Ubiquity of Biological Ice Nucleators in Snowfall

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Aerosols play an integral role in climate by directly scattering or absorbing solar radiation, and indirectly these particles serve as condensation and ice nuclei in clouds. Ice formation in tropospheric clouds is required for snow and most rainfall. At temperatures > −40°C, ice formation is not spontaneous, and diverse substrates can act as catalysts of ice nucleation (1). Biological ice nucleators (IN) are the most active IN in nature, and some bacterial plant pathogens can catalyze ice formation at temperatures near −2°C. Although biological particles are important components of aerosols (2), existing technologies are unable to detect IN active at temperatures > −10°C (3), and the impact of biological IN on atmospheric processes remains ambiguous (4).

We examined the concentration and nature of IN active at temperatures > −10°C in 19 fresh snowfalls collected at mid- and high-latitude locations from October 2005 to June 2006 (5) and found the lowest cumulative concentration of IN at −9°C in Antarctic snowfall (four IN per liter of snow melt) and highest in samples from Montana and France (110 and 120 IN per liter, respectively). DNA-containing cells between 0.3 and 15 μm in diameter (1.5 × 10^4 to 5.4 × 10^6 cells per liter) accounted for <1% of the total particles in the snow samples (1.1 × 10^3 to 3.9 × 10^3 particles per liter), and the concentration of biological IN was independent of the particle and cell concentration (Fig. 1). The data presented in Fig. 1B are for the warmest temperature for which concentrations of IN could be calculated for all samples (−7°C), thereby facilitating comparison.

The activity of most known biological IN is mediated by proteins or proteinaceous compounds (4). Therefore, we reasoned that IN of biological origin would be inactivated by heat, whereas IN of mineral origin would not. Heat treatment inactivated 69% to 100% of the IN active at temperatures ≥ −7°C and ≤ −4°C (Fig. 1B). In bacteria, ice-nucleation activity depends on an intact cell wall and is conferred by a 120- to 180-kD protein embedded in the outer membrane (6). Disruption of the cell wall reduces ice-nucleation activity in bacteria (7). Hence, we postulated that lysozyme would decrease the ice-nucleation activity if bacteria were involved and found that 0 to 85% of the IN were susceptible to lysozyme (Fig. 1B). Incomplete hydrolysis and resistance to lysozyme could allow some bacterial cells to remain intact in the presence of lysozyme, and, therefore, our results are probably underestimating the concentration of bacterial IN in snowfall.

The samples analyzed were collected during seasons and in locations (e.g., Antarctica) devoid of deciduous plants, making it likely that the biological IN we observed were transported from long distances and maintained their ice-nucleating activity in the atmosphere. Assuming that all biological IN in the snow samples were DNA-containing cells, we estimate that about 0.4% of the cells in mid-latitude snowfalls were ice-nucleating active at temperatures between −7°C and −4°C. Although extrapolating the absolute composition and quantity of biological IN in clouds on the basis of concentrations in precipitation is difficult, our results indicate that these particles are widely dispersed in the atmosphere, and, if present in clouds, they may have an important role in the initiation of ice formation, especially when minimum cloud temperatures are relatively warm.

Biological IN were first described nearly 40 years ago (1); however, there are no estimates for the biological component of IN in the atmosphere. Our results reveal that biological IN active at warm temperatures are abundant in fresh snow samples and are ubiquitous in precipitation from worldwide locations. Unearthing a role for biological IN in the precipitation cycle has implications for deciphering feedbacks between the biosphere and climate, improving climate forecast models, and understanding atmospheric dissemination strategies of plant pathogens (4) and other microorganisms.

Fig. 1. Particle, cell, and freeze immersion IN concentrations in fresh snowfall collected from Montana, France, the Yukon, and Antarctica. (A) Quantity of particles and DNA-containing cells between 0.3 and 15 μm in diameter. (B) The cumulative concentration of IN (≥0.2 μm) per liter of snow melt that were active at −7°C and sensitive to lysozyme and heat denaturation. From left to right, samples were ranked on the basis of increasing cumulative concentrations of IN. Samples 2, 5 to 8, 10 to 12, 17, and 18 were collected in the vicinity of Bozeman, Montana; samples 3, 9, 13 to 16, and 19 from the French Alps and Pyrenees in southern France; and samples 1 and 4 from Ross Island, Antarctica and the Wheaton Glacier, Yukon, respectively. See table S1 for numerical data and detail on sampling dates and site locations.

References and Notes
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Table S1
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