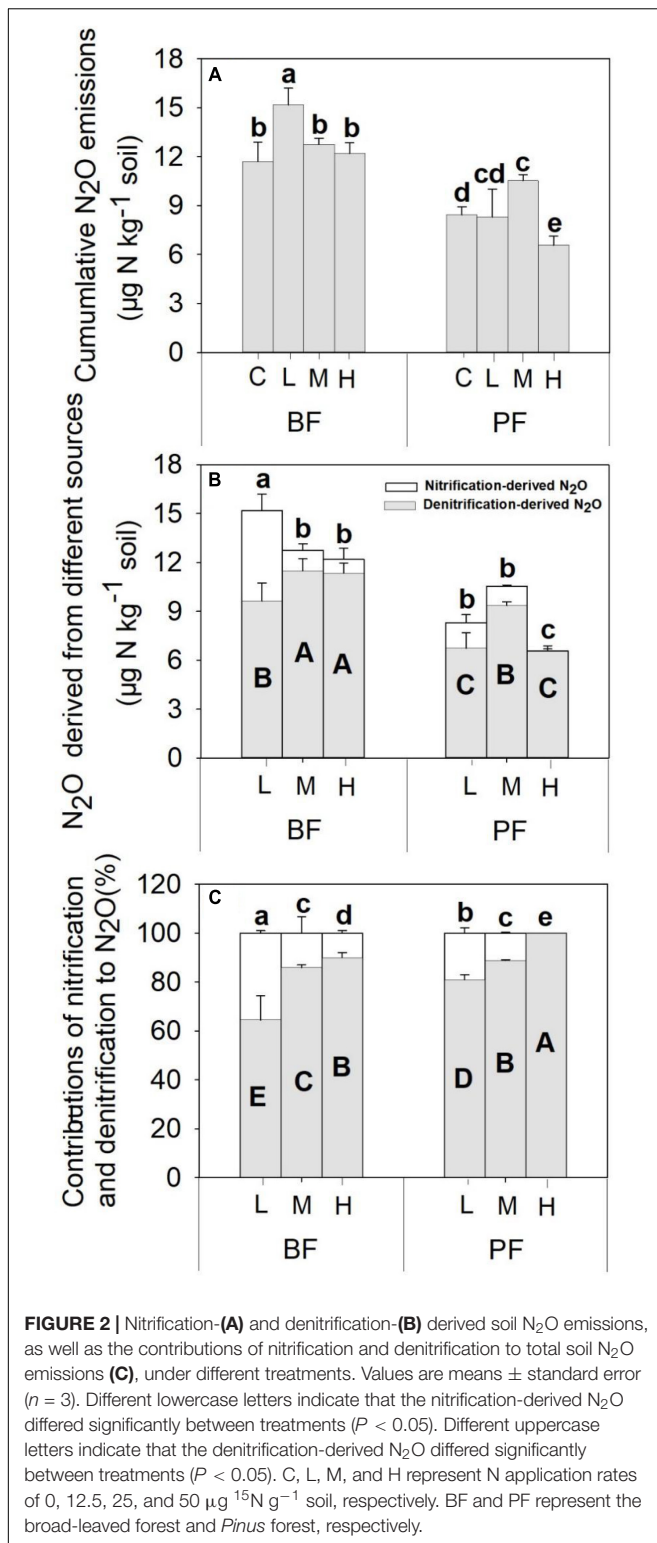


FIGURE 2 | Nitrification-(A) and denitrification-(B) derived soil N₂O emissions, as well as the contributions of nitrification and denitrification to total soil N₂O emissions (C), under different treatments. Values are means ± standard error ($n = 3$). Different lowercase letters indicate that the nitrification-derived N₂O differed significantly between treatments ($P < 0.05$). Different uppercase letters indicate that the denitrification-derived N₂O differed significantly between treatments ($P < 0.05$). C, L, M, and H represent N application rates of 0, 12.5, 25, and 50 μg N g⁻¹ soil, respectively. BF and PF represent the broad-leaved forest and *Pinus* forest, respectively.

in the PF (12%) was greater than that in the BF (5%), whereas the increase in abundance of *nirS* genes in the BF (15%) was slightly higher compared with that in the PF (14%). Inputs of N had no significant effect on *nosZ* gene abundance (Figure 5G).

Controlling Factors of Nitrification- and Denitrification-Derived N₂O Emissions

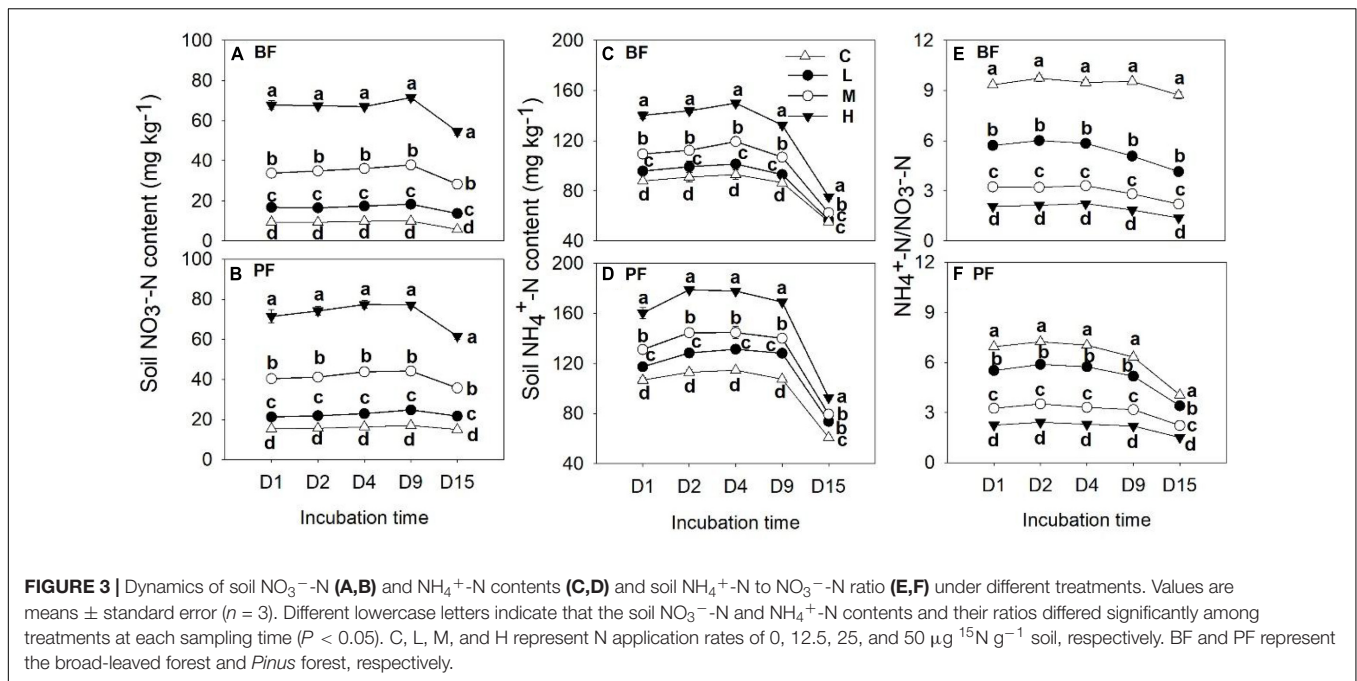
According to the SEM, soil properties (soil NH₄⁺-N, NO₃⁻-N, DON, and pH) and nitrification and denitrification functional genes (*AOA-amoA*, *AOB-amoA*, *nirK*, *nirS*, and *nosZ*) explained 99 and 95% of the variance observed in nitrification-derived N₂O emissions in the BF and PF, respectively (Figures 6A,B). These variables also could explain 55 and 88% of the variance of denitrification-derived N₂O emissions in the BF and PF, respectively (Figures 6C,D). The dominant factor in regulating soil nitrification-derived N₂O emission in the BF and PF was soil NH₄⁺-N (standardized total coefficient = -0.93, $P < 0.001$) and soil pH (standardized total coefficient = -0.74, $P < 0.001$), respectively, (Table 3). The dominant factors in regulating soil denitrification-derived N₂O in BF and PF were significantly differently (Figures 6C,D and Table 3). In the BF, it was soil pH. However, in the PF, the dominant regulatory factor was *nirS* (Table 3).

DISCUSSION

Effects of N Addition on Nitrification- and Denitrification-Derived N₂O Emissions

In the present study, soil N₂O fluxes responded non-linearly to N addition. Low- and middle-level N application rates increased soil N₂O emissions which might be due to N application and increased N availability (Figure 3) for nitrifying and denitrifying microorganisms. However, high N input had no significant effect on soil N₂O emission from BF or even decreased soil N₂O emission from PF (Figure 2B). Similarly, previous studies carried out in a subtropical montane forest also reported that short-term low and medium N addition favored N₂O production but this effect became weaker or even decreased N₂O emissions at high N addition (Han et al., 2018). A possible reason may be that the positive effect of high N application on denitrification-derived N₂O in both forests was partially offset by its negative effect on nitrification-derived N₂O (Figure 2B). Another explanation may be that the consumption of N₂O may be greater than the production of N₂O at high N treatment (Senbayram et al., 2012).

The application of N not only affects total soil N₂O emissions but also the contribution of nitrification and denitrification processes to soil N₂O production. In the present study, the average nitrification-derived N₂O emissions under different N treatments ranged from 0 to 0.37 μg N kg⁻¹ day⁻¹ (Figures 1C,D), which is similar to a previous study in subtropical forests [0.06–0.40 μg N kg⁻¹ day⁻¹, Zhang et al. (2011)]. Over 15 days of incubation, nitrification-derived N₂O contributed 0 to 36% of the total N₂O emissions (Figure 2). This range is within the range (<50%) that has been documented in previous studies carried out in the subtropical forest with similar N application rates (Zhang et al., 2011; Han et al., 2018). In contrast to our first hypothesis, our result showed a decrease in the contribution of the nitrification process to the total N₂O emissions following N application (Figure 3), and the magnitude of decrease was about 80 to 100% across the two forests. Deppe



et al. (2017) also found that N addition can significantly decrease the contribution of nitrification to N₂O production in a Haplic Luvisol soil, but the magnitude of decrease (7 to 21%) was not as pronounced as the one observed in the present study. One possible explanation for this phenomenon may be that N application decreased soil pH (Figure 4A), and soil nitrification rates have been demonstrated to be relatively low in acidic soils and decrease with decreasing soil pH (Cheng et al., 2013; Wang et al., 2020).

In contrast to the response of nitrification-derived N₂O emissions to N input as explained above, the highest denitrification-derived N₂O emissions occurred at the middle N application rate in both BF and PF, and the denitrification process is the dominant pathway for the production of soil N₂O emission, accounting for 64 to 100% of the total N₂O fluxes (Figure 3C). Previous studies carried out across subtropical forests reported that 53 to 100% of the total N₂O emissions were derived from denitrification (Zhang et al., 2011; Zhu et al., 2013a; Han et al., 2018). Moreover, the relative contributions of denitrification to N₂O emissions in forest soils were found to increase significantly with increasing N application (Zhu et al., 2013a; Morse et al., 2015; Niu et al., 2016), with the magnitude of increase ranging from 51 to 130% in a subtropical forest (Zhu et al., 2013a). The significantly higher relative contribution of denitrification to total N₂O production in the PF compared with the BF in the present study can be partly attributed to the greater decrease in nitrification-derived N₂O emissions in the PF (100%) than in the BF (80%).

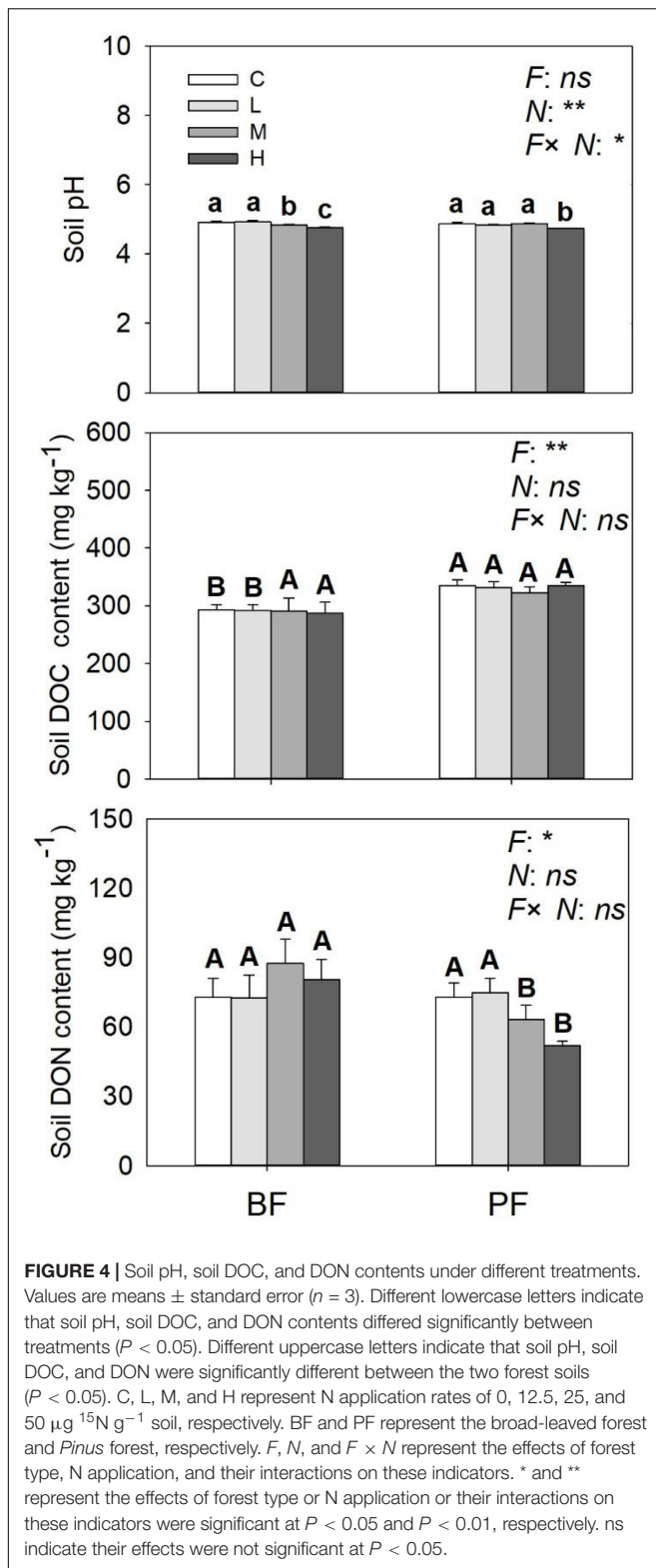
Moreover, both nitrification- and denitrification-derived N₂O emissions were significantly higher in the BF relative to the PF, which indicated that the BF was a more important N₂O emission source than the PF. This is consistent with our third hypothesis. A study by Zhang et al. (2008) in southern

China also reported that the emissions of N₂O from a mature monsoon evergreen BF were significantly higher compared to those from a PF. The authors attributed their observation to the higher soil N availability in the monsoon evergreen BF. However, their explanation cannot be extended to our findings because the BF had significantly lower soil available N content than the PF (Figure 3). Considerable differences in the amount of N₂O emissions among the investigated forest soils might be attributed to the differences in tree species, or more specifically, their influences on the soil's abiotic and/or biotic properties (Butterbach-Bahl et al., 1997; Qin et al., 2019). It has been reported that trees in BF are associated with arbuscular mycorrhizal fungi, which can promote soil microbial communities with higher N cycling potential and activity relative to microbial communities in soils dominated by the trees associated with ectomycorrhizal fungi in the PF (Mushinski et al., 2021).

Overall, our results supported part of our first hypothesis as the denitrification process was the dominant pathway of soil N₂O production. However, the denitrification-derived N₂O emissions were observed as non-linear responses to N application in both forests. In addition, the nitrification-derived N₂O emissions decreased with increasing N application in both BF and PF, which is in contrast with our first hypothesis. An explanation for this phenomenon is presented in section “Controlling Factors Related to Nitrification-Derived N₂O Emissions in Forest Soils.”

Controlling Factors Related to Nitrification-Derived N₂O Emissions in Forest Soils

Soil NH₄⁺-N content and soil pH contributed the most to the variations of nitrification-derived N₂O emissions in the BF and



PF, respectively (Figure 6 and Table 3). Their effect was more pronounced than the effects of AOB-*amoA* on nitrification-derived N₂O emission. Therefore, in contrast to our second

hypothesis, our results suggest that N-induced changes in soil environmental factors were more important than the abundance of *amoA* genes in regulating nitrification-derived N₂O emissions. Similarly, Tang et al. (2019) found that soil environmental factors (especially soil NH₄⁺ content) accounted for 50 to 77% of the variation in potential nitrification activities, which was better explained than nitrification functional genes in fertilized soils. In the BF, high soil NH₄⁺-N concentrations not only had a direct negative effect on nitrification-derived N₂O emission but also had an indirect negative effect *via* influencing soil pH and the abundance of AOB (Figure 6A). The reasons for high soil NH₄⁺-N contents reducing nitrification-derived N₂O emission may be as follows: ammonia oxidizers are unable to tolerate high NH₄⁺-N contents (> 100 mg N kg⁻¹ soil) in acidic soils (Gubry-Rangin et al., 2010; Carey et al., 2016; Farquharson, 2016; Breuillin-Sessoms et al., 2017; Li et al., 2018; Liu et al., 2018). In the current study, the average soil NH₄⁺-N content in N-amended soils was much higher than not only the aforementioned 100 mg N kg⁻¹ soil but also other soil NH₄⁺-N content averages that have been reported in previous studies in various subtropical fertilized forest soils (ranging from 1 to 50 mg N kg⁻¹ soil) (Han et al., 2018; Tang et al., 2018, 2019; Wang et al., 2018). This indicates the lack of efficient oxidation of NH₄⁺ to NO₃⁻ during nitrification that has led to lower nitrification-derived N₂O emissions in N-amended soils (Farquharson, 2016; Wang et al., 2020).

Soil NH₄⁺-N content also had a direct significant effect on nitrification-derived N₂O emission in the BF, but its total effect was offset by its indirect effect on soil pH (Figure 6B). This resulted in soil pH playing a more important role in regulating nitrification-derived N₂O emissions in the PF (Table 3). Previous studies have reported that soil with a lower pH support less microbial diversity and lower N-cycle potential (Cheng et al., 2019; Mushinski et al., 2021). A meta-analysis reports that N addition dramatically decreases the ratios of fungi to bacteria and microbial C to N on an average by 10 and 8%, respectively (Zhou et al., 2017). In addition, the nitrification rate is generally decreased with decreasing soil pH, and its rate can be inhibited in soil with a pH lower than 5 (Zhao et al., 2018; Cheng et al., 2019). This could explain why the nitrification-derived N₂O emission at high N addition in the PF is close to zero (Figure 2B). Overall, in terms of nitrification-derived N₂O emissions, changes in soil environmental factors (e.g., soil NH₄⁺-N and pH) induced by N application play more important roles than changes in the abundance of AOA and AOB in controlling its emission.

Controlling Factors Related to Denitrification-Derived N₂O Emissions in Forest Soils

Soil pH is the most important controller of denitrification-derived N₂O emission in the BF (Figure 6C and Table 3), and high N application significantly decreased soil pH (Figure 4). The reasons for soil pH regulating denitrification-derived N₂O emissions in BF may be as follows. Soil pH can exert a direct and an indirect effect on biological properties (e.g., reductase activities, the abundance, and expression of functional genes,

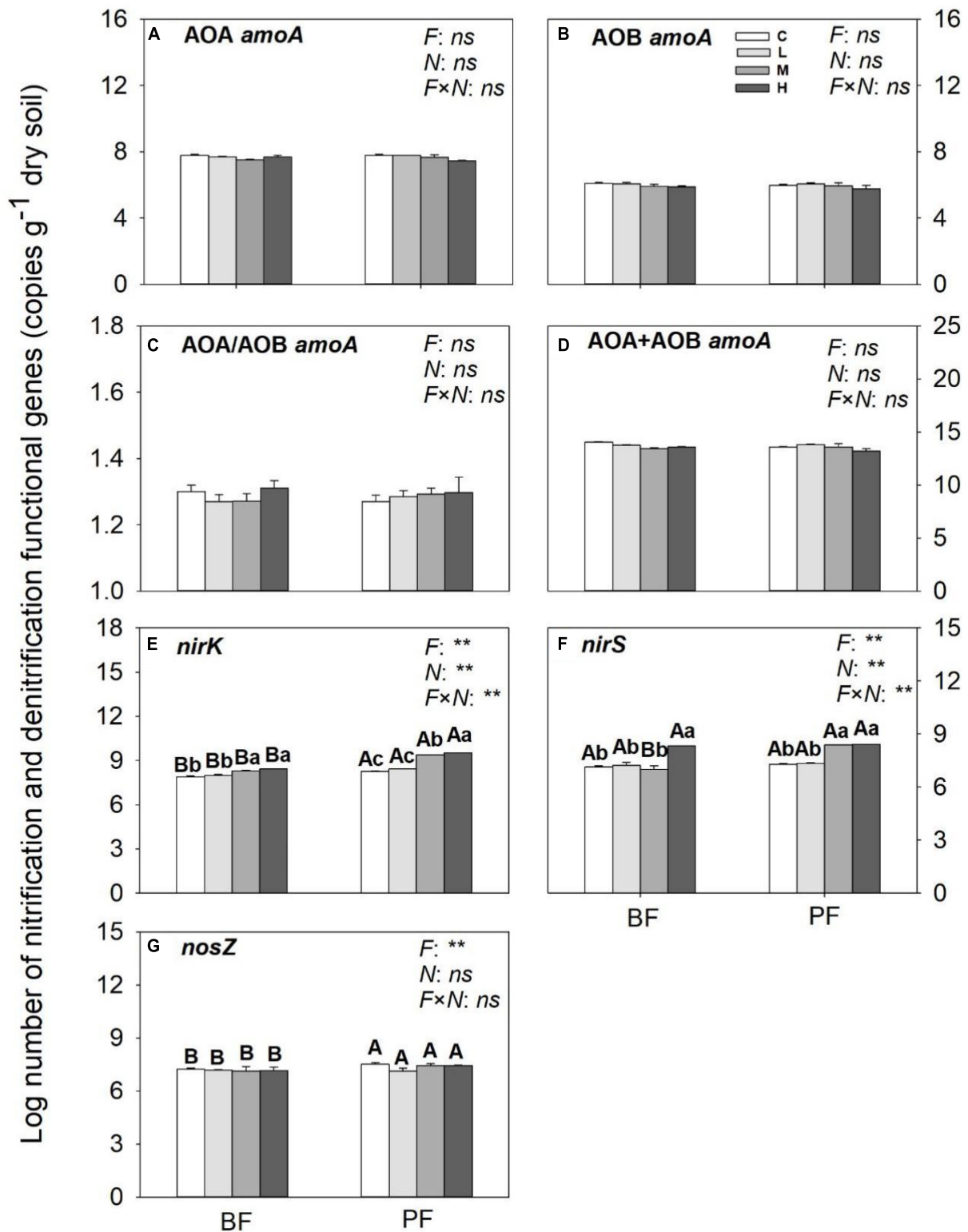


FIGURE 5 | Abundances of nitrification (A–D) and denitrification (E,F) functional genes under different treatments. Values are means ± standard error ($n = 3$). Different lowercase letters indicate that the functional gene abundance differed significantly among treatments ($P < 0.05$). C, L, M, and H represent N application rates of 0, 12.5, 25, and 50 $\mu\text{g } ^{15}\text{N g}^{-1}$ soil, respectively. BF and PF represent the broad-leaved forest and *Pinus* forest, respectively. *F*, *N*, and *F* × *N* represent the effects of forest type, N application, and their interactions on these indicators. ** represent the effects of forest type or N application or their interactions on these indicators were significant at $P < 0.05$ and $P < 0.01$, respectively. ns indicate their effects were not significant at $P < 0.05$.

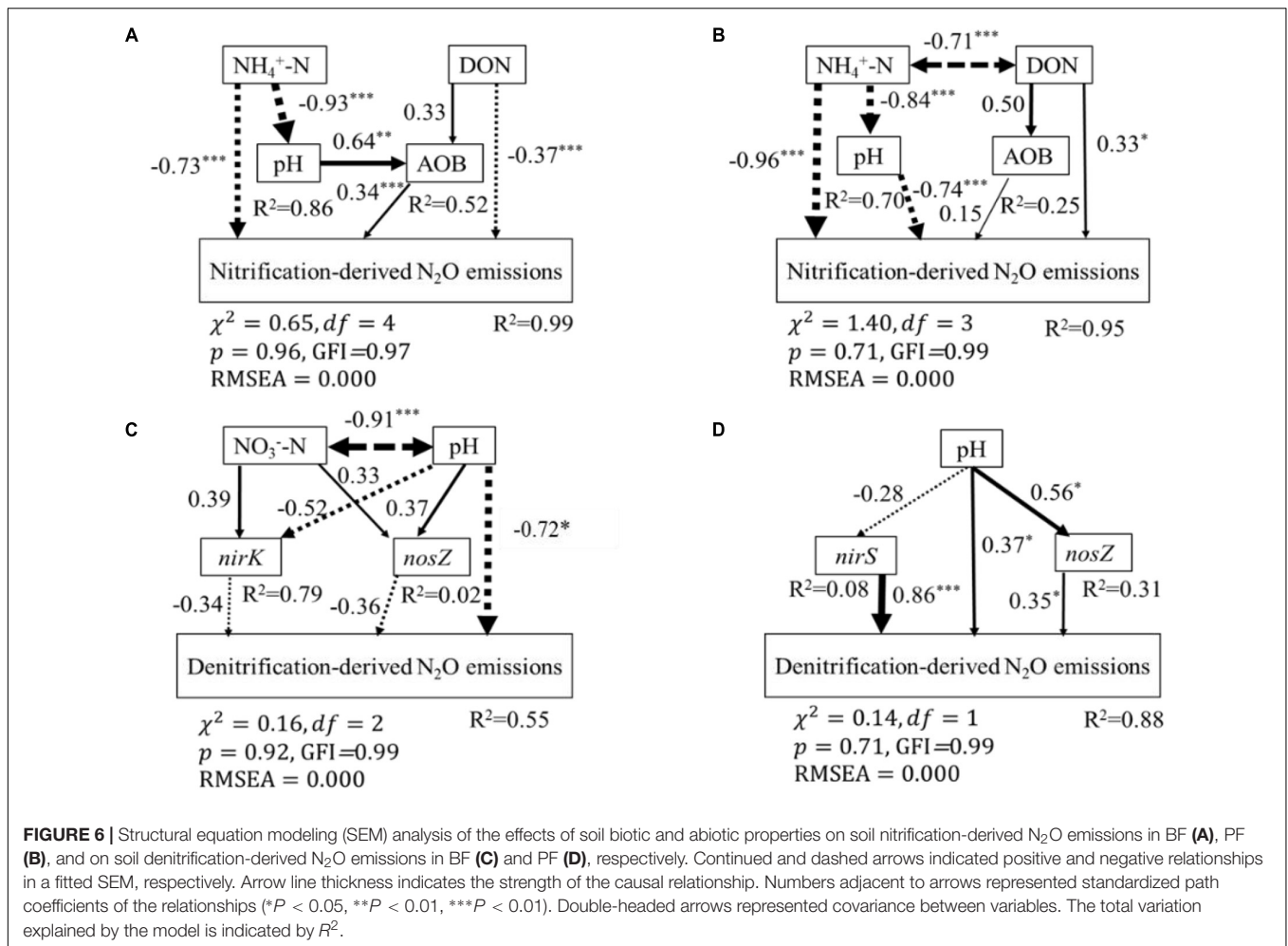


TABLE 3 | Standardized total effects of each explanatory variable on nitrification- and denitrification-derived N₂O emissions.

	Nitrification-derived N ₂ O emissions		Denitrification-derived N ₂ O emissions	
	Explanatory variable	Standardized total effects	Explanatory variable	Standardized total effects
BF	NH ₄ ⁺ -N	-0.93	pH	-0.85
	AOB	0.34	<i>nosZ</i>	-0.36
	DON	-0.26	<i>nirK</i>	-0.34
	pH	0.22	NO ₃ ⁻ -N	-0.25
PF	pH	-0.74	<i>nirS</i>	0.86
	NH ₄ ⁺ -N	-0.54	<i>nosZ</i>	0.35
	DON	0.41	pH	0.33
	AOB	0.15		

etc.) (Liu et al., 2010; Hu et al., 2015; Han et al., 2019). The reductases for nitrate, nitrite, and nitric oxide are more active at pH < 7 (Gillam et al., 2008), while the N₂O reductase activities are inhibited in soil with low pH (Levy-Booth et al., 2014). A review covering 50 years of research on the effects of soil pH on denitrification also reports that there is a consistent negative relationship between denitrification-derived N₂O production and soil pH (Šimek and Cooper, 2002). A close relationship between soil pH and denitrification-derived N₂O in the BF suggested that soil pH is a strong predictor of

denitrifier activity in the BF. In the PF, the abundance of *nirS* is the dominant factor controlling denitrification-derived N₂O emissions. Previous studies have also reported that the abundance of denitrification genes correlated with *nirS* gene abundance (Morales et al., 2010).

The dominant factors in regulating soil denitrification-derived N₂O emission were different in these two forests. This may be because differences in the tree species lead to variations in soil environment and the community composition of functional microbes, and changes in these variations induced the differences

in the nitrification- and denitrification-derived N₂O emissions in these acidic forest soils (Barberán et al., 2015; Soper et al., 2018). Moreover, regardless of nitrification- or denitrification-derived N₂O emissions, soil environmental factors are more important than nitrifier and denitrifier abundance in regulating soil N₂O emissions in both forests (except denitrification-derived N₂O emission in PF). Previous studies also demonstrate that tree species-induced changes in community composition and environmental factors play more important roles in regulating soil N₂O emissions than the abundance of nitrifier/denitrifier functional genes (Qin et al., 2019). In the present study, we still do not know how the species diversity or the nature of individual species affect nitrification- and denitrification-derived N₂O production in the field, even though several studies have investigated N₂O emission from plantation forest soils (Arai et al., 2008; Wang et al., 2014; Kou-Giesbrecht and Menge, 2021). Further study needs to focus on the role of the identities of specific species in the production processes of N₂O under different N inputs.

CONCLUSION

Our results demonstrated clearly that nitrification-derived N₂O emissions decreased with N application rates, while denitrification-derived N₂O emissions were non-linear responses to N addition. The total soil N₂O emissions, nitrification-, and denitrification-derived N₂O emissions were significantly higher in the broadleaf forest than in the conifers forest. Changes in soil environmental factors (i.e., pH and soil NH₄⁺-N content) induced by N addition are more important than the abundance of nitrification and denitrification functional genes in regulating the production process of soil N₂O. The incorporation of soil nitrogen substrates, soil pH, and forest type into the nitrogen cycling model will more precisely predict the global N₂O from soil under elevated N deposition.

REFERENCES

- Arai, S., Ishizuka, S., Ohta, S., Ansori, S., Tokuchi, N., Tanaka, N., et al. (2008). Potential N₂O emissions from leguminous tree plantation soils in the humid tropics. *Glob. Biogeochem. Cycles* 22:GB2028. doi: 10.1029/2007GB002965
- Barberán, A., McGuire, K. L., Wolf, J. A., Jones, F. A., Wright, S. J., Turner, B. L., et al. (2015). Relating belowground microbial composition to the taxonomic, phylogenetic, and functional trait distributions of trees in a tropical forest. *Ecol. Lett.* 18, 1397–1405. doi: 10.1111/ele.12536
- Breuilin-Sessoms, F., Venterea, R. T., Sadowsky, M. J., Coulter, J. A., Clough, T. J., and Wang, P. (2017). Nitrification gene ratio and free ammonia explain nitrite and nitrous oxide production in urea-amended soils. *Soil Biol. Biochem.* 111, 143–153. doi: 10.1016/j.soilbio.2017.04.007
- Butterbach-Bahl, K., Gasche, R., Breuer, L., and Papen, H. (1997). Fluxes of NO and N₂O from temperate forest soils: impact of forest type, N deposition and of liming on the NO and N₂O emissions. *Nutr. Cycl. Agroecosyst.* 48, 79–90. doi: 10.1023/A:1009785521107
- Cao, L., Liang, Y., Wang, Y., and Lu, H. (2015). Runoff and soil loss from Pinus massoniana forest in southern China after simulated rainfall. *Catena* 129, 1–8. doi: 10.1016/j.catena.2015.02.009
- Carey, C. J., Dove, N. C., Beman, J. M., Hart, S. C., and Aronson, E. L. (2016). Meta-analysis reveals ammonia-oxidizing bacteria respond more strongly to nitrogen

DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

AUTHOR CONTRIBUTIONS

QQ: conceptualization, methodology, software, investigation, supervision, validation, formal analysis, writing—original draft, writing—reviewing and editing, visualization, data curation, and resources. AM: investigation and writing—reviewing and editing. SJ: writing—original draft, writing—reviewing and editing, visualization, and supervision. YH: conceptualization, methodology, software, validation, formal analysis, writing—original draft, writing—reviewing and editing, visualization, and supervision. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fpls.2022.950367/full#supplementary-material>

- addition than ammonia-oxidizing archaea. *Soil Biol. Biochem.* 99, 158–166. doi: 10.1016/j.soilbio.2016.05.014
- Chen, W. B., and Peng, S. L. (2020). Land-use legacy effects shape microbial contribution to N₂O production in three tropical forests. *Geoderma* 358, 113979. doi: 10.1016/j.geoderma.2019.113979
- Cheng, Y., Wang, J., Chang, S. X., Cai, Z., Müller, C., and Zhang, J. (2019). Nitrogen deposition affects both net and gross soil nitrogen transformations in forest ecosystems: a review. *Environ. Pollut.* 244, 608–616. doi: 10.1016/j.envpol.2018.10.054
- Cheng, Y., Wang, J., Mary, B., Zhang, J. B., Cai, Z. C., and Chang, S. X. (2013). Soil pH has contrasting effects on gross and net nitrogen mineralizations in adjacent forest and grassland soils in central Alberta, Canada. *Soil Biol. Biochem.* 57, 848–857. doi: 10.1016/j.soilbio.2012.08.021
- Deppe, M., Well, R., Giesemann, A., Spott, O., and Flessa, H. (2017). Soil N₂O fluxes and related processes in laboratory incubations simulating ammonium fertilizer depots. *Soil Biol. Biochem.* 104, 68–80. doi: 10.1016/j.soilbio.2016.10.005
- Fang, Y., Koba, K., Makabe, A., Takahashi, C., Zhu, W., Hayashi, T., et al. (2015). Microbial denitrification dominates nitrate losses from forest ecosystems. *Proc. Natl. Acad. Sci. U.S.A.* 112, 1470. doi: 10.1073/pnas.1416776112
- Farquharson, R. (2016). Nitrification rates and associated nitrous oxide emissions from agricultural soils – a synopsis. *Soil Res.* 54, 469–480. doi: 10.1071/SR15304

- Gillam, K., Zebarth, B., and Burton, D. (2008). Nitrous oxide emissions from denitrification and the partitioning of gaseous losses as affected by nitrate and carbon addition and soil aeration. *Can. J. Soil Sci.* 88, 133–143. doi: 10.4141/CJSS06005
- Gubry-Rangin, C., Nicol, G. W., and Prosser, J. I. (2010). Archaea rather than bacteria control nitrification in two agricultural acidic soils. *FEMS Microbiol. Ecol.* 74, 566–574. doi: 10.1111/j.1574-6941.2010.00971.x
- Gurmesa, G. A., Wang, A., Li, S., Peng, S., de Vries, W., Gundersen, P., et al. (2022). Retention of deposited ammonium and nitrate and its impact on the global forest carbon sink. *Nat. Commun.* 13, 1–9. doi: 10.1038/s41467-022-28345-1
- Han, X., Shen, W., Zhang, J., and Müller, C. (2018). Microbial adaptation to long-term N supply prevents large responses in N dynamics and N losses of a subtropical forest. *Sci. Total Environ.* 626, 1175–1187. doi: 10.1016/j.scitotenv.2018.01.132
- Han, X., Xu, C., Nie, Y., He, J., Wang, W., Deng, Q., et al. (2019). Seasonal variations in N₂O emissions in a subtropical forest with exogenous nitrogen enrichment are predominately influenced by the abundances of soil nitrifiers and denitrifiers. *J. Geophys. Res. Biogeosci.* 124, 3635–3651. doi: 10.1029/2019JG005477
- Hu, H. W., Chen, D., and He, J. Z. (2015). Microbial regulation of terrestrial nitrous oxide formation: understanding the biological pathways for prediction of emission rates. *FEMS Microbiol. Rev.* 39, 729–749. doi: 10.1093/fems/rev/fuv021
- IPCC (2013). “Climate change 2013: the physical science basis,” in *Contribution of Working Group I to the Fifth Assessment Report of the Intergovernmental Panel on Climate Change*, eds T. F. Stocker, D. Qin, G. K. Plattner, M. Tignor, S. K. Allen, J. Boschung, et al. (Cambridge: Cambridge University Press), 1535.
- Jones, D. L., and Willett, V. B. (2006). Experimental evaluation of methods to quantify dissolved organic nitrogen (DON) and dissolved organic carbon (DOC) in soil. *Soil Biol. Biochem.* 38, 991–999. doi: 10.1016/j.soilbio.2005.08.012
- Kou-Giesbrecht, S., and Menge, D. N. (2021). Nitrogen-fixing trees increase soil nitrous oxide emissions: a meta-analysis. *Ecology* 102:e03415. doi: 10.1002/ecy.3415
- Levy-Booth, D. J., Prescott, C. E., and Grayston, S. J. (2014). Microbial functional genes involved in nitrogen fixation, nitrification and denitrification in forest ecosystems. *Soil Biol. Biochem.* 75, 11–25. doi: 10.1016/j.soilbio.2014.03.021
- Li, D., Zhang, X., Green, S. M., Dungait, J. A. J., Wen, X., Tang, Y., et al. (2018). Nitrogen functional gene activity in soil profiles under progressive vegetative recovery after abandonment of agriculture at the Puding Karst Critical Zone Observatory, SW China. *Soil Biol. Biochem.* 125, 93–102. doi: 10.1016/j.soilbio.2018.07.004
- Liu, B., Morkved, P. T., Frostegård, Å, and Bakken, L. R. (2010). Denitrification gene pools, transcription and kinetics of NO, N₂O and N₂ production as affected by soil pH. *FEMS Microbiol. Ecol.* 72, 407–417. doi: 10.1111/j.1574-6941.2010.00856.x
- Liu, H., Ding, Y., Zhang, Q., Liu, X., Xu, J., Li, Y., et al. (2018). Heterotrophic nitrification and denitrification are the main sources of nitrous oxide in two paddy soils. *Plant Soil* 445, 39–53. doi: 10.1007/s11104-018-3860-x
- Liu, X., Zhang, Y., Han, W., Tang, A., Shen, J., Cui, Z., et al. (2013). Enhanced nitrogen deposition over China. *Nature* 494:459. doi: 10.1038/nature11917
- Lu, M., Yang, Y., Luo, Y., Fang, C., Zhou, X., Chen, J., et al. (2011). Responses of ecosystem nitrogen cycle to nitrogen addition: a meta-analysis. *New Phytol.* 189, 1040–1050. doi: 10.1111/j.1469-8137.2010.03563.x
- Lyu, M., Nie, Y., Giardina, C. P., Vadeboncoeur, M. A., Ren, Y., Fu, Z., et al. (2019). Litter quality and site characteristics interact to affect the response of priming effect to temperature in subtropical forests. *Funct. Ecol.* 33, 2226–2238. doi: 10.1111/1365-2435.13428
- Morales, S. E., Cosart, T., and Holben, W. E. (2010). Bacterial gene abundances as indicators of greenhouse gas emission in soils. *ISME J.* 4, 799–808. doi: 10.1038/ismej.2010.8
- Morse, J. L., Durán, J., Beall, F., Enanga, E. M., Creed, I. F., Fernandez, I., et al. (2015). Soil denitrification fluxes from three northeastern North American forests across a range of nitrogen deposition. *Oecologia* 177, 17–27. doi: 10.1007/s00442-014-3117-1
- Mushinski, R. M., Payne, Z. C., Raff, J. D., Craig, M. E., Pusede, S. E., Rusch, D. B., et al. (2021). Nitrogen cycling microbiomes are structured by plant mycorrhizal associations with consequences for nitrogen oxide fluxes in forests. *Glob. Change Biol.* 27, 1068–1082. doi: 10.1111/gcb.15439
- Niu, S., Classen, A. T., Dukes, J. S., Kardol, P., Liu, L., Luo, Y., et al. (2016). Global patterns and substrate-based mechanisms of the terrestrial nitrogen cycle. *Ecol. Lett.* 19, 697–709. doi: 10.1111/ele.12591
- Pett-Ridge, J., Petersen, D. G., Nuccio, E., and Firestone, M. K. (2013). Influence of oxic/anoxic fluctuations on ammonia oxidizers and nitrification potential in a wet tropical soil. *FEMS Microbiol. Ecol.* 85, 179–194. doi: 10.1111/1574-6941.12111
- Qin, H. L., Xing, X. Y., Tang, Y. F., Hou, H. J., Yang, J., Shen, R., et al. (2019). Linking soil N₂O emissions with soil microbial community abundance and structure related to nitrogen cycle in two acid forest soils. *Plant Soil* 435, 95–109. doi: 10.1007/s11104-018-3863-7
- Senbayram, M., Chen, R., Budai, A., Bakken, L., and Dittert, K. (2012). N₂O emission and the N₂O/(N₂O+N₂) product ratio of denitrification as controlled by available carbon substrates and nitrate concentrations. *Agric. Ecosyst. Environ.* 147, 4–12. doi: 10.1016/j.agee.2011.06.022
- Shcherbak, I., Millar, N., and Robertson, G. P. (2014). Global metaanalysis of the nonlinear response of soil nitrous oxide (N₂O) emissions to fertilizer nitrogen. *Proc. Natl. Acad. Sci. U.S.A.* 111, 9199. doi: 10.1073/pnas.1322434111
- Šimek, M., and Cooper, J. E. (2002). The influence of soil pH on denitrification: progress towards the understanding of this interaction over the last 50 years. *Eur. J. Soil Sci.* 53, 345–354. doi: 10.1046/j.1365-2389.2002.00461.x
- Solomon, S. (2007). *Climate Change 2007 – The Physical Science Basis: Working Group I Contribution to the Fourth Assessment Report of the IPCC*. Cambridge: Cambridge University Press.
- Soper, F. M., Sullivan, B. W., Nasto, M. K., Osborne, B. B., Bru, D., Balzotti, C. S., et al. (2018). Remotely sensed canopy nitrogen correlates with nitrous oxide emissions in a lowland tropical rainforest. *Ecology* 99, 2080–2089. doi: 10.1002/ecy.2434
- Stevens, R., Laughlin, R., Burns, L., Arah, J., and Hood, R. (1997). Measuring the contributions of nitrification and denitrification to the flux of nitrous oxide from soil. *Soil Biol. Biochem.* 29, 139–151. doi: 10.1016/S0038-0717(96)00303-3
- Tang, W., Chen, D., Phillips, O. L., Liu, X., Zhou, Z., Li, Y., et al. (2018). Effects of long-term increased N deposition on tropical montane forest soil N₂ and N₂O emissions. *Soil Biol. Biochem.* 126, 194–203. doi: 10.1016/j.soilbio.2018.08.027
- Tang, Y., Yu, G., Zhang, X., Wang, Q., Tian, D., Tian, J., et al. (2019). Environmental variables better explain changes in potential nitrification and denitrification activities than microbial properties in fertilized forest soils. *Sci. Total Environ.* 647, 653–662. doi: 10.1016/j.scitotenv.2018.07.437
- Wang, G., Zhou, Y., Xu, X., Ruan, H., and Wang, J. (2013). Temperature sensitivity of soil organic carbon mineralization along an elevation gradient in the Wuyi Mountains, China. *PLoS One* 8:e53914. doi: 10.1371/journal.pone.0053914
- Wang, H., Liu, S., Zhang, X., Mao, Q., Li, X., You, Y., et al. (2018). Nitrogen addition reduces soil bacterial richness, while phosphorus addition alters community composition in an old-growth N-rich tropical forest in southern China. *Soil Biol. Biochem.* 127, 22–30. doi: 10.1016/j.soilbio.2018.08.022
- Wang, J., Tu, X. S., Zhang, H. M., Cui, J. Y., Ni, K., Chen, J. L., et al. (2020). Effects of ammonium-based nitrogen addition on soil nitrification and nitrogen gas emissions depend on fertilizer-induced changes in pH in a tea plantation soil. *Sci. Total Environ.* 747:8. doi: 10.1016/j.scitotenv.2020.141340
- Wang, Y., Cheng, S., Fang, H., Yu, G., Xu, M., Dang, X., et al. (2014). Simulated nitrogen deposition reduces CH₄ uptake and increases N₂O emission from a subtropical plantation forest soil in southern China. *PLoS One* 9:e93571. doi: 10.1371/journal.pone.0093571
- Xu, W., Li, W., Chen, S., Yang, D., Zhang, Y., Xue, J., et al. (2016). Quantification of nitrogen dry and wet deposition in Fujian tobacco planting area. *Acta Ecol. Sin.* 36, 424–432. doi: 10.1016/j.chnaes.2016.09.007
- Xu, X., Zhou, Y., Ruan, H., Luo, Y., and Wang, J. (2010). Temperature sensitivity increases with soil organic carbon recalcitrance along an elevational gradient in the Wuyi Mountains, China. *Soil Biol. Biochem.* 42, 1811–1815. doi: 10.1016/j.soilbio.2010.06.021
- Zhang, J., Cai, Z., and Zhu, T. (2011). N₂O production pathways in the subtropical acid forest soils in China. *Environ. Res.* 111, 643–649. doi: 10.1016/j.envres.2011.04.005
- Zhang, K., Zhu, Q., Liu, J., Wang, M., Zhou, X., Li, M., et al. (2019). Spatial and temporal variations of N₂O emissions from global forest and grassland

- ecosystems. *Agric. For. Meteorol.* 266–267, 129–139. doi: 10.1016/j.agrformet.2018.12.011
- Zhang, W., Mo, J., Yu, G., Fang, Y., Li, D., Lu, X., et al. (2008). Emissions of nitrous oxide from three tropical forests in Southern China in response to simulated nitrogen deposition. *Plant Soil* 306, 221–236. doi: 10.1007/s11104-008-9575-7
- Zhao, W., Zhang, J., Müller, C., and Cai, Z. (2018). Effects of pH and mineralisation on nitrification in a subtropical acid forest soil. *Soil Res.* 56, 275–283. doi: 10.1071/SR17087
- Zhou, G., Peng, C., Li, Y., Liu, S., Zhang, Q., Tang, X., et al. (2013). A climate change-induced threat to the ecological resilience of a subtropical monsoon evergreen broad-leaved forest in Southern China. *Glob. Change Biol.* 19, 1197–1210. doi: 10.1111/gcb.12128
- Zhou, Z., Wang, C., Zheng, M., Jiang, L., and Luo, Y. (2017). Patterns and mechanisms of responses by soil microbial communities to nitrogen addition. *Soil Biol. Biochem.* 115, 433–441. doi: 10.1016/j.soilbio.2017.09.015
- Zhu, J., He, N., Wang, Q., Yuan, G., Wen, D., Yu, G., et al. (2015). The composition, spatial patterns, and influencing factors of atmospheric wet nitrogen deposition in Chinese terrestrial ecosystems. *Sci. Total Environ.* 511, 777–785. doi: 10.1016/j.scitotenv.2014.12.038
- Zhu, J., Mulder, J., Bakken, L., and Dörsch, P. (2013a). The importance of denitrification for N₂O emissions from an N-saturated forest in SW China: results from in situ ¹⁵N labeling experiments. *Biogeochemistry* 116, 103–117. doi: 10.1007/s10533-013-9883-8
- Zhu, J., Mulder, J., Solheimslid, S. O., and Dörsch, P. (2013b). Functional traits of denitrification in a subtropical forest catchment in China with high atmospheric N deposition. *Soil Biol. Biochem.* 57, 577–586. doi: 10.1016/j.soilbio.2012.09.017
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