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Dear Editor,

Please find attached a point-by-point summary of our replies to the referee comments, which details our changes to the manuscript initially submitted to ACPD, along with a marked-up version of the manuscript highlighting changes. In addition to changes adopted based on the referee comments, we have also made a minor amendment to the caption of Figure 5, which now reads:

“After each washing cycle, ~10 % of the protein was observed to be lost to the supernatant, suggesting that the adsorption process is partially reversible. This is in addition to a maximum of 4 % loss in each wash, due to the amount removed to conduct the droplet freezing experiments.”

This addition relates to a comment raised to us by a colleague, on whether we had taken this loss due to the experimental procedure into account in the ACPD manuscript. Indeed, we had, but not explicitly stated the fact and as such wish to add this statement for clarity.

We hope you find our changes satisfactory.

Many thanks for your time, and consideration of our manuscript.

Best Wishes,

A handwritten signature in black ink that reads "Daniel O'Sullivan".

Daniel O'Sullivan

Response to Referee #1

The Manuscript entitled “The adsorption of fungal ice-nucleating proteins on mineral dusts: a terrestrial reservoir of atmospheric ice-nucleating particles” deals with the potential of proteins to serve as cell free and available IN in soils. In detail, clay and cell free ice nuclei washed from the mycelial cell wall from *Fusarium avenaceum* are used in laboratory experiments as demonstration. Basically, the manuscript is very well written, and the background is precisely elaborated. Therefore I can easily give the manuscript the excellent for the form. The topic of organic and biological residues is of major interest in this field of science. As stated in the manuscript, most mineral particles are covered with carbonaceous matter originating from organic and/or biological sources. I agree here with the authors these might be of significant impact and also might have relevance in water cycling.

Fusarium has spread to the whole World. Especially *Fusarium avenaceum* as generalist in plant pathogenicity commonly found to be IN active contribute a huge amount of biological particles on our Earth. However, there would be a gain for the data. It is obvious, that there is a chemical or biochemical modification of the clay upon incubation with *F.avenaceum* washing water. But there is no experiment showing it to be proteins. A protein extraction to recover proteins from the clay or incubated clay used in a protein assay (e.g. BCA or Bradford) could show proteins or proteinaceous compounds bound to the clay. Although it is known that proteins adsorb to minerals, this would fortify the interpretations, and could also be related to the impact of the different cations in either protein or proteinaceous IN adsorption to the clay. Because of the complex composition of biological samples, an experiment showing protein being enriched in the treated clay would be a gain. Anyhow, the results are related to the conclusion as well as I generally agree with the interpretation. The protein assay is not mandatory for the manuscript and the conclusions are concise. Therefore I recommend publication of the manuscript in ACP after the open discussion.

We thank referee #1 for both taking the time to review our manuscript and for his/her insightful comments. Our replies are given in blue, while changes to the text are listed in *blue italic*.

With regards to there being no experiment showing that the modification of the clay upon incubation with *Fusarium* washing waters is caused by a protein, on the basis of evidence in the literature, it is very unlikely that proteinaceous compounds are not involved. The proteinaceous characteristics of INPs from *F. avenaceum* include the thermal lability of INPs, and susceptibility to treatment with proteinase K (*Hasegawa et al (1994)*). Agents such as phenylboric acid and chloroform did not affect activity, suggesting that lipids and saccharides are not central to its nucleating ability (*Hasegawa et al (1994)*). Also of note, for the ice nucleating fungus *Fusarium acuminatum*, genes coding for the ice-nucleating protein have reportedly been expressed in *E. coli* (*Lagzian et al., 2014*).

On the basis of this, within the manuscript we referred to the INPs as “proteinaceous-INPs”, consistent with the literature. We agree that further purification, isolation and characterisation of the macromolecule(s) responsible for activity in *F. avenaceum* is certainly of interest, but we suggest that this is an area of future research, rather than being critical for the conclusions drawn in this article.

We have highlighted the previous literature which supports INPs in the washing waters as being proteinaceous in the introduction for the interested reader to gain an understanding of the current state of knowledge:

*“Similarly, a single gram of *Fusarium avenaceum* harbours 10⁸ nano-sized INPs active at temperatures above -7 °C (O’Sullivan et al., 2015). These INP appear to be proteinaceous as inferred from their thermal lability, combined with their susceptibility to treatment with Proteinase K (Hasegawa et al (1994))”*

References

Hasegawa, Y., Ishihara, Y., and Tokuyama, T.: Characteristics of ice-nucleation activity in *Fusarium avenaceum* IFO 7158, *Biosci., Biotechnol., Biochem.*, 58, 2273-2274, 10.1271/bbb.58.2273, 1994.

Lagzian, M., Latifi, A. M., Bassami, M. R., and Mirzaei, M.: An ice nucleation protein from *Fusarium acuminatum*: cloning, expression, biochemical characterization and computational modeling, *Biotechnol. Lett.*, 36, 2043-2051, 2014.

Response to Referee #2, Dr. Tom Hill

We would like to thank Dr. Hill for taking the time to review our article and for his helpful comments. Our replies are given in blue, while changes to the text are listed in *blue italic*.

The ability of organic INPs (“leaf-derived nuclei”) to adsorb to and confer enhanced IN activity to minerals (esp. clays) was first proposed and demonstrated by Russ Schnell almost 40 years ago. Recent research has generated an awareness of the potential importance of this process, especially following recent studies showing that organic soil INPs are often very small (ie, would not be detected when bound to mineral particles). This paper is an important and overdue advancement of this topic. Its main new contribution is that IN proteins are readily adsorbed onto clays while maintaining their activity. Secondly, it shows that ionic concentration (in the range found in soil water) and, to lesser extent, composition are important positive variables enhancing binding, but that paper pH is not. It is also useful and intriguing to be shown that the adsorption is reversible, which to me suggests that the binding process is relatively gentle and, hence, doesn’t deform and potentially inactivate the proteins.

I have some suggestion for minor amendments.

P. 3 L. 18: “released” would be better than “lost”.

Agreed- changed.

P. 3 L. 19: date missing for Pouleur.

Corrected

P. 3 L. 21: no comma.

Corrected

P. 4 L. 15: “much higher”.

Corrected

P. 5 L. 4-14: Very nice background to this aspect.

Many thanks!

P. 6 L. 22: I would add “(collected in the filtrate)” for those who need reminding of their small size.

Added “*after collection in the filtrate*” here

P. 8, L. 22: Can you remind us what K is here?

Added “*(no. of ice-nucleating sites per ml of suspension, eq. 1)*”

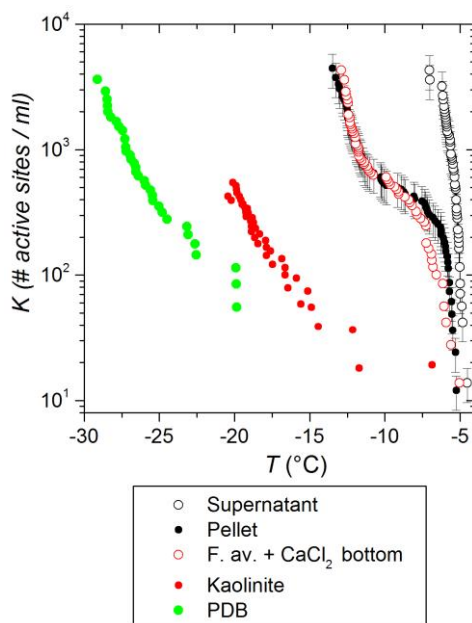
P. 8 Clay-protein interactions in the absence of electrolytes section: In relation to Fig 2., I notice that in Fig. 3 the 1 mM NaCl treatment was also apparently unimpressive after 2 h, but given time was very effective. The same may have held true for the no electrolytes case. If you didn't test these ones for 48 h then you should acknowledge this even though it doesn't change the underlying story, and even though in a soil solution you would seldom encounter such a lack of ions. IN Fig. 2, it would also be nice to know what the underlying kaolinite INP profile was. I assume it's the log-linear line that would exist of the hump starting from -11 C and warmer was removed. Could this be mentioned?

Thanks for these points. The aim of this section was purely to show that regardless of pH, adsorption was not observed to occur within a contact time typical of when adsorption has been observed to occur for a wide variety of other proteins on minerals. To add further context to this section, and justify why we chose the 2 hour contact time in the first place, we have added to section 3.1:

“As adsorption is a time-dependent process, it is possible that extending the contact time could result in uptake of the protein to the clay. Nevertheless, the results here show that on timescales characteristic of protein adsorption to clays (Yu et al., 2013), uptake of the P-INP was minor in the absence of electrolytes.”

With relation to Fig. 2, the kink below -11 °C does not appear to be related to the presence of kaolinite, which is active at far lower temperatures. We also note that it is not related to the potato dextrose broth in which the fungus was grown. To illustrate this we have added a supplementary figure where we show the K values for PDB and kaolinite. We have also added the text in section 3.1:

“Measured K values for kaolinite on its own (with no adjustment to pH) are far below those of either the pellet or supernatant (Fig. S1).”



*Fig. S1: Results of supernatant depletion test at pH 5.7 (as per Fig. 2, no added salt ions), compared to tests for ice nucleation by potato dextrose broth (PDB, diluted 1/10 from the original broth) and a 1 wt% kaolinite suspension (Whale et al. (2015)). Note that only a trace of PDB will adhere to the mycelium, and hence make it into the supernatant depletion suspensions; accordingly there is no significant contribution to the observed ice nucleation activity from PDB. Nucleation from kaolinite is many orders lower than *F. avenaceum*. Also shown is the bottom 0.1 mL fraction from a control supernatant depletion test performed on *F. avenaceum* in 1 mM CaCl_2 , where no kaolinite was added. No enrichment of the protein in the bottom fraction can be seen, consistent with a lack of protein flocculation.*

P. 9 L.7: Couldn't it simply be that the ionic strength is just so low that the tendency was for surface ions of any type to remain in the bulk phase, just as K^+ is rapidly stripped off K-feldspar when that is put in DI?

On the contrary, it would be expected that the release of exchangeable surface cations in deionized water for kaolinite Kga 1-b would be diminishingly small, as there are no cations in solution to replace those lost from the surface to the solution. The equilibrium between ions at the surface and those solvated in solution will lie heavily towards those remaining at the surface, suggesting that ions being lost from the surface play at most only a very minor role in explaining the adsorption dynamics. In this case we prefer to keep the discussion around that of Tombácz and Szekeres, 2006 which is already cited in the paper.

P. 9 L.11: Surely it's not just the charge, but also the presence of ions in solution to replace any that leave the surface of the clay, in a dynamic equilibrium? It would be useful, too, to provide a typical range for ionic strengths in soil solutions. I see Edmeades et al (1985; Aust J of Soil Res., 23, 151) gives a range of 5-16 mM for NZ grassland soil, which is very relevant to your Fig. 4., supporting the case that under normal conditions the INPs would rapidly bind to the clays.

Agreed- we've added "and promote cation exchange at the surface of the clays" here.

With regards to typical ionic strengths of soil solutions, we have added further info for the interested reader: *“Ionic strengths of 1 and 10 mM were chosen to approximate those found in soil solutions (Campbell et al., 1989;Griffin and Jurinak, 1973;Edmeades et al., 1985;Dolling and Ritchie, 1985).”*

P. 9 L.17: Date for Yu et al.

Fixed.

P. 9 L.18: Fig. 3 is impressive.

We were also intrigued to see how strong an effect the addition of salts had!

P. 11 L. 9: Also, divalent cations may cause flocculation of both clay particles and proteins, producing clumps of these. The natural pH of the kaolinite soln (5.7) would promote this.

For solutions of *F. avenaceum* washing waters only, 1-10 mM ionic strength is a very low concentration in terms of that required to cause proteins to flocculate on their own. This was verified with control supernatant depletion experiments where kaolinite was omitted- no increases in the bottom 0.1 mL of the suspension were observed (see Fig. S1). When the clay is present, yes, this perhaps encourages the clay particles to stick together, but this does not affect the conclusion that the proteins preferentially adsorb to the clay. We have modified the pertinent line in the methods section, which now refers to the new Fig S1:

“Control experiments were performed in the absence of kaolinite with Milli-Q® purified water alone, or with CaCl₂ showing that there was no significant preferential sedimentation of the protein (see Fig S1).”

P. 12. L.24: Maybe add the caveat that adsorption to kaolinite did not cause deactivation. During the process of drying, for example, the adsorption/binding may become stronger and so deform and affect the IN activity and propensity to be desorbed.

We have added the caveat:

“although further environmental processing, such as repeated wetting and drying of soils, could potentially perturb protein activity”

P. 13 L. 7: Nicely put.

P. 13. L. 14: I don't think all the efficient ones are proteins. Or, at least, it's premature to generalize?

Agreed- this statement was too strong. Changed to:

“Indeed, the characteristics of the organo-clay INPs examined in this study are similar to the efficient INPs which have been identified in topsoils (O'Sullivan et al., 2015;Conen et al., 2011), a substantial proportion of which were shown to be thermally labile and exhibit exceptional ice-nucleating abilities.”

P. 16. L. 16: Fusarium are common in soils yes, but not the most common. They tend to be pathogens. And only a few species are IN active. This sentence is overstating their abundance I think. Since this work would apply equally, I assume, to other IN fungi, such as *Mortierella alpina*, and other as-yet undiscovered IN fungi (the most dominant species, in terms of vegetative biomass, tend to be the Basidiomycetes, which are notoriously difficult to grow in pure culture), this section could be broadened to include adsorption of IN proteins released by many soil organisms.

Thanks for drawing our attention to this, we had worded this poorly. This has been reworded as below.

With regards to this work applying equally to other sources of proteinaceous INPs, it is correct that the conclusion that P-INPs from other sources could also confer activity to soils, although this must be caveated by noting that the extent of adsorption may differ. Nonetheless, this is an important point, and to this end we have broadened our discussion. The section now reads:

“In addition to Fusaria, other organisms can also contribute to the reservoir of cell free P-INPs in soils, such as the fungus M. alpina (Fröhlich-Nowoisky et al., 2015), certain lichens (Kieft and Ruscetti, 1990), and even ice-nucleating bacteria (Phelps et al., 1986). For Fusaria, ice-nucleating activity has been observed in a number of species, and species such as Fusaria are widespread in soils, occurring across the globe wherever crops are grown (Pitt et al., 2009)”

References

Campbell, D., Kinniburgh, D., and Beckett, P.: The soil solution chemistry of some Oxfordshire soils: temporal and spatial variability, *J. Soil Sci.*, 40, 321-339, 1989.

Conen, F., Morris, C., Leifeld, J., Yakutin, M., and Alewell, C.: Biological residues define the ice nucleation properties of soil dust, *Atmos. Chem. Phys*, 11, 9643-9648, 2011.

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Kieft, T. L., and Ruscetti, T.: Characterization of biological ice nuclei from a lichen, *J. Bacteriol.*, 172, 3519, 1990.

O'Sullivan, D., Murray, B. J., Ross, J. F., Whale, T. F., Price, H. C., Atkinson, J. D., Umo, N. S., and Webb, M. E.: The relevance of nanoscale biological fragments for ice nucleation in clouds, *Sci Rep*, 5, 8082, 10.1038/srep08082, 2015.

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1 The adsorption of fungal ice-nucleating proteins on mineral dusts: a terrestrial
2 reservoir of atmospheric ice-nucleating particles

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Abstract

The occurrence of ice-nucleating particles (INPs) in our atmosphere has a profound impact on the properties and lifetime of supercooled clouds. To date, the identities, sources and abundances of particles capable of nucleating ice at relatively low supercoolings ($T > -15\text{ }^{\circ}\text{C}$) remain enigmatic. While biomolecules such as proteins and carbohydrates have been implicated as important high-temperature INPs, the lack of knowledge on the environmental fates of these species makes it difficult to assess their potential atmospheric impacts. Here we show that such nanoscale ice nucleating proteins from a common soil borne fungus (*Fusarium avenaceum*) preferentially bind to and confer their ice-nucleating properties to kaolinite. The ice-nucleating activity of the proteinaceous INPs is unaffected by adsorption to the clay, and once bound the proteins do not readily desorb, retaining much of the activity even after multiple washings with pure water. The atmospheric implications of the finding that biological residues can confer their ice-nucleating ability to dust particles are discussed.

1 1. Introduction

2 Aerosols in our atmosphere can modify cloud properties by acting as cloud condensation nuclei (CCN) and
3 ice-nucleating particles (INPs). While atmospheric INPs are rare in comparison to CCN (DeMott et al.,
4 2010;Pruppacher and Klett, 1997), they exert a disproportionate effect on cloud properties such as albedo,
5 extent, lifetime, dynamics and conversion to precipitation (Rosenfeld et al., 2011;Storelvmo et al.,
6 2011;Boucher et al., 2013). Biological particles in our atmosphere are the most efficient known catalysts
7 relevant for ice formation in supercooled cloud water droplets (Murray et al., 2012;Hoose and Möhler,
8 2012). Certain bacteria, fungal spores and pollen grains can facilitate the freezing of supercooled cloud
9 water droplets at temperatures where common inorganic ice-nucleating particles such as mineral dusts or
10 black carbon are comparably inactive ($T > -15\text{ °C}$) (Murray et al., 2012;Hoose and Möhler, 2012;Fröhlich-
11 Nowoisky et al., 2015;Pummer et al., 2015;O'Sullivan et al., 2015). However, number densities of these
12 supermicron biological particles are typically many orders of magnitude lower than those of their inorganic
13 counterparts at altitudes relevant for supercooled clouds (Hoose et al., 2010).

14
15 The majority of past studies on the atmospheric implications of biological ice nucleators have focused on
16 examining the activities of supermicron cellular entities such as bacteria, fungal spores and pollen grains.
17 However, there is evidence that subcellular portions of biological INPs, such as proteins and carbohydrates,
18 can be readily ~~lost~~released from terrestrial INPs such as fungal mycelia or pollen grains on contact with
19 water and continue to exhibit the ice-nucleating abilities of the parent particles (Pouleur et al.,
20 1992;Augustin et al., 2013;O'Sullivan et al., 2015;Fröhlich-Nowoisky et al., 2015;Pummer et al., 2012).
21 Similarly, in the oceans, subcellular INPs thought to be associated with phytoplankton, have also been
22 identified (Wilson et al., 2015;Schnell and Vali, 1975). This is important because the number densities of
23 these nanoscale fragments can far outnumber those of the parent cellular particles. For instance, each
24 single grain of pollen has roughly 10,000 INPs at its surface, which are easily separable on contact with
25 water (Augustin et al., 2013;O'Sullivan et al., 2015). Similarly, a single gram of *Fusarium avenaceum*
26 harbours 10^8 nano-sized INPs active at temperatures above -7 °C (O'Sullivan et al., 2015). These INP

1 appear to be proteinaceous as inferred from their thermal lability, combined with their
2 susceptibility to treatment with Proteinase K-(Hasegawa et al (1994)). -However, given their size, these
3 nanoscale INPs (nano-INPs) are unlikely to be directly aerosolized in abundance from surfaces. While
4 readily lost from cells on wetting, significant questions remain regarding the environmental fates of
5 biological nano-INPs. In particular, do these efficient INPs adsorb to larger particles which are more readily
6 aerosolized such as soil mineral dusts, and critically, is activity retained upon adsorption?

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8 Several recent studies have indicated that biogenic matter co-transported with soil dusts has an
9 exceptional ability to catalyze the freezing of supercooled water droplets. Pratt and co-workers (2009)
10 demonstrated that in mixed phase clouds above Wyoming, mineral dust particles were the dominant ice
11 crystal residues sampled, while 60% of these mineral dust particles were internally mixed with an
12 observable amount of biogenic and/or humic-like substances. Similarly, co-located mineral dusts and
13 biological materials have been demonstrated by Creamean et al. (2013) to represent a major contributor
14 in the generation of precipitation over the Sierra Nevada Mountains on the West Coast of the U.S.

15
16 Laboratory-based studies have shown that organic matter in soils is a potent source of INPs with activities
17 greatly much higher than those of the mineral components in the soils (Conen et al., 2011;O'Sullivan et al.,
18 2014;Tobo et al., 2014;Hill et al., 2016). In the 1970s, Schnell (1977) performed experiments where distilled
19 water was passed through a column of kaolinite capped with decaying leaf litter. Nuclei in the leaf litter
20 were passed down through the column, leading to a dramatic increase in the ice nucleation activity of the
21 kaolinite. Schnell suggested that biogenic materials became associated with the clay particles, imparting
22 their ice-nucleating activity to the clay. Since this early study, it has been posited that the exceptional ice-
23 nucleating abilities of fertile soil dust may stem from the incorporation of residues from bacterial, fungal
24 and pollenaceous ice-nucleating particles (Conen et al., 2011;O'Sullivan et al., 2015;Fröhlich-Nowoisky et
25 al., 2015). Very recently, it has been reported that nanoscale pollenaceous *INPs*, which are thought to be
26 carbohydrates, retain their nucleating activity when internally mixed with mineral dusts particles

1 (Augustin-Bauditz et al., 2016). However, to date it remains unclear as to whether cell-free ice-nucleating
2 proteins, such as those derived from *Fusarium* fungi, can adsorb to and confer their exceptional nucleating
3 activities to mineral dust particles in soils.

4

5 Once adsorbed to mineral surfaces, degradation of biomolecules such as proteins, peptides and amino
6 acids by soil microbes and enzymes is generally reduced (Gianfreda et al., 1991; Fiorito et al., 2008; Koskella
7 and Stotzky, 1997; Calamai et al., 2000; Lozzi et al., 2001). Indeed, proteins and carbohydrates in soils can
8 exhibit exceptionally long lifetimes, on the order of up to half a century (Schmidt et al., 2011).
9 Furthermore, adsorbed biomolecules can continue to exhibit activity when attached to clay surfaces. For
10 instance, bound enzymes continue to convert substrates (Leprince and Quiquampoix, 1996; Quiquampoix,
11 1987; Rao et al., 2000; Calamai et al., 2000), and bound insecticidal proteins from *B. thuringensis* retain
12 their ability to kill larvae (Koskella and Stotzky, 1997). However, once adsorbed to mineral surfaces, the
13 magnitude of enzyme activities is typically reduced, and in some cases completely lost (Zimmerman and
14 Ahn, 2011). Hence, it is necessary to study if the activity of ice-active proteins is retained when bound to
15 clay minerals.

16

17 Given recent findings that extracellular proteinaceous INPs (P-INPs) from common, globally distributed soil
18 fungi are readily separated from the parent fungus upon contact with water (O'Sullivan et al.,
19 2015; Fröhlich-Nowoisky et al., 2015), examination of their environmental fates is required to understand
20 how they can add to the budget of atmospheric INPs. In this study, we examine the interactions of
21 nanoscale ice-nucleating proteins from the soil-borne fungus *Fusarium avenaceum* with the common soil
22 clay mineral kaolinite. It is demonstrated that, in the presence of cations commonly found in soil solutions,
23 the P-INPs are rapidly adsorbed to the clay mineral, with the majority remaining bound even after
24 successive washes. The implications of the finding that the proteins continue to exhibit their exceptional
25 ice-nucleating abilities once they adsorb to mineral dust particles are discussed.

26

2. Methods

2.1 Sample Preparation

Fusarium avenaceum was sourced from the Commonwealth Agricultural Bureau (CABI) and grown by incubation in sterile potato dextrose broth for three days at 28 °C. A portion of mycelium was separated from the broth and resuspended in 50 mL of Milli-Q® grade water. Residual potato dextrose broth on the mycelium was minor. Regardless of this, the potato dextrose broth itself was tested for ice nucleation activity, and no activity above the background (freezing of Milli-Q® purified water) was observed (Fig. S1). The mycelial suspension was then agitated by hand, and the proteinaceous ice-nucleating particles separated by filtration using 0.2 µm cellulose acetate filters (Sartorius Stedim®, Minisart 16534) and oil-free syringes.

11

To examine the affinity of cell-free proteinaceous ice-nucleating particles (P-INPs) to the 1:1 (Si:Al) clay mineral kaolinite, the untreated clay was mixed together with an aqueous suspension containing P-INPs recovered by 0.2 µm filtration of a *Fusarium avenaceum* suspension. Using the supernatant depletion method (Ralla et al., 2010; Ding and Henrichs, 2002; Lepoitevin et al., 2014), the clay was removed from the suspension by centrifugation, and resuspended in an equivalent volume of water (Fig. 1). Any freely suspended proteins would be expected to remain in the supernatant, while those bound to clay mineral particles would collect as a pellet at the bottom of the centrifuge tube. The ice-nucleating activity of both the resuspended clay, and particles which remained in the supernatant were then determined using a microliter droplet freezing technique (see section 2.2).

21

For these adsorption experiments, 4 ml of a stock P-INP suspension (derived from an initial concentration of 2×10^{-2} g/ml of mycelium prior to filtration, followed by a 1/20 dilution after ~~filtration~~ collection in the filtrate) was mixed with 1 ml of a stock suspension of Clay Mineral Society kaolinite Kga-1b (0.125 g/ml) and diluted to 10 ml total. The kaolinite was dispersed in water by ultra-sonication and stirred magnetically for about one hour to produce the stock suspension, before being mixed with the stock P-INP suspension.

1 The effects of solution acidity were examined by modifying the initial *pH* with either HCl or NaOH, and
2 monitoring using a Hanna instruments glass *pH* meter. The effects of electrolytes was also probed, by
3 adjusting the solution ionic strength to either 1 or 10 mM with sodium chloride (Fisher, S/3160/5), calcium
4 chloride dihydrate (Sigma, C3306) or magnesium chloride hexahydrate (Sigma, M2670). After a prescribed
5 contact time (either 2 hours or 2 days), 1 ml of the suspension was centrifuged at 10,000 ×g for 4 minutes
6 to pellet the clay. The supernatant was then carefully removed by gentle decanting, and the use of a 50 µL
7 pipette if necessary. Following this, the pellet was resuspended in 1 ml of water and vortexed to disperse.
8 Control experiments were performed in the absence of kaolinite with Milli-Q® purified water alone, or
9 with CaCl₂ showing that there was no significant preferential sedimentation of the protein (see Fig S1). For
10 desorption experiments, the clay pellet (and associated P-INPs) was repeatedly washed in Milli-Q® purified
11 water, resuspended and centrifuged again.

12

13 **2.2 Measurement of ice-nucleating activities**

14 The measurement of ice-nucleating activities was performed using the Leeds microliter nucleation by
15 immersed particles instrument (µL-NIPI). The apparatus and experimental procedures have been
16 described in depth elsewhere (O'Sullivan et al., 2014; O'Sullivan et al., 2015; Whale et al., 2015) and so are
17 only briefly discussed here. 40-50 1 µL droplets were pipetted onto a 22 mm diameter silanized slide
18 (Hampton Scientific), which was cooled from the underside using Stirling-engine-chilled aluminium stage
19 (Grant-Asymptote, EF600). The headspace above the droplets was enclosed in a Perspex shield, and gently
20 purged with 200 cm³ min⁻¹ of nitrogen gas to prevent unwanted condensation. Nucleation events were
21 detected using a USB camera embedded inside the external humidity control chamber by manually
22 analyzing the recorded video. For all experiments, a 1 °C min⁻¹ cooling rate was employed. The
23 temperature error associated with the platinum resistance thermistor which monitored the aluminium
24 cold stage was estimated as ±0.4 K. Each experiment was repeated twice, and the cumulative fraction
25 frozen (*F*) of droplets compiled as a function of temperature. Full crystallization at such high temperatures

1 could take up to a few seconds; nucleation points were taken as the first observable instance of ice
2 formation in the droplets.

3 4 **2.3 Data analysis**

5 From the cumulative fraction of droplets frozen (F) as a function of temperature, the time independent
6 singular formulation can be used to describe the number of nucleating sites active at a given temperature
7 contained within a given volume of water (V) (Vali, 1971):

$$K = \frac{-\ln(1 - F)}{V} \quad (1)$$

10

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12 Accordingly, the difference in the number of ice-nucleating sites between the supernatant and the
13 resuspended clay pellet could be used as an indicator for the relative affinity of the free protein to the clay
14 surfaces. Uncertainties in the calculation of the number of nucleation sites are estimated by propagating
15 those stemming from the mass of mycelium in the initial suspensions, dilution errors, droplet volume
16 errors together and uncertainties which arise due to background freezing of the Milli-Q® water.
17 Background nucleation, stemming from impurities in the Milli-Q® water and the silanized-glass substrate
18 were subtracted out in the calculation of K (further details of this procedure can be found elsewhere
19 (O'Sullivan et al., 2015)), although in practice these nucleation sites were many orders of magnitude less
20 common than those of the P-INP in suspension.

21

22 **3. Results and Discussion**

23 **3.1. Clay-protein interactions in the absence of electrolytes**

24 Initially, we investigated adsorption of P-INPs to the clay in the absence of added salts and the results are
25 shown in Fig. 2. Values of K (no. of ice-nucleating sites per ml of suspension, eq. 1) are presented for

1 suspensions where the initial *pH* was not adjusted (5.7) as well as where it was adjusted to 7.4 and 4.7 by
2 addition of base or acid; this range of *pH* is relevant for typical topsoils. Measured *K* values for kaolinite on
3 its own (with no adjustment to *pH*) are far below those of either the pellet or supernatant (Fig. S1). The
4 open symbols in Fig. 2 are for the freezing of the supernatant after centrifuging the suspension, whereas
5 the closed symbols are for the resuspended clay pellets. It is clear that the majority of the P-INPs (around
6 90 % of those active at -7°C) remained in the supernatant (Fig. 2) and did not strongly adsorb to the clay
7 particles. This result was independent of *pH* within the range investigated. Hence, in the absence of added
8 salts, adsorption to the clay was minimal. As adsorption is a time-dependent process, it is possible that
9 extending the contact time could result in uptake of the protein to the clay. Nevertheless, the results here
10 show that on timescales characteristic of protein adsorption to clays (Yu et al., 2013), uptake of the P-INP
11 was minor in the absence of electrolytes.

12
13 Given the tendency of proteins to adsorb readily to clays (Zimmerman and Ahn, 2011; Yu et al., 2013), the
14 absence of protein uptake to the clay observed here suggests that, in this *pH* range, repulsive interactions
15 may inhibit adsorption. In kaolinite heterogeneous electrostatic charges exist at various points on the
16 surface (Tombácz and Szekeres, 2006). Permanent negative charges develop in kaolinite due to isomorphic
17 substitutions, along with *pH*-dependent charges due to surface silanol and aluminol groups. In the range
18 of *pH* studied, the silanol groups will likely be deprotonated, while aluminol groups become protonated
19 below about *pH* 6.5 (Tombácz and Szekeres, 2006). The presence of salt ions which are present in soils can
20 significantly alter the nature of the electrostatic interactions, and promote cation exchange at the surface
21 of the clays. Hence, the effects of adding salt ions such as Na⁺ and Ca²⁺ on the adsorption behavior was
22 further probed.

23

24 **3.2 Clay-protein interactions in the presence of electrolytes**

25 Solution ionic strength is known to be important in regulating the adsorption of proteins to solid surfaces
26 (Yu et al., 2013; Rabe et al., 2011). To examine the impacts of ionic strength and the identity of salts on the

1 uptake of the protein to the clay, salts were added to the suspension and the results are shown in Fig. 3.
2 Ionic strengths of 1 and 10 mM were chosen to approximate those found in soil solutions (Campbell et al.,
3 1989;Griffin and Jurinak, 1973;Edmeades et al., 1985;Dolling and Ritchie, 1985). Control experiments in
4 the absence of added kaolinite suggested no preferential settling of the protein in the electrolyte solution
5 under centrifugation. After a contact time of 2 hours, only a minor enhancement in the extent of
6 partitioning of P-INPs from the solution to the clay was observed when 1 mM NaCl was added (Fig. 3(a),
7 blue circles). However, for 1 mM CaCl₂ a large increase in the uptake of the protein to the clay was found
8 (Fig. 3(a), red circles). When the contact time was increased to 2 days (Fig. 3(b)), a dramatic increase in the
9 extent of adsorption to the clay was observed, with around 90% of INPs active at -7°C now being
10 associated with the kaolinite for both 1 mM CaCl₂ and NaCl solutions. These observations indicate that
11 electrolytes can enhance the binding of ice active fungal proteins to kaolinite and that the identity of the
12 cations in solution influences the adsorption process.

13
14 On increasing the ionic strength of the solution to 10 mM with NaCl (for a 2 hour contact time), the amount
15 of protein depleted from the supernatants was found to increase relative to when 1 mM was used such
16 that the activity of the clay matched that of the supernatant (Fig 4 (a) blue circles). Once again, in the
17 presence of divalent ions, such as Ca²⁺ or Mg²⁺, an even larger depletion of the free protein in solution was
18 found, with the amount of P-INP in the pellet now exceeding that in the supernatant (Fig. 4(a)). Similar to
19 the case for an ionic strength of 1 mM, allowing an extended contact time of 2 days was found to lead to
20 a further depletion of the free protein in the suspension. As can be seen in Fig. 4 (b), after 2 days the
21 majority of the protein is found to be associated with the clay mineral, regardless of the salt used. The
22 relatively rapid uptake of the ice-nucleating protein to the clay mineral kaolinite observed here is also
23 consistent with previous studies of the adsorption of a variety of proteins, where adsorption has been
24 found to reach equilibrium on timescales of minutes to hours (Ding and Henrichs, 2002;Lepoitevin et al.,
25 2014;Ralla et al., 2010;Venkateswerlu and Stotzky, 1992). Such time scales are short on environmental
26 time scales where cell-free proteins can have extended contact times with wet soil minerals.

1
2 The effect of adding salts to the suspensions here likely involves altering the extent of electrostatic
3 interactions between proteins in the suspension and the kaolinite adsorbent, as electrolytes act to screen
4 Coulombic interactions (Roth and Lenhoff, 1995;Rabe et al., 2011). Variations in ionic strength impact upon
5 both the kinetics and equilibrium position of protein adsorption on solid surfaces (Rabe et al., 2011). With
6 increasing ionic strength, adsorption of proteins (or protein domains) to oppositely charged substrates is
7 diminished, while adsorption between proteins and substrates with the same charge is enhanced (Rabe et
8 al., 2011). Furthermore, the presence of electrolytes can also enhance adsorption by reducing unfavorable
9 lateral interactions between adsorbed proteins (Rabe et al., 2011). Together with the observed lack of
10 adsorption across the *pH* range probed (Figure 2) the impacts of ionic strength in the current work suggest
11 that in the presence of added salts, repulsive electrostatic interactions are diminished allowing shorter
12 range attractive forces to dominate.

13
14 The increase in adsorption when a divalent cation is used rather than a monovalent one suggests that the
15 character of the ion may also impact upon the adsorption process through mechanisms other than just
16 the screening of negative charges. Similar results have previously been reported by Van Dulm et al. (1981)
17 for the adsorption of proteins to negatively charged polystyrene surfaces. Effects such as relative abilities
18 of different ions to act as bridges between proteins and surfaces of the same charge were suggested by
19 the authors as possible reasons for preferential protein uptake in the presence of divalent salts.

20
21 In a wide variety of past studies of protein adsorption on clays, adsorption has been found to be
22 accompanied by concurrent changes in activities of the proteins (Zimmerman and Ahn, 2011). While no
23 effects, or even an enhancement in activity has been observed for certain enzymes, reduction in activity
24 is a common feature of protein adsorption to mineral dusts (Zimmerman and Ahn, 2011;Calamai et al.,
25 2000;Gianfreda et al., 1991;Hughes and Simpson, 1978). Such impairment of activities has been suggested
26 to arise from conformational changes of the protein or the blocking of active sites upon adsorption (Rao

1 et al., 2000;Leprince and Quiquampoix, 1996;Quiquampoix, 1987;Lozzi et al., 2001). In the current study,
2 the continued activity of the adsorbed protein suggests that rapid inactivation in soils due to
3 conformational changes upon adsorption does not take place and activity is likely to be transferred to
4 surrounding soil particles upon release of *Fusarium* P-INPs.

5
6

7 **3.3. Reversibility of Adsorption**

8 To examine the extent to which adsorption was reversible, the clay pellet was repeatedly washed,
9 centrifuged and resuspended in Milli-Q® purified water, with the results indicated in Figure 5. The first
10 thing to note is that the clay pellet retains a substantial amount of fungal P-INPs even after five cycles of
11 washing and centrifuging. But, with each wash the activity of the clay was observed to reduce with around
12 10% of the protein lost to the supernatant on each cycle. The implication of this finding is that in soils the
13 adsorption is at least partially reversible, which may have important consequences for the biodegradability
14 of *Fusarium* P-INPs by soil microbes. The fact that a portion of the protein could be removed by washing
15 further suggests that while the majority of the fungal P-INPs are likely to be bound to clay minerals in soils,
16 they are not so tightly bound so as to prevent diffusion of the protein in soils over time. Accordingly, fungal
17 P-INPs may leach into surface and ground waters, leading to the migration of fungal P-INPs to other soil
18 surfaces spatially removed from the original source of the ice-nucleating fungus (Helassa et al., 2009).

19

20 **4. Conclusions**

21 Although certain biological particles harbor the most efficient naturally-occurring ice nucleation sites
22 known, the roles biological particles play in cloud ice formation remains poorly constrained. Currently,
23 there is a growing awareness that subcellular biological fragments from organisms, such as bacteria, pollen
24 and fungi may be important atmospheric INPs (O'Sullivan et al., 2015;Pummer et al., 2012;Augustin et al.,
25 2013;Fröhlich-Nowoisky et al., 2015;Pummer et al., 2015;Šantl-Temkiv et al., 2015;Rangel - Alvarado et
26 al., 2015). However, our knowledge of the environmental processing of these biomolecular INPs, such as

1 occurs in soils, currently limits our abilities to assess their contributions to airborne particles possessing
2 efficient ice-nucleating sites. Whether these nanoscale fragments may add to the reservoir of INPs found
3 in topsoils depends critically on whether activity is preserved upon adsorption to soil minerals.

4
5 In this study we have demonstrated that proteinaceous INPs from a common soil-borne fungus are
6 preferentially partitioned to clay surfaces in the presence of water containing electrolytes commonly
7 found in soil solutions. Upon adsorption, the exceptional ice-nucleating activity of the fungal P-INPs was
8 maintained, suggesting the adsorption process does not inactivate the efficient nucleating abilities of the
9 protein. Accordingly, fungal P-INPs are likely to confer activity to soil particles once removed from mycelia
10 by wetting, although further environmental processing, such as repeated wetting and drying of soils, could
11 potentially perturb protein activity. The fact that a minor amount of the protein could be removed from
12 the clay suggests that adsorption to the clay was reversible, which is an important observation for the
13 environmental fates of fungal P-INPs. With adsorption being reversible, the protein may be leachable by
14 surface waters and can potentially migrate to other soil surfaces distant from the original source fungi
15 (Helassa et al., 2009). Furthermore, the fact that a minor portion of the protein could be desorbed upon
16 washing suggests that the protein may be accessible as a nutrient source for soil microbes, making it more
17 susceptible to biodegradation (Fiorito et al., 2008; Gianfreda et al., 1991; Dashman and Stotzky, 1986).
18 Consequently, future work should be aimed at addressing the issue of ice active protein lifetime with
19 respect to biodegradation in soils.

20
21 The persistence of P-INP activity following adsorption to soil minerals has important implications for our
22 understandings of the roles which biological entities may play as sources of atmospheric INPs. As recently
23 suggested, biogenic matter residues within soil dusts are potentially the largest contributor to the burden
24 of atmospheric INPs at temperatures above -15 °C on a global basis (O'Sullivan et al., 2014; Conen et al.,
25 2011). Indeed, the characteristics of the organo-clay INPs examined in this study are similar to the efficient
26 INPs which have been identified in topsoils (O'Sullivan et al., 2015; Conen et al., 2011), being proteinaceous

1 ~~a substantial proportion of which were shown to be thermally labile in nature~~ and exhibit exceptional ice-
2 nucleating abilities. ~~In addition to *Fusaria*, other organisms can also contribute to the reservoir of cell free~~
3 ~~P-INPs in soils, such as the fungus *M. alpina* (Fröhlich-Nowoisky et al., 2015), certain lichens (Kieft and~~
4 ~~Ruscetti, 1990), and even ice-nucleating bacteria (Phelps et al., 1986). For *Fusaria*, ice-nucleating activity~~
5 ~~has been observed in a number of species, and species such as *Fusaria* are widespread in soils, occurring~~
6 ~~across the globe wherever crops are grown (Pitt et al., 2009).~~ Ice nucleating activity has been observed
7 ~~in a variety of *Fusarium* species, and *Fusaria* are one of the most common soil fungi, occurring across the~~
8 ~~globe wherever crops are grown (Pitt et al., 2009).~~ This, together with the fact that the contribution of
9 dusts from agricultural regions to the total dust emissions is estimated to be in the region of 25% (Ginoux
10 et al., 2012), means mineral-bound proteins from ~~fungi-*Fusaria* and other fungi~~ may be a major source of
11 the rare particles capable of nucleating cloud ice at temperatures above -15 °C. However, not every strain
12 of the five known ice nucleation active *Fusarium* species produce efficient INPs (Pouleur et al.,
13 1992;Richard et al., 1996). While *Fusaria* such as *F. avenaceum* are globally distributed, the relative
14 abundances of each species is governed by factors such as climate and land use (West et al., 2012). The
15 impacts of these factors on the distributions of P-INP producing *Fusaria* has yet to be fully discerned. This,
16 together with studies of the biodegradability of fungal P-INPs will be required to further constrain the net
17 contributions these particles make to the source strength of biological INPs lofted to the atmosphere.

18
19

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21

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25

26 **Author Contributions**

1 D.O.S. designed and conducted the ice nucleation experiments, analyzed the data and authored the
2 manuscript. B.J.M. and M.E.W. oversaw the project and helped to write the manuscript. J.F.R. assisted
3 with the culturing of the fungi and with the data analysis.

4

5 The authors declare no competing financial interests.

6

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1 Figures and Figure Legends

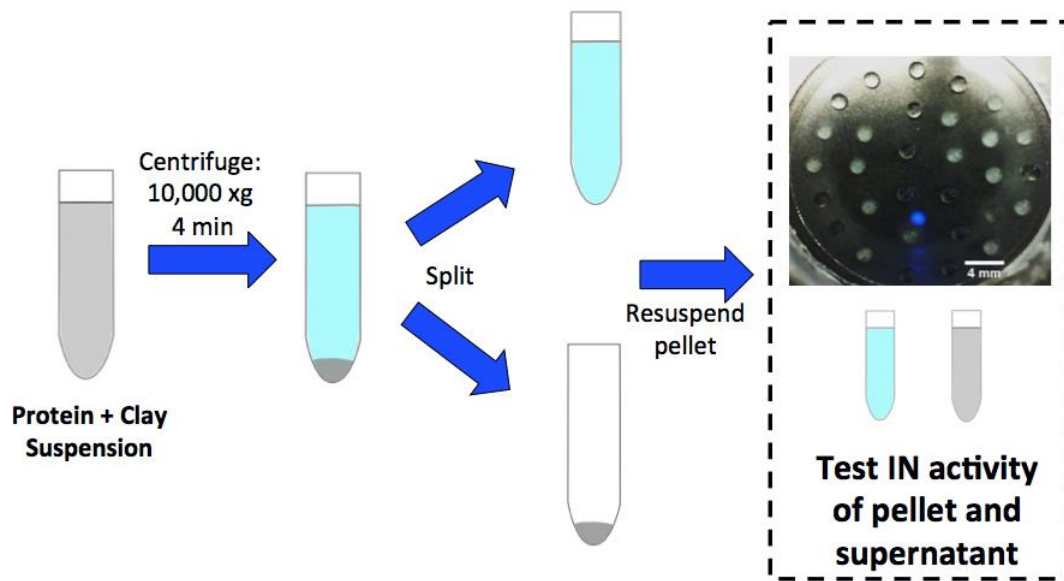


Figure 1: Schematic of the depletion method used to evaluate interactions between proteinaceous ice-nucleating particles (P-INPs) and the clay mineral kaolinite. Initially, suspensions containing the clay and protein were allowed to mix for a prescribed contact time. Following this, the clay was pelleted by centrifugation, the supernatant carefully removed and the remaining pellet resuspended in an equivalent volume of water (1 ml). The ice-nucleating activity of both the resuspended clay, and the supernatant is then tested to examine for the presence of P-INPs.

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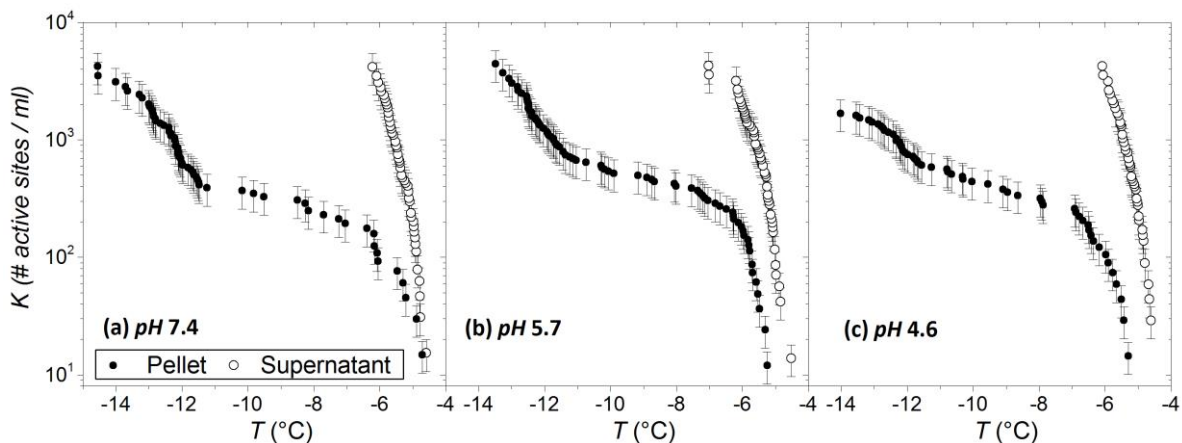


Figure 2: Effect of clay surfaces on fungal P-INPs in solution in the absence of added salts. The ice nucleating activities of the supernatant and the resuspended clay removed by centrifugation of a suspension of fungal P-INPs and kaolinite are illustrated. The contact time was 2 hours. Regardless of the initial *pH* used, ice-nucleating activity is predominantly found in the supernatant, suggesting that adsorption only occurs to a minor extent, if at all.

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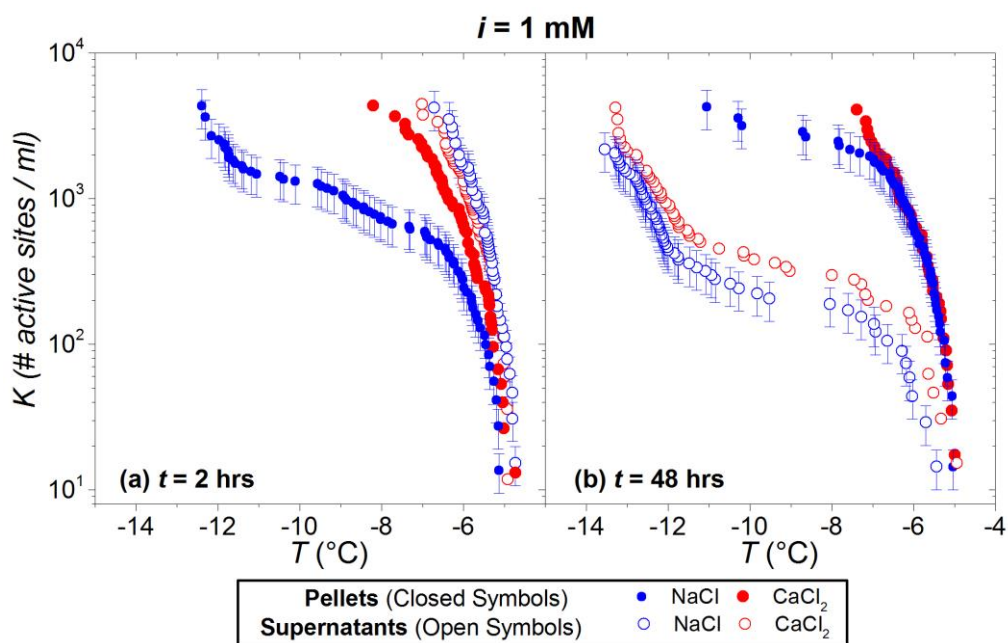


Figure 3: Effect of electrolytes at an ionic strength of 1 mM on the uptake of P-INPs to the clay surface with a contact time of (a) 2 hours and (b) 2 days. Initially, in (a) it can be seen that uptake to the clay is greater in the presence of Ca^{2+} (closed red circles) compared with Na^+ (closed blue circles), but after 2 days (b), the extent of adsorption is the similar regardless of the salt used. After 2 days, depletion of the free protein is extensive, with on the order of 90 % of INPs active at -7°C found to be associated with the clay.

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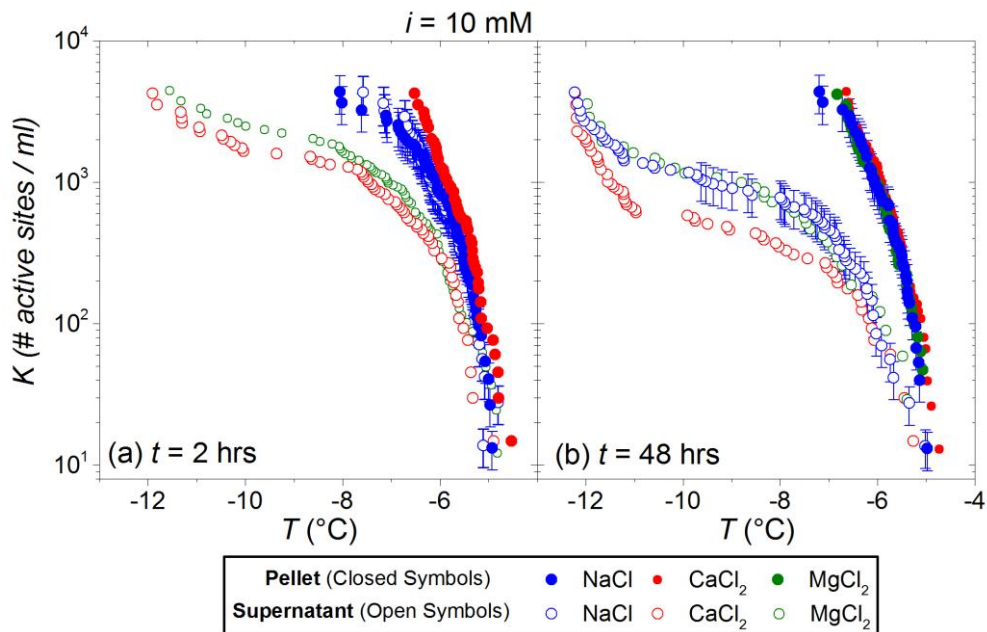


Figure 4: Effect of increasing ionic strength to 10 mM on the uptake of P-INPs to the clay surface. Illustrated are the activities of the clay (closed symbols) and the supernatant (open symbols) after contact times of (a) 2 hours and (b) 2 days. After 2 hours, the extent of adsorption is observed to be greater than the case of when an ionic strength of 1 mM is used (Figure 3(a)). On increasing the contact time adsorption continues to proceed, where it can be seen that after 2 days, greater than 90% of the protein is adsorbed to the clay pellet (panel (b)).

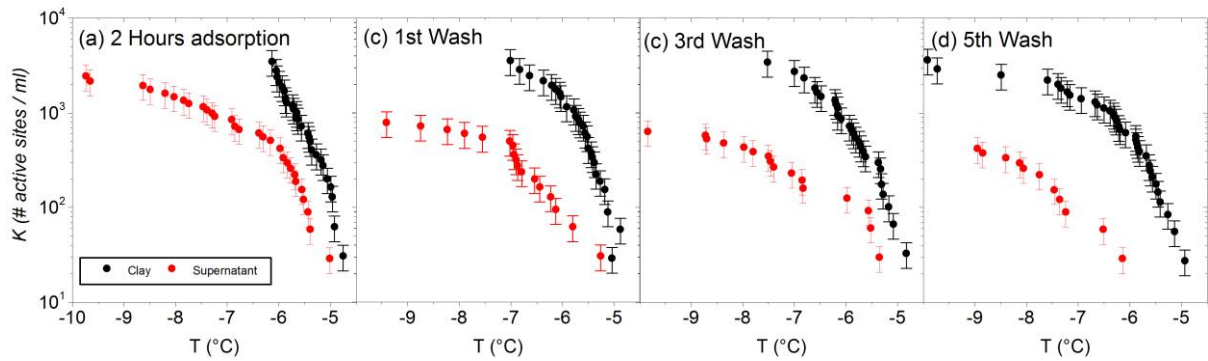


Figure 5: Effect of repeatedly washing the clay pellet by successive resuspension and centrifugation (see Figure 1). Illustrated are (a) activities of the clay and supernatant after adsorption is allowed to proceed for two hours, and (b) – (d) the effects of successive washing of the clay pellet with pure water. After each washing cycle, on the order of 10 % of the protein was observed to be lost to the supernatant, suggesting that the adsorption process is partially reversible. This is in addition to a maximum of 4 % loss in each wash, due to the amount removed to conduct the droplet freezing experiments.

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