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Fungal denitrification: Bipolaris sorokiniana exclusively denitrifies inorganic nitrogen in the presence and absence of oxygen

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One sentence summary: Inorganic nitrogen addition alters fungal denitrification and respiration in the presence and absence of oxygen. Editor: Matthias Brock

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ABSTRACT

Fungi may play an important role in the production of the greenhouse gas nitrous oxide (N_2O). Bipolaris sorokiniana is a ubiquitous saprobe found in soils worldwide, yet denitrification by this fungal strain has not previously been reported. We aimed to test if B. sorokiniana would produce N_2O and CO_2 in the presence of organic and inorganic forms of nitrogen (N) under microaerobic and anaerobic conditions. Nitrogen source (organic-N, inorganic-N, no-N control) significantly affected N_2O and CO_2 production both in the presence and absence of oxygen, which contrasts with bacterial denitrification. Inorganic N addition increased denitrification of N_2O (from 0 to 0.3 μ g N_2O -N h⁻¹ g⁻¹ biomass) and reduced respiration of N_2O (from 0.1 to 0.02 mg N_2O h⁻¹ g⁻¹ biomass). Isotope analyses indicated that nitrite, rather than ammonium or glutamine, was transformed to N_2O . Results suggest the source of N may play a larger role in fungal N_2O production than oxygen status.

Keywords: nitrite; glutamine; nitrous oxide; carbon dioxide; organic nitrogen; co-denitrification

INTRODUCTION

Fungal denitrification may contribute more to soil emissions of the greenhouse gas nitrous oxide (N_2O) than bacterial denitrification (Laughlin and Stevens 2002; Chen, Mothapo and Shi 2015), but physicochemical factors potentially altering fungal nitrogen (N) cycling require investigation. While many soil bacteria contain the N_2O reductase gene (nosZ) (required for N_2O to N_2 conversion), this gene is not found in fungi. Nonetheless, N_2 production by soil fungi has been reported (Shoun et al. 2012; Long

et al. 2013). Some fungi, particularly for genera in the Hypocreales order (such as Fusarium), are capable of producing N₂O (Maeda et al. 2015) under anaerobic (Zumft 1997; Morozkina and Kurakov 2007; Shoun et al. 2012) or microaerobic conditions (Zhou et al. 2001; Morozkina and Kurakov 2007; Takaya 2009). Two pathways of fungal denitrification have been reported: (a) classical denitrification of nitrate or nitrite when oxygen is limited and insufficient to support aerobic respiration and (b) hybrid formation of N₂O, where two N sources are combined (Spott, Russow

and Stange 2011). It is not clear if induction of anaerobiosis or changes in N source (Takaya 2002; Wei et al. 2014) cause a shift from one pathway to another. Hybrid N₂O and/or N₂ formation (also referred to as co-denitrification) has been observed in the presence of inorganic and organic N (Su, Takaya and Shoun 2004; Spott, Russow and Stange 2011; Long et al. 2013) but may also occur in the presence of two inorganic forms of N (Spott, Russow and Stange 2011). Data are lacking that indicate how inorganic and organic sources of N affect fungal denitrification and co-denitrification to N_2O pre- and postinduction of anaerobiosis. Addition of two N sources (one enriched with 15N) would allow us to determine if N2O formation could be attributed to denitrification or co-denitrification.

While a number of fungal strains have been tested for denitrification potential (Maeda et al. 2015), many strains, such as Bipolaris sorokiniana, have not been tested. Bipolaris sorokiniana is the asexual form (anamorph) of the fungus Cochliobolus sativus, a common plant pathogen in the order Pleosporales with a wide range of plant hosts. It is particularly common on cereals in the Poaceae family and is widely known as the causal agent for the common root rot. It can also survive as thick-walled conidia or as saprotrophic mycelium in soil or crop debris. Geographic distribution of B. sorokiniana is worldwide, particularly near midlatitudes, in grasslands and agricultural fields (Kumar et al. 2002). It is not known if this soil fungus will denitrify inorganic or organic forms of N and if aerobic respiration and denitrification co-occur under controlled conditions. While a large body of research has reported bacterial N2O and CO2 production under microaerobic and anaerobic conditions (Butterbach-Bahl et al. 2013), few reports are available indicating if rates of fungal CO2 and N2O production vary with inorganic and/or organic N. It is unclear if fungi incubated with O2 will respond to induction of anaerobiosis in a manner similar to denitrifying bacteria, with increased rates of N2O production occurring within hours of induction (Firestone and Tiedje 1979; Smith and Tiedje 1979). The goal of this research was to test how N source (organic or inorganic) influences production of N2O and CO2 by B. sorokiniana pre and post anaerobiosis. Results add to the emerging body of knowledge regarding controls on fungal denitrification and codenitrification processes.

MATERIALS AND METHODS

We used pure culture of B. sorokiniana (Sacc.) Shoemaker [telemorph: C. sativus (S. Ito & Kurib.) Drechsler ex Dastur] to test how organic N [glutamine (C5H10N2O3)] and inorganic N [ammonium sulfate ((NH₄)₂SO₄) and sodium nitrite $(NaNO_2)$] sources might influence fungal production of N_2O and CO_2 under sterile conditions. We controlled O_2 status with air-tight laboratory incubation vessels to evaluate rates of N2O and CO2 production pre and post anaerobiosis on the same samples (Firestone and Tiedje 1979; Smith and Tiedje 1979; Zhou et al. 2001; Butterbach-Bahl et al. 2013). We obtained isolate ICMP 6809 from the ICMP culture collection (http://www.landcareresearch.co.nz/resources/collections/icmp) in its anamorphic state. It was grown under aerobic conditions in the same media previously used by Rohe et al. (2014a,b) and Shoun et al. (1992), containing 1% glucose, 0.2% peptone, 0.02% MgSO₄·7 H₂O, 2 ppm CaCl₂·6 H₂O, 2 ppm FeSO₄ ·7 H₂O and 0.01 mol potassium phosphate (pH 7.4). This media, which contained N as peptone, was used only for fungal growth. Another media, where the N source was omitted, was used for the incubations and referred to here as no-N media. After 7 days, cultures were washed, drained and stored in a reduced volume of the no-N media. Using the no-N media, four solutions were prepared for fungal inoculation: (a) no-N media (b) 0.5 mmols N as $C_5H_{10}N_2O_3$, (c) 0.25 mmols N as $Na^{15}NO_2$ (99.5 atom%; Cambridge Isotope Laboratory, Andover, MA) and 0.25 mmols N as $(NH_4)_2SO_4$, or (d) 0.25 mmols N as $Na^{15}NO_2$ and 0.25 mmols N as $C_5H_{10}N_2O_3$. The $Na^{15}NO_2$ was used to determine if these fungi would preferentially use NO2 to form N2O (thus forming ⁴⁶N₂O) or if other sources of N would be utilized.

Approximately 10 ml of fungal biomass were transferred to 125 ml serum bottles, and four replicate bottles were inoculated with 1 ml of each sterile solution and 1 ml of no-N media. Additional bottles containing media solutions without fungi were also prepared. All bottles were sealed and the headspace evacuated and flushed with ultrapure helium, and then injected with O2 to achieve 0.4% O2 headspace before setting up on a robotic gas chromatograph (GC) fitted with electron capture and thermal conductivity detectors (Phillips et al. 2014; McMillan et al. 2015). Bottles were placed on the GC and measured every 6 hours for N2O, CO2 and O2 at 19°C. Following this 48-h incubation experiment, bottles were then evacuated and flushed three times to create an anaerobic headspace for the second experiment, where the only difference between experiments was headspace O2 concentration. Sterility was maintained and conditions remained constant during both incubations, including pH (ranged from 6.2 to 6.9). Data were normalized to the corresponding dry weight of fungal biomass in each bottle. The N isotopes for N₂O gas in the headspace of each sample at the end of the incubations were measured on a continuous-flow isotope ratio mass spectrometer (Thermo Finnigan Delta V; Thermo Scientific, Waltham, MA) in line with an automated gas bench interface (Thermo Gas Bench II) to determine if ${}^{45}N_2O$ or ${}^{46}N_2O$ were present. Precision of the isotopic analysis was <1 atom%. Data were analysed for effects of N source treatment on N2O and CO2 production rates for microaerobic and anaerobic experiments separately with a generalized linear model and means were compared with Tukey's test.

RESULTS AND DISCUSSION

Table 1 and Fig. 1 show effects of treatment on CO2 and N2O production for the microaerobic incubations were similar to effects of treatment for the anaerobic incubations. Fungi inoculated with only organic-N (C5H10N2O3) or no-N media only linearly produced CO_2 (1–2 mg CO_2 h⁻¹ g⁻¹) but not N_2O . Fungi inoculated with inorganic-N (Na15NO2 or ((NH4)2SO4) or inorganic-N plus organic-N (Na15NO2 and C5H10N2O3) linearly produced N_2O (0.2–0.3 μg N_2O -N h^{-1} g^{-1}) but an order of magnitude less CO_2 . The mass $^{46}N_2O$ comprised 94.7%–97.3% of all N_2O in the headspace, yielding an equivalent 15 N of 98 to 99 (± 0.3) atom%. This enrichment was effectively equivalent to enrichment of the $^{15}\text{NO}_2$ used in the experiment. The lack of headspace $^{45}\text{N}_2\text{O}$ indicated hybrid formation of N2O by B. sorokiniana did not occur under any of these conditions, and NO_2^- was the only N form used to form N2O. Headspace O2 (Fig. 2) over the aerobic time course indicated rapid declines for the no-N media and organic-N treatments but not the inorganic-N treatments. Minimal rates of CO₂ respiration (Table 1) in the presence of inorganic-N were consistent with lack of O2 utilization by B. sorokiniana (Fig. 2). We also observed chemodenitrification (6%-8% of total biologically produced N2O) for media amended with inorganic-N only (van Cleemput 1998; Kampschreur et al. 2011; Jones et al. 2015).

Table 1. Rates of N₂O and CO₂ produced during (1) aerobic and (2) anaerobic incubation experiments by treatment per gram of fungal biomass. Average rates followed by different letters for each experiment represent significant differences between treatments. The same letters indicate no differences between treatments.

Experiment	Treatment	Mean ng N_2 O-N h^{-1} g^{-1} biomass	Mean		
			Std. Dev.	μ g CO ₂ h ⁻¹ g ⁻¹ biomass	Std. Dev.
Anaerobic	No N	0.51ª	0.26	896.45ª	206.08
	Organic N	0.29 ^a	0.11	728.46 ^a	148.66
	Inorganic N	311.30 ^b	18.98	6.17 ^b	1.34
	Inorganic N + organic N	211.79 ^c	14.44	7.87 ^b	1.48
Aerobic	No N	0.57ª	0.23	1179.89ª	262.62
	Organic N	0.32ª	0.23	988.19 ^a	222.95
	Inorganic N	354.55 ^b	26.00	18.26 ^b	1.48
	Inorganic N + organic N	306.33 ^c	28.61	31.52 ^b	4.03

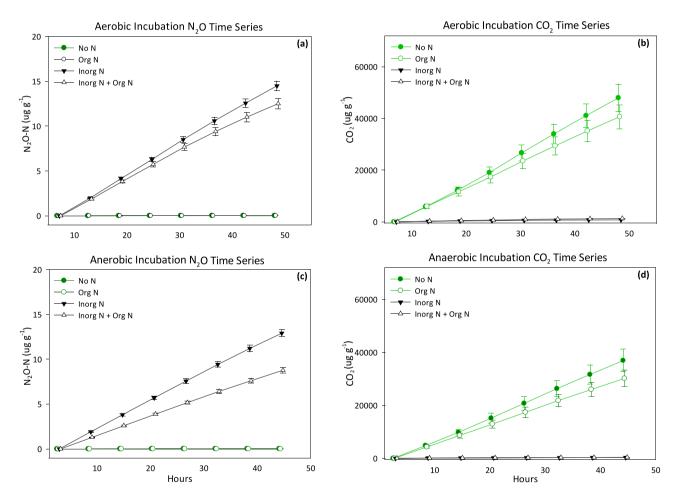


Figure 1. Average (std. dev; n = 4) N₂O and CO₂ measured in the headspace per g of fungal biomass by treatment during aerobic (a, b) and anaerobic (c, d) incubations.

Evidence of chemodenitrification, where N2O was produced under sterile conditions, may be due to reactions between NO₂and reduced metals in the media (van Cleemput 1998; Kampschreur et al. 2011). We chose the media for this experiment because it is commonly used for fungal denitrification investigations (Shoun et al. 1992; Rohe et al. 2014a,b). Kampschreur et al. (2011) used a chemostat to demonstrate that NO₂ - to NO is an equilibrium-based reaction, where emissions of NO and N2O were coupled to iron oxidation under anoxic conditions. A recent review by Medinets et al. (2015) argued that NO2-, a precursor of NO, is central to processes associated with chemodenitrification. Additional research is needed to determine the reactants involved in this specific media and the prevalence of chemodenitrification in an oxygenated headspace.

Evidence of biological denitrification coupled with reduced CO2 respiration in the presence of inorganic-N may be due to B. sorokiniana preferentially denitrifying as a defence against accumulation of NO₂⁻ (Geets, Boon and Verstraete 2006; Clark et al. 2012). Fungi are capable of converting NO₂⁻ to NO, the precursor to soil N2O emissions (Russow, Stange and Nueue 2009), through expression of the NO₂⁻ reductase gene (nirK), and NO is a regulator of O2 consumption in eukaryotes (Thomas et al.

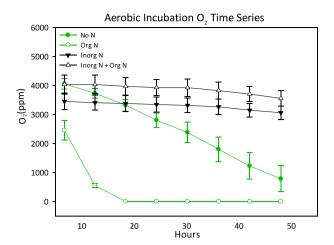


Figure 2. Average (std. dev; n = 4) O₂ measured in the headspace by treatment during aerobic incubation

2001). We suspect that some NO₂⁻ was converted to NO (both abiotic and biotic), which inhibited mitochondrial respiration (Thomas et al. 2001; Medinets et al. 2015) and therefore CO2 production, as shown in Fig. 1 and Table 1. One way of discerning the importance of NO_2 reduction to NO and N_2O by B. sorokiniana would be to identify nirK gene expression in conjunction with NO measurements. If B. sorokiniana does not reduce NO₂⁻ to NO, then chemical transformation of NO₂⁻ may be the source of NO (van Cleemput 1998; Kampschreur et al. 2011). In either event, there is a need to identify chemical versus biological sources of NO, the precursor to N2O (Russow, Stange and

Denitrification in soils and sediments is referred to as a keystone ecosystem service with positive water quality implications. However, when N2O is the final end product of denitrification, there are negative impact on stratospheric ozone and radiative forcing in the troposphere (Erisman et al. 2013). Here, B. sorokiniana, a ubiquitous soil fungus, demonstrated the capacity to transform dissolved NO_2^- to gaseous N_2O at the expense of aerobic respiration, a more energetically favourable pathway. Denitrification and aerobic respiration for B. sorokiniana were closely linked via NO₂⁻. Additional investigations into alternative respiration pathways and effects of inorganic N on C assimilation are warranted, particularly with respect to O₂ status. Here, we focused on how inorganic and organic N sources influence pathways to N₂O production before and after onset of anaerobiosis (Firestone and Tiedje 1979; Smith and Tiedje 1979; Zhou et al. 2001). Future work should be designed with sample in parallel to test how O2 and N interact to affect fungal denitrification phenotype and genotype. Our results suggest that fungal denitrification may not be driven by anaerobiosis but instead by form of N; however, additional strains need to be tested to determine if this is widespread among soil fungi or unique to B. sorokiniana.

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