#### **ORIGINAL PAPER**



# Effects of soil water content at freezing, thaw temperature, and snowmelt infiltration on $N_2O$ emissions and denitrifier gene and transcript abundance during a single freeze-thaw event

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#### Abstract

Climate change-related warming and increased precipitation may alter winter snow cover and thawing events, and therefore, may carry significant consequences for nitrous oxide (N<sub>2</sub>O) production pathways such as denitrification, and the abundance and expression of denitrifying microorganisms. We used a soil microcosm study to investigate the combined effect of soil thaw temperature, initial water filled pore space (WFPS) prior to soil freezing, and snowmelt infiltration simulated by the addition of water on N<sub>2</sub>O emission and denitrification rates, soil respiration rate, and the abundance and transcription of denitrifying (*nirK*, *nirS*, and *nosZ*) bacteria during a single freeze-thaw event. Soil respiration rate was primarily controlled by an increase in soil thaw temperature, whereas soil N<sub>2</sub>O emission and denitrification rates were generally greater in soils with a higher initial WFPS and soil thaw temperature. In contrast, snowmelt infiltration generally had a negligible effect on these rates, which may be related to pre-existing soil conditions that were already conducive to denitrification. Unexpectedly, the *nosZ* transcript/*nosZ* gene abundance ratio was lower in soils thawed at 8.0 °C compared to 1.5 °C; however, this may have resulted in a lower N<sub>2</sub>O reduction, thus explaining the greater levels of N<sub>2</sub>O emitted from soils thawed at 8.0 °C. Overall, this study demonstrated that increased N<sub>2</sub>O production during a single freeze-thaw event was primarily linked to antecedent conditions of high initial WFPS, soil thaw temperature, and a synergistic interplay between these two environmental parameters, and provides evidence that an increase in annual temperature and precipitation, along with the timing of precipitation, may further stimulate N<sub>2</sub>O production pathways.

Keywords Water-filled pore space · Soil microcosms · Nitrous oxide · Denitrifier abundance · Denitrifier transcription

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## Introduction

In Canada, agriculture accounts for 75% of total anthropogenic annual emissions of the potent greenhouse gas nitrous oxide (N<sub>2</sub>O) (Environment and Climate Change Canada 2023), of which non-growing season fluxes associated with freeze-thaw cycles represent 29–73% of the annual N<sub>2</sub>O emissions (Risk et al. 2013; Wagner-Riddle et al. 2017; Hung et al. 2021). Snow cover, which insulates soil from changes in air temperature, can affect winter soil nitrogen (N) cycling and soil gaseous emissions (Brin et al. 2018, 2019; Jia et al. 2021). However, recent climate-related increases in annual temperature and precipitation may alter the amount, timing, and continuity of winter snow cover and thawing events (Huntington et al. 2004; Hayhoe et al. 2007; Brooks et al. 2011; Blankinship and Hart 2012). Subsequently, these changes may lead to later snow onset,

increased mid-winter snowmelt events, and early spring snowmelt, and therefore, may carry implications for denitrification and  $N_2O$  production.

Although increased N<sub>2</sub>O production from a snowmelt event was shown to be partially due to the physical release of trapped gas build-up under ice layers within the soil (Bremner et al. 1980; Burton and Beauchamp 1994; Teepe et al. 2001), substantial emissions arise from *de novo*  $N_2O$ production (Wagner-Riddle et al. 2008; Németh et al. 2014; Risk et al. 2014). Snowmelt infiltration increases the soil water content, which subsequently lowers soil oxygen  $(O_2)$ concentrations and creates anoxic microsites that are suitable for denitrifier activity and N<sub>2</sub>O production (Teepe et al. 2000; Koponen and Martikainen 2004; Congreves et al. 2018). Snowmelt infiltration also increases substrate availability for N<sub>2</sub>O production pathways through increased nutrient transport, cell lysis from excessive influx of water into microbial cells, and cytoplasmic solute secretion from osmotic regulation (Christensen and Christensen 1991; Fierer and Schimel 2003; Congreves et al. 2018).

The effect of snowmelt infiltration on  $N_2O$  emissions during a freeze-thaw event is, however, complex and may be confounded by several factors (Congreves et al. 2018). Intermittent  $N_2O$  fluxes during freeze-thaw events are also related to changes in soil temperature (Congreves et al. 2018; Brin et al. 2018, 2019; Badewa et al. 2022). Soil warming increases microbial metabolism and anoxic microsite formation that promote  $N_2O$  producing pathways such as denitrification (Henry 2008, 2013; Chantigny et al. 2019) and increases soil water content that further stimulates microbial activity and denitrification (Phillips 2008; Risk et al. 2013; Congreves et al. 2018).

Soil moisture is another primary control of  $N_2O$  production during freeze-thaw events, where high soil moisture leads to greater  $N_2O$  pulses during spring thaw (Chen et al. 2018, 2021; Brin et al. 2019; Li et al. 2021; Badewa et al. 2022). It has been shown that high spring thaw  $N_2O$  fluxes were mainly due to pre-winter conditions (i.e., soil moisture at the time of soil freezing) rather than spring soil conditions (Banerjee et al. 2016; Brin et al. 2018). These findings were attributed to a greater disruption of soil aggregates and microbes in moist soils during soil freezing that resulted in an increased release of substrates for denitrification (Brin et al. 2018, 2019; Congreves et al. 2018).

Given the interconnection between snowmelt infiltration, soil warming, and soil moisture, there is reason to believe a concomitant change in these biophysical factors will exacerbate  $N_2O$  production during a freeze-thaw event. Few studies have investigated the combined effect of soil moisture prior to soil freezing and soil thaw temperature on  $N_2O$  emissions (Koponen and Martikainen 2004; Chen et al. 2021; Yang et al. 2022); however, there are currently no studies that have investigated the interaction between snowmelt infiltration, soil thaw temperature, and soil moisture prior to soil freezing on  $N_2O$  production and denitrifier abundance and expression during a freeze-thaw event. A better understanding of the mechanisms by which thawing leads to  $N_2O$  production will help predict changes that may occur with climate change-altered winter snow cover.

The objective of this study was to investigate the effects of soil moisture prior to soil freezing (initial water-filled pore space [WFPS] of 15% or 30%), soil thaw temperature (1.5-8.0 °C), and snowmelt infiltration (simulated by with or without water addition) on soil inorganic N availability, soil respiration, denitrification rate, N2O emissions and denitrifier gene abundance and transcription. We hypothesized that soil under conditions that are more conducive to denitrification, i.e., with a combination of greater WFPS before soil freezing, higher soil thaw temperature, and snowmelt infiltration, would have greater N<sub>2</sub>O emissions, denitrification rates, denitrifier abundance and denitrifier gene transcription, compared to soils with lower WFPS, lower soil thaw temperature, and no snowmelt infiltration. We also hypothesized that the timing of soil water additions would influence N2O production during thaw, where 'prewinter' soil water conditions (initial WFPS) would have a priming effect on the denitrifier communities, resulting in a greater impact on N<sub>2</sub>O production than 'spring' soil water conditions (snowmelt infiltration).

## Materials and methods

## **Experimental design**

This microcosm experiment consisted of a  $2 \times 2 \times 2$  factorial arrangement of treatments in a completely randomized design (n=4). Factors included two initial WFPS treatments before soil freezing [15% ( $W_{15}$ ) or 30% ( $W_{30}$ )], two soil thaw temperatures [1.5 °C ( $T_{1.5}$ ) or 8.0 °C ( $T_{8.0}$ )], and two snowmelt infiltration treatments simulated by water addition [no snowmelt (SM<sup>-</sup>) or snowmelt (SM<sup>+</sup>)]. In total, there were eight treatment combinations:  $W_{15}T_{1.5}SM^-$ ,  $W_{15}T_{8.0}SM^-$ ,  $W_{15}T_{8.0}SM^+$ ,  $W_{30}T_{1.5}SM^-$ ,  $W_{30}T_{1.5}SM^+$ ,  $W_{30}T_{8.0}SM^-$ , and  $W_{30}T_{8.0}SM^+$ .

The initial WFPS treatments of 15% and 30% were selected based on the range of volumetric water content observed in snow-covered plots in a field during winter freezing (late December 2013-January 2014) (Brin et al. 2018). Soil thaw temperature treatments of 1.5 °C and 8.0 °C were selected based on the range of mean daily temperatures observed in snow-covered plots in a field during spring thaw (late February-April 2014) (Brin et al. 2018). To implement the snowmelt infiltration treatments, soils were treated with

deionized water (0.08 mL dH<sub>2</sub>O g<sup>-1</sup> dry soil; temperature of 0.5 °C). This snowmelt volume was selected based on the volumetric water content observed in snow-covered plots in a field during spring thaw (late February-April 2014) (Brin et al. 2018). The deionized water was added at once, as this was representative of a rapid snow melt frequently observed in the spring (greater atmospheric temperature under sunny or rainy conditions) or during the coldest months of winter, January and February (sudden increase in atmospheric temperature accompanied by abundant rainfall).

#### Soil sampling

Soil (0–20 cm depth) was collected from a field under a potato (*Solanum tuberosum* L.)-barley (*Hordeum vulgare* L.) rotation at the Fredericton Research and Development Centre, Agriculture and Agri-Food Canada located in Fredericton, New Brunswick, Canada (45°55'23"N, 66°36'25"W). The soil texture was a sandy loam (680 g sand kg<sup>-1</sup>, 314 g silt kg<sup>-1</sup>, and 66 g clay kg<sup>-1</sup>) (pipette method with organic matter removal, Kroetsch and Wang 2007). Soil organic C and total N concentrations were 10.3 g C kg<sup>-1</sup> and 1.2 g N kg<sup>-1</sup>, respectively, as determined by dry combustion (Elementar varioMACRO, Skjemstad and Baldock 2007). Soil pH (1:1 water dilution) was 6.0. The soil was collected in early December 2014, air-dried, passed through a 4.75 mm sieve, and held at 4.0 °C until microcosm preparation in late February 2015.

#### Microcosm preparation and sampling

A schematic of the microcosm preparation and sampling is presented in Fig. 1. While working in a walk-in refrigerator set at 4.0 °C, soil for each replicate was divided into bins designated for the two initial WFPS treatments. To each bin, ground red clover residues were added at a rate of 1000 mg C kg<sup>-1</sup> dry soil, which falls within the range of expected organic C additions in agricultural soils after the plow-down of plant residues (Bolinder et al. 2002; Miller et al. 2012). Soil nitrate  $(NO_3)$  solution was added to the soil to obtain a concentration of 15 mg N kg<sup>-1</sup> dry soil. The red clover residues had a C/N ratio of 14:1 and represented a forage crop that is frequently incorporated into agricultural soils in autumn in eastern Canada as a source of organic matter. Concentrations of added NO<sub>3</sub><sup>-</sup> and organic C were in the range of concentrations encountered under field conditions following autumn crop residue incorporation (Bolinder et al. 2002). The WFPS of soil was determined using the gravimetric water content (GWC) and the following calculation: 100 x (GWC x bulk density)/total soil porosity, where total soil porosity = 1-(bulk density/2.65 g cm<sup>-3</sup> (soil particle density). The soil was at an initial WFPS of 15%. The amount of water required to obtain a WFPS of 30% was determined using the same calculation and deionized water was added to the bins for the W<sub>30</sub> treatment. Soils of W<sub>15</sub> and W<sub>30</sub> treatments were well-homogenized in the bins by gently mixing.

A 500 mL glass canning jar served as an individual soil microcosm. The soil (276 g equivalent of oven-dry soil) was added to the microcosms and then hand-packed to the target bulk density of 1.03 Mg m<sup>-3</sup> which is similar to field

	Soil pre-treatment	7-day soil incubation	Measurements
STEP 1	STEP 2	STEP 3	STEP 4
WFPS 15 - Red clover - NO <sub>3</sub> <sup>-</sup> solution OR	To simulate winter conditions: Incubated at -1.0°C for 4 weeks	<ul> <li>T<sub>pre-thaw</sub> = before soil thaw or snowmelt infiltration</li> <li>T<sub>0-7days</sub> = snowmelt infiltration incubated at 1.5°C or 8.0°C</li> </ul>	Jar set 1: N <sub>2</sub> O and CO <sub>2</sub> emissions Jar set 2: Soil properties and denitrifier gene and transcript abundance
		Two sampling approaches:	
WFPS 30 - Red clover - NO <sub>3</sub> <sup>-</sup> solution		Jar set 1: 20 non-destructive sampling time points Jar set 2 & 3: 3 destructive sampling time points	Jar set 3: Denitrification rate using the acetylene blockage method
- H <sub>2</sub> O			

Fig. 1 Schematic overview of treatment implementation and experimental set-up of the single freeze-thaw event used in this study. Soils were adjusted to an initial water filled pore space (WFPS) of either 15% or 30% (W<sub>15</sub> or W<sub>30</sub>) and incubated at -1.0 °C for four-weeks.

Soils were then incubated for a seven-day period at either 1.5 or  $8.0 \,^{\circ}$ C and were amended with or without water to simulate snowmelt infiltration (SM<sup>+</sup> or SM<sup>-</sup>). The measurements performed in each jar set are presented under Step 4.

conditions and ensures similar oxygen concentration and substrate availability among soil microcosms. Microcosms were covered with Parafilm pierced with holes to reduce water evaporation and allow for gas exchange, then placed in incubators (model MIR-553, Sanyo Scientific, Japan) set at -1.0 °C for four weeks. The soil freeze temperature of -1.0 °C was selected based on the maintained soil temperature observed in snow-covered plots in a field during winter freezing until spring thaw (late December 2013-late February 2014) (Brin et al. 2018). After four weeks of winter freezing, microcosms were then moved to an incubator set at 1.5-8.0 °C to implement the soil thaw temperature treatments  $(T_{1,5} \text{ or } T_{8,0})$  and were treated with or without deionized water to implement the snowmelt infiltration treatments (SM<sup>+</sup> or SM<sup>-</sup>). Deionized water was added all at once in the soil microcosms. This is representative of field conditions; a sudden increase in atmospheric temperature accompanied by abundant rainfall resulting in an intense snow melt is frequently observed in the spring but can even occur during the coldest months of winter (January and February) in the Maritimes. The addition of snowmelt to  $W_{15}$  and  $W_{30}$ treatments resulted in a WFPS of 30% and 44%, respectively. The WFPS within treatments was stable throughout the incubation (Fig. 2). Soil temperatures were monitored in



**Fig. 2** Percent water filled pore space (WFPS) at the time of soil freezing (15% or 30%) and throughout the seven-day incubation following the onset of soil thawing (at 1.5 or 8.0 °C) and with or without the addition of water to simulate snowmelt infiltration (SM<sup>+</sup> or SM<sup>-</sup>).

soil microcosms that were not used for sampling throughout the incubation using iButton technology (Thermochron, Whitewater, WI, USA) by placing an iButton data logger in the soil, and temperature data can be found in the supplemental information (Figure S1a-d).

Soil and gas samples were collected at day 0 (i.e., the end of the four-week freezing period at -1.0 °C) and immediately before the thaw temperature and snowmelt infiltration treatments were imposed. The day 0 time point was used as baseline data for examining the temporal changes in N<sub>2</sub>O emissions, denitrification rates, soil inorganic N concentration, soil moisture, and denitrifier gene abundance and transcription from day = 0 to the end of the incubation period. Following this, the thaw temperature and snowmelt infiltration treatments were implemented, and three sets of microcosms were used to quantify all desired parameters. The headspace of the first set was frequently non-destructively sampled throughout the incubation period to quantify soil respiration and N<sub>2</sub>O emissions every 4.8 h (with 9.6 h overnight) from 0 to 3 days and then every 12 h from 3 to 7 days. The second and third sets were destructively sampled at 1, 3, and 7 days, where the second set was used to quantify soil inorganic N concentration, soil WFPS, and denitrifier gene abundance and transcription and the third set to quantify denitrification rate. The sampling frequency and incubation length were selected based on previous studies that observed that the majority of CO<sub>2</sub> and N<sub>2</sub>O emissions occurred within 72 h following soil thawing/ re-wetting and lasted for 7 days of incubation (Priemé et al. 2001; Koponen and Martikainen 2004; Teepe et al. 2004).

# Soil respiration, N<sub>2</sub>O emissions and denitrification rate

At each sampling time point, the first set of microcosms designated for frequent gas sampling were sealed with a screw-top lid fitted with a rubber septum, and 10% of the headspace volume was replaced with compressed air to provide a sufficient volume of gas for sampling. To quantify soil respiration (i.e., emissions of  $CO_2$ ) and  $N_2O$  emission rates, headspace gas (20 mL) was collected at 0, 30, and 60 min and transferred to a pre-evacuated 12 mL glass vial (Exetainers; Labco, High Wycombe, U.K.). After gas sampling, the microcosms were re-covered with Parafilm pierced with holes.

Rates of total denitrification  $(N_2O + N_2)$  were quantified using the third set of microcosms which were sealed with a screw-top lid at the time of sampling. 10% of the headspace volume was replaced with acetylene  $(C_2H_2)$  to quantify total denitrification. The presence of  $C_2H_2$  inhibits  $N_2O$  reductase activity; consequently,  $N_2O$  emissions quantified in the presence of  $C_2H_2$  reflect total denitrification as specified by Groffman et al. (2006). A 60-min  $C_2H_2$  diffusion period (Miller et al. 2008) occurred before the removal of the initial gas sample, and therefore 20 mL of headspace gas was collected at 60, 120, 180, and 240 min after sealing and transferred to a pre-evacuated 12 mL glass vial (Exetainers; Labco, High Wycombe, U.K.).

Headspace gas samples were analyzed for  $CO_2$  and  $N_2O$  concentrations using a Varian Star 3800 gas chromatograph (Varian, Mississauga, ON), as described by Burton et al. (2008). Soil respiration,  $N_2O$  emissions, and denitrification rates were calculated as the change in the mass of  $CO_2$ -C or  $N_2O$ -N in the microcosm headspace over 60 min (frequent sampling) or 180 min (destructive sampling). Cumulative gas fluxes of  $N_2O$  and  $CO_2$  measurements (jar set 1) and denitrification gas flux (jar set 3) were calculated over the 7-day incubation period using the linear trapezoidal method of integration between sampling time points (Burton et al. 2008). For the cumulative denitrification gas flux, it was assumed that the gas flux measured on the sampling date was representative of the average daily flux.

#### Soil inorganic N concentration and moisture

Soil sub-samples were collected from the second set of microcosms.. Soil inorganic N was extracted by shaking 25 g moist soil with 50 mL 0.5M K<sub>2</sub>SO<sub>4</sub> for 30 min at room temperature (~20 °C). Extracts were filtered using vacuum filtration and then stored at -20 °C until analysis. Concentrations of soil  $NO_2^{-}N + NO_3^{-}N$  (expressed as  $NO_3^{-}N$ ) and ammonium (NH4<sup>+</sup>-N) were measured using a Technicon AutoAnalyzer II system following Technicon Industrial Methods #100-70 W and #98-70 W, respectively (Technicon Industrial Systems, Terrytown, MA). Ammonium fixation is not a concern in this soil given the low amount of clay (66 g kg<sup>-1</sup> dry soil) and that it consists mostly of kaolinite, a 1:1 clay mineral. Soil moisture (GWC) was also determined in soils from the second set of microcosms. Sub-samples of moist soil were collected and dried in an oven set at 105 °C for 3 days.

#### **DNA and RNA extractions**

Soil sub-samples (~10 g) were taken from the second set of microcosms, flash frozen in liquid N<sub>2</sub> then stored at -80 °C. Both DNA and RNA were extracted from 1 g of freezedried soil using the method of Griffiths et al. (2000) and were further purified using the Power Clean DNA Cleanup kit (Mo-Bio, Carlsbad, CA) for DNA using a DNase I treatment and the RNeasy mini kit (Qiagen, Toronto, ON, Canada) for RNA. DNA and RNA were quantified using Picogreen and Ribogreen kits (Invitrogen, Burlington, ON, Canada). Reverse transcription of RNA to cDNA was performed using the SuperScript VILO cDNA Synthesis Kit (Invitrogen) and 0.2 mg mL<sup>-1</sup> of BSA using the following conditions: 25 °C for 10 min, 42 °C for 72 min, and 85 °C for 5 min.

#### **Quantitative PCR**

Copy numbers of nirS, nirK, and nosZ clade I and the transcript abundance of nosZ clade I were measured by qPCR on DNA and cDNA samples as described by Dandie et al. (2011) with concentration of templates ranging from 2 to 14 ng and 0.2 to 0.4 ng for DNA and cDNA, respectively, using a StepOne Plus Real-time PCR system (Applied Biosystems, Streetsvilles, ON, Canada). Absence of DNA contamination in RNA samples were verified by qPCR on non-reverse transcribed RNA for a subset of samples. Standards were linearized plasmids containing cloned nirS, nirK, and nosZ gene sequences as described in Dandie et al. (2011) and curve descriptors were as follows: for *nirS* copy numbers: slope = -3.7 to -3.9, efficiency (*E*) = 80.9 to 85.8%, intercept (Y)=46.5 to 47.6, and  $R^2 = 0.998$ ; for *nirK* copy numbers: slope = -3.5 to -3.7, efficiency (*E*) = 83.3 to 90.9%, intercept (Y)=43.6 to 48.6, and  $R^2$ =0.998 to 1.00; for *nosZ* clade I copy numbers: slope = -3.4 to -3.7, efficiency (E) = 87.5 to 88.3%, intercept (Y) = 43.6 to 44.6, and  $R^2 = 0.991$  to 0.997; and for *nosZ* clade I transcript numbers: slope = -3.5 to -3.8, efficiency (E) = 83.7 to 91.3%, intercept (Y) = 43.9 to 44.7, and  $R^2 = 0.994$  to 0.997.

## **Statistical analyses**

The RStudio base statistic software was used to conduct statistical analyses (v.4.1.3). Data were assessed for normality (i.e., normal and independent distribution) using homoscedasticity diagnostic plots and Shapiro-Wilks tests. For the frequent gaseous emission rates, a repeated measures ANOVA was performed using initial WFPS, thaw temperature, snowmelt, and time (repeated measure; 20 sampling time points) as fixed factors. ANOVAs for destructive denitrification rate, soil inorganic N concentrations, gene abundances, and the ratio of nosZ transcript / nosZ gene copy number were performed using initial WFPS, thaw temperature, snowmelt, and time (three sampling time points) as fixed factors. Treatment, time, and interaction means were tested using post-hoc Tukey Honest Significant Difference (HSD) tests. The treatment means and standard errors presented in figures were calculated from untransformed data. Significance was accepted at  $P \le 0.05$ .

# Results

#### Soil inorganic N concentrations

Soil NH<sub>4</sub><sup>+</sup>-N concentration did not vary with initial WFPS at day 0 i.e., at the end of the four-week freezing period and before the thaw temperature and snowmelt infiltration treatments were imposed (Fig. 3a). There was a significant initial WFPS x thaw temperature x time interaction on soil NH<sub>4</sub><sup>+</sup>-N concentration (P=0.021; Table S1). Soil NH<sub>4</sub><sup>+</sup>-N concentration increased over time for all treatments, and the increase over time was greater for the T<sub>8.0</sub> thaw temperature than for the T<sub>1.5</sub> thaw temperature, whereas the effect of initial WFPS was somewhat inconsistent, with high initial WFPS increasing NH<sub>4</sub><sup>+</sup>-N concentration only on day 7 for the T<sub>1.5</sub> thaw temperature and only on day 3 for the T<sub>8.0</sub> thaw temperature (Fig. 3a). There was no significant effect of snowmelt on soil NH<sub>4</sub><sup>+</sup>-N concentration (Table S1).

Soil NO<sub>3</sub><sup>-</sup>-N concentration at day 0 was lower for the  $W_{30}$  treatments compared with the  $W_{15}$  treatments (Fig. 3b). There was a significant initial WFPS x thaw temperature interaction on soil NO<sub>3</sub><sup>-</sup>-N concentration (*P*=0.021; Table S1). The decrease in soil nitrate due to high initial WFPS

was greater at  $T_{8.0}$  thaw temperature than at  $T_{1.5}$  thaw temperature (Fig. 3b). There was also a significant initial WFPS x snowmelt interaction on NO<sub>3</sub><sup>-</sup>-N concentration (P=0.003; Table S1) where there was a greater decrease in NO<sub>3</sub><sup>-</sup>-N concentration for the low initial WFPS than for the high initial WFPS in the presence of snowmelt (Fig. 3b). In addition, there was a significant main effect of time on soil NO<sub>3</sub><sup>-</sup>-N concentration (P < 0.001; Table S1), in which soil nitrate concentrations significantly decreased over time for all treatments (Fig. 3b).

# Soil respiration, N<sub>2</sub>O emissions, and denitrification rate

# Non-destructive sampling for soil respiration and N<sub>2</sub>O emissions

There was a significant initial WFPS x thaw temperature x time interaction on soil respiration (P < 0.001; Table S2). Soil respiration at T<sub>8.0</sub> thaw temperature was approximately double that of the T<sub>1.5</sub> thaw temperature and was generally similar regardless of initial WFPS (Fig. 4a). Soil respiration for the W<sub>30</sub> treatment at the T<sub>8.0</sub> thaw temperature was greater





**Fig. 3** Soil ammonium (mg N kg<sup>-1</sup> dry soil) and nitrate (mg N kg<sup>-1</sup> dry soil) concentration throughout the seven-day incubation of a single-freeze thaw event. Soils were adjusted to an initial water filled pore space (WFPS) of either 15% or 30% (W<sub>15</sub> or W<sub>30</sub>) and incubated at -1.0 °C for four-weeks. Soils were sampled at day 0 i.e. at the end

of the freezing period and immediately before the treatments were imposed. Soils were then incubated for a seven-day period at either 1.5 or 8.0 °C and were amended with or without water to simulate snowmelt infiltration (SM<sup>+</sup> or SM<sup>-</sup>). Values represent the mean (n=4) and the standard error.

than for the  $W_{15}$  treatment for about the first two days, after which soil respiration was generally similar across initial WFPS treatments and relatively consistent over time. Soil respiration at the  $T_{1.5}$  thaw temperature changed little over time (Fig. 4a). There was also a significant main effect of snowmelt, where the addition of snowmelt increased soil respiration rate (P < 0.001; Table S2).

There was a significant interaction between initial WFPS x thaw temperature on N<sub>2</sub>O emissions (P < 0.001, Table S2). Soil N<sub>2</sub>O emission rates were significantly greater in the W<sub>30</sub>T<sub>8.0</sub> treatment compared to the W<sub>30</sub>T<sub>1.5</sub>, W<sub>15</sub>T<sub>1.5</sub>, and W<sub>15</sub>T<sub>8.0</sub> treatments, and in the W<sub>30</sub>T<sub>1.5</sub> treatment compared to the W<sub>15</sub>T<sub>1.5</sub> and W<sub>15</sub>T<sub>8.0</sub> treatments (Fig. 4b). There was also a significant interaction between initial WFPS x time (P < 0.001; Table S2) where N<sub>2</sub>O emission rates increased over time from 2 to 7 days in the W<sub>30</sub> treatment but not the W<sub>15</sub> treatment.

#### Destructive sampling for denitrification rate

Denitrification rate at the end of the frozen period was 95% greater across the  $W_{30}$  treatments compared to the  $W_{15}$  treatments (Fig. 5). There was a significant interaction between initial WFPS x thaw temperature on denitrification rate (P < 0.001, Table S3). Denitrification rate for the  $T_{8.0}$  treatment increased over the duration of the incubation for the  $W_{30}$  treatment but remained low over the duration

of the incubation for the  $W_{15}$  treatment. For the  $T_{1.5}$  treatment, denitrification rate decreased between day 1 and 3 and remained low thereafter for the  $W_{30}$  treatment but remained low for the entire duration of incubation for the  $W_{15}$  treatment (Fig. 5). Snowmelt did not significantly affect denitrification rate (Table S3).

#### **Cumulative gas emissions**

Soil thaw temperature, initial WFPS and snowmelt all significantly affected cumulative soil respiration (both  $P \le 0.001$ ; Table S4). When averaged across other treatments, cumulative respiration was almost twice as great for  $T_{8.0}$  treatments compared to  $T_{1.5}$  treatments, 14% greater for SM<sup>+</sup> treatments compared to the SM<sup>-</sup> treatments, and 8% greater for  $W_{30}$  treatments compared to  $W_{15}$  treatments (Table 1).

There was a significant initial WFPS x thaw temperature x snowmelt interaction on cumulative N<sub>2</sub>O emissions (P < 0.001; Table S4). Cumulative N<sub>2</sub>O emissions were low for the W<sub>15</sub> treatments regardless of thaw temperature and snowmelt treatments, whereas for the W<sub>30</sub> treatment, cumulative N<sub>2</sub>O emissions were over two-fold greater for T<sub>8.0</sub> treatments compared to T<sub>1.5</sub> treatments, but 17% lower for SM<sup>+</sup> treatments compared to SM<sup>-</sup> treatments (Table 1).

There was also a significant initial WFPS x thaw temperature x snowmelt interaction on cumulative denitrification emissions (P < 0.001; Table S4). Cumulative denitrification





**Fig. 4** Soil respiration rate (CO<sub>2</sub> mg C kg<sup>-1</sup> dry soil day<sup>-1</sup>) and nitrous oxide emission rate (N<sub>2</sub>O  $\mu$ g N kg<sup>-1</sup> dry soil day<sup>-1</sup>) throughout the seven-day incubation of a single-freeze thaw event. Soils were adjusted to an initial water filled pore space (WFPS) of either 15% or

30% (W<sub>15</sub> or W<sub>30</sub>) and incubated at -1.0 °C for four-weeks. Soils were then incubated for a seven-day period at either 1.5 or 8.0 °C and were amended with or without water to simulate snowmelt infiltration (SM<sup>+</sup> or SM<sup>-</sup>). Values represent the mean (n=4) and the standard error.



**Fig. 5** Denitrification rate (N<sub>2</sub>O  $\mu$ g N kg<sup>-1</sup> dry soil day<sup>-1</sup>) using the acetylene inhibition method throughout the seven-day incubation of a single-freeze thaw event. Soils were adjusted to an initial water filled pore space (WFPS) of either 15% or 30% (W<sub>15</sub> or W<sub>30</sub>) and incubated at -1.0 °C for four-weeks. Soils were sampled at day 0 i.e. at the end of the freezing period and immediately before the treatments were imposed. Soils were then incubated for a seven-day period at either 1.5 or 8.0 °C and were amended with or without water to simulate snowmelt infiltration (SM<sup>+</sup> or SM<sup>-</sup>). Values represent the mean (*n*=4) and the standard error.

 Table 1 Cumulative soil respiration, nitrous oxide, and denitrification

 emissions after 7 days of incubation

Treatment	Cumulative emissions			
	Soil respiration (mg CO <sub>2</sub> -C kg <sup>-1</sup> dry soil)	Nitrous oxide ( $\mu$ g N <sub>2</sub> O-N kg <sup>-1</sup> dry soil)	Denitrifica- tion $(\mu g N_2 O-N kg^{-1} dry soil)$	
W <sub>15</sub> T <sub>1.5</sub> SM <sup>-</sup>	57.7±5.18	$6.66 \pm 0.34$	$4.06 \pm 7.79$	
$W_{15}T_{1.5}SM^{+}$	$66.3 \pm 5.44$	$7.73 \pm 0.19$	$4.03 \pm 0.57$	
$W_{30}T_{1.5}SM^{-}$	$68.3 \pm 2.67$	$51.2 \pm 18.0$	$116.8 \pm 16.5$	
$W_{30}T_{1.5}SM^+$	$84.3 \pm 7.53$	$40.1 \pm 9.05$	$67.8 \pm 9.66$	
$W_{15}T_{8.0}SM^{-}$	$126 \pm 2.03$	$6.65 \pm 0.32$	$4.08 \pm 8.13$	
$W_{15}T_{8.0}SM^+$	$143 \pm 5.28$	$7.25 \pm 0.63$	$3.08 \pm 3.21$	
W <sub>30</sub> T <sub>8.0</sub> SM <sup>-</sup>	$129 \pm 6.39$	$117 \pm 18.3$	$469 \pm 91.9$	
$W_{30}T_{80}SM^{+}$	$141 \pm 2.03$	$99.6 \pm 3.93$	$832 \pm 117$	

emissions were minimal for the  $W_{15}$  treatments regardless of thaw temperature or snowmelt treatments, whereas for the  $W_{30}$  treatments, cumulative denitrification was about 7 times greater for  $T_{8.0}$  treatments compared  $T_{1.5}$  treatments and the presence of snowmelt decreased denitrification only for the  $W_{30}T_{8,0}$  treatments (Table 1).

#### **Denitrifier abundance and transcription**

The abundance of *nirK*, *nirS*, and *nosZ* genes and the ratio of *nosZ* transcript abundance/*nosZ* gene abundance did not statistically vary with initial WFPS at the day 0. There were significant main effects of soil thaw temperature and time on *nirK* abundance (both P < 0.001; Table S5). The abundance of *nirK* genes was significantly greater in the T<sub>8.0</sub> treatments compared to the T<sub>1.5</sub> treatments and increased over time during the incubation (Fig. 6a). In contrast, while *nirS* abundance also increased over time, there were no significant effects of initial WFPS, thaw temperature or snowmelt on *nirS* abundance (P = 0.019; Table S5) (Fig. 6b).

The *nosZ* gene abundance was significantly greater in the  $W_{15}$  treatments compared to the  $W_{30}$  treatments (P < 0.001; Table S6) (Fig. 6c). In addition, there were significant main effects of soil thaw temperature and time on the *nosZ* transcript abundance/*nosZ* gene abundance ratio (both P < 0.01; Fig. 6d; Table S6). The *nosZ* transcript abundance/*nosZ* gene abundance ratio was significantly greater in the  $T_{1.5}$  treatments compared to the  $T_{8.0}$  treatments and decreased over time.

# Discussion

This study investigated the effects of initial WFPS prior to soil freezing, soil thaw temperature, and snowmelt infiltration simulated by water addition on soil respiration, soil  $N_2O$  emissions, denitrification, and denitrifier abundance and expression during a single soil freeze-thaw event. It was hypothesized that the combination of high initial WFPS before soil freezing, high soil thaw temperature, and snowmelt infiltration during soil thawing would concomitantly create soil conditions conducive to increasing  $N_2O$  emissions, denitrification, denitrifier abundance and transcription, and that pre-freezing soil water conditions would have a greater influence of  $N_2O$  production than water added to the soil via snowmelt infiltration during soil thaw.

# Effects of initial WFPS, thaw temperature and snowmelt on soil respiration and soil inorganic N concentrations

In this study, respiration rate and cumulative respiration were most affected by thaw temperature and were almost twice as great when thawed at 8.0°C than when thawed at 1.5°C. Increased soil water content, either through the addition of water to simulate snowmelt or increased initial



**Fig. 6** Soil gene abundance (# of copies  $g^{-1}$  dry soil) of *nirK* (**a**), *nirS* (**b**), and *nosZ* clade I (**c**), and the ratio of *nosZ* clade I cDNA to *nosZ* clade I DNA (**d**) throughout the seven-day incubation of a single-freeze thaw event. Soils were adjusted to an initial water filled pore space (WFPS) of either 15% or 30% (W<sub>15</sub> or W<sub>30</sub>) and incubated at -1.0 °C for four-weeks. Soils were sampled at day 0 i.e. at the end

of the freezing period and immediately before the treatments were imposed. Soils were then incubated for a seven-day period at either 1.5 or 8.0 °C and were amended with or without water to simulate snowmelt infiltration (SM<sup>+</sup> or SM<sup>-</sup>). Values represent the mean (n=4) and the standard error.

WFPS, also resulted in increased cumulative respiration. These findings are similar to previous incubation and fieldbased studies that observed greater CO2 emissions from soils that had a greater soil water content and temperature at the time of soil thawing or experienced snow accumulation during the winter (Ouyang et al. 2015; Brin et al. 2018; Wertz et al. 2018; Badewa et al. 2022). The increased respiration rate observed in the current study during soil thawing was likely primarily a result of increased heterotrophic metabolism in response to soil warming (Sharma et al. 2006; Feng et al. 2007; Henry 2008, 2013; Chantigny et al. 2019), and increased C substrate availability through a high initial WFPS prior to soil freezing and snowmelt infiltration that further stimulated microbial activity (Christensen and Christensen 1991; Fierer and Schimel 2003; Brooks et al. 2011; Congreves et al. 2018).

Soil NH<sub>4</sub>-N concentration was primarily affected by soil thaw temperature and was ~4-fold greater at the end of the incubation when thawed at 8.0°C than at 1.5°C. Similarly, previous studies also found that increased thaw temperature resulted in increased soil NH<sub>4</sub>-N concentration during a freeze-thaw event (Wertz et al. 2016, 2018; Song et al. 2017; Li et al. 2021). Increased NH<sub>4</sub>-N concentration in response to a higher thaw temperature was likely associated with an increased net N mineralization following increased microbial metabolism in response to soil warming (Contosta et al. 2011; Grant 2014; Ruan and Roberston, 2017; Song et al. 2017), which is consistent with the observed increase in microbial activity inferred by soil respiration in response to soil warming.

Treatments that resulted in increased soil water content generally resulted in decreased soil NO3-N concentration in soils, likely due to reduced soil oxygen availability creating more anoxic soil conditions, and thereby increasing the rate of denitrification and subsequently NO<sub>3</sub><sup>-</sup> consumption (Firestone et al. 1980; Brin et al. 2018; Congreves et al. 2018). This is supported by the lower NO<sub>3</sub>-N concentration for the initial WFPS of 30% compared to soil with an initial WFPS of 15%, both at day 0 and throughout the 7 days of incubation. It is also consistent with a reduction in soil NO<sub>3</sub>-N concentration at 8.0 °C compared to 1.5 °C, as the increased soil thaw temperature would be expected to further promote anoxic microsite formation (Henry 2008, 2013; Chantigny et al. 2019). Interestingly, snowmelt infiltration only decreased NO3-N concentration in soils with an initial WFPS of 15%; this may be due to conditions that were already conducive to denitrification in soils with an initial WFPS of 30% and thus were not significantly enhanced by additional water infiltration.

# Effects of initial WFPS, thaw temperature and snowmelt on $N_2O$ and denitrification emissions

Soil N<sub>2</sub>O emission and denitrification rates were generally greater when the soil oxygen supply was more limited at an initial WFPS of 30% and a soil thaw temperature of 8.0 °C, most notably resulting in a~100-fold increase in denitrification rates compared to soils with an initial WFPS of 15% and a soil thaw temperature of 1.5 °C. This was likely related to a combination of interrelated mechanisms that increased substrate availability and anaerobiosis (Fig. 7). This includes the release of substrates through microbial cell lysis, microbial cytoplasmic release, and soil pore connectivity via increased soil moisture (Christensen and Christensen 1991; Fierer and Schimel 2003; Wagner-Riddle, 2017; Congreves et al. 2018); increased microbial metabolism via increased substrate availability and soil warming (Henry 2008, 2013; Chantigny et al. 2019); and increased anoxic microsite formation via increased microbial metabolism and soil moisture (Phillips 2008; Risk et al. 2013; Congreves et al. 2018).

Interestingly, while the addition of water to simulate snowmelt infiltration would also be expected to reduce oxygen supply and thus increase denitrification and N2O emissions, this was not observed in the current study. Instead, it was primarily the antecedent conditions of high soil moisture prior to soil freezing that gave rise to increased N<sub>2</sub>O emissions. This supports our original hypothesis that it is not just soil moisture conditions but also the timing of water additions to the soil over the non-growing season that influences N<sub>2</sub>O fluxes. This is of particular interest when considering the expected increased frequency of freeze-thaw cycles in response to climate warming. For example, if snowmelt infiltration occurred during a thaw event, it may not significantly influence N2O emissions; however, if snowmelt infiltration occurred during a mid-winter thaw event in which the soil re-freezes, then it will further increase soil moisture at the time of soil freezing and carry a significant impact for N<sub>2</sub>O emissions during spring thaw. Previous investigations have also demonstrated that it is specifically the antecedent conditions of high soil moisture prior to soil freezing that led to increased N2O emissions during a thaw event (Banerjee et al. 2016; Brin et al. 2018), even after 165 freeze-thaw cycles (Chen et al. 2021). A legacy effect of high initial WFPS leading to greater N2O emissions during soil thawing may be related to a greater disruption of soil microbes or soil aggregates upon soil freezing in response to volume expansion that occurs during the phase change from liquid water to ice (~9%), subsequently increasing macropore connectivity and nutrient availability for N2O production pathways (Bullock et al. 1988; Christensen and Christensen 1991; Six et al. 2004; Chai et al. 2014; Congreves et al. 2018).



**Fig. 7** Schematic of the interrelated factors influencing the production of nitrous oxide (N<sub>2</sub>O), carbon dioxide (CO<sub>2</sub>), and denitrification rate in soil with an initial soil water-filled pore space (WFPS) of 30% and a soil thaw temperature of 8.0 °C compared to soil with an initial WFPS of 15% and soil thaw temperature of 1.5 °C. Soils with an initial WFPS prior to soil freezing of 15% or 30% were amended with red clover (1000 mg C kg<sup>-1</sup> dry soil) and KNO<sub>3</sub><sup>-</sup> (15 mg N kg<sup>-1</sup> dry soil) and frozen at -1.0 °C for four weeks to simulate winter conditions. The soils were then treated with or without water to simulate the presence (or absence) of snowmelt infiltration and were incubated at 1.5 or 8.0 °C for 7 days. Soil N<sub>2</sub>O, CO<sub>2</sub>, and denitrification rates were generally greater in soils with an initial WFPS of 30% and a soil thaw temperature of 8.0 °C. This was likely related to a combination

of interrelated mechanisms that increased substrate availability and anaerobiosis. This includes the release of substrates through microbial cell lysis, microbial cytoplasmic release, and soil pore connectivity via increased soil moisture, increased microbial metabolism via increased substrate availability and soil warming, and increased anoxic microsite formation via increased microbial metabolism and soil moisture. The addition of water to simulate snowmelt infiltration did not contribute largely to increasing denitrification, N<sub>2</sub>O, and CO<sub>2</sub> emissions, while increased soil thaw temperature had the greatest role on increased CO<sub>2</sub> emissions. However, antecedent conditions of high soil moisture prior to soil freezing was the most important factor in the increased N<sub>2</sub>O and denitrification rates, indicating that it is not just about soil moisture conditions but also about the timing of water additions to the soil over the non-growing season that influences N<sub>2</sub>O fluxes.

The negligible or insignificant effect of snowmelt infiltration on N<sub>2</sub>O production may at first be related to pre-freezing soil conditions that may limit snowmelt infiltration. For example, snowmelt infiltration can occur in the open pores of the frozen soil layer; however, if those pores are plugged with ice from high pre-freezing soil water content, then snowmelt infiltration would be restricted or limited during early stages of soil thawing (Gray et al. 1985; Bayard et al. 2005; He et al. 2015; Congreves et al. 2018). However, this can only be the case for the first 24 h (during soil thawing) and not for the remainder of the incubation. Perhaps the negligible effect of snowmelt infiltration on denitrification rate from 1 to 7 days of incubation was due to soil conditions that were already anoxic such that denitrification would not be greatly enhanced by additional water infiltration, or due to snowmelt water containing negligible nitrate or available C, which may act to dilute substrate availability at anoxic microsites.

The C<sub>2</sub>H<sub>2</sub> blockage method may lead to an underestimation of denitrification due to an oxidation reaction between nitric oxide (NO), an intermediate of the denitrification process, and O2 in the presence of C2H2 (Bollmann and Conrad 1997; McKenney et al. 1996). Although there is a possibility of reduced soil O<sub>2</sub> concentrations following soil thaw and snowmelt, and thus a reduced possibility of NO oxidation, it is noteworthy that low  $O_2$  concentrations have still been shown to increase NO oxidation. (McKenney et al. 1997). Similarly, studies that have added organic C and  $NO_3^{-}$  to soils reported a significant (McKenney et al. 1996) or minimal (Murray and Knowles 2003) NO oxidation by C<sub>2</sub>H<sub>2</sub> under anoxic conditions. However, lower NO oxidation has been observed in soils incubated at lower temperatures (2 to 10°C) compared to soils incubated at 25°C (Dunfield and Knowles 1997). The lower incubation temperatures used in our study might have reduced the oxidation of NO by  $C_2H_2$ ; however, it cannot be ruled out that the denitrification rates were potentially underestimated.

# Effects of initial WFPS, thaw temperature and snowmelt on denitrifier abundance and transcription

Similar to denitrification rate, snowmelt infiltration did not affect denitrifier gene abundance or transcription, supporting the theory that water infiltration did not further stimulate denitrification due to the presence of soil conditions already conducive to denitrification. The *nirK* gene abundance did, however, increase in response to an increase in soil temperature, which is similar to a previous study (Braker et al. 2010), and suggested an increase in microbial metabolism resulting in anoxic soil conditions that were more favorable for denitrification. In contrast, *nirS* gene abundance

did not change in response to the experimental parameters investigated in this study. Our findings are similar to those of Németh et al. (2014) who also observed consistent levels of *nirS* gene and transcript abundance throughout a springthaw event in a field-based study. An increase in *nirK* but not *nirS* gene abundance may indicate that *nirK*-harbouring denitrifiers played a more important role in soil denitrification in our study, which has also been observed in other studies (Yoshida et al. 2009; Kou et al. 2021).

We had hypothesized that greater initial WFPS prior to soil freezing and increased soil thaw temperature would result in a depletion of soil  $O_2$  concentration and increased denitrification substrate availability thus favoring the growth of denitrifiers and inducing denitrification. The opposite result was observed: there was a decrease in *nosZ* gene abundance and transcription in soil with greater initial WFPS prior to soil freezing and increased soil thaw temperature. Our findings contrast those of previous studies, where *nosZ* gene abundance and transcription increased in response to increased soil moisture or soil warming (Brin et al. 2019; Wu et al. 2020).

In our study, decreased nosZ transcription in warmer soils may have resulted in lower levels of N2O reduction to N<sub>2</sub> and thus explain the greater N<sub>2</sub>O emissions observed in soils thawed at 8.0 °C compared to 1.5 °C. However, decreased nosZ transcription in soils with a greater initial WFPS was inconsistent with the lower proportion of denitrification end-products emitted as N2O in soils with an initial WFPS of 30% compared to 15%. This may reflect high levels of nosZ transcriptional activity in the initial WFPS30 treatments within the first 24 h of the study (i.e., before the first sampling time point), subsequently resulting in a pool of active NOS enzymes and thus low levels of detectable nosZ transcription after 24 h of the incubation. The *nosZ* gene primer set used in this study might not have captured as much diversity compared to the new primer set developed by Zhang et al. (2021); however, the same subset of the total nosZ denitrifier communities is measured among treatments, thus the results are still informative in understanding how denitrifier abundance is influenced by soil thawing temperature, snow melt and WFPS. It is also worth noting that N<sub>2</sub>O may have originated from pathways other than denitrification (Gao et al. 2023a, b), potentially explaining the absent relationship between nosZ gene and transcript abundance and increased N2O emissions in our study.

# Conclusions

Our study is the first to investigate the synergistic effect of soil moisture at the time of freezing, soil thaw temperature, and snowmelt infiltration on N2O emissions during a freezethaw event. In our experimental system, a synergistic effect was observed where a greater response of N<sub>2</sub>O emissions to increased thaw temperature occurred at a greater initial WFPS. In addition, we observed that snowmelt infiltration only had a modest to negligible effect on N2O emissions and that changes in denitrifier gene or transcript abundance were not associated with N<sub>2</sub>O emissions or denitrification. As climate change continues to increase annual temperatures and precipitation, it is predicted that winter snow cover and thawing events will become increasingly erratic and discontinuous, leading to increased rain events and reduced snow cover. Our findings unequivocally demonstrate that greater soil moisture at the onset of freezing, caused by amplified rainfall or mid-winter thawing, and higher soil surface thaw temperatures owing to decreased snow cover and elevated atmospheric temperatures, can considerably increase N<sub>2</sub>O production during a freeze-thaw event. The interactive effects of several factors including a range of soil initial WFPS, freezing temperatures, and snowmelt timing and intensity on denitrifying communities, denitrification rate, and N<sub>2</sub>O emissions in soil undergoing multiple freeze-thaw events need to be further investigated to better understand how the warmer winters predicted by climate change might influence the soil N dynamics and the production of N<sub>2</sub>O emissions in the future.

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#### Declarations

Competing interests The authors declare no competing interests.

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