



## Sources of organic ice nucleating particles in soils

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**Abstract.** Soil organic matter (SOM) may be a significant source of atmospheric ice nucleating particles (INPs), especially of those active  $> -15^{\circ}\text{C}$ . However, due to both a lack of investigations and the complexity of the SOM itself, the identities of these INPs remain unknown. To more comprehensively characterize organic INPs we tested locally representative soils in Wyoming and Colorado for total organic INPs, INPs in the heat-labile fraction, ice nucleating (IN) bacteria, IN fungi, IN fulvic and humic acids, IN plant tissue, and ice nucleation by monolayers of aliphatic alcohols. All soils contained  $\approx 10^6$  to  $\approx 5 \times 10^7$  INPs  $\text{g}^{-1}$  dry soil active at  $-10^{\circ}\text{C}$ . Removal of SOM with  $\text{H}_2\text{O}_2$  removed  $\geq 99\%$  of INPs active  $> -18^{\circ}\text{C}$  (the limit of testing), while heating of soil suspensions to  $105^{\circ}\text{C}$  showed that labile INPs increasingly predominated  $> -12^{\circ}\text{C}$  and comprised  $\geq 90\%$  of INPs active  $> -9^{\circ}\text{C}$ . Papain protease, which inactivates IN proteins produced by the fungus *Mortierella alpina*, common in the region's soils, lowered INPs active at  $\geq -11^{\circ}\text{C}$  by  $\geq 75\%$  in two arable soils and in sagebrush shrubland soil. By contrast, lysozyme, which digests bacterial cell walls, only reduced INPs active at  $\geq -7.5$  or  $\geq -6^{\circ}\text{C}$ , depending on the soil. The known IN bacteria were not detected in any soil, using PCR for the *ina* gene that codes for the active protein. We directly isolated and photographed two INPs from soil, using repeated cycles of freeze testing and subdivision of droplets of dilute soil suspensions; they were complex and apparently organic entities. Ice nucleation activity was not

affected by digestion of Proteinase K-susceptible proteins or the removal of entities composed of fulvic and humic acids, sterols, or aliphatic alcohol monolayers. Organic INPs active colder than  $-10$  to  $-12^{\circ}\text{C}$  were resistant to all investigations other than heat, oxidation with  $\text{H}_2\text{O}_2$ , and, for some, digestion with papain. They may originate from decomposing plant material, microbial biomass, and/or the humin component of the SOM. In the case of the latter then they are most likely to be a carbohydrate. Reflecting the diversity of the SOM itself, soil INPs have a range of sources which occur with differing relative abundances.

### 1 Introduction

We have known for more than 4 decades that soils are reservoirs of organic ice nucleating particles (INPs) (Vali, 1968; Schnell and Vali, 1972, 1973, 1976). Decaying leaf litter was found to produce up to  $10^{10}$  INPs per gram active at  $-10^{\circ}\text{C}$ , with an onset of activity at up to  $\approx -1.5^{\circ}\text{C}$  (Schnell and Vali, 1972, 1973). Rich top soils and peat were shown to be ice nucleating (IN) at  $-4^{\circ}\text{C}$  (Vali, 1968). The component responsible for the highest temperatures of freezing, an IN bacterium, was soon isolated (Maki et al., 1974; Vali et al., 1976). However, the proposal (Schnell and Vali, 1972) that particles composed of soil organic matter (SOM) may act as INPs was contested (Rosinski et al., 1974) and then neglected

as a topic for investigation, even though the aerial release of INPs from litters (Schnell and Vali, 1976) suggested that this may be a significant atmospheric source, especially of INPs active  $\geq -10^\circ\text{C}$ . INPs influence cloud microphysical and radiative properties, the release of latent heat, and the triggering of precipitation.

Recently there has been a resurgence of related research, with the general aims of correlating abundances of organic INPs in the boundary layer with local ecotypes (Prenni et al., 2009; Bowers et al., 2010; Garcia et al., 2012; Huffman et al., 2013; Tobo et al., 2013; Mason et al., 2015), gauging the stimulation of their release by humidity, rainfall, and harvesting (Garcia et al., 2012; Huffman et al., 2013; Prenni et al., 2013; Wright et al., 2014; Bigg et al., 2015), and measuring organic INP abundance in clouds and precipitation (Christner et al., 2008; Delort et al., 2010; Joly et al., 2014; Monteil et al., 2014; Morris et al., 2008; Petters and Wright, 2015; Šantl-Temkiv et al., 2015).

By contrast, limited progress has been made in characterizing the nature of these “bio-INPs”. Heating to  $90\text{--}100^\circ\text{C}$  to deactivate IN macromolecules and digestion with  $\text{H}_2\text{O}_2$  to entirely remove the SOM confirmed their dominance of the INP population active  $\approx > -15^\circ\text{C}$  in diverse sieved soils (Conen et al., 2011; O’Sullivan et al., 2014). Further, in dusts generated from Wyoming agricultural soils, treatment with heat ( $300^\circ\text{C}$ ) and  $\text{H}_2\text{O}_2$  digestion revealed that organic-rich INPs predominated at all tested temperatures  $> -36^\circ\text{C}$  (Tobo et al., 2014). Recent investigations into specific sources of organic INPs found that the soil-inhabiting fungus *Mortierella alpina*, which is common in many of the soils studied here, is an ice nucleator: all strains tested initiated freezing at  $-5$  to  $-6^\circ\text{C}$  and released large numbers of cell-free INPs (Fröhlich-Nowoisky et al., 2015). Of potential widespread relevance, too, is the recent discovery of ice nucleation by cellulose (Hiranuma et al., 2015a): activity was detected starting at  $-16^\circ\text{C}$ .

To more comprehensively assess the identities of organic INPs in soils, we tested soils representative of local ecotypes in Wyoming and Colorado for the presence of INPs from a wide range of potential sources. We tested for total organic INPs, INPs in the heat-labile fraction, IN bacteria (using PCR of the *ina* gene that codes for the Ina protein; Warren, 1995), IN *M. alpina* (by digestion with papain protease), IN fulvic and humic acids (via acid and alkaline extractions), ice nucleating entities comprised of crystals of sterols or monolayers of aliphatic alcohols (by removal with chloroform), and IN plant tissue. We also demonstrated that cycles of freeze-testing and subdivision of droplets of dilute soil suspensions can be used to directly isolate individual INPs from soil for subsequent analyses.

## 2 Materials and methods

### 2.1 Soil sampling

Topsoils were sampled from beneath representative natural ecotypes and land uses, mostly in Wyoming. These included conventionally and organically cropped land and pasture at the University of Wyoming’s Sustainable Agricultural Research and Extension Center (SAREC), near Lingle, Wyoming, sagebrush shrubland, native glacial scree grassland and lodgepole pine forest in south-east Wyoming, and ponderosa pine forest at the Manitou Experimental Forest Observatory (MEFO) (Ortega et al., 2014) in Colorado. Details of sites and vegetation are given in Table 1. At each site, three replicate soil samples were obtained. Each was obtained from a separate  $10 \times 10\text{ m}$  area, and within each area three cores, each 5 cm deep and  $\approx 10\text{ cm}$  in diameter, were retrieved and mixed together on site. Samples were thoroughly mixed, any stones (no soils had smaller, gravel-sized pieces) and roots were removed, and then the samples were stored at  $4^\circ\text{C}$  before testing. Sub-samples were frozen in airtight 15 mL screw-cap tubes at  $-20^\circ\text{C}$  for DNA extractions and papain digestions.

Dry weights were obtained by heating a sub-sample of each soil to  $105^\circ\text{C}$  for 24 h. For analysis of soil organic carbon and total nitrogen only, sub-samples of soils with a  $\text{pH} > 7$  were initially immersed in 0.1 M HCl overnight to remove carbonates and then filtered (no. 42 paper, Whatman) and dried as above. Samples were ground and sieved through a 0.5 mm sieve and analysed using a CHN analyser.

### 2.2 Measurement of INPs

Concentrations of INPs in soil suspensions were derived using the immersion freezing method. Samples were made initially into a 10-fold slurry with INP-free diluent, mixed on an orbital shaker at 200 rpm for 20 min and then a series of 10-fold dilutions constructed. INP estimates were made using 32 aliquots of  $50\text{--}80\ \mu\text{L}$  of suspension dispensed into sterile 96-well polypropylene PCR trays (Life Science Products Inc.). Full details of the immersion freezing protocol are given in Garcia et al. (2012) but are summarized here. INP numbers were estimated by counting the number of frozen aliquots in the PCR tray as it was cooled in a thermal cycler (PTC-200, MJ Research). The cycler was programmed to descend in  $0.5$  or  $1^\circ\text{C}$  increments with a 5 min dwell time at each temperature. Once at  $-9^\circ\text{C}$  (the limit of this device) the tray was transferred to a 96-well aluminium incubation block (VWR) cooled to  $\approx -12^\circ\text{C}$  inside a foam box in a freezer. A thermistor was inserted into a side well and after 10 min of further incubation in the freezer the block temperature and number of frozen wells was recorded. The tray was then transferred to two further blocks at sequentially colder temperatures (down to  $\approx -20^\circ\text{C}$ ). Holding times at each temperature were chosen based on observations of the time required for essentially

Table 1. Descriptions of sites sampled.

Site	Sampling date(s)	Lat.	Long.	Vegetation	Soil type	C <sub>org</sub> (%)	N (%)	pH
Agricultural ecotypes								
Grass/alfalfa fallow <sup>a</sup>	2 Mar 2011 and 17 May 2011	42.12266	-104.38585	Dead material on surface from previous year's sowing of alfalfa, orchard grass, and meadow brome. Organically managed field.	Haverson and McCook light brownish-grey loams. <sup>b</sup>	0.55	0.076	8.1
Sugar beet <sup>a</sup>	2 Mar 2011 and 17 May 2011	42.12878	-104.39516	Bare at sampling. Previous year was Roundup Ready sugar beet.	Haverson and McCook light brownish-grey loams. <sup>b</sup>	0.75	0.11	8.15
Pasture <sup>a</sup>	2 Mar 2011 and 17 May 2011	42.13243	-104.39428	Roadside, low-input pasture of smooth and downy bromes.	Haverson and McCook light brownish-grey loams. <sup>b</sup>	2.7	0.46	7.85
Native ecotypes								
Native grass-land	24 May 2011	41.2881	-106.11124	Bluebunch wheatgrass, Idaho fescue, western wheatgrass, and three-tip sagebrush.	Greyback very cobbly sandy loam; outwash from alluvial fan. Surface layer greyish brown to brown very cobbly sandy loam. <sup>c</sup>	2.2	0.27	6.45
Sagebrush shrubland	2 Dec 2012 and 17 Feb 2013	41.03159	-106.00170	Semi-arid rolling Wyoming big sagebrush shrubland.	Fine, loose sand with some surface cobbles.	1.5 <sup>e</sup>	-	7.15
Lodgepole pine forest	24 May 2011 and 2 Dec 2012	41.32439	-106.16062	Lodgepole pine, with understory of elk sedge, low sedge, creeping juniper, Oregon grape, kinnikinnick, woods rose, and heartleaf arnica.	Ansile-Granite gravelly sandy loam. 5 cm layer of needles and bark residue. <sup>c</sup>	31 <sup>e</sup>	2.0	5.9
Ponderosa pine forest	22 Jul 2012	39.10330	-105.10395	Dominated by ponderosa pine, with sparse grasses and sedge ground cover.	Gravelly alluvium, classified as loamy, mixed Eutroboralfs or Aridic Haploborolls. <sup>d</sup>	11.6	0.6	5.05

<sup>a</sup> At the University of Wyoming's Sustainable Agricultural Research and Extension Center (SAREC) near Lingle, Wyoming. <sup>b</sup> United States Department of Agriculture (1971). <sup>c</sup> United States Department of Agriculture (1998). <sup>d</sup> Moore (1992). <sup>e</sup> Soil organic carbon obtained by loss-on-ignition at 550 °C for 3 h and then multiplying values by 0.58 (De Vos et al., 2005).

full expression of ice nucleation activity, and they may produce slightly higher concentrations than methods using faster ramp rates.

Results were adjusted for INPs in the diluent (INP-free until  $< -15^{\circ}\text{C}$ ). The effect of papain digestion was assessed using CSU's ice spectrometer, a method that eliminates the transfer procedures used in the above method (details of the method are given in Hiranuma et al., 2015b). Numbers of INPs in soil suspensions were estimated using Eq. (1):

$$\text{INPs} [\text{mL}^{-1}] = \frac{-\ln(f)}{V}, \quad (1)$$

where  $f$  is the proportion of droplets not frozen and  $V$  is the volume of each aliquot (Vali, 1971), and INPs per mL of soil suspension were converted to INPs  $\text{g}^{-1}$  of dry soil. Confidence intervals (95 %) were derived using the formula (no. 2) recommended by Agresti and Coull (1998).

### 2.3 Removal of SOM

Removal of SOM was achieved by digestion with hydrogen peroxide. Fifty millilitres of 15 %  $\text{H}_2\text{O}_2$  were added to 2.5–10 g fresh soil; the slurry was steeped until initial foaming subsided and then boiled while topping up with  $\text{H}_2\text{O}_2$  as required until all SOM and then all residual  $\text{H}_2\text{O}_2$  were decomposed (but see Mikutta et al., 2005, for a review of the inefficiencies inherent in the use of oxidizers to remove SOM from soils). Suspensions were left overnight to settle, the pH restored if necessary (drop-wise addition of 1 M NaOH to the pasture soil and 1 M HCl to the kaolinite), and the volume made up to a 10-fold dilution with deionized water. Ten-fold serial dilutions were then made using 0.45  $\mu\text{m}$  pore diameter filtered 10 mM sodium phosphate buffer (pH 7) or deionized water as the diluents (comparison tests showed no difference).

For the sugar beet and organic grass/alfalfa fallow soils and kaolinite, a comparison of INP concentrations was made for the untreated and  $\text{H}_2\text{O}_2$  digested samples with comparably treated aerosolized soil dusts (0.6  $\mu\text{m}$  diameter) measured using a continuous flow diffusion chamber (Tobo et al., 2014). INP fraction data presented by Tobo et al. (2014) were converted to per gram values, assuming the particles were spherical and had a density of  $2.6 \text{ g cm}^{-3}$ .

### 2.4 Removal of humics

Testing for INPs of humic origin was performed on the SAREC pasture soil since it contained  $\approx 5\%$  SOM and a commensurately high level of INPs. Fulvic and humic acids were removed using the International Humic Substance Society method as described in Swift (1996). Briefly, 1 M HCl and deionized water were added to a sample until a 10 : 1 liquid : dry-soil weight ratio until a pH of  $\approx 1.5$  was obtained. The suspension was shaken for 1 h, settled, and the lightly coloured supernatant decanted off. The sediment was neutralized with 1 M NaOH and then 0.1 M NaOH was added

until a 10 : 1 liquid : dry-soil weight ratio was reached. This was shaken intermittently for 4 h, settled overnight, and the now dark brown supernatant containing the humic acids removed by decanting. Suspensions of the residual soil were then made in 0.45  $\mu\text{m}$  pore diameter filtered 10 mM sodium phosphate buffer at a pH value approximating the soil sample and tested for INP content.

### 2.5 Removal of sterols and aliphatics

Two contrasting soils, the organic horizon from the lodgepole pine forest and the sagebrush shrubland soil, were tested for ice nucleating entities sourced from monolayers of alcohols, sterol crystals, and other chloroform-soluble organics. This was assessed by dissolving the organics and testing for any decrease in ice nucleation activity. Soils were air-dried and then  $\approx 5 \text{ g}$  added to 30 mL chloroform; the mix was shaken and then steeped for 3 days before removing the chloroform by vacuum filtration using a glass filter holder fitted with a 0.45  $\mu\text{m}$  pore diameter cellulose filter (Gelman). Samples were then rinsed three times with fresh chloroform and dried overnight. Suspensions of the soils were made in buffer (10 mM sodium phosphate buffer at pH values approximating the original soil sample and filtered through a 0.1  $\mu\text{m}$  pore diameter syringe filter (Nalgene, Thermo Scientific) and tested for INP content.

### 2.6 Heat treatments

Ice nucleating particles produced by IN bacteria and fungi (probably including lichen symbionts) are typically proteins (see Pummer et al., 2015). Protein function is controlled by secondary and tertiary structures and in IN bacteria also by the formation of complexes (e.g. Kozloff et al., 1991; Schmid et al., 1997; Garnham et al., 2011). Heat is a simple but effective tool to disrupt and denature these proteinaceous INP. Two temperatures, 60 and 105  $^{\circ}\text{C}$ , were used.

Sixty degrees Celsius is a moderate heat that will fragment molecular complexes and denature heat-labile proteins. Accordingly, in tests on diverse IN bacteria isolated from crops and grasses, incubation at 60  $^{\circ}\text{C}$  for 20 min degraded INPs from class A aggregates nucleating  $\geq -3^{\circ}\text{C}$  to class C oligomers typically freezing at  $-8$  to  $-9^{\circ}\text{C}$  (Turner et al., 1990; Hill et al., 2014), while 60  $^{\circ}\text{C}$  for 60 min deactivated the INPs of the IN fungi *Acremonium implicatum* and *Isaria farinosa* (Pummer et al., 2015) isolated from boundary layer air in Colorado (Huffman et al., 2013). INPs produced by bacteria and fungi are, however, often unaffected by this temperature. This includes leaf-derived ice nuclei (Schnell and Vali, 1973), most strains of IN *M. alpina* (Fröhlich-Nowoisky et al., 2015), IN lichen (Kieft and Ruscetti, 1990), and IN *Fusarium* spp. (Pouleur et al., 1992), although the onset temperature of activity of cell-free INPs of *F. avenaceum* was lowered when heated  $> 40^{\circ}\text{C}$  (Hasegawa et al., 1994). It should also be noted that immobilization of proteins onto

clays or tannins can significantly affect their sensitivity (either increase or decrease) to denaturation by 60 °C heat (Rao et al., 2000). To test the sensitivity of soil INPs to moderate heat, PCR trays were thawed after initial INP testing and then heated in the thermal cycler to 60 °C for 20 min and retested.

To test the effect of a more denaturing heat, 1.8 mL of a 100-fold dilution of soil suspension was aliquoted into a 2 mL screw-cap microcentrifuge tube and immersed in mineral oil at 105 °C for 20 min (plus 7 min for equilibration). Serial dilutions were made with 0.45 µm pore size filtered 10 mM sodium phosphate buffer (at pH values approximating the original soil sample).

## 2.7 Enzymatic digestions

The enzyme lysozyme lyses all bacteria by digesting the cell wall polymer peptidoglycan. It also hydrolyses fungal chitin oligosaccharides but not the chitin polymer itself. Gram-negative bacteria such as the IN bacteria are less susceptible to lysozyme digestion of their cell walls than Gram-positives because their outer membrane acts as a barrier and because their cell walls have less peptidoglycan. Their efficient lysis requires EDTA which, by compromising the integrity of the outer membrane, assists the passage of lysozyme. Tests on pure cultures of diverse IN bacterial isolates grown in nutrient broth, with cell densities measured using absorbance at 600 nm, confirmed that addition of EDTA was required (optimal at 5 mM) for efficient lysis with lysozyme at 4 mg mL<sup>-1</sup> (Pooley and Brown, 1990), a concentration comparable to that used by Christner et al. (2008). A series of soil suspensions ranging in dilution from 10<sup>-2</sup> to 10<sup>-7</sup> were constructed in 0.45 µm pore diameter filtered 10 mM Tris (pH 8), 5 mM EDTA (pH 8), and 4 mg mL<sup>-1</sup> lysozyme (L7651, Sigma-Aldrich) and incubated at 16 °C for 24 h. The control comprised soil suspensions in Tris and EDTA incubated under the same conditions.

To test soils for the presence of INPs susceptible to digestion by Proteinase K, such as those possessed by IN *Fusarium* species (Hasegawa et al., 1994; Humphreys et al., 2001), suspensions of the agricultural soils were digested with 5 U mL<sup>-1</sup> Proteinase K (Qiagen) using the same buffer, incubation conditions and negative controls as used for lysozyme (also optimal for this enzyme). Soil suspensions (1 in 20 dilution) were also digested with papain protease (AppliChem) at concentrations of 10 or 20 mg mL<sup>-1</sup> in 10 mM sodium phosphate buffer (pH 6.3) at 60 °C for 20 min. Controls were again processed in the same manner except without the enzyme. The INP-free buffer was used for serial dilutions for INP determinations.

Adsorption of enzymes onto the surfaces of minerals, with consequent deactivation of their IN sites, as observed by Zolles et al. (2015), does not appear to have been a factor in the soils tested; lysozyme and Proteinase K had no effect upon INP counts below their expected impacts at relatively warm temperatures.

## 2.8 PCR of the *ina* gene in IN bacteria

For DNA extraction, 2 g of soil was mixed into a thick slurry with deionized water (18.2 MΩ and 0.2 µm filtered) and 50 µL transferred to the DNA extraction tube. For the lodgepole pine forest litter, 2 g was mixed with 40 mL deionized water, shaken for 20 min at 250 rpm, then 0.5 mL transferred to a 1.5 mL microfuge tube, spun at 22 500 g for 5 min, and had all but 50 µL of the supernatant removed. This was then transferred to the DNA extraction tube using solution MD1 of the DNA extraction kit. DNA extraction was performed using the recommended protocol of the PowerLyzer™ UltraClean® Microbial DNA Isolation Kit (MO BIO Laboratories Inc.), with homogenization on a FastPrep® bead beater (BIO 101 Inc.) at setting 4 for 5 min.

Quantitative PCR of the *ina* gene was initially attempted using the method described in Hill et al. (2014). However, a lack of detectable amplicons in any samples combined with the high diversity of soil genomic DNA led to significant mispriming, which limited the method's sensitivity. We therefore spiked standard *ina* gene PCRs with a range of known *ina* gene copies and used detection of a discernible amplicon on gels to gauge the approximate limits of detection.

Each PCR contained 1 U of GoTaq® Hot Start Polymerase (Promega) in 1 × Colorless GoTaq® Flexi Buffer (no Mg in buffer), 0.9 µM forward primer 3308f (5 GGCGAT-MGVAGCAAactsac 3), 0.9 µM reverse primer 3462r1 (5 TGTAVCKTTTSCCGTCCcag 3), 0.2 mM each dNTP, 1.25 mM MgCl<sub>2</sub>, 4 % DMSO, 2 µL of DNA extract, 0.5 µL *ina* gene spike, and deionized water to a total volume of 25 µL. Spikes ranged from 60 to 1185 *ina* gene copies using genomic DNA extracted from a pure culture of *P. syringae* Cit7 (see Hill et al., 2014, for details of the isolate and culturing and preparation of standards).

Cycling conditions were an initial denaturation at 95 °C for 2 min, followed by 40 cycles of 94 °C for 15 s and 54 °C for 25 s (combined primer annealing and extension). Amplification was performed on a Bio-Rad DNAEngine® (Hercules). After amplification, products (194 bp) were electrophoresed in 1.5 % MetaPhor® agarose gels (Cambrex) in 1 × sodium borate buffer at 200 V for 35 min, using ethidium bromide for visualization. A 50 bp ladder (G4521, Promega) was used for sizing. Presence of an unambiguous amplicon on gels was used to estimate the limit of detection of *ina* genes for each soil's DNA and converted to theoretical upper limits of IN bacteria per gram dry soil (there is one copy of the *ina* gene per bacterium).

## 2.9 Direct isolation of INPs from soil

Freezing triggered by an INP can be exploited as a means to isolate it. Initially, a 10<sup>-5</sup> dilution of sagebrush soil was prepared in 0.2 µm pore diameter filtered deionized water. Then, 50 µL aliquots were dispensed into a 96-well PCR tray and the tray cooled to -7.0 °C. Frozen wells were noted and the

plate thawed. The contents of a well that froze was transferred to a sterile microplate lid (Nunc, Thermo Scientific) and divided into an array of 1  $\mu\text{L}$  droplets. The plate was covered, placed on a cold block at  $\approx -15^\circ\text{C}$  for 10 min, uncovered, and, using a stereo microscope, the single frozen droplet was noted. The plate was warmed, 2–4  $\mu\text{L}$  of deionized water added to the frozen droplet, and the droplet then divided into an array of 0.1 or 0.2  $\mu\text{L}$  droplets for freeze testing. This cycle was repeated 2–3 further times. The lid was then transferred to a microscope, the droplet containing the INP observed as it evaporated, and the residual photographed.

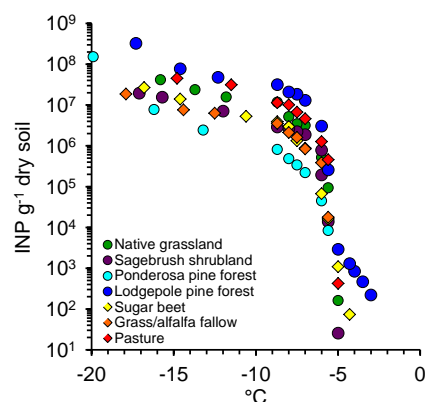
## 2.10 Ice nucleation by sagebrush tissues

The semi-arid sagebrush shrubland soil had a particularly pronounced onset of ice nucleation activity, increasing from  $\approx 30$  to almost a million INPs  $\text{g}^{-1}$  soil between  $-5$  and  $-6^\circ\text{C}$ . This suggests a single class of INP was responsible. To test whether sagebrush tissues were the INPs responsible, shoots (leaves and small stems) and roots up to 1 mm diameter were collected at the sagebrush shrubland site in February 2013 and tested.

Foliage and roots were washed to remove loose soil, placed in fresh plastic bags filled with deionized water and sonicated for several minutes to further dislodge soil particles. The water was replaced and the process repeated five times for a total of 30 min of sonication. A subsample of each tissue type was then ground under liquid nitrogen in a pestle and mortar (cleaned by soaking in 5 %  $\text{H}_2\text{O}_2$  for several hours followed by repeated rinses in deionized water). The ground root powder was examined under a microscope and contained root cell fragments plus bacteria and occasional pieces of fungal hyphae. Serial dilutions in deionized water were tested for INP content. A sample of the initial 50-fold dilution was re-tested after heating to  $105^\circ\text{C}$  for 15 min to determine INP heat sensitivity.

## 3 Results and discussion

All soils presented contained abundant INPs active at warm temperatures (Fig. 1). While the onset of nucleation ranged over  $2.5^\circ\text{C}$ , all soils demonstrated a pronounced increase below  $-5^\circ\text{C}$ , while below approximately  $-8^\circ\text{C}$  the rate of increase lessened to become progressively log-linear. INP concentrations at  $-10^\circ\text{C}$  ranged from  $\approx 10^6$  to  $\approx 5 \times 10^7$   $\text{g}^{-1}$  dry soil, with levels in the native grassland and agricultural soils comparable to those found in grassland and agricultural topsoils in Germany, Hungary, and England (Conen et al., 2011; O'Sullivan et al., 2014, soils A, C, and D). INPs in the two pine forest litters were lower than reported for comparable USA High Plains litters by Schnell and Vali (1976); they noted they were similar to values of  $\approx 10^9$  INPs  $\text{g}^{-1}$  fresh litter found in D-type vegetation. However, this High Plains



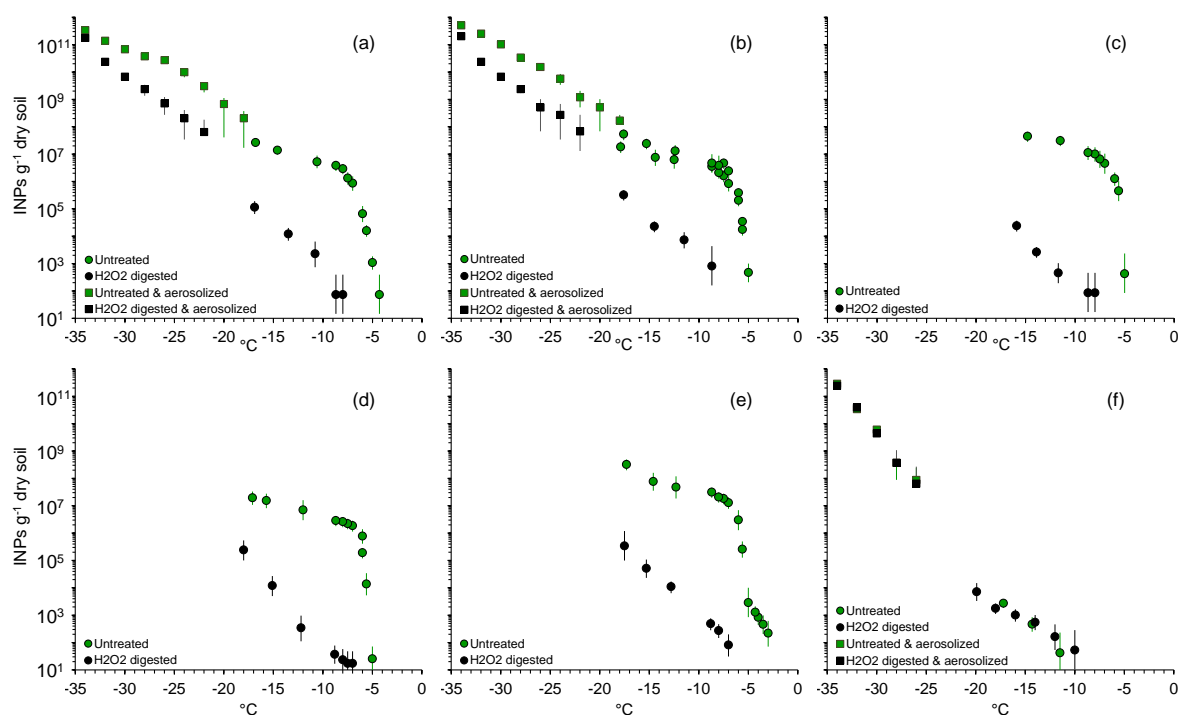
**Figure 1.** Ice nucleating particle concentrations in four natural ecotype and three agricultural soils.

litter may also have included deciduous plant litter (Russell Schnell, personal communication, 2016). Unexpectedly, the lowest concentrations were found in ponderosa pine forest litter, even though the organic matter content of this sample was high ( $\approx 20\%$ , Table 1). INP production by soils and litters is thus not only a function of climatic region (Schnell and Vali, 1976) and SOM content (Conen et al., 2011) but also dependent upon the resource quality of the SOM supplied by the vegetation, among other factors.

## 3.1 INPs in the total soil organic matter

Reductions in INPs caused by SOM removal with  $\text{H}_2\text{O}_2$  (Fig. 2) were similar to or greater than those found by Conen et al. (2011) and greater than those found by O'Sullivan et al. (2014), who used a gentler peroxide treatment. At the lower temperature limit ( $-15$  to  $-18^\circ\text{C}$ ), INPs were reduced by 2–3 orders of magnitude, comparable to the findings of Conen et al. (2011; soils B–D), but more than a 10-fold greater impact than observed by O'Sullivan et al. (2014). By contrast, no effect of  $\text{H}_2\text{O}_2$  digestion was observed for kaolinite, in agreement with the results of Tobo et al. (2014) and as also found for the minerals montmorillonite and K-feldspar (Conen et al., 2011; O'Sullivan et al., 2014).

There was reasonable agreement between results obtained here, using immersion freezing on bulk soil suspensions, with INP spectra below  $-18^\circ\text{C}$  obtained from aerosolized soil dusts (0.6  $\mu\text{m}$  diameter) of the sugar beet and organic alfalfa fallow soils, and measured using the continuous flow diffusion chamber (Tobo et al., 2014) (Fig. 2). The closeness of the correspondence of spectra in agricultural soils may be partly fortuitous (due to several biases cancelling out each other's effects), since it is unlikely that 0.6  $\mu\text{m}$  diameter particles will possess the same INP-active fraction as whole soil. Of greater importance are the large reductions in INPs following  $\text{H}_2\text{O}_2$  digestions in both sets of tests, suggesting that organic forms would also comprise the bulk of INPs active



**Figure 2.** Effect of removal of SOM by  $\text{H}_2\text{O}_2$  digestion upon INP concentrations of soils. Soils are (a) sugar beet, (b) organic grass/alfalfa fallow, (c) pasture, (d) sagebrush shrubland, (e) lodgepole pine forest, and (f) pure kaolinite. The sugar beet, organic grass/alfalfa, and kaolinite figures include data of aerosolized soil dusts ( $0.6\ \mu\text{m}$  diameter) from Tobo et al. (2014) measured using a continuous flow diffusion chamber.

over a wide range of temperatures in dusts raised from these agricultural soils.

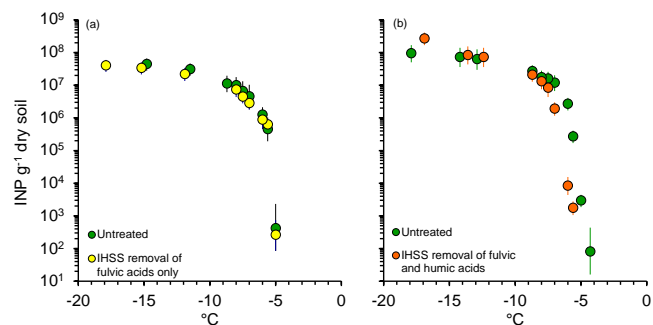
### 3.2 INPs in fulvic and humic acids

Soil organic matter comprises a spectrum of forms ranging from fresh plant inputs to ancient, refractory organic matter. It is often divided into the broad classes of decomposable plant material, resistant plant material, microbial biomass, humus, and inert organic matter (Coleman and Jenkinson, 1999). The humus fraction, a complex mixture of biologically transformed plant and microbial debris, usually predominates and has itself been classically sub-divided into three groups: fulvic acids, humic acids, and humin. Fulvic acids are the fraction that dissolve in  $0.1\ \text{M HCl}$ , humic acids the fraction that subsequently solubilize in  $0.1\ \text{M NaOH}$ , and humin the residue (Powlson et al., 2013). These three groups delineate broad and “operationally defined” – as opposed to “ecologically defined” – pools of humic substances (Wander, 2004).

Humic and fulvic acids are of particular interest because of their apparent similarity to atmospheric HULIS (HUMic-Like Substances), the often significant constituent in water-soluble organic carbon extracted from aerosols (Graber and Rudich, 2006; Wang and Knopf, 2011). However, Graber and Rudich (2006) concluded that HULIS are both smaller

( $\text{MW} < 1000$ ) and less aromatic than terrestrial humic substances and more likely to be the product of the oligomerization of smaller molecules while airborne or the breakdown and reassembly of fulvic and humic acids lofted from soil surfaces. HULIS thus appears to be similar to, but not the same as, humus. Studies using commercially available fulvic and humic acids as surrogates for HULIS found they possessed unremarkable ice nucleation activities (Kanji et al., 2008; O’Sullivan et al., 2014; Wang and Knopf, 2011). O’Sullivan et al. (2014) recorded the warmest activity with sieved suspensions: onset of activity for Suwannee River fulvic acid was  $\approx -14\ ^\circ\text{C}$  and for Leonardite humic acid it was  $\approx -19\ ^\circ\text{C}$ . Fornea et al. (2009) measured onset of freezing temperatures of  $-10.5\ ^\circ\text{C}$  for Pahokee peat, but they used individual particles of the intact soil. It should be noted that the standard extraction method used would also solubilize and remove most atmospheric HULIS forms from the SOM (Feczko et al., 2007).

In the pasture soil, removal of fulvic acids had no apparent impact upon INPs, while removal of humic acids only reduced INPs active  $\geq -7\ ^\circ\text{C}$  (Fig. 3), and these may have been biological macromolecular INPs denatured by the pH 13 extractant. The small molecular weights and the diversity of forms of fulvic and humic acids make it improbable that they possess significant ice nucleation activity. By contrast, the residual humin is compromised in large part of



**Figure 3.** Effect of removal of fulvic (a) and both fulvic and humic (b) acids upon the INP content of the pasture soil.

macromolecules or aggregates of peptides, aliphatics, carbohydrates, lignin, and the bacterial cell wall polymer peptidoglycan (Simpson et al., 2007) and hence is a more likely source of IN molecules.

### 3.3 INPs from sterols and monolayers of aliphatic alcohols

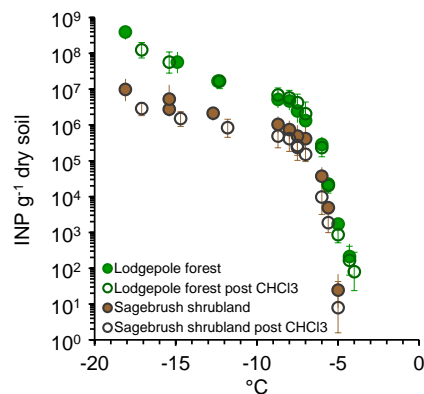
Several classes of naturally occurring organic compounds are effective INPs (Bashkirova and Krasikov, 1957; Head, 1961, 1962; Fukuta, 1966). Crystals of some sterols, for example, trigger nucleation  $> -5^{\circ}\text{C}$ , although the fungal sterol ergosterol and the plant sterol  $\beta$ -sitosterol were found to be inactive (Head, 1962; Fukuta and Mason, 1963).

Under optimal conditions, monolayers of aliphatic alcohols can also induce ice nucleation at temperatures ranging from  $\approx -14^{\circ}\text{C}$  for  $\text{C}_{14}\text{H}_{29}\text{OH}$  to just below  $-1^{\circ}\text{C}$  for  $\text{C}_{31}\text{H}_{63}\text{OH}$  (Gavish et al., 1990; Popovitz-Biro et al., 1994; see also Rosinski, 1980). Nucleation is facilitated by alcohols self-assembling into two-dimensional crystalline clusters on air: liquid or liquid: liquid interfaces, with their hydroxyl groups embedded into the water surface and arrayed with a spacing closely matching hexagonal ice (Gavish et al., 1990; Popovitz-Biro et al., 1991). In the two soils tested here, the removal of organics with chloroform had no significant effect upon the INP spectra (Fig. 4), suggesting that IN organic crystals or long-chain alcohols were not present or did not occur in sufficient quantities or the required purity.

### 3.4 Heat-labile INPs

As shown in Fig. 5, the  $60^{\circ}\text{C}$  treatment had essentially no impact upon the soils' INP activation temperature spectra, apart from some reduction  $\geq -6^{\circ}\text{C}$  in the sugar beet and lodgepole pine soils. This general insensitivity suggests that labile IN proteins or protein complexes from IN bacteria or other microbial or plant sources were not present, or not significant contributors, of high-temperature INPs to the soils.

Heating suspensions of IN bacteria, isolated from crops at the agricultural research centre, to  $105^{\circ}\text{C}$  denatured and deactivated their Ina proteins (Hill et al., 2014). Likewise,



**Figure 4.** Effect of removal of organics, such as sterols and aliphatic alcohols, with chloroform upon INP temperature spectra of forest and shrubland soils.

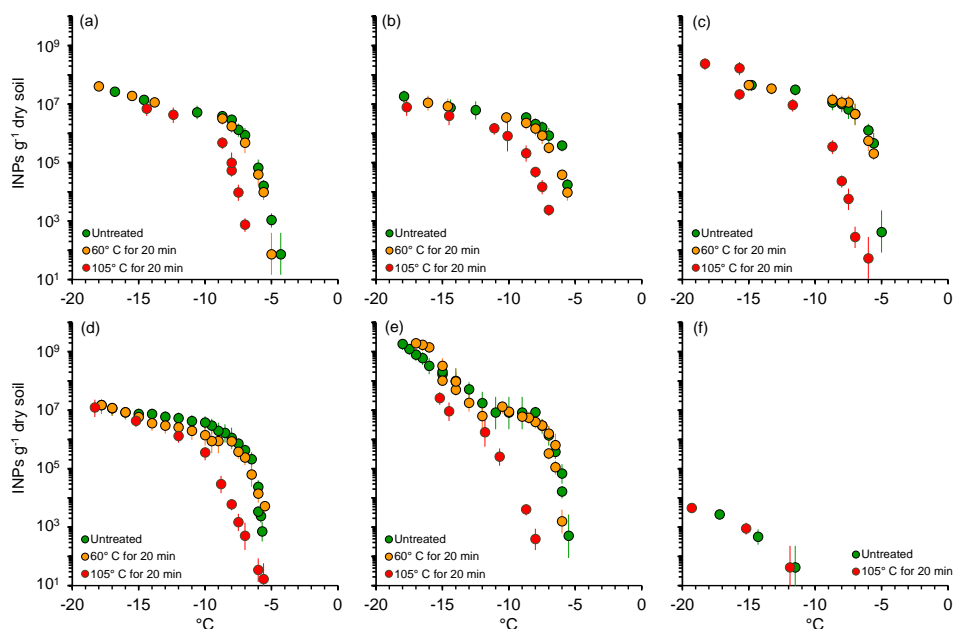
$98^{\circ}\text{C}$  treatment almost eliminated the ice nucleation activity of all but a couple of local isolates of IN *M. alpina* (Fröhlich-Nowoisky et al., 2015), while  $90^{\circ}\text{C}$  heating lowered the median activity of IN *F. avenaceum* by  $10^{\circ}\text{C}$ . Correspondingly, the  $105^{\circ}\text{C}$  treatment reduced the INP concentration active at  $-9^{\circ}\text{C}$  in all soils by  $\geq 10$ -fold. In the lodgepole pine litter this reduction extended to at least  $-15^{\circ}\text{C}$ . Interestingly, for boundary layer aerosols collected above crops in northern Colorado (Garcia et al., 2012), the reduction in INPs following  $98^{\circ}\text{C}$  heating was similar to that found in these soils but, like with the lodgepole pine litter, extended further, to  $\approx -18^{\circ}\text{C}$  (the limit of the measures in that study). In three out of the four soils tested by Conen et al. (2011) heating to  $100^{\circ}\text{C}$  reduced INPs active at  $-12^{\circ}\text{C}$  by 70–98 % while in three of the four agricultural soils tested by O'Sullivan et al. (2014) a  $90^{\circ}\text{C}$  treatment reduced INPs active at  $-10$  to  $-12^{\circ}\text{C}$  by around 10-fold.

### 3.5 IN bacteria

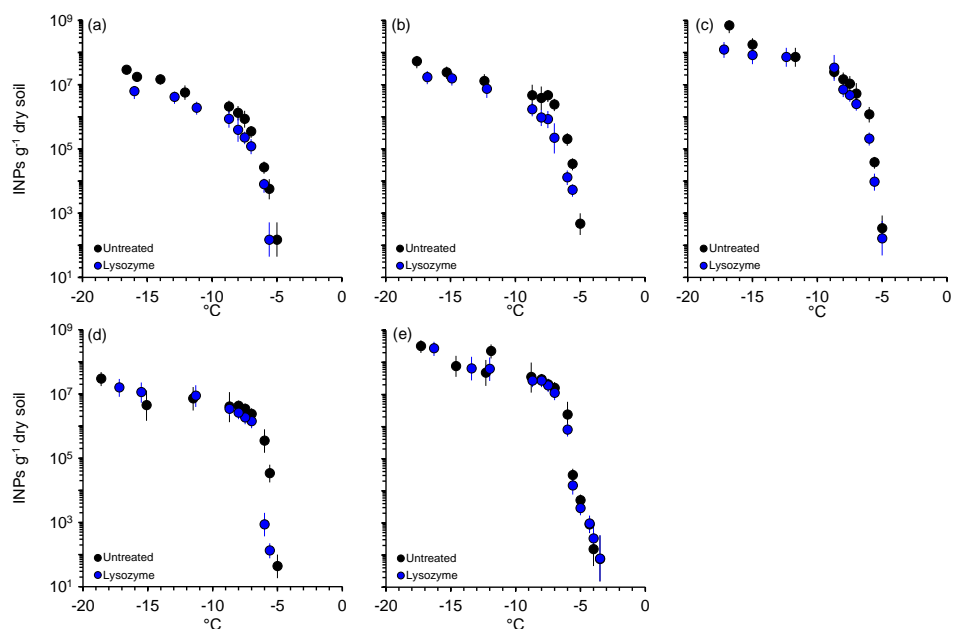
Lysozyme had a minimal overall impact upon INP concentrations but did reduce INPs active at  $\geq -6$  or  $\geq -7.5^{\circ}\text{C}$ , apart from in the lodgepole pine soil (Fig. 6). In this respect – its affect on high-temperature-active INPs – it generally corresponds to the findings of Christner et al. (2008), of lysozyme-sensitive material accounting for 50 and 25 % of INPs active  $\geq -9^{\circ}\text{C}$  in precipitation samples from Montana and Louisiana respectively. The bacteria deactivated by the lysozyme may have included the known IN bacteria at levels below the detection limits of the PCR test (see below) and/or other IN bacterial species.

Evidence exists both for and against the occurrence of IN bacteria (species of *Pseudomonas*, *Pantoea*, and *Xanthomonas*) as natural components of the soil microflora. They have been isolated at  $\approx 10^5\text{ g}^{-1}$  from surface field soil (Lindemann et al., 1982), and IN *Ps. borealis* was reported as a natural component of Arctic tundra topsoil (Wilson et al.,





**Figure 5.** Effect of heating in water to 105 °C upon INPs in soils. Soils are (a) sugar beet, (b) organic grass/alfalfa fallow, (c) pasture, (d) sagebrush shrubland, (e) lodgepole pine forest, and (f) pure kaolinite.

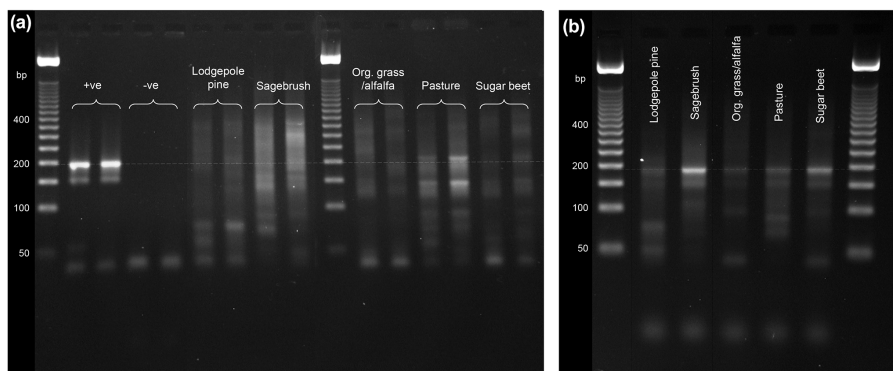


**Figure 6.** Effect of lysozyme digestion upon soil INPs. Soils are (a) sugar beet, (b) organic grass/alfalfa fallow, (c) pasture, (d) sagebrush shrubland, and (e) lodgepole pine forest.

2006). Garcia et al. (2012) also found  $\approx 10^5$  *ina* genes  $g^{-1}$  in the surface soil of a just-harvested corn field, but it is possible these came from pulverized corn tissue dusting the surface. Vali et al. (1976) also demonstrated the ability of a locally isolated *Ps. syringae* to colonize leaf litter. By contrast, Conen et al. (2011) predicted that the contribution of IN bacteria to soil INPs would be negligible based on previous stud-

ies showing that *Ps. syringae* typically did not survive more than a few weeks after addition to soils (e.g. McCarter et al., 1983; Goodnow et al., 1990). Even so, it is possible that their IN activity could be longer-lived if their Ina proteins are adsorbed onto clays.

Direct testing for the presence of IN bacteria, using PCR with primers designed to amplify most alleles of the *ina* gene



**Figure 7.** Test for *ina* gene presence in soil DNA. (a) No detectable *ina* gene amplicons (194 bp, as indicated by the faint dashed line) were discernable in any of the soils. (b) Soil DNA spiked with 60 *ina* gene copies. Amplicons were visible in all samples, although differing in intensity due to sub-optimal amplification in most.

(Hill et al., 2014), which codes for the ice-active Ina protein, did not detect them in any of the soil DNA extracts (Fig. 7a). When soil DNA was spiked with DNA from *IN Ps. syringae* Cit7, the *ina* genes were readily amplified, although with somewhat limited sensitivity due to mis-priming and co-amplification of non-target DNA (Fig. 7b). Using spiking tests with a range of *ina* gene copies, the lower detection limits were determined. When these were expressed as a fraction of INPs, the IN bacteria accounted for 0.5 to 4 % of INPs active at  $-12^{\circ}\text{C}$  in the pasture and organic grass/alfalfa soils respectively ( $-12^{\circ}\text{C}$  is the temperature at which a single Ina protein nucleates; Govindarajan and Lindow, 1988). Under optimal conditions, the proportion of IN bacteria with ice nucleation activity at  $-10^{\circ}\text{C}$  can be 10 % or higher (Kim et al., 1987; Nemecek-Marshall et al., 1993; Attard et al., 2012). Even if the populations possessed a 10 % frequency of nucleation, the IN bacteria would then only account for < 0.5 % of INPs at  $-12^{\circ}\text{C}$ .

It is possible that Ina proteins could accumulate in soil by being adsorbed onto clay and/or organic matter (Schnell, 1977; O'Sullivan et al., 2016), thereby being protected from enzymatic digestion. Accordingly, even a very small IN bacterial population could accumulate to exert a disproportionate influence upon the INP content of soils. This mechanism could also apply to other cell-free INPs.

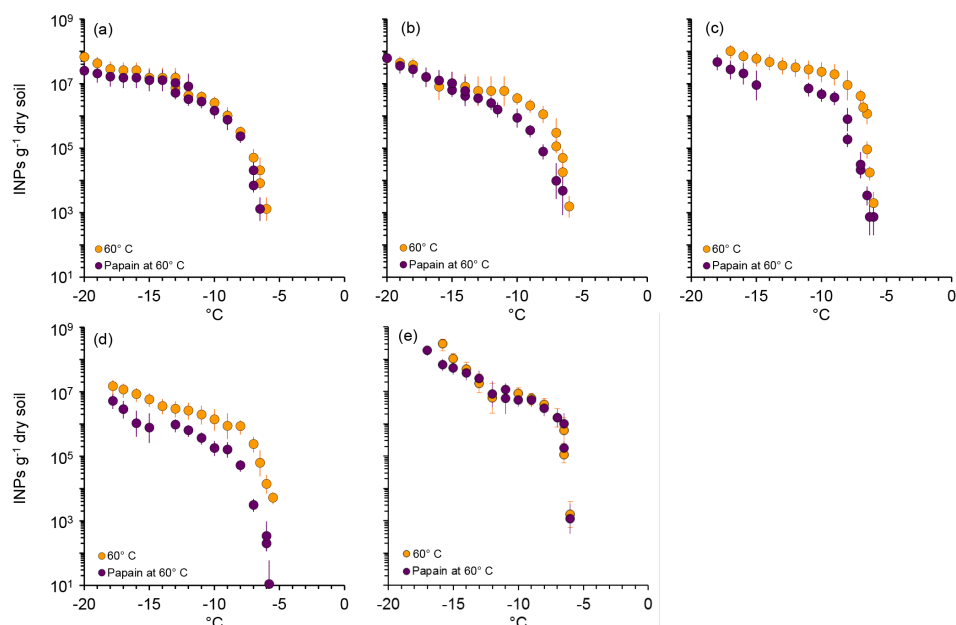
### 3.6 IN fungi

Screening of fungi isolated from agricultural and natural ecotype topsoils in Wyoming (including several of the soils used in this study) found a single species of IN fungus, *Mortierella alpina*, to be widespread and also abundant in soils with fresh inputs of plant material (Fröhlich-Nowoisky et al., 2015). *Mortierella alpina* is a saprobe utilizing decaying organic matter (Wagner et al., 2013), and due to its ability to solubilize phosphorus also develops a mutualistic association with another fungal genus that forms mycorrhizal symbioses with sagebrush (Wicklow-Howard, 1994; Zhang

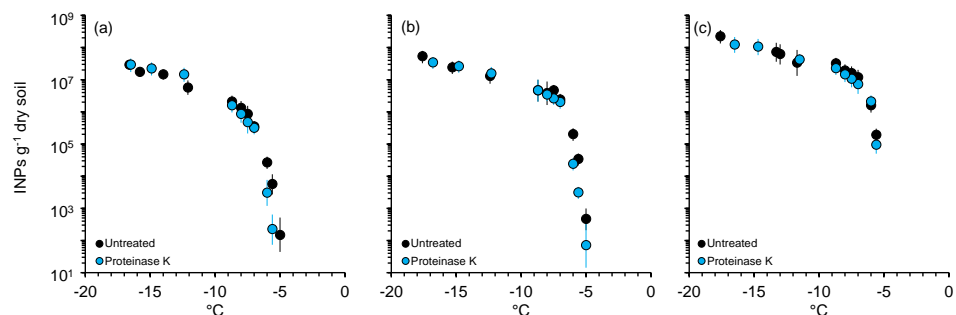
et al., 2011). All 39 isolates tested by Fröhlich-Nowoisky et al. (2015) initiated freezing at  $-5$  to  $-6^{\circ}\text{C}$ . They also typically released  $10^8$  to  $10^9$  small ( $< 10\text{ nm}$ ) INPs per gram fresh weight of mycelium that nucleated between  $-5$  and  $-8^{\circ}\text{C}$  (see O'Sullivan et al., 2015). For the three most common clades, which occurred in agricultural and lodgepole pine forest soils, digestion of isolates with papain lowered the INP concentration by  $\approx 3$ –4 orders of magnitude (Fröhlich-Nowoisky et al., 2015), suggesting the INP was a protein.

In this study, digestion with papain lowered INPs active  $\geq -11^{\circ}\text{C}$  in the organic grass/alfalfa, the pasture, and the sagebrush soils by  $\geq 75\%$  (Fig. 8). Indeed, in these soils its effect was comparable to that of  $105^{\circ}\text{C}$  heating. By contrast, papain only deactivated INPs active  $\geq -7^{\circ}\text{C}$  in the sugar beet soil and had no impact on lodgepole pine forest soil, even though IN *M. alpina* was readily isolated from both (Fröhlich-Nowoisky et al., 2015). This may have been because these soil samples had been stored frozen and had been taken at different times from those in the Fröhlich-Nowoisky et al. (2015) study, and because a high relative abundance obtained via dilution plating, which is heavily biased towards spores, does not automatically equate to a significant vegetative biomass (Warcup, 1957). Alternatively, the papain-susceptible INPs in the other three soils were not from *M. alpina* but from another source.

Several species of fungi from the genus *Fusarium* are also known to be IN (Pouleur et al., 1992; Richard et al., 1996). These species contain strains that are plant pathogens as well as saprophytes and are promoted by agricultural practices. Digestion of cultures of *F. avenaceum* and *F. acuminatum* with Proteinase K significantly reduced ice nucleation activity, suggesting that a protein is responsible (Hasegawa et al., 1994; Humphreys et al., 2001). In the agricultural soils tested, incubation with Proteinase K had a minimal effect upon overall INP concentrations. In the organic grass/alfalfa and sugar beet soils it reduced INPs active  $\geq -6^{\circ}\text{C}$  by  $\approx 10$ -fold (Fig. 9), as would be expected for deactivation



**Figure 8.** Effect of papain protease digestion upon INPs in soils. Soils are (a) sugar beet, (b) organic grass/alfalfa fallow, (c) pasture, (d) sagebrush shrubland, and (e) lodgepole pine forest.



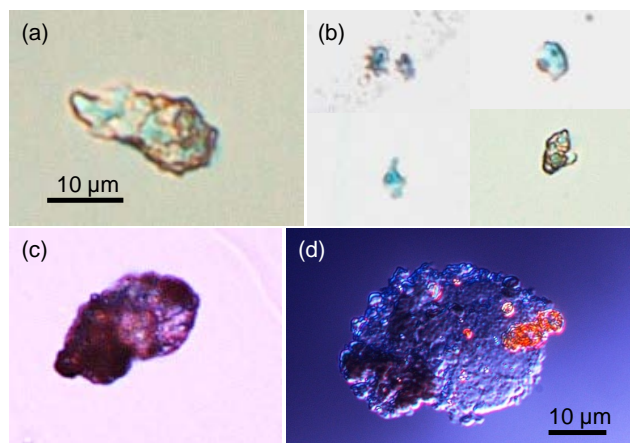
**Figure 9.** Effect of Proteinase K digestion upon INPs in soils. Soils are (a) sugar beet, (b) organic grass/alfalfa fallow, and (c) pasture.

of IN *Fusarium* (Hasegawa et al., 1994; Richard et al., 1996; Pouleur et al., 1992; Humphreys et al., 2001) and *Mortierella*, but it had no effect on the roadside pasture soil. The cultivated soils thus contained some Proteinase K-sensitive IN microflora.

### 3.7 Direct isolation of INPs from sagebrush soil

We present the first images of probable INPs directly isolated from soil. INPs active at  $-7^{\circ}\text{C}$  in the sagebrush shrubland soil were isolated by repeated cycles cooling an array of droplets to identify the INP-containing droplet, addition of de-ionized water to dilute that droplet, and its further subdivision and subsequent freeze testing. Starting with a  $10^{-5}$  dilution of the soil suspension, a further 5–6 cycles of freezing and droplet subdivision were required to obtain a single visible particle in the evaporated droplet residue.

Two INPs were isolated. INP1 was roughly  $20 \times 5 \mu\text{m}$  and appeared to be a complex entity (Fig. 10a). After isolation and photography, INP1 was re-suspended in 1  $\mu\text{L}$  water, which was divided into four droplets and these freeze tested. This time two wells froze, and in all four droplets there were numerous fragments of the original piece (Fig. 10b). In sagebrush surface soils, freeze/thaw cycles occur frequently from mid-autumn to mid-spring, providing multiple opportunities for such fracturing and multiplication of INPs. INP2 appeared to be a roughly  $30 \times 20 \mu\text{m}$  piece of gel (Fig. 10c, d). It was partly squashed by capillary forces when mounted between slide and coverslip (Fig. 10d) and was semi-transparent. Although the matrix may have been mucilage secreted by roots (Foster et al., 1983), exopolysaccharide and alginate gels are also produced by microbes, and the two often intermix forming mucigels (Foster, 1986). The actual INP was potentially one of many things: either a component of the gel itself or one of its many inclusions.

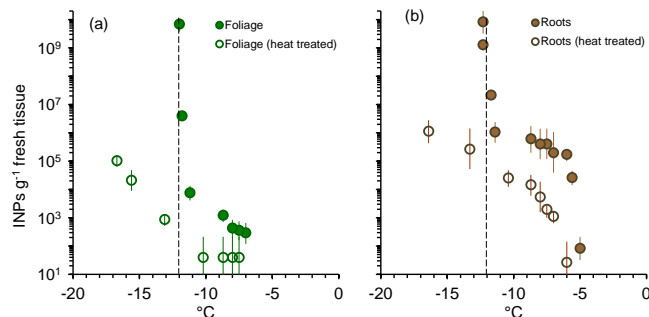


**Figure 10.** Ice nucleating particle, INP1, isolated from sagebrush soil (a). After six freeze-thaw cycles it fragmented into many small pieces, the larger fragments of which are shown in (b), creating at least two INPs (i.e. two droplets froze). Ice nucleating particle, INP2, in white light (c) and illuminated using differential interference contrast microscopy (blue polarized light was used simply to accentuate structural details, but the rust colour of the inclusions is natural, d).

Filtration tests of a sagebrush soil suspension showed that 68 % of INPs active at  $-7^{\circ}\text{C}$  were  $> 5\ \mu\text{m}$  diameter and 88 %  $> 0.45\ \mu\text{m}$ . Hence, the randomly selected INPs isolated here were also large. It is possible that the true INP was an accompanying component too small to be visible. However, the fragmentation of INP1 after six freeze/thaw cycles leading to the production of two INPs suggests that the large particle was indeed the source.

### 3.8 Ice nucleation by sagebrush tissues

The particularly pronounced onset of ice nucleation activity in sagebrush shrubland soil (Fig. 2) suggested a single class of IN material was responsible. Since Wyoming big sagebrush (*Artemisia tridentata*) contributes organic matter directly to the soil via leaf fall and the shedding of fine roots and root hairs, its tissues may be a direct source of these INPs. Indeed, powdered sagebrush shoots and roots contained a distinct and abundant class of INPs that nucleated at  $\approx -12^{\circ}\text{C}$  (Fig. 11). The relatively warm temperature of activity of sagebrush tissue INPs suggests it may play a functional and protective role for the plant, although its onset temperature is much lower than that possessed by IN peach wood tissue ( $\approx -2^{\circ}\text{C}$ , Gross et al., 1988). Interestingly, *A. tridentata* seedlings from comparable altitudes exhibited significant damage ( $> 50\%$  loss of potential function of Photosystem II) when chilled to between  $-12.5$  and  $-15^{\circ}\text{C}$  (Loik and Redar, 2003), suggesting that sagebrush tissues may naturally supercool until being nucleated by these endogenous INPs. Heating to  $105^{\circ}\text{C}$  completely deactivated the INPs active at  $\approx -12^{\circ}\text{C}$ . Likewise, they appeared to be lost upon



**Figure 11.** Ice nucleating particles in sagebrush shoot (a) and root (b) tissues ground under liquid nitrogen and after heating to  $105^{\circ}\text{C}$ . The dashed line at  $-12^{\circ}\text{C}$  is a guide to indicate the temperature of freezing of the tissues.

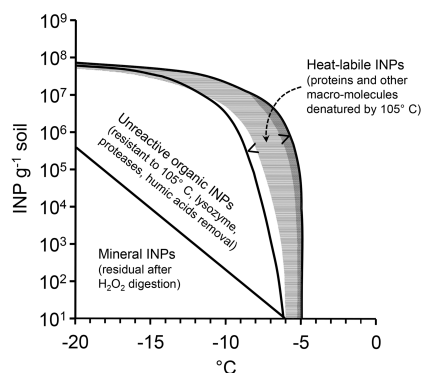
decomposition, since there was no evidence of an increase in INPs active around this temperature in the soil (Fig. 2). Cellulose may also have contributed to the residual INP activity of the sagebrush tissue that remained after heating (Hiranuma et al., 2015a).

In addition to the highly active entity at  $\approx -12^{\circ}\text{C}$ , sagebrush roots also had a lesser class of INPs with a profile similar to that of the soil (Fig. 11b). This high-temperature group may have been contributed by the diverse microflora that colonizes roots, from soil biological material still attached to the epidermis of the roots after washing, or from arbuscular mycorrhizal fungi and other microflora inhabiting the root cortex (Watson, 1987). Notably, one of the few fungal genera that form a mycorrhizal symbiosis with sagebrush also develops a mutualistic association with *M. alpina* (Wicklow-Howard, 1994; Zhang et al., 2011).

## 4 Conclusions

Ice nucleating particles denatured by heating ( $105^{\circ}\text{C}$ ) accounted for most INPs active  $> -10^{\circ}\text{C}$ . A range of entities both within the soil microflora and SOM will be heat-labile. Proteins will be denatured; molecular complexes, organelles and structures (e.g. ribosomes, flagella, membranes) entirely disrupted; crystals dissolved; and aggregates of storage polymers (e.g. polyhydroxyalkanoates) and lipid bodies dispersed. Immersion in hot water could also affect the IN ability of minerals. For example, the IN activity of K-feldspar (Atkinson et al., 2013) may be altered by a change in composition of surface ions (Zolles et al., 2015). However, the lack of any effect upon heating samples to  $60^{\circ}\text{C}$  suggests this was not the case (see also O'Sullivan et al., 2015).

While lysozyme and Proteinase K had minimal overall impacts upon INP concentrations, papain protease reduced INP numbers significantly in two agricultural soils and in the sagebrush shrubland soil, suggesting that in these a protein was responsible. This may have been due to the presence of *M. alpina* (Fröhlich-Nowoisky et al., 2015), but it equally



**Figure 12.** Sources of ice nucleating particles in the pasture soil. Within the heat-labile segment, horizontal lines indicate IN proteins digested by papain (e.g. from *Mortierella alpina*), and diagonal lines INPs in live bacteria (sensitive to lysis with lysozyme). The mineral INPs wedge is inherently approximate since  $\text{H}_2\text{O}_2$  oxidation may not remove all SOM and cannot be assumed not to deactivate IN sites on some minerals.

may have been due to IN proteins from other microbes and/or the soil fauna (e.g. protozoa, mites, nematodes). The three enzymes have very specific sites of activity within their target molecules (lysozyme also hydrolyses fungal chitin oligosaccharides, but does not digest the chitin polymer itself) but may have also had non-target influences. For example, enzymes may bind to and block INPs on mineral surfaces (see Zolles et al., 2015). Further, while reaction conditions for all enzymes were theoretically optimal, and ample time given for completion, their efficiencies are not guaranteed in such a complex milieu as soil; their substrates may have been protected by adsorption to clays (O’Sullivan et al., 2016) or tannins or by location within flocs or biofilms, and they themselves may have been inactivated by similar mechanisms or by cleavage by native proteases.

Two probable INPs were isolated by exploiting their IN activity (initially at  $-7^\circ\text{C}$ ). Since both were complex entities, the specific IN component was not able to be identified. Freeze fracturing of one into at least two INPs demonstrated that freeze–thaw cycles in topsoils could cause multiplication of INPs.

Figure 12 summarizes the classes of INPs in the pasture soil. With variations in proportions it can be used as a general summary of likely INP composition in arable and grassland topsoils. Notable is the segment of organic INPs active below  $-10$  to  $-12^\circ\text{C}$  that was unaffected by any challenge short of oxidation with  $\text{H}_2\text{O}_2$  (labelled “Unreactive organic INPs”). Their refractoriness can be used to rule out some potential sources:

- they are not proteins or peptides, due to their resistance to  $105^\circ\text{C}$  heating and insensitivity to papain and Proteinase K (they could be proteins protected by adsorp-

tion onto clay, but this would not explain their presence in the lodgepole pine litter);

- they are not peptidoglycan from bacteria since they survived lysozyme digestion;
- they are unlikely to be chitin and related fungal cell wall polysaccharides since Pummer et al. (2013) found no appreciable ice nucleation activity above  $-26^\circ\text{C}$  in fungal spores from a range of common fungi (other than in a known IN *Fusarium*);
- they do not appear to be fulvic or humic acids.

Hence, they are some component/s of plant material, microbial biomass, and/or humin. Humin is comprised in large part of macromolecules or aggregates of peptides, aliphatics, peptidoglycan, carbohydrates, and lignin. Ruling out the first three leaves lignin or carbohydrates as possibilities. Lignin seems an unlikely candidate as an INP due to its disordered structure and relative hydrophobicity. Cellulose, a carbohydrate, may account for some of the activity colder than  $-15^\circ\text{C}$  (Hiranuma et al., 2015a).

The variable effectiveness of the challenge tests in different soils indicates that soil INPs come from a range of sources, occur with differing relative abundances, and may be protected by different mechanisms. For example, the complete insensitivity of INPs in the lodgepole pine forest litter to any tests other than heat and  $\text{H}_2\text{O}_2$  digestion suggests that this population differs fundamentally from those in non-forest soils.

Since SOM comprises a diverse continuum of forms, ranging from fresh plant inputs to ancient inert organic matter, it should be expected that both the compositions and the intrinsic temperatures of nucleation of the INPs will be similarly broad. This can be inferred from the smooth curves of the INP temperature spectra, each being the sum of many underlying activity distributions. Organic INPs will also occur as organo-mineral complexes, which will presumably affect their activity and protect them by spatial isolation and adsorption onto clays (Schnell, 1977; Kögel-Knabner et al., 2008; O’Sullivan et al., 2016).

All soils, with the relative exception of ponderosa pine litter, were rich potential sources of organic, warm-temperature-active IN aerosols. Agricultural intensification in semi-arid regions of the western USA over the past 200 years has led to a more than 4-fold increase in atmospheric dustiness (Neff et al., 2008). The increased wind erosion also selectively removes the fine and light fractions, which are enriched in particles  $< 10\ \mu\text{m}$  in diameter and organic matter approximately 6-fold (Van Pelt and Zobeck, 2007). Biological particles, predominantly bacteria and fungi, accounted for an average of 40 % of the organic carbon in particles  $< 10\ \mu\text{m}$  at Storm Peak Laboratory in western Colorado (Wiedinmyer et al., 2009) and were present in dust events intersecting the site (Hallar et al., 2011). INP emissions by these western

US landscapes will thus be influenced by soils that are more prone to destabilization by both previous and current human activities.

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